ANTIGEN BINDING PROTEINS TO PROPROTEIN CONVERTASE SUBTILISIN KEXIN TYPE 9 (PCSK9)

RELATED APPLICATIONS

[0001] This application claims priority to U.S. Provisional Applications Ser. No. 60/957,668, filed August 23, 2007, Ser No. 61/008,965, filed December 21, 2007, and Ser. No. 61/010,630, filed January 9, 2008, hereby incorporated by reference in their entireties.

FIELD OF THE INVENTION

[0002] The present invention relates to antigen binding proteins that bind to proprotein convertase subtilisin kexin type 9 (PCSK9) and methods of using and making the antigen binding proteins.

BACKGROUND OF VARIOUS EMBODIMENTS

[0003] Proprotein convertase subtilisin kexin type 9 (PCSK9) is a serine protease involved in regulating the levels of the low density lipoprotein receptor (LDLR) protein (Horton *et al.*, 2007; Seidah and Prat, 2007). *In vitro* experiments have shown that adding PCSK9 to HepG2 cells lowers the levels of cell surface LDLR (Benjannet *et al.*, 2004; Lagace *et al.*, 2006; Maxwell *et al.*, 2005; Park *et al.*, 2004). Experiments with mice have shown that increasing PCSK9 protein levels decreases levels of LDLR protein in the liver (Benjannet *et al.*, 2004; Lagace *et al.*, 2006; Maxwell *et al.*, 2005; Park *et al.*, 2004), while PCSK9 knockout mice have increased levels of LDLR in the liver (Rashid *et al.*, 2005). Additionally, various human PCSK9 mutations that result in either increased or decreased levels of plasma LDL have been identified (Kotowski *et al.*, 2006; Zhao *et al.*, 2006). PCSK9 has been shown to directly interact with the LDLR protein, be endocytosed along with the LDLR, and co-immunofluoresce with the LDLR throughout the endosomal pathway (Lagace *et al.*, 2006). Degradation of the LDLR by PCSK9 has not been observed and the mechanism through which it lowers extracellular LDLR protein levels is uncertain.

[0004] PCSK9 is a prohormone-proprotein convertase in the subtilisin (S8) family of serine proteases (Seidah *et al.*, 2003). Humans have nine prohormone-proprotein convertases

that can be divided between the S8A and S8B subfamilies (Rawlings *et al.*, 2006). Furin, PC1/PC3, PC2, PACE4, PC4, PC5/PC6 and PC7/PC8/LPC/SPC7 are classified in subfamily S8B. Crystal and NMR structures of different domains from mouse furin and PC1 reveal subtilisin-like pro- and catalytic domains, and a P domain directly C-terminal to the catalytic domain (Henrich *et al.*, 2003; Tangrea *et al.*, 2002). Based on the amino acid sequence similarity within this subfamily, all seven members are predicted to have similar structures (Henrich *et al.*, 2005). SKI-1/S1P and PCSK9 are classified in subfamily S8A. Sequence comparisons with these proteins also suggest the presence of subtilisin-like pro- and catalytic domains (Sakai *et al.*, 1998; Seidah *et al.*, 2003; Seidah *et al.*, 1999). In these proteins the amino acid sequence C-terminal to the catalytic domain is more variable and does not suggest the presence of a P domain.

[0005] Prohormone-proprotein convertases are expressed as zymogens and they mature through a multi step process. The function of the pro-domain in this process is two-fold. The pro-domain first acts as a chaperone and is required for proper folding of the catalytic domain (Ikemura *et al.*, 1987). Once the catalytic domain is folded, autocatalysis occurs between the pro-domain and catalytic domain. Following this initial cleavage reaction, the pro-domain remains bound to the catalytic domain where it then acts as an inhibitor of catalytic activity (Fu *et al.*, 2000). When conditions are correct, maturation proceeds with a second autocatalytic event at a site within the pro-domain (Anderson *et al.*, 1997). After this second cleavage event occurs the pro-domain and catalytic domain dissociate, giving rise to an active protease.

[0006] Autocatalysis of the PCSK9 zymogen occurs between Gln152 and Ser153 (VFAQ|SIP) (Naureckiene *et al.*, 2003), and has been shown to be required for its secretion from cells (Seidah *et al.*, 2003). A second autocatalytic event at a site within PCSK9's pro-domain has not been observed. Purified PCSK9 is made up of two species that can be separated by non-reducing SDS-PAGE; the pro-domain at 17 Kd, and the catalytic plus C-terminal domains at 65 Kd. PCSK9 has not been isolated without its inhibitory pro-domain, and measurements of PCSK9's catalytic activity have been variable (Naureckiene *et al.*, 2003; Seidah *et al.*, 2003).

SUMMARY OF VARIOUS EMBODIMENTS

[0007] In some embodiments, the invention comprises an antigen binding protein to PCSK9.

[0008] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9 comprising: A) one or more heavy chain complementary determining regions (CDRHs) selected from the group consisting of: (i) a CDRH1 from a CDRH1 in a sequence selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; (ii) a CDRH2 from a CDRH2 in a sequence selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; (iii) a CDRH3 from a CDRH3 in a sequence selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; and (iv) a CDRH of (i), (ii), and (iii) that contains one or more amino acid substitutions, deletions or insertions of no more than 4 amino acids; B) one or more light chain complementary determining regions (CDRLs) selected from the group consisting of: (i) a CDRL1 from a CDRL1 in a sequence selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; (ii) a CDRL2 from a CDRL2 in a sequence selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; (iii) a CDRL3 from a CDRL3 in a sequence selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; and (iv) a CDRL of (i), (ii) and (iii) that contains one or more amino acid substitutions, deletions or insertions of no more than 4 amino acids; or C) one or more heavy chain CDRHs of A) and one or more light chain CDRLs of B). In some embodiments, the isolated antigen binding protein comprises at least one CDRH of A) and at least one CDRL of B). In some embodiments, the isolated antigen binding protein comprises at least two CDRH of A) and at least two CDRL of B). In some embodiments, the isolated antigen binding protein comprises said CDRH1, CDRH2, CDRH3, CDRL1, CDRL2 and CDRL3. In some embodiments, the CDRH of A) is selected from at least one of the group consisting of: (i) a CDRH1 amino acid sequence selected from the CDRH1 in a sequence selected from the group consisting of SEQ ID NO: 67, 79, 89, and 49; (ii) a CDRH2 amino acid

sequence selected from the CDRH2 in a sequence selected from the group consisting of SEQ ID NO: 67, 79, 89, and 49; (iii) a CDRH3 amino acid sequence selected from the CDRH3 in a sequence selected from the group consisting of SEQ ID NO: 67, 79, 89, and 49; and (iv) a CDRH of (i), (ii) and (iii) that contains one or more amino acid substitutions, deletions or insertions of no more than 2 amino acids. In addition, the CDRL of B) is selected from at least one of the group consisting of: (i) a CDRL1 amino acid sequence selected from the CDRL1 in a sequence selected from the group consisting of SEQ ID NO: 12, 35, 32, and 23; (ii) a CDRL2 amino acid sequence selected from the CDRL2 in a sequence selected from the group consisting of SEQ ID NO: 12, 35, 32, and 23; (iii) a CDRL3 amino acid sequence selected from the CDRL3 in a sequence selected from the group consisting of SEQ ID NO: 12, 35, 32, and 23; and (iv) a CDRL of (i), (ii) and (iii) that contains one or more amino acid substitutions, deletions or insertions of no more than 2 amino acids; or C) one or more heavy chain CDRHs of A) and one or more light chain CDRLs of B. In some embodiments, the CDRH of A) is selected from at least one of the group consisting of: (i) a CDRH1 amino acid sequence of the CDRH1 amino acid sequence in SEQ ID NO: 67; (ii) a CDRH2 amino acid sequence of the CDRH2 amino acid sequence in SEQ ID NO: 67; (iii) a CDRH3 amino acid sequence of the CDRH3 amino acid sequence in SEQ ID NO: 67; and (iv) a CDRH of (i), (ii) and (iii) that contains one or more amino acid substitutions, deletions or insertions of no more than 2 amino acids; said CDRL of B) is selected from at least one of the group consisting of: (i) a CDRL1 amino acid sequence of the CDRL1 amino acid sequence in SEQ ID NO: 12; (ii) a CDRL2 amino acid sequence of the CDRL2 amino acid sequence in SEQ ID NO: 12; (iii) a CDRL3 amino acid sequence of the CDRL3 amino acid sequence in SEQ ID NO: 12; and (iv) a CDRL of (i), (ii) and (iii) that contains one or more amino acid substitutions, deletions or insertions of no more than 2 amino acids; or C) one or more heavy chain CDRHs of A) and one or more light chain CDRLs of B). In some embodiments, the antigen binding protein comprises A) a CDRH1 of the CDRH1 sequence in SEQ ID NO: 67, a CDRH2 of the CDRH2 sequence in SEQ ID NO: 67, and a CDRH3 of the CDRH3 sequence in SEQ ID NO: 67, and B) a CDRL1 of the CDRL1 sequence in SEQ ID NO: 12, a CDRL2 of the CDRL2 sequence in SEQ ID NO: 12, and a CDRL3 of the CDRL3 sequence in SEQ ID NO: 12. In some embodiments, the antigen binding protein comprises a heavy chain variable region (VH) having at least 80% sequence identity with an amino acid sequence selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87,

58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60, and/or a light chain variable region (VL) having at least 80% sequence identity with an amino acid sequence selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46. In some embodiments, the VH has at least 90% sequence identity with an amino acid sequence selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60, and/or the VL has at least 90% sequence identity with an amino acid sequence selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46. In some embodiments, the VH is selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60, and/or the VL is selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46.

[0009] In some aspects, the invention comprises an isolated antigen binding protein that specifically binds to an epitope that is bound by any of the ABPs disclosed herein.

[0010] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9, wherein the antigen binding protein comprises: A) one or more heavy chain CDRs (CDRHs) selected from at least one of the group consisting of: (i) a CDRH1 with at least 80% sequence identity to a CDRH1 in one of the sequences selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; (ii) a CDRH2 with at least 80% sequence identity to a CDRH2 in one of the sequences selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; and (iii) a CDRH3 with at least 80% sequence identity to a CDRH3 in one of the sequences selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; B) one or more light chain CDRs (CDRLs) selected from at least one of the group consisting of: (i) a CDRL1 with at least 80% sequence identity to a CDRL1 in one of the sequences selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; (ii)a CDRL2 with at least 80% sequence

identity to a CDRL2 in one of the sequences selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; and (iii) a CDRL3 with at least 80% sequence identity to a CDRL3 in one of the sequences selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; or C) one or more heavy chain CDRHs of A) and one or more light chain CDRLs of B). In some embodiments, the antigen binding protein comprises: A) one or more CDRHs selected from at least one of the group consisting of: (i) a CDRH1 with at least 90% sequence identity to a CDRH1 in one of the sequences selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; (ii) a CDRH2 with at least 90% sequence identity to a CDRH2 in one of the sequences selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; and (iii) a CDRH3 with at least 90% sequence identity to a CDRH3 in one of the sequences selected from the group consisting of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60; B) one or more CDRLs selected from at least one of the group consisting of: (i) a CDRL1 with at least 90% sequence identity to a CDRL1 in one of the sequences selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; (ii) a CDRL2 with at least 90% sequence identity to a CDRL2 in one of the sequences selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; and (iii) a CDRL3 with at least 90% sequence identity to a CDRL3 in one of the sequences selected from the group consisting of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46; or C) one or more heavy chain CDRHs of A) and one or more light chain CDRLs of B).

[0011] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9, the antigen binding protein comprises: A) a heavy chain complementary determining region (CDRH) selected from at least one of the group consisting of: (i) a CDRH3 selected from the CDRH3 within the sequences selected from the group consisting of SEQ ID NOs: 67, 79, and 49, (ii) a CDRH3 that differs in amino acid sequence from the CDRH3 of (i) by

an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}X_{12}X_{13}X_{14}$ (SEQ ID NO: 404), wherein X_1 is selected from the group consisting of D, A, R, and not amino acid, X_2 is selected from the group consisting of Y, I, G, and no amino acid, X_3 is selected from the group consisting of D, A, G, and no amino acid, X_4 is selected from the group consisting of F, A, L, and no amino acid, X_5 is selected from the group consisting of W, L, A, and no amino acid, X_6 is selected from the group consisting of S, Y, A, and no amino acid, X_7 is selected from the group consisting of A, Y, R, and no amino acid, X_8 is selected from the group consisting of Y, P, and no amino acid, X_9 is selected from the group consisting of Y, G, and no amino acid, X_{10} is selected from the group consisting of D, G, and no amino acid, X_{11} is selected from the group consisting of A, M, and no amino acid, X_{12} is selected from the group consisting of F, D, and no amino acid, X_{13} is selected from the group consisting of D, V, and no amino acid, X_{14} is selected from the group consisting of V and no amino acid; B) a light chain complementary determining region (CDRL) selected from at least one of the group consisting of: (i) a CDRL3 selected from the CDRL3 within the sequences selected from the group consisting of SEQ ID NOs: 12, 35, and 23, (ii) a CDRL3 that differs in amino acid sequence from the CDRL3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRL3 amino acid sequence selected from the group consisting of: $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}$ (SEQ ID NO: 405), wherein X_1 is selected from the group consisting of Q and G, X_2 is selected from the group consisting of S, T, A, and no amino acid, X_3 is selected from the group consisting of Y, no amino acid, and W, X_4 is selected from the group consisting of D and no amino acid, X_5 is selected from the group consisting of S and no amino acid, X_6 is selected from the group consisting of S and no amino acid, X_7 is selected from the group consisting of L, T, and no amino acid, X_8 is selected from the group consisting of no amino acid, A, and S, X_9 is selected from the group consisting of no amino acid, G, A, and V, X_{10} is selected from the group consisting of no amino acid, S, Y, and V, X_{11} is selected from the group consisting of no amino acid and V.

[0012] In some aspects, the invention comprises an isolated antigen binding protein comprising a light chain having the amino acid sequence selected from the group consisting of: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, 46, and some combination thereof.

[0013] In some embodiments, the antigen binding protein specifically binds to an epitope that is bound by at least one of the antigen binding proteins disclosed herein. In some embodiments, the isolated antigen binding protein further comprises a heavy chain having the amino acid sequence selected from the group consisting of: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, 60, and some combination thereof. In some embodiments, the amino acid sequence of the ABP is selected from the group consisting of SEQ ID NO: 12, 35, 23, and some combination thereof. In some embodiments, the heavy chain of the ABP comprises a CDRH3 of SEQ ID NO: 67, a CDRH2 of SEQ ID NO: 67, and a CDRH1 of SEQ ID NO:67, and said light chain comprises a CDRL3 of SEQ ID NO: 12, a CDRL2 of SEQ ID NO: 12, and a CDRL1 of SEQ ID NO: 12. In some embodiments, the isolated antigen binding protein is a monoclonal antibody, a polyclonal antibody, a recombinant antibody, a human antibody, a humanized antibody, a chimeric antibody, a multispecific antibody, or an antibody fragment thereof. In some embodiments, the isolated antigen binding protein is a Fab fragment, a Fab' fragment, a $F(ab')_2$ fragment, a Fv fragment, a diabody, or a single chain antibody molecule. In some embodiments, the isolated antigen binding protein is a human antibody. In some embodiments, the isolated antigen binding protein is a monoclonal antibody. In some embodiments, the isolated antigen binding protein is of the IgG1-, IgG2- IgG3- or IgG4-type. In some embodiments, the isolated antigen binding protein is of the IgG4- or IgG2-type. In some embodiments, the isolated antigen binding protein is coupled to a labeling group. In some embodiments, the isolated antigen binding protein competes for binding to PCSK9 with an antigen binding protein described herein. In some embodiments, the isolated antigen binding protein is a monoclonal antibody, a polyclonal antibody, a recombinant antibody, a human antibody, a humanized antibody, a chimeric antibody, a multispecific antibody, or an antibody fragment thereof. In some embodiments, the isolated antigen binding protein is a Fab fragment, a Fab' fragment, a $F(ab')_2$ fragment, a Fv fragment, a diabody, or a single chain antibody molecule. In some embodiments, the isolated antigen binding protein is coupled to a labeling group. In some embodiments, the isolated antigen binding protein reduces binding of PCSK9 to LDLR. In some embodiments, the isolated antigen binding protein the antigen binding protein decreases an amount of LDL present in a subject when administered to the subject. In some embodiments, the isolated antigen binding protein decreases an amount of serum cholesterol present in a subject when administered to the subject. In some embodiments, the isolated antigen

binding protein increases an amount of LDLR present in a subject when administered to the subject.

[0014] In some aspects, the invention comprises a vector comprising a nucleic acid molecule as described herein. In some embodiments, the invention comprises a host cell comprising a nucleic acid molecule as described herein.

[0015] In some aspects, the invention comprises an isolated antigen binding protein that competes for binding to PCSK9 with an antigen binding protein disclosed herein.

[0016] In some aspects, the invention comprises a nucleic acid molecule encoding the antigen binding protein according disclosed herein.

[0017] In some aspects, the invention comprises a pharmaceutical composition comprising at least one antigen binding protein described herein.

[0018] In some aspects, the invention comprises a method for treating or preventing a condition associated with elevated serum cholesterol levels in a patient, comprising administering to a patient in need thereof an effective amount of at least one isolated antigen binding protein disclosed herein.

[0019] In some aspects, the invention comprises a method of inhibiting binding of PCSK9 to LDLR in a subject comprising administering an effective amount of at least one antigen binding protein disclosed herein.

[0020] In some aspects, the invention comprises an antigen binding protein that selectively binds to PCSK9, wherein the antigen binding protein binds to PCSK9 with a K_d that is smaller than 100 pM.

[0021] In some aspects, the invention comprises a method for treating or preventing a condition associated with elevated serum cholesterol levels in a subject, the method comprising administering to a subject in need thereof an effective amount of at least one isolated antigen binding protein disclosed herein simultaneously or sequentially with an agent that elevates the availability of LDLR protein.

[0022] In some aspects, the invention comprises a method of lowering serum cholesterol level in a subject, the method comprising administering to a subject an effective amount of at least one isolated antigen binding protein as disclosed herein.

[0023] In some aspects, the invention comprises a method of lowering serum cholesterol level in a subject, the method comprising administering to a subject an effective

amount of at least one isolated antigen binding protein as disclosed herein, simultaneously or sequentially with an agent that elevates the availability of LDLR protein.

[0024] In some aspects, the invention comprises a method of increasing LDLR protein level in a subject, the method comprising administering to a subject an effective amount of at least one isolated antigen binding protein as disclosed herein.

[0025] In some aspects, the invention comprises a method of increasing LDLR protein levels in a subject, the method comprising administering to a subject an effective amount of at least one isolated antigen binding protein as disclosed herein simultaneously or sequentially with an agent that elevates the availability of LDLR protein.

[0026] In some aspects, the invention comprises a pharmaceutical composition comprising an ABP as disclosed herein and an agent that elevates the availability of LDLR protein levels. In some embodiments, the agent that elevates the availability of LDLR protein comprises a statin. In some embodiments, the statin is selected from the group consisting of atorvastatin, cerivastatin, fluvastatin, lovastatin, mevastatin, pitavastatin, pravastatin, rosuvastatin, simvastatin, and some combination thereof.

[0027] In some aspect, the invention comprises a method of making the antigen binding protein as described herein, comprising the step of preparing said antigen binding protein from a host cell that secretes said antigen binding protein.

[0028] In some aspect, the invention comprises a pharmaceutical composition comprising at least one antigen binding protein as described herein and a pharmaceutically acceptable excipient. In some embodiments, the pharmaceutical composition further comprises an additional active agent. In some embodiments, said additional active agent is selected from the group consisting of a radioisotope, radionuclide, a toxin, or a therapeutic and a chemotherapeutic group.

[0029] In some aspects, the invention comprises a method for treating or preventing a condition associated with an elevated serum cholesterol level in a patient. The method comprises administering to a patient in need thereof an effective amount of at least one isolated antigen binding protein as disclosed herein. In some embodiments, the condition is hypercholesterolemia.

[0030] In some aspects, the invention comprises a method of inhibiting binding of PCSK9 to LDLR in a patient comprising administering an effective amount of at least one antigen binding protein according as described herein.

[0031] In some aspect, the invention comprises an antigen binding protein that binds to PCSK9 with a K_d that is smaller than 100 pM. In some embodiments, the antigen binding protein binds with a K_d that is smaller than 10 pM. In some embodiments, the antigen binding protein binds with a K_d that is less than 5 pM.

[0032] In some aspects, the invention comprises a method for treating or preventing a condition associated with elevated serum cholesterol levels in a subject, said method comprising administering to a subject in need thereof an effective amount of at least one isolated antigen binding protein described herein simultaneously or sequentially with an agent that elevates the availability of LDLR protein. In some embodiments, the agent that elevates the availability of LDLR protein comprises a statin. In some embodiments, the statin is selected from the group consisting of atorvastatin, cerivastatin, fluvastatin, lovastatin, mevastatin, pitavastatin, pravastatin, rosuvastatin, simvastatin, and some combination thereof.

[0033] In some aspects, the invention comprises a method of lowering the serum cholesterol level in a subject. The method comprises administering to a subject an effective amount of at least one isolated antigen binding protein as described herein.

[0034] In some aspects, the invention comprises a method of lowering serum cholesterol levels in a subject comprising administering to a subject an effective amount of at least one isolated antigen binding protein, as described herein, simultaneously or sequentially with an agent that elevates the availability of LDLR protein. In some embodiments, the agent that elevates the availability of LDLR protein comprises a statin. In some embodiments, the statin is selected from the group consisting of atorvastatin, cerivastatin, fluvastatin, lovastatin, mevastatin, pitavastatin, pravastatin, rosuvastatin, simvastatin, and some combination thereof.

[0035] In some aspects, the invention comprises a method of increasing LDLR protein levels in a subject by administering to a subject an effective amount of at least one isolated antigen binding protein as provided herein.

[0036] In some aspects, the invention comprises a method of increasing LDLR protein levels in a subject by administering to a subject an effective amount of at least one isolated antigen binding protein, as described herein, simultaneously or sequentially with an agent that elevates the availability of LDLR protein. In some embodiments, the agent that elevates the availability of LDLR protein levels comprises a statin. In some embodiments, the

statin is selected from the group consisting of atorvastatin, cerivastatin, fluvastatin, lovastatin, mevastatin, pitavastatin, pravastatin, rosuvastatin, simvastatin, and some combination thereof.

[0037] In some aspects, the invention comprises a neutralizing antibody that binds to PCSK9 and reduces a low density lipoprotein receptor (LDLR) lowering effect of PCSK9 on LDLR. In some embodiments, the antibody specifically binds to PCSK9. In some embodiments, the antibody binds to the catalytic domain of PCSK9. In some embodiments, the antibody binds to an epitope within residues 31-447 of SEQ ID NO: 3. In some embodiments, the antibody binds to PCSK9 having an amino acid sequence that is at least 90% identical to SEQ ID NO: 3.

[0038] In some aspects, the invention comprises a neutralizing antigen binding protein that binds to PCSK9, wherein the antigen binding protein binds to PCSK9 at a location within residues 31-447 of SEQ ID NO: 3. In some embodiments, when the antigen binding protein is bound to PCSK9, the antibody is positioned 8 angstroms or less from at least one of the following residues of PCSK9: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, S381, W156, N157, L158, E159, H193, E195, H229, R237, G240, K243, D367, I368, G370, A371, S373, S376, Q382, W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, G384, K69, D70, P71, S148, V149, D186, T187, E211, D212, G213, R218, Q219, C223, D224, G227, H229, L253, N254, G259, P288, A290, G291, G316, R319, Y325, V346, G352, T353, G365, I368, I369, S372, S373, C378, F379, T385, S386, Q387, S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, F379, I154, T187, H193, E195, I196, M201, V202, C223, T228, S235, G236, A239, G244, M247, I369, S372, C375, or C378. In some embodiments, the antibody is positioned 8 angstroms or less from at least one of the following residues of PCSK9: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, S381, W156, N157, L158, E159, H193, E195, H229, R237, G240, K243, D367, I368, G370, A371, S373, S376, or Q382. In some embodiments, the antibody is positioned 5 angstroms or less from at least one of the following residues of PCSK9: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, or S381. In some embodiments, the antibody is positioned 5 angstroms or less from at least two of the following residues of PCSK9: S153, I154, P155, R194, D238, A239, I369, S372, D374,

C375, T377, C378, F379, V380, or S381. In some embodiments, the antibody is 5 angstroms or less from at least four of the following residues of PCSK9: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, or S381. In some embodiments, the antibody is positioned 8 angstroms or less from at least one of the following residues of PCSK9: W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, G384, K69, D70, P71, S148, V149, D186, T187, E211, D212, G213, R218, Q219, C223, D224, G227, H229, L253, N254, G259, P288, A290, G291, G316, R319, Y325, V346, G352, T353, G365, I368, I369, S372, S373, C378, F379, T385, S386, or Q387. In some embodiments, the antibody is positioned 5 angstroms or less from at least one of the following residues of PCSK9: W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, or G384. In some embodiments, the antibody is positioned 5 angstroms or less from at least two of the following residues of PCSK9: W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, or G384. In some embodiments, the antibody is positioned 5 angstroms or less from at least four of the following residues of PCSK9: W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, or G384. In some embodiments, the antibody is positioned 8 angstroms or less from at least one of the following residues of PCSK9: S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, F379, I154, T187, H193, E195, I196, M201, V202, C223, T228, S235, G236, A239, G244, M247, I369, S372, C375, or C378. In some embodiments, the antibody is positioned 5 angstroms or less from at least one of the following residues of PCSK9: S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, or F379. In some embodiments, the antibody is positioned 5 angstroms or less from at least two of the following residues of PCSK9: S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, or F379. In some embodiments, the antibody is positioned 5 angstroms or less from at least four of the following residues of PCSK9: S153, S188, I189,

Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, or F379 .

[0039] In some aspects, the invention comprises a neutralizing antibody that binds to PCSK9, wherein the antibody binds to PCSK9 and reduces the likelihood that PCSK9 binds to LDLR.

[0040] In some embodiments, an antibody or antigen binding molecule that binds to PCSK9 is contemplated. The antibody binds to PCSK9 at a location within residues 31-447 of SEQ ID NO: 3. In some embodiments, the antibody or antigen binding molecule, when bound to PCSK9, is positioned 8 angstroms or less from at least one of the following residues of PCSK9: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, S381, W156, N157, L158, E159, H193, E195, H229, R237, G240, K243, D367, I368, G370, A371, S373, S376, Q382, W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, G384, K69, D70, P71, S148, V149, D186, T187, E211, D212, G213, R218, Q219, C223, D224, G227, H229, L253, N254, G259, P288, A290, G291, G316, R319, Y325, V346, G352, T353, G365, I368, I369, S372, S373, C378, F379, T385, S386, Q387, S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, F379, I154, T187, H193, E195, I196, M201, V202, C223, T228, S235, G236, A239, G244, M247, I369, S372, C375, or C378.

[0041] In some embodiments, an isolated antibody or antigen binding molecule that blocks an antibody to PCSK9 from binding within 8 angstroms of a residue of PCSK9 is provided. In some embodiments the residue of PCSK9 is selected from at least one of the following PCSK9 residues: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, S381, W156, N157, L158, E159, H193, E195, H229, R237, G240, K243, D367, I368, G370, A371, S373, S376, Q382, W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, G384, K69, D70, P71, S148, V149, D186, T187, E211, D212, G213, R218, Q219, C223, D224, G227, H229, L253, N254, G259, P288, A290, G291, G316, R319, Y325, V346, G352, T353, G365, I368, I369, S372, S373, C378, F379, T385, S386, Q387, S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, F379, I154, T187,

H193, E195, I196, M201, V202, C223, T228, S235, G236, A239, G244, M247, I369, S372, C375, or C378.

[0042] In some embodiments, an isolated antibody or antigen binding molecule that binds to PCSK9 at a location that overlaps with a location that LDLR binds to PCSK9 is provided. In some embodiments, the location that LDLR binds to PCSK9 includes at least one amino acid residue selected from the group consisting of: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, and S381.

[0043] In some embodiments, an isolated antibody or antigen binding molecule that binds to PCSK9 is provided. In some embodiments, the antibody or antigen binding molecule reduces the likelihood that EGFa will bind to PCSK9 within 8 angstroms of at least one of the following residues on PCSK9: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, S381, W156, N157, L158, E159, H193, E195, H229, R237, G240, K243, D367, I368, G370, A371, S373, S376, or Q382.

[0044] In some embodiments, an antibody, antigen binding protein, or antigen binding molecule that binds to a surface of PCSK9 that overlaps with a surface that EGFa binds, Ab 21B12 binds, and/or 31H4 binds is provided. In some embodiments, an antibody, antigen binding protein, or antigen binding molecule that binds to PCSK9 in a manner that is similar to that depicted in the figures is provided.

[0045] In some embodiments, the above embodiments are neutralizing antibodies or antigen binding proteins. In some embodiments, the antigen binding protein is not LDLR or a fragment thereof (such as EGFa).

[0046] In some aspects, the invention comprises an isolated neutralizing antibody, wherein when the antibody is bound to PCSK9, the antibody is positioned 8 angstroms or less from at least one of the following residues of PCSK9: T468, R469, M470, A471, T472, R496, R499, E501, A502, Q503, R510, H512, F515, P540, P541, A542, E543, H565, W566, E567, V568, E569, R592, E593, S465, G466, P467, A473, I474, R476, G497, E498, M500, G504, K506, L507, V508, A511, N513, A514, G516, V536, T538, A539, A544, T548, D570, L571, H591, A594, S595, and H597 of SEQ ID NO: 3. In some embodiments, the antibody is positioned 5 angstroms or less from at least one of the following residues of PCSK9: T468, R469, M470, A471, T472, R496, R499, E501, A502, Q503, R510, H512, F515, P540, P541, A542, E543, H565, W566, E567, V568, E569, R592, and E593 of SEQ ID NO: 3.

[0047] In some aspects, the invention comprises an isolated antigen binding protein. The antigen binding protein comprises: A) a CDRH1 of the CDRH1 sequence in SEQ ID NO: 89, a CDRH2 of the CDRH2 sequence in SEQ ID NO: 89, and a CDRH3 of the CDRH3 sequence in SEQ ID NO: 89, and B) a CDRL1 of the CDRL1 sequence in SEQ ID NO:32, a CDRL2 of the CDRL2 sequence in SEQ ID NO:32, and a CDRL3 of the CDRL3 sequence in SEQ ID NO:32.

[0048] In some aspects, the invention comprises an isolated antigen binding protein that binds to a PCSK9 protein of SEQ ID NO: 1 where the binding between said isolated antigen binding protein and a variant PCSK9 protein is less than 50% of the binding between the isolated antigen binding protein and the PCSK9 protein of SEQ ID NO: 1 and/or SEQ ID NO: 303. In some embodiments, the variant PCSK9 protein comprises at least one mutation of a residue at a position selected from the group consisting or comprising 207, 208, 185, 181, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554, as shown in SEQ ID NO: 1. In some embodiments, the at least one mutation selected from the group comprising or consisting of R207E, D208R, E181R, R185E, R439E, E513R, V538R, E539R, T132R, S351R, A390R, A413R, and E582R. In some embodiments, the at least one mutation is selected from the group consisting of D162R, R164E, E167R, S123R, E129R, A311R, D313R, D337R, R519E, H521R,and Q554R.

[0049] In some aspects, the invention comprises an antigen binding protein that binds to a PCSK-9 protein of SEQ ID NO: 303 in a first manner and binds to a variant of PCSK9 in a second manner. The PCSK9 variant has at least one point mutation at a position selected from the group comprising or consisting of: 207, 208, 185, 181, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554 of SEQ ID NO: 303 and/or SEQ ID NO: 1. In some embodiments, the first manner comprises a first EC50, a first Bmax, or a first EC50 and a first Bmax. In some embodiments, the second manner comprises a second EC50, a second Bmax, or a second EC50 and a second Bmax. The value for the first manner is different from the value for the second manner. In some embodiments, the first manner comprises a first EC50, wherein the second manner involves a second EC50, and wherein the point mutation is selected from the group consisting or comprising: R207E, D208R, E181R, R185E, R439E, E513R, V538R, E539R, T132R, S351R, A390R, A413R, and E582R. In some embodiments, the first EC50 is at least 20% different from the second EC50. In some

embodiments, the first EC50 is at least 50% different from the second EC50. In some embodiments, the second EC50 is a larger numerical value than the first EC50. In some embodiments, the first EC50 is determined by a multiplex bead binding assay. In some embodiments, the second EC50 is greater than 1 um. In some embodiments, the antigen binding protein is a neutralizing antigen binding protein. In some embodiments, the neutralizing antigen binding protein is a competitive neutralizing antigen binding protein. In some embodiments, the neutralizing antigen binding protein is a non-competitive neutralizing antigen binding protein. In some embodiments, the first manner comprises a first Bmax and the second manner comprises a second Bmax that is different from the first Bmax. The PCSK9 variant has at least one point mutation selected from the group consisting or comprising: D162R, R164E, E167R, S123R, E129R, A311R, D313R, D337R, R519E, H521R,and Q554R. In some embodiments, the second Bmax is about 10% of the first Bmax. In some embodiments, the first Bmax is at least 20% different from the second Bmax. In some embodiments, the first Bmax is at least 50% different from the second Bmax.

[0050] In some aspects, the invention comprises an isolated antigen binding protein that binds to a PCSK9 protein of SEQ ID NO: 3, wherein the epitope of the antigen binding protein includes at least one of the following amino acids of SEQ ID NO: 1: 207, 208, 181, 185, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554.

[0051] In some aspects, the invention comprises an isolated neutralizing antigen binding protein that binds to a PCSK9 protein comprising the amino acid sequence of SEQ ID NO: 1, wherein the neutralizing antigen binding protein decreases the LDLR lowering effect of PCSK9 on LDLR. In some embodiments, the antigen binding protein is a LDLR non-competitive neutralizing antigen binding protein. In some embodiments, the antigen binding protein is a LDLR competitive neutralizing antigen binding protein.

[0052] In some aspects, the invention comprises an isolated antigen binding protein, wherein said antigen binding protein comprises: A) a CDRH1 of the CDRH1 sequence in SEQ ID NO: 49, a CDRH2 of the CDRH2 sequence in SEQ ID NO: 49, and a CDRH3 of the CDRH3 sequence in SEQ ID NO: 49, and B) a CDRL1 of the CDRL1 sequence in SEQ ID NO:23, a CDRL2 of the CDRL2 sequence in SEQ ID NO:23, and a CDRL3 of the CDRL3 sequence in SEQ ID NO:23.

[0053] In some aspects, the invention comprises a composition comprising a crystallized PCSK9 protein and an antigen binding protein that binds to PCSK9. The composition comprises the crystallized PCSK9 protein is such that the three dimensional structure of the PCSK9 protein can be determined to a resolution of about 2.2 angstroms or better. In some embodiments, the antigen binding protein is an antibody or a fragment thereof.

[0054] In some aspects, the invention comprises a crystallized PCSK9 protein and at least an EGFa section of a LDLR protein, wherein the EGFa section of the LDLR protein is bound by a PCSK9 protein, wherein said crystallized PCSK9 protein is such that the three dimensional structure of the PCSK9 protein can be determined to a resolution of about 2.2 angstroms or better. In some embodiments, the molecular model is on a computer readable medium.

[0055] In some aspects, the invention comprises the use of an antigen binding protein as described herein, in the preparation of a medicament for the lowering of serum cholesterol.

[0056] In some aspects, the invention comprises the use of an antigen binding protein as described herein, in the preparation of a medicament for treating or preventing a condition associated with elevated serum cholesterol levels in a subject.

[0057] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9, the antigen binding protein comprising: A) a heavy chain complementary determining region (CDRH) selected from at least one of the group consisting of: (i) a CDRH1 selected from the CDRH1 within the sequences selected from the group consisting of SEQ ID NOs: 67, 79, 89, and 49, (ii) a CDRH1 that differs in amino acid sequence from the CDRH1 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRH1 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}$ (SEQ ID NO: 406), wherein X_1 is selected from the group consisting of G, X_2 is selected from the group consisting of Y, F, and G, X_3 is selected from the group consisting of T and S, X_4 is selected from the group consisting of L and F, X_5 is selected from the group consisting of T, S, and N, X_6 is selected from the group consisting of S and A, X_7 is selected from the group consisting of Y and F, X_8 is selected from the group consisting of G, S, and Y, X_9 is selected from the group consisting of I, M, and W, X_{10} is selected from the group consisting of S, N and H, B) a light chain complementary determining region (CDRL) selected from at least one of the group consisting of: (i) a CDRL1 selected from the CDRL1 within the sequences selected from

the group consisting of SEQ ID NOs: 12, 32, 35, and 23, (ii) a CDRL1 that differs in amino acid sequence from the CDRL3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRL1 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}X_{12}X_{13}X_{14}$ (SEQ ID NO: 407), wherein X_1 is selected from the group consisting of T and no amino acid, X_2 is selected from the group consisting of G and S, X_3 is selected from the group consisting of S, T, and G, X_4 is selected from the group consisting of S, X_5 is selected from the group consisting of S, X_6 is selected from the group consisting of N, D, and S, X_7 is selected from the group consisting of I, V, and N, X_8 is selected from the group consisting of G and I, X_9 is selected from the group consisting of A and G, X_{10} is selected from the group consisting of G, Y, S, and N, X_{11} is selected from the group consisting of Y and N, X_{12} is selected from the group consisting of D, S, T, and F, X_{13} is selected from the group consisting of V, X_{14} is selected from the group consisting of S, N, and H. One of skill in the art will appreciate that a single ABP or antibody can meet one or more of the above options and still fall within the described invention for this embodiment.

[0058] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9, the antigen binding protein comprising: A) a heavy chain complementary determining region (CDRH) selected from at least one of the group consisting of the following: (i) a CDRH2 selected from the CDRH2 within the sequences selected from the group consisting of SEQ ID NOs: 67, 79, 89, and 49, (ii) a CDRH2 that differs in amino acid sequence from the CDRH2 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRH2 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}X_{12}X_{13}X_{14}$ $X_{15}X_{16}X_{17}$ (SEQ ID NO: 408), wherein X_1 is selected from the group consisting of W, S, L and no amino acid, X_2 is selected from the group consisting of V, I, and E, X_3 is selected from the group consisting of S, W, and I, X_4 is selected from the group consisting of F, S, and N, X_5 is selected from the group consisting of Y, S, D, and H, X_6 is selected from the group consisting of N, S, and G, X_7 is selected from the group consisting of S and G, X_8 is selected from the group consisting of N, Y, D, and R, X_9 is selected from the group consisting of T, I, and E, X_{10} is selected from the group consisting of N, S, Y, and D, X_{11} is selected from the group consisting of Y, X_{12} is selected from the group consisting of A and N, X_{13} is selected from the group consisting of Q, D, and P, X_{14} is selected from the group consisting of K and S, X_{15} is selected from the group consisting of L, and V, X_{16} is selected from

the group consisting of Q and K, X_{17} is selected from the group consisting of G and S, B) a light chain complementary determining region (CDRL) selected from at least one of the group consisting of the following: (i) a CDRL2 selected from the CDRL3 within the sequences selected from the group consisting of SEQ ID NOs: 12, 32, 35, and 23, (ii) a CDRL2 that differs in amino acid sequence from the CDRL3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRL2 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7$ (SEQ ID NO: 409), wherein X_1 is selected from the group consisting of G, E, S, and D, X_2 is selected from the group consisting of N, V, and Y, X_3 is selected from the group consisting of S and N, X_4 is selected from the group consisting of N, Q, and K, X_5 is selected from the group consisting of R, X_6 is selected from the group consisting of P, X_7 is selected from the group consisting of S.

[0059] In some aspects, the invention comprises An isolated antigen binding protein that binds PCSK9, the antigen binding protein comprising: A) a heavy chain complementary determining region (CDRH) selected from at least one of the group consisting of the following: (i) a CDRH3 selected from the CDRH3 within the sequences selected from the group consisting of SEQ ID NOs: 67, 79, 89, and 49, (ii) a CDRH3 that differs in amino acid sequence from the CDRH3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRH3 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}X_{12}X_{13}X_{14}$ (SEQ ID NO: 410), wherein X_1 is selected from the group consisting of D, and no amino acid, X_2 is selected from the group consisting of Y, A, and no amino acid, X_3 is selected from the group consisting of D, I, and no amino acid, X_4 is selected from the group consisting of F, A, and no amino acid, X_5 is selected from the group consisting of W, A, and no amino acid, X_6 is selected from the group consisting of S, L, and no amino acid, X_7 is selected from the group consisting of A, Y, G, and no amino acid, X_8 is selected from the group consisting of Y, Q, and no amino acid, X_9 is selected from the group consisting of G, Y, and L, X_{10} is selected from the group consisting of Y, D, and V, X_{11} is selected from the group consisting of G, A, and P, X_{12} is selected from the group consisting of M and F, X_{13} is selected from the group consisting of D, X_{14} is selected from the group consisting of V and Y, and B) a light chain complementary determining region (CDRL) selected from at least one of the group consisting of the following: (i) a CDRL3 selected from the CDRL3 within the sequences selected from the group consisting of SEQ ID NOs: 12, 32, 35, and 23, (ii) a CDRL3 that differs in amino

acid sequence from the CDRL3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRL3 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}$ (SEQ ID NO: 411), wherein X_1 is selected from the group consisting of Q, A, G, and no amino acid, X_2 is selected from the group consisting of S, V, T, and no amino acid, X_3 is selected from the group consisting of Y, N, and W, X_4 is selected from the group consisting of S and D, X_5 is selected from the group consisting of S, Y, and D, X_6 is selected from the group consisting of S and T, X_7 is selected from the group consisting of L and S, X_8 is selected from the group consisting of S, T, and N, X_9 is selected from the group consisting of G, S, and A, X_{10} is selected from the group consisting of S, M, W, and Y, and X_{11} is selected from the group consisting of V. In some embodiments, any of the above amino acids can be replaced by a conservative amino acid substitution.

[0060] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9, the antigen binding protein comprises A) a heavy chain complementary determining region (CDRH) selected from at least one of the group consisting of (i) a CDRH1 selected from the CDRH1 within the sequences selected from the group consisting of SEQ ID NOs: 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, and 58, (ii) a CDRH1 that differs in amino acid sequence from the CDRH1 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRH1 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}$ (SEQ ID NO: 412), wherein X_1 is selected from the group consisting of G, P, and A, X_2 is selected from the group consisting of Y, W, F, T, and S, X_3 is selected from the group consisting of T, P, S and A, C, V, L, and I, X_4 is selected from the group consisting of L, F, I, V, M, A, and Y, X_5 is selected from the group consisting of T, P, S, and A, X_6 is selected from the group consisting of S, T, A, and C, X_7 is selected from the group consisting of Y, W, F, T, and S, X_8 is selected from the group consisting of G, P, and A, X_9 is selected from the group consisting of I, L, V, M, A, and F, X_{10} is selected from the group consisting of S, T, A, and C, B) a light chain complementary determining region (CDRL) selected from at least one of the group consisting of: (i) a CDRL1 selected from the CDRL1 within the sequences selected from the group consisting of SEQ ID NOs: 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, and 24, (ii) a CDRL1 that differs in amino acid sequence from the CDRL3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRL1 amino acid sequence selected from the group consisting of

$X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}X_{12}X_{13}X_{14}$ (SEQ ID NO: 413), wherein, X_1 is selected from the group consisting of T and S, X_2 is selected from the group consisting of G, P, and A, X_3 is selected from the group consisting of T, and S, X_4 is selected from the group consisting of S N, T, A, C, and Q, X_5 is selected from the group consisting of S, T, A, and C, X_6 is selected from the group consisting of D, and E, X_7 is selected from the group consisting of V, I, M, L, F, and A, X_8 is selected from the group consisting of G, P, and A, X_9 is selected from the group consisting of G, A, R, P, V, L, I, K, Q, and N, X_{10} is selected from the group consisting of Y, W, F, T, and S, X_{11} is selected from the group consisting of N, and Q, X_{12} is selected from the group consisting of Y, S, W, F, T, A, and C, X_{13} is selected from the group consisting of V, I, M, L, F, and A, X_{14} is selected from the group consisting of S, T, A, and C.

[0061] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9, the antigen binding protein comprising: A) a heavy chain complementary determining region (CDRH) selected from at least one of the group consisting of: (i) a CDRH2 selected from the CDRH2 within the sequences selected from the group consisting of SEQ ID NOs: 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, and 58, (ii) a CDRH2 that differs in amino acid sequence from the CDRH2 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRH2 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9X_{10}X_{11}X_{12}X_{13}X_{14}$ $X_{15}X_{16}X_{17}$, (SEQ ID NO: 414), wherein X_1 is selected from the group consisting of W, Y, and F, X_2 is selected from the group consisting of V, I, M, L, F, and A, X_3 is selected from the group consisting of S, T, A, and C, X_4 is selected from the group consisting of A, F, V, L, I, Y, and M, X_5 is selected from the group consisting of Y, W, F, T, and S, X_6 is selected from the group consisting of N and Q, X_7 is selected from the group consisting of G, P, and A, X_8 is selected from the group consisting of N, and Q, X_9 is selected from the group consisting of T, and S, X_{10} is selected from the group consisting of N, and Q, X_{11} is selected from the group consisting of Y, W, F, T, and S, X_{12} is selected from the group consisting of A, V, L, and I, X_{13} is selected from the group consisting of Q, E, N, and D, X_{14} is selected from the group consisting of K, R, Q, and N, X_{15} is selected from the group consisting of L, F, V, I, M, A, and Y, X_{16} is selected from the group consisting of Q, and N, X_{17} is selected from the group consisting of G, P, and A, B) a light chain complementary determining region (CDRL) selected from at least one of the group consisting of: (i) a CDRL2 selected from the CDRL3 within the sequences selected from the group consisting of SEQ ID NOs: 14, 15, 16,

17, 18, 19, 20, 21, 22, 23, and 24, (ii) a CDRL2 that differs in amino acid sequence from the CDRL3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRL2 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7$ (SEQ ID NO: 415), wherein X_1 is selected from the group consisting of E, and D, X_2 is selected from the group consisting of V, I, M, L, F, and A, X_3 is selected from the group consisting of S, T, A, and C, X_4 is selected from the group consisting of N, and Q, X_5 is selected from the group consisting of R, K, Q, and N, X_6 is selected from the group consisting of P, and A, X_7 is selected from the group consisting of S, T, A, and C.

[0062] In some aspects, the invention comprises an isolated antigen binding protein that binds PCSK9, the antigen binding protein comprising: A) a heavy chain complementary determining region (CDRH) selected from at least one of the group consisting of (i) a CDRH3 selected from the CDRH3 within the sequences selected from the group consisting of SEQ ID NOs: 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, and 58, (ii) a CDRH3 that differs in amino acid sequence from the CDRH3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRH3 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6$ (SEQ ID NO: 416), wherein X_1 is selected from the group consisting of G, P, A and no amino acid, X_2 is selected from the group consisting of Y, W, F, T, and S, X_3 is selected from the group consisting of G, V, P, A, I, M, L, and F, X_4 is selected from the group consisting of M, L, F, and I, X_5 is selected from the group consisting of D, and E, X_6 is selected from the group consisting of V, I, M, L, F, and A, B) a light chain complementary determining region (CDRL) selected from at least one of the group consisting of: (i) a CDRL3 selected from the CDRL3 within the sequences selected from the group consisting of SEQ ID NOs: 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, and 24, (ii) a CDRL3 that differs in amino acid sequence from the CDRL3 of (i) by an amino acid addition, deletion or substitution of not more than two amino acids; and (iii) a CDRL3 amino acid sequence selected from the group consisting of $X_1X_2X_3X_4X_5X_6X_7X_8X_9$ (SEQ ID NO: 417), wherein X_1 is selected from the group consisting of S, N, T, A, C, and Q, X_2 is selected from the group consisting of S, T, A, and C, X_3 is selected from the group consisting of Y, W, F, T, and S, X_4 is selected from the group consisting of T, and S, X_5 is selected from the group consisting of S, T, A, and C, X_6 is selected from the group consisting of S, T, A, and C, X_7 is selected from the group consisting of N, S, Q, T, A, and C, X_8

is selected from the group consisting of M, V, L, F, I, and A, X_9 is selected from the group consisting of V, I, M, L, F, and A.

BRIEF DESCRIPTION OF THE FIGURES

[0063] FIG. 1A depicts an amino acid sequence of the mature form of the PCSK9 with the pro-domain underlined.

[0064] FIGs. $1B_1$-$1B_4$ depict amino acid and nucleic acid sequences of PCSK9 with the pro-domain underlined and the signal sequence in bold.

[0065] FIGs. 2A-2D are sequence comparison tables of various light chains of various antigen binding proteins. FIG. 2C continues the sequence started in FIG. 2A. FIG. 2D continues the sequence started on FIG. 2B.

[0066] FIGs. 3A-3D are sequence comparison tables of various heavy chains of various antigen binding proteins. FIG. 3C continues the sequence started in FIG. 3A. FIG. 3D continues the sequence started on FIG. 3B.

[0067] FIGs. 3E-3JJ depict the amino acid and nucleic acid sequences for the variable domains of some embodiments of the antigen binding proteins.

[0068] FIG. 3KK depicts the amino acid sequences for various constant domains.

[0069] FIGs. 3LL-3BBB depict the amino acid and nucleic acid sequences for the variable domains of some embodiments of the antigen binding proteins.

[0070] FIGs. 3CCC-3JJJ are sequence comparison tables of various heavy and light chains of some embodiments of the antigen binding proteins.

[0071] FIG. 4A is a binding curve of an antigen bindng protein to human PCSK9.

[0072] FIG. 4B is a binding curve of an antigen bindng protein to human PCSK9.

[0073] FIG. 4C is a binding curve of an antigen bindng protein to cynomolgus PCSK9.

[0074] FIG. 4D is a binding curve of an antigen bindng protein to cynomolgus PCSK9.

[0075] FIG. 4E is a binding curve of an antigen bindng protein to mouse PCSK9.

[0076] FIG. 4F is a binding curve of an antigen bindng protein to mouse PCSK9.

[0077] FIG. 5A depicts the results of an SDS PAGE experiement involving PCSK9 and various antigen binding proteins demonstrating the relative purity and concentration of the proteins.

[0078] FIG. 5B and 5C depict graphs from biacore solution equilibrium assays for 21B12.

[0079] FIG. 5D depicts the graph of the kinetics from a biacore capture assay.

[0080] FIG. 5E depicts a bar graph depicting binning results for three ABPs.

[0081] FIG. 6A is an inhibition curve of antigen binding protein 31H4 IgG2 to PCSK9 in an in vitro PCSK9:LDLR binding assay

[0082] FIG. 6B is an inhibition curve of antigen binding protein 31H4 IgG4 to PCSK9 in an in vitro PCSK9:LDLR binding assay.

[0083] FIG. 6C is an inhibition curve of antigen binding protein 21B12 IgG2 to PCSK9 in an in vitro PCSK9:LDLR binding assay.

[0084] FIG. 6D is an inhibition curve of antigen binding protein 21B12 IgG4 to PCSK9 in an in vitro PCSK9:LDLR binding assay.

[0085] FIG. 7A is an inhibition curve of antigen binding protein 31H4 IgG2 in the cell LDL uptake assay showing the effect of the ABP to reduce the LDL uptake blocking effects of PCSK9

[0086] FIG. 7B is an inhibition curve of antigen binding protein 31H4 IgG4 in the cell LDL uptake assay showing the effect of the ABP to reduce the LDL uptake blocking effects of PCSK9

[0087] FIG. 7C is an inhibition curve of antigen binding protein 21B12 IgG2 in the cell LDL uptake assay showing the effect of the ABP to reduce the LDL uptake blocking effects of PCSK9

[0088] FIG. 7D is an inhibition curve of antigen binding protein 21B12 IgG4 in the cell LDL uptake assay showing the effect of the ABP to reduce the LDL uptake blocking effects of PCSK9

[0089] FIG. 8A is a graph depicting the serum cholesterol lowering ability in mice of ABP 31H4, changes relative to the IgG control treated mice (* $p < 0.01$).

[0090] FIG. 8B is a graph depicting the serum cholesterol lowering ability in mice of ABP 31H4, changes relative to time = zero hours (# p, 0.05).

[0091] FIG. 8C is a graph depicting the effect of ABP 31H4 on HDL cholesterol levels in C57B1/6 mice (* $p < 0.01$).

[0092] FIG. 8D is a graph depicting the effect of ABP 31H4 on HDL cholesterol levels in C57Bl/6 mice (# $p < 0.05$).

[0093] FIG. 9 depicts a western blot analysis of the ability of ABP 31H4 to enhance the amount of liver LDLR protein present after various time points.

[0094] FIG. 10A is a graph depicting the ability of an antigen binding protein 31H4 to lower total serum cholesterol in wild type mice, relative.

[0095] FIG. 10B is a graph depicting the ability of an antigen binding protein 31H4 to lower HDL in wild type mice.

[0096] FIG. 10C is a graph depicting the serum cholesterol lowering ability of various antigen binding proteins 31H4 and 16F12.

[0097] FIG. 11A depicts an injection protocol for testing the duration and ability of antigen binding proteins to lower serum cholesterol.

[0098] FIG. 11B is a graph depicting the results of the protocol in FIG. 11A.

[0099] FIG. 12A depicts LDLR levels in response to the combination of a statin and ABP 21B12 in HepG2 cells.

[0100] FIG. 12B depicts LDLR levels in response to the combination of a statin and ABP 31H4 in HepG2 cells.

[0101] FIG. 12C depicts LDLR levels in response to the combination of a statin and ABP 25A7.1, a nonneutralizing antibody, (in contrast the "25A7" a neutralizing antibody) in HepG2 cells.

[0102] FIG. 12D depicts LDLR levels in response to the combination of a statin and ABP 21B12 in HepG2 cells overexpressing PCSK9.

[0103] FIG. 12E depicts LDLR levels in response to the combination of a statin and ABP 31H4 in HepG2 cells overexpressing PCSK9.

[0104] FIG. 12F depicts LDLR levels in response to the combination of a statin and ABP 25A7.1, a nonneutralizing antibody, (in contrast the "25A7" a neutralizing antibody) in HepG2 cells overexpressing PCSK9.

[0105] FIG. 13A depicts the various light chain amino acid sequences of various ABPs to PCSK9. The dots (.) indicate no amino acid.

[0106] FIG. 13B depicts a light chain cladogram for various ABPs to PCSK9.

[0107] FIG. 13C depicts the various heavy chain amino acid sequences of various ABPs to PCSK9. The dots (.) indicate no amino acid.

[0108] FIG. 13D depicts a heavy chain dendrogram for various ABPs to PCSK9.

[0109] FIG. 13E depicts a comparison of light and heavy CDRs and designation of groups from which to derive consensus.

[0110] FIG. 13F depicts the consensus sequences for Groups 1 and 2.

[0111] FIG. 13G depicts the consensus sequences for Groups 3 and 4.

[0112] FIG. 13H depicts the consensus sequences for Groups 1 and 2. The dots (.) indicated identical residues.

[0113] FIG. 13I depicts the consensus sequences for Group 2. The dots (.) indicated identical residues.

[0114] FIG. 13J depicts the consensus sequences for Groups 3 and 4. The dots (.) indicated identical residues.

[0115] FIG. 14A is a graph depicting *in vivo* LDL lowering ability of various ABPs (at 10 mg/kg).

[0116] FIG. 14B is a graph depicting *in vivo* LDL lowering ability of various ABPs (at 30 mg/kg).

[0117] FIG. 15A and FIG. 15B are sequence comparison tables of various light chains of various embodiments of antigen binding proteins. FIG. 15B continues the sequence started in FIG. 15A.

[0118] FIG. 15C and FIG. 15D are sequence comparison tables of various light chains of various embodiments of antigen binding proteins. FIG. 15D continues the sequence started in FIG. 15C.

[0119] FIG. 16A is a depiction of a gel used to test the ability of Ab 21B12 to bind to the ProCat or VD sections of PCSK9.

[0120] FIG. 16B is a depiction of a gel used to test the ability of Ab 31H4 to bind to the ProCat or VD sections of PCSK9.

[0121] FIG. 17 is a depiction of the structure of PCSK9 and the EGFa section of LDLR.

[0122] FIG. 18A is a depiction of the structure of PCSK9 and the 31H4 Ab.

[0123] FIG. 18B is a depiction of the structure of PCSK9 and the 31H4 Ab.

[0124] FIG. 19A is a depiction of the structure of PCSK9, the 31H4 Ab, and the 21B12 Ab.

[0125] FIG. 19B is a depiction of the structure of PCSK9 and the 21B12 Ab.

[0126] FIG. 20A is a depiction of the structure of PCSK9 and EGFa from the LDLR superimposed with the structure of antibodies 31H4 and 21B12 bound to PCSK9.

[0127] FIG. 20B is a depiction of the structural model of PCSK9 and LDLR.

[0128] FIG. 20C is a depiction of the structural model of PCSK9 and LDLR from an alternative perspective.

[0129] FIG. 20D is a depiction of the structural model of PCSK9 and LDLR with structural representations of 31H4 and 21B12 included.

[0130] FIG. 20E is a depiction of the structural model in FIG. 20D, rotated 90 degrees about the noted axis.

[0131] FIG. 20F is a depiction of the structural model in FIG. 20D rotated 180 degrees about the noted axis.

[0132] FIG. 21A is a depiction of the structure of PCSK9 and 31A4.

[0133] FIG. 21B is a depiction of the structure of PCSK9 and 31A4.

[0134] FIG. 21C is a depiction of the structure of PCSK9 and 31A4.

[0135] FIG. 21D is a depiction of the structural model of full length PCSK9 and 31A4.

[0136] FIG. 22 is a set of ABP sequences identifying various differences between the human ABP sequences and the ABP sequences that were raised in E. coli and used for the crystal structures.

[0137] FIG. 23 is a table of the various binning results.

[0138] FIG. 23A is a first part of a table depicting the various binning results.

[0139] FIG. 23B is a second part of a table depicting the various binning results.

[0140] FIG. 23C is a third part of a table depicting the various binning results.

[0141] FIG. 23D is a fourth part of a table depicting the various binning results.

[0142] FIG. 24A is a depiction of a western blot under non-reduced conditions.

[0143] FIG. 24B is a depiction of a western blot under reduced conditions.

[0144] FIG. 25A is a depiction of the surface coverage of PCSK9.

[0145] FIG. 25B is a depiction of the surface coverage of PCSK9.

[0146] FIG. 25C is a depiction of the surface coverage of PCSK9.

[0147] FIG. 25D is a depiction of the surface coverage of PCSK9.

[0148] FIG. 25E is a depiction of the surface coverage of PCSK9.

[0149] FIG. 25F is a depiction of the surface coverage of PCSK9.

[0150] FIG. 26 is a sequence comparison of the PCSK9 amino acid sequence and all of the residues that were mutated in PCSK9 variants to examine the epitopes of the various antibodies.

[0151] FIG. 27A depicts the 21B12 epitope hits, as mapped onto a crystal structure of PCSK9 with the 21B12.

[0152] FIG. 27B depicts the 31H4 epitope hits, as mapped ont a crystal structure of PCSK9 with 31H4 and 21B1.

[0153] FIG. 27C depicts the 31A4 epitope hits, as mapped onto a crystal structure of PCSK9 with 31H4 and 21B12.

[0154] FIG. 27D depicts the 12H11 epitope hits, as mapped onto the crystal structure of PCSK9 with 31H4 and 21B12.

[0155] FIG. 27E depicts the 3C4 epitope hits, as mapped onto the crystal structure of PCSK9 with 31H4 and 21B12.

[0156] FIG. 28A is a graph demonstrating the binding ability of the various ABPs to various parts of PCSK9.

[0157] FIG. 28B is a graph demonstrating the binding ability of the various ABPs to various parts of PCSK9.

[0158] FIG. 28C is a graph comparing the LDLR binding ability of two ABPs.

[0159] FIG. 28D is a graph comparing the cell LDL uptake activity of two ABPs.

DETAILED DESCRIPTION OF CERTAIN EXEMPLARY EMBODIMENTS

[0160] Antigen binding proteins (such as antibodies and functional binding fragments thereof) that bind to PCSK9 are disclosed herein. In some embodiments, the antigen binding proteins bind to PCSK9 and prevent PCSK9 from functioning in various ways. In some embodiments, the antigen binding proteins block or reduce the ability of PCSK9 to interact with other substances. For example, in some embodiments, the antigen binding protein binds to

PCSK9 in a manner that prevents or reduces the likelihood that PCSK9 will bind to LDLR. In other embodiments, antigen binding proteins bind to PCSK9 but do not block PCSK9's ability to interact with LDLR. In some embodiments, the antigen binding proteins are human monoclonal antibodies.

[0161] As will be appreciated by one of skill in the art, in light of the present disclosure, altering the interactions between PCSK9 and LDLR can increase the amount of LDLR available for binding to LDL, which in turn decreases the amount of serum LDL in a subject, resulting in a reduction in the subject's serum cholesterol level. As such, antigen binding proteins to PCSK9 can be used in various methods and compositions for treating subjects with elevated serum cholesterol levels, at risk of elevated serum cholesterol levels, or which could benefit from a reduction in their serum cholesterol levels. Thus, various methods and techniques for lowering, maintaining, or preventing an increase in serum cholesterol are also described herein. In some embodiments, the antigen binding protein allows for binding between PCSK9 and LDLR, but the antigen binding protein prevents or reduces the adverse activity of PCSK9 on LDLR. In some embodiments, the antigen binding protein prevents or reduces the binding of PCSK9 to LDLR.

[0162] For convenience, the following sections generally outline the various meanings of the terms used herein. Following this discussion, general aspects regarding antigen binding proteins are discussed, followed by specific examples demonstrating the properties of various embodiments of the antigen binding proteins and how they can be employed.

Definitions and Embodiments

[0163] It is to be understood that both the foregoing general description and the following detailed description are exemplary and explanatory only and are not restrictive of the invention as claimed. In this application, the use of the singular includes the plural unless specifically stated otherwise. In this application, the use of "or" means "and/or" unless stated otherwise. Furthermore, the use of the term "including", as well as other forms, such as "includes" and "included", is not limiting. Also, terms such as "element" or "component" encompass both elements and components comprising one unit and elements and components that comprise more than one subunit unless specifically stated otherwise. Also, the use of the term "portion" can include part of a moiety or the entire moiety.

[0164] The section headings used herein are for organizational purposes only and are not to be construed as limiting the subject matter described. All documents, or portions of documents, cited in this application, including but not limited to patents, patent applications, articles, books, and treatises, are hereby expressly incorporated by reference in their entirety for any purpose. As utilized in accordance with the present disclosure, the following terms, unless otherwise indicated, shall be understood to have the following meanings:

[0165] The term "proprotein convertase subtilisin kexin type 9" or "PCSK9" refers to a polypeptide as set forth in SEQ ID NO: 1 and/or 3 or fragments thereof, as well as related polypeptides, which include, but are not limited to, allelic variants, splice variants, derivative variants, substitution variants, deletion variants, and/or insertion variants including the addition of an N-terminal methionine, fusion polypeptides, and interspecies homologs. In certain embodiments, a PCSK9 polypeptide includes terminal residues, such as, but not limited to, leader sequence residues, targeting residues, amino terminal methionine residues, lysine residues, tag residues and/or fusion protein residues. "PCSK9" has also been referred to as FH3, NARC1, HCHOLA3, proprotein convertase subtilisin/kexin type 9, and neural apoptosis regulated convertase 1. The PCSK9 gene encodes a proprotein convertase protein that belongs to the proteinase K subfamily of the secretory subtilase family. The term "PCSK9" denotes both the proprotein and the product generated following autocatalysis of the proprotein. When only the autocatalyzed product is being referred to (such as for an antigen binding protein that selectively binds to the cleaved PCSK9), the protein can be referred to as the "mature," "cleaved", "processed" or "active" PCSK9. When only the inactive form is being referred to, the protein can be referred to as the "inactive", "pro-form", or "unprocessed" form of PCSK9. The term PCSK9 as used herein also includes naturally occurring alleles, such as the mutations D374Y, S127R and F216L. The term PCSK9 also encompasses PCSK9 molecules incorporating post-translational modifications of the PCSK9 amino acid sequence, such as PCSK9 sequences that have been glycosylated, PEGylated, PCSK9 sequences from which its signal sequence has been cleaved, PCSK9 sequence from which its pro domain has been cleaved from the catalytic domain but not separated from the catalytic domain (e.g., FIGs. 1A and 1B).

[0166] The term "PCSK9 activity" includes any biological effect of PCSK9. In certain embodiments, PCSK9 activity includes the ability of PCSK9 to interact or bind to a substrate or receptor. In some embodiments, PCSK9 activity is represented by the ability of

PCSK9 to bind to a LDL receptor (LDLR). In some embodiments, PCSK9 binds to and catalyzes a reaction involving LDLR. In some embodiments, PCSK9 activity includes the ability of PCSK9 to alter (e.g., reduce) the availability of LDLR. In some embodiments, PCSK9 activity includes the ability of PCSK9 to increase the amount of LDL in a subject. In some embodiments, PCSK9 activity includes the ability of PCSK9 to decrease the amount of LDLR that is available to bind to LDL. In some embodiments, "PCSK9 activity" includes any biological activity resulting from PCSK9 signaling. Exemplary activities include, but are not limited to, PCSK9 binding to LDLR, PCSK9 enzyme activity that cleaves LDLR or other proteins, PCSK9 binding to proteins other than LDLR that facilitate PCSK9 action, PCSK9 altering APOB secretion (Sun X-M et al, "Evidence for effect of mutant PCSK9 on apoliprotein B secretion as the cause of unusually severe dominant hypercholesterolemia, Human Molecular Genetics 14: 1161-1169, 2005 and Ouguerram K et al, "Apolipoprotein B100 metabolism in autosomal-dominant hypercholesterolemia related to mutations in PCSK9, Arterioscler thromb Vasc Biol. 24: 1448-1453, 2004), PCSK9's role in liver regeneration and neuronal cell differentiation (Seidah NG et al, "The secretory proprotein convertase neural apoptosis-regulated convertase 1 (NARC-1): Liver regeneration and neuronal differentiation" PNAS 100: 928-933, 2003), and PCSK9s role in hepatic glucose metabolism (Costet et al., "Hepatic PCSK9 expression is regulated by nutritional status via insulin and sterol regulatory element-binding protein 1c" J. Biol. Chem. 281(10):6211-18, 2006).

[0167] The term "hypercholesterolemia," as used herein, refers to a condition in which cholesterol levels are elevated above a desired level. In some embodiments, this denotes that serum cholesterol levels are elevated. In some embodiments, the desired level takes into account various "risk factors" that are known to one of skill in the art (and are described or referenced herein).

[0168] The term "polynucleotide" or "nucleic acid" includes both single-stranded and double-stranded nucleotide polymers. The nucleotides comprising the polynucleotide can be ribonucleotides or deoxyribonucleotides or a modified form of either type of nucleotide. Said modifications include base modifications such as bromouridine and inosine derivatives, ribose modifications such as 2',3'-dideoxyribose, and internucleotide linkage modifications such as phosphorothioate, phosphorodithioate, phosphoroselenoate, phosphorodiselenoate, phosphoroanilothioate, phoshoraniladate and phosphoroamidate.

[0169] The term "oligonucleotide" means a polynucleotide comprising 200 or fewer nucleotides. In some embodiments, oligonucleotides are 10 to 60 bases in length. In other embodiments, oligonucleotides are 12, 13, 14, 15, 16, 17, 18, 19, or 20 to 40 nucleotides in length. Oligonucleotides can be single stranded or double stranded, *e.g.*, for use in the construction of a mutant gene. Oligonucleotides can be sense or antisense oligonucleotides. An oligonucleotide can include a label, including a radiolabel, a fluorescent label, a hapten or an antigenic label, for detection assays. Oligonucleotides can be used, for example, as PCR primers, cloning primers or hybridization probes.

[0170] An "isolated nucleic acid molecule" means a DNA or RNA of genomic, mRNA, cDNA, or synthetic origin or some combination thereof which is not associated with all or a portion of a polynucleotide in which the isolated polynucleotide is found in nature, or is linked to a polynucleotide to which it is not linked in nature. For purposes of this disclosure, it should be understood that "a nucleic acid molecule comprising" a particular nucleotide sequence does not encompass intact chromosomes. Isolated nucleic acid molecules "comprising" specified nucleic acid sequences can include, in addition to the specified sequences, coding sequences for up to ten or even up to twenty other proteins or portions thereof, or can include operably linked regulatory sequences that control expression of the coding region of the recited nucleic acid sequences, and/or can include vector sequences.

[0171] Unless specified otherwise, the left-hand end of any single-stranded polynucleotide sequence discussed herein is the 5' end; the left-hand direction of double-stranded polynucleotide sequences is referred to as the 5' direction. The direction of 5' to 3' addition of nascent RNA transcripts is referred to as the transcription direction; sequence regions on the DNA strand having the same sequence as the RNA transcript that are 5' to the 5' end of the RNA transcript are referred to as "upstream sequences;" sequence regions on the DNA strand having the same sequence as the RNA transcript that are 3' to the 3' end of the RNA transcript are referred to as "downstream sequences."

[0172] The term "control sequence" refers to a polynucleotide sequence that can affect the expression and processing of coding sequences to which it is ligated. The nature of such control sequences can depend upon the host organism. In particular embodiments, control sequences for prokaryotes can include a promoter, a ribosomal binding site, and a transcription termination sequence. For example, control sequences for eukaryotes can include promoters

comprising one or a plurality of recognition sites for transcription factors, transcription enhancer sequences, and transcription termination sequence. "Control sequences" can include leader sequences and/or fusion partner sequences.

[0173] The term "vector" means any molecule or entity (*e.g.*, nucleic acid, plasmid, bacteriophage or virus) used to transfer protein coding information into a host cell.

[0174] The term "expression vector" or "expression construct" refers to a vector that is suitable for transformation of a host cell and contains nucleic acid sequences that direct and/or control (in conjunction with the host cell) expression of one or more heterologous coding regions operatively linked thereto. An expression construct can include, but is not limited to, sequences that affect or control transcription, translation, and, if introns are present, affect RNA splicing of a coding region operably linked thereto.

[0175] As used herein, "operably linked" means that the components to which the term is applied are in a relationship that allows them to carry out their inherent functions under suitable conditions. For example, a control sequence in a vector that is "operably linked" to a protein coding sequence is ligated thereto so that expression of the protein coding sequence is achieved under conditions compatible with the transcriptional activity of the control sequences.

[0176] The term "host cell" means a cell that has been transformed, or is capable of being transformed, with a nucleic acid sequence and thereby expresses a gene of interest. The term includes the progeny of the parent cell, whether or not the progeny is identical in morphology or in genetic make-up to the original parent cell, so long as the gene of interest is present.

[0177] The term "transfection" means the uptake of foreign or exogenous DNA by a cell, and a cell has been "transfected" when the exogenous DNA has been introduced inside the cell membrane. A number of transfection techniques are well known in the art and are disclosed herein. *See, e.g.*, Graham *et al.*, 1973, *Virology* 52:456; Sambrook *et al.*, 2001, Molecular Cloning: A Laboratory Manual, *supra*; Davis *et al.*, 1986, *Basic Methods in Molecular Biology*, Elsevier; Chu *et al.*, 1981, *Gene* 13:197. Such techniques can be used to introduce one or more exogenous DNA moieties into suitable host cells.

[0178] The term "transformation" refers to a change in a cell's genetic characteristics, and a cell has been transformed when it has been modified to contain new DNA or RNA. For example, a cell is transformed where it is genetically modified from its native state by

introducing new genetic material *via* transfection, transduction, or other techniques. Following transfection or transduction, the transforming DNA can recombine with that of the cell by physically integrating into a chromosome of the cell, or can be maintained transiently as an episomal element without being replicated, or can replicate independently as a plasmid. A cell is considered to have been "stably transformed" when the transforming DNA is replicated with the division of the cell.

[0179] The terms "polypeptide" or "protein" means a macromolecule having the amino acid sequence of a native protein, that is, a protein produced by a naturally-occurring and non-recombinant cell; or it is produced by a genetically-engineered or recombinant cell, and comprise molecules having the amino acid sequence of the native protein, or molecules having deletions from, additions to, and/or substitutions of one or more amino acids of the native sequence. The term also includes amino acid polymers in which one or more amino acids are chemical analogs of a corresponding naturally-occurring amino acid and polymers. The terms "polypeptide" and "protein" specifically encompass PCSK9 antigen binding proteins, antibodies, or sequences that have deletions from, additions to, and/or substitutions of one or more amino acid of antigen-binding protein. The term "polypeptide fragment" refers to a polypeptide that has an amino-terminal deletion, a carboxyl-terminal deletion, and/or an internal deletion as compared with the full-length native protein. Such fragments can also contain modified amino acids as compared with the native protein. In certain embodiments, fragments are about five to 500 amino acids long. For example, fragments can be at least 5, 6, 8, 10, 14, 20, 50, 70, 100, 110, 150, 200, 250, 300, 350, 400, or 450 amino acids long. Useful polypeptide fragments include immunologically functional fragments of antibodies, including binding domains. In the case of a PCSK9-binding antibody, useful fragments include but are not limited to a CDR region, a variable domain of a heavy and/or light chain, a portion of an antibody chain or just its variable region including two CDRs, and the like.

[0180] The term "isolated protein" referred means that a subject protein (1) is free of at least some other proteins with which it would normally be found, (2) is essentially free of other proteins from the same source, *e.g.*, from the same species, (3) is expressed by a cell from a different species, (4) has been separated from at least about 50 percent of polynucleotides, lipids, carbohydrates, or other materials with which it is associated in nature, (5) is operably associated (by covalent or noncovalent interaction) with a polypeptide with which it is not associated in

nature, or (6) does not occur in nature. Typically, an "isolated protein" constitutes at least about 5%, at least about 10%, at least about 25%, or at least about 50% of a given sample. Genomic DNA, cDNA, mRNA or other RNA, of synthetic origin, or any combination thereof can encode such an isolated protein. Preferably, the isolated protein is substantially free from proteins or polypeptides or other contaminants that are found in its natural environment that would interfere with its therapeutic, diagnostic, prophylactic, research or other use.

[0181] The term "amino acid" includes its normal meaning in the art.

[0182] A "variant" of a polypeptide (*e.g.*, an antigen binding protein, or an antibody) comprises an amino acid sequence wherein one or more amino acid residues are inserted into, deleted from and/or substituted into the amino acid sequence relative to another polypeptide sequence. Variants include fusion proteins.

[0183] The term "identity" refers to a relationship between the sequences of two or more polypeptide molecules or two or more nucleic acid molecules, as determined by aligning and comparing the sequences. "Percent identity" means the percent of identical residues between the amino acids or nucleotides in the compared molecules and is calculated based on the size of the smallest of the molecules being compared. For these calculations, gaps in alignments (if any) are preferably addressed by a particular mathematical model or computer program (*i.e.*, an "algorithm"). Methods that can be used to calculate the identity of the aligned nucleic acids or polypeptides include those described in *Computational Molecular Biology*, (Lesk, A. M., ed.), 1988, New York: Oxford University Press; Biocomputing Informatics and Genome Projects, (Smith, D. W., ed.), 1993, New York: Academic Press; Computer Analysis of Sequence Data, Part I, (Griffin, A. M., and Griffin, H. G., eds.), 1994, New Jersey: Humana Press; von Heinje, G., 1987, Sequence Analysis in Molecular Biology, New York: Academic Press; Sequence Analysis Primer, (Gribskov, M. and Devereux, J., eds.), 1991, New York: M. Stockton Press; and Carillo *et al.*, 1988, *SIAM J. Applied Math.* 48:1073.

[0184] In calculating percent identity, the sequences being compared are typically aligned in a way that gives the largest match between the sequences. One example of a computer program that can be used to determine percent identity is the GCG program package, which includes GAP (Devereux *et al.*, 1984, *Nucl. Acid Res.* 12:387; Genetics Computer Group, University of Wisconsin, Madison, WI). The computer algorithm GAP is used to align the two polypeptides or polynucleotides for which the percent sequence identity is to be determined. The

sequences are aligned for optimal matching of their respective amino acid or nucleotide (the "matched span", as determined by the algorithm). A gap opening penalty (which is calculated as 3x the average diagonal, wherein the "average diagonal" is the average of the diagonal of the comparison matrix being used; the "diagonal" is the score or number assigned to each perfect amino acid match by the particular comparison matrix) and a gap extension penalty (which is usually 1/10 times the gap opening penalty), as well as a comparison matrix such as PAM 250 or BLOSUM 62 are used in conjunction with the algorithm. In certain embodiments, a standard comparison matrix (*see*, Dayhoff *et al.*, 1978, *Atlas of Protein Sequence and Structure* 5:345-352 for the PAM 250 comparison matrix; Henikoff *et al.*, 1992, *Proc. Natl. Acad. Sci. U.S.A.* 89:10915-10919 for the BLOSUM 62 comparison matrix) is also used by the algorithm.

[0185] Examples of parameters that can be employed in determining percent identity for polypeptides or nucleotide sequences using the GAP program are the following:

- Algorithm: Needleman *et al.*, 1970, *J. Mol. Biol.* 48:443-453
- Comparison matrix: BLOSUM 62 from Henikoff *et al.*, 1992, *supra*
- Gap Penalty: 12 (but with no penalty for end gaps)
- Gap Length Penalty: 4
- Threshold of Similarity: 0

[0186] Certain alignment schemes for aligning two amino acid sequences may result in matching of only a short region of the two sequences, and this small aligned region may have very high sequence identity even though there is no significant relationship between the two full-length sequences. Accordingly, the selected alignment method (GAP program) can be adjusted if so desired to result in an alignment that spans at least 50 or other number of contiguous amino acids of the target polypeptide.

[0187] As used herein, the twenty conventional (*e.g.*, naturally occurring) amino acids and their abbreviations follow conventional usage. See *Immunology--A Synthesis* (2nd Edition, E. S. Golub and D. R. Gren, Eds., Sinauer Associates, Sunderland, Mass. (1991)), which is incorporated herein by reference for any purpose. Stereoisomers (e.g., D-amino acids) of the twenty conventional amino acids, unnatural amino acids such as α-, α-disubstituted amino acids, N-alkyl amino acids, lactic acid, and other unconventional amino acids can also be suitable components for polypeptides of the present invention. Examples of unconventional amino acids include: 4-hydroxyproline, γ-carboxyglutamate, ε-N,N,N-trimethyllysine, ε-N-acetyllysine, O-

phosphoserine, N-acetylserine, N-formylmethionine, 3-methylhistidine, 5-hydroxylysine, σ-N-methylarginine, and other similar amino acids and imino acids (e.g., 4-hydroxyproline). In the polypeptide notation used herein, the left-hand direction is the amino terminal direction and the right-hand direction is the carboxy-terminal direction, in accordance with standard usage and convention.

[0188] Similarly, unless specified otherwise, the left-hand end of single-stranded polynucleotide sequences is the 5' end; the left-hand direction of double-stranded polynucleotide sequences is referred to as the 5' direction. The direction of 5' to 3' addition of nascent RNA transcripts is referred to as the transcription direction; sequence regions on the DNA strand having the same sequence as the RNA and which are 5' to the 5' end of the RNA transcript are referred to as "upstream sequences"; sequence regions on the DNA strand having the same sequence as the RNA and which are 3' to the 3' end of the RNA transcript are referred to as "downstream sequences."

[0189] Conservative amino acid substitutions can encompass non-naturally occurring amino acid residues, which are typically incorporated by chemical peptide synthesis rather than by synthesis in biological systems. These include peptidomimetics and other reversed or inverted forms of amino acid moieties.

[0190] Naturally occurring residues can be divided into classes based on common side chain properties:

1) hydrophobic: norleucine, Met, Ala, Val, Leu, Ile;
2) neutral hydrophilic: Cys, Ser, Thr, Asn, Gln;
3) acidic: Asp, Glu;
4) basic: His, Lys, Arg;
5) residues that influence chain orientation: Gly, Pro; and
6) aromatic: Trp, Tyr, Phe.

For example, non-conservative substitutions can involve the exchange of a member of one of these classes for a member from another class. Such substituted residues can be introduced, for example, into regions of a human antibody that are homologous with non-human antibodies, or into the non-homologous regions of the molecule.

[0191] In making changes to the antigen binding protein or the PCSK9 protein, according to certain embodiments, the hydropathic index of amino acids can be considered.

Each amino acid has been assigned a hydropathic index on the basis of its hydrophobicity and charge characteristics. They are: isoleucine (+4.5); valine (+4.2); leucine (+3.8); phenylalanine (+2.8); cysteine/cystine (+2.5); methionine (+1.9); alanine (+1.8); glycine (-0.4); threonine (-0.7); serine (-0.8); tryptophan (-0.9); tyrosine (-1.3); proline (-1.6); histidine (-3.2); glutamate (-3.5); glutamine (-3.5); aspartate (-3.5); asparagine (-3.5); lysine (-3.9); and arginine (-4.5).

[0192] The importance of the hydropathic amino acid index in conferring interactive biological function on a protein is understood in the art. Kyte *et al.*, J. Mol. Biol., 157:105-131 (1982). It is known that certain amino acids can be substituted for other amino acids having a similar hydropathic index or score and still retain a similar biological activity. In making changes based upon the hydropathic index, in certain embodiments, the substitution of amino acids whose hydropathic indices are within ±2 is included. In certain embodiments, those which are within ±1 are included, and in certain embodiments, those within ±0.5 are included.

[0193] It is also understood in the art that the substitution of like amino acids can be made effectively on the basis of hydrophilicity, particularly where the biologically functional protein or peptide thereby created is intended for use in immunological embodiments, as in the present case. In certain embodiments, the greatest local average hydrophilicity of a protein, as governed by the hydrophilicity of its adjacent amino acids, correlates with its immunogenicity and antigenicity, i.e., with a biological property of the protein.

[0194] The following hydrophilicity values have been assigned to these amino acid residues: arginine (+3.0); lysine (+3.0); aspartate (+3.0 ± 1); glutamate (+3.0 ± 1); serine (+0.3); asparagine (+0.2); glutamine (+0.2); glycine (0); threonine (-0.4); proline (-0.5 ± 1); alanine (-0.5); histidine (-0.5); cysteine (-1.0); methionine (-1.3); valine (-1.5); leucine (-1.8); isoleucine (-1.8); tyrosine (-2.3); phenylalanine (-2.5) and tryptophan (-3.4). In making changes based upon similar hydrophilicity values, in certain embodiments, the substitution of amino acids whose hydrophilicity values are within ±2 is included, in certain embodiments, those which are within ±1 are included, and in certain embodiments, those within ±0.5 are included. One can also identify epitopes from primary amino acid sequences on the basis of hydrophilicity. These regions are also referred to as "epitopic core regions."

[0195] Exemplary amino acid substitutions are set forth in Table 1.

TABLE 1

Amino Acid Substitutions

Original Residues	Exemplary Substitutions	Preferred Substitutions
Ala	Val, Leu, Ile	Val
Arg	Lys, Gln, Asn	Lys
Asn	Gln	Gln
Asp	Glu	Glu
Cys	Ser, Ala	Ser
Gln	Asn	Asn
Glu	Asp	Asp
Gly	Pro, Ala	Ala
His	Asn, Gln, Lys, Arg	Arg
Ile	Leu, Val, Met, Ala, Phe, Norleucine	Leu
Leu	Norleucine, Ile, Val, Met, Ala, Phe	Ile
Lys	Arg, 1,4 Diamino-butyric Acid, Gln, Asn	Arg
Met	Leu, Phe, Ile	Leu
Phe	Leu, Val, Ile, Ala, Tyr	Leu
Pro	Ala	Gly
Ser	Thr, Ala, Cys	Thr
Thr	Ser	Ser
Trp	Tyr, Phe	Tyr
Tyr	Trp, Phe, Thr, Ser	Phe
Val	Ile, Met, Leu, Phe, Ala, Norleucine	Leu

[0196] The term "derivative" refers to a molecule that includes a chemical modification other than an insertion, deletion, or substitution of amino acids (or nucleic acids). In certain embodiments, derivatives comprise covalent modifications, including, but not limited

to, chemical bonding with polymers, lipids, or other organic or inorganic moieties. In certain embodiments, a chemically modified antigen binding protein can have a greater circulating half-life than an antigen binding protein that is not chemically modified. In certain embodiments, a chemically modified antigen binding protein can have improved targeting capacity for desired cells, tissues, and/or organs. In some embodiments, a derivative antigen binding protein is covalently modified to include one or more water soluble polymer attachments, including, but not limited to, polyethylene glycol, polyoxyethylene glycol, or polypropylene glycol. *See, e.g.*, U.S. Patent Nos: 4,640,835, 4,496,689, 4,301,144, 4,670,417, 4,791,192 and 4,179,337. In certain embodiments, a derivative antigen binding protein comprises one or more polymer, including, but not limited to, monomethoxy-polyethylene glycol, dextran, cellulose, or other carbohydrate based polymers, poly-(N-vinyl pyrrolidone)-polyethylene glycol, propylene glycol homopolymers, a polypropylene oxide/ethylene oxide co-polymer, polyoxyethylated polyols (e.g., glycerol) and polyvinyl alcohol, as well as mixtures of such polymers.

[0197] In certain embodiments, a derivative is covalently modified with polyethylene glycol (PEG) subunits. In certain embodiments, one or more water-soluble polymer is bonded at one or more specific position, for example at the amino terminus, of a derivative. In certain embodiments, one or more water-soluble polymer is randomly attached to one or more side chains of a derivative. In certain embodiments, PEG is used to improve the therapeutic capacity for an antigen binding protein. In certain embodiments, PEG is used to improve the therapeutic capacity for a humanized antibody. Certain such methods are discussed, for example, in U.S. Patent No. 6,133,426, which is hereby incorporated by reference for any purpose.

[0198] Peptide analogs are commonly used in the pharmaceutical industry as non-peptide drugs with properties analogous to those of the template peptide. These types of non-peptide compound are termed "peptide mimetics" or "peptidomimetics." Fauchere, J., Adv. Drug Res., 15:29 (1986); Veber & Freidinger, TINS, p.392 (1985); and Evans *et al.*, J. Med. Chem., 30:1229 (1987), which are incorporated herein by reference for any purpose. Such compounds are often developed with the aid of computerized molecular modeling. Peptide mimetics that are structurally similar to therapeutically useful peptides can be used to produce a similar therapeutic or prophylactic effect. Generally, peptidomimetics are structurally similar to a paradigm polypeptide (i.e., a polypeptide that has a biochemical property or pharmacological activity), such as human antibody, but have one or more peptide linkages optionally replaced by

a linkage selected from: --CH_2 NH--, --CH_2 S--, --CH_2 -CH_2 --, --CH=CH-(cis and trans), --$COCH_2$ --, --CH(OH)CH_2 --, and --CH_2 SO--, by methods well known in the art. Systematic substitution of one or more amino acids of a consensus sequence with a D-amino acid of the same type (e.g., D-lysine in place of L-lysine) can be used in certain embodiments to generate more stable peptides. In addition, constrained peptides comprising a consensus sequence or a substantially identical consensus sequence variation can be generated by methods known in the art (Rizo and Gierasch, Ann. Rev. Biochem., 61:387 (1992), incorporated herein by reference for any purpose); for example, by adding internal cysteine residues capable of forming intramolecular disulfide bridges which cyclize the peptide.

[0199] The term "naturally occurring" as used throughout the specification in connection with biological materials such as polypeptides, nucleic acids, host cells, and the like, refers to materials which are found in nature or a form of the materials that is found in nature.

[0200] An "antigen binding protein" ("ABP") as used herein means any protein that binds a specified target antigen. In the instant application, the specified target antigen is the PCSK9 protein or fragment thereof. "Antigen binding protein" includes but is not limited to antibodies and binding parts thereof, such as immunologically functional fragments. Peptibodies are another example of antigen binding proteins. The term "immunologically functional fragment" (or simply "fragment") of an antibody or immunoglobulin chain (heavy or light chain) antigen binding protein, as used herein, is a species of antigen binding protein comprising a portion (regardless of how that portion is obtained or synthesized) of an antibody that lacks at least some of the amino acids present in a full-length chain but which is still capable of specifically binding to an antigen. Such fragments are biologically active in that they bind to the target antigen and can compete with other antigen binding proteins, including intact antibodies, for binding to a given epitope. In some embodiments, the fragments are neutralizing fragments. In some embodiments, the fragments can block or reduce the likelihood of the interaction between LDLR and PCSK9. In one aspect, such a fragment will retain at least one CDR present in the full-length light or heavy chain, and in some embodiments will comprise a single heavy chain and/or light chain or portion thereof. These biologically active fragments can be produced by recombinant DNA techniques, or can be produced by enzymatic or chemical cleavage of antigen binding proteins, including intact antibodies. Immunologically functional immunoglobulin fragments include, but are not limited to, Fab, a diabody (heavy chain variable

domain on the same polypeptide as a light chain variable domain, connected via a short peptide linker that is too short to permit pairing between the two domains on the same chain), Fab', $F(ab')_2$, Fv, domain antibodies and single-chain antibodies, and can be derived from any mammalian source, including but not limited to human, mouse, rat, camelid or rabbit. It is further contemplated that a functional portion of the antigen binding proteins disclosed herein, for example, one or more CDRs, could be covalently bound to a second protein or to a small molecule to create a therapeutic agent directed to a particular target in the body, possessing bifunctional therapeutic properties, or having a prolonged serum half-life. As will be appreciated by one of skill in the art, an antigen bindng protein can include nonprotein components. In some sections of the present disclosure, examples of ABPs are described herein in terms of "number/letter/number" (*e.g.*, 25A7). In these cases, the exact name denotes a specific antibody. That is, an ABP named 25A7 is not necessarily the same as an antibody named 25A7.1, (unless they are explicitly taught as the same in the specification, *e.g.*, 25A7 and 25A7.3). As will be appreciated by one of skill in the art, in some embodiments LDLR is not an antigen binding protein. In some embodiments, binding subsections of LDLR are not antigen binding proteins, *e.g.*, EGFa. In some embodiments, other molecules through which PCSK9 signals *in vivo* are not antigen binding proteins. Such embodiments will be explicitly identified as such.

[0201] Certain antigen binding proteins described herein are antibodies or are derived from antibodies. In certain embodiments, the polypeptide structure of the antigen binding proteins is based on antibodies, including, but not limited to, monoclonal antibodies, bispecific antibodies, minibodies, domain antibodies, synthetic antibodies (sometimes referred to herein as "antibody mimetics"), chimeric antibodies, humanized antibodies, human antibodies, antibody fusions (sometimes referred to herein as "antibody conjugates"), and fragments thereof, respectively. In some embodiments, the ABP comprises or consists of avimers (tightly binding peptide). These various antigen binding proteins are further described herein.

[0202] An "Fc" region comprises two heavy chain fragments comprising the C_H1 and C_H2 domains of an antibody. The two heavy chain fragments are held together by two or more disulfide bonds and by hydrophobic interactions of the C_H3 domains.

[0203] A "Fab fragment" comprises one light chain and the C_H1 and variable regions of one heavy chain. The heavy chain of a Fab molecule cannot form a disulfide bond with another heavy chain molecule.

[0204] A "Fab' fragment" comprises one light chain and a portion of one heavy chain that contains the VH domain and the C_H1 domain and also the region between the C_H1 and C_H2 domains, such that an interchain disulfide bond can be formed between the two heavy chains of two Fab' fragments to form an $F(ab')_2$ molecule.

[0205] A "$F(ab')_2$ fragment" contains two light chains and two heavy chains containing a portion of the constant region between the C_H1 and C_H2 domains, such that an interchain disulfide bond is formed between the two heavy chains. A $F(ab')_2$ fragment thus is composed of two Fab' fragments that are held together by a disulfide bond between the two heavy chains.

[0206] The "Fv region" comprises the variable regions from both the heavy and light chains, but lacks the constant regions.

[0207] "Single-chain antibodies" are Fv molecules in which the heavy and light chain variable regions have been connected by a flexible linker to form a single polypeptide chain, which forms an antigen binding region. Single chain antibodies are discussed in detail in International Patent Application Publication No. WO 88/01649 and United States Patent Nos. 4,946,778 and No. 5,260,203, the disclosures of which are incorporated by reference.

[0208] A "domain antibody" is an immunologically functional immunoglobulin fragment containing only the variable region of a heavy chain or the variable region of a light chain. In some instances, two or more V_H regions are covalently joined with a peptide linker to create a bivalent domain antibody. The two V_H regions of a bivalent domain antibody can target the same or different antigens.

[0209] A "bivalent antigen binding protein" or "bivalent antibody" comprises two antigen binding sites. In some instances, the two binding sites have the same antigen specificities. Bivalent antigen binding proteins and bivalent antibodies can be bispecific, *see, infra*. A bivalent antibody other than a "multispecific" or "multifunctional" antibody, in certain embodiments, typically is understood to have each of its binding sites identical.

[0210] A "multispecific antigen binding protein" or "multispecific antibody" is one that targets more than one antigen or epitope.

[0211] A "bispecific," "dual-specific" or "bifunctional" antigen binding protein or antibody is a hybrid antigen binding protein or antibody, respectively, having two different antigen binding sites. Bispecific antigen binding proteins and antibodies are a species of

multispecific antigen binding protein antibody and can be produced by a variety of methods including, but not limited to, fusion of hybridomas or linking of Fab' fragments. *See, e.g.*, Songsivilai and Lachmann, 1990, *Clin. Exp. Immunol.* 79:315-321; Kostelny *et al.*, 1992, *J. Immunol.* 148:1547-1553. The two binding sites of a bispecific antigen binding protein or antibody will bind to two different epitopes, which can reside on the same or different protein targets.

[0212] An antigen binding protein is said to "specifically bind" its target antigen when the dissociation constant (K_d) is $\leq 10^{-7}$ M. The ABP specifically binds antigen with "high affinity" when the K_d is $\leq 5 \times 10^{-9}$ M, and with "very high affinity" when the K_d is $\leq 5 \times 10^{-10}$ M. In one embodiment, the ABP has a K_d of $\leq 10^{-9}$ M. In one embodiment, the off-rate is $< 1 \times 10^{-5}$. In other embodiments, the ABPs will bind to human PCSK9 with a K_d of between about 10^{-9} M and 10^{-13} M, and in yet another embodiment the ABPs will bind with a $K_d \leq 5 \times 10^{-10}$. As will be appreciated by one of skill in the art, in some embodiments, any or all of the antigen binding fragments can specifically bind to PCSK9.

[0213] An antigen binding protein is "selective" when it binds to one target more tightly than it binds to a second target.

[0214] "Antigen binding region" means a protein, or a portion of a protein, that specifically binds a specified antigen (*e.g.*, a paratope). For example, that portion of an antigen binding protein that contains the amino acid residues that interact with an antigen and confer on the antigen binding protein its specificity and affinity for the antigen is referred to as "antigen binding region." An antigen binding region typically includes one or more "complementary binding regions" ("CDRs"). Certain antigen binding regions also include one or more "framework" regions. A "CDR" is an amino acid sequence that contributes to antigen binding specificity and affinity. "Framework" regions can aid in maintaining the proper conformation of the CDRs to promote binding between the antigen binding region and an antigen. Structurally, framework regions can be located in antibodies between CDRs. Examples of framework and CDR regions are shown in FIGs. 2A-3D, 3CCC-3JJJ, and 15A-15D. In some embodiments, the sequences for CDRs for the light chain of antibody 3B6 are as follows: CDR1 TLSSGYSSYEVD (SEQ ID NO: 279); CDR2 VDTGGIVGSKGE (SEQ ID NO: 280); CDR3 GADHGSGTNFVVV (SEQ ID NO: 281), and the FRs are as follows: FR1 QPVLTQPLFASASLGASVTLTC (SEQ ID NO: 282); FR2 WYQQRPGKGPRFVMR (SEQ

ID NO: 283); FR3 GIPDRFSVLGSGLNRYLTIKNIQEEDESDYHC (SEQ ID NO: 284); and FR4 FGGGTKLTVL (SEQ ID NO: 285).

[0215] In certain aspects, recombinant antigen binding proteins that bind PCSK9, for example human PCSK9, are provided. In this context, a "recombinant antigen binding protein" is a protein made using recombinant techniques, *i.e.*, through the expression of a recombinant nucleic acid as described herein. Methods and techniques for the production of recombinant proteins are well known in the art.

[0216] The term "antibody" refers to an intact immunoglobulin of any isotype, or a fragment thereof that can compete with the intact antibody for specific binding to the target antigen, and includes, for instance, chimeric, humanized, fully human, and bispecific antibodies. An "antibody" is a species of an antigen binding protein. An intact antibody will generally comprise at least two full-length heavy chains and two full-length light chains, but in some instances can include fewer chains such as antibodies naturally occurring in camelids which can comprise only heavy chains. Antibodies can be derived solely from a single source, or can be "chimeric," that is, different portions of the antibody can be derived from two different antibodies as described further below. The antigen binding proteins, antibodies, or binding fragments can be produced in hybridomas, by recombinant DNA techniques, or by enzymatic or chemical cleavage of intact antibodies. Unless otherwise indicated, the term "antibody" includes, in addition to antibodies comprising two full-length heavy chains and two full-length light chains, derivatives, variants, fragments, and muteins thereof, examples of which are described below. Furthermore, unless explicitly excluded, antibodies include monoclonal antibodies, bispecific antibodies, minibodies, domain antibodies, synthetic antibodies (sometimes referred to herein as "antibody mimetics"), chimeric antibodies, humanized antibodies, human antibodies, antibody fusions (sometimes referred to herein as "antibody conjugates"), and fragments thereof, respectively. In some embodiments, the term also encompasses peptibodies.

[0217] Naturally occurring antibody structural units typically comprise a tetramer. Each such tetramer typically is composed of two identical pairs of polypeptide chains, each pair having one full-length "light" (in certain embodiments, about 25 kDa) and one full-length "heavy" chain (in certain embodiments, about 50-70 kDa). The amino-terminal portion of each chain typically includes a variable region of about 100 to 110 or more amino acids that typically is responsible for antigen recognition. The carboxy-terminal portion of each chain typically

defines a constant region that can be responsible for effector function. Human light chains are typically classified as kappa and lambda light chains. Heavy chains are typically classified as mu, delta, gamma, alpha, or epsilon, and define the antibody's isotype as IgM, IgD, IgG, IgA, and IgE, respectively. IgG has several subclasses, including, but not limited to, IgG1, IgG2, IgG3, and IgG4. IgM has subclasses including, but not limited to, IgM1 and IgM2. IgA is similarly subdivided into subclasses including, but not limited to, IgA1 and IgA2. Within full-length light and heavy chains, typically, the variable and constant regions are joined by a "J" region of about 12 or more amino acids, with the heavy chain also including a "D" region of about 10 more amino acids. *See, e.g., Fundamental Immunology*, Ch. 7 (Paul, W., ed., 2nd ed. Raven Press, N.Y. (1989)) (incorporated by reference in its entirety for all purposes). The variable regions of each light/heavy chain pair typically form the antigen binding site.

[0218] The variable regions typically exhibit the same general structure of relatively conserved framework regions (FR) joined by three hyper variable regions, also called complementarity determining regions or CDRs. The CDRs from the two chains of each pair typically are aligned by the framework regions, which can enable binding to a specific epitope. From N-terminal to C-terminal, both light and heavy chain variable regions typically comprise the domains FR1, CDR1, FR2, CDR2, FR3, CDR3 and FR4. The assignment of amino acids to each domain is typically in accordance with the definitions of Kabat Sequences of Proteins of Immunological Interest (National Institutes of Health, Bethesda, Md. (1987 and 1991)), or Chothia & Lesk, J. Mol. Biol., 196:901-917 (1987); Chothia *et al.*, Nature, 342:878-883 (1989).

[0219] In certain embodiments, an antibody heavy chain binds to an antigen in the absence of an antibody light chain. In certain embodiments, an antibody light chain binds to an antigen in the absence of an antibody heavy chain. In certain embodiments, an antibody binding region binds to an antigen in the absence of an antibody light chain. In certain embodiments, an antibody binding region binds to an antigen in the absence of an antibody heavy chain. In certain embodiments, an individual variable region specifically binds to an antigen in the absence of other variable regions.

[0220] In certain embodiments, definitive delineation of a CDR and identification of residues comprising the binding site of an antibody is accomplished by solving the structure of the antibody and/or solving the structure of the antibody-ligand complex. In certain embodiments, that can be accomplished by any of a variety of techniques known to those skilled

in the art, such as X-ray crystallography. In certain embodiments, various methods of analysis can be employed to identify or approximate the CDR regions. Examples of such methods include, but are not limited to, the Kabat definition, the Chothia definition, the AbM definition and the contact definition.

[0221] The Kabat definition is a standard for numbering the residues in an antibody and is typically used to identify CDR regions. *See, e.g.*, Johnson & Wu, Nucleic Acids Res., 28: 214-8 (2000). The Chothia definition is similar to the Kabat definition, but the Chothia definition takes into account positions of certain structural loop regions. *See, e.g.*, Chothia *et al.*, J. Mol. Biol., 196: 901-17 (1986); Chothia *et al.*, Nature, 342: 877-83 (1989). The AbM definition uses an integrated suite of computer programs produced by Oxford Molecular Group that model antibody structure. *See, e.g.*, Martin *et al.*, Proc Natl Acad Sci (USA), 86:9268-9272 (1989); "AbM™, A Computer Program for Modeling Variable Regions of Antibodies," Oxford, UK; Oxford Molecular, Ltd. The AbM definition models the tertiary structure of an antibody from primary sequence using a combination of knowledge databases and *ab initio* methods, such as those described by Samudrala *et al.*, "*Ab Initio* Protein Structure Prediction Using a Combined Hierarchical Approach," in PROTEINS, Structure, Function and Genetics Suppl., 3:194-198 (1999). The contact definition is based on an analysis of the available complex crystal structures. *See, e.g.*, MacCallum *et al.*, J. Mol. Biol., 5:732-45 (1996).

[0222] By convention, the CDR regions in the heavy chain are typically referred to as H1, H2, and H3 and are numbered sequentially in the direction from the amino terminus to the carboxy terminus. The CDR regions in the light chain are typically referred to as L1, L2, and L3 and are numbered sequentially in the direction from the amino terminus to the carboxy terminus.

[0223] The term "light chain" includes a full-length light chain and fragments thereof having sufficient variable region sequence to confer binding specificity. A full-length light chain includes a variable region domain, V_L, and a constant region domain, C_L. The variable region domain of the light chain is at the amino-terminus of the polypeptide. Light chains include kappa chains and lambda chains.

[0224] The term "heavy chain" includes a full-length heavy chain and fragments thereof having sufficient variable region sequence to confer binding specificity. A full-length heavy chain includes a variable region domain, V_H, and three constant region domains, C_H1, C_H2, and C_H3. The V_H domain is at the amino-terminus of the polypeptide, and the C_H domains

are at the carboxyl-terminus, with the C_H3 being closest to the carboxy-terminus of the polypeptide. Heavy chains can be of any isotype, including IgG (including IgG1, IgG2, IgG3 and IgG4 subtypes), IgA (including IgA1 and IgA2 subtypes), IgM and IgE.

[0225] A bispecific or bifunctional antibody typically is an artificial hybrid antibody having two different heavy/light chain pairs and two different binding sites. Bispecific antibodies can be produced by a variety of methods including, but not limited to, fusion of hybridomas or linking of Fab' fragments. *See, e.g.,* Songsivilai *et al.*, Clin. Exp. Immunol., 79: 315-321 (1990); Kostelny *et al.*, J. Immunol., 148:1547-1553 (1992).

[0226] Some species of mammals also produce antibodies having only a single heavy chain.

[0227] Each individual immunoglobulin chain is typically composed of several "immunoglobulin domains," each consisting of roughly 90 to 110 amino acids and having a characteristic folding pattern. These domains are the basic units of which antibody polypeptides are composed. In humans, the IgA and IgD isotypes contain four heavy chains and four light chains; the IgG and IgE isotypes contain two heavy chains and two light chains; and the IgM isotype contains five heavy chains and five light chains. The heavy chain C region typically comprises one or more domains that can be responsible for effector function. The number of heavy chain constant region domains will depend on the isotype. IgG heavy chains, for example, contain three C region domains known as C_H1, C_H2 and C_H3. The antibodies that are provided can have any of these isotypes and subtypes. In certain embodiments of the present invention, an anti-PCSK9 antibody is of the IgG2 or IgG4 subtype.

[0228] The term "variable region" or "variable domain" refers to a portion of the light and/or heavy chains of an antibody, typically including approximately the amino-terminal 120 to 130 amino acids in the heavy chain and about 100 to 110 amino terminal amino acids in the light chain. In certain embodiments, variable regions of different antibodies differ extensively in amino acid sequence even among antibodies of the same species. The variable region of an antibody typically determines specificity of a particular antibody for its target

[0229] The term "neutralizing antigen binding protein" or "neutralizing antibody" refers to an antigen binding protein or antibody, respectively, that binds to a ligand and prevents or reduces the biological effect of that ligand. This can be done, for example, by directly blocking a binding site on the ligand or by binding to the ligand and altering the ligand's ability

to bind through indirect means (such as structural or energetic alterations in the ligand). In some embodiments, the term can also denote an antigen binding protein that prevents the protein to which it is bound from performing a biological function. In assessing the binding and/or specificity of an antigen binding protein, *e.g.*, an antibody or immunologically functional fragment thereof, an antibody or fragment can substantially inhibit binding of a ligand to its binding partner when an excess of antibody reduces the quantity of binding partner bound to the ligand by at least about 1-20, 20-30%, 30-40%, 40-50%, 50-60%, 60-70%, 70-80%, 80-85%, 85-90%, 90-95%, 95-97%, 97-98%, 98-99% or more (as measured in an *in vitro* competitive binding assay). In some embodiments, in the case of PCSK9 antigen binding proteins, such a neutralizing molecule can diminish the ability of PCSK9 to bind the LDLR. In some embodiments, the neutralizing ability is characterized and/or described via a competition assay. In some embodiments, the neutralizing ability is described in terms of an IC_{50} or EC_{50} value. In some embodiments, ABPs 27B2, 13H1, 13B5 and 3C4 are non-neutralizing ABPs, 3B6, 9C9 and 31A4 are weak neutralizers, and the remaining ABPs in Table 2 are strong neutralizers. In some embodiments, the antibodies or antigen binding proteins neutralize by binding to PCSK9 and preventing PCSK9 from binding to LDLR (or reducing the ability of PCSK9 to bind to LDLR). In some embodiments, the antibodies or ABPs neutralize by binding to PCSK9, and while still allowing PCSK9 to bind to LDLR, preventing or reducing the PCSK9 mediated degradation of LDLR. Thus, in some embodiments, a neutralizing ABP or antibody can still permit PCSK9/LDLR binding, but will prevent (or reduce) subsequent PCSK9 involved degradation of LDLR.

[0230] The term "target" refers to a molecule or a portion of a molecule capable of being bound by an antigen binding protein. In certain embodiments, a target can have one or more epitopes. In certain embodiments, a target is an antigen. The use of "antigen" in the phrase "antigen binding protein" simply denotes that the protein sequence that comprises the antigen can be bound by an antibody. In this context, it does not require that the protein be foreign or that it be capable of inducing an immune response.

[0231] The term "compete" when used in the context of antigen binding proteins (*e.g.*, neutralizing antigen binding proteins or neutralizing antibodies) that compete for the same epitope means competition between antigen binding proteins as determined by an assay in which the antigen binding protein (*e.g.*, antibody or immunologically functional fragment thereof)

being tested prevents or inhibits (*e.g.*, reduces) specific binding of a reference antigen binding protein (*e.g.*, a ligand, or a reference antibody) to a common antigen (*e.g.*, PCSK9 or a fragment thereof). Numerous types of competitive binding assays can be used to determine if one antigen binding protein competes with another, for example: solid phase direct or indirect radioimmunoassay (RIA), solid phase direct or indirect enzyme immunoassay (EIA), sandwich competition assay (*see, e.g.*, Stahli *et al.*, 1983, *Methods in Enzymology* 9:242-253); solid phase direct biotin-avidin EIA (*see, e.g.*, Kirkland *et al.*, 1986, *J. Immunol.* 137:3614-3619) solid phase direct labeled assay, solid phase direct labeled sandwich assay (*see, e.g.*, Harlow and Lane, 1988, *Antibodies, A Laboratory Manual*, Cold Spring Harbor Press); solid phase direct label RIA using I-125 label (*see, e.g.*, Morel *et al.*, 1988, *Molec. Immunol.* 25:7-15); solid phase direct biotin-avidin EIA (*see, e.g.*, Cheung, *et al.*, 1990, *Virology* 176:546-552); and direct labeled RIA (Moldenhauer *et al.*, 1990, *Scand. J. Immunol.* 32:77-82). Typically, such an assay involves the use of purified antigen bound to a solid surface or cells bearing either of these, an unlabelled test antigen binding protein and a labeled reference antigen binding protein. Competitive inhibition is measured by determining the amount of label bound to the solid surface or cells in the presence of the test antigen binding protein. Usually the test antigen binding protein is present in excess. Antigen binding proteins identified by competition assay (competing antigen binding proteins) include antigen binding proteins binding to the same epitope as the reference antigen binding proteins and antigen binding proteins binding to an adjacent epitope sufficiently proximal to the epitope bound by the reference antigen binding protein for steric hindrance to occur. Additional details regarding methods for determining competitive binding are provided in the examples herein. Usually, when a competing antigen binding protein is present in excess, it will inhibit (*e.g.*, reduce) specific binding of a reference antigen binding protein to a common antigen by at least 40-45%, 45-50%, 50-55%, 55-60%, 60-65%, 65-70%, 70-75% or 75% or more. In some instances, binding is inhibited by at least 80-85%, 85-90%, 90-95%, 95-97%, or 97% or more.

[0232] The term "antigen" refers to a molecule or a portion of a molecule capable of being bound by a selective binding agent, such as an antigen binding protein (including, *e.g.*, an antibody or immunological functional fragment thereof). In some embodiments, the antigen is capable of being used in an animal to produce antibodies capable of binding to that antigen. An

antigen can possess one or more epitopes that are capable of interacting with different antigen binding proteins, *e.g.*, antibodies.

[0233] The term "epitope" includes any determinant capable being bound by an antigen binding protein, such as an antibody or to a T-cell receptor. An epitope is a region of an antigen that is bound by an antigen binding protein that targets that antigen, and when the antigen is a protein, includes specific amino acids that directly contact the antigen binding protein. Most often, epitopes reside on proteins, but in some instances can reside on other kinds of molecules, such as nucleic acids. Epitope determinants can include chemically active surface groupings of molecules such as amino acids, sugar side chains, phosphoryl or sulfonyl groups, and can have specific three dimensional structural characteristics, and/or specific charge characteristics. Generally, antibodies specific for a particular target antigen will preferentially recognize an epitope on the target antigen in a complex mixture of proteins and/or macromolecules.

[0234] As used herein, "substantially pure" means that the described species of molecule is the predominant species present, that is, on a molar basis it is more abundant than any other individual species in the same mixture. In certain embodiments, a substantially pure molecule is a composition wherein the object species comprises at least 50% (on a molar basis) of all macromolecular species present. In other embodiments, a substantially pure composition will comprise at least 80%, 85%, 90%, 95%, or 99% of all macromolecular species present in the composition. In other embodiments, the object species is purified to essential homogeneity wherein contaminating species cannot be detected in the composition by conventional detection methods and thus the composition consists of a single detectable macromolecular species.

[0235] The term "agent" is used herein to denote a chemical compound, a mixture of chemical compounds, a biological macromolecule, or an extract made from biological materials.

[0236] As used herein, the terms "label" or "labeled" refers to incorporation of a detectable marker, e.g., by incorporation of a radiolabeled amino acid or attachment to a polypeptide of biotin moieties that can be detected by marked avidin (e.g., streptavidin containing a fluorescent marker or enzymatic activity that can be detected by optical or colorimetric methods). In certain embodiments, the label or marker can also be therapeutic. Various methods of labeling polypeptides and glycoproteins are known in the art and can be used. Examples of labels for polypeptides include, but are not limited to, the following:

radioisotopes or radionuclides (e.g., ^{3}H, ^{14}C, ^{15}N, ^{35}S, ^{90}Y, ^{99}Tc, ^{111}In, ^{125}I, ^{131}I), fluorescent labels (e.g., FITC, rhodamine, lanthanide phosphors), enzymatic labels (e.g., horseradish peroxidase, β-galactosidase, luciferase, alkaline phosphatase), chemiluminescent, biotinyl groups, predetermined polypeptide epitopes recognized by a secondary reporter (e.g., leucine zipper pair sequences, binding sites for secondary antibodies, metal binding domains, epitope tags). In certain embodiments, labels are attached by spacer arms of various lengths to reduce potential steric hindrance.

[0237] The term "biological sample", as used herein, includes, but is not limited to, any quantity of a substance from a living thing or formerly living thing. Such living things include, but are not limited to, humans, mice, monkeys, rats, rabbits, and other animals. Such substances include, but are not limited to, blood, serum, urine, cells, organs, tissues, bone, bone marrow, lymph nodes, and skin.

[0238] The term "pharmaceutical agent composition" (or agent or drug) as used herein refers to a chemical compound, composition, agent or drug capable of inducing a desired therapeutic effect when properly administered to a patient. It does not necessarily require more than one type of ingredient.

[0239] The term "therapeutically effective amount" refers to the amount of a PCSK9 antigen binding protein determined to produce a therapeutic response in a mammal. Such therapeutically effective amounts are readily ascertained by one of ordinary skill in the art.

[0240] The term "modulator," as used herein, is a compound that changes or alters the activity or function of a molecule. For example, a modulator can cause an increase or decrease in the magnitude of a certain activity or function of a molecule compared to the magnitude of the activity or function observed in the absence of the modulator. In certain embodiments, a modulator is an inhibitor, which decreases the magnitude of at least one activity or function of a molecule. Certain exemplary activities and functions of a molecule include, but are not limited to, binding affinity, enzymatic activity, and signal transduction. Certain exemplary inhibitors include, but are not limited to, proteins, peptides, antibodies, peptibodies, carbohydrates or small organic molecules. Peptibodies are described in, e.g., U.S. Patent No. 6,660,843 (corresponding to PCT Application No. WO 01/83525).

[0241] The terms "patient" and "subject" are used interchangeably and include human and non-human animal subjects as well as those with formally diagnosed disorders, those

without formally recognized disorders, those receiving medical attention, those at risk of developing the disorders, etc.

[0242] The term "treat" and "treatment" includes therapeutic treatments, prophylactic treatments, and applications in which one reduces the risk that a subject will develop a disorder or other risk factor. Treatment does not require the complete curing of a disorder and encompasses embodiments in which one reduces symptoms or underlying risk factors.

[0243] The term "prevent" does not require the 100% elimination of the possibility of an event. Rather, it denotes that the likelihood of the occurrence of the event has been reduced in the presence of the compound or method.

[0244] Standard techniques can be used for recombinant DNA, oligonucleotide synthesis, and tissue culture and transformation (e.g., electroporation, lipofection). Enzymatic reactions and purification techniques can be performed according to manufacturer's specifications or as commonly accomplished in the art or as described herein. The foregoing techniques and procedures can be generally performed according to conventional methods well known in the art and as described in various general and more specific references that are cited and discussed throughout the present specification. *See, e.g.,* Sambrook *et al., Molecular Cloning: A Laboratory Manual* (2d ed., Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y. (1989)), which is incorporated herein by reference for any purpose. Unless specific definitions are provided, the nomenclatures utilized in connection with, and the laboratory procedures and techniques of, analytical chemistry, synthetic organic chemistry, and medicinal and pharmaceutical chemistry described herein are those well known and commonly used in the art. Standard techniques can be used for chemical syntheses, chemical analyses, pharmaceutical preparation, formulation, and delivery, and treatment of patients.

Antigen Binding Proteins to PCSK9

[0245] Proprotein convertase subtilisin kexin type 9 (PCSK9) is a serine protease involved in regulating the levels of the low density lipoprotein receptor (LDLR) protein (Horton *et al.*, 2007; Seidah and Prat, 2007). PCSK9 is a prohormone-proprotein convertase in the subtilisin (S8) family of serine proteases (Seidah *et al.*, 2003). An exemplary human PCSK9 amino acid sequence is presented as SEQ ID NOs: 1 and 3. in FIG. 1A (depicting the "pro" domain of the protein as underlined) and FIG. 1B (depicting the signal sequence in bold and the

pro domain underlined). An exemplary human PCSK9 coding sequence is presented as SEQ ID NO: 2 (FIG. 1B). As described herein, PCSK9 proteins can also include fragments of the full length PCSK9 protein. The structure of the PCSK9 protein has recently been solved by two groups (Cunningham et al., Nature Structural & Molecular Biology, 2007, and Piper et al., Structure, 15:1-8, 2007), the entireties of both of which are herein incorporated by reference. PCSK9 includes a signal sequence, a N-terminal prodomain, a subtilisin-like catalytic domain and a C-terminal domain.

[0246] Antigen binding proteins (ABPs) that bind PCSK9, including human PCSK9, are provided herein. In some embodiments, the antigen binding proteins provided are polypeptides which comprise one or more complementary determining regions (CDRs), as described herein. In some antigen binding proteins, the CDRs are embedded into a "framework" region, which orients the CDR(s) such that the proper antigen binding properties of the CDR(s) is achieved. In some embodiments, antigen binding proteins provided herein can interfere with, block, reduce or modulate the interaction between PCSK9 and LDLR. Such antigen binding proteins are denoted as "neutralizing." In some embodiments, binding between PCSK9 and LDLR can still occur, even though the antigen binding protein is neutralizing and bound to PCSK9. For example, in some embodiments, the ABP prevents or reduces the adverse influence of PCSK9 on LDLR without blocking the LDLR binding site on PCSK9. Thus, in some embodiments, the ABP modulates or alters PCSK9's ability to result in the degradation of LDLR, without having to prevent the binding interaction between PCSK9 and LDLR. Such ABPs can be specifically described as "non-competitively neutralizing" ABPs. In some embodiments, the neutralizing ABP binds to PCSK9 in a location and/or manner that prevents PCSK9 from binding to LDLR. Such ABPs can be specifically described as "competitively neutralizing" ABPs. Both of the above neutralizers can result in a greater amount of free LDLR being present in a subject, which results in more LDLR binding to LDL (thereby reducing the amount of LDL in the subject). In turn, this results in a reduction in the amount of serum cholesterol present in a subject.

[0247] In some embodiments, the antigen binding proteins provided herein are capable of inhibiting PCSK9-mediated activity (including binding). In some embodiments, antigen binding proteins binding to these epitopes inhibit, *inter alia*, interactions between PCSK9

and LDLR and other physiological effects mediated by PCSK9. In some embodiments, the antigen binding proteins are human, such as fully human antibodies to PCSK9.

[0248] In some embodiments, the ABP binds to the catalytic domain of PCSK9. In some embodiments, the ABP binds to the mature form of PCSK9. In some embodiments the ABP binds in the prodomain of PCSK9. In some embodiments, the ABP selectively binds to the mature form of PCSK9. In some embodiments, the ABP binds to the catalytic domain in a manner such that PCSK9 cannot bind or bind as efficiently to LDLR. In some embodiments, the antigen binding protein does not bind to the c-terminus of the cataylytic domain. In some embodiments, the antigen binding protein does not bind to the n-terminus of the catalytic domain. In some embodiments, the ABP does not bind to the n- or c-terminus of the PCSK9 protein. In some embodiments, the ABP binds to any one of the epitopes bound by the antibodies discussed herein. In some embodiments, this can be determined by competition assays between the antibodies disclosed herein and other antibodes. In some embodiments, the ABP binds to an epitope bound by one of the antibodies described in Table 2. In some embodiments, the antigen binding proteins bind to a specific conformational state of PCSK9 so as to prevent PCSK9 from interacting with LDLR. In some embodiments, the ABP binds to the V domain of PCSK9. In some embodiments, the ABP binds to the V domain of PCSK9 and prevents (or reduces) PCSK9 from binding to LDLR. In some embodiments, the ABP binds to the V domain of PCSK9, and while it does not prevent (or reduce) the binding of PCSK9 to LDLR, the ABP prevents or reduces the adverse activities mediated through PCSK9 on LDLR.

[0249] The antigen binding proteins that are disclosed herein have a variety of utilities. Some of the antigen binding proteins, for instance, are useful in specific binding assays, affinity purification of PCSK9, in particular human PCSK9 or its ligands and in screening assays to identify other antagonists of PCSK9 activity. Some of the antigen binding proteins are useful for inhibiting binding of PCSK9 to LDLR, or inhibiting PCSK9-mediated activities.

[0250] The antigen binding proteins can be used in a variety of therapeutic applications, as explained herein. For example, in some embodiments the PCSK9 antigen binding proteins are useful for treating conditions associated with PCSK9, such as cholesterol related disorders (or "serum cholesterol related disorders") such as hypercholesterolemia, as further described herein. Other uses for the antigen binding proteins include, for example, diagnosis of PCSK9-associated diseases or conditions and screening assays to determine the

presence or absence of PCSK9. Some of the antigen binding proteins described herein are useful in treating consequences, symptoms, and/or the pathology associated with PCSK9 activity.

[0251] In some embodiments, the antigen binding proteins that are provided comprise one or more CDRs (*e.g.*, 1, 2, 3, 4, 5 or 6 CDRs). In some embodiments, the antigen binding protein comprises (a) a polypeptide structure and (b) one or more CDRs that are inserted into and/or joined to the polypeptide structure. The polypeptide structure can take a variety of different forms. For example, it can be, or comprise, the framework of a naturally occurring antibody, or fragment or variant thereof, or can be completely synthetic in nature. Examples of various polypeptide structures are further described below.

[0252] In certain embodiments, the polypeptide structure of the antigen binding proteins is an antibody or is derived from an antibody, including, but not limited to, monoclonal antibodies, bispecific antibodies, minibodies, domain antibodies, synthetic antibodies (sometimes referred to herein as "antibody mimetics"), chimeric antibodies, humanized antibodies, antibody fusions (sometimes referred to as "antibody conjugates"), and portions or fragments of each, respectively. In some instances, the antigen binding protein is an immunological fragment of an antibody (*e.g.*, a Fab, a Fab', a $F(ab')_2$, or a scFv). The various structures are further described and defined herein.

[0253] Certain of the antigen binding proteins as provided herein specifically and/or selectively bind to human PCSK9. In some embodiments, the antigen binding protein specifically and/or selectively binds to human PCSK9 protein having and/or consisting of residues 153-692 of SEQ ID NO: 3. In some embodiments the ABP specifically and/or selectively binds to human PCSK9 having and/or consiting of residues 31-152 of SEQ ID NO: 3. In some embodiments, the ABP selectively binds to a human PCSK9 protein as depicted in FIG. 1A (SEQ ID NO: 1). In some embodiments, the antigen binding protein specifically binds to at least a fragment of the PCSK9 protein and/or a full length PCSK9 protein, with or without a signal sequence.

[0254] In embodiments where the antigen binding protein is used for therapeutic applications, an antigen binding protein can inhibit, interfere with or modulate one or more biological activities of PCSK9. In one embodiment, an antigen binding protein binds specifically to human PCSK9 and/or substantially inhibits binding of human PCSK9 to LDLR by at least about 20%-40%, 40-60%, 60-80%, 80-85%, or more (for example, by measuring binding

in an *in vitro* competitive binding assay). Some of the antigen binding proteins that are provided herein are antibodies. In some embodiments, the ABP has a K_d of less (binding more tightly) than 10^{-7}, 10^{-8}, 10^{-9}, 10^{-10}, 10^{-11}, 10^{-12}, 10^{-13} M. In some embodiments, the ABP has an IC_{50} for blocking the binding of LDLR to PCSK9 (D374Y, high affinity variant) of less than 1 microM, 1000 nM to 100 nM, 100nM to 10 nM, 10nM to 1 nM, 1000pM to 500pM, 500 pM to 200 pM, less than 200 pM, 200 pM to 150 pM, 200 pM to 100 pM, 100 pM to 10 pM, 10 pM to 1 pM.

[0255] One example of an IgG2 heavy chain constant domain of an anti-PCSK9 antibody of the present invention has the amino acid sequence as shown in SEQ ID NO: 154, FIG. 3KK.

[0256] One example of an IgG4 heavy chain constant domain of an anti-PCSK9 antibody of the present invention has the amino acid sequence as shown in SEQ ID NO: 155, FIG. 3KK.

[0257] One example of a kappa light chain constant domain of an anti-PCSK9 antibody has the amino acid sequence as shown in SEQ ID NO: 157, FIG. 3KK.

[0258] One example of a lambda light chain constant domain of an anti-PCSK9 antibody has the amino acid sequence as shown in SEQ ID NO: 156, FIG. 3KK.

[0259] Variable regions of immunoglobulin chains generally exhibit the same overall structure, comprising relatively conserved framework regions (FR) joined by three hypervariable regions, more often called "complementarity determining regions" or CDRs. The CDRs from the two chains of each heavy chain/light chain pair mentioned above typically are aligned by the framework regions to form a structure that binds specifically with a specific epitope on the target protein (*e.g.*, PCSK9). From N-terminal to C-terminal, naturally-occurring light and heavy chain variable regions both typically conform with the following order of these elements: FR1, CDR1, FR2, CDR2, FR3, CDR3 and FR4. A numbering system has been devised for assigning numbers to amino acids that occupy positions in each of these domains. This numbering system is defined in Kabat Sequences of Proteins of Immunological Interest (1987 and 1991, NIH, Bethesda, MD), or Chothia & Lesk, 1987, *J. Mol. Biol.* 196:901-917; Chothia *et al.*, 1989, *Nature* 342:878-883.

[0260] Various heavy chain and light chain variable regions are provided herein and are depicted in FIGs. 2A-3JJ and 3LL-3BBB. In some embodiments, each of these variable regions can be attached to the above heavy and light chain constant regions to form a complete

antibody heavy and light chain, respectively. Further, each of the so generated heavy and light chain sequences can be combined to form a complete antibody structure.

[0261] Specific examples of some of the variable regions of the light and heavy chains of the antibodies that are provided and their corresponding amino acid sequences are summarized in TABLE 2.

TABLE 2

Exemplary Heavy and Light Chain Variable Regions

Antibody	Light/Heavy SEQ ID NO
30A4	5/74
3C4	7/85
23B5	9/71
25G4	10/72
31H4	12/67
27B2	13/87
25A7	15/58
27H5	16/52
26H5	17/51
31D1	18/53
20D10	19/48
27E7	20/54
30B9	21/55
19H9	22/56
26E10	23/49
21B12	23/49
17C2	24/57
23G1	26/50
13H1	28/91
9C9	30/64
9H6	31/62
31A4	32/89
1A12	33/65
16F12	35/79
22E2	36/80
27A6	37/76
28B12	38/77
28D6	39/78
31G11	40/83
13B5	42/69
31B12	44/81
3B6	46/60

[0262] Again, each of the exemplary variable heavy chains listed in Table 2 can be combined with any of the exemplary variable light chains shown in Table 2 to form an antibody. Table 2 shows exemplary light and heavy chain pairings found in several of the antibodies

disclosed herein. In some instances, the antibodies include at least one variable heavy chain and one variable light chain from those listed in Table 2. In other instances, the antibodies contain two identical light chains and two identical heavy chains. As an example, an antibody or antigen binding protein can include a heavy chain and a light chain, two heavy chains, or two light chains. In some embodiments the antigen binding protein comprises (and/or consists) of 1, 2, and/or 3 heavy and/or light CDRs from at least one of the sequences listed in Table 2 (CDRs for the sequences are outlined in FIGs. 2A-3D, and other embodiments in FIGs. 3CCC-3JJJ and 15A-15D). In some embodiments, all 6 CDRs (CDR1-3 from the light (CDRL1, CDRL2, CDRL3) and CDR1-3 from the heavy (CDRH1, CDRH2, and CDRH3)) are part of the ABP. In some embodiments, 1, 2, 3, 4, 5, or more CDRs are included in the ABP. In some embodiments, one heavy and one light CDR from the CDRs in the sequences in Table 2 is included in the ABP (CDRs for the sequences in table 2 are outlined in FIGs. 2A-3D). In some embodiments, additional sections (e.g., as depicted in FIG. 2A-2D, 3A-3D, and other embodiments in 3CCC-3JJJ and 15A-15D) are also included in the ABP. Examples of CDRs and FRs for the heavy and light chains noted in Table 2 are outlined in FIGs. 2A-3D (and other embodiments in FIGs. 3CCC-3JJJ and 15A-15D). Optional light chain variable sequences (including CDR1, CDR2, CDR3, FR1, FR2, FR3, and FR4) can be selected from the following: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46. Optional heavy chain variable sequences (including CDR1, CDR2, CDR3, FR1, FR2, FR3, and FR4) can be selected from the following: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60. In some of the entries in FIG. 2A-3D, variations of the sequences or alternative boundaries of the CDRs and FRs are identified. These alternatives are identified with a "v1" following the ABP name. As most of these alternatives are minor in nature, only sections with differences are displayed in the table. It is understood that the remaining section of the light or heavy chain is the same as shown for the base ABP in the other panels. Thus, for example, 19H9v1 in FIG. 2C has the same FR1, CDR1, and FR2 as 19H9 in FIG. 2A as the only difference is noted in FIG. 2C. For three of the nucleic acid sequences (ABPs 26E10, 30B9, and 31B12), additional alternative nucleic acid sequences are provided in the figures. As will be appreciated by one of skill in the art, no more than one such sequence need actually be used in the creation of an antibody or ABP. Indeed, in some

embodiments, only one or neither of the specific heavy or light chain nucleic acids need be present.

[0263] In some embodiments, the ABP is encoded by a nucleic acid sequence that can encode any of the protein sequences in Table 2.

[0264] In some embodiments, the ABP binds selectively to the form of PCSK9 that binds to LDLR (*e.g.*, the autocatalyzed form of the molecule). In some embodiments, the antigen binding protein does not bind to the c-terminus of the cataylytic domain (*e.g.*, the 5. 5-10, 10-15, 15-20, 20-25, 25-30, 30-40 most amino acids in the c-terminus). In some embodiments, the antigen binding protein does not bind to the n-terminus of the catalytic domain (*e.g.*, the 5. 5-10, 10-15, 15-20, 20-25, 25-30, 30-40 most amino acids in the n-terminus). In some embodiments, the ABP binds to amino acids within amino acids 1-100 of the mature form of PCSK9. In some embodiments, the ABP binds to amino acids within (and/or amino acid sequences consisting of) amino acids 31-100, 100-200, 31-152, 153-692, 200-300, 300-400, 452-683, 400-500, 500-600, 31-692, 31-449, and/or 600-692. In some embodiments, the ABP binds to the catalytic domain. In some embodiments, the neutralizing and/or non-neutralizing ABP binds to the prodomain. In some embodiments, the ABP binds to both the catalytic and pro domains. In some embodiments, the ABP binds to the catalytic domain so as to obstruct an area on the catalytic domain that interacts with the pro domain. In some embodiments, the ABP binds to the catalytic domain at a location or surface that the pro-domain interacts with as outlined in Piper et al. (Structure 15:1-8 (2007), the entirety of which is hereby incorporated by reference, including the structural representations therein). In some embodiments, the ABP binds to the catalytic domain and restricts the mobility of the prodomain. In some embodiments, the ABP binds to the catalytic domain without binding to the pro-domain. In some embodiments, the ABP binds to the catalytic domain, without binding to the pro-domain, while preventing the pro-domain from reorienting to allow PCSK9 to bind to LDLR. In some embodiments, the ABP binds in the same epitope as those surrounding residues 149-152 of the pro-domain in Piper et al. In some embodiments, the ABPs bind to the groove (as outlined in Piper et al.) on the V domain. In some embodiments, the ABPs bind to the histidine-rich patch proximal to the groove on the V domain. In some embodiments, such antibodies (that bind to the V domain) are not neutralizing. In some embodiments, antibodies that bind to the V domain are neutralizing. In some embodiments, the neutralizing ABPs prevent the binding of PCSK9 to LDLR. In some

embodiments, the neturalizing ABPs, while preventing the PCSK9 degradation of LDLR, do not prevent the binding of PCSK9 to LDLR (for example ABP 31A4). In some embodiments, the ABP binds to or blocks at least one of the histidines depicted in Figure 4 of the Piper et al. paper. In some embodiments, the ABP blocks the catalytic triad in PCSK9.

[0265] In some embodiments, the antibody binds selectively to variant PCSK9 proteins, *e.g.*, D374Y over wild type PCSK9. In some embodiments, these antibodies bind to the variant at least twice as strongly as the wild type, and preferably 2-5, 5-10, 10-100, 100-1000, 1000-10,000 fold or more to the mutant than the wild type (as measured via a K_d). In some embodiments, the antibody selectively inhibits variant D374Y PCSK9 from interacting with LDLR over wild type PCSK9's ability to interact with LDLR. In some embodiments, these antibodies block the variant's ability to bind to LDLR more strongly than the wild type's ability, e.g., at least twice as strongly as the wild type, and preferably 2-5, 5-10, 10-100, 100-1000 fold or more to the mutant than the wild type (as measured via an IC_{50}). In some embodiments, the antibody binds to and neutralizes both wild type PCSK9 and variant forms of PCSK9, such as D374Y at similar levels. In some embodiments, the antibody binds to PCSK9 to prevent variants of LDLR from binding to PCSK9. In some embodiments, the variants of LDLR are at least 50% identical to human LDLR. It is noted that variants of LDLR are known to those of skill in the art (*e.g.*, Brown MS et al, "Calcium cages, acid baths and recycling receptors" Nature 388: 629-630, 1997). In some embodiments, the ABP can raise the level of effective LDLR in heterozygote familial hypercholesterolemia (where a loss-of function variant of LDLR is present).

[0266] In some embodiments, the ABP binds to (but does not block) variants of PCSK9 that are at least 50%, 50-60, 60-70, 70-80, 80-90, 90-95, 95-99, or greater percent identity to the form of PCSK9 depicted in FIG. 1A and/or FIG. 1B. In some embodiments, the ABP binds to (but does not block) variants of PCSK9 that are at least 50%, 50-60, 60-70, 70-80, 80-90, 90-95, 95-99, or greater percent identity to the mature form of PCSK9 depicted in FIG. 1A and/or FIG. 1B. In some embodiments, the ABP binds to and prevents variants of PCSK9 that are at least 50%, 50-60, 60-70, 70-80, 80-90, 90-95, 95-99, or greater percent identity to the form of PCSK9 depicted in FIG. 1A and/or FIG. 1B from interacting with LDLR. In some embodiments, the ABP binds to and prevents variants of PCSK9 that are at least 50, 50-60, 60-70, 70-80, 80-90, 90-95, 95-99, or greater percent identity to the mature form of PCSK9 depicted in FIG. 1B from interacting with LDLR. In some embodiments, the variant of PCSK9 is a

human variant, such as variants at position 474, E620G, and/or E670G. In some embodiments, the amino acid at position 474 is valine (as in other humans) or threonine (as in cyno and mouse). Given the cross-reactivity data presented herein, it is believed that the present antibodies will readily bind to the above variants.

[0267] In some embodiments, the ABP binds to an epitope bound by one of the antibodies described in Table 2. In some embodiments, the antigen binding proteins bind to a specific conformational state of PCSK9 so as to prevent PCSK9 from interacting with LDLR.

Humanized Antigen Binding Proteins (*e.g.*, Antibodies)

[0268] As described herein, an antigen binding protein to PCSK9 can comprise a humanized antibody and/or part thereof. An important practical application of such a strategy is the "humanization" of the mouse humoral immune system.

[0269] In certain embodiments, a humanized antibody is substantially non-immunogenic in humans. In certain embodiments, a humanized antibody has substantially the same affinity for a target as an antibody from another species from which the humanized antibody is derived. *See, e.g.*, U.S. Patent 5,530,101, U.S. Patent 5,693,761; U.S. Patent 5,693,762; U.S. Patent 5,585,089.

[0270] In certain embodiments, amino acids of an antibody variable domain that can be modified without diminishing the native affinity of the antigen binding domain while reducing its immunogenicity are identified. *See, e.g.*, U.S. Patent Nos. 5,766,886 and 5,869,619.

[0271] In certain embodiments, modification of an antibody by methods known in the art is typically designed to achieve increased binding affinity for a target and/or to reduce immunogenicity of the antibody in the recipient. In certain embodiments, humanized antibodies are modified to eliminate glycosylation sites in order to increase affinity of the antibody for its cognate antigen. *See, e.g.*, Co *et al.*, Mol. Immunol., 30:1361-1367 (1993). In certain embodiments, techniques such as "reshaping," "hyperchimerization," or "veneering/resurfacing" are used to produce humanized antibodies. *See, e.g.*, Vaswami *et al.*, Annals of Allergy, Asthma, & Immunol. 81:105 (1998); Roguska *et al.*, Prot. Engineer., 9:895-904 (1996); and U.S. Patent No. 6,072,035. In certain such embodiments, such techniques typically reduce antibody immunogenicity by reducing the number of foreign residues, but do not prevent anti-idiotypic and anti-allotypic responses following repeated administration of the antibodies. Certain other

methods for reducing immunogenicity are described, e.g., in Gilliland *et al.*, J. Immunol ., 62(6): 3663-71 (1999).

[0272] In certain instances, humanizing antibodies results in a loss of antigen binding capacity. In certain embodiments, humanized antibodies are "back mutated." In certain such embodiments, the humanized antibody is mutated to include one or more of the amino acid residues found in the donor antibody. *See, e.g.,* Saldanha *et al.*, *Mol Immunol* 36:709-19 (1999).

[0273] In certain embodiments the complementarity determining regions (CDRs) of the light and heavy chain variable regions of an antibody to PCSK9 can be grafted to framework regions (FRs) from the same, or another, species. In certain embodiments, the CDRs of the light and heavy chain variable regions of an antibody to PCSK9 can be grafted to consensus human FRs. To create consensus human FRs, in certain embodiments, FRs from several human heavy chain or light chain amino acid sequences are aligned to identify a consensus amino acid sequence. In certain embodiments, the FRs of an antibody to PCSK9 heavy chain or light chain are replaced with the FRs from a different heavy chain or light chain. In certain embodiments, rare amino acids in the FRs of the heavy and light chains of an antibody to PCSK9 are not replaced, while the rest of the FR amino acids are replaced. Rare amino acids are specific amino acids that are in positions in which they are not usually found in FRs. In certain embodiments, the grafted variable regions from an antibody to PCSK9 can be used with a constant region that is different from the constant region of an antibody to PCSK9. In certain embodiments, the grafted variable regions are part of a single chain Fv antibody. CDR grafting is described, e.g., in U.S. Patent Nos. 6,180,370, 6,054,297, 5,693,762, 5,859,205, 5,693,761, 5,565,332, 5,585,089, and 5,530,101, and in Jones *et al.*, Nature, 321: 522-525 (1986); Riechmann *et al.*, Nature, 332: 323-327 (1988); Verhoeyen *et al.*, Science, 239:1534-1536 (1988), Winter, FEBS Letts., 430:92-94 (1998), which are hereby incorporated by reference for any purpose.

Human Antigen Binding Proteins *(e.g.,* Antibodies)

[0274] As described herein, an antigen binding protein that binds to PCSK9 can comprise a human (*i.e.*, fully human) antibody and/or part thereof. In certain embodiments, nucleotide sequences encoding, and amino acid sequences comprising, heavy and light chain immunoglobulin molecules, particularly sequences corresponding to the variable regions are provided. In certain embodiments, sequences corresponding to complementarity determining

regions (CDR's), specifically from CDR1 through CDR3, are provided. According to certain embodiments, a hybridoma cell line expressing such an immunoglobulin molecule is provided. According to certain embodiments, a hybridoma cell line expressing such a monoclonal antibody is provided. In certain embodiments a hybridoma cell line is selected from at least one of the cell lines described in Table 2, *e.g.*, 21B12, 16F12 and 31H4. In certain embodiments, a purified human monoclonal antibody to human PCSK9 is provided.

[0275] One can engineer mouse strains deficient in mouse antibody production with large fragments of the human Ig loci in anticipation that such mice would produce human antibodies in the absence of mouse antibodies. Large human Ig fragments can preserve the large variable gene diversity as well as the proper regulation of antibody production and expression. By exploiting the mouse machinery for antibody diversification and selection and the lack of immunological tolerance to human proteins, the reproduced human antibody repertoire in these mouse strains can yield high affinity fully human antibodies against any antigen of interest, including human antigens. Using the hybridoma technology, antigen-specific human MAbs with the desired specificity can be produced and selected. Certain exemplary methods are described in WO 98/24893, U.S. Patent No. 5,545,807, EP 546073, and EP 546073.

[0276] In certain embodiments, one can use constant regions from species other than human along with the human variable region(s).

[0277] The ability to clone and reconstruct megabase sized human loci in yeast artificial chromosomes (YACs) and to introduce them into the mouse germline provides an approach to elucidating the functional components of very large or crudely mapped loci as well as generating useful models of human disease. Furthermore, the utilization of such technology for substitution of mouse loci with their human equivalents could provide insights into the expression and regulation of human gene products during development, their communication with other systems, and their involvement in disease induction and progression.

[0278] Human antibodies avoid some of the problems associated with antibodies that possess murine or rat variable and/or constant regions. The presence of such murine or rat derived proteins can lead to the rapid clearance of the antibodies or can lead to the generation of an immune response against the antibody by a patient. In order to avoid the utilization of murine or rat derived antibodies, fully human antibodies can be generated through the introduction of

functional human antibody loci into a rodent, other mammal or animal so that the rodent, other mammal or animal produces fully human antibodies.

[0279] Humanized antibodies are those antibodies that, while initially starting off containing antibody amino acid sequences that are not human, have had at least some of these nonhuman antibody amino acid sequences replaced with human antibody sequences. This is in contrast with human antibodies, in which the antibody is encoded (or capable of being encoded) by genes possessed a human.

Antigen Binding Protein Variants

[0280] Other antibodies that are provided are variants of the ABPs listed above formed by combination or subparts of the variable heavy and variable light chains shown in Table 2 and comprise variable light and/or variable heavy chains that each have at least 50%, 50-60, 60-70, 70-80%, 80-85%, 85-90%, 90-95%, 95-97%, 97-99%, or above 99% identity to the amino acid sequences of the sequences in Table 2 (either the entire sequence or a subpart of the sequence, *e.g.*, one or more CDR). In some instances, such antibodies include at least one heavy chain and one light chain, whereas in other instances the variant forms contain two identical light chains and two identical heavy chains (or subparts thereof). In some embodiments, the sequence comparison in FIG. 2A-3D (and 13A-13J and other embodiments in 15A-15D) can be used in order to identify sections of the antibodies that can be modified by observing those variations that impact binding and those variations that do not appear to impact binding. For example, by comparing similar sequences, one can identify those sections (e.g., particular amino acids) that can be modified and how they can be modified while still retaining (or improving) the functionality of the ABP. In some embodiments, variants of ABPs include those consensus groups and sequences depicted in FIGs. 13A, 13C, 13F, 13G, 13H, 13I and/or 13J and variations are allowed in the positions identified as variable in the figures. The CDRs shown in FIGs. 13A, 13C, 13F, and 13G were defined based upon a hybrid combination of the Chothia method (based on the location of the structural loop regions, *see, e.g.*, "Standard conformations for the canonical structures of immunoglobulins," Bissan Al-Lazikani, Arthur M. Lesk and Cyrus Chothia, *Journal of Molecular Biology*, 273(4): 927-948, 7 November (1997)) and the Kabat method (based on sequence variability, *see, e.g., Sequences of Proteins of Immunological Interest*, Fifth

Edition. NIH Publication No. 91-3242, Kabat et al., (1991)). Each residue determined by either method, was included in the final list of CDR residues (and are presented in FIGs. 13A, 13C, 13F, and 13G). The CDRs in FIGs. 13H, 13I, and 13J were obtained by the Kabat method alone. Unless specified otherwise, the defined consensus sequences, CDRs, and FRs in FIGs. 13H-13J will define and control the noted CDRs and FRs for the referenced ABPs in FIG. 13.

[0281] In certain embodiments, an antigen binding protein comprises a heavy chain comprising a variable region comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from at least one of the sequences of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60. In certain embodiments, an antigen binding protein comprises a heavy chain comprising a variable region comprising an amino acid sequence at least 95% identical to an amino acid sequence selected from at least one of the sequences of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60. In certain embodiments, an antigen binding protein comprises a heavy chain comprising a variable region comprising an amino acid sequence at least 99% identical to an amino acid sequence selected from at least one of the sequences of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60.

[0282] In some embodiments, the antigen binding protein comprises a sequence that is at least 90%, 90-95%, and/or 95-99% identical to one or more CDRs from the CDRs in at least one of sequences of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60. In some embodiments, 1, 2, 3, 4, 5, or 6 CDR (each being at least 90%, 90-95%, and/or 95-99% identical to the above sequences) is present.

[0283] In some embodiments, the antigen binding protein comprises a sequence that is at least 90%, 90-95%, and/or 95-99% identical to one or more FRs from the FRs in at least one of sequences of SEQ ID NO: 74, 85, 71, 72, 67, 87, 58, 52, 51, 53, 48, 54, 55, 56, 49, 57, 50, 91, 64, 62, 89, 65, 79, 80, 76, 77, 78, 83, 69, 81, and 60. In some embodiments, 1, 2, 3, or 4 FR (each being at least 90%, 90-95%, and/or 95-99% identical to the above sequences) is present.

[0284] In certain embodiments, an antigen binding protein comprises a light chain comprising a variable region comprising an amino acid sequence at least 90% identical to an amino acid sequence selected from at least one of the sequences of SEQ ID NO: 5, 7, 9, 10, 12,

13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46. In certain embodiments, an antigen binding protein comprises a light chain comprising a variable region comprising an amino acid sequence at least 95% identical to an amino acid sequence selected from at least one of the sequences of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46. In certain embodiments, an antigen binding protein comprises a light chain comprising a variable region comprising an amino acid sequence at least 99% identical to an amino acid sequence selected from at least one of the sequences of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46.

[0285] In some embodiments, the antigen binding protein comprises a sequence that is at least 90%, 90-95%, and/or 95-99% identical to one or more CDRs from the CDRs in at least one of sequences of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46. In some embodiments, 1, 2, 3, 4, 5, or 6 CDR (each being at least 90%, 90-95%, and/or 95-99% identical to the above sequences) is present.

[0286] In some embodiments, the antigen binding protein comprises a sequence that is at least 90%, 90-95%, and/or 95-99% identical to one or more FRs from the FRs in at least one of sequences of SEQ ID NO: 5, 7, 9, 10, 12, 13, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 26, 28, 30, 31, 32, 33, 35, 36, 37, 38, 39, 40, 42, 44, and 46. In some embodiments, 1, 2, 3, or 4 FR (each being at least 90%, 90-95%, and/or 95-99% identical to the above sequences) is present.

[0287] In light of the present disclosure, a skilled artisan will be able to determine suitable variants of the ABPs as set forth herein using well-known techniques. In certain embodiments, one skilled in the art can identify suitable areas of the molecule that may be changed without destroying activity by targeting regions not believed to be important for activity. In certain embodiments, one can identify residues and portions of the molecules that are conserved among similar polypeptides. In certain embodiments, even areas that can be important for biological activity or for structure can be subject to conservative amino acid substitutions without destroying the biological activity or without adversely affecting the polypeptide structure.

[0288] Additionally, one skilled in the art can review structure-function studies identifying residues in similar polypeptides that are important for activity or structure. In view

of such a comparison, one can predict the importance of amino acid residues in a protein that correspond to amino acid residues which are important for activity or structure in similar proteins. One skilled in the art can opt for chemically similar amino acid substitutions for such predicted important amino acid residues.

[0289] One skilled in the art can also analyze the three-dimensional structure and amino acid sequence in relation to that structure in similar ABPs. In view of such information, one skilled in the art can predict the alignment of amino acid residues of an antibody with respect to its three dimensional structure. In certain embodiments, one skilled in the art can choose not to make radical changes to amino acid residues predicted to be on the surface of the protein, since such residues can be involved in important interactions with other molecules. Moreover, one skilled in the art can generate test variants containing a single amino acid substitution at each desired amino acid residue. The variants can then be screened using activity assays known to those skilled in the art. Such variants can be used to gather information about suitable variants. For example, if one discovered that a change to a particular amino acid residue resulted in destroyed, undesirably reduced, or unsuitable activity, variants with such a change can be avoided. In other words, based on information gathered from such routine experiments, one skilled in the art can readily determine the amino acids where further substitutions should be avoided either alone or in combination with other mutations.

[0290] A number of scientific publications have been devoted to the prediction of secondary structure. See Moult J., Curr. Op. in Biotech., 7(4):422-427 (1996), Chou *et al.*, Biochemistry, 13(2):222-245 (1974); Chou *et al.*, Biochemistry, 113(2):211-222 (1974); Chou *et al.*, Adv. Enzymol. Relat. Areas Mol. Biol., 47:45-148 (1978); Chou *et al.*, Ann. Rev. Biochem., 47:251-276 and Chou *et al.*, Biophys. J., 26:367-384 (1979). Moreover, computer programs are currently available to assist with predicting secondary structure. One method of predicting secondary structure is based upon homology modeling. For example, two polypeptides or proteins which have a sequence identity of greater than 30%, or similarity greater than 40% often have similar structural topologies. The recent growth of the protein structural database (PDB) has provided enhanced predictability of secondary structure, including the potential number of folds within a polypeptide's or protein's structure. *See* Holm *et al.*, Nucl. Acid. Res., 27(1):244-247 (1999). It has been suggested (Brenner *et al.*, Curr. Op. Struct. Biol., 7(3):369-376 (1997)) that there are a limited number of folds in a given polypeptide or protein and that once a critical

number of structures have been resolved, structural prediction will become dramatically more accurate.

[0291] Additional methods of predicting secondary structure include "threading" (Jones, D., Curr. Opin. Struct. Biol., 7(3):377-87 (1997); Sippl *et al.*, Structure, 4(1):15-19 (1996)), "profile analysis" (Bowie *et al.*, Science, 253:164-170 (1991); Gribskov *et al.*, Meth. Enzym., 183:146-159 (1990); Gribskov *et al.*, Proc. Nat. Acad. Sci. USA, 84(13):4355-4358 (1987)), and "evolutionary linkage" (See Holm, *supra* (1999), and Brenner, *supra* (1997)).

[0292] In certain embodiments, antigen binding protein variants include glycosylation variants wherein the number and/or type of glycosylation site has been altered compared to the amino acid sequences of a parent polypeptide. In certain embodiments, protein variants comprise a greater or a lesser number of N-linked glycosylation sites than the native protein. An N-linked glycosylation site is characterized by the sequence: Asn-X-Ser or Asn-X-Thr, wherein the amino acid residue designated as X can be any amino acid residue except proline. The substitution of amino acid residues to create this sequence provides a potential new site for the addition of an N-linked carbohydrate chain. Alternatively, substitutions which eliminate this sequence will remove an existing N-linked carbohydrate chain. Also provided is a rearrangement of N-linked carbohydrate chains wherein one or more N-linked glycosylation sites (typically those that are naturally occurring) are eliminated and one or more new N-linked sites are created. Additional preferred antibody variants include cysteine variants wherein one or more cysteine residues are deleted from or substituted for another amino acid (e.g., serine) as compared to the parent amino acid sequence. Cysteine variants can be useful when antibodies must be refolded into a biologically active conformation such as after the isolation of insoluble inclusion bodies. Cysteine variants generally have fewer cysteine residues than the native protein, and typically have an even number to minimize interactions resulting from unpaired cysteines.

[0293] According to certain embodiments, amino acid substitutions are those which: (1) reduce susceptibility to proteolysis, (2) reduce susceptibility to oxidation, (3) alter binding affinity for forming protein complexes, (4) alter binding affinities, and/or (4) confer or modify other physicochemical or functional properties on such polypeptides. According to certain embodiments, single or multiple amino acid substitutions (in certain embodiments, conservative amino acid substitutions) can be made in the naturally-occurring sequence (in certain

embodiments, in the portion of the polypeptide outside the domain(s) forming intermolecular contacts). In certain embodiments, a conservative amino acid substitution typically may not substantially change the structural characteristics of the parent sequence (e.g., a replacement amino acid should not tend to break a helix that occurs in the parent sequence, or disrupt other types of secondary structure that characterizes the parent sequence). Examples of art-recognized polypeptide secondary and tertiary structures are described in *Proteins, Structures and Molecular Principles* (Creighton, Ed., W. H. Freeman and Company, New York (1984)); *Introduction to Protein Structure* (C. Branden & J. Tooze, eds., Garland Publishing, New York, N.Y. (1991)); and Thornton *et al.*, Nature, 354:105 (1991), which are each incorporated herein by reference.

[0294] In some embodiments, the variants are variants of the nucleic acid sequences of the ABPs disclosed herein. One of skill in the art will appreciate that the above discussion can be used for identifying, evaluating, and/creating ABP protein variants and also for nucleic acid sequences that can encode for those protein variants. Thus, nucleic acid sequences encoding for those protein variants (as well as nucleic acid sequences that encode for the ABPs in Table 2, but are different from those explicitly disclosed herein) are contemplated. For example, an ABP variant can have at least 80, 80-85, 85-90, 90-95, 95-97, 97-99 or greater identity to at least one nucleic acid sequence described in SEQ ID NOs: 152, 153, 92, 93, 94, 95, 96, 97, 98, 99, 100, 101, 102, 103, 104, 105, 106, 107, 108, 109, 110, 111, 112, 113, 114, 115, 116, 117, 118, 119, 120, 121, 122, 123, 124, 125, 126, 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138, 139, 140, 141, 142, 143, 144, 145, 146, 147, 148, 149, 150, 151 or at least one to six (and various combinations thereof) of the CDR(s) encoded by the nucleic acid sequences in SEQ ID NOs: 152, 153, 92, 93, 94, 95, 96, 97, 98, 99, 100, 101, 102, 103, 104, 105, 106, 107, 108, 109, 110, 111, 112, 113, 114, 115, 116, 117, 118, 119, 120, 121, 122, 123, 124, 125, 126, 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138, 139, 140, 141, 142, 143, 144, 145, 146, 147, 148, 149, 150, and 151.

[0295] In some embodiments, the antibody (or nucleic acid sequence encoding it) is a variant if the nucleic acid sequence that encodes the particular ABP (or the nucleic acid sequence itself) can selectively hybridize to any of the nucleic acid sequences that encode the proteins in Table 2 (such as, but not limited to SEQ ID NO: 152, 153, 92, 93, 94, 95, 96, 97, 98, 99, 100, 101, 102, 103, 104, 105, 106, 107, 108, 109, 110, 111, 112, 113, 114, 115, 116, 117, 118, 119, 120, 121, 122, 123, 124, 125, 126, 127, 128, 129, 130, 131, 132, 133, 134, 135, 136, 137, 138,

139, 140, 141, 142, 143, 144, 145, 146, 147, 148, 149, 150, and 151) under stringent conditions. In one embodiment, suitable moderately stringent conditions include prewashing in a solution of 5XSSC; 0.5% SDS, 1.0 mM EDTA (pH 8:0); hybridizing at 50° C, -65° C, 5xSSC, overnight or, in the event of cross-species homology, at 45° C with 0.5xSSC; followed by washing twice at 65° C for 20 minutes with each of 2x, 0.5x and 0.2xSSC containing 0.1% SDS. Such hybridizing DNA sequences are also within the scope of this invention, as are nucleotide sequences that, due to code degeneracy, encode an antibody polypeptide that is encoded by a hybridizing DNA sequence and the amino acid sequences that are encoded by these nucleic acid sequences. In some embodiments, variants of CDRs include nucleic acid sequences and the amino acid sequences encoded by those sequences, that hybridize to one or more of the CDRs within the sequences noted above (individual CDRs can readily be determined in light of FIGs. 2A-3D, and other embodiments in FIGs. 3CCC-3JJJ and 15A-15D). The phrase "selectively hybridize" referred to in this context means to detectably and selectively bind. Polynucleotides, oligonucleotides and fragments thereof in accordance with the invention selectively hybridize to nucleic acid strands under hybridization and wash conditions that minimize appreciable amounts of detectable binding to nonspecific nucleic acids. High stringency conditions can be used to achieve selective hybridization conditions as known in the art and discussed herein. Generally, the nucleic acid sequence homology between the polynucleotides, oligonucleotides, and fragments of the invention and a nucleic acid sequence of interest will be at least 80%, and more typically with preferably increasing homologies of at least 85%, 90%, 95%, 99%, and 100%. Two amino acid sequences are homologous if there is a partial or complete identity between their sequences. For example, 85% homology means that 85% of the amino acids are identical when the two sequences are aligned for maximum matching. Gaps (in either of the two sequences being matched) are allowed in maximizing matching; gap lengths of 5 or less are preferred with 2 or less being more preferred. Alternatively and preferably, two protein sequences (or polypeptide sequences derived from them of at least 30 amino acids in length) are homologous, as this term is used herein, if they have an alignment score of at more than 5 (in standard deviation units) using the program ALIGN with the mutation data matrix and a gap penalty of 6 or greater. See Dayhoff, M. O., in Atlas of Protein Sequence and Structure, pp. 101-110 (Volume 5, National Biomedical Research Foundation (1972)) and Supplement 2 to this volume, pp. 1-10. The two sequences or parts thereof are more preferably homologous if their amino acids are

greater than or equal to 50% identical when optimally aligned using the ALIGN program. The term "corresponds to" is used herein to mean that a polynucleotide sequence is homologous (i.e., is identical, not strictly evolutionarily related) to all or a portion of a reference polynucleotide sequence, or that a polypeptide sequence is identical to a reference polypeptide sequence. In contradistinction, the term "complementary to" is used herein to mean that the complementary sequence is homologous to all or a portion of a reference polynucleotide sequence. For illustration, the nucleotide sequence "TATAC" corresponds to a reference sequence "TATAC" and is complementary to a reference sequence "GTATA".

Preparation of Antigen Binding Proteins (*e.g.*, Antibodies)

[0296] In certain embodiments, antigen binding proteins (such as antibodies) are produced by immunization with an antigen (*e.g.*, PCSK9). In certain embodiments, antibodies can be produced by immunization with full-length PCSK9, a soluble form of PCSK9, the catalytic domain alone, the mature form of PCSK9 shown in FIG. 1A, a splice variant form of PCSK9, or a fragment thereof. In certain embodiments, the antibodies of the invention can be polyclonal or monoclonal, and/or can be recombinant antibodies. In certain embodiments, antibodies of the invention are human antibodies prepared, for example, by immunization of transgenic animals capable of producing human antibodies (see, for example, PCT Published Application No. W0 93/12227).

[0297] In certain embodiments, certain strategies can be employed to manipulate inherent properties of an antibody, such as the affinity of an antibody for its target. Such strategies include, but are not limited to, the use of site-specific or random mutagenesis of the polynucleotide molecule encoding an antibody to generate an antibody variant. In certain embodiments, such generation is followed by screening for antibody variants that exhibit the desired change, e.g. increased or decreased affinity.

[0298] In certain embodiments, the amino acid residues targeted in mutagenic strategies are those in the CDRs. In certain embodiments, amino acids in the framework regions of the variable domains are targeted. In certain embodiments, such framework regions have been shown to contribute to the target binding properties of certain antibodies. *See, e.g.*, Hudson, Curr. Opin. Biotech., 9:395-402 (1999) and references therein.

[0299] In certain embodiments, smaller and more effectively screened libraries of antibody variants are produced by restricting random or site-directed mutagenesis to hyper-mutation sites in the CDRs, which are sites that correspond to areas prone to mutation during the somatic affinity maturation process. *See, e.g.*, Chowdhury & Pastan, Nature Biotech., 17: 568-572 (1999) and references therein. In certain embodiments, certain types of DNA elements can be used to identify hyper-mutation sites including, but not limited to, certain direct and inverted repeats, certain consensus sequences, certain secondary structures, and certain palindromes. For example, such DNA elements that can be used to identify hyper-mutation sites include, but are not limited to, a tetrabase sequence comprising a purine (A or G), followed by guainine (G), followed by a pyrimidine (C or T), followed by either adenosine or thymidine (A or T) (i.e., A/G-G-C/T-A/T). Another example of a DNA element that can be used to identify hyper-mutation sites is the serine codon, A-G-C/T.

Preparation of Fully Human ABPs (*e.g.*, Antibodies)

[0300] In certain embodiments, a phage display technique is used to generate monoclonal antibodies. In certain embodiments, such techniques produce fully human monoclonal antibodies. In certain embodiments, a polynucleotide encoding a single Fab or Fv antibody fragment is expressed on the surface of a phage particle. *See, e.g.*, Hoogenboom *et al.*, J. Mol. Biol., 227: 381 (1991); Marks *et al.*, *J Mol Biol* 222: 581 (1991); U.S. Patent No. 5,885,793. In certain embodiments, phage are "screened" to identify those antibody fragments having affinity for target. Thus, certain such processes mimic immune selection through the display of antibody fragment repertoires on the surface of filamentous bacteriophage, and subsequent selection of phage by their binding to target. In certain such procedures, high affinity functional neutralizing antibody fragments are isolated. In certain such embodiments (discussed in more detail below), a complete repertoire of human antibody genes is created by cloning naturally rearranged human V genes from peripheral blood lymphocytes. *See, e.g.*, Mullinax *et al.*, Proc Natl Acad Sci (USA), 87: 8095-8099 (1990).

[0301] According to certain embodiments, antibodies of the invention are prepared through the utilization of a transgenic mouse that has a substantial portion of the human antibody producing genome inserted but that is rendered deficient in the production of endogenous, murine antibodies. Such mice, then, are capable of producing human immunoglobulin molecules

and antibodies and are deficient in the production of murine immunoglobulin molecules and antibodies. Technologies utilized for achieving this result are disclosed in the patents, applications and references disclosed in the specification, herein. In certain embodiments, one can employ methods such as those disclosed in PCT Published Application No. WO 98/24893 or in Mendez *et al.*, Nature Genetics, 15:146-156 (1997), which are hereby incorporated by reference for any purpose.

[0302] Generally, fully human monoclonal ABPs (*e.g.*, antibodies) specific for PCSK9 can be produced as follows. Transgenic mice containing human immunoglobulin genes are immunized with the antigen of interest, *e.g.* PCSK9, lymphatic cells (such as B-cells) from the mice that express antibodies are obtained. Such recovered cells are fused with a myeloid-type cell line to prepare immortal hybridoma cell lines, and such hybridoma cell lines are screened and selected to identify hybridoma cell lines that produce antibodies specific to the antigen of interest. In certain embodiments, the production of a hybridoma cell line that produces antibodies specific to PCSK9 is provided.

[0303] In certain embodiments, fully human antibodies are produced by exposing human splenocytes (B or T cells) to an antigen *in vitro*, and then reconstituting the exposed cells in an immunocompromised mouse, *e.g.* SCID or nod/SCID. *See, e.g.*, Brams *et al.*, J.Immunol. 160: 2051-2058 (1998); Carballido *et al.*, Nat. Med., 6: 103-106 (2000). In certain such approaches, engraftment of human fetal tissue into SCID mice (SCID-hu) results in long-term hematopoiesis and human T-cell development. *See, e.g.*, McCune *et al.*, Science, 241:1532-1639 (1988); Ifversen *et al.*, Sem. Immunol., 8:243-248 (1996). In certain instances, humoral immune response in such chimeric mice is dependent on co-development of human T-cells in the animals. *See, e.g.*, Martensson *et al.*, Immunol., 83:1271-179 (1994). In certain approaches, human peripheral blood lymphocytes are transplanted into SCID mice. *See, e.g.*, Mosier *et al.*, Nature, 335:256-259 (1988). In certain such embodiments, when such transplanted cells are treated either with a priming agent, such as Staphylococcal Enterotoxin A (SEA), or with anti-human CD40 monoclonal antibodies, higher levels of B cell production is detected. *See, e.g.*, Martensson *et al.*, Immunol., 84: 224-230 (1995); Murphy *et al.*, Blood, 86:1946-1953 (1995).

[0304] Thus, in certain embodiments, fully human antibodies can be produced by the expression of recombinant DNA in host cells or by expression in hybridoma cells. In other embodiments, antibodies can be produced using the phage display techniques described herein.

[0305] The antibodies described herein were prepared through the utilization of the XenoMouse® technology, as described herein. Such mice, then, are capable of producing human immunoglobulin molecules and antibodies and are deficient in the production of murine immunoglobulin molecules and antibodies. Technologies utilized for achieving the same are disclosed in the patents, applications, and references disclosed in the background section herein. In particular, however, a preferred embodiment of transgenic production of mice and antibodies therefrom is disclosed in U.S. Patent Application Serial No. 08/759,620, filed December 3, 1996 and International Patent Application Nos. WO 98/24893, published June 11, 1998 and WO 00/76310, published December 21, 2000, the disclosures of which are hereby incorporated by reference. *See also* Mendez *et al.*, Nature Genetics, 15:146-156 (1997), the disclosure of which is hereby incorporated by reference.

[0306] Through the use of such technology, fully human monoclonal antibodies to a variety of antigens have been produced. Essentially, XenoMouse® lines of mice are immunized with an antigen of interest (*e.g.* PCSK9), lymphatic cells (such as B-cells) are recovered from the hyper-immunized mice, and the recovered lymphocytes are fused with a myeloid-type cell line to prepare immortal hybridoma cell lines. These hybridoma cell lines are screened and selected to identify hybridoma cell lines that produced antibodies specific to the antigen of interest. Provided herein are methods for the production of multiple hybridoma cell lines that produce antibodies specific to PCSK9 Further, provided herein are characterization of the antibodies produced by such cell lines, including nucleotide and amino acid sequence analyses of the heavy and light chains of such antibodies.

[0307] The production of the XenoMouse® strains of mice is further discussed and delineated in U.S. Patent Application Serial Nos. 07/466,008, filed January 12, 1990, 07/610,515, filed November 8, 1990, 07/919,297, filed July 24, 1992, 07/922,649, filed July 30, 1992, 08/031,801, filed March 15, 1993, 08/112,848, filed August 27, 1993, 08/234,145, filed April 28, 1994, 08/376,279, filed January 20, 1995, 08/430, 938, filed April 27, 1995, 08/464,584, filed June 5, 1995, 08/464,582, filed June 5, 1995, 08/463,191, filed June 5, 1995, 08/462,837, filed June 5, 1995, 08/486,853, filed June 5, 1995, 08/486,857, filed June 5, 1995, 08/486,859, filed June 5, 1995, 08/462,513, filed June 5, 1995, 08/724,752, filed October 2, 1996, 08/759,620, filed December 3, 1996, U.S. Publication 2003/0093820, filed November 30, 2001 and U.S. Patent Nos. 6,162,963, 6,150,584, 6,114,598, 6,075,181, and 5,939,598 and

Japanese Patent Nos. 3 068 180 B2, 3 068 506 B2, and 3 068 507 B2. *See also* European Patent No., EP 0 463 151 B1, grant published June 12, 1996, International Patent Application No., WO 94/02602, published February 3, 1994, International Patent Application No., WO 96/34096, published October 31, 1996, WO 98/24893, published June 11, 1998, WO 00/76310, published December 21, 2000. The disclosures of each of the above-cited patents, applications, and references are hereby incorporated by reference in their entirety.

[0308] In an alternative approach, others, including GenPharm International, Inc., have utilized a "minilocus" approach. In the minilocus approach, an exogenous Ig locus is mimicked through the inclusion of pieces (individual genes) from the Ig locus. Thus, one or more V_H genes, one or more D_H genes, one or more J_H genes, a mu constant region, and usually a second constant region (preferably a gamma constant region) are formed into a construct for insertion into an animal. This approach is described in U.S. Patent No. 5,545,807 to Surani *et al.* and U.S. Patent Nos. 5,545,806, 5,625,825, 5,625,126, 5,633,425, 5,661,016, 5,770,429, 5,789,650, 5,814,318, 5,877,397, 5,874,299, and 6,255,458 each to Lonberg & Kay, U.S. Patent No. 5,591,669 and 6,023.010 to Krimpenfort & Berns, U.S. Patent Nos. 5,612,205, 5,721,367, and 5,789,215 to Berns *et al.*, and U.S. Patent No. 5,643,763 to Choi & Dunn, and GenPharm International U.S. Patent Application Serial Nos. 07/574,748, filed August 29, 1990, 07/575,962, filed August 31, 1990, 07/810,279, filed December 17, 1991, 07/853,408, filed March 18, 1992, 07/904,068, filed June 23, 1992, 07/990,860, filed December 16, 1992, 08/053,131, filed April 26, 1993, 08/096,762, filed July 22, 1993, 08/155,301, filed November 18, 1993, 08/161,739, filed December 3, 1993, 08/165,699, filed December 10, 1993, 08/209,741, filed March 9, 1994, the disclosures of which are hereby incorporated by reference. *See also* European Patent No. 0 546 073 B1, International Patent Application Nos. WO 92/03918, WO 92/22645, WO 92/22647, WO 92/22670, WO 93/12227, WO 94/00569, WO 94/25585, WO 96/14436, WO 97/13852, and WO 98/24884 and U.S. Patent No. 5,981,175, the disclosures of which are hereby incorporated by reference in their entirety. *See further* Taylor *et al.*, 1992, Chen *et al.*, 1993, Tuaillon *et al.*, 1993, Choi *et al.*, 1993, Lonberg *et al.*, (1994), Taylor *et al.*, (1994), and Tuaillon *et al.*, (1995), Fishwild *et al.*, (1996), the disclosures of which are hereby incorporated by reference in their entirety.

[0309] Kirin has also demonstrated the generation of human antibodies from mice in which, through microcell fusion, large pieces of chromosomes, or entire chromosomes, have

been introduced. *See* European Patent Application Nos. 773 288 and 843 961, the disclosures of which are hereby incorporated by reference. Additionally, KM™ mice, which are the result of cross-breeding of Kirin's Tc mice with Medarex's minilocus (Humab) mice have been generated. These mice possess the human IgH transchromosome of the Kirin mice and the kappa chain transgene of the Genpharm mice (Ishida *et al.*, Cloning Stem Cells, (2002) 4:91-102).

[0310] Human antibodies can also be derived by *in vitro* methods. Suitable examples include but are not limited to phage display (CAT, Morphosys, Dyax, Biosite/Medarex, Xoma, Symphogen, Alexion (formerly Proliferon), Affimed) ribosome display (CAT), yeast display, and the like.

[0311] In some embodiments, the antibodies described herein possess human IgG4 heavy chains as well as IgG2 heavy chains. Antibodies can also be of other human isotypes, including IgG1. The antibodies possessed high affinities, typically possessing a Kd of from about 10^{-6} through about 10^{-13} M or below, when measured by various techniques.

[0312] As will be appreciated, antibodies can be expressed in cell lines other than hybridoma cell lines. Sequences encoding particular antibodies can be used to transform a suitable mammalian host cell. Transformation can be by any known method for introducing polynucleotides into a host cell, including, for example packaging the polynucleotide in a virus (or into a viral vector) and transducing a host cell with the virus (or vector) or by transfection procedures known in the art, as exemplified by U.S. Patent Nos. 4,399,216, 4,912,040, 4,740,461, and 4,959,455 (which patents are hereby incorporated herein by reference). The transformation procedure used depends upon the host to be transformed. Methods for introducing heterologous polynucleotides into mammalian cells are well known in the art and include dextran-mediated transfection, calcium phosphate precipitation, polybrene mediated transfection, protoplast fusion, electroporation, encapsulation of the polynucleotide(s) in liposomes, and direct microinjection of the DNA into nuclei.

[0313] Mammalian cell lines available as hosts for expression are well known in the art and include many immortalized cell lines available from the American Type Culture Collection (ATCC), including but not limited to Chinese hamster ovary (CHO) cells, HeLa cells, baby hamster kidney (BHK) cells, monkey kidney cells (COS), human hepatocellular carcinoma cells (e.g., Hep G2), human epithelial kidney 293 cells, and a number of other cell lines. Cell

lines of particular preference are selected through determining which cell lines have high expression levels and produce antibodies with constitutive PCSK9 binding properties.

[0314] In certain embodiments, antibodies and/or ABP are produced by at least one of the following hybridomas: 21B12, 31H4, 16F12, any the other hybridomas listed in Table 2 or disclosed in the examples. In certain embodiments, antigen binding proteins bind to PCSK9 with a dissociation constant (K_D) of less than approximately 1 nM, *e.g.*, 1000pM to 100 pM, 100 pM to 10 pM, 10 pM to 1 pM, and/or 1 pM to 0.1 pM or less.

[0315] In certain embodiments, antigen binding proteins comprise an immunoglobulin molecule of at least one of the IgG1, IgG2, IgG3, IgG4, Ig E, IgA, IgD, and IgM isotype. In certain embodiments, antigen binding proteins comprise a human kappa light chain and/or a human heavy chain. In certain embodiments, the heavy chain is of the IgG1, IgG2, IgG3, IgG4, IgE, IgA, IgD, or IgM isotype. In certain embodiments, antigen binding proteins have been cloned for expression in mammalian cells. In certain embodiments, antigen binding proteins comprise a constant region other than any of the constant regions of the IgG1, IgG2, IgG3, IgG4, IgE, IgA, IgD, and IgM isotype.

[0316] In certain embodiments, antigen binding proteins comprise a human lambda light chain and a human IgG2 heavy chain. In certain embodiments, antigen binding proteins comprise a human lambda light chain and a human IgG4 heavy chain. In certain embodiments, antigen binding proteins comprise a human lambda light chain and a human IgG1, IgG3, IgE, IgA, IgD or IgM heavy chain. In other embodiments, antigen binding proteins comprise a human kappa light chain and a human IgG2 heavy chain. In certain embodiments, antigen binding proteins comprise a human kappa light chain and a human IgG4 heavy chain. In certain embodiments, antigen binding proteins comprise a human kappa light chain and a human IgG1, IgG3, IgE, IgA, IgD or IgM heavy chain. In certain embodiments, antigen binding proteins comprise variable regions of antibodies ligated to a constant region that is neither the constant region for the IgG2 isotype, nor the constant region for the IgG4 isotype. In certain embodiments, antigen binding proteins have been cloned for expression in mammalian cells.

[0317] In certain embodiments, conservative modifications to the heavy and light chains of antibodies from at least one of the hybridoma lines: 21B12, 31H4 and 16F12 (and corresponding modifications to the encoding nucleotides) will produce antibodies to PCSK9 having functional and chemical characteristics similar to those of the antibodies from the

hybridoma lines: 21B12, 31H4 and 16F12. In contrast, in certain embodiments, substantial modifications in the functional and/or chemical characteristics of antibodies to PCSK9 can be accomplished by selecting substitutions in the amino acid sequence of the heavy and light chains that differ significantly in their effect on maintaining (a) the structure of the molecular backbone in the area of the substitution, for example, as a sheet or helical conformation, (b) the charge or hydrophobicity of the molecule at the target site, or (c) the bulk of the side chain.

[0318] For example, a "conservative amino acid substitution" can involve a substitution of a native amino acid residue with a nonnative residue such that there is little or no effect on the polarity or charge of the amino acid residue at that position. Furthermore, any native residue in the polypeptide can also be substituted with alanine, as has been previously described for "alanine scanning mutagenesis."

[0319] Desired amino acid substitutions (whether conservative or non-conservative) can be determined by those skilled in the art at the time such substitutions are desired. In certain embodiments, amino acid substitutions can be used to identify important residues of antibodies to PCSK9, or to increase or decrease the affinity of the antibodies to PCSK9 as described herein.

[0320] In certain embodiments, antibodies of the present invention can be expressed in cell lines other than hybridoma cell lines. In certain embodiments, sequences encoding particular antibodies can be used for transformation of a suitable mammalian host cell. According to certain embodiments, transformation can be by any known method for introducing polynucleotides into a host cell, including, for example packaging the polynucleotide in a virus (or into a viral vector) and transducing a host cell with the virus (or vector) or by transfection procedures known in the art, as exemplified by U.S. Pat. Nos. 4,399,216, 4,912,040, 4,740,461, and 4,959,455 (which patents are hereby incorporated herein by reference for any purpose). In certain embodiments, the transformation procedure used can depend upon the host to be transformed. Methods for introduction of heterologous polynucleotides into mammalian cells are well known in the art and include, but are not limited to, dextran-mediated transfection, calcium phosphate precipitation, polybrene mediated transfection, protoplast fusion, electroporation, encapsulation of the polynucleotide(s) in liposomes, and direct microinjection of the DNA into nuclei.

[0321] Mammalian cell lines available as hosts for expression are well known in the art and include, but are not limited to, many immortalized cell lines available from the American

Type Culture Collection (ATCC), including but not limited to Chinese hamster ovary (CHO) cells, HeLa cells, baby hamster kidney (BHK) cells, monkey kidney cells (COS), human hepatocellular carcinoma cells (e.g., Hep G2), and a number of other cell lines. In certain embodiments, cell lines can be selected through determining which cell lines have high expression levels and produce antibodies with constitutive HGF binding properties. Appropriate expression vectors for mammalian host cells are well known.

[0322] In certain embodiments, antigen binding proteins comprise one or more polypeptides. In certain embodiments, any of a variety of expression vector/host systems can be utilized to express polynucleotide molecules encoding polypeptides comprising one or more ABP components or the ABP itself. Such systems include, but are not limited to, microorganisms, such as bacteria transformed with recombinant bacteriophage, plasmid, or cosmid DNA expression vectors; yeast transformed with yeast expression vectors; insect cell systems infected with virus expression vectors (e.g., baculovirus); plant cell systems transfected with virus expression vectors (e.g., cauliflower mosaic virus, CaMV, tobacco mosaic virus, TMV) or transformed with bacterial expression vectors (e.g., Ti or pBR322 plasmid); or animal cell systems.

[0323] In certain embodiments, a polypeptide comprising one or more ABP components or the ABP itself is recombinantly expressed in yeast. Certain such embodiments use commercially available expression systems, e.g., the *Pichia* Expression System (Invitrogen, San Diego, CA), following the manufacturer's instructions. In certain embodiments, such a system relies on the pre-pro-alpha sequence to direct secretion. In certain embodiments, transcription of the insert is driven by the alcohol oxidase (AOX1) promoter upon induction by methanol.

[0324] In certain embodiments, a secreted polypeptide comprising one or more ABP components or the ABP itself is purified from yeast growth medium. In certain embodiments, the methods used to purify a polypeptide from yeast growth medium is the same as those used to purify the polypeptide from bacterial and mammalian cell supernatants.

[0325] In certain embodiments, a nucleic acid encoding a polypeptide comprising one or more ABP components or the ABP itself is cloned into a baculovirus expression vector, such as pVL1393 (PharMingen, San Diego, CA). In certain embodiments, such a vector can be used according to the manufacturer's directions (PharMingen) to infect *Spodoptera frugiperda* cells in

sF9 protein-free media and to produce recombinant polypeptide. In certain embodiments, a polypeptide is purified and concentrated from such media using a heparin-Sepharose column (Pharmacia).

[0326] In certain embodiments, a polypeptide comprising one or more ABP components or the ABP itself is expressed in an insect system. Certain insect systems for polypeptide expression are well known to those of skill in the art. In one such system, *Autographa californica* nuclear polyhedrosis virus (AcNPV) is used as a vector to express foreign genes in *Spodoptera frugiperda* cells or in *Trichoplusia* larvae. In certain embodiments, a nucleic acid molecule encoding a polypeptide can be inserted into a nonessential gene of the virus, for example, within the polyhedrin gene, and placed under control of the promoter for that gene. In certain embodiments, successful insertion of a nucleic acid molecule will render the nonessential gene inactive. In certain embodiments, that inactivation results in a detectable characteristic. For example, inactivation of the polyhedrin gene results in the production of virus lacking coat protein.

[0327] In certain embodiments, recombinant viruses can be used to infect *S. frugiperda* cells or *Trichoplusia* larvae. *See, e.g.*, Smith *et al.*, J. Virol., 46: 584 (1983); Engelhard *et al.*, Proc. Nat. Acad. Sci. (USA), 91: 3224-7 (1994).

[0328] In certain embodiments, polypeptides comprising one or more ABP components or the ABP itself made in bacterial cells are produced as insoluble inclusion bodies in the bacteria. In certain embodiments, host cells comprising such inclusion bodies are collected by centrifugation; washed in 0.15 M NaCl, 10 mM Tris, pH 8, 1 mM EDTA; and treated with 0.1 mg/ml lysozyme (Sigma, St. Louis, MO) for 15 minutes at room temperature. In certain embodiments, the lysate is cleared by sonication, and cell debris is pelleted by centrifugation for 10 minutes at 12,000 X g. In certain embodiments, the polypeptide-containing pellet is resuspended in 50 mM Tris, pH 8, and 10 mM EDTA; layered over 50% glycerol; and centrifuged for 30 minutes at 6000 X g. In certain embodiments, that pellet can be resuspended in standard phosphate buffered saline solution (PBS) free of Mg^{++} and Ca^{++}. In certain embodiments, the polypeptide is further purified by fractionating the resuspended pellet in a denaturing SDS polyacrylamide gel (*See, e.g.*, Sambrook *et al.*, *supra*). In certain embodiments, such a gel can be soaked in 0.4 M KCl to visualize the protein, which can be excised and electroeluted in gel-running buffer lacking SDS. According to certain embodiments, a

Glutathione-S-Transferase (GST) fusion protein is produced in bacteria as a soluble protein. In certain embodiments, such GST fusion protein is purified using a GST Purification Module (Pharmacia).

[0329] In certain embodiments, it is desirable to "refold" certain polypeptides, e.g., polypeptides comprising one or more ABP components or the ABP itself. In certain embodiments, such polypeptides are produced using certain recombinant systems discussed herein. In certain embodiments, polypeptides are "refolded" and/or oxidized to form desired tertiary structure and/or to generate disulfide linkages. In certain embodiments, such structure and/or linkages are related to certain biological activity of a polypeptide. In certain embodiments, refolding is accomplished using any of a number of procedures known in the art. Exemplary methods include, but are not limited to, exposing the solubilized polypeptide agent to a pH typically above 7 in the presence of a chaotropic agent. An exemplary chaotropic agent is guanidine. In certain embodiments, the refolding/oxidation solution also contains a reducing agent and the oxidized form of that reducing agent. In certain embodiments, the reducing agent and its oxidized form are present in a ratio that will generate a particular redox potential that allows disulfide shuffling to occur. In certain embodiments, such shuffling allows the formation of cysteine bridges. Exemplary redox couples include, but are not limited to, cysteine/cystamine, glutathione/dithiobisGSH, cupric chloride, dithiothreitol DTT/dithiane DTT, and 2-mercaptoethanol (bME)/dithio-bME. In certain embodiments, a co-solvent is used to increase the efficiency of refolding. Exemplary cosolvents include, but are not limited to, glycerol, polyethylene glycol of various molecular weights, and arginine.

[0330] In certain embodiments, one substantially purifies a polypeptide comprising one or more ABP components or the ABP itself. Certain protein purification techniques are known to those of skill in the art. In certain embodiments, protein purification involves crude fractionation of polypeptide fractionations from non-polypeptide fractions. In certain embodiments, polypeptides are purified using chromatographic and/or electrophoretic techniques. Exemplary purification methods include, but are not limited to, precipitation with ammonium sulphate; precipitation with PEG; immunoprecipitation; heat denaturation followed by centrifugation; chromatography, including, but not limited to, affinity chromatography (e.g., Protein-A-Sepharose), ion exchange chromatography, exclusion chromatography, and reverse phase chromatography; gel filtration; hydroxyapatite chromatography; isoelectric focusing;

polyacrylamide gel electrophoresis; and combinations of such and other techniques. In certain embodiments, a polypeptide is purified by fast protein liquid chromatography or by high pressure liquid chromotography (HPLC). In certain embodiments, purification steps can be changed or certain steps can be omitted, and still result in a suitable method for the preparation of a substantially purified polypeptide.

[0331] In certain embodiments, one quantitates the degree of purification of a polypeptide preparation. Certain methods for quantifying the degree of purification are known to those of skill in the art. Certain exemplary methods include, but are not limited to, determining the specific binding activity of the preparation and assessing the amount of a polypeptide within a preparation by SDS/PAGE analysis. Certain exemplary methods for assessing the amount of purification of a polypeptide preparation comprise calculating the binding activity of a preparation and comparing it to the binding activity of an initial extract. In certain embodiments, the results of such a calculation are expressed as "fold purification." The units used to represent the amount of binding activity depend upon the particular assay performed.

[0332] In certain embodiments, a polypeptide comprising one or more ABP components or the ABP itself is partially purified. In certain embodiments, partial purification can be accomplished by using fewer purification steps or by utilizing different forms of the same general purification scheme. For example, in certain embodiments, cation-exchange column chromatography performed utilizing an HPLC apparatus will generally result in a greater "fold purification" than the same technique utilizing a low-pressure chromatography system. In certain embodiments, methods resulting in a lower degree of purification can have advantages in total recovery of polypeptide, or in maintaining binding activity of a polypeptide.

[0333] In certain instances, the electrophoretic migration of a polypeptide can vary, sometimes significantly, with different conditions of SDS/PAGE. *See, e.g.*, Capaldi *et al.*, Biochem. Biophys. Res. Comm., 76: 425 (1977). It will be appreciated that under different electrophoresis conditions, the apparent molecular weights of purified or partially purified polypeptide can be different.

Exemplary Epitopes

[0334] Epitopes to which anti-PCSK9 antibodies bind are provided. In some embodiments, epitopes that are bound by the presently disclosed antibodies are particularly

useful. In some embodiments, antigen binding proteins that bind to any of the epitopes that are bound by the antibodies described herein are useful. In some embodiments, the epitopes bound by any of the antibodies listed in Table 2 and FIGs. 2 and 3 are especially useful. In some embodiments, the epitope is on the catalytic domain PCSK9.

[0335] In certain embodiments, a PCSK9 epitope can be utilized to prevent (*e.g.*, reduce) binding of an anti-PCSK9 antibody or antigen binding protein to PCSK9. In certain embodiments, a PCSK9 epitope can be utilized to decrease binding of an anti-PCSK9 antibody or antigen binding protein to PCSK9. In certain embodiments, a PCSK9 epitope can be utilized to substantially inhibit binding of an anti-PCSK9 antibody or antigen binding protein to PCSK9.

[0336] In certain embodiments, a PCSK9 epitope can be utilized to isolate antibodies or antigen binding proteins that bind to PCSK9. In certain embodiments, a PCSK9 epitope can be utilized to generate antibodies or antigen binding proteins which bind to PCSK9. In certain embodiments, a PCSK9 epitope or a sequence comprising a PCSK9 epitope can be utilized as an immunogen to generate antibodies or antigen binding proteins that bind to PCSK9. In certain embodiments, a PCSK9 epitope can be administered to an animal, and antibodies that bind to PCSK9 can subsequently be obtained from the animal. In certain embodiments, a PCSK9 epitope or a sequence comprising a PCSK9 epitope can be utilized to interfere with normal PCSK9-mediated activity, such as association of PCSK9 with the LDLR.

[0337] In some embodiments, antigen binding proteins disclosed herein bind specifically to N-terminal prodomain, a subtilisin-like catalytic domain and/or a C-terminal domain. In some embodiments, the antigen binding protein binds to the substrate-binding groove of PCSK-9 (described in Cunningham et al., incorporated herein in its entirety by reference).

[0338] In some embodiments, the domain(s)/region(s) containing residues that are in contact with or are buried by an antibody can be identified by mutating specific residues in PCSK9 (e.g., a wild-type antigen) and determining whether the antigen binding protein can bind the mutated or variant PCSK9 protein. By making a number of individual mutations, residues that play a direct role in binding or that are in sufficiently close proximity to the antibody such that a mutation can affect binding between the antigen binding protein and antigen can be identified. From a knowledge of these amino acids, the domain(s) or region(s) of the antigen that contain residues in contact with the antigen binding protein or covered by the antibody can be

elucidated. Such a domain can include the binding epitope of an antigen binding protein. One specific example of this general approach utilizes an arginine/glutamic acid scanning protocol (see, e.g., Nanevicz, T., *et al.*, 1995, *J. Biol. Chem.*, 270:37, 21619-21625 and Zupnick, A., *et al.*, 2006, *J. Biol. Chem.*, 281:29, 20464-20473). In general, arginine and glutamic acids are substituted (typically individually) for an amino acid in the wild-type polypeptide because these amino acids are charged and bulky and thus have the potential to disrupt binding between an antigen binding protein and an antigen in the region of the antigen where the mutation is introduced. Arginines that exist in the wild-type antigen are replaced with glutamic acid. A variety of such individual mutants are obtained and the collected binding results analyzed to determine what residues affect binding.

[0339] Example 39 describes one such arginine/glutamic acid scanning of PCSK9 for PCSK9 antigen binding proteins provided herein. A series of mutant PCSK9 antigens were created, with each mutant antigen having a single mutation. Binding of each mutant PCSK9 antigen with various PCSK9 ABPs was measured and compared to the ability of the selected ABPs to bind wild-type PCSK9 (SEQ ID NO: 303).

[0340] An alteration (for example a reduction or increase) in binding between an antigen binding protein and a variant PCSK9 as used herein means that there is a change in binding affinity (*e.g.*, as measured by known methods such as Biacore testing or the bead based assay described below in the examples), EC_{50}, and/or a change (for example a reduction) in the total binding capacity of the antigen binding protein (for example, as evidenced by a decrease in Bmax in a plot of antigen binding protein concentration versus antigen concentration). A significant alteration in binding indicates that the mutated residue is directly involved in binding to the antigen binding protein or is in close proximity to the binding protein when the binding protein is bound to antigen.

[0341] In some embodiments, a significant reduction in binding means that the binding affinity, EC50, and/or capacity between an antigen binding protein and a mutant PCSK9 antigen is reduced by greater than 10%, greater than 20%, greater than 40 %, greater than 50 %, greater than 55 %, greater than 60 %, greater than 65 %, greater than 70 %, greater than75 %, greater than 80 %, greater than 85 %, greater than 90% or greater than 95% relative to binding between the antigen binding protein and a wild type PCSK9 (e.g., shown in SEQ ID NO: 1 and/or SEQ ID NO: (303). In certain embodiments, binding is reduced below detectable limits.

In some embodiments, a significant reduction in binding is evidenced when binding of an antigen binding protein to a variant PCSK9 protein is less than 50% (for example, less than 40%, 35%, 30%, 25%, 20%, 15% or 10%) of the binding observed between the antigen binding protein and a wild-type PCSK9 protein (for example, the protein of SEQ ID NO: 1 and/or SEQ ID NO; (303). Such binding measurements can be made using a variety of binding assays known in the art. A specific example of one such assay is described in Example 39.

[0342] In some embodiments, antigen binding proteins are provided that exhibit significantly lower binding for a variant PCSK9 protein in which a residue in a wild-type PCSK9 protein (e.g., SEQ ID NO: 1 or SEQ ID NO: 303 is substituted with arginine or glutamic acid. In some embodiments, binding of an antigen binding protein is significantly reduced or increased for a variant PCSK9 protein having any one or more (e.g., 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or 244) of the following mutations: R207E, D208R, R185E, R439E, E513R, V538R, E539R, T132R, S351R, A390R, A413R, E582R, D162R, R164E, E167R, S123R, E129R, A311R, D313R, D337R, R519E, H521R, and Q554R as compared to a wild-type PCSK9 protein (*e.g.*, SEQ ID NO: 1 or SEQ ID NO: 303. In the shorthand notation used here, the format is: Wild type residue: Position in polypeptide: Mutant residue, with the numbering of the residues as indicated in SEQ ID NO: 1or SEQ ID NO: 303.

[0343] In some embodiments, binding of an antigen binding protein is significantly reduced or increased for a mutant PCSK9 protein having one or more (e.g., 1, 2, 3, 4, 5, or more) mutations at the following positions: 207, 208, 185, 181, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554, as shown in SEQ ID NO: 1 as compared to a wild-type PCSK9 protein (e.g., SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, binding of an antigen binding protein is reduced or increased for a mutant PCSK9 protein having one or more (e.g., 1, 2, 3, 4, 5, or more) mutations at the following positions: 207, 208, 185, 181, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554, as shown in SEQ ID NO: 1 as compared to a wild-type PCSK9 protein (e.g., SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, binding of an antigen binding protein is substantially reduced or increased for a mutant PCSK9 protein having one or more (e.g., 1, 2, 3, 4, 5, or more) mutations at the following positions: 207, 208, 185, 181, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554,

within SEQ ID NO: 1 as compared to a wild-type PCSK9 protein (e.g., SEQ ID NO: 1 or SEQ ID NO: 303.

[0344] In some embodiments, binding of an ABP is significantly reduced or increased for a mutant PCSK9 protein having one or more (e.g., 1, 2, 3, 4, 5, etc.) of the following mutations: R207E, D208R, R185E, R439E, E513R, V538R, E539R, T132R, S351R, A390R, A413R, E582R, D162R, R164E, E167R, S123R, E129R, A311R, D313R, D337R, R519E, H521R, and Q554R within SEQ ID NO: 1 or SEQ ID NO: 303, as compared to a wild-type PCSK9 protein (e.g., SEQ ID NO: 1 or SEQ ID NO: 303).

[0345] In some embodiments, binding of an ABP is significantly reduced or increased for a mutant PCSK9 protein having one or more (e.g., 1, 2, 3, 4, 5, etc.) of the following mutations: R207E, D208R, R185E, R439E, E513R, V538R, E539R, T132R, S351R, A390R, A413R, and E582R within SEQ ID NO: 1 or SEQ ID NO: 303, as compared to a wild-type PCSK9 protein (*e.g.*, SEQ ID NO: 1 or SEQ ID NO: 303). In some embodiments, the binding is reduced. In some embodiments, the reduction in binding is observed as a change in EC50. In some embodiments, the change in EC50 is an increase in the numerical value of the EC50 (and thus is a decrease in binding).

[0346] In some embodiments, binding of an ABP is significantly reduced or increased for a mutant PCSK9 protein having one or more (e.g., 1, 2, 3, 4, 5, etc.) of the following mutations: D162R, R164E, E167R, S123R, E129R, A311R, D313R, D337R, R519E, H521R, and Q554R within SEQ ID NO: 1, as compared to a wild-type PCSK9 protein (e.g., SEQ ID NO: 1 or SEQ ID NO: 303). In some embodiments, the binding is reduced. In some embodiments, the reduction in binding is observed as a change in Bmax. In some embodiments, the shift in Bmax is a reduction of the maximum signal generated by the ABP. In some embodiments, for an amino acid to be part of an epitope, the Bmax is reduced by at least 10%, for example, reductions of at least any of the following amounts: 20, 30, 40, 50, 60, 70, 80, 90, 95, 98, 99, or 100 percent can, in some embodiments, indicate that the residue is part of the epitope.

[0308] Although the variant forms just listed are referenced with respect to the wild-type sequence shown in SEQ ID NO: 1 or SEQ ID NO: 303, it will be appreciated that in an allelic variant of PCSK9 the amino acid at the indicated position could differ. Antigen binding proteins showing significantly lower binding for such allelic forms of PCSK9 are also

contemplated. Accordingly, in some embodiments, any of the above embodiments can be compared to an allelic sequence, rather than purely the wild-type sequence shown in FIG. 1A

[0347] In some embodiments, binding of an antigen binding protein is significantly reduced for a variant PCSK9 protein in which the residue at a selected position in the wild-type PCSK9 protein is mutated to any other residue. In some embodiments, the herein described arginine/glutamic acid replacements are used for the identified positions. In some embodiments, alanine is used for the identified positions.

[0348] As noted above, residues directly involved in binding or covered by an antigen binding protein can be identified from scanning results. These residues can thus provide an indication of the domains or regions of SEQ ID NO: 1 (or SEQ ID NO: 303 or SEQ ID NO: 3) that contain the binding region(s) to which antigen binding proteins bind. As can be seen from the results summarized in Example 39, in some embodiments an antigen binding protein binds to a domain containing at least one of amino acids: 207, 208, 185, 181, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554 of SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, the antigen binding protein binds to a region containing at least one of amino acids 207, 208, 185, 181, 439, 513, 538, 539, 132, 351, 390, 413, 582, 162, 164, 167, 123, 129, 311, 313, 337, 519, 521, and 554 of SEQ ID NO: 1 or SEQ ID NO: 303.

[0349] In some embodiments, the antigen binding protein binds to a region containing at least one of amino acids 162, 164, 167, 207 and/or 208 of SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, more than one (e.g., 2, 3, 4, or 5) of the identified residues are part of the region that is bound by the ABP. In some embodiments, the ABP competes with ABP 21B12.

[0350] In some embodiments, the antigen binding protein binds to a region containing at least one of amino acid 185 of SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, the ABP competes with ABP 31H4.

[0351] In some embodiments, the antigen binding protein binds to a region containing at least one of amino acids 439, 513, 538, and/or 539 of SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, more than one (e.g., 2, 3, or 4) of the identified residues are part of the region that is bound by the ABP. In some embodiments, the ABP competes with ABP 31A4.

[0352] In some embodiments, the antigen binding protein binds to a region containing at least one of amino acids 123, 129, 311, 313, 337, 132, 351, 390, and/or 413 of SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, more than one (e.g., 2, 3, 4, 5, 6, 7, 8, or 9) of the identified residues are part of the region that is bound by the ABP. In some embodiments, the ABP competes with ABP 12H11.

[0353] In some embodiments, the antigen binding protein binds to a region containing at least one of amino acid 582, 519, 521, and/or 554 of SEQ ID NO: 1 or SEQ ID NO: 303. In some embodiments, more than one (e.g., 2, 3, or 4) of the identified residues are part of the region that is bound by the ABP. In some embodiments, the ABP competes with ABP 3C4.

[0354] In some embodiments, the antigen binding proteins binds to the foregoing regions within a fragment or the full length sequence of SEQ ID NO: 1 or SEQ ID NO: 303. In other embodiments, antigen binding proteins bind to polypeptides consisting of these regions. The reference to "SEQ ID NO: 1 or SEQ ID NO: 303" denotes that one or both of these sequences can be employed or relevant. The phrase does not denote that only one should be employed.

[0355] As noted above, the above description references specific amino acid positions with reference to SEQ ID NO: 1. However, throughout the specification generally, reference is made to a Pro/Cat domain that commences at position 31, which is provided in SEQ ID NO: 3. As noted below, SEQ ID NO: 1 and SEQ ID NO: 303 lack the signal sequence of PCSK9. As such, any comparison between these various disclosures should take this difference in numbering into account. In particular, any amino acid position in SEQ ID NO: 1, will correspond to an amino acid position 30 amino acids further into the protein in SEQ ID NO: 3. For example, position 207 of SEQ ID NO: 1, corresponds to position 237 of SEQ ID NO: 3 (the full length sequence, and the numbering system used in the present specification generally). Table 39.6 outlines how the above noted positions, which reference SEQ ID NO: 1 (and/or SEQ ID NO: 303) correspond to SEQ ID NO: 3 (which includes the signal sequence). Thus, any of the above noted embodiments that are described in regard to SEQ ID NO: 1 (and/or SEQ ID NO: 303), are described in reference to SEQ ID NO: 3, by the noted corresponding positions.

[0356] In some embodiments, ABP 21B12 binds to an epitope including residues 162-167 (e.g., residues D162-E167 of SEQ ID NO: 1). In some embodiments, ABP 12H11

binds to an epitope that includes residues 123-132 (*e.g.*, S123-T132 of SEQ ID NO: 1). In some embodiments, ABP 12H11 binds to an epitope that includes residues 311-313 (*e.g.*, A311-D313 of SEQ ID NO: 1). In some embodiments, ABPs can bind to an epitope that includes any one of these strands of sequences.

Competing Antigen Binding Proteins

[0357] In another aspect, antigen binding proteins are provided that compete with one of the exemplified antibodies or functional fragments binding to the epitope described herein for specific binding to PCSK9. Such antigen binding proteins can also bind to the same epitope as one of the herein exemplified antigen binding proteins, or an overlapping epitope. Antigen binding proteins and fragments that compete with or bind to the same epitope as the exemplified antigen binding proteins are expected to show similar functional properties. The exemplified antigen binding proteins and fragments include those described above, including those with the heavy and light chains, variable region domains and CDRs included in TABLE 2 And/or FIGs. 2-3 and 15 . Thus, as a specific example, the antigen binding proteins that are provided include those that compete with an antibody or antigen binding protein having:

(a) all 6 of the CDRs listed for an antibody listed in FIGs. 2-3 and 15;

(b) a VH and a VL listed for an antibody listed in Table 2; or

(c) two light chains and two heavy chains as specified for an antibody listed in Table 2.

Certain Therapeutic Uses and Pharmaceutical Compositions

[0358] In certain instances, PCSK9 activity correlates with a number of human disease states. For example, in certain instances, too much or too little PCSK9 activity correlates with certain conditions, such as hypercholesterolemia. Therefore, in certain instances, modulating PCSK9 activity can be therapeutically useful. In certain embodiments, a neutralizing antigen binding protein to PCSK9 is used to modulate at least one PCSK9 activity (*e.g.*, binding to LDLR). Such methods can treat and/or prevent and/or reduce the risk of disorders that relate to elevated serum cholesterol levels or in which elevated cholesterol levels are relevant.

[0359] As will be appreciated by one of skill in the art, in light of the present disclosure, disorders that relate to, involve, or can be influenced by varied cholesterol, LDL, or LDLR levels can be addressed by various embodiments of the antigen binding proteins. In some

embodiments, a "cholesterol related disorder" (which includes "serum cholesterol related disorders") includes any one or more of the following: hypercholesterolemia, heart disease, metabolic syndrome, diabetes, coronary heart disease, stroke, cardiovascular diseases, Alzheimers disease and generally dyslipidemias, which can be manifested, for example, by an elevated total serum cholesterol, elevated LDL, elevated triglycerides, elevated VLDL, and/or low HDL. Some non-limiting examples of primary and secondary dyslipidemias that can be treated using an ABP, either alone, or in combination with one or more other agents include the metabolic syndrome, diabetes mellitus, familial combined hyperlipidemia, familial hypertriglyceridemia, familial hypercholesterolemias, including heterozygous hypercholesterolemia, homozygous hypercholesterolemia, familial defective apoplipoprotein B-100; polygenic hypercholesterolemia; remnant removal disease, hepatic lipase deficiency; dyslipidemia secondary to any of the following: dietary indiscretion, hypothyroidism, drugs including estrogen and progestin therapy, beta-blockers, and thiazide diuretics; nephrotic syndrome, chronic renal failure, Cushing's syndrome, primary biliary cirrhosis, glycogen storage diseases, hepatoma, cholestasis, acromegaly, insulinoma, isolated growth hormone deficiency, and alcohol-induced hypertriglyceridemia. ABP can also be useful in preventing or treating atherosclerotic diseases, such as, for example, coronary heart disease, coronary artery disease, peripheral arterial disease, stroke (ischaemic and hemorrhagic), angina pectoris, or cerebrovascular disease and acute coronary syndrome, myocardial infarction. In some embodiments, the ABP is useful in reducing the risk of: nonfatal heart attacks, fatal and non-fatal strokes, certain types of heart surgery, hospitalization for heart failure, chest pain in patients with heart disease, and/or cardiovascular events because of established heart disease such as prior heart attack, prior heart surgery, and/or chest pain with evidence of clogged arteries. In some embodiments, the ABP and methods can be used to reduce the risk of recurrent cardiovascular events.

[0360] As will be appreciated by one of skill in the art, diseases or disorders that are generally addressable (either treatable or preventable) through the use of statins can also benefit from the the application of the instant antigen binding proteins. In addition, in some embodiments, disorders or disease that can benefit from the prevention of cholesterol synthesis or increased LDLR expression can also be treated by various embodiments of the antigen binding proteins. In addition, as will be appreciated by one of skill in the art, the use of the anti-

PCSK9 antibodies can be especially useful in the treatment of Diabetes. Not only is Diabetes a risk factor for coronary heart disease, but insulin increases the expression of PCSK9. That is, people with Diabetes have elevated plasma lipid levels (which can be related to high PCSK9 levels) and can benefit from lowering those levels. This is generally discussed in more detail in Costet et al. ("Hepatic PCSK9 Expression is Regulated by Nutirtional Status via Insulin and Sterol Regulatiory Element-binding Protein 1C", J. Biol. Chem., 281: 6211-6218, 2006), the entirety of which is incorporated herein by reference.

[0361] In some embodiments, the antigen binding protein is administered to those who have diabetes mellitus, abdominal aortic aneurysm, atherosclerosis and/or peripheral vascular disease in order to decrease their serum cholesterol levels to a safer range. In some embodiments, the antigen binding protein is administered to patients at risk of developing any of the herein described disorders. In some embodiments, the ABPs are administered to subjects that smoke, have hypertension or a familial history of early heart attacks.

[0362] In some embodiments, a subject is administered an ABP if they are at a moderate risk or higher on the 2004 NCEP treatment goals. In some embodiments, the ABP is admininstered to a subject if the subject's LDL cholesterol level is greater than 160 mg/dl. In some embodiments, the ABP is administered if the subjects LDL cholesterol level is greater than 130 (and they have a moderate or moderately high risk according to the 2004 NCEP treatment goals). In some embodiments, the ABP is administered if the subjects LDL cholesterol level is greater than 100 (and they have a high or very high risk according to the 2004 NCEP treatment goals).

[0363] A physician will be able to select an appropriate treatment indications and target lipid levels depending on the individual profile of a particular patient. One well-accepted standard for guiding treatment of hyperlipidemia is the Third Report of the National Cholesterol Education Program (NCEP) Expert Panel on Detection, Evaluation, and Treatment of the High Blood Cholesterol in Adults (Adult Treatment Panel III) Final Report, National Institutes of Health, NIH Publication No. 02-5215 (2002), the printed publication of which is hereby incorporated by reference in its entirety.

[0364] In some embodiments, antigen binding proteins to PCSK9 are used to decrease the amount of PCSK9 activity from an abnormally high level or even a normal level. In some embodiments, antigen binding proteins to PCSK9 are used to treat or prevent

hypercholesterolemia and/or in the preparation of medicaments therefore and/or for other cholesterol related disorders (such as those noted herein). In certain embodiments, an antigen binding protein to PCSK9 is used to treat or prevent conditions such as hypercholesterolemia in which PCSK9 activity is normal. In such conditions, for example, reduction of PCSK9 activity to below normal can provide a therapeutic effect.

[0365] In some embodiments, more than one antigen binding protein to PCSK9 is used to modulate PCSK9 activity.

[0366] In certain embodiments, methods are provided of treating a cholesterol related disorder, such as hypercholesterolemia comprising administering a therapeutically effective amount of one or more antigen binding proteins to PCSK9 and another therapeutic agent.

[0367] In certain embodiments, an antigen binding protein to PCSK9 is administered alone. In certain embodiments, an antigen binding protein to PCSK9 is administered prior to the administration of at least one other therapeutic agent. In certain embodiments, an antigen binding protein to PCSK9 is administered concurrent with the administration of at least one other therapeutic agent. In certain embodiments, an antigen binding protein to PCSK9 is administered subsequent to the administration of at least one other therapeutic agent. In other embodiments, an antigen binding protein to PCSK9 is administered prior to the administration of at least one other therapeutic agent. Therapeutic agents (apart from the antigen binding protein), include, but are not limited to, at least one other cholesterol-lowering (serum and/or total body cholesterol) agent or an agent. In some embodiments, the agent increases the expression of LDLR, have been observed to increase serum HDL levels, lower LDL levels or lower triglyceride levels. Exemplary agents include, but are not limited to, statins (atorvastatin, cerivastatin, fluvastatin, lovastatin, mevastatin, pitavastatin, pravastatin, rosuvastatin, simvastatin), Nicotinic acid (Niacin) (NIACOR, NIASPAN (slow release niacin), SLO-NIACIN (slow release niacin)), Fibric acid (LOPID (Gemfibrozil), TRICOR (fenofibrate), Bile acid sequestrants (QUESTRAN (cholestyramine), colesevelam (WELCHOL), COLESTID (colestipol)), Cholesterol absorption inhibitors (ZETIA (ezetimibe)), Combining nicotinic acid with statin (ADVICOR (LOVASTATIN and NIASPAN), Combining a statin with an absorption inhibitor (VYTORIN (ZOCOR and ZETIA) and/or lipid modifying agents. In some embodiments, the ABP is combined with PPAR gamma agonsits, PPAR alpha/gamma agonists, squalene synthase inhibitors, CETP inhibitors, anti-hypertensives, anti-diabetic agents (such as sulphonyl ureas,

insulin, GLP-1 analogs, DDPIV inhibitors), ApoB modulators, MTP inhibitoris and /or arteriosclerosis obliterans treatments. In some embodiments, the ABP is combined with an agent that increases the level of LDLR protein in a subject, such as statins, certain cytokines like oncostatin M, estrogen, and/or certain herbal ingredients such as berberine. In some embodiments, the ABP is combined with an agent that increases serum cholesterol levels in a subject (such as certain anti-psycotic agents, certain HIV protease inhibitors, dietary factors such as high fructose, sucrose, cholesterol or certain fatty acids and certain nuclear receptor agonists and antagonists for RXR, RAR, LXR, FXR). In some embodiments, the ABP is combined with an agent that increases the level of PCSK9 in a subject, such as statins and/or insulin. The combination of the two can allow for the undesireable side-effects of other agents to be mitigated by the ABP. As will be appreciated by one of skill in the art, in some embodiments, the ABP is combined with the other agent/compound. In some embodiments, the ABP and other agent are administered concurrently. In some embodiments, the ABP and other agent are not administered simultaneously, with the ABP being administered before or after the agent is administered. In some embodiments, the subject receives both the ABP and the other agent (that increases the level of LDLR) during a same period of prevention, occurrence of a disorder, and/or period of treatment.

[0368] Pharmaceutical compositions of the invention can be administered in combination therapy, *i.e.*, combined with other agents. In certain embodiments, the combination therapy comprises an antigen binding protein capable of binding PCSK9, in combination with at least one anti-cholesterol agent. Agents include, but are not limited to, in vitro synthetically prepared chemical compositions, antibodies, antigen binding regions, and combinations and conjugates thereof. In certain embodiments, an agent can act as an agonist, antagonist, alllosteric modulator, or toxin. In certain embodiments, an agent can act to inhibit or stimulate its target (e.g., receptor or enzyme activation or inhibition), and thereby promote increased expression of LDLR or decrease serum cholesterol levels.

[0369] In certain embodiments, an antigen binding protein to PCSK9 can be administered prior to, concurrent with, and subsequent to treatment with a cholesterol-lowering (serum and/or total cholesterol) agent. In certain embodiments, an antigen binding protein to PCSK9 can be administered prophylactially to prevent or mitigate the onset of hypercholesterolemia, heart disease, diabetes, and/or any of the cholesterol related disorder. In

certain embodiments, an antigen binding protein to PCSK9 can be administered for the treatment of an existing hypercholesterolemia condition. In some embodiments, the ABP delays the onset of the disorder and/or symptoms associated with the disorder. In some embodiments, the ABP is provided to a subject lacking any sympotoms of any one of the cholesterol related disorders or a subset thereof.

[0370] In certain embodiments, an antigen binding protein to PCSK9 is used with particular therapeutic agents to treat various cholesterol related disorders, such as hypercholesterolemia. In certain embodiments, in view of the condition and the desired level of treatment, two, three, or more agents can be administered. In certain embodiments, such agents can be provided together by inclusion in the same formulation. In certain embodiments, such agent(s) and an antigen binding protein to PCSK9 can be provided together by inclusion in the same formulation. In certain embodiments, such agents can be formulated separately and provided together by inclusion in a treatment kit. In certain embodiments, such agents and an antigen binding protein to PCSK9 can be formulated separately and provided together by inclusion in a treatment kit. In certain embodiments, such agents can be provided separately. In certain embodiments, when administered by gene therapy, the genes encoding protein agents and/or an antigen binding protein to PCSK9 can be included in the same vector. In certain embodiments, the genes encoding protein agents and/or an antigen binding protein to PCSK9 can be under the control of the same promoter region. In certain embodiments, the genes encoding protein agents and/or an antigen binding protein to PCSK9 can be in separate vectors.

[0371] In certain embodiments, the invention provides for pharmaceutical compositions comprising an antigen binding protein to PCSK9 together with a pharmaceutically acceptable diluent, carrier, solubilizer, emulsifier, preservative and/or adjuvant.

[0372] In certain embodiments, the invention provides for pharmaceutical compositions comprising an antigen binding protein to PCSK9 and a therapeutically effective amount of at least one additional therapeutic agent, together with a pharmaceutically acceptable diluent, carrier, solubilizer, emulsifier, preservative and/or adjuvant.

[0373] In certain embodiments, an antigen binding protein to PCSK9 can be used with at least one therapeutic agent for inflammation. In certain embodiments, an antigen binding protein to PCSK9 can be used with at least one therapeutic agent for an immune disorder. Exemplary therapeutic agents for inflammation and immune disorders include, but are not

limited to cyclooxygenase type 1 (COX-1) and cyclooxygenase type 2 (COX-2) inhibitors small molecule modulators of 38 kDa mitogen-activated protein kinase (p38-MAPK); small molecule modulators of intracellular molecules involved in inflammation pathways, wherein such intracellular molecules include, but are not limited to, jnk, IKK, NF-κB, ZAP70, and lck. Certain exemplary therapeutic agents for inflammation are described, e.g., in C.A. Dinarello & L.L. Moldawer *Proinflammatory and Anti-Inflammatory Cytokines in Rheumatoid Arthritis: A Primer for Clinicians* Third Edition (2001) Amgen Inc. Thousand Oaks, CA.

[0374] In certain embodiments, pharmaceutical compositions will include more than one different antigen binding protein to PCSK9. In certain embodiments, pharmaceutical compositions will include more than one antigen binding protein to PCSK9 wherein the antigen binding proteins to PCSK9 bind more than one epitope. In some embodiments, the various antigen binding proteins will not compete with one another for binding to PCSK9. In some embodiments, any of the antigen binding proteins depicted in Table 2 and FIGs. 2 and/or 3 can be combined together in a pharmaceutical composition.

[0375] In certain embodiments, acceptable formulation materials preferably are nontoxic to recipients at the dosages and concentrations employed. In some embodiments, the formulation material(s) are for s.c. and/or I.V. administration. In certain embodiments, the pharmaceutical composition can contain formulation materials for modifying, maintaining or preserving, for example, the pH, osmolarity, viscosity, clarity, color, isotonicity, odor, sterility, stability, rate of dissolution or release, adsorption or penetration of the composition. In certain embodiments, suitable formulation materials include, but are not limited to, amino acids (such as glycine, glutamine, asparagine, arginine or lysine); antimicrobials; antioxidants (such as ascorbic acid, sodium sulfite or sodium hydrogen-sulfite); buffers (such as borate, bicarbonate, Tris-HCl, citrates, phosphates or other organic acids); bulking agents (such as mannitol or glycine); chelating agents (such as ethylenediamine tetraacetic acid (EDTA)); complexing agents (such as caffeine, polyvinylpyrrolidone, beta-cyclodextrin or hydroxypropyl-beta-cyclodextrin); fillers; monosaccharides; disaccharides; and other carbohydrates (such as glucose, mannose or dextrins); proteins (such as serum albumin, gelatin or immunoglobulins); coloring, flavoring and diluting agents; emulsifying agents; hydrophilic polymers (such as polyvinylpyrrolidone); low molecular weight polypeptides; salt-forming counterions (such as sodium); preservatives (such as benzalkonium chloride, benzoic acid, salicylic acid, thimerosal, phenethyl alcohol,

methylparaben, propylparaben, chlorhexidine, sorbic acid or hydrogen peroxide); solvents (such as glycerin, propylene glycol or polyethylene glycol); sugar alcohols (such as mannitol or sorbitol); suspending agents; surfactants or wetting agents (such as pluronics, PEG, sorbitan esters, polysorbates such as polysorbate 20, polysorbate 80, triton, tromethamine, lecithin, cholesterol, tyloxapal); stability enhancing agents (such as sucrose or sorbitol); tonicity enhancing agents (such as alkali metal halides, preferably sodium or potassium chloride, mannitol sorbitol); delivery vehicles; diluents; excipients and/or pharmaceutical adjuvants. (*Remington's Pharmaceutical Sciences*, 18th Edition, A.R. Gennaro, ed., Mack Publishing Company (1995). In some embodiments, the formulation comprises PBS; 20mM NaOAC, pH 5.2, 50mM NaCl; and/or 10mM NAOAC, pH 5.2, 9% Sucrose.

[0376] In certain embodiments, an antigen binding protein to PCSK9 and/or a therapeutic molecule is linked to a half-life extending vehicle known in the art. Such vehicles include, but are not limited to, polyethylene glycol, glycogen (e.g., glycosylation of the ABP), and dextran. Such vehicles are described, e.g., in U.S. Application Serial No. 09/428,082, now US Patent No. 6,660,843 and published PCT Application No. WO 99/25044, which are hereby incorporated by reference for any purpose.

[0377] In certain embodiments, the optimal pharmaceutical composition will be determined by one skilled in the art depending upon, for example, the intended route of administration, delivery format and desired dosage. See, for example, *Remington's Pharmaceutical Sciences, supra*. In certain embodiments, such compositions may influence the physical state, stability, rate of *in vivo* release and rate of *in vivo* clearance of the antibodies of the invention.

[0378] In certain embodiments, the primary vehicle or carrier in a pharmaceutical composition can be either aqueous or non-aqueous in nature. For example, in certain embodiments, a suitable vehicle or carrier can be water for injection, physiological saline solution or artificial cerebrospinal fluid, possibly supplemented with other materials common in compositions for parenteral administration. In some embodiments, the saline comprises isotonic phosphate-buffered saline. In certain embodiments, neutral buffered saline or saline mixed with serum albumin are further exemplary vehicles. In certain embodiments, pharmaceutical compositions comprise Tris buffer of about pH 7.0-8.5, or acetate buffer of about pH 4.0-5.5, which can further include sorbitol or a suitable substitute therefore. In certain embodiments, a

composition comprising an antigen binding protein to PCSK9, with or without at least one additional therapeutic agents, can be prepared for storage by mixing the selected composition having the desired degree of purity with optional formulation agents (*Remington's Pharmaceutical Sciences, supra*) in the form of a lyophilized cake or an aqueous solution. Further, in certain embodiments, a composition comprising an antigen binding protein to PCSK9, with or without at least one additional therapeutic agents, can be formulated as a lyophilizate using appropriate excipients such as sucrose.

[0379] In certain embodiments, the pharmaceutical composition can be selected for parenteral delivery. In certain embodiments, the compositions can be selected for inhalation or for delivery through the digestive tract, such as orally. The preparation of such pharmaceutically acceptable compositions is within the ability of one skilled in the art.

[0380] In certain embodiments, the formulation components are present in concentrations that are acceptable to the site of administration. In certain embodiments, buffers are used to maintain the composition at physiological pH or at a slightly lower pH, typically within a pH range of from about 5 to about 8.

[0381] In certain embodiments, when parenteral administration is contemplated, a therapeutic composition can be in the form of a pyrogen-free, parenterally acceptable aqueous solution comprising a desired antigen binding protein to PCSK9, with or without additional therapeutic agents, in a pharmaceutically acceptable vehicle. In certain embodiments, a vehicle for parenteral injection is sterile distilled water in which an antigen binding protein to PCSK9, with or without at least one additional therapeutic agent, is formulated as a sterile, isotonic solution, properly preserved. In certain embodiments, the preparation can involve the formulation of the desired molecule with an agent, such as injectable microspheres, bio-erodible particles, polymeric compounds (such as polylactic acid or polyglycolic acid), beads or liposomes, that can provide for the controlled or sustained release of the product which can then be delivered via a depot injection. In certain embodiments, hyaluronic acid can also be used, and can have the effect of promoting sustained duration in the circulation. In certain embodiments, implantable drug delivery devices can be used to introduce the desired molecule.

[0382] In certain embodiments, a pharmaceutical composition can be formulated for inhalation. In certain embodiments, an antigen binding protein to PCSK9, with or without at least one additional therapeutic agent, can be formulated as a dry powder for inhalation. In

certain embodiments, an inhalation solution comprising an antigen binding protein to PCSK9, with or without at least one additional therapeutic agent, can be formulated with a propellant for aerosol delivery. In certain embodiments, solutions can be nebulized. Pulmonary administration is further described in PCT application no. PCT/US94/001875, which describes pulmonary delivery of chemically modified proteins.

[0383] In certain embodiments, it is contemplated that formulations can be administered orally. In certain embodiments, an antigen binding protein to PCSK9, with or without at least one additional therapeutic agents, that is administered in this fashion can be formulated with or without those carriers customarily used in the compounding of solid dosage forms such as tablets and capsules. In certain embodiments, a capsule can be designed to release the active portion of the formulation at the point in the gastrointestinal tract when bioavailability is maximized and pre-systemic degradation is minimized. In certain embodiments, at least one additional agent can be included to facilitate absorption of an antigen binding protein to PCSK9 and/or any additional therapeutic agents. In certain embodiments, diluents, flavorings, low melting point waxes, vegetable oils, lubricants, suspending agents, tablet disintegrating agents, and binders can also be employed.

[0384] In certain embodiments, a pharmaceutical composition can involve an effective quantity of an antigen binding protein to PCSK9, with or without at least one additional therapeutic agents, in a mixture with non-toxic excipients which are suitable for the manufacture of tablets. In certain embodiments, by dissolving the tablets in sterile water, or another appropriate vehicle, solutions can be prepared in unit-dose form. In certain embodiments, suitable excipients include, but are not limited to, inert diluents, such as calcium carbonate, sodium carbonate or bicarbonate, lactose, or calcium phosphate; or binding agents, such as starch, gelatin, or acacia; or lubricating agents such as magnesium stearate, stearic acid, or talc.

[0385] Additional pharmaceutical compositions will be evident to those skilled in the art, including formulations involving antigen binding proteins to PCSK9, with or without at least one additional therapeutic agent(s), in sustained- or controlled-delivery formulations. In certain embodiments, techniques for formulating a variety of other sustained- or controlled-delivery means, such as liposome carriers, bio-erodible microparticles or porous beads and depot injections, are also known to those skilled in the art. See for example, PCT Application No. PCT/US93/00829 which describes the controlled release of porous polymeric microparticles for

the delivery of pharmaceutical compositions. In certain embodiments, sustained-release preparations can include semipermeable polymer matrices in the form of shaped articles, *e.g.* films, or microcapsules. Sustained release matrices can include polyesters, hydrogels, polylactides (U.S. 3,773,919 and EP 058,481), copolymers of L-glutamic acid and gamma ethyl-L-glutamate (Sidman *et al.*, Biopolymers, 22:547-556 (1983)), poly (2-hydroxyethyl-methacrylate) (Langer *et al.*, J. Biomed. Mater. Res., 15:167-277 (1981) and Langer, Chem. Tech., 12:98-105 (1982)), ethylene vinyl acetate (Langer *et al.*, *supra*) or poly-D(-)-3-hydroxybutyric acid (EP 133,988). In certain embodiments, sustained release compositions can also include liposomes, which can be prepared by any of several methods known in the art. *See, e.g.*, Eppstein *et al.*, Proc. Natl. Acad. Sci. USA, 82:3688-3692 (1985); EP 036,676; EP 088,046 and EP 143,949.

[0386] The pharmaceutical composition to be used for *in vivo* administration typically is sterile. In certain embodiments, this can be accomplished by filtration through sterile filtration membranes. In certain embodiments, where the composition is lyophilized, sterilization using this method can be conducted either prior to or following lyophilization and reconstitution. In certain embodiments, the composition for parenteral administration can be stored in lyophilized form or in a solution. In certain embodiments, parenteral compositions generally are placed into a container having a sterile access port, for example, an intravenous solution bag or vial having a stopper pierceable by a hypodermic injection needle.

[0387] In certain embodiments, once the pharmaceutical composition has been formulated, it can be stored in sterile vials as a solution, suspension, gel, emulsion, solid, or as a dehydrated or lyophilized powder. In certain embodiments, such formulations can be stored either in a ready-to-use form or in a form (*e.g.*, lyophilized) that is reconstituted prior to administration.

[0388] In certain embodiments, kits are provided for producing a single-dose administration unit. In certain embodiments, the kit can contain both a first container having a dried protein and a second container having an aqueous formulation. In certain embodiments, kits containing single and multi-chambered pre-filled syringes (*e.g.*, liquid syringes and lyosyringes) are included.

[0389] In certain embodiments, the effective amount of a pharmaceutical composition comprising an antigen binding protein to PCSK9, with or without at least one additional

therapeutic agent, to be employed therapeutically will depend, for example, upon the therapeutic context and objectives. One skilled in the art will appreciate that the appropriate dosage levels for treatment, according to certain embodiments, will thus vary depending, in part, upon the molecule delivered, the indication for which an antigen binding protein to PCSK9, with or without at least one additional therapeutic agent, is being used, the route of administration, and the size (body weight, body surface or organ size) and/or condition (the age and general health) of the patient. In certain embodiments, the clinician can titer the dosage and modify the route of administration to obtain the optimal therapeutic effect. In certain embodiments, a typical dosage can range from about 0.1 µg/kg to up to about 100 mg/kg or more, depending on the factors mentioned above. In certain embodiments, the dosage can range from 0.1 µg/kg up to about 100 mg/kg; or 1 µg/kg up to about 100 mg/kg; or 5 µg/kg up to about 100 mg/kg.

[0390] In certain embodiments, the frequency of dosing will take into account the pharmacokinetic parameters of an antigen binding protein to PCSK9and/or any additional therapeutic agents in the formulation used. In certain embodiments, a clinician will administer the composition until a dosage is reached that achieves the desired effect. In certain embodiments, the composition can therefore be administered as a single dose, or as two or more doses (which may or may not contain the same amount of the desired molecule) over time, or as a continuous infusion via an implantation device or catheter. Further refinement of the appropriate dosage is routinely made by those of ordinary skill in the art and is within the ambit of tasks routinely performed by them. In certain embodiments, appropriate dosages can be ascertained through use of appropriate dose-response data. In some embodiments, the amount and frequency of administration can take into account the desired cholesterol level (serum and/or total) to be obtained and the subject's present cholesterol level, LDL level, and/or LDLR levels, all of which can be obtained by methods that are well known to those of skill in the art.

[0391] In certain embodiments, the route of administration of the pharmaceutical composition is in accord with known methods, *e.g.* orally, through injection by intravenous, intraperitoneal, intracerebral (intra-parenchymal), intracerebroventricular, intramuscular, subcutaneously, intra-ocular, intraarterial, intraportal, or intralesional routes; by sustained release systems or by implantation devices. In certain embodiments, the compositions can be administered by bolus injection or continuously by infusion, or by implantation device.

[0392] In certain embodiments, the composition can be administered locally via implantation of a membrane, sponge or another appropriate material onto which the desired molecule has been absorbed or encapsulated. In certain embodiments, where an implantation device is used, the device can be implanted into any suitable tissue or organ, and delivery of the desired molecule can be via diffusion, timed-release bolus, or continuous administration.

[0393] In certain embodiments, it can be desirable to use a pharmaceutical composition comprising an antigen binding protein to PCSK9, with or without at least one additional therapeutic agent, in an *ex vivo* manner. In such instances, cells, tissues and/or organs that have been removed from the patient are exposed to a pharmaceutical composition comprising an antigen binding protein to PCSK9, with or without at least one additional therapeutic agent, after which the cells, tissues and/or organs are subsequently implanted back into the patient.

[0394] In certain embodiments, an antigen binding protein to PCSK9 and/or any additional therapeutic agents can be delivered by implanting certain cells that have been genetically engineered, using methods such as those described herein, to express and secrete the polypeptides. In certain embodiments, such cells can be animal or human cells, and can be autologous, heterologous, or xenogeneic. In certain embodiments, the cells can be immortalized. In certain embodiments, in order to decrease the chance of an immunological response, the cells can be encapsulated to avoid infiltration of surrounding tissues. In certain embodiments, the encapsulation materials are typically biocompatible, semi-permeable polymeric enclosures or membranes that allow the release of the protein product(s) but prevent the destruction of the cells by the patient's immune system or by other detrimental factors from the surrounding tissues.

[0395] Based on the ability of ABPs to significantly neutralize PCSK9 activity (as demonstrated in the Examples below), these ABPs will have therapeutic effects in treating and preventing symptoms and conditions resulting from PCSK9-mediated activity, such as hypercholesterolemia.

Diagnostic Applications

[0396] In some embodiments, the ABP is used as a diagnostic tool. The ABP can be used to assay the amount of PCSK9 present in a sample and/or subject. As will be appreciated by one of skill in the art, such ABPs need not be neutralizing ABPs. In some embodiments, the

diagnostic ABP is not a neutralizing ABP. In some embodiments, the diagnostic ABP binds to a different epitope than the neutralizing ABP binds to. In some embodiments, the two ABPs do not compete with one another.

[0397] In some embodiments, the ABPs disclosed herein are used or provided in an assay kit and/or method for the detection of PCSK9 in mammalian tissues or cells in order to screen/diagnose for a disease or disorder associated with changes in levels of PCSK9. The kit comprises an ABP that binds PCSK9 and means for indicating the binding of the ABP with PCSK9, if present, and optionally PCSK9 protein levels. Various means for indicating the presence of an ABP can be used. For example, fluorophores, other molecular probes, or enzymes can be linked to the ABP and the presence of the ABP can be observed in a variety of ways. The method for screening for such disorders can involve the use of the kit, or simply the use of one of the disclosed ABPs and the determination of whether the ABP binds to PCSK9 in a sample. As will be appreciated by one of skill in the art, high or elevated levels of PCSK9 will result in larger amounts of the ABP binding to PCSK9 in the sample. Thus, degree of ABP binding can be used to determine how much PCSK9 is in a sample. Subjects or samples with an amount of PCSK9 that is greater than a predetermined amount (*e.g.*, an amount or range that a person without a PCSK9 related disorder would have) can be characterized as having a PCSK9 mediated disorder. In some embodiments, the ABP is administered to a subject taking a statin, in order to determine if the statin has increased the amount of PCSK9 in the subject.

[0398] In some embodiments, the ABP is a non-neutralizing ABP and is used to determine the amount of PCSK9 in a subject receiving an ABP and/or statin treatment.

EXAMPLES

[0399] The following examples, including the experiments conducted and results achieved, are provided for illustrative purposes only and are not to be construed as limiting the present invention.

EXAMPLE 1

Immunization and Titering

Generation of Anti-PCSK9 Antibodies and Hybridomas

[0400] Antibodies to the mature form of PCSK9 (depicted as the sequence in FIG. 1A, with the pro-domain underlined), were raised in XenoMouse® mice (Abgenix, Fremont, CA), which are mice containing human immunoglobulin genes. Two groups of XenoMouse® mice, group 1 and 2, were used to produce antibodies to PCSK9. Group 1 included mice of the XenoMouse® strain XMG2-KL, which produces fully human $IgG2_\kappa$ and $IgG2\lambda$ antibodies. Group 1 mice were immunized with human PCSK9. PCSK9 was prepared using standard recombinant techniques using the GenBank sequence as reference (NM_174936). Group 2 involved mice of the XenoMouse® strain XMG4-KL, which produce fully human $IgG4_\kappa$ and $IgG4\lambda$ antibodies. Group 2 mice were also immunized with human PCSK9.

[0401] The mice of both groups were injected with antigen eleven times, according to the schedule in Table 3. In the initial immunizations, each mouse was injected with a total of 10 μg of antigen delivered intraperitoneally into the abdomen. Subsequent boosts are 5ug doses and injection method is staggered between intraperitoneal injections into the abdomen and sub-cutaneous injections at the base of the tail. For intraperitoneal injections antigen is prepared as an emulsion with TiterMax® Gold (Sigma, Cat # T2684) and for subcutaneous injections antigen is mixed with Alum (aluminum phosphate) and CpG oligos. In injections 2 through 8 and 10, each mouse was injected with a total of 5 μg of antigen in the adjuvant alum gel. A final injection of 5 μg of antigen per mouse is delivered in Phospho buffered saline and delivered into 2 sites 50% IP into the abdomen and 50% SQ at the base of tail. The immunization programs are summarized in Table 3, shown below.

TABLE 3

mouse strain	XMG2/kl	XMG4/kl
# of animals	10	10
immunogen	PCSK9-V5/His	PCSK9-V5/His
1st boost	IP injection 10ug each	IP injection 10ug each
	Titermax Gold	Titermax Gold
2nd boost	tail injection 5ug each	tail injection 5ug each
	Alum/CpG ODN	Alum/CpG ODN
3rd boost	IP injection 5ug each	IP injection 5ug each

	Titermax Gold	Titermax Gold
4th boost	tail injection 5ug each	tail injection 5ug each
	Alum/CpG ODN	Alum/CpG ODN
5th boost	IP injection 5ug each	IP injection 5ug each
	Titermax Gold	Titermax Gold
6th boost	tail injection 5ug each	tail injection 5ug each
	Alum/CpG ODN	Alum/CpG ODN
7th boost	IP injection 5ug each	IP injection 5ug each
	Titermax Gold	Titermax Gold
8th boost	tail injection 5ug each	tail injection 5ug each
	Alum/CpG ODN	Alum/CpG ODN
bleed		
9th boost	IP injection 5ug each	IP injection 5ug each
	Titermax Gold	Titermax Gold
10th boost	tail injection 5ug each	tail injection 5ug each
	Alum/CpG ODN	Alum/CpG ODN
11th boost	BIP 5ug each	BIP 5ug each
	PBS	PBS
harvest		

[0402] The protocol used to titer the XenoMouse animals was as follows: Costar 3368 medium binding plates were coated with neutravadin @ 8ug/ml (50ul/well) and incubated at 4°C in 1XPBS/0.05% azide overnight. They were washed using TiterTek 3-cycle wash with RO water. Plates were blocked using 250ul of 1XPBS/1%milk and incubated for at least 30 minutes at RT. Block was washed off using TiterTek 3-cycle wash with RO water. One then captured b-human PCSK9 @ 2ug/ml in 1XPBS/1%milk/10mM Ca2+ (assay diluent) 50ul/well and incubated for 1hr at RT. One then washed using TiterTek 3-cycle wash with RO water. For the primary antibody, sera was titrated 1:3 in duplicate from 1:100. This was done in assay diluent 50ul/well and incubated for 1hr at RT. One then washed using TiterTek 3-cycle wash with RO water. The secondary antibody was goat anti Human IgG Fc HRP @ 400 ng/ml in assay diluent at 50ul/well. This was incubated for 1hr at RT. This was then washed using TiterTek 3-cycle wash with RO water and patted dry on paper towels. For the substrate, one-step

TMB solution (Neogen, Lexington, Kentucky) was used (50ul/well) and it was allowed to develop for 30 min at RT.

[0403] The protocols followed in the ELISA assays was as follows: For samples comprising b-PCSK9 with no V5His tag the following protocol was employed: Costar 3368 medium binding plates (Corning Life Sciences) were employed. The plates were coated with neutravadin at 8 µg/ml in 1XPBS/0.05%Azide, (50 µl/well). The plates were incubated at 4°C overnight. The plates were then washed using a Titertek M384 plate washer (Titertek, Huntsville, AL). A 3-cycle wash was perfomed. The plates were blocked with 250 µl of 1XPBS/1% milk and incubated approximately 30 minutes at room temperature. The plates were then washed using the M384 plate washer. A 3-cycle wash was perfomed. The capture was b-hu PCSK9, without a V5 tag, and was added at 2 µg/ml in 1XPBS/1%milk/10mM Ca^{2+} (40 µl/well). The plates were then incubated for 1 hour at room temperature. A 3-cycle wash was perfomed. Sera were titrated 1:3 in duplicate from 1:100, and row H was blank for sera. The titration was done in assay diluent, at a volume of 50 µl/well. The plates were incubated for 1 hour at room temperature. Next, a 3-cycle wash was perfomed. Goat anti Human IgG Fc HRP at 100 ng/ml (1:4000) in 1XPBS/1%milk/10mM Ca^{2+} (50 µl/well) was added to the plate and was incubated 1 hour at room temperature. The plates were washed once again, using a 3-cycle wash. The plates were then patted dry with paper towel. Finally, 1 step TMB (Neogen, Lexington, Kentucky) (50 µl/well) was added to the plate and was quenched with 1N hydrochloric acid (50 µl/well) after 30 minutes at room temperature. OD's were read immediately at 450 nm using a Titertek plate reader.

[0404] Positive controls to detect plate bound PCSK9 were soluble LDL receptor (R&D Systems, Cat #2148LD/CF) and a polyclonal rabbit anti-PCSK9 antibody (Caymen Chemical #10007185) titrated 1:3 in duplicate from 3 µg/ml in assay diluent. LDLR was detected with goat anti LDLR (R&D Systems, Cat #AF2148) and rabbit anti goat IgGFc HRP at a concentration of 400 ng/ml; the rabbit polyclonal was detected with goat anti-rabbit IgG Fc at a concentration of 400 ng/ml in assay diluent. Negative control was naive XMG2-KL and XMG4-KL sera titrated 1:3 in duplicate from 1:100 in assay diluent.

[0405] For samples comprising b-PCSK9 with a V5His tag the following protocol was employed: Costar 3368 medium binding plates (Corning Life Sciences) were employed. The plates were coated with neutravadin at 8 µg/ml in 1XPBS/0.05%Azide, (50 µl/well). The

plates were incubated at 4°C overnight. The plates were then washed using a Titertek M384 plate washer (Titertek, Huntsville, AL). A 3-cycle wash was perfomed. The plates were blocked with 250 µl of 1XPBS/1% milk and incubated approximately 30 minutes at room temperature. The plates were then washed using the M384 plate washer. A 3-cycle wash was perfomed. The capture was b-hu PCSK9, with a V5 tag, and was added at 2 µg/ml in 1XPBS/1%milk/10mM Ca^{2+} (40 µl/well). The plates were then incubated for 1 hour at room temperature. A 3-cycle wash was perfomed. Sera were titrated 1:3 in duplicate from 1:100, and row H was blank for sera. The titration was done in assay diluent, at a volume of 50 µl/well. The plates were incubated for 1 hour at room temperature. Next, the plates were washed using the M384 plate washer operated using a 3-cycle wash. Goat anti Human IgG Fc HRP at 400 ng/ml in 1XPBS/1%milk/10mM Ca^{2+} was added at 50 µl/well to the plate and the plate was incubated 1 hour at room temperature. The plates were washed once again, using a 3-cycle wash. The plates were then patted dry with paper towel. Finally, 1 step TMB (Neogen, Lexington, Kentucky) (50 µl/well) was added to the plate and the plate was quenched with 1N hydrochloric acid (50 µl/well) after 30 minutes at room temperature. OD's were read immediately at 450 nm using a Titertek plate reader.

[0406] Positive control was LDLR, rabbit anti-PCSK9 titrated 1:3 in duplicate from 3 µg/ml in assay diluent. LDLR detect with goat anti-LDLR (R&D Systems, Cat #AF2148) and rabbit anti-goat IgG Fc HRP at a concentration of 400 ng/ml; rabbit poly detected with goat anti-rabbit IgG Fc at a concentration of 400 ng/ml in assay diluent. Human anti-His 1.2,3 and anti-V5 1.7.1 titrated 1:3 in duplicate from 1 µg/ml in assay diluent; both detected with goat anti-human IgG Fc HRP at a concentration of 400 ng/ml in assay diluent. Negative control was naive XMG2-KL and XMG4-KL sera titrated 1:3 in duplicate from 1:100 in assay diluent.

[0407] Titers of the antibody against human PCSK9 were tested by ELISA assay for mice immunized with soluble antigen as described. Table 4 summarizes the ELISA data and indicates that there were some mice which appeared to be specific for PCSK9. *See, e.g.*, Table 4. Therefore, at the end of the immunization program, 10 mice (in bold in Table 4) were selected for harvest, and splenocytes and lymphocytes were isolated from the spleens and lymph nodes respectively, as described herein.

TABLE 4

Summary of ELISA Results

	Animal ID	Titer b-hu PCSK9 (V5His) @ 2ug/ml	Titer b-hu PCSK9 @ 2ug/ml
Group 1 - IgG2k/l	P175807	>72900 @ OD 2.2	68359
	P175808	>72900 @ OD 2.3	>72900 @ OD 2.5
	P175818	>72900 @ OD 3.2	**>72900 @ OD 3.0**
	P175819	>72900 @ OD 3.4	**>72900 @ OD 3.2**
	P175820	>72900 @ OD 2.4	>72900 @ OD 2.5
	P175821	>72900 @ OD 3.4	**>72900 @ OD 3.0**
	P175830	>72900 @ OD 2.6	>72900 @ OD 2.5
	P175831	>72900 @ OD 3.1	**>72900 @ OD 3.1**
	P175832	>72900 @ OD 3.8	**>72900 @ OD 3.6**
	P175833	>72900 @ OD 2.6	>72900 @ OD 2.3
Group 2 - IgG4k/l	P174501	19369	17109
	P174503	31616	**23548**
	P174508	48472	**30996**
	P174509	23380	21628
	P174510	15120	9673
	P175773	19407	15973
	P175774	54580	**44424**
	P175775	60713	**55667**
	P175776	30871	**22899**
	P175777	16068	12532
	Naïve G2	< 100 @ OD 0.54	< 100 @ OD 0.48
	Naïve G4	< 100 @ OD 1.57	< 100 @ OD 1.32

EXAMPLE 2

Recovery of Lymphocytes, B-cell Isolations, Fusions and Generation of Hybridomas

[0408] This example outlines how the immune cells were recovered and the hybridomas were generated. Selected immunized mice were sacrificed by cervical dislocation and the draining lymph nodes were harvested and pooled from each cohort. The B cells were dissociated from lymphoid tissue by grinding in DMEM to release the cells from the tissues, and the cells were suspended in DMEM. The cells were counted, and 0.9 ml DMEM per 100 million lymphocytes was added to the cell pellet to resuspend the cells gently but completely.

[0409] Lymphocytes were mixed with nonsecretory myeloma P3X63Ag8.653 cells purchased from ATCC, cat.# CRL 1580 (Kearney *et al.*, (1979) *J. Immunol.* 123, 1548-1550) at a ratio of 1:4. The cell mixture was gently pelleted by centrifugation at 400 x g 4 min. After decanting of the supernatant, the cells were gently mixed using a 1 ml pipette. Preheated

PEG/DMSO solution from Sigma (cat# P7306) (1 ml per million of B-cells) was slowly added with gentle agitation over 1 min followed by 1 min of mixing. Preheated IDMEM (2 ml per million of B cells) (DMEM without glutamine, L-glutamine, pen/strep, MEM non-essential amino acids (all from Invitrogen), was then added over 2 minutes with gentle agitation. Finally preheated IDMEM (8 ml per 10^6 B-cells) was added over 3 minutes.

[0410] The fused cells were spun down 400 x g 6 min and resuspended in 20 ml selection media (DMEM (Invitrogen), 15 % FBS (Hyclone), supplemented with L-glutamine, pen/strep, MEM Non-essential amino acids, Sodium Pyruvate, 2-Mercaptoethanol (all from Invitrogen), HA-Azaserine Hypoxanthine and OPI (oxaloacetate, pyruvate, bovine insulin) (both from Sigma) and IL-6 (Boehringer Mannheim)) per million B-cells. Cells were incubated for 20-30 min at 37C and then resuspended in 200 ml selection media and cultured for 3-4 days in T175 flask prior to 96 well plating. Thus, hybridomas that produced antigen binding proteins to PCSK9 were produced.

EXAMPLE 3

Selection of PCSK9 Antibodies

[0411] The present example outlines how the various PCSK9 antigen binding proteins were characterized and selected. The binding of secreted antibodies (produced from the hybridomas produced in Examples 1 and 2) to PCSK9 was assessed. Selection of antibodies was based on binding data and inhibition of PCSK9 binding to LDLR and affinity. Binding to soluble PCSK9 was analyzed by ELISA, as described below. BIAcore® (surface plasmon resonance) was used to quantify binding affinity.

Primary Screen

[0412] A primary screen for antibodies which bind to wild-type PCSK9 was performed. The primary screen was performed on two harvests. The primary screen comprised an ELISA assay and was performed using the following protocol:

[0413] Costar 3702 medium binding 384 well plates (Corning Life Sciences) were employed. The plates were coated with neutravadin at a concentration of 4 µg/ml in 1XPBS/0.05%Azide, at a volume of 40 µl/well. The plates were incubated at 4°C overnight. The plates were then washed using a Titertek plate washer (Titertek, Huntsville, AL). A 3-cycle wash was perfomed. The plates were blocked with 90 µl of 1XPBS/1%milk and incubated

approximately 30 minutes at room temperature. The plates were then washed. Again, a 3-cycle wash was perfomed. The capture sample was biotinylated-PCSK9, without a V5 tag, and was added at 0.9 μg/ml in 1XPBS/1%milk/10mM Ca^{2+} at a volume of 40 μl/well. The plates were then incubated for 1 hour at room temperature. Next, the plates were washed using the Titertek plate washer operated using a 3-cycle wash. 10 μl of supernatant was transferred into 40 μl of 1XPBS/1%milk/10mM Ca^{2+} and incubated 1.5 hours at room temperature. Again the plates were washed using the Titertek plate washer operated using a 3-cycle wash. 40 μl/well of Goat anti-Human IgG Fc POD at a concentration of 100 ng/ml (1:4000) in 1XPBS/1%milk/10mM Ca^{2+} was added to the plate and was incubated 1 hour at room temperature. The plates were washed once again, using a 3-cycle wash. Finally, 40 μl/well of One-step TMB (Neogen, Lexington, Kentucky) was added to the plate and quenching with 40 μl/well of 1N hydrochloric acid was performed after 30 minutes at room temperature. OD's were read immediately at 450 nm using a Titertek plate reader.

[0414] The primary screen resulted in a total of 3104 antigen specific hybridomas being identified from the two harvests. Based on highest ELISA OD, 1500 hybridomas per harvest were advanced for a total of 3000 positives.

Confirmatory Screen

[0415] The 3000 positives were then rescreened for binding to wild-type PCSK9 to confirm stable hybridomas were established. The screen was performed as follows: Costar 3702 medium binding 384 well plates (Corning Life Sciences) were employed. The plates were coated with neutravadin at 3 μg/ml in 1XPBS/0.05%Azide at a volume of 40 μl/well. The plates were incubated at 4°C overnight. The plates were then washed using a Titertek plate washer (Titertek, Huntsville, AL). A 3-cycle wash was perfomed. The plates were blocked with 90 μl of 1XPBS/1%milk and incubated approximately 30 minutes at room temperature. The plates were then washed using the M384 plate washer. A 3-cycle wash was perfomed. The capture sample was b-PCSK9, without a V5 tag, and was added at 0.9 μg/ml in 1XPBS/1%milk/10mM Ca^{2+} at a volume of 40 μl/well. The plates were then incubated for 1 hour at room temperature. Next, the plates were washed using a 3-cycle wash. 10 μl of supernatant was transferred into 40 μl of 1XPBS/1%milk/10mM Ca^{2+} and incubated 1.5 hours at room temperature. Again the

plates were washed using the Titertek plate washer operated using a 3-cycle wash. 40 µl/well of Goat anti-Human IgG Fc POD at a concentration of 100 ng/ml (1:4000) in 1XPBS/1%milk/10mM Ca^{2+} was added to the plate, and the plate was incubated 1 hour at room temperature. The plates were washed once again, using the Titertek plate washer operated using a 3-cycle wash. Finally, 40 µl/well of One-step TMB (Neogen, Lexington, Kentucky) was added to the plate and was quenched with 40 µl/well of 1N hydrochloric acid after 30 minutes at room temperature. OD's were read immediately at 450 nm using a Titertek plate reader. A total of 2441 positives repeated in the second screen. These antibodies were then used in the subsequent screenings.

Mouse Cross-reactivity Screen

[0416] The panel of hybridomas was then screened for cross-reactivity to mouse PCSK9 to make certain that the antibodies could bind to both human and mouse PCSK9. The following protocol was employed in the cross-reactivity screen: Costar 3702 medium binding 384 well plates (Corning Life Sciences) were employed. The plates were coated with neutravadin at 3 µg/ml in 1XPBS/0.05%Azide at a volume of 40 µl/well. The plates were incubated at 4°C overnight. The plates were then washed using a Titertek plate washer (Titertek, Huntsville, AL). A 3-cycle wash was perfomed. The plates were blocked with 90 µl of 1XPBS/1%milk and incubated approximately 30 minutes at room temperature. The plates were then washed using the Titertek plate washer. A 3-cycle wash was perfomed. The capture sample was biotinylated-mouse PCSK9, and was added at 1 µg/ml in 1XPBS/1%milk/10mM Ca^{2+} at a volume of 40 µl/well. The plates were then incubated for 1 hour at room temperature. Next, the plates were washed using the Titertek plate washer operated using a 3-cycle wash. 50 µl of supernatant was transferred to the plates and incubated 1 hour at room temperature. Again the plates were washed using a 3-cycle wash. 40 µl/well of Goat anti-Human IgG Fc POD at a concentration of 100 ng/ml (1:4000) in 1XPBS/1%milk/10mM Ca^{2+} was added to the plate and the plate was incubated 1 hour at room temperature. The plates were washed once again, using a 3-cycle wash. Finally, 40 µl/well One-step TMB (Neogen, Lexington, Kentucky) was added to the plate and was quenched with 40 µl/well of 1N hydrochloric acid after 30 minutes at room temperature. OD's were read immediately at 450 nm using a Titertek plate reader. 579

antibodies were observed to cross-react with mouse PCSK9. These antibodies were then used in the subsequent screenings.

D374Y Mutant Binding Screen

[0417] The D374Y mutation in PCSK9 has been documented in the human population (*e.g.*, Timms KM et al, "A mutation in PCSK9 causing autosomal-dominant hypercholesterolemia in a Utah pedigree", Hum. Genet. 114: 349-353, 2004). In order to determine if the antibodies were specific for the wild type or also bound to the D374Y form of PCSK9, the samples were then screened for binding to the mutant PCSK9 sequence comprising the mutation D374Y. The protocol for the screen was as follows: Costar 3702 medium binding 384 well plates (Corning Life Sciences) were employed in the screen. The plates were coated with neutravadin at 4 µg/ml in 1XPBS/0.05% Azide at a volume of 40 µl/well. The plates were incubated at 4°C overnight. The plates were then washed using a Titertek plate washer (Titertek, Huntsville, AL). A 3-cycle wash was perfomed. The plates were blocked with 90 µl of 1XPBS/1%milk and incubated approximately 30 minutes at room temperature. The plates were then washed using the Titertek plate washer. A 3-cycle wash was perfomed. The plates were coated with biotinylated human PCSK9 D374Y at a concentration of 1 µg/ml in 1XPBS/1%milk/10mMCa^{2+} and incubated for 1 hour at room temperature. The plates were then washed using a Titertek plate washer. A 3-cycle wash was perfomed. Late exhaust hybridoma culture supernatant was diluted 1:5 in PBS/milk/Ca^{2+} (10 ml plus 40 ml) and incubated for 1 hour at room temperature. Next, 40 µl/well of rabbit anti-human PCSK9 (Cayman Chemical) and human anti-His 1.2.3 1:2 at 1ug/ml in 1XPBS/1%milk/10mMCa^{2+} was titrated onto the plates, which were then incubated for 1 hour at room temperature. The plates were then washed using a Titertek plate washer. A 3-cycle wash was perfomed. 40 µl/well of Goat anti-Human IgG Fc HRP at a concentration of 100 ng/ml (1:4000) in 1XPBS/1%milk/10mM Ca^{2+} was added to the plate and the plate was incubated 1 hour at room temperature. 40 µl/well of Goat anti-rabbit IgG Fc HRP at a concentration of 100 ng/ml (1:4000) in 1XPBS/1%milk/10mM Ca^{2+} was added to the plate and the plate was incubated 1 hour at room temperature. The plates were then washed using a Titertek plate washer. A 3-cycle wash was perfomed. Finally, 40 µl/well of One-step TMB (Neogen, Lexington, Kentucky) was added to the plate and was quenched with 40 µl/well of 1N hydrochloric acid after 30 minutes at room temperature. OD's were read

immediately at 450 nm using a Titertek plate reader. Over 96% of the positive hits on the wild-type PCSK9 also bound mutant PCSK9.

Large Scale Receptor Ligand Blocking Screen

[0418] To screen for the antibodies that block PCSK9 binding to LDLR an assay was developed using the D374Y PCSK9 mutant. The mutant was used for this assay because it has a higher binding affinity to LDLR allowing a more sensitive receptor ligand blocking assay to be developed. The following protocol was employed in the receptor ligand blocking screen: Costar 3702 medium binding 384 well plates (Corning Life Sciences) were employed in the screen. The plates were coated with goat anti-LDLR (R&D Cat #AF2148) at 2 µg/ml in 1XPBS/0.05%Azide at a volume of 40 µl/well. The plates were incubated at 4°C overnight. The plates were then washed using a Titertek plate washer (Titertek, Huntsville, AL). A 3-cycle wash was performed. The plates were blocked with 90 µl of 1XPBS/1% milk and incubated approximately 30 minutes at room temperature. The plates were then washed using the Titertek plate washer. A 3-cycle wash was performed. The capture sample was LDLR (R&D, Cat #2148LD/CF), and was added at 0.4 µg/ml in 1XPBS/1%milk/10mM Ca^{2+} at a volume of 40 µl/well. The plates were then incubated for 1 hour and 10 minutes at room temperature. Contemporaneously, 20 ng/ml of biotinylated human D374Y PCSK9 was incubated with 15 microliters of hybridoma exhaust supernatant in Nunc polypropylene plates and the exhaust supernatant concentration was diluted 1:5. The plates were then pre-incubated for about 1 hour and 30 minutes at room temperature. Next, the plates were washed using the Titertek plate washer operated using a 3-cycle wash. 50 µl/well of the pre-incubated mixture was transferred onto the LDLR coated ELISA plates and incubated for 1 hour at room temperature. To detect LDLR-bound b-PCSK9, 40 µl/well streptavidin HRP at 500 ng/ml in assay diluent was added to the plates. The plates were incubated for 1 hour at room temperature. The plates were again washed using a Titertek plate washer. A 3-cycle wash was performed. Finally, 40 µl/well of One-step TMB (Neogen, Lexington, Kentucky) was added to the plate and was quenched with 40 µl/well of 1N hydrochloric acid after 30 minutes at room temperature. OD's were read immediately at 450 nm using a Titertek plate reader. The screen identified 384 antibodies that blocked the interaction between PCSK9 and the LDLR well, 100 antibodies blocked the interaction strongly (OD < 0.3).

These antibodies inhibited the binding interaction of PCSK9 and LDLR greater than 90% (greater than 90% inhibition).

Receptor Ligand Binding Assay on Blocker Subset

[0419] The receptor ligand assay was then repeated using the mutant enzyme on the 384 member subset of neutralizers identified in the first large scale receptor ligand inhibition assay. The same protocol was employed in the screen of the 384 member blocker subset assay as was done in the large scale receptor ligand blocking screen. This repeat screen confirmed the initial screening data.

[0420] This screen of the 384 member subset identified 85 antibodies that blocked interaction between the PCSK9 mutant enzyme and the LDLR greater than 90%.

Receptor Ligand Binding Assay of Blockers that Bind the Wild Type PCSK9 but not the D374Y Mutant

[0421] In the initial panel of 3000 sups there were 86 antibodies shown to specifically bind to the wild-type PCSK9 and not to the huPCSK9(D374Y) mutant. These 86 sups were tested for the ability to block wild-type PCSK9 binding to the LDLR receptor. The following protocol was employed: Costar 3702 medium binding 384 well plates (Corning Life Sciences) were employed in the screen. The plates were coated with anti-His 1.2.3 at 10 µg/ml in 1XPBS/0.05% Azide at a volume of 40 µl/well. The plates were incubated at 4°C overnight. The plates were then washed using a Titertek plate washer (Titertek, Huntsville, AL). A 3-cycle wash was perfomed. The plates were blocked with 90 µl of 1XPBS/1%milk and incubated approximately 30 minutes at room temperature. The plates were then washed using the Titertek plate washer. A 3-cycle wash was perfomed. LDLR (R&D Systems, #2148LD/CF or R&D Systems, #2148LD) was added at 5 µg/ml in 1XPBS/1%milk/10mM Ca^{2+} at a volume of 40 µl/well. The plates were then incubated for 1 hour at room temperature. Next, the plates were washed using the Titertek plate washer operated using a 3-cycle wash. Contemporaneously, biotinylated human wild-type PCSK9 was pre-incubated with hybridoma exhaust supernatant in Nunc polypropylene plates. 22 µl of hybridoma sup was transferred into 33ul of b-PCSK9 at a concentration of 583 ng/ml in 1XPBS/1%milk/10mMCa2+, giving a final b-PCSK9 concentration = 350 ng/ml and the exhaust supernatant at a final dilution of 1:2.5. The plates

were pre-incubated for approximately 1 hour and 30 minutes at room temperature. 50 µl/well of the preincubated mixture was transferred onto LDLR captured ELISA plates and incubated for 1 hour at room temperature. The plates were then washed using the Titertek plate washer. A 3-cycle wash was perfomed. 40 µl/well streptavidin HRP at 500 ng/ml in assay diluent was added to the plates. The plates were incubated for 1 hour at room temperature. The plates were then washed using a Titertek plate washer. A 3-cycle wash was perfomed. Finally, 40 µl/well of One-step TMB (Neogen, Lexington, Kentucky) was added to the plate and was quenched with 40 µl/well of 1N hydrochloric acid after 30 minutes at room temperature. OD's were read immediately at 450 nm using a Titertek plate reader.

Screening Results

[0422] Based on the results of the assays described, several hybridoma lines were identified as producing antibodies with desired interactions with PCSK9. Limiting dilution was used to isolate a manageable number of clones from each line. The clones were designated by hybridoma line number (e.g. 21B12) and clone number (e.g. 21B12.1). In general, no difference among the different clones of a particular line were detected by the functional assays described herein. In a few cases, clones were identified from a particular line that behaved differently in the functional assays, for example, 25A7.1 was found not to block PCSK9/LDLR but 25A7.3 (referred to herein as 25A7) was neutralizing. The isolated clones were each expanded in 50-100 ml of hybridoma media and allowed to grow to exhaustion, (i.e., less than about 10% cell viability). The concentration and potency of the antibodies to PCSK9 in the supernatants of those cultures were determined by ELISA and by *in vitro* functional testing, as described herein. As a result of the screening described herein, the hybridomas with the highest titer of antibodies to PCSK9 were identified. The selected hybridomas are shown in FIGS 2A-3D and Table 2.

EXAMPLE 4.1

Production of Human 31H4 IgG4 Antibodies from Hybridomas

[0423] This example generally describes how one of the antigen binding proteins was produced from a hybridoma line. The production work used 50ml exhaust supernatant generation followed by protein A purification. Integra production was for scale up and was performed later. Hybridoma line 31H4 was grown in T75 flasks in 20 ml of media (Integra

Media, Table 5). When the hybridoma was nearly confluent in the T75 flasks, it was transferred to an Integra flask (Integra Biosciences, Integra CL1000, cat# 90 005).

[0424] The Integra flask is a cell culture flask that is divided by a membrane into two chambers, a small chamber and a large chamber. A volume of 20-30 ml hybridoma cells at a minimum cell density of $1x10^6$ cells per ml from the 31H4 hybridoma line was placed into the small chamber of an Integra flask in Integra media (see Table 5 for components of Integra media). Integra media alone (1L) was placed in the large chambers of the Integra flasks. The membrane separating the two chambers is permeable to small molecular weight nutrients but is impermeable to hybridoma cells and to antibodies produced by those cells. Thus, the hybridoma cells and the antibodies produced by those hybridoma cells were retained in the small chamber.

[0425] After one week, media was removed from both chambers of the Integra flask and was replaced with fresh Integra media. The collected media from the small chambers was separately retained. After a second week of growth, the media from the small chamber was again collected. The collected media from week 1 from the hybridoma line was combined with the collected media from week 2 from the hybridoma line. The resulting collected media sample from the hybridoma line was spun to remove cells and debris (15 minutes at 3000rpm) and the resulting supernatant was filtered (0.22um). Clarified conditioned media was loaded onto a Protein A-Sepharose column. Optionally, the media can be first concentrated and then loaded onto a Protein A Sepharose column. Non-specific bindings were removed by an extensive PBS wash. Bound antibody proteins on the Protein A column were recovered by standard acidic antibody elution from Protein A columns (such as 50 mM Citrate, pH 3.0). Aggregated antibody proteins in the Protein A Sepharose pool were removed by size exclusion chromatography or binding ion exchange chromatography on anion exchanger resin such as Q Sepharose resin. The specific IEX conditions for the 31H4 proteins are Q-Sepharose HP at pH 7.8-8.0. Antibody was eluted with a NaCl gradient of 10 mM-500 mM in 25 column volumes.

TABLE 5

Composition of Media

INTEGRA MEDIA
HSFM
10% Ultra Low IgG serum
2mmol/L L-glutamine

1% NEAA
4g/L glucose

EXAMPLE 4.2

Production of Recombinant 31H4 Human IgG2 Antibodies From Transfected Cells

[0426] The present example outlines how 31H4 IgG2 antibodies were produced from transfected cells. 293 cells for transient expression and CHO cells for stable expression were transfected with plasmids that encode 31H4 heavy and light chains. Conditioned media from transfected cells was recovered by removing cells and cell debris. Clarified conditioned media was loaded onto a Protein A-Sepharose column. Optionally, the media can first be concentrated and then loaded onto a Protein A Sepharose column. Non-specific bindings were removed by extensive PBS wash. Bound antibody proteins on the Protein A column were recovered by standard acidic antibody elution from Protein A columns (such as 50 mM citrate, pH 3.0). Aggregated antibody proteins in the Protein A Sepharose pool were removed by size exclusion chromatography or binding ion exchange chromatography on anion exchanger resin such as Q Sepharose resin. The specific IEX conditions for the 31H4 proteins are Q-Sepharose HP at pH 7.8-8.0. The antibody was eluted with a NaCl gradient of 10 mM-500 mM in 25 column volumes.

EXAMPLE 5

Production of Human 21B12 IgG4 Antibodies from Hybridomas

[0427] The present example outlines how antibody 21B12 IgG4 was produced from hybridomas. Hybridoma line 21B12 was grown in T75 flasks in media (Integra Media, Table 5). When the hybridomas were nearly confluent in the T75 flasks, they were transferred to Integra flasks (Integra Biosciences, Integra CL1000, cat# 90 005).

[0428] The Integra flask is a cell culture flask that is divided by a membrane into two chambers, a small chamber and a large chamber. A volume of 20-30 ml hybridoma cells at a minimum cell density of $1x10^6$ cells per ml from the 31H4 hybridoma line was placed into the small chamber of an Integra flask in Integra media (see Table 5 for components of Integra media). Integra media alone (1L) was placed in the large chambers of the Integra flasks. The membrane separating the two chambers is permeable to small molecular weight nutrients but is

impermeable to hybridoma cells and to antibodies produced by those cells. Thus, the hybridoma cells and the antibodies produced by those hybridoma cells were retained in the small chamber. After one week, media was removed from both chambers of the Integra flask and was replaced with fresh Integra media. The collected media from the small chambers was separately retained. After a second week of growth, the media from the small chamber was again collected. The collected media from week 1 from the hybridoma line was combined with the collected media from week 2 from the hybridoma line. The resulting collected media sample from the hybridoma line was spun to remove cells and debris (15 minutes at 3000 rpm) and the resulting supernatant was filtered (0.22 μm). Clarified conditioned media were loaded onto a Protein A Sepharose column. Optionally, the media are first concentrated and then loaded onto a Protein A Sepharose column. Non-specific bindings were removed by an extensive PBS wash. Bound antibody proteins on the Protein A column were recovered by standard acidic antibody elution from Protein A columns (such as 50 mM Citrate, pH 3.0). Aggregated antibody proteins in the Protein A Sepharose pool were removed by size exclusion chromatography or binding ion exchange chromatography on anion exchanger resin such as Q Sepharose resin. The specific IEX conditions for the 21B12 proteins are Q-Sepharose HP at pH 7.8-8.0. The antibody was eluted with a NaCl gradient of 10 mM-500 mM in 25 column volumes.

EXAMPLE 6

Production of Human 21B12 IgG2 Antibodies From Transfected Cells

[0429] The present example outlines how 21B12 IgG2 antibodies were produced from transfected cells. Cells (293 cells for transient expression and CHO cells for stable expression) were transfected with plasmids that encode 21B12 heavy and light chains. Conditioned media from hybridoma cells were recovered by removing cells and cell debris. Clarified conditioned media were loaded onto a Protein A-Sepharose column. Optionally, the media can first be concentrated and then loaded onto a Protein A Sepharose column. Non-specific bindings were removed by extensive PBS wash. Bound antibody proteins on the Protein A column were recovered by standard acidic antibody elution from Protein A columns (50 mM Citrate, pH 3.0). Aggregated antibody proteins in the Protein A Sepharose pool were removed by size exclusion chromatography or binding ion exchange chromatography on cation exchanger

resin such as SP-Sepharose resin. The specific IEX conditions for the 21B12 proteins were SP-Sepharose HP at pH 5.2. Antibodies were eluted with 25 column volumes of buffer that contains a NaCl gradient of 10 mM-500 mM in 20 mM sodium acetate buffer.

EXAMPLE 7

Production of Human 16F12 IgG4 Antibodies from Hybridomas

[0430] The present example outlines how antibody 16F12 IgG4 was produced from hybridomas. Hybridoma line 16F12 was grown in T75 flasks in media (see Table 5). When the hybridomas were nearly confluent in the T75 flasks, they were transferred to Integra flasks (Integra Biosciences, Integra CL1000, cat# 90 005).

[0431] The Integra flask is a cell culture flask that is divided by a membrane into two chambers, a small chamber and a large chamber. A volume of 20-30 ml Hybridoma cells at a minimum cell density of $1x10^6$ cells per ml from the 31H4 hybridoma line was placed into the small chamber of an Integra flask in Integra media (see Table 5 for components of Integra media). Integra media alone (1L) was placed in the large chambers of the Integra flasks. The membrane separating the two chambers is permeable to small molecular weight nutrients but is impermeable to hybridoma cells and to antibodies produced by those cells. Thus, the hybridoma cells and the antibodies produced by those hybridoma cells were retained in the small chamber.

[0432] After one week, media was removed from both chambers of the Integra flask and was replaced with fresh Integra media. The collected media from the small chambers was separately retained. After a second week of growth, the media from the small chamber was again collected. The collected media from week 1 from the hybridoma line was combined with the collected media from week 2 from the hybridoma line. The resulting collected media sample from the hybridoma line were spun to remove cells and debris (15 minutes at 3000 rpm) and the resulting supernatants were filtered (0.22 μm). Clarified conditioned media were loaded onto a Protein A Sepharose column. Optionally, the media can be first concentrated and then loaded onto a Protein A Sepharose column. Non-specific bindings were removed by extensive PBS wash. Bound antibody proteins on the Protein A column were recovered by standard acidic antibody elution from Protein A columns (50 mM Citrate, pH 3.0). Aggregated antibody proteins in the Protein A Sepharose pool were removed by size exclusion chromatography or binding ion exchange chromatography on anion exchanger resin such as Q Sepharose resin. The

specific IEX conditions for the 16F12 proteins are Q Sepharose HP at pH 7.8-8.0. Antibody was eluted with a NaCl gradient of 10 mM-500 mM in 25 column volumes.

EXAMPLE 8

Production of Human 16F12 IgG2 Antibodies From Transfected Cells

[0433] The present example outlines how 16F12 IgG2 antibodies were produced from transfected cells. Cells (293 cells for transient expression and CHO cells for stable expression) were transfected with plasmids that encode 16F12 heavy and light chains. Conditioned media from hybridoma cells were recovered by removing cells and cell debris. Clarified conditioned media were loaded onto a Protein A-Sepharose. Optionally, the media can be first concentrated and then loaded onto a Protein A Sepharose column. Non-specific bindings were removed by extensive PBS wash. Bound antibody proteins on the Protein A column were recovered by standard acidic antibody elution from Protein A columns (50 mM Citrate, pH 3.0). Aggregated antibody proteins in the Protein A Sepharose pool were removed by size exclusion chromatography or binding ion exchange chromatography on cation exchanger resin such as SP Sepharose resin. The specific IEX conditions for the 16F12 proteins are SP Sepharose HP at pH 5.2. Antibody is eluted with 25 column volumes of buffer that contains a NaCl gradient of 10 mM-500 mM in 20 mM sodium acetate buffer.

EXAMPLE 9

Sequence Analysis of Antibody Heavy and Light Chains

[0434] The nucleic acid and amino acid sequences for the light and heavy chains of the above antibodies were then deteremined by Sanger (dideoxy) nucleotide sequencing. Amino acid sequences were then deduced for the nucleic acid sequences. The nucleic acid sequences for the variable domains are depicted in FIG.s 3E-3JJ.

[0435] The cDNA sequences for the lambda light chain variable regions of 31H4, 21B12, and 16F12 were determined and are disclosed as SEQ ID NOs: 153, 95, and 105 respectively.

[0436] The cDNA sequences for the heavy chain variable regions of 31H4, 21B12, and 16F12 were determined and are disclosed as SEQ ID NOs: 152, 94, and 104 respectively.

[0437] The lambda light chain constant region (SEQ ID NO: 156), and the IgG2 and IgG4 heavy chain constant regions (SEQ ID NOs: 154 and 155) are shown in FIG. 3KK.

[0438] The polypeptide sequences predicted from each of those cDNA sequences were determined. The predicted polypeptide sequences for the lambda light chain variable regions of 31H4, 21B12, and 16F12 were predicted and are disclosed as SEQ ID NOs: 12, 23, and 35 respectively, the lambda light chain constant region (SEQ ID NO: 156), the heavy chain variable regions of 31H4, 21B12, and 16F12 were predicted and are disclosed as (SEQ. ID NOs. 67, 49, and 79 respectively. The IgG2 and IgG4 heavy chain constant regions (SEQ ID NOs: 154 and 155).

[0439] The FR1, CDR1, FR2, CDR2, FR3, CDR3, FR4 divisions are shown in FIG 2A-3D.

[0440] Based on the sequence data, the germline genes from which each heavy chain or light chain variable region was derived was determined. The identity of the germline genes are indicated next to the corresponding hybridoma line in FIGs. 2A-3D and each is represented by a unique SEQ ID NO. FIGs. 2A-3D also depict the determined amino acid sequences for additional antibodies that were characterized.

EXAMPLE 8

Determination of Isoelectric Points of Three Antibodies

[0441] The theoretical pIs of the antibodies based on amino acid sequence were determined to be 7.36 for 16F12; 8.47 for 21B12; and 6.84 for 31H4.

EXAMPLE 9

Characterization of Binding of Antibodies to PCSK9

[0442] Having identified a number of antibodies that bind to PCSK9, several approaches were employed to quantify and further characterize the nature of the binding. In one aspect of the study, a Biacore affinity analysis was performed. In another aspect of the study a KinExA® affinity analysis was performed. The samples and buffers employed in these studies are presented in Table 6 below.

TABLE 6

sample	[sample] mg/ml	Buffer	[sample] uM
hPCSK9	1.26	PBS	16.6
mPCSK9-8xHIS	1.44	PBS	18.9
cPCSK9-V5-6xHIS	0.22	PBS	2.9
16F12, anti-PCSK9 huIgG4	4.6	20mM NaOAC, pH 5.2, 50mM NaCl	31.9
21B12, anti-PCSK9 huIgG4	3.84	10mM NAOAC, pH 5.2, 9% Sucrose	27.0
31H4, anti-PCSK9 huIgG4	3.3	10mM NAOAC, pH 5.2, 9% Sucrose	22.9

BIAcore® Affinity Measurements

[0443] A BIAcore® (surface plasmon resonance device, Biacore, Inc., Piscataway, NJ) affinity analysis of the 21B12 antibodies to PCSK9 described in this Example was performed according to the manufacturer's instructions.

[0444] Briefly, the surface plasmon resonance experiments were performed using Biacore 2000 optical biosensors (Biacore, GE Healthcare, Piscataway, NJ). Each individual anti-PCSK9 antibody was immobilized to a research-grade CM5 biosensor chip by amine-coupling at levels that gave a maximum analyte binding response (Rmax) of no more than 200 resonance units (RU). The concentration of PCSK9 protein was varied at 2 fold intervals (the analyte) and was injected over the immobilized antibody surface (at a flow rate of 100 µl/min for 1.5 minutes). Fresh HBS-P buffer (pH 7.4, 0.01 M Hepes, 0.15 M NaCl, 0.005% surfactant P-20, Biacore) supplemented with 0.01% BSA was used as binding buffer. Binding affinities of each anti-PCSK9 antibody were measured in separate experiments against each of the human, mouse, and cynomolgus monkey PCSK9 proteins at pH 7.4 (the concentrations used were 100, 50, 25, 12.5, 6.25, 3.125, and 0 nM).

[0445] In addition, the binding affinities of antibody to human PCSK9 were also measured at pH 6.0 with the pH 6.0 HBS-P buffer (pH 6.0, 0.01 M Hepes, 0.15 M NaCl, 0.005% surfactant P-20, Biacore) supplemented with 0.01% BSA. The binding signal obtained was

proportional to the free PCSK9 in solution. The dissociation equilibrium constant (K_D) was obtained from nonlinear regression analysis of the competition curves using a dual-curve one-site homogeneous binding model (KinExA® software, Sapidyne Instruments Inc., Boise, ID) (n=1 for the 6.0 pH runs). Interestingly, the antibodies appeared to display a tighter binding affinity at the lower pH (where the Kd was 12.5, 7.3, and 29 pM for 31H4, 21B12, and 16F12 respectively).

[0446] Antibody binding kinetic parameters including k_a (association rate constant), k_d (dissociation rate constant), and K_D (dissociation equilibrium constant) were determined using the BIA evaluation 3.1 computer program (BIAcore, Inc. Piscataway, NJ). Lower dissociation equilibrium constants indicate greater affinity of the antibody for PCSK9. The K_D values determined by the BIAcore® affinity analysis are presented in Table 7.1, shown below.

TABLE 7.1

Antibody	hPCSK9	CynoPCSK9	mPCSK9
31H4	210 pM	190 pM	6 nM
21B12	190 pM	360 pM	460 nM
16F12	470 pM	870 pM	6.4 nM

Table 7.2 depicts the k_{on} and k_{off} rates.

TABLE 7.2

-	K_{on} (M-1 s-1)	K_{off} (s-1)	K_D
31H4.1, pH 7.4	2.45 e+5	5.348 e-5	210 pM
31H4.1, pH 6	5.536 e+6	6.936 e-5	12.5 pM
21B12.1, pH 7.4	3.4918 e+4	6.634 e-6	190 pM
21B12.1, pH 6	2.291 e+6	1.676 e-5	7.3 pM
16F12.1, pH 7.4	1.064 e+5	4.983 e-5	470 pM
16F12.1, pH 6	2.392 e+6	7.007 e-5	29 pM

KinExA® Affinity Measurements

[0447] A KinExA® (Sapidyne Instruments, Inc., Boise, ID) affinity analysis of 16F12 and 31H4 was performed according to the manufacturer's instructions. Briefly, Reacti-Gel™ (6x) (Pierce) was pre-coated with one of human, V5-tagged cyno or His-tagged mouse PCSK9 proteins and blocked with BSA. 10 or 100 pM of either antibody 16F12 or antibody 31H4 and one of the PCSK9 proteins was then incubated with various concentrations (0.1 pM – 25 nM) of PCSK9 proteins at room temperature for 8 hours before being passed through the PCSK9-coated beads. The amount of the bead-bound 16F12 or 31H4 was quantified by fluorescently (Cy5) labeled goat anti-human IgG (H+L) antibody (Jackson Immuno Research). The binding signal is proportional to the concentration of free 16F12 or 31H4 at binding equilibrium. Equilibrium dissociation constant (K_D) were obtained from nonlinear regression of the two sets of competition curves using a one-site homogeneous binding model. The KinExA® Pro software was employed in the analysis. Binding curves generated in this analysis are presented as FIGs. 4A-4F.

[0448] Both the 16F12 and 31H4 antibodies showed similar affinity to human and cyno PCSK9, but approximately 10-250 fold lower affinity to mouse PCSK9. Of the two antibodies tested using the KinExA® system, antibody 31H4 showed higher affinity to both human and cyno PCSK9 with 3 and 2 pM K_D, respectively. 16F12 showed slightly weaker affinity at 15pM K_D to human PCSK9 and 16 pM K_D to cyno PCSK9.

[0449] The results of the KinExA® affinity analysis are summarized in Table 8.1, shown below.

TABLE 8.1

	hPCSK9		cPCSK		mPCSK	
Sample	**KD (pM)**	95% CI	**KD (pM)**	95% CI	**KD (pM)**	95% CI
16F12	**15**	11 ~22	**16**	14~19	**223**	106~410
31H4.1	**3**	1~5	**2**	1~3	**500**	400~620

[0450] In addition, a SDS PAGE was run to check the quality and quantity of the samples and is shown in FIG. 5A. cPCSK9 showed around 50% less on the gel and also from the active binding concentration calculated from KinExA® assay. Therefore, the K_D of the mAbs to cPCSK9 was adjusted as 50% of the active cPCSK9 in the present.

[0451] A BIAcore solution equilibirium binding assay was used to measure the Kd values for ABP 21B12. 21B12.1 showed little signal using KinExA assay, therefore, biacore solution equilibrium assay was applied. Since no significant binding was observed on binding of antibodies to immobilized PCSK9 surface, 21B12 antibody was immobilized on the flow cell 4 of a CM5 chip using amine coupling with density around 7000 RU. Flow cell 3 was used as a background control. 0.3, 1, and 3 nM of human PCSK9 or cyno PCSK9 were mixed with a serial dilutions of 21B12.1 antibody samples (ranged from 0.001 ~ 25 nM) in PBS plus 0.1mg/ml BSA, 0.005% P20. Binding of the free PCSK9 in the mixed solutions were measured by injecting over the 21B12.1 antibody surface. 100% PCSK9 binding signal on 21B12.1 surface was determined in the absence of mAb in the solution. A decreased PCSK9 binding response with increasing concentrations of mAb indicated that PCSK9 binding to mAb in solution, which blocked PCSK9 from binding to the immobilized peptibody surface. Plotting the PCSK9 binding signal versus mAb concentrations, K_D was calculated from three sets of curves (0.3, 1 and 3nM fixed PCSK9 concentration) using a one-site homogeneous binding model in KinExA Pro™ software. Although cPCSK9 has lower protein concentration observed from KinExA assay and SDS-gel, its concentration was not adjusted here since the concentration of cPCSK9 was not used for calculation of K_D. The results are displayed in Table 8.2 below and in FIGs. 5B-5D. FIG. 5B depicts the results from the solution equilibrium assay at three different hPCSK9 concentrations for hPCSK9. FIG. 5C depicts a similar set of results for mPCSK9. FIG. 5D depicts the results from the above biacore capture assay.

TABLE 8.2

	hPCSK9		cPCSK		mPCSK	
Sample	KD (pM)	95% CI	KD (pM)	95% CI	KD (pM)	95% CI
21B12.1	15	9~23	11	7~16	17000	-

EXAMPLE 10

Epitope Binning

[0452] Competition ELISA was used for anti-PCSK9 antibody binning. Briefly, to determine if two antibodies belong to the same epitope bin, one of the antibodies (mAb1) was first coated onto an ELISA plate (NUNC) at 2 μg/ml by overnight incubation. The plate was

then washed and blocked with 3% BSA. Meanwhile, 30 ng/ml of biotinylated hPCSK9 was incubated with the second antibody (mAb2) for 2 hours at room temperature. The mixture was applied to coated mAb1 and incubated for 1 hour at room temperature. The ELISA plate was then washed and incubated with Neutravidin-HRP (Pierce) at 1:5000 dilutions for 1 hour. After another wash, the plate was incubated with TMB substrate and signal was detected at 650 nm using a Titertek plate reader. Antibodies with the same binding profiles were grouped together into the same epitope bin. The results of the antibody binning studies are presented in Table 8.3.

TABLE 8.3

Clone	Bin
21B12.2	1
31H4	3
20D10	1
25A7.1	2
25A7.3	1
23G1	1
26H5	1
31D1	1
16F12	3
28D6	3
27A6	3
31G11	3
27B2	ND
28B12	3
22E2	3
1A12.2	1
3B6	1
3C4	4
9C9	1
9H6	1
13B5	6
13H1	7
17C2	1
19H9.2	1
23B5	1
25G4	1
26E10	1
27E7	1
27H5	1
30A4	1
30B9	1

Clone	Bin
31A4	5
31B12	5

[0453] Additional examination of the epitope binning was performed using BIAcore. Three mAbs, 16F12, 21B12 and 31H4, were immobilized on flow cells 2, 3 and 4 with density around 8000 RU. 5 nM PCSK9 from human, mouse and cyno were injected over the mAb surfaces to reach around 100 to 500 RU. 10nM mAbs were then injected over the PCSK9 surface. Binding of three mAbs to three different PCSK9 proteins over the three mAbs were then recorded.

[0454] If the two mAbs had a similar epitope on the antigen, mAb 1 will not show the binding to the antigen already bound to the mAb 2. If the two mAbs have the different epitope on the antigen, mAb1 will show the binding to the antigen bound to the mAb2. FIG. 5E depicts these epitope binning results in graph form for three mAbs on human PCSk9. A similar pattern was observed for mPCSK9 and cPCSK9. As shown in the graph, 16F12 and 31H4 appear to share a similar epitope, while 21B12 appears to have a different epitope.

EXAMPLE 11

Efficacy of 31H4 and 21B12 for Blocking D374Y PCSK9/LDLR Binding

[0455] This example provides the IC50 values for two of the antibodies in blocking PCSK9 D374Y's ability to bind to LDLR. Clear 384 well plates (Costar) were coated with 2 micrograms/ml of goat anti-LDL receptor antibody (R&D Systems) diluted in buffer A (100 mM sodium cacodylate, pH 7.4). Plates were washed thoroughly with buffer A and then blocked for 2 hours with buffer B (1% milk in buffer A). After washing, plates were incubated for 1.5 hours with 0.4 micrograms/ml of LDL receptor (R&D Systems) diluted in buffer C (buffer B supplemented with 10 mM CaCl2). Concurrent with this incubation, 20 ng/ml of biotinylated D374Y PCSK9 was incubated with various concentrations of the 31H4 IgG2, 31H4 IgG4, 21B12 IgG2 or 21B12 IgG4 antibody, which was diluted in buffer A, or buffer A alone (control). The LDL receptor containing plates were washed and the biotinylated D374Y PCSK9/antibody mixture was transferred to them and incubated for 1 hour at room temperature. Binding of the biotinylated D374Y to the LDL receptor was detected by incubation with streptavidin-HRP (Biosource) at 500 ng/ml in buffer C followed by TMB substrate (KPL). The signal was quenched with 1N HCl and the absorbance read at 450 nm.

[0456] The results of this binding study are shown in FIGs. 6A-6D. Summarily, IC_{50} values were determined for each antibody and found to be 199 pM for 31H4 IgG2 (FIG. 6A), 156 pM for 31H4 IgG4 (FIG. 6B), 170 pM for 21B12 IgG2 (FIG. 6C), and 169 pM for 21B12 IgG4 (FIG. 6D).

[0457] The antibodies also blocked the binding of wild-type PCSK9 to the LDLR in this assay.

EXAMPLE 12

Cell LDL Uptake Assay

[0458] This example demonstrates the ability of various antigen binding proteins to reduce LDL uptake by cells. Human HepG2 cells were seeded in black, clear bottom 96-well plates (Costar) at a concentration of $5x10^5$ cells per well in DMEM medium (Mediatech, Inc) supplemented with 10% FBS and incubated at 37°C (5% CO2) overnight. To form the PCSK9 and antibody complex, 2 µg/ml of D374Y human PCSK9 was incubated with various concentrations of antibody diluted in uptake buffer (DMEM with 1% FBS) or uptake buffer alone (control) for 1 hour at room temperature. After washing the cells with PBS, the D374Y PCSK9/antibody mixture was transferred to the cells, followed by LDL-BODIPY (Invitrogen) diluted in uptake buffer at a final concentration of 6 µg/ml. After incubation for 3 hours at 37°C (5% CO2), cells were washed thoroughly with PBS and the cell fluorescence signal was detected by Safire™ (TECAN) at 480-520nm (excitation) and 520-600nm (emission).

[0459] The results of the cellular uptake assay are shown in FIGs. 7A-7D. Summarily, IC_{50} values were determined for each antibody and found to be 16.7 nM for 31H4 IgG2 (FIG. 7A), 13.3 nM for 31H4 IgG4 (FIG. 7B), 13.3 nM for 21B12 IgG2 (FIG. 7C), and 18 nM for 21B12 IgG4 (FIG. 7D). These results demonstrate that the applied antigen binding proteins can reduce the effect of PCSK9 (D374Y) to block LDL updtake by cells The antibodies also blocked the effect of wild-type PCSK9 in this assay.

EXAMPLE 13

Serum cholesterol Lowering Effect of the 31H4 Antibody in 6 Day Study

[0460] In order to assess total serum cholesterol (TC) lowering in wild type (WT) mice via antibody therapy against PCSK9 protein, the following procedure was performed.

[0461] Male WT mice (C57BL/6 strain, aged 9-10 weeks, 17-27 g) obtained from Jackson Laboratory (Bar Harbor, ME) were fed a normal chow (Harland-Teklad, Diet 2918) through out the duration of the experiment. Mice were administered either anti-PCSK9 antibody 31H4 (2 mg/ml in PBS) or control IgG (2 mg/ml in PBS) at a level of 10mg/kg through the mouse's tail vein at T=0. Naïve mice were also set aside as a naïve control group. Dosing groups and time of sacrifice are shown in Table 9.

TABLE 9

Group	Treatment	Time point after dosing	Number
1	IgG	8 hr	7
2	31H4	8 hr	7
3	IgG	24 hr	7
4	31H4	24 hr	7
5	IgG	72 hr	7
6	31H4	72 hr	7
7	IgG	144 hr	7
8	31H4	144 hr	7
9	Naïve	n/a	7

[0462] Mice were sacrificed with CO2 asphyxiation at the pre-determined time points shown in Table 9. Blood was collected via vena cava into eppendorf tubes and was allowed to clot at room temperature for 30 minutes. The samples were then spun down in a table top centrifuge at 12,000xg for 10 minutes to separate the serum. Serum total cholesterol and HDL-C were measured using Hitachi 912 clinical analyzer and Roche/Hitachi TC and HDL-C kits.

[0463] The results of the experiment are shown in FIGs. 8A-8D. Summarily, mice to which antibody 31H4 was administered showed decreased serum cholesterol levels over the course of the experiment (FIG. 8A and FIG. 8B). In addition, it is noted that the mice also showed decreased HDL levels (FIG. 8C and FIG. 8D). For FIG. 8A and FIG. 8C, the percentage change is in relation to the control IgG at the same time point (*P<0.01, # P<0.05). For FIG. 8B and FIG 8D, the percentage change is in relation to total serum cholesterol and HDL levels measured in naïve animals at t=0 hrs (*P<0.01, # P<0.05).

[0464] In respect to the lowered HDL levels, it is noted that one of skill in the art will appreciate that the decrease in HDL in mice is not indicative that an HDL decrease will occur in humans and merely further reflects that the serum cholesterol level in the organism has decreased. It is noted that mice transport the majority of serum cholesterol in high density lipoprotein (HDL) particles which is different to humans who carry most serum cholesterol on LDL particles. In mice the measurement of total serum cholesterol most closely resembles the level of serum HDL-C. Mouse HDL contains apolipoprotein E (apoE) which is a ligand for the LDL receptor (LDLR) and allows it to be cleared by the LDLR. Thus, examining HDL is an appropriate indicator for the present example, in mice (with the understanding that a decrease in HDL is not expected for humans). For example, human HDL, in contrast, does not contain apoE and is not a ligand for the LDLR. As PCSK9 antibodies increase LDLR expression in mouse, the liver can clear more HDL and therefore lowers serum HDL-C levels.

EXAMPLE 14

Effect of Antibody 31H4 on LDLR Levels in a 6 Day Study

[0465] The present example demonstrates that an antigen binding protein alters the level of LDLR in a subject, as predicted, over time. A Western blot analysis was performed in order to ascertain the effect of antibody 31H4 on LDLR levels. 50-100 mg of liver tissue obtained from the sacrified mice described in Example 13 was homogenized in 0.3 ml of RIPA buffer (Santa Cruz Biotechnology Inc.) containing complete protease inhibitor (Roche). The homogenate was incubated on ice for 30 minutes and centrifuged to pellet cellular debris. Protein concentration in the supernatant was measured using BioRad protein assay reagents (BioRad laboratories). 100µg of protein was denatured at 70°C for 10 minutes and separated on 4-12% Bis-Tris SDS gradient gel (Invitrogen). Proteins were transferred to a 0.45 µm PVDF membrane (Invitrogen) and blocked in washing buffer (50mM Tris PH7.5, 150mM NaCL, 2mM $CaCl_2$ and 0.05% Tween 20) containing 5% non-fat milk for 1 hour at room temperature. The blot was then probed with goat anti-mouse LDLR antibody (R&D system) 1:2000 or anti-ß actin (sigma) 1:2000 for 1 hour at room temperature. The blot was washed briefly and incubated with bovine anti-goat IgG-HRP (Santa Cruz Biotechnology Inc.) 1:2000 or goat anti-mouse IgG-HRP (Upstate) 1:2000. After a 1 hour incubation at room temperature, the blot was washed thoroughly and immunoreactive bands were detected using ECL plus kit (Amersham

biosciences). The Western blot showed an increase in LDLR protein levels in the presence of antibody 31H4, as depicted in FIG. 9.

EXAMPLE 15

Serum cholesterol Lowering Effect of Antibody 31H4 in a 13 Day Study

[0466] In order to assess total serum cholesterol (TC) lowering in wild type (WT) mice via antibody therapy against PCSK9 protein in a 13 day study, the following procedure was performed.

[0467] Male WT mice (C57BL/6 strain, aged 9-10 weeks, 17-27 g) obtained from Jackson Laboratory (Bar Harbor, ME) were fed a normal chow (Harland-Teklad, Diet 2918) through out the duration of the experiment. Mice were administered either anti-PCSK9 antibody 31H4 (2 mg/ml in PBS) or control IgG (2 mg/ml in PBS) at a level of 10 mg/kg through the mouse's tail vein at T=0. Naïve mice were also set aside as naïve control group.

[0468] Dosing groups and time of sacrifice are shown in Table 10. Animals were sacrificed and livers were extracted and prepared as in Example 13.

TABLE 10

Group	Treatment	Time point after dosing	Number	Dose
1	IgG	72 hr	6	10mg/kg
2	31H4	72 hr	6	10mg/kg
3	31H4	72 hr	6	1mg/kg
4	IgG	144 hr	6	10mg/kg
5	31H4	144 hr	6	10mg/kg
6	31H4	144 hr	6	1mg/kg
7	IgG	192 hr	6	10mg/kg
8	31H4	192 hr	6	10mg/kg
9	31H4	192 hr	6	1mg/kg
10	IgG	240 hr	6	10mg/kg
11	31H4	240hr	6	10mg/kg
12	31H4	240hr	6	1mg/kg
13	IgG	312 hr	6	10mg/kg

Group	Treatment	Time point after dosing	Number	Dose
14	31H4	312 hr	6	10mg/kg
15	31H4	312 hr	6	1mg/kg
16	Naive	n/a	6	n/a

[0469] When the 6 day experiment was extended to a 13 day study, the same serum cholesterol lowering effect observed in the 6 day study was also observed in the 13 day study. More specifically, animals dosed at 10 mg/kg demonstrated a 31% decrease in serum cholesterol on day 3, which gradually returned to pre-dosing levels by day 13. FIG. 10A depicts the results of this experiment. FIG. 10C depicts the results of repeating the above procedure with the 10mg/kg dose of 31H4, and with another antibody, 16F12, also at 10mg/kg. Dosing groups and time of sacrifice are shown in Table 11.

TABLE 11

Group	Treatment	Time point after dosing	Number	Dose
1	IgG	24 hr	6	10mg/kg
2	16F12	24 hr	6	10mg/kg
3	31H4	24 hr	6	10mg/kg
4	IgG	72 hr	6	10mg/kg
5	16F12	72 hr	6	10mg/kg
6	31H4	72 hr	6	10mg/kg
7	IgG	144 hr	6	10mg/kg
8	16F12	144 hr	6	10mg/kg
9	31H4	144 hr	6	10mg/kg
10	IgG	192 hr	6	10mg/kg
11	16F12	192 hr	6	10mg/kg
12	31H4	192 hr	6	10mg/kg
13	IgG2	240 hr	6	10mg/kg
14	16F12	240hr	6	10mg/kg

Group	Treatment	Time point after dosing	Number	Dose
15	31H4	240hr	6	10mg/kg
16	IgG2	312 hr	6	10mg/kg
17	16F12	312 hr	6	10mg/kg
18	31H4	312 hr	6	10mg/kg
19	Naive	n/a	6	10mg/kg

[0470] As shown in FIG. 10C both 16F12 and 31H4 resulted in significant and substantial decreases in total serum cholesterol after just a single dose and provided benefits for over a week (10 days or more). The results of the repeated 13 day study were consistent with the results of the first 13 day study, with a decrease in serum cholesterol levels of 26% on day 3 being observed. For FIG. 10A and FIG. 10B, the percentage change is in relation to the control IgG at the same time point (*P<0.01). For FIG. 10C, the percentage change is in relation to the control IgG at the same time point (*P<0.05).

EXAMPLE 16

Effect of Antibody 31H4 on HDL Levels in a 13 Day Study

[0471] The HDL levels for the animals in Example 15 were also examined. HDL levels decreased in the mice. More specifically, animals dosed at 10 mg/kg demonstrated a 33% decrease in HDL levels on day 3, which gradually returned to pre-dosing levels by day 13. FIG. 10B depicts the results of the experiment. There was a decrease in HDL levels of 34% on day 3. FIG. 10B depicts the results of the repeated 13 day experiment.

[0472] As will be appreciated by one of skill in the art, while the antibodies will lower mouse HDL, this is not expected to occur in humans because of the differences in HDL in humans and other organisms (such as mice). Thus, the decrease in mouse HDL is not indicative of a decrease in human HDL.

EXAMPLE 17

Repeated Administration of Antibodies Produce Continued Benefits of Antigen Binding Peptides

[0473] In order to verify that the results obtained in the Examples above can be prolonged for further benefits with additional doses, the Experiments in Examples 15 and 16 were repeated with the dosing schedule depicted in FIG. 11A. The results are displayed in FIG. 11B. As can be seen in the graph in FIG. 11B, while both sets of mice displayed a significant decrease in total serum cholesterol because all of the mice received an initial injection of the 31H4 antigen binding protein, the mice that received additional injections of the 31H4 ABP displayed a continued reduction in total serum cholesterol, while those mice that only received the control injection eventually displayed an increase in their total serum cholesterol. For FIG. 11, the percentage change is in relation to the naïve animals at t=0 hours (*P<0.01, **P<0.001).

[0474] The results from this example demonstrate that, unlike other cholesterol treatment methods, in which repeated applications lead to a reduction in efficacy because of biological adjustments in the subject, the present approach does not seem to suffer from this issue over the time period examined. Moreover, this suggests that the return of total serum cholesterol or HDL cholesterol levels to baseline, observed in the previous exampless is not due to some resistance to the treatment being developed by the subject, but rather the depletion of the antibody availability in the subject.

EXAMPLE 18

Epitope Mapping of Human Anti PCSK9 Antibodies

[0475] This example outlines methods for determining which residues in PCSK9 are involved in forming or part of the epitope for the antigen binding proteins disclosed herein to PCSK9.

[0476] In order to determine the epitopes to which certain of the ABPs of the present invention bind, the epitopes of the ABPs can be mapped using synthetic peptides derived from the specific PCSK9 peptide sequence.

[0477] A SPOTs peptide array (Sigma Genosys) can be used to study the molecular interaction of the human anti-PCSK9 antibodies with their peptide epitope. SPOTs technology is based on the solid-phase synthesis of peptides in a format suitable for the systematic analysis of antibody epitopes. Synthesis of custom arrayed oligopeptides is commerically available from Sigma-Genosys. A peptide array of overlapping oligopeptides derived from the amino-acid sequence of the PCSK9 peptide can be obtained. The array can comprise a series of 12-mer

peptides as spots on a polypropylene membrane sheets. The peptide array can span the entire length of the PCSK9 mature sequence. Each consecutive peptide can be offset by 1 residue from the previous one, yielding a nested, overlapping library of arrayed oligopeptides. The membrane carrying the peptides can be reacted with different anti-PCSK9 antibodies (1 micrograms/ml). The binding of the mAbs to the membrane-bound peptides can be assessed by an enzyme-linked immunosorbent assay using HRP-conjugated secondary antibody followed by enhanced chemiluminescence (ECL).

[0478] In addition, functional epitopes can be mapped by combinatorial alanine scanning. In this process, a combinatorial alanine-scanning strategy can be used to identify amino acids in the PCSK9 protein that are necessary for interaction with anti-PCSK9 ABPs. To accomplish this, a second set of SPOTs arrays can be used for alanine scanning. A panel of variant peptides with alanine substitutions in each of the 12 residues can be scanned as above. This will allow for the epitopes for the ABPs to the human PCSK9 to be mapped and identified.

[0479] In the alternative, given that it is possible that the epitope is conformational, a combination of alanine scanning and/or arginine scanning, antibody FAB/PCSK9 co-crystalization, and limited proteolysis/LC-MS (liquid chromatography mass spec.) can be employed to indentify the epitopes.

Example 19

Uses of PCSK9 Antibodies for the Treatment of Cholesterol Related Disorders

[0480] A human patient exhibiting a Cholesterol Related Disorder (in which a reduction in cholesterol (such as serum cholesterol) can be beneficial) is administered a therapeutically effective amount of PCSK9 antibody, 31H4 (or, for example, 21B12 or 16F12). At periodic times during the treatment, the patient is monitored to determine whether the symptoms of the disorder has subsided. Following treatment, it is found that patients undergoing treatment with the PCSK9 antibody have reduced serum cholesterol levels, in comparison to patients that are not treated.

EXAMPLE 20

Uses of PCSK9 Antibodies for the Treatment of Hypercholesterolemia

[0481] A human patient exhibiting symptoms of hypercholesterolemia is administered a therapeutcially effective amount of PCSK9 antibody, such as 31H4 (or, for example, 21B12 or 16F12). At periodic times during the treatment, the human patient is monitored to determine whether the serum cholesterol level has declined. Following treatment, it is found that the patient receiving the treatment with the PCSK9 antibodies has reduced serum cholesterol levels in comparison to arthritis patients not receiving the treatment.

EXAMPLE 21

Uses of PCSK9 Antibodies for the Prevention of Coronary Heart Disease and/or Recurrent Cardiovascular Events

[0482] A human patient at risk of developing coronary haeart disease is identified. The patient is administered a therapeutically effective amount of PCSK9 antibody, such as 31H4 (or, for example, 21B12 or 16F12), either alone, concurrently or sequentially with a statin, e.g., simvastatin. At periodic times during the treatment, the human patient is monitored to determine whether the patient's total serum cholesterol level changes. Throughout the preventative treatment, it is found that the patient receiving the treatment with the PCSK9 antibodies has reduced serum cholesterol thereby reducing their risk to coronary heart disases or recurrent cardiovascular events in comparison to patients not receiving the treatment.

EXAMPLE 22

Use of PCSK9 Antibodies as a Diagnostic Agent

[0483] An Enzyme-Linked Immunosorbent Assay (ELISA) for the detection of PCSK9 antigen in a sample can used to diagnose patients exhibiting high levels of PCSK9 production. In the assay, wells of a microtiter plate, such as a 96-well microtiter plate or a 384-well microtiter plate, are adsorbed for several hours with a first fully human monoclonal antibody directed against PCSK9. The immobilized antibody serves as a capture antibody for any of the PCSK9 that may be present in a test sample. The wells are rinsed and treated with a blocking agent such as milk protein or albumin to prevent nonspecific adsorption of the analyte.

[0484] Subsequently the wells are treated with a test sample suspected of containing the PCSK9, or with a solution containing a standard amount of the antigen. Such a sample may be, for example, a serum sample from a subject suspected of having levels of circulating antigen considered to be diagnostic of a pathology.

[0485] After rinsing away the test sample or standard, the wells are treated with a second fully human monoclonal PCSK9 antibody that is labeled by conjugation with biotin. A monoclonal or mouse or other species origin can also be used. The labeled PCSK9 antibody serves as a detecting antibody. After rinsing away excess second antibody, the wells are treated with avidin-conjugated horseradish peroxidase (HRP) and a suitable chromogenic substrate. The concentration of the antigen in the test samples is determined by comparison with a standard curve developed from the standard samples.

[0486] This ELISA assay provides a highly specific and very sensitive assay for the detection of the PCSK9 antigen in a test sample.

Determination of PCSK9 Protein Concentration in Subjects

[0487] A sandwich ELISA can quantify PCSK9 levels in human serum. Two fully human monoclonal PCSK9 antibodies from the sandwich ELISA, recognize different epitopes on the PCSK9 molecule. Alternatively, monoclonal antibodies of mouse or other species origin may be used. The ELISA is performed as follows: 50 µL of capture PCSK9 antibody in coating buffer (0.1 M $NaHCO_3$, pH 9.6) at a concentration of 2 µg/mL is coated on ELISA plates (Fisher). After incubation at 4° C. overnight, the plates are treated with 200 µL of blocking buffer (0.5% BSA, 0.1% Tween 20, 0.01% Thimerosal in PBS) for 1 hour at 25° C. The plates are washed (3x) using 0.05% Tween 20 in PBS (washing buffer, WB). Normal or patient sera (Clinomics, Bioreclaimation) are diluted in blocking buffer containing 50% human serum. The plates are incubated with serum samples overnight at 4° C, washed with WB, and then incubated with 100 µL/well of biotinylated detection PCSK9 antibody for 1 hour at 25° C After washing, the plates are incubated with HRP-Streptavidin for 15 minutes, washed as before, and then treated with 100 µL/well of o-phenylenediamine in H_2O_2 (Sigma developing solution) for color generation. The reaction is stopped with 50 µL/well of H_2SO_4 (2M) and analyzed using an ELISA plate reader at 492 nm. Concentration of PCSK9 antigen in serum samples is calculated by comparison to dilutions of purified PCSK9 antigen using a four parameter curve fitting program.

Determination of PCSK9 Variant Protein Concentration in Subjects

[0488] The steps outlined above can be performed using antibodies noted herein that bind to both the wild type PCSK9 and the variant PCSK9 (D374Y). Next, antibodies that bind to

the wild type but not the mutant can be used (again using a similar protocol as outlined above) to determine if the PCSK9 present in the subject is wild type or the D374Y variant. As will be appreciatedy by one of skill in the art, results that are positive for both rounds will be wild-type, while those that are positive for the first round, but not the second round of antibodies, will include the D374Y mutation. There are high frequency mutations in the population that are known and the could benefit particularly from an agent such as the ABPs disclosed herein.

EXAMPLE 23

Use of PCSK9 Antigen Binding Protein for the Prevention of Hypercholesterolemia

[0489] A human patient exhibiting a risk of developing hypercholesterolemia is identified via family history analysis and/or lifestyle, and/or current cholesterol levels. The subject is regularly administered (e.g., one time weekly) a therapeutically effective amount of PCSK9 antibody, 31H4 (or, for example, 21B12 or 16F12). At periodic times during the treatment, the patient is monitored to determine whether serum cholesterol levels have decreased. Following treatment, it is found that subjects undergoing preventative treatment with the PCSK9 antibody have lowered serum cholesterol levels, in comparison to subjects that are not treated.

EXAMPLE 24

PCSK9 ABPs Further Upregulated LDLR in the Presence of Statins

[0490] This example demonstrates that ABPs to PCSK9 produced further increases in LDLR availability when used in the presence of statins, demonstrating that further benefits can be achieved by the combined use of the two.

[0491] HepG2 cells were seeded in DMEM with 10% fetal bovine serum (FBS) and grown to ~90% confluence. The cells were treated with indicated amounts of mevinolin (a statin, Sigma) and PCSK9 ABPs (FIGs. 12A-12C) in DMEM with 3% FBS for 48 hours. Total cell lysates were prepared. 50 mg of total proteins were separated by gel electrophoresis and transferred to PVDF membrane. Immunoblots were performed using rabbit anti-human LDL receptor antibody (Fitzgerald) or rabbit anti-human b-actin antibody. The enhanced chemiluminescent results are shown in the top panels of FIGs. 12A-12C. The intensity of the bands were quantified by ImageJ software and normalized by b-actin. The relative levels of

LDLR are shown in the lower panels of FIGs. 12A-12C. ABPs 21B12 and 31H4 are PCSK9 neutralizing antibodies, while 25A7.1 is a non-neutralizing antibody.

[0492] HepG2-PCSK9 cells were also created. These were stable HepG2 cell line transfected with human PCSK9. The cells were seeded in DMEM with 10% fetal bovine serum (FBS) and grew to ~90% confluence. The cells were treated with indicated amounts of mevinolin (Sigma) and PCSK9 ABPs (FIGs. 12D-12F) in DMEM with 3% FBS for 48 hours. Total cell lysates were prepared. 50 mg of total proteins were separated by gel electrophoresis and transferred to PVDF membrane. Immunoblots were performed using rabbit anti-human LDL receptor antibody (Fitzgerald) or rabbit anti-human b-actin antibody. The enhanced chemiluminescent results are shown in the top panels. The intensity of the bands were quantified by ImageJ software and normalized by b-actin.

[0493] As can be seen in the results depicted in FIGs. 12A-12F, increasing amounts of the neutralizing antibody and increasing amounts of the statin generally resulted in increases in the level of LDLR. This increase in effectiveness for increasing levels of the ABP is especially evident in FIGs. 12D-12F, in which the cells were also transfected with PCSK9, allowing the ABPs to demonstrate their effectiveness to a greater extent.

[0494] Interestingly, as demonstrated by the results in the comparison of FIGs. 12D-12F to 12A-12C, the influence of the ABP concentrations on LDLR levels increased dramatically when PCSK9 was being produced by the cells. In addition, it is clear that the neutralizing ABPs (21B12 and 31H4) resulted in a greater increase in LDLR levels, even in the presence of statins, than the 25A7.1 ABP (a non-neutralizer), demonstrating that additional benefits can be achieved by the use of both statins and ABPs to PCSK9.

EXAMPLE 25

Consensus Sequences

[0495] Consensus sequences were determined using standard phylogenic analyses of the CDRs corresponding to the V_H and V_L of anti-PCSK9 ABPs. The consensus sequences were determined by keeping the CDRs contiguous within the same sequence corresponding to a V_H or V_L. Briefly, amino acid sequences corresponding to the entire variable domains of either V_H or V_L were converted to FASTA formatting for ease in processing comparative alignments and inferring phylogenies. Next, framework regions of these sequences were replaced with an

artificial linker sequence ("bbbbbbbbbb" placeholders, non-specific nucleic acid construct) so that examination of the CDRs alone could be performed without introducing any amino acid position weighting bias due to coincident events (*e.g.*, such as unrelated antibodies that serendipitously share a common germline framework heritage) while still keeping CDRs contiguous within the same sequence corresponding to a V_H or V_L. V_H or V_L sequences of this format were then subjected to sequence similarity alignment interrogation using a program that employs a standard ClutalW-like algorithm (*see*, Thompson *et al.*, 1994, *Nucleic Acids Res.* 22:4673-4680). A gap creation penalty of 8.0 was employed along with a gap extension penalty of 2.0. This program likewise generated phylograms (phylogenic tree illustrations) based on sequence similarity alignments using either UPGMA (unweighted pair group method using arithmetic averages) or Neighbor-Joining methods (*see*, Saitou and Nei, 1987, *Molecular Biology and Evolution* 4:406-425) to construct and illustrate similarity and distinction of sequence groups *via* branch length comparison and grouping. Both methods produced similar results but UPGMA-derived trees were ultimately used as the method employs a simpler and more conservative set of assumptions. UPGMA-derived trees were generated where similar groups of sequences were defined as having fewer than 15 substitutions per 100 residues (*see*, legend in tree illustrations for scale) amongst individual sequences within the group and were used to define consensus sequence collections. The results of the comparisons are depicted in FIGs. 13A-13J. In FIG. 13E, the groups were chosen so that sequences in the light chain that clade are also a clade in the heavy chain and have fewer than 15 substitutions.

[0496] As will be appreciated by one of skill in the art, the results presented in FIGs. 13A-13J present a large amount of guidance as to the importance of particular amino acids (for example, those amino acids that are conserved) and which amino acid positions can likely be altered (for example, those positions that have different amino acids for different ABPs).

EXAMPLE 26

Mouse Model for PCSK9 and ABP ability to lower LDL *in vivo*

[0497] To generate mice which over-expressed human PCSK9, three week old WT C57Bl/6 mice were injected via tail vein administration with various concentrations of adenoassociated virus (AAV), recombinantly modified to express human PCSK9, to determine the correct titer which would provide a measurable increase of LDL-cholesterol in the

mice. Using this particular virus that expressed human PCSK9, it was determined that 4.5 x 10E12 pfu of virus would result in an LDL-cholesterol level of approximately 40mg/dL in circulating blood (normal levels of LDL in a WT mice are approximately 10mg/dL). The human PCSK9 levels in these animals was found to be approximately 13ug/mL. A colony of mice were generated using this injection criteria.

[0498] One week after injection, mice were assessed for LDL-cholesterol levels, and randomized into different treatment groups. Animals were then administered, via tail vein injection, a single bolus injection of either 10mg/kg or 30mg/kg of 16F12, 21B12, or 31H4 antigen binding proteins. IgG2 ABP was administered in a separate group of animals as a dosing control. Subgroups of animals (n=6-7) were then euthanized at 24 and 48 hours after ABP administration. There were no effects on LDL-cholesterol levels following IgG2 administration at either dose. Both 31H4 and 21B12 demonstrated significant LDL-cholesterol lowering up to and including 48 hours post-administration, as compared to IgG2 control (shown in FIG. 14A and 14B at two different doses). 16F12 shows an intermediary LDL-cholesterol lowering response, with levels returning to baseline of approximately 40mg/dL by the 48 hour time point. This data is consistent with *in vitro* binding data (Biacore and Kinexa), which shows near equivalent binding affinity between 31H4 and 21B12, and a lesser affinity of 16F12 to human PCSK9.

[0499] As can be seen in the results, total cholesterol and HDL-cholesterol were reduced by the PCSK9 ABPs in the model (both total and HDL-C are elevated above WT mice due to the overexpression of PCSK9). While cholesterol lowering in this model appears to occur over a relatively short period of time, this is believed to be due to the levels of human PCSK9 that are present, which are supraphysiologically high in this model. In addition, given that the expression is governed by AAV, there is no regulation of PCSK9 expression. In these figures, (*) denotes a $P<0.05$, and (**) denotes a $P<0.005$ as compared to LDL-cholesterol levels observed in IgG2 control injected animals at the same time point. The 13 microgram/ml level of serum human PCSK9 in the mice corresponds to an approximately 520-fold increase above the endogenous mouse PCSK9 levels (~ 25 ng/ml), and an approximately 75-fold increase above average human serum levels (~ 175 ng/ml). Thus, the antigen binding proteins should be even more effective in humans.

[0500] As will be appreciated by one of skill in the art, the above results demonstrate that appropriateness of the mouse model for testing the antigen binding protein's ability to alter serum cholesterol in a subject. One of skill in the art will also recognize that the use of mouse HDL to monitor serum cholesterol levels in a mouse, while useful for monitoring mouse serum cholesterol levels, is not indicative of the ABPs impact on human HDL in humans. For example, Cohen et al. ("Sequence variations in PCSK9, low LDL, and protection against coronary heart disease", N Engl J Med, 354:1264-1272, 2006) demonstrated the lack of any effect of the PCSK9 loss-of-function mutations on human HDL levels (the entirety of which is incorporated by reference). Thus, one of skill in the art will appreciate that the ability of the ABP to lower mouse HDL (which lack LDL) is not indicative of the ABP's ability to lower human HDL. Indeed, as shown by Cohen, this is unlikely to occur for neutralizing antibodies in humans.

EXAMPLE 27

31H4 and 21B12 bind to the ProCat region of PCSK9

[0501] The present example describes one method for determining where various antibodies bind to PCSK9.

[0502] The ProCat (31-449 of SEQ ID NO: 3) or V domain (450-692 of SEQ ID NO: 3) of the PCSK9 protein was combined with either antibody 31H4 or 21B12. The samples were analyzed by Native PAGE for complex formation. As can be seen in FIG. 16A and FIG. 16B, gel shifts were present for the ProCat / 31H4 and ProCat / 21B12 samples, demonstrating that the antibodies bound to the ProCat domain.

EXAMPLE 28

The LDLR EGFa domain binds to the catalytic domain of PCSK9

[0503] The present example presents the solved crystal structure of PCSK9 ProCat (31-454 of SEQ ID NO: 3) bound to the LDLR EGFa domain (293-334) at 2.9 Å resolution (the conditions for which are described in the below Examples).

[0504] A representation of the structure of PCSK9 bound to EGFa is shown in FIG. 17. The crystal structure (and its depiction in FIG. 17) reveals that the EGFa domain of LDLR binds to the catalytic domain of PCSK9. In addition, the interaction of PCSK9 and EGFa

appears to occur across a surface of PCSK9 that is between residues D374 and S153 in the structure depicted in FIG. 17.

[0505] Specific core PCSK9 amino acid residues of the interaction interface with the LDLR EGFa domain were defined as PCSK9 residues that are within 5 Å of the EGFa domain. The core residues are as follows: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, and S381.

[0506] Boundary PCSK9 amino acid residues of the interaction interface with the LDLR EGFa domain were defined as PCSK9 residues that are 5-8 Å from the EGFa domain. The boundary residues are as follows: W156, N157, L158, E159, H193, E195, H229, R237, G240, K243, D367, I368, G370, A371, S373, S376, and Q382. Residues that are underlined are nearly or completely buried within PCSK9.

[0507] As will be appreciated by one of skill in the art, the results from this example demonstrate where PCSK9 and EGFa interact. Thus, antibodies that interact with or block any of these residues can be useful as antibodies that inhibit the interaction between PCSK9 and the EGFa domain of LDLR (and/or LDLR generally). In some embodiments, antibodies that, when bound to PCSK9, interact with or block any of the above residues or are within 15-8, 8, 8-5, or 5 angstroms of the above residues are contemplated to provide useful inhibition of PCSK9 binding to LDLR.

EXAMPLE 29

31H4 interacts with amino acid residues from both the pro- and catalytic domains of PCSK9

[0508] The present example presents the crystal structure of full length PCSK9 (N533A mutant of SEQ ID NO: 3) bound to the Fab fragment of 31H4, determined to 2.3 Å resolution (the conditions for which are described in the below Examples). This structure, depicted in FIG. 18A and 18B, shows that 31H4 binds to PCSK9 in the region of the catalytic site and makes contacts with amino acid residues from both the prodomain and catalytic domain.

[0509] The depicted structure also allows one to identify specific core PCSK9 amino acid residues for the interaction interface of 31H4 with PCSK9. This was defined as residues that are within 5 Å of the 31H4 protein. The core residues are as follows: W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258,

N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, and G384.

[0510] The structures were also used to identify boundary PCSK9 amino acid residues for the interaction interface with 31H4. These residues were PCSK9 residues that were 5-8 Å from the 31H4 protein. The boundary residues are as follows: K69, D70, P71, S148, V149, D186, T187, E211, D212, G213, R218, Q219, C223, D224, G227, H229, L253, N254, G259, P288, A290, G291, G316, R319, Y325, V346, G352, T353, G365, I368, I369, S372, S373, C378, F379, T385, S386, and Q387. Amino acid residues completely buried within the PCSK9 protein are underlined.

[0511] As will be appreciated by one of skill in the art, FIG. 18B depicts the interaction between the CDRs on the antigen binding protein and PCSK9. As such, the model allows one of skill in the art to identify the residues and/or CDRs that are especially important in the paratope, and which residues are less critical to the paratope. As can be seen in FIG. 18B, the heavy chain CDR1, CDR2, and CDR3 are most directly involved in the antigen binding protein's binding to the epitope, with the CDRs from the light chain being relatively far away from the epitope. As such, it is probable that larger variations in the light chain CDRs are possible, without unduly interfering with the binding of the antigen binding protein to PCSK9. In some embodiments, residues in the structures that directly interact are conserved (or alternatively conservatively replaced) while residues that are not directly interacting with one another can be altered to a greater extent. As such, one of skill in the art, given the present teachings, can predict which residues and areas of the antigen binding proteins can be varied without unduly interfering with the antigen binding protein's ability to bind to PCSK9. For example, those residues that are located closest to PCSK9 when the antigen binding protein is bound to PCSK9 are those that likely play a more important role in the binding of the antigen binding protein to PCSK9. As above, these residues can be divided into those that are within 5 angstroms of PCSK9 and those that are between 5 and 8 angstroms. Specific core 31H4 amino acid residues of the interaction interface with PCSK9 were defined as 31H4 residues that are within 5 Å of the PCSK9 protein. For the heavy chain, the residues that are within 5 angstroms include the following: T28, S30, S31, Y32, S54, S55, S56, Y57, I58, S59, Y60, N74, A75, R98, Y100, F102, W103, S104, A105, Y106, Y107, D108, A109, and D111. For the light chain,

those residues that are within 5 angstroms include the following: L48, S51, Y93, and S98. For the heavy chain, those residues that are 5-8 Å from the PCSK9 protein include the following: G26, F27, F29, W47, S50, I51, S52, S53, K65, F68, T69, I70, S71, R72, D73, K76, N77, D99, D101, F110, and V112. For the light chain, those residues that are within 5-8 angstroms of PCSK9 include A31, G32, Y33, D34, H36, Y38, I50, G52, N55, R56, P57, S58, D94, S95, S96, L97, G99, and S100.

[0512] As will be appreciated by one of skill in the art, the results from Example 29 demonstrate where antibodies to PCSK9 can interact on PCSK9 and still block PCSK9 from interacting with EGFa (and thus LDLR). Thus, antigen binding proteins that interact with any of these PCSK9 residues, or that block any of these residues (*e.g.*, from other antigen binding proteins that bind to these residues), can be useful as antibodies that inhibit the interaction of PCSK9 and EGFa (and LDLR accordingly). Thus, in some embodiments, antigen binding proteins that interact with any of the above residues or interact with residues that are within 5 Å of the above residues are contemplated to provide useful inhibition PCSK9 binding to LDLR. Similarly, antigen binding proteins that block any of the above residues (which can be determined, for example, via a competition assay) can also be useful for inhibition of the PCSK9/LDLR interaction.

EXAMPLE 30

21B12 binds to the catalytic domain of PCSK9, has a distinct binding site from 31H4 and can bind to PCSK9 simultaneously with 31H4

[0513] The present example presents the crystal structure of PCSK9 ProCat (31-449 of SEQ ID NO: 3) bound to the Fab fragments of 31H4 and 21B12, determined at 2.8 Å resolution (the conditions for which are described in the below Examples). This crystal structure, depicted in FIG. 19A and FIG. 19B, shows that 31H4 and 21B12 have distinct binding sites on PCSK9 and that both antigen binding proteins can bind to PCSK9 simultaneously. The structure shows that 21B12 interacts with amino acid residues from PCSK9's catalytic domain. In this structure, the interaction between PCSK9 and 31H4 is similar to what was observed above.

[0514] Specific core PCSK9 amino acid residues of the interaction interface with 21B12 were defined as PCSK9 residues that are within 5 Å of the 21B12 protein. The core

residues are as follows: S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, and F379.

[0515] Boundary PCSK9 amino acid residues of the interaction interface with 21B12 were defined as PCSK9 residues that were 5-8 Å from the 21B12 protein. The boundary residues are as follows: I154, T187, H193, E195, I196, M201, V202, C223, T228, S235, G236, A239, G244, M247, I369, S372, C375, and C378. Amino acid residues nearly or completely buried within the PCSK9 protein are underlined.

[0516] As will be appreciated by one of skill in the art, FIG. 19B depicts the interaction between the CDRs on the antigen binding protein and PCSK9. As such, the model allows one of skill in the art to identify the residues and/or CDRs which are especially important for the paratope and which residues are less critical to the paratope. As can be seen in the structure, heavy chain CDR2 and light chain CDR1 appear to closely interact with the epitope. Next, heavy chain CDR1, heavy chain CDR3 and light chain CDR3, appear to be close to the epitope, but not as close as the first set of CDRs. Finally, light chain CDR2 appears to be some distance from the epitope. As such, it is probable that larger variations in the more distant CDRs are possible without unduly interfering with the binding of the antigen binding protein to PCSK9. In some embodiments, residues in the structures that directly interact are conserved (or alternatively conservatively replaced) while residues that are not directly interacting with one another can be altered to a greater extent. As such, one of skill in the art, given the present teachings, can predict which residues and areas of the antigen binding proteins can be varied without unduly interfering with the antigen binding protein's ability to bind to PCSK9. For example, those residues that are located closest to PCSK9 when the antigen binding protein is bound to PCSK9 are those that likely play a more important role in the binding of the antigen binding protein to PCSK9. As above, these residues can be divided into those that are within 5 angstroms of PCSK9 and those that are between 5 and 8 angstroms. Specific core 21B12 amino acid residues of the interaction interface with PCSK9 were defined as 21B12 residues that are within 5 Å of the PCSK9 protein. For the heavy chain, the residues that are within 5 angstroms include the following: T30, S31, Y32, G33, W50, S52, F53, Y54, N55, N57, N59, R98, G99, Y100, and G101. For the light chain, those residues that are within 5 angstroms include the following: G30, G31, Y32, N33, S34, E52, Y93, T94, S95, T96, and S97. For the heavy chain,

those residues that are 5-8 Å from the PCSK9 protein include the following: T28, L29, I34, S35, W47, V51, G56, T58, Y60, T72, M102, and D103. For the light chain, those residues that are within 5-8 angstroms of PCSK9 include the following: S26, V29, V35, Y51, N55, S92, M98, and V99.

[0517] As will be appreciated by one of skill in the art, the results from Example 30 demonstrate where antigen binding proteins to PCSK9 can interact on PCSK9 and still block PCSK9 from interacting with EGFa (and thus LDLR). Thus, antigen binding proteins that interact with any of these PCSK9 residues or that block any of these residues can be useful as antibodies that inhibit the interaction of PCSK9 and EGFa (and LDLR accordingly). Thus, in some embodiments, antibodies that interact with any of the above residues or interact with residues that are within 5 Å of the above residues are contemplated to provide useful inhibition PCSK9 binding to LDLR. Similarly, antigen binding proteins that block any of the above residues (which can be determined, for example, via a competition assay) can also be useful for inhibition of PCSK9/LDLR interaction.

EXAMPLE 31

Interaction between EGFa, PCSK9, and the Antibodies

[0518] The structure of the ternary complex (PCSK9 / 31H4 / 21B12) from the above example was overlaid on the PCSK9 / EGFa structure (determined as described in Example 28) and the result of this combination is depicted in FIG. 20A. This figure demonstrates areas on PCSK9 which can be usefully targeted to inhibit PCSK9 interaction with EGFa. The figure shows that both 31H4 and 21B12 partially overlap with the position of the EGFa domain of LDLR and sterically interfere with its binding to PCSK9. In addition, as can be seen in the structures, 21B12 directly interacts with a subset of amino acid residues that are specifically involved in binding to the LDLR EGFa domain.

[0519] As noted above, analysis of the crystal structures identified specific amino acids involved in the interaction between PCSK9 and the partner proteins (the core and boundary regions of the interface on the PCSK9 surface) and the spatial requirements of these partner proteins to interact with PCSK9. The structures suggest ways to inhibit the interaction between PCSK9 and the LDLR. First, as noted above, binding an agent to PCSK9 where it shares residues in common with the binding site of the EGFa domain of the LDLR would inhibit the

interaction between PCSK9 and the LDLR. Second, an agent that binds outside of the residues in common can sterically interfere with the EGFa domain or regions of the LDLR that are either N- or C-terminal to the EGFa domain to prevent the interaction between PCSK9 and the LDLR.

[0520] In some embodiments, the residues that are involved in both EGFa binding and are close to the areas where the above noted antigen binding proteins bind are especially useful for manipulating PCSK9 binding to LDLR. For example, amino acid residues from interfaces in common in both the core region and boundary region for the different binding partners are listed in Table 12 below. Amino acid residues completely buried within the PCSK9 protein are underlined.

Table 12

Parameters	Amino acid position(s)
31H4 / EGFa both under 5 Å	D374, V380, S381
31H4 under 5 Å / EGFa 5-8 Å	D367, Q382
31H4 at 5-8 Å / EGFa under 5 Å	I369, S372, C378, F379
31H4 / EGFa both at 5-8 Å	H229, S373
21B12 / EGFa both under 5 Å	S153, R194, D238, D374, T377, F379
21B12 under 5 Å / EGFa 5-8 Å	R237, K243, S373, S376
21B12 at 5-8 Å / EGFa under 5 Å	I154, A239, I369, S372, C375, C378
21B12 / EGFa both at 5-8 Å	H193, E195

[0521] As will be appreciated by one of skill in the art, in some embodiments, the antigen binding proteins bind to and/or block at least one of the above noted residues.

EXAMPLE 32

Structural interaction of LDLR and PCSK9

[0522] A model of full length PCSK9 bound to a full length representation of the LDLR was made using the PCSK9 ProCat (31-454 of SEQ ID NO: 3)/EGFa complex structure.

The structure of full length PCSK9[1] (Piper, D.E. et al. The crystal structure of PCSK9: a regulator of plasma LDL-cholesterol. *Structure* 15, 545-52 (2007)) was overlaid onto the PCSK9 ProCat 31-454 from the complex and the structure of the LDLR in its low pH conformation (Rudenko, G. et al. Structure of the LDL receptor extracellular domain at endosomal pH. *Science* 298, 2353-8 (2002)) was overlaid onto the EGFa domain from the complex. Depictions of the model are shown in FIGs. 20B and 20C. The EGFa domain is indicated by the box in the figure. The figures show regions of the LDLR outside of the immediate EGFa binding domain that lie in close proximity to PCSK9. FIGs. 20D-20F show the above interaction, along with mesh surface representations of antibody 31H4 and 21B12 from three different angles. As is clear from the depictions, not only can the antibody interact and/or interfere with LDLR's interaction with PCSK9 at the actual binding site, but other steric interactions appear to occur as well.

[0523] In light of the above results, it is clear that antigen binding proteins that bind to PCSK9 can also inhibit the interaction between PCSK9 and the LDLR by clashing with various regions of the LDLR (not just the site at which LDLR and PCSK9 interact). For example, it can clash with repeat 7 (R7), the EGFb domain, and/or the β-propeller domain.

Embodiments of Antigen Binding Molecules that bind to or block EGFa interaction with PCSK9

[0524] As will be appreciated by one of skill in the art, Examples 28-32, and their accompanying figures, provide a detailed description of how and where EGFa interacts with PCSK9 and how two representative neutralizing antigen binding proteins, 21B12 and 31H4 interact with PCSK9 and produce their neutralizing effect. As such, one of skill in the art will readily be able to identify antigen binding molecules that can similarly reduce the binding between EGFa (including LDLR) and PCSK9 by identifying other antigen binding molecules that bind at or near at least one of the same locations on PCSK9. While the relevant locations (or epitopes) on PCSK9 are identified in the figures and the present description, it can also be advantageous to describe these sites as being within a set distance from residues that have been identified as close to the EGFa binding site. In some embodiments, an antigen binding molecule will bind to or within 30 angstroms of one or more of the following residues (numbering in reference to SEQ ID NO: 3): S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, S381, W156, N157, L158, E159, H193, E195, H229, R237, G240,

K243, D367, I368, G370, A371, S373, S376, Q382, W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, G384, K69, D70, P71, S148, V149, D186, T187, E211, D212, G213, R218, Q219, C223, D224, G227, H229, L253, N254, G259, P288, A290, G291, G316, R319, Y325, V346, G352, T353, G365, I368, I369, S372, S373, C378, F379, T385, S386, Q387, S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, F379, I154, T187, H193, E195, I196, M201, V202, C223, T228, S235, G236, A239, G244, M247, I369, S372, C375, or C378. In some embodiments, the antigen binding molecule binds within 30 angstroms of one or more of the following residues (numbering in reference to SEQ ID NO: 3): S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, S381, W156, N157, L158, E159, H193, E195, H229, R237, G240, K243, D367, I368, G370, A371, S373, S376, or Q382. In some embodiments, the antigen binding molecule binds within 30 angstroms of one or more of the following residues (numbering in reference to SEQ ID NO: 3): W72, F150, A151, Q152, T214, R215, F216, H217, A220, S221, K222, S225, H226, C255, Q256, G257, K258, N317, F318, T347, L348, G349, T350, L351, E366, D367, D374, V380, S381, Q382, S383, G384, K69, D70, P71, S148, V149, D186, T187, E211, D212, G213, R218, Q219, C223, D224, G227, H229, L253, N254, G259, P288, A290, G291, G316, R319, Y325, V346, G352, T353, G365, I368, I369, S372, S373, C378, F379, T385, S386, or Q387. In some embodiments, the antigen binding molecule binds within 30 angstroms of one or more of the following residues (numbering in reference to SEQ ID NO: 3): S153, S188, I189, Q190, S191, D192, R194, E197, G198, R199, V200, D224, R237, D238, K243, S373, D374, S376, T377, F379, I154, T187, H193, E195, I196, M201, V202, C223, T228, S235, G236, A239, G244, M247, I369, S372, C375, or C378.

[0525] In some embodiments, the antigen binding molecule binds within 30, 30-25, 25-20, 20-15, 15-8, 8, 8-5, 5, 5-4, 4 or less angstroms from one or more of the above residues. In some embodiments, the antigen binding molecule, when bound to PCSK9, is within at least one of the above distances, for more than one of the above noted residues. For example, in some embodiments, the antigen binding molecule is within one of the recited distances (e.g., 30, 30-25, 25-20, 20-15, 15-8, 8, 8-5, 5, 5-4, 4 or less) for at least 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 20-25, 25-30, 30-35, 35-40, 40-45, 45-50, 50-55, 55-60, 60-65, 65-70, 70-75

or more of the above residues. In some embodiments, the antigen binding molecule is within one of the recited distances for at least 1-10, 10-20, 20-30, 30-40, 40-50, 50-60, 60-70, 70-80, 80-90, 90-95, 95-99, 99-100% of the residues identified in each group of subgroup thereof (such as only those surface residues in the group). Unless specifically stated otherwise, the distance between the antigen binding molecule and PCSK9 is the shortest distance between the covalently bonded atom on PCSK9 and the covalently bonded atom of the antigen binding molecule that are the closest atoms of PCSK9 and the antigen binding molecule. Similarly, unless specifically stated otherwise, the distance between a residue (on the antigen binding molecule or PCSK9) and another protein (either PCSK9 or the antigen binding molecule respectively), is the distance from the closest point on the identified residue to the closest covalently bonded part of the other protein. In some embodiments, the distance can be measured from the backbone of the amino acid chains. In some embodiments, the distance can be measured between an edge of the paratope and an edge (closest to one another) of the epitope. In some embodiments, the distance can be measured between the center of the surface of the paratope and the center of the surface of the epitope. As will be appreciated by one of skill in the art, the present description is applicable for each of the individual sets of residues listed herein. For example, the above ranges are contemplated generally and specifically for the 8 angstrom residues listed in Examples 28-32 and the 5 angstrom residues listed in Examples 28-32.

[0526] In some embodiments, the antigen binding molecule binds to a surface on PCSK9 that is bound by at least one of EGFa, 21B12, or 31H4. In some embodiments, the antigen binding molecule binds to PCSK9 at a location that overlaps with the interaction locations between PCSK9 and EFGa, Ab 31H4, and/or Ab 21B12 (as described in the above examples and figures). In some embodiments, the antigen binding molecule binds to PCSK9 at a position that is further away from one of the above recited residues. In some embodiments, such an antigen binding molecule can still be an effective neutralizing antigen binding molecule.

[0527] In some embodiments, the structure of the catalytic domain of PCSK9 can be described as generally being triangular (as shown in FIG. 19A). The first side of the triangle is shown as being bound by 31H4. The second side of the triangle is shown as being bound by 21B12, and the third side of the triangle is positioned toward the bottom of the page, immediately above the "FIG. 19A" label. In some embodiments, antigen binding molecules that bind to the first and/or second sides of the catalytic domain of PCSK9 can be useful as

neutralizing antibodies as they can either directly or sterically interfere with EGFa's binding to PCSK9. As will be appreciated by one of skill in the art, when the antigen binding molecules are large enough, such as a full antibody, the antigen binding molecule need not directly bind to the EGFa binding site in order to interfere with the binding of EGFa to PCSK9.

[0528] As will be appreciated by one of skill in the art, while the EGFa domain of the LDLR has been used in many of the examples, the models and structures are still applicable to how the full length LDLR protein will interact with PCSK9. Indeed, the additional structure present on the full length LDLR protein presents additional protein space that can further be blocked by one of the antigen binding molecules. As such, if the antigen binding molecule blocks or inhibits binding of EGFa to PCSK9, it will likely be at least as, if not more, effective with the full length LDLR protein. Similarly, antigen binding molecules that are within a set distance or block various residues that are relevant for inhibiting EGFa binding, will likely be as effective, if not more effective, for the full length LDLR.

[0529] As will be appreciated by one of skill in the art, any molecule that blocks or binds to the above noted PCSK9 residues (or within the recited distances), or that inhibits one or more of the interactions noted in the above examples and figures, can be used to inhibit the interaction of EGFa (or LDLR generally) and PCSK9. As such, the molecule need not be limited to an antigen binding "protein," as any antigen binding molecule can also serve the required purpose. Examples of antigen binding molecules include aptamers, which can be either oligonucleic acid or peptide molecules. Other examples of antigen binding molecules include avimers, peptibodies, small molecules and polymers, and modified versions of EGFa that can increase its affinity to PCSK9 and/or half-life, such as mutation of amino acids, glycosylation, pegylation, Fc fusions, and avimer fusions. As will be appreciated by one of skill in the art, in some embodiments LDLR is not an antigen binding molecule. In some embodiments, binding subsections of LDLR are not antigen binding molecules, *e.g.*, EGFa. In some embodiments, other molecules through which PCSK9 signals *in vivo* are not antigen binding molecules. Such embodiments will be explicitly identified as such.

EXAMPLE 33

Expression and purification of protein samples

[0530] The present example describes some embodiments for how the various embodiments of the PCSK9 proteins/variants were made and purified (including the LDLR EGFa domain). PCSK9 proteins/variants (*e.g.*, PSCK9 31-692 N533A, PCSK9 449TEV and PCSK9 ProCat 31-454) were expressed in baculovirus infected Hi-5 insect cells with an N-terminal honeybee melittin signal peptide followed by a His_6 tag. The PCSK9 proteins were purified by nickel affinity chromatography, ion exchange chromatography and size exclusion chromatography. The melittin-His_6 tag was removed during purification by cleavage with TEV protease. The construct PCSK9 449TEV was used to generate PCSK9 ProCat (31-449) and V domain (450-692) samples. This construct had a TEV protease cleavage site inserted between PCSK9 residues 449 and 450. For the full length N555A variant for crystallography, the PCSK9 31-454 fragment, and the PCSK9 449TEV variant for crystalography, the post rTEV protein product also included an initial GAMG sequence. Thus, post rTEV cleavage, these proteins were GAMG-PCSK9. Furthermore, the PCSK9 449TEV protein included the sequence "ENLYFQ" (SEQ ID NO: 403) inserted between positions H449 and G450 of SEQ ID NO: 3. After cleavage with rTEV, the PCSK9 ProCat protein generated from this construct was GAMG-PCSK9 (31-449)-ENLYFQ and the V domain generated from this construct was PCSK9 (450-692) of SEQ ID NO: 3.

[0531] The 21B12 and 31H4 Fab fragments were expressed in *E. coli*. These proteins were purified by nickel affinity chromatography, size exclusion chromatography and ion exchange chromatography.

[0532] The LDLR EGFa domain (293-334) was expressed as a GST fusion protein in *E. coli*. The EGFa domain was purified by ion exchange chromatography, glutathione sepharose affinity chromatography and size exclusion chromatography. The GST protein was removed during the purification by cleavage with PreScission protease.

EXAMPLE 34

Complex formation and crystallization

[0533] The present example describes how complexes and crystals used in the above structure examination Examples were made.

[0534] The PCSK9 31-692 N533A / 31H4 complex was made by mixing a 1.5 molar excess of the 31H4 Fab with PCSK9. The complex was purified by size exclusion

chromatography to remove excess 31H4 Fab. The PCSK9 31-692 N533A / 31H4 complex crystallizes in 0.1 M Tris pH 8.3, 0.2 M sodium acetate, 15% PEG 4000, 6% dextran sulfate sodium salt (Mr 5000).

[0535] The PCSK9 ProCat 31-449 / 31H4 / 21B12 complex was made by first mixing a 1.5 molar excess of 31H4 Fab with PCSK9 31-449. The complex was separated from excess 31H4 by purification on a size exclusion chromatography column. A 1.5 molar excess of 21B12 Fab was then added to the PCSK9 31-449 / 31H4 complex. The ternary complex was separated from excess 21B12 by purification on a size exclusion chromatography column. The PCSK9 ProCat 31-449 / 31H4 / 21B12 complex crystallizes in 0.1 M Tris pH 8.5, 0.2 M ammonium phosphate monobasic, 50% MPD.

[0536] The PCSK9 ProCat 31-454 / EGFa complex was made by mixing a 1.2 molar excess of EGFa domain with PCSK9 31-454. The PCSK9 ProCat 31-454 / EGFa domain complex crystallizes in 0.2 M potassium formate, 20% PEG 3350.

EXAMPLE 35

Data collection and structure determination

[0537] The present example describes how the datasets were collected and the structures determined for the above structure examination Examples.

[0538] Initial datasets for the PCSK9 31-692 N533A / 31H4 and PCSK9 ProCat 31-449 / 31H4 / 21B12 crystals were collected on a Rigaku FR-E X-ray source. The PCSK9 ProCat 31-454 / EGFa dataset and higher resolution datasets for the PCSK9 31-692 N533A / 31H4 and PCSK9 ProCat 31-449 / 31H4 / 21B12 crystals were collected at the Berkeley Advanced Light Source beamline 5.0.2. All datasets were processed with denzo/scalepack or HKL2000 (Otwinowski, Z., Borek, D., Majewski, W. & Minor, W. Multiparametric scaling of diffraction intensities. *Acta Crystallogr A* 59, 228-34 (2003)).

[0539] PCSK9 / 31H4 crystals grew in the C2 space group with unit cell dimensions a=264.9, b=137.4, c=69.9 Å, β=102.8° and diffract to 2.3 Å resolution. The PCSK9 / 31H4 structure was solved by molecular replacement with the program MOLREP (The CCP4 suite: programs for protein crystallography. *Acta Crystallogr D Biol Crystallogr* 50, 760-3 (1994) using the PCSK9 structure (Piper, D.E. et al. The crystal structure of PCSK9: a regulator of plasma LDL-cholesterol. *Structure* 15, 545-52 (2007)) as the starting search model. Keeping the

PCSK9 31-692 solution fixed, an antibody variable domain was used as a search model. Keeping the PCSK9 31-692 / antibody variable domain solution fixed, an antibody constant domain was used as a search model. The complete structure was improved with multiple rounds of model building with Quanta and refinement with cnx. (Brunger, A.T. et al. Crystallography & NMR system: A new software suite for macromolecular structure determination. *Acta Crystallogr D Biol Crystallogr* 54, 905-21 (1998)).

[0540] PCSK9 / 31H4 / 21B12 crystals grew in the $P2_12_12$ space group with unit cell dimensions a=138.7, b=246.2, c=51.3 Å and diffract to 2.8 Å resolution. The PCSK9 / 31H4 / 21B12 structure was solved by molecular replacement with the program MOLREP using the PCSK9 ProCat / 31H4 variable domain as the starting search model. Keeping the PCSK9 ProCat / 31H4 variable domain fixed, a search for antibody constant domain was performed. Keeping the PCSK9 ProCat / 31H4 / 21B12 constant domain fixed, an antibody variable domain was used as a search model. The complete structure was improved with multiple rounds of model building with Quanta and refinement with cnx.

[0541] PCSK9 / EGFa domain crystals grew in the space group $P6_522$ with unit cell dimensions a=b=70.6, c=321.8 Å and diffract to 2.9 Å resolution. The PCSK9 / EGFa domain structure was solved by molecular replacement with the program MOLREP using the PCSK9 ProCat as the starting search model. Analysis of the electron density maps showed clear electron density for the EGFa domain. The LDLR EGFa domain was fit by hand and the model was improved with multiple rounds of model building with Quanta and refinement with cnx.

[0542] Core interaction interface amino acids were determined as being all amino acid residues with at least one atom less than or equal to 5 Å from the PCSK9 partner protein. 5 Å was chosen as the core region cutoff distance to allow for atoms within a van der Waals radius plus a possible water-mediated hydrogen bond. Boundary interaction interface amino acids were determined as all amino acid residues with at least one atom less than or equal to 8 Å from the PCSK9 partner protein but not included in the core interaction list. Less than or equal to 8 Å was chosen as the boundary region cutoff distance to allow for the length of an extended arginine amino acid. Amino acids that met these distance criteria were calculated with the program PyMOL. (DeLano, W.L. The PyMOL Molecular Graphics System. (Palo Alto, 2002)).

EXAMPLE 36

Cyrstal Structure of PCSK9 and 31A4

[0543] The crystal structure of the 31A4/PCSK9 complex was deteremined.

Expression and purification of protein samples

[0544] PCSK9 449TEV (a PCSK9 construct with a TEV protease cleavage site inserted between residue 449 and 450, numbering according to SEQ ID NO: 3) was expressed in baculovirus infected Hi-5 insect cells with an N-terminal honeybee melittin signal peptide followed by a His_6 tag. The PCSK9 protein was purified by first by nickel affinity chromatography. TEV protease was used to remove the melittin-His_6 tag and cleave the PCSK9 protein between the catalytic domain and V domain. The V domain was further purified by ion exchange chromatography and size exclusion chromatography. The 31A4 Fab fragment was expressed in *E. coli*. This protein was purified by nickel affinity chromatography, size exclusion chromatography and ion exchange chromatography.

Complex formation and crystallization

[0545] The PCSK9 V domain / 31A4 complex was made by mixing a 1.5 molar excess of PCSK9 V domain with 31A4 Fab. The complex was separated from excess PCSK9 V domain by purification on a size exclusion chromatography column. The PCSK9 V domain / 31A4 complex crystallized in 1.1 M Succinic acid pH 7, 2% PEG MME 2000.

Data collection and structure determination

[0546] The dataset for the PCSK9 V domain / 31A4 crystal was collected on a Rigaku FR-E x-ray source and processed with denzo/scalepack (Otwinowski, Z., Borek, D., Majewski, W. & Minor, W. Multiparametric scaling of diffraction intensities. *Acta Crystallogr A* 59, 228-34 (2003)).

[0547] PCSK9 V domain / 31A4 crystals grow in the $P2_12_12_1$ space group with unit cell dimensions a=74.6, b=131.1, c=197.9 Å with two complex molecules per asymmetric unit, and diffract to 2.2 Å resolution. The PCSK9 V domain / 31A4 structure was solved by molecular replacement with the program MOLREP (CCP4. The CCP4 suite: programs for protein crystallography. *Acta Crystallogr D Biol Crystallogr* 50, 760-3 (1994)) using the V domain of the PCSK9 structure (Piper, D.E. et al. The crystal structure of PCSK9: a regulator of plasma LDL-cholesterol. *Structure* 15, 545-52 (2007)) as the starting search model. Keeping the PCSK9 450-692 solution fixed, an antibody variable domain was used as a search model. After initial refinement, the antibody constant domains were fit by hand. The complete structure was

improved with multiple rounds of model building with Quanta and refinement with cnx (Brunger, A.T. et al. Crystallography & NMR system: A new software suite for macromolecular structure determination. *Acta Crystallogr D Biol Crystallogr* 54, 905-21 (1998)).

[0548] Core interaction interface amino acids were determined as being all amino acid residues with at least one atom less than or equal to 5 Å from the PCSK9 partner protein. 5 Å was chosen as the core region cutoff distance to allow for atoms within a van der Waals radius plus a possible water-mediated hydrogen bond. Boundary interaction interface amino acids were determined as all amino acid residues with at least one atom less than or equal to 8 Å from the PCSK9 partner protein but not included in the core interaction list. Less than or equal to 8 Å was chosen as the boundary region cutoff distance to allow for the length of an extended arginine amino acid. Amino acids that met these distance criteria were calculated with the program PyMOL (DeLano, W.L. The PyMOL Molecular Graphics System. (Palo Alto, 2002)). Distances were calculated using the V domain "A" and 31A4 "L1,H1" complex.

[0549] The crystal structure of the PCSK9 V domain bound to the Fab fragment of 31A4 was determined at 2.2 Å resolution. The depictions of the crystal structure are provided in FIGs. 21A-21D. FIGs. 21A-21C shows that the 31A4 Fab binds to the PCSK9 V domain in the region of subdomains 1 and 2.

[0550] A model of full length PCSK9 bound the 31A4 Fab was made. The structure of full length PCSK9 was overlaid onto the PCSK9 V domain from the complex. A figure of this model is shown in FIG. 21D. The site of the interaction between the EGFa domain of the LDLR and PCSK9 is highlighted.

[0551] Analysis of the structure shows where this antibody interacts with PCSK9 and demonstrated that antibodies that do not bind to the LDLR binding surface of PCSK9 can still inhibit the degradation of LDLR that is mediated through PCSK9 (when the results are viewed in combination with Example 40 and 41 below). In addition, analysis of the crystal structure allows for identification of specific amino acids involved in the interaction between PCSK9 and the 31A4 antibody. Furthermore, the core and boundary regions of the interface on the PCSK9 surface were also determined. Specific core PCSK9 amino acid residues of the interaction interface with 31A4 were defined as PCSK9 residues that are within 5 Å of the 31A4 protein. The core residues are T468, R469, M470, A471, T472, R496, R499, E501, A502, Q503, R510, H512, F515, P540, P541, A542, E543, H565, W566, E567, V568, E569, R592, and E593.

Boundary PCSK9 amino acid residues of the interaction interface with 31A4 were defined as PCSK9 residues that are 5-8 Å from the 31A4 protein. The boundary residues are as follows: S465, G466, P467, A473, I474, R476, G497, E498, M500, G504, K506, L507, V508, A511, N513, A514, G516, V536, T538, A539, A544, T548, D570, L571, H591, A594, S595, and H597. Amino acid residues nearly or completely buried within the PCSK9 protein are highlighted by underline. As noted herein, the numbering references the amino acid positions of SEQ ID NO: 3 (adjusted as noted herein).

[0552] Specific core 31A4 amino acid residues of the interaction interface with PCSK9 were defined as 31A4 residues that are within 5 Å of the PCSK9 protein. The core residues for the 31A4 antibody are as follows: Heavy Chain: G27, S28, F29, S30, A31, Y32, Y33, E50, N52, H53, R56, D58, K76, G98, Q99, L100, and V101; Light Chain: S31, N32, T33, Y50, S51, N52, N53, Q54, W92, and D94. Boundary 31A4 amino acid residues of the interaction interface with PCSK9 were defined as 31A4 residues that are 5-8 Å from the PCSK9 protein. The boundary residues for 31A4 are as follows: Heavy Chain: V2, G26, W34, N35, W47, I51, S54, T57, Y59, A96, R97, P102, F103, and D104; Light Chain: S26, S27, N28, G30, V34, N35, R55, P56, K67, V91, D93, S95, N97, G98, and W99.

[0553] The crystal structure also displayed the spatial requirements of this ABP in its interaction with PCSK9. As shown in this structure, surprisingly, antibodies that bind to PCSK9 without directly preventing PCSK9's interaction with the LDLR can still inhibit PCSK9's function.

[0554] In some embodiments, any antigen binding protein that binds to, covers, or prevents 31A4 from interacting with any of the above residues can be employed to bind to or neutralize PCSK9. In some embodiments, the ABP binds to or interacts with at least one of the following PCSK9 (SEQ ID NO: 3) residues: T468, R469, M470, A471, T472, R496, R499, E501, A502, Q503, R510, H512, F515, P540, P541, A542, E543, H565, W566, E567, V568, E569, R592, and E593. In some embodiments, the ABP is within 5 angstroms of one or more of the above residues. In some embodiments, the ABP binds to or interacts with at least one of the following PCSK9 (SEQ ID NO: 3) residues: S465, G466, P467, A473, I474, R476, G497, E498, M500, G504, K506, L507, V508, A511, N513, A514, G516, V536, T538, A539, A544, T548, D570, L571, H591, A594, S595, and H597. In some embodiments, the ABP is 5 to 8 angstroms

from one or more of the above residues. In some embodiments, the ABP interacts, blocks, or is within 8 angstroms of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, or 50 of the above residues.

[0555] The coordinates for the crystal structures discussed in the above Examples are are presented in Table 35.1 (full length PCSK9 and 31H4), Table 35.2 (PCSK9 and EGFa), Table 35.3 (PCSK9, 31H4, and 21B12), and Table 35.4 (PCSK9 and 31A4). Antigen binding proteins and molecules that interact with the relevant areas or residues of the structure of PCSK9 (including those areas or residues within 15, 15-8, 8, 8-5, 5, or fewer angstroms from where EGFa, or the antibodies, interact with PCSK9) depicted in the figures and/or their corresponding positions on the structures from the coordinates are also contemplated.

[0556] The antibodies that are described in the coordinates were raised in E. coli and thus possess some minor amino acid differences from the fully human antibodies. The first residue in the variable region was a glutamic acid instead of a glutamine for the heavy and light chains of 21B12 and for the light chain for 31H4. In addition to the differences in the sequence of variable region, there were also some differences in the constant region of the antibodies described by the coordinates (again due to the fact that the antibody was raised in E. coli). FIG. 22 highlights (via underlining shading, or bold) the differences between the constant regions of the 21B12, 31H4, and 31A4 Fabs (raised in E. coli) when compared to SEQ ID NOs: 156, and 155. For 21B12 31H4, and 31A4, the light chain constant sequence is similar to human lambda (SEQ ID NO: 156). The underlined glycine residue is an insertion between where the 21B12 and 31H4 variable sequences stop and the lambda sequence starts.

[0557] For both 21B12 and 31H4, the heavy chain constant is similar to human IgG4 (SEQ ID NO: 155). The highlighted differences in FIG. 22 are shown in Table 36.1:

Table 36.1

Crystal SEQ	ID NO: 155
S	C
K	R
G	E
G	S
Q	K
I	T
N	D
K	R
P	S

[0558] In regard to 31A4, while it also has the same distinctions noted above, there are three additional differences. As shown in FIG. 22, there are two additional amino acids at the start, *which comes from incomplete processing of the signal peptide in E. coli expression.* In addition, there is one additional substitution in the 31A4 heavy chain constant region when compared to SEQ ID NO: 155, which is the adjustment of a L (in SEQ ID NO: 155) to a H. Finally, 31A4 does have a glutamine as the initial amino acid of the Fab, rather than the adjustment to glutamic acid noted above for 21B12 and 31H4.

[0559] For all three antibodies, the end of the heavy chain (boxed in dark grey) differs as well, but the amino acids are not ordered in the structure so they do not appear in the cooridnates. As will be appreciated by one of skill in the art, his-tags are not a required part of the ABP and should not be considered as part of the ABP's sequence, unless explicitly called out by reference to a specific SEQ ID NO that includes a histidine tag and a statement that the ABP sequence "includes the Histidine tag."

EXAMPLE 37
Epitope Mapping--Binning

[0560] An alternative set of binning experiments was conducted in addition to the set in Example 10. As in Example 10, ABPs that compete with each other can be thought of as binding to the same site on the target and in common parlance are said to "bin" together.

[0561] A modification of the Multiplexed Binning method described by Jia, et al (J. Immunological Methods, 288 (2004) 91-98) was used. Individual bead codes of streptavidin-coated Luminex beads was incubated in 100ul 0.5 ug/ml biotinylated monovalent mouse-anti-human IgG capture antibody (BD Pharmingen, #555785) for 1 hour at room temperature in the dark, then washed 3x with PBSA, phosphate buffered saline (PBS) plus 1% bovine serum albumin (BSA). Each bead code was separately incubated with 100 ul 2 ug/ml anti-PCSK9 antibody (Coating Antibody) for 1 hour then washed 3x with PBSA. The beads were pooled then dispensed to a 96-well filter plate (Millipore, #MSBVN1250). 100ul of 2 ug/ml purified PCSK9 protein was added to half the wells. Buffer was added to the other half as control. The reaction was incubated for 1 hour then washed. 100 ul of a 2 ug/ml anti-PCSK9 antibody (Detection Ab) was added to all the wells, incubated for 1 hour then washed. An irrelevant human-IgG (Jackson, #009-000-003) was run as another control. 20ul PE-conjugated

monovalent mouse-anti-human IgG (BD Pharmingen, #555787) was added to each well and incubated for 1 hour then washed. Beads were resuspended in 100ul PBSA and a minimum of 100 events/bead code were collected on the BioPlex instrument (BioRad).

[0562] Median Fluorescent Intensity (MFI) of the antibody pair without PCSK9 was subtracted from signal of the corresponding reaction containing PCSK9. For the antibody pair to be considered bound simultaneously, and therefore in different bins, the subtracted signal had to be greater than 3 times the signal of the antibody competing with itself and the 3 times the signal of the antibody competing with the irrelevant antibody.

[0563] The data from the above is depicted in FIGs. 23A-23D. The ABPs fell into five bins. The shaded boxes indicate ABPs that can bind simultaneously to PCSK9. The nonshaded boxes indicate those ABPs that compete with each other for binding. A summary of the results is shown in Table 37.1.

Table 37.1.

BIN 1	BIN 2	BIN 3	BIN 4	BIN 5
01A12.2	27B2.1	16F12.1	11G1.5	30A4.1
03B6.1	27B2.5	22E2.1	03C4.1	13B5.1
09C9.1	12H11.1	27A6.1		13H1.1
17C2.1		28B12.1		31A4.1
21B12.2		28D6.1		31B12.1
23G1.1		31G11.1		
25G4.1		31H4.1		
26E10.1		08A1.2		
11H4.1		08A3.1		
11H8.1		11F1.1		
19H9.2				
26H5.1				
27E7.1				
27H5.1				
30B9.1				
02B5.1				
23B5.1				
27B2.6				
09H6.1				

[0564] Bins 1 (competes with ABP 21B12) and 3 (competes with 31H4) are exclusive of each other; bin 2 competes with bins 1 and 3; and Bin 4 does not compete with bins 1 and 3. Bin 5, in this example, is presented as a "catch all" bin to describe those ABPs that do not fit into the other bins. Thus, the above identified ABPs in each of the binds are representative of different types of epitope locations on PCSK9, some of which overlap with each other.

[0565] As will be appreciated by one of skill in the art, if the reference ABP prevents the binding of the probe ABP then the antibodies are said to be in the same bin. The order in which the ABPs are employed can be important. If ABP A is employed as the reference ABP and blocks the binding of ABP B the converse is not always true: ABP B used as the reference ABP will not necessarily block ABP A. There are a number of factors in play here: the binding of an ABP can cause conformational changes in the target which prevent the binding of the

second ABP, or epitopes which overlap but do not completely occlude each other may allow for the second ABP to still have enough high-affinity interactions with the target to allow binding. ABPs with a much higher affinity may have a greater ability to bump a blocking ABP out of the way. In general, if competition is observed in either order the ABPs are said to bin together, and if both ABPs can block each other then it is likely that the epitopes overlap more completely.

EXAMPLE 38

Epitope Mapping—Western Blot

[0566] The present example demonstrates whether or not the epitopes for the examined ABPs were linear or conformational. Denaturing reducing and denaturing non-reducing western blots were run to determine which antibodies have a conformational epitope. Antibodies that bind to a denaturing reducing western blot have a linear epitope and are not conformational. The results are presented in FIG. 24A and FIG. 24B. For the blot, 0.5 ug/lane of purified full-length human PCSK9 was run on a 4-12% NuPAGE Bis-Tris gel and MES SDS Running Buffer. 1 ug/ml anti-PCSK9 antibodies, except 0.5 ug/ml 31G11, were used to probe the blot. 1:5000 donkey-anti-human-IR700 secondary was used and read on a LiCOR instrument. Antibody 13H1 bound to a linear epitope on the pro-domain of PCSK9. All other antibodies displayed results that were consistent with conformational epitopes. These gels split apart the pro-domain from the rest of the protein, and the pro domain ran at about 15kDa. In addition, 3C4 and 31A4 appeared to bind to conformational epitopes which were preserved by disulfide bonds, as these antibodies bound to PCSK-9 under denaturing conditions where the disulfide bonds had been preserved (left) but reducing the samples (right) eliminated binding.

EXAMPLE 39

Epitope Mapping--Arginine / Glutamic Acid Scanning

[0567] Representative ABPs from each bin (from Example 37) were selected for further epitope analysis. An arginine/glutamic acid-scanning strategy was performed for mapping ABP binding to PCSK9. By way of background, this method determines if a residue is part of the structural epitope, meaning those residues in the antigen which contact or are buried by the antibody. Arginine and glutamic acid sidechains are charged and bulky and can disrupt antibody binding even if the mutated residue is not directly involved in antibody binding.

Residue Selection

[0568] The crystal structure of PCSK9 was used to select the residues to be mutated for epitope mapping. The method used to choose residues to mutate involved both computational mechanisms and interactive structure analysis. The PCSK9 structure contained gaps of missing residues and was missing 30 amino acids in the N- (i.e., the signal sequence) and 10 amino acids in the C-termini. The internal missing residues were modeled onto the structure, but the N- and C-terminal missing residues were not. The solvent exposure ratio for each residue was calculated: the surface area of each residue in the context of the protein (SA1) was divided by the surface area of the residue in a trimer with flanking glycines (SA2) with a conserved backbone structure. Residues with solvent exposure ratio greater than 10% (R10) were selected as well as the 40 missing terminal residues. From these, prolines and glycines with positive Φ angles were excluded to reduce the possibility of misfolding. The number of residues to be mutated in the V domain was reduced by using a solvent exposure ratio of 37% along with visual inspection of the entire protein to bring the total number of mutations to 285. Various orientations of the surface of PCSK9 with these various classes identifies are shown in FIG. 25A-25F. In these figures, lightest gray denotes areas that were not selected or were deselected. darker gray denotes those residues selected).

Cloning and Expression

[0569] Once the residues to be altered were identified, the various residues were altered. Human PCSK9 was cloned into the pTT5 vector with a C-terminal Flag-His tag. Mutants were made from this original construct by site-directed mutagenesis using a QuikChange II kit from Stratagene. Sense and anti-sense oligonucleotides used for mutagenesis were designed using Amgen's MutaGenie software. All PCSK9 constructs were expressed in transiently-transfected 293-6E cells in 24-well plates and re-racked into three 96-well plates with a non-mutated PCSK9 control (wild-type, WT) in each plate. Expression levels and integrity of the recombinant proteins in conditioned media were checked by Western blot. Of the 285 mutants originally selected, 41 failed in cloning or expression. 244 mutants were used for epitope mapping. An alignment of the PCSK9 parent sequence and a representative PCSK9 sequence with the 244 mutated residues is shown in FIG. 26. Separate constructs were made containing a single mutation. For the purposes of the epitope sequences and the epitope based

inventions involving changes in binding, the sequences are provided in reference to SEQ ID NO: 1 and/or SEQ ID NO: 303. The sequences in FIG. 26 were the sequences used for the present binding epitope studies. One of skill in the art will appreciate that the present results apply to other PCSK9 variants disclosed herein as well (*e.g.*, SEQ ID NO: 1 and 3, as well as the other allelic variants).

[0570] Five antibodies, a representative of each bin, were chosen for fine epitope mapping. They were 21B12, 31H4, 12H11, 31A4, 3C4. All conformational epitope antibodies. Three, 21B12, 31H4, and 31A4 were also crystallized with PCSK9, as described above.

Structural and functional epitopes

[0571] Epitopes can be further defined as structural or functional. Functional epitopes are generally a subset of the structural epitopes and have those residues that directly contribute to the affinity of the interaction (e.g. hydrogen bonds, ionic interactions). Structural epitopes can be thought of as the patch of the target which is covered by the antibody.

[0572] The scanning mutagenesis employed was an arginine and glutamic acid scan. These two sidechains were chosen due to their large steric bulk and their charge, which allows mutations that occur in the structural epitope to have a greater effect on antibody binding. Arginine was generally employed except when the WT reside was arginine, and in these cases the residue was mutated to glutamic acid to switch the charge.

[0573] For the purpose of epitope mapping, a bead-based multiplexed assay was used to measure antibody binding to PCSK9 and PCSK9 mutants simultaneously. Antibody binding to mutants was then compared to its binding to the wild-type in the same well. The variants were split into three groups: Group 1: 81 variants + 2 wt controls + 1 negative control + 1 other PCSK9 supernatant; Group 2: 81 variants + 2 wt controls + 2 negative controls; and Group 3: 82 variants + 2 wt control + 1 negative control.

[0574] The assay was run as follows: 85 sets of color-coded strepavidin-coated LumAvidin beads (Luminex) were bound with biotinylated anti-pentaHis antibody (Qiagen, #1019225) for 1 hour at room temperature (RT) then washed three times in PBS, 1% BSA, 0.1% Tween 20. Each color-coded bead set was then allowed to bind to a PCSK9 mutant, wild-type, or negative control in 150 ul supernatant overnight at 4˚C

[0575] The color-coded bead sets, each associated to a specific protein, were washed and pooled. At this point, there were 3 pools of 85 bead sets, one pool for each group of mutants and controls. The beads from each pool were aliquoted to 24 wells (3 columns) of a 96-well filter plate (Millipore, #MSBVN1250). 100 ul of anti-PCSK9 antibodies in 4-fold dilutions were added to nine columns for triplicate points and incubated for 1 hour at RT and washed. 100ul of 1:200 dilution phycoerythrin (PE)-conjugated anti-human IgG Fc (Jackson Immunoresearch, #109-116-170) was added to each well and incubated for 1 hour at RT and washed.

[0576] Beads were resuspended in 1% BSA in PBS, shaken for 10mins and read on the BioPlex instrument (Bio-Rad). The instrument identifies each bead by its color-code thereby identifying the specific protein associated with the color code. At the same time, it measures the amount of antibody bound to the beads by fluorescence intensity of the PE dye. Antibody binding to each mutant can then be compared directly its binding to the wild type in the same pool. IL-17R chimera E was used as a negative control. A summary of all of the mutants examined is shown in Table 39.1 (with reference to the sequence numbering used in FIG. 1A and 26).

Table 39.1

	1	2	3	4	5	6	7	8	9	10	11	12
A	WT PCSK9	Y8R	E18R	P26R	A38R	T56R	A70R	H83R	E102R	L128R	D145R	
B	Q1R	E9R	E19R	E27R	K39R	H57R	Q71R	V84R	L105R	E129R	S148R	
C	E2R	E10R	D20R	G29R	D40R	L58R	A73R	H86R	K106R	R130E	pcsk9 supe test	
D	D3R	L11R	G21R	T30R	L44R	Q60R	R74E	K95R	H109R	T132R	IL17R chimera E	
E	E4R	V12R	L22R	T31R	T47R	E62R	R75E	S97R	D111R	D139R	WT PCSK9	
F	D5R	A14R	A23R	A32R	K53R	R63E	Y77R	G98R	A121R	E140R		
G	G6R	L15R	E24R	T33R	E54R	R66E	L78R	D99R	S123R	Y141R		
H	D7R	S17R	A25R	H35R	E55R	R67E	L82R	L101R	W126R	Q142R		

	1	2	3	4	5	6	7	8	9	10	11	12
A	WT PCSK9	M171R	E181R	Q189R	K213R	R242E	G251R	L294R	L321R	Q352R	E380R	
B	L149R	V172R	D182R	A190R	G214R	K243R	G252R	A311R	E336R	M366R	R384E	
C	S158R	T173R	G183R	S191R	S216R	S244R	R265E	Q312R	D337R	S371R	IL17R chimera E	
D	Q160R	D174R	T184R	K192R	R221E	Q245R	A269R	D313R	D344R	A372R	IL17R chimera E	
E	S161R	E176R	R185E	S195R	Q226R	L246R	Q272R	Q314R	T347R	E373R	WT PCSK9	
F	D162R	N177R	F186R	H196R	K228R	V247R	R276E	T317R	F349R	E375R		
G	R164E	V178R	H187R	R207E	T230R	Q248R	A277R	L318R	V350R	T377R		
H	E167R	E180R	R188E	D208R	F240R	V250R	R289E	T320R	S351R	L378R		

	1	2	3	4	5	6	7	8	9	10	11	12
A	WT PCSK9	N395R	V405R	W423R	R446E	E513R	Q525R	Q554R	Q589R	S632R	A641R	
B	I386R	E396R	N409R	Q424R	D460R	A514R	E537R	N556R	Q591R	T633R	R650E	
C	H387R	A397R	A413R	A433R	A472R	S515R	V538R	K579R	A595R	T634R	R652E	
D	F388R	W398R	S417R	H434R	F485R	M516R	E539R	V580R	E597R	G635R	IL17R chimera E	
E	A390R	E401R	T418R	T438R	G486R	R519E	L541R	K581R	E598R	S636R	WT PCSK9	
F	K391R	D402R	H419R	R439E	E488R	H521R	H544R	E582R	V620R	T637R		
G	D392R	Q403R	G420R	M440R	N503R	H523R	V548R	H583R	R629E	S638R		
H	V393R	R404E	A421R	T442R	T508R	Q524R	R552E	G584R	V631R	E639R		

Bead Variability Study

[0577] Before running the epitope mapping binding assay, a validation experiment was conducted to assess the "bead region" to "bead region" (B-B) variability. In the validation experiment, all beads were conjugated with the same wild type control protein. Therefore, the difference between beads regions was due to purely B-B variance and was not confounded by difference between wild type and mutant proteins. The titration of antibody was run with twelve replications in different wells.

[0578] The objective of this statistical analysis was to estimate the B-B variability of the estimated EC50 of binding curves. The estimated B-B standard deviation (SD) was then used to build the EC50 confidence intervals of wild type and mutant proteins during curve comparison experiments.

[0579] A four-parameter logistic model was fitted to the binding data for each bead region. The resulting file, containing curve quality control (QC) results and parameter estimates

for top (max), bottom (min), Hillslope (slope), and natural log of EC50 (xmid) of the curves, was used as the raw data for the analysis. B-B variability for each parameter was then estimated by fitting mixed effect model using SAS PROC MIXED procedure. Only curves with "good" QC status were included in the analysis. The final mixed effect model included only residual (i.e. individual bead regions) as random effect. Least squares means (LS-mean) for each parameter were estimated by the mixed effect model as well. B-B SD was calculated by taking square root of B-B variance. Fold change between LS-mean + 2SD and LS-mean - 2SD, which represent approximately upper and lower 97.5 percentile of the population, was also calculated. The results are displayed in Table 39.2

Table 39.2 Least square mean and bead-to-bead variance estimations

Assay ID	parname	Ls Mean	B-B Variance	-2SD	+2SD	Fold Change*
PCSK9	max	15000	997719	13002.3	16997.7	1.3
PCSK9	min	162.09	1919.66	74.5	249.7	3.4
PCSK9	slope	0.8549	0.000599	0.8	0.9	1.1
PCSK9	xmid	3.1715	0.002098	3.1	3.3	1.2

* xmid is natural log of the EC50. Fold change for xmid was converted back to original scale.

Identifying Residues in the Structural Epitope

[0580] A residue was considered part of the structural epitope (a "hit") when mutating it to arginine or glutamic acid alters antibody binding. This is seen as a shift in the EC50 or a reduction of maximum signal compared to antibody binding to wild type. Statistical analyses of antibody binding curves to wild type and mutants were used to identify statistically significant EC50 shifts. The analysis takes into consideration variation in the assay and curve fitting.

Hit Identification based on EC50 Comparison

[0581] The EC50 and Bmax values were generated from a Weighted 4-Parameter Logistical model fitted to the binding data using S-PLUS with VarPower software (Insightful Corporation, Seattle WA). The EC50s of the mutant binding curves and wild type binding curves were compared. Statistically significant differences were identified as hits for further consideration. The curves with "nofit" or "badfit" flags were excluded from the analysis.

The Variations in EC50 Estimates

[0582] Two sources of variations were considered in the comparison of EC50 estimates, variation from the curve fit and the bead-bead variation. Wild types and mutants were linked to different beads, hence their difference are confounded with the bead-bead difference (described above). The curve fit variation was estimated by the standard error of the log EC50 estimates. Bead-bead variation was experimentally determined using an experiment where wild type controls were linked to each one of the beads (described above). The bead variation in EC50 estimates of wild type binding curve from this experiment was used to estimate the bead-bead variation in the actual epitope mapping experiment.

Testing for EC50 Shift between Mutants and Wild Type

[0583] The comparisons of two EC50s (in log scale) was conducted using Student's t-test. The t-statistic was calculated as the ratio between delta (the absolute differences between EC50 estimates) and the standard deviation of delta. The variance of delta was estimated by the sum of the three components, variance estimate of EC50 for mutant and wild type curves in the nonlinear regression and two times the bead-bead variance estimated from a separate experiment. The multiple of two for the bead-bead variance was due to the assumption that both mutant and wild type beads had the same variance. The degree of freedom of the standard deviation of delta was calculated using the Satterthwaite's (1946) approximation. Individual p-values and confidence intervals (95% and 99%) were derived based on Student's t distribution for each comparison. In the case of multiple wild type controls, a conservative approach was taken by picking the wild type control that was most similar to the mutant, i.e., picking the ones with the largest p-values.

[0584] Multiplicity adjustments were important to control the false positive(s) while conducting a large number of tests simultaneously. Two forms of multiplicity adjustment were implemented for this analysis: family wise error (FWE) control and false discovery rate (FDR) control. The FWE approach controls the probability that one or more hits are not real; FDR approach controls the expected proportion of false positive among the selected hits. The former approach is more conservative and less powerful than the latter one. There are many methods available for both approaches, for this analysis, the Hochberg's (1988) method for FWE analysis

and Benjamini-Hochberg's (1995) FDR method for FDR analysis were selected. Adjusted p-values for both approaches were calculated.

Results

EC50 Shift

[0585] Mutations whose EC50 is significantly different from wild type, e.g., having a False Discovery Rate adjusted p-value for the whole assay of 0.01 or less, were considered part of the structural epitope. All the hits also had a Familywise type I error rate adjusted p-value for each antibody of less than 0.01 except residue R185E for antibody 31H4 which had an FWE adjusted p-value per antibody of 0.0109. The residues in the structural epitope of the various antibodies determined by EC50 shift are shown in Table 39.3 (point mutations are with reference to SEQ ID NO: 1 and 303)

Table 39.3

Antibody	Mutation	FDR Adjusted Pval	FWE Adjusted By Pval	Low99	Low95	FoldChange	High95	High99	RawPval
21B12	D208R	0.0000	0.0000	0.3628	0.3844	0.4602	0.5509	0.5837	0.0000
21B12	R207E	0.0000	0.0000	1.7148	1.8488	2.3191	2.9090	3.1364	0.0000
31H4	R185E	0.0024	0.0109	1.2444	1.3525	1.7421	2.2439	2.4388	0.0000
31A4	E513R	0.0001	0.0003	1.4764	1.6219	2.1560	2.8660	3.1485	0.0000
31A4	E539R	0.0000	0.0000	1.6014	1.7461	2.2726	2.9578	3.2252	0.0000
31A4	R439E	0.0000	0.0000	3.1565	3.6501	5.5738	8.5113	9.8420	0.0000
31A4	V536R	0.0004	0.0013	1.4225	1.5700	2.1142	2.8471	3.1423	0.0000
12H11	A390R	0.0000	0.0001	1.4140	1.5286	1.9389	2.4594	2.6588	0.0000
12H11	A413R	0.0009	0.0028	1.2840	1.3891	1.7653	2.2434	2.4269	0.0000
12H11	S351R	0.0009	0.0028	1.2513	1.3444	1.6761	2.0896	2.2452	0.0000
12H11	T132R	0.0000	0.0001	1.3476	1.4392	1.7631	2.1599	2.3068	0.0000
3C4	E582R	0.0016	0.0069	1.3523	1.5025	2.0642	2.8359	3.1509	0.0000

Maximum Signal Reduction

[0586] The percent maximum signal was calculated using the maximum signal from the curve fitting (BmaxPerWT) and raw data point (RawMaxPerWT). Mutations that reduced the antibody binding maximum signal by ≥ 70% as compared to to wild type signal or that reduced the signal of one antibody compared to other antibodies by >50% when all other antibodies are at least 40% of wild type were considered hits and part of the epitope. Table

39.4 displays the residues that are in the structural epitope (*italics*) as determined by reduction of maximum signal.

Table 39.4

antibody	Mutants	BmaxPerWT	RawMaxPerWT
21B12	A311R	141.6388	139.7010
31H4	A311R	145.2189	147.8244
31A4	A311R	103.4377	96.2214
12H11	*A311R*		*14.9600*
3C4	A311R	129.0460	131.2060
21B12	*D162R*		*7.0520*
31H4	D162R	108.8308	112.4904
31A4	D162R	98.8873	95.9268
12H11	D162R	94.6280	97.4928
3C4	D162R	101.4281	100.1586
21B12	D313R	45.8356	45.0011
31H4	D313R	45.6242	44.9706
31A4	D313R	47.9726	44.7741
12H11	*D313R*	*16.1811*	*18.4262*
3C4	D313R	58.5269	57.6032
21B12	D337R	61.9070	62.2852
31H4	D337R	63.1604	64.1029
31A4	D337R	62.9124	59.4852
12H11	*D337R*		*10.8443*
3C4	D337R	73.0326	73.9961
21B12	E129R	139.9772	138.9671
31H4	E129R	141.6792	139.1764
31A4	E129R	77.3005	74.8946
12H11	*E129R*	*28.6398*	*29.3751*
3C4	E129R	85.7701	85.7802

antibody	Mutants	BmaxPerWT	RawMaxPerWT
21B12	*E167R*		*15.1082*
31H4	E167R	127.4479	128.2698
31A4	E167R	115.3403	112.6951
12H11	E167R	111.0979	109.6813
3C4	E167R	109.3223	108.7864
21B12	H521R	133.8480	133.9791
31H4	H521R	130.2068	128.4879
31A4	H521R	124.5091	129.3218
12H11	H521R	130.7979	134.4355
3C4	*H521R*		*22.1077*
21B12	Q554R	125.9594	125.2103
31H4	Q554R	122.2045	128.7304
31A4	Q554R	113.6769	121.3369
12H11	Q554R	116.1789	118.4170
3C4	*Q554R*		*31.8416*
21B12	*R164E*	*17.3807*	*19.8505*
31H4	R164E	97.8218	99.6673
31A4	R164E	98.2595	96.3352
12H11	R164E	88.0067	89.8807
3C4	R164E	105.0589	105.7286
21B12	R519E	139.4598	141.2949
31H4	R519E	135.5609	140.0000
31A4	R519E	134.2303	137.1110
12H11	R519E	135.4756	137.0824
3C4	*R519E*		*44.0091*
21B12	S123R	87.6431	88.1356
31H4	S123R	85.5312	84.7668
31A4	S123R	68.4371	66.8131
12H11	*S123R*	*20.8560*	*20.6910*
3C4	S123R	73.6475	71.5959

(Point mutations are with reference to SEQ ID NO: 1 and FIG. 26).

[0587] Table 39.5 displays a summary of all of the hits for the various antibodies.

Table 39.5

EC50 shift hits					Bmax shift hits				
21B12	31H4	31A4	12H11	3C4	21B12	31H4	31A4	12H11	3C4
R207E	R185E	R439E	T132R	E582R	D162R			S123R	R519E
D208R*		E513R	S351R		R164E			E129R	H521R
		V538R	A390R		E167R			A311R	Q554R
		E539R	A413R					D313R	
								D337R	

* decreases EC50

[0588] To further examine how these residues form part of or all of the relevant epitopes, the above noted positions were mapped onto various crystal structure models, the results are shown in FIG. 27A through 27E. FIG. 27A depicts the 21B12 epitope hits, as mapped onto a crystal structure of PCSK9 with the 21B12 antibody. The structure identifies PCSK9 residues as follows: light gray indicates those residues that were not mutated (with the exception of those residues that are explicitly indicated on the structure) and darker gray indicates those residues mutated (a minority of which failed to express). Residues that are explicitly indicated were tested (regardless of the shading indicated on the figure) and resulted in a significant change in EC50 and/or Bmax The epitope hits were based on Bmax shift. In this figure, 31H4 is behind 21 B12.

[0589] FIG. 27B depicts the 31H4 epitope hits, as mapped onto a crystal structure of PCSK9 with 31H4 and 21B12 antibodies. The structure identifies PCSK9 residues as follows: light gray indicates those residues that were not mutated (with the exception of those residues that are explicitly indicated on the structure) and darker gray indicates those residues mutated (a minority of which failed to express). Residues that are explicitly indicated were tested (regardless of the shading indicated on the figure) and resulted in a significant change in EC50 and/or Bmax . The epitope hits were based on the EC50 shift.

[0590] FIG. 27C depicts the 31A4 epitope hits, as mapped onto a crystal structure of PCSK9 with 31H4 and 21B12 antibodies. The structure identifies PCSK9 residues as follows: light gray indicates those residues that were not mutated (with the exception of those residues that are explicitly indicated on the structure) and darker gray indicates those residues mutated (a minority of which failed to express). Residues that are explicitly indicated were tested

(regardless of the shading indicated on the figure) and resulted in a significant change in EC50 and/or Bmax. The epitope hits were based on the EC50 shift. 31A4 antibody is known to bind to the V-domain of PCSK9, which appears consistent with the results presented in FIG. 27C.

[0591] FIG. 27D depicts the 12H11 epitope hits, as mapped onto the crystal structure of PCSK9 with 31H4 and 21B12 antibodies. The structure identifies PCSK9 residues as follows: light gray indicates those residues that were not mutated (with the exception of those residues that are explicitly indicated on the structure) and darker gray indicates those residues mutated (a minority of which failed to express). Residues that are explicitly indicated were tested (regardless of the shading indicated on the figure) and resulted in a significant change in EC50 and/or Bmax. 12H11 competes with 21B12 and 31H4 in the binning assay described above.

[0592] FIG. 27E depicts the 3C4 epitope hits, as mapped onto the crystal structure of PCSK9 with 31H4 and 21B12 antibodies. The structure identifies PCSK9 residues as follows: light gray indicates those residues that were not mutated (with the exception of those residues that are explicitly indicated on the structure) and darker gray indicates those residues mutated (a minority of which failed to express). Residues that are explicitly indicated were tested (regardless of the shading indicated on the figure) and resulted in a significant change in EC50 and/or Bmax.

[0593] 3C4 does not compete with 21B12 and 31H4 in the binning assay. 3C4 binds to the V-domain in the domain binding assay (see results from Example 40, FIGs. 28A and 28B).

[0594] While there were approximately a dozen mutants that could have been expected to have an effect on binding (based upon the crystal structure), the present experiment demonstrated that, surprisingly, they did not. As will be appreciated by one of skill in the art, the results presented above are in good agreement with the crystal structures and PCSK-9's binding of these antibodies. This demonstrates that the provided structural and corresponding functional data adequately identifies the key residues and areas of interaction of the neutralizing ABPs and PCSK9. Thus, variants of the ABPs that possess the ability to bind to the above noted areas are adequately provided by the present description.

[0595] As will be appreciated by one of skill in the art, while the B-max drop and EC50 shift hits can be considered manifestations of the same phenomenon, strictly speaking, a B-max drop alone does not reflect a loss of affinity per se but, rather, the destruction of some percentage of the epitope of an antibody. Although there is no overlap in the hits determined by

B-max and EC50, mutations with a strong affect on binding may not allow for the generation of a useful binding curve and hence, no EC50 can be determined for such variants.

[0596] As will be appreciated by one of skill in the art, ABPs in the same bin (with the exception of bin 5, which as noted above, is a general catch all bin) likely bind to overlapping sites on the target protein. As such, the above epitopes and relevant residues can generally be extended to all such ABPs in the same bin.

[0597] To further examine the above results in regard to ABP 31H4, position E181R, which, according to the above crystal structure, was predicted to interact with R185 to form part of the surface that interacts with the ABP, was also altered (E181R). The results, while not statistically significant on their own, were, when combined with the crystal structure, demonstrative of 31H4 interacting with E181R (data not shown). Thus, position 181 also appears to form part of the epitope for the 31H4 ABP.

[0598] As noted above, the above binding data and epitope characterization references a PCSK9 sequence (SEQ ID NO: 1) that does not include the first 30 amino acids of PCSK9. Thus, the numbering system of this protein fragment, and the SEQ ID NO:s that refer to this fragment, are shifted by 30 amino acids compared to the data and experiments that used a full length PCSK9 numbering system(such as that used in the crystal study data described above). Thus, to compare these results, an extra 30 amino acids should be added to the positions in each of the above epitope mapping results. For example, position 207 of SEQ ID NO: 1 (or SEQ ID NO: 303), correlates to position 237 of SEQ ID NO: 3 (the full length sequence, and the numbering system used throughout the rest of the specification). Table 39.6 outlines how the above noted positions, which reference SEQ ID NO: 1 (and/or SEQ ID NO: 303) correlate with SEQ ID NO: 3 (which includes the signal sequence).

TABLE 39.6

AMINO ACID POSITION IN SEQ ID NO: 1 (EPITOPE DATA)	AMINO ACID POSITION IN SEQ ID NO: 3 (EPITOPE DATA)
207	237
208	238
185	215
181	211
439	469
513	543
538	568

539	569
132	162
351	381
390	420
413	443
582	612
162	192
164	194
167	197
123	153
129	159
311	341
313	343
337	367
519	549
521	551
554	584

[0599] Thus, those embodiments described herein with reference to SEQ ID NO: 1 can also be described, by their above noted corresponding position with reference to SEQ ID NO: 3.

EXAMPLE 40

PCSK9 Domain Binding Assay

[0600] The present example examined where on PCSK9 the various ABPs bound.

[0601] Clear, 96 well maxisorp plates (Nunc) were coated overnight with 2 ug/ml of various anti-PCSK9 antibodies diluted in PBS. Plates were washed thoroughly with PBS/.05% Tween-20 and then blocked for two hours with 3% BSA/PBS. After washing, plates were incubated for two hours with either full length PCSK9 (aa 31-692 SEQ ID NO: 3, procat PCSK9 (aa 31-449 SEQ ID NO: 3) or v-domain PCSK9 (aa 450-692 of SEQ ID NO: 3) diluted in general assay diluent (Immunochemistry Technologies, LLC). Plates were washed and a rabbit polyclonal biotinylated anti-PCSK9 antibody (D8774), which recognizes the procat and v-domain as well as full-length PCSK9, was added at 1 ug/ml (in 1%BSA/PBS). Bound full-length, procat or v-domain PCSK9 was detected by incubation with neutravidin-HRP (Thermo Scientific) at 200 ng/ml (in 1% BSA/PBS) followed by TMB substrate (KPL) and absorbance measurement at 650 nm. The results, presented in FIGS. 28A and 28B, demonstrate the ability of the various ABS to bind to various parts of PCSK9. As shown in FIG. 28B, ABP 31A4 binds to the V domain of PCSK9.

EXAMPLE 41

Neutralizing, non-competitive antigen binding proteins

[0602] The present example demonstrates how to identify and characterize an antigen binding protein that is non-competitive with LDLR for binding with PCSK9, but is still neutralizing towards PCSK9 activity. In other words, such an antigen binding protein will not block PCSK9 from binding to LDLR, but will prevent or reduce PCSK9 mediated LDLR degradation.

[0603] Clear, 384 well plates (Costar) were coated with 2 ug/ml of goat anti-LDL receptor antibody (R&D Systems) diluted in buffer A (100 mM sodium cacodylate, pH 7.4). Plates were washed thoroughly with buffer A and then blocked for 2 hours with buffer B (1% milk in buffer A). After washing, plates were incubated for 1.5 hours with 0.4 ug/ml of LDL receptor (R&D Systems) diluted in buffer C (buffer B supplemented with 10 mM $CaCl_2$). Concurrent with this incubation, 20 ng/ml of biotinylated D374Y PCSK9 was incubated with 100 ng/ml of antibody diluted in buffer A or buffer A alone (control). The LDL receptor containing plates were washed and the biotinylated D374Y PCSK9/antibody mixture was transferred to them and incubated for 1 hour at room temperature. Binding of the biotinylated D374Y to the LDL receptor was detected by incubation with streptavidin-HRP (Biosource) at 500 ng/ml in buffer C followed by TMB substrate (KPL). The signal was quenched with 1N HCl and the absorbance read at 450 nm. The results are presented in FIG. 28C, which shows that while ABP 31H4 inhibits LDLR binding, ABP 31A4 does not inhibit LDLR binding to PCSK9. In combination with the results from Example 40 and shown in FIGs. 28A and 28B, it is clear that 31A4 ABP binds to the V domain of PCSK9 and does not block the interaction of PCSK9 with LDLR.

[0604] Next, the Ability of ABP 31A4 to serve as a neutralizing ABP was further confirmed via a cell LDL uptake assay (as described in the examples above). The results of this LDL uptake assay are presented in FIG. 28D. As shown in FIG. 28D, ABP 31A4 displays significant PCSK9 neutralizing ability. Thus, in light of Example 40 and the present results, it is clear that ABPs can bind to PCSK9 without blocking the PCSK9 and LDLR binding interaction, while still being useful as neutralizing PCSK9 ABPs.

Incorporation by Reference

[0605] All references cited herein, including patents, patent applications, papers, text books, and the like, and the references cited therein, to the extent that they are not already, are hereby incorporated herein by reference in their entirety. To the extent that any of the definitions or terms provided in the references incorporated by reference differ from the terms and discussion provided herein, the present terms and definitions control.

Equivalents

[0606] The foregoing written specification is considered to be sufficient to enable one skilled in the art to practice the invention. The foregoing description and examples detail certain preferred embodiments of the invention and describe the best mode contemplated by the inventors. It will be appreciated, however, that no matter how detailed the foregoing may appear in text, the invention may be practiced in many ways and the invention should be construed in accordance with the appended claims and any equivalents thereof.

TABLE 35.1

ATOM	1	CB	THR	61	-65.324	19.274	-35.379	1.00	66.96	A	C
ATOM	2	OG1	THR	61	-64.490	20.386	-35.733	1.00	67.86	A	O
ATOM	3	CG2	THR	61	-65.574	19.285	-33.870	1.00	66.57	A	C
ATOM	4	C	THR	61	-63.283	17.835	-35.088	1.00	62.73	A	C
ATOM	5	O	THR	61	-63.080	16.945	-34.257	1.00	63.03	A	O
ATOM	6	N	THR	61	-65.516	16.775	-35.528	1.00	66.08	A	N
ATOM	7	CA	THR	61	-64.635	17.950	-35.808	1.00	65.41	A	C
ATOM	8	N	ALA	62	-62.364	18.740	-35.417	1.00	59.54	A	N
ATOM	9	CA	ALA	62	-61.013	18.712	-34.866	1.00	55.44	A	C
ATOM	10	CB	ALA	62	-60.100	19.581	-35.709	1.00	54.24	A	C
ATOM	11	C	ALA	62	-60.988	19.182	-33.414	1.00	53.41	A	C
ATOM	12	O	ALA	62	-61.570	20.211	-33.075	1.00	53.36	A	O
ATOM	13	N	THR	63	-60.309	18.421	-32.561	1.00	50.60	A	N
ATOM	14	CA	THR	63	-60.219	18.743	-31.141	1.00	48.08	A	C
ATOM	15	CB	THR	63	-60.538	17.503	-30.272	1.00	48.75	A	C
ATOM	16	OG1	THR	63	-59.717	16.402	-30.683	1.00	48.97	A	O
ATOM	17	CG2	THR	63	-61.997	17.104	-30.426	1.00	47.89	A	C
ATOM	18	C	THR	63	-58.831	19.261	-30.766	1.00	46.77	A	C
ATOM	19	O	THR	63	-57.853	19.004	-31.465	1.00	46.12	A	O
ATOM	20	N	PHE	64	-58.754	19.999	-29.662	1.00	45.98	A	N
ATOM	21	CA	PHE	64	-57.476	20.471	-29.136	1.00	44.11	A	C
ATOM	22	CB	PHE	64	-57.537	21.980	-28.894	1.00	42.47	A	C
ATOM	23	CG	PHE	64	-56.352	22.529	-28.150	1.00	41.54	A	C
ATOM	24	CD1	PHE	64	-55.113	22.627	-28.764	1.00	39.91	A	C
ATOM	25	CD2	PHE	64	-56.484	22.964	-26.839	1.00	40.63	A	C
ATOM	26	CE1	PHE	64	-54.024	23.149	-28.085	1.00	39.85	A	C
ATOM	27	CE2	PHE	64	-55.399	23.489	-26.151	1.00	40.22	A	C
ATOM	28	CZ	PHE	64	-54.166	23.582	-26.776	1.00	39.77	A	C
ATOM	29	C	PHE	64	-57.110	19.744	-27.841	1.00	44.16	A	C
ATOM	30	O	PHE	64	-57.966	19.506	-26.982	1.00	43.98	A	O
ATOM	31	N	HIS	65	-55.834	19.388	-27.711	1.00	43.51	A	N
ATOM	32	CA	HIS	65	-55.348	18.640	-26.554	1.00	42.21	A	C
ATOM	33	CB	HIS	65	-55.015	17.204	-26.964	1.00	42.39	A	C
ATOM	34	CG	HIS	65	-56.168	16.477	-27.581	1.00	45.11	A	C
ATOM	35	CD2	HIS	65	-56.648	16.477	-28.848	1.00	46.30	A	C
ATOM	36	ND1	HIS	65	-57.005	15.656	-26.855	1.00	45.17	A	N
ATOM	37	CE1	HIS	65	-57.951	15.183	-27.648	1.00	45.62	A	C
ATOM	38	NE2	HIS	65	-57.757	15.666	-28.863	1.00	45.65	A	N
ATOM	39	C	HIS	65	-54.115	19.297	-25.941	1.00	42.51	A	C
ATOM	40	O	HIS	65	-53.276	19.865	-26.643	1.00	41.98	A	O
ATOM	41	N	ARG	66	-54.013	19.220	-24.622	1.00	42.62	A	N
ATOM	42	CA	ARG	66	-52.863	19.765	-23.916	1.00	43.48	A	C
ATOM	43	CB	ARG	66	-53.152	21.202	-23.471	1.00	45.30	A	C
ATOM	44	CG	ARG	66	-54.358	21.305	-22.561	1.00	51.92	A	C
ATOM	45	CD	ARG	66	-54.333	22.550	-21.702	1.00	57.80	A	C
ATOM	46	NE	ARG	66	-55.193	22.390	-20.530	1.00	63.34	A	N
ATOM	47	CZ	ARG	66	-54.755	22.114	-19.302	1.00	66.07	A	C
ATOM	48	NH1	ARG	66	-55.620	21.983	-18.304	1.00	68.14	A	N
ATOM	49	NH2	ARG	66	-53.457	21.979	-19.063	1.00	66.65	A	N
ATOM	50	C	ARG	66	-52.570	18.890	-22.698	1.00	42.14	A	C
ATOM	51	O	ARG	66	-53.427	18.129	-22.246	1.00	41.73	A	O
ATOM	52	N	CYS	67	-51.358	18.999	-22.172	1.00	40.93	A	N
ATOM	53	CA	CYS	67	-50.965	18.224	-21.004	1.00	41.21	A	C
ATOM	54	CB	CYS	67	-49.500	18.505	-20.678	1.00	41.47	A	C
ATOM	55	SG	CYS	67	-48.844	17.576	-19.295	1.00	40.60	A	S
ATOM	56	C	CYS	67	-51.843	18.605	-19.813	1.00	43.29	A	C
ATOM	57	O	CYS	67	-52.072	19.789	-19.555	1.00	43.45	A	O
ATOM	58	N	ALA	68	-52.331	17.606	-19.088	1.00	43.43	A	N
ATOM	59	CA	ALA	68	-53.144	17.871	-17.907	1.00	45.62	A	C
ATOM	60	CB	ALA	68	-53.809	16.579	-17.416	1.00	43.28	A	C
ATOM	61	C	ALA	68	-52.315	18.501	-16.783	1.00	46.90	A	C
ATOM	62	O	ALA	68	-52.852	19.232	-15.949	1.00	46.15	A	O
ATOM	63	N	LYS	69	-51.010	18.227	-16.767	1.00	48.41	A	N
ATOM	64	CA	LYS	69	-50.132	18.747	-15.715	1.00	50.02	A	C
ATOM	65	CB	LYS	69	-48.974	17.773	-15.454	1.00	52.87	A	C
ATOM	66	CG	LYS	69	-49.388	16.305	-15.385	1.00	58.02	A	C
ATOM	67	CD	LYS	69	-49.184	15.722	-13.990	1.00	61.71	A	C
ATOM	68	CE	LYS	69	-50.035	14.472	-13.783	1.00	63.38	A	C
ATOM	69	NZ	LYS	69	-50.285	14.221	-12.334	1.00	64.13	A	N
ATOM	70	C	LYS	69	-49.575	20.110	-16.119	1.00	48.59	A	C
ATOM	71	O	LYS	69	-48.626	20.200	-16.891	1.00	49.36	A	O
ATOM	72	N	ASP	70	-50.160	21.169	-15.576	1.00	48.35	A	N
ATOM	73	CA	ASP	70	-49.948	22.512	-16.099	1.00	47.47	A	C
ATOM	74	CB	ASP	70	-50.744	23.526	-15.268	1.00	52.53	A	C

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ATOM     75  CG  ASP   70     -51.508  24.519 -16.137  1.00 57.41      A    C
ATOM     76  OD1 ASP   70     -52.644  24.189 -16.562  1.00 58.27      A    O
ATOM     77  OD2 ASP   70     -50.971  25.623 -16.399  1.00 57.80      A    O
ATOM     78  C   ASP   70     -48.488  22.966 -16.200  1.00 44.86      A    C
ATOM     79  O   ASP   70     -48.098  23.599 -17.180  1.00 45.04      A    O
ATOM     80  N   PRO   71     -47.665  22.667 -15.187  1.00 41.90      A    N
ATOM     81  CD  PRO   71     -48.023  22.174 -13.845  1.00 40.54      A    C
ATOM     82  CA  PRO   71     -46.258  23.083 -15.281  1.00 39.08      A    C
ATOM     83  CB  PRO   71     -45.723  22.898 -13.857  1.00 39.03      A    C
ATOM     84  CG  PRO   71     -46.688  21.975 -13.186  1.00 41.03      A    C
ATOM     85  C   PRO   71     -45.402  22.349 -16.324  1.00 37.94      A    C
ATOM     86  O   PRO   71     -44.280  22.757 -16.608  1.00 36.99      A    O
ATOM     87  N   TRP   72     -45.925  21.270 -16.893  1.00 36.09      A    N
ATOM     88  CA  TRP   72     -45.201  20.539 -17.933  1.00 34.40      A    C
ATOM     89  CB  TRP   72     -45.456  19.034 -17.806  1.00 32.15      A    C
ATOM     90  CG  TRP   72     -44.904  18.427 -16.551  1.00 29.66      A    C
ATOM     91  CD2 TRP   72     -45.042  17.066 -16.126  1.00 28.28      A    C
ATOM     92  CE2 TRP   72     -44.381  16.947 -14.884  1.00 29.32      A    C
ATOM     93  CE3 TRP   72     -45.659  15.938 -16.674  1.00 28.21      A    C
ATOM     94  CD1 TRP   72     -44.183  19.061 -15.580  1.00 28.83      A    C
ATOM     95  NE1 TRP   72     -43.866  18.178 -14.574  1.00 27.11      A    N
ATOM     96  CZ2 TRP   72     -44.322  15.742 -14.181  1.00 28.10      A    C
ATOM     97  CZ3 TRP   72     -45.600  14.745 -15.978  1.00 29.75      A    C
ATOM     98  CH2 TRP   72     -44.935  14.655 -14.741  1.00 30.43      A    C
ATOM     99  C   TRP   72     -45.622  21.004 -19.327  1.00 33.51      A    C
ATOM    100  O   TRP   72     -45.074  20.554 -20.330  1.00 34.06      A    O
ATOM    101  N   ARG   73     -46.599  21.903 -19.379  1.00 33.72      A    N
ATOM    102  CA  ARG   73     -47.088  22.437 -20.643  1.00 35.14      A    C
ATOM    103  CB  ARG   73     -48.370  23.237 -20.407  1.00 37.00      A    C
ATOM    104  CG  ARG   73     -49.543  22.389 -19.975  1.00 41.40      A    C
ATOM    105  CD  ARG   73     -50.786  23.229 -19.825  1.00 45.06      A    C
ATOM    106  NE  ARG   73     -50.898  24.206 -20.902  1.00 50.54      A    N
ATOM    107  CZ  ARG   73     -51.976  24.951 -21.127  1.00 54.08      A    C
ATOM    108  NH1 ARG   73     -53.045  24.829 -20.347  1.00 55.58      A    N
ATOM    109  NH2 ARG   73     -51.983  25.826 -22.126  1.00 54.14      A    N
ATOM    110  C   ARG   73     -46.042  23.327 -21.304  1.00 34.68      A    C
ATOM    111  O   ARG   73     -45.294  24.025 -20.619  1.00 35.28      A    O
ATOM    112  N   LEU   74     -45.986  23.294 -22.633  1.00 32.04      A    N
ATOM    113  CA  LEU   74     -45.086  24.166 -23.385  1.00 32.47      A    C
ATOM    114  CB  LEU   74     -43.966  23.341 -24.031  1.00 31.53      A    C
ATOM    115  CG  LEU   74     -42.990  22.623 -23.087  1.00 32.34      A    C
ATOM    116  CD1 LEU   74     -42.183  21.578 -23.863  1.00 27.72      A    C
ATOM    117  CD2 LEU   74     -42.061  23.647 -22.437  1.00 28.10      A    C
ATOM    118  C   LEU   74     -45.846  24.934 -24.468  1.00 33.33      A    C
ATOM    119  O   LEU   74     -45.677  24.677 -25.662  1.00 33.93      A    O
ATOM    120  N   PRO   75     -46.687  25.897 -24.064  1.00 34.33      A    N
ATOM    121  CD  PRO   75     -46.820  26.440 -22.698  1.00 33.65      A    C
ATOM    122  CA  PRO   75     -47.519  26.621 -25.039  1.00 34.35      A    C
ATOM    123  CB  PRO   75     -48.351  27.573 -24.174  1.00 35.25      A    C
ATOM    124  CG  PRO   75     -47.545  27.746 -22.915  1.00 35.46      A    C
ATOM    125  C   PRO   75     -46.688  27.369 -26.083  1.00 33.52      A    C
ATOM    126  O   PRO   75     -45.553  27.768 -25.817  1.00 32.98      A    O
ATOM    127  N   GLY   76     -47.249  27.547 -27.275  1.00 32.91      A    N
ATOM    128  CA  GLY   76     -46.513  28.227 -28.328  1.00 32.08      A    C
ATOM    129  C   GLY   76     -45.937  27.286 -29.371  1.00 31.14      A    C
ATOM    130  O   GLY   76     -45.480  27.726 -30.424  1.00 31.40      A    O
ATOM    131  N   THR   77     -45.947  25.989 -29.080  1.00 30.23      A    N
ATOM    132  CA  THR   77     -45.584  24.986 -30.073  1.00 30.51      A    C
ATOM    133  CB  THR   77     -44.197  24.368 -29.776  1.00 32.27      A    C
ATOM    134  OG1 THR   77     -43.199  25.398 -29.816  1.00 33.45      A    O
ATOM    135  CG2 THR   77     -43.840  23.311 -30.819  1.00 32.84      A    C
ATOM    136  C   THR   77     -46.647  23.894 -30.107  1.00 30.00      A    C
ATOM    137  O   THR   77     -47.129  23.441 -29.064  1.00 30.45      A    O
ATOM    138  N   TYR   78     -47.024  23.492 -31.317  1.00 28.44      A    N
ATOM    139  CA  TYR   78     -48.156  22.597 -31.514  1.00 27.71      A    C
ATOM    140  CB  TYR   78     -49.396  23.395 -31.939  1.00 27.62      A    C
ATOM    141  CG  TYR   78     -49.730  24.496 -30.967  1.00 29.17      A    C
ATOM    142  CD1 TYR   78     -49.199  25.768 -31.128  1.00 29.25      A    C
ATOM    143  CE1 TYR   78     -49.408  26.757 -30.183  1.00 31.96      A    C
ATOM    144  CD2 TYR   78     -50.494  24.241 -29.837  1.00 28.93      A    C
ATOM    145  CE2 TYR   78     -50.709  25.225 -28.880  1.00 32.16      A    C
ATOM    146  CZ  TYR   78     -50.157  26.479 -29.057  1.00 32.47      A    C
ATOM    147  OH  TYR   78     -50.305  27.440 -28.082  1.00 32.98      A    O
ATOM    148  C   TYR   78     -47.840  21.552 -32.570  1.00 27.76      A    C
ATOM    149  O   TYR   78     -47.154  21.834 -33.559  1.00 25.34      A    O
ATOM    150  N   VAL   79     -48.343  20.343 -32.341  1.00 26.43      A    N
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ATOM    151  CA  VAL    79     -48.337  19.311 -33.357  1.00 27.54      A    C
ATOM    152  CB  VAL    79     -48.010  17.921 -32.760  1.00 28.88      A    C
ATOM    153  CG1 VAL    79     -48.012  16.871 -33.868  1.00 27.66      A    C
ATOM    154  CG2 VAL    79     -46.666  17.959 -32.059  1.00 28.19      A    C
ATOM    155  C   VAL    79     -49.726  19.259 -33.961  1.00 28.85      A    C
ATOM    156  O   VAL    79     -50.712  18.983 -33.264  1.00 28.23      A    O
ATOM    157  N   VAL    80     -49.807  19.534 -35.255  1.00 28.01      A    N
ATOM    158  CA  VAL    80     -51.073  19.430 -35.960  1.00 28.18      A    C
ATOM    159  CB  VAL    80     -51.194  20.529 -37.028  1.00 28.78      A    C
ATOM    160  CG1 VAL    80     -52.524  20.404 -37.761  1.00 28.21      A    C
ATOM    161  CG2 VAL    80     -51.061  21.904 -36.362  1.00 26.47      A    C
ATOM    162  C   VAL    80     -51.126  18.063 -36.617  1.00 30.35      A    C
ATOM    163  O   VAL    80     -50.352  17.769 -37.533  1.00 30.14      A    O
ATOM    164  N   VAL    81     -52.024  17.215 -36.128  1.00 30.97      A    N
ATOM    165  CA  VAL    81     -52.139  15.869 -36.657  1.00 32.26      A    C
ATOM    166  CB  VAL    81     -52.423  14.850 -35.534  1.00 33.61      A    C
ATOM    167  CG1 VAL    81     -52.529  13.441 -36.121  1.00 31.68      A    C
ATOM    168  CG2 VAL    81     -51.316  14.915 -34.487  1.00 31.98      A    C
ATOM    169  C   VAL    81     -53.271  15.838 -37.674  1.00 34.32      A    C
ATOM    170  O   VAL    81     -54.400  16.235 -37.373  1.00 33.98      A    O
ATOM    171  N   LEU    82     -52.955  15.387 -38.883  1.00 34.05      A    N
ATOM    172  CA  LEU    82     -53.938  15.329 -39.961  1.00 37.51      A    C
ATOM    173  CB  LEU    82     -53.260  15.638 -41.300  1.00 34.52      A    C
ATOM    174  CG  LEU    82     -52.581  17.011 -41.323  1.00 34.04      A    C
ATOM    175  CD1 LEU    82     -52.087  17.337 -42.720  1.00 33.14      A    C
ATOM    176  CD2 LEU    82     -53.566  18.057 -40.857  1.00 33.19      A    C
ATOM    177  C   LEU    82     -54.595  13.950 -40.009  1.00 39.40      A    C
ATOM    178  O   LEU    82     -54.057  12.980 -39.475  1.00 38.11      A    O
ATOM    179  N   LYS    83     -55.764  13.864 -40.634  1.00 44.40      A    N
ATOM    180  CA  LYS    83     -56.464  12.586 -40.745  1.00 49.45      A    C
ATOM    181  CB  LYS    83     -57.752  12.756 -41.550  1.00 50.92      A    C
ATOM    182  CG  LYS    83     -58.854  13.495 -40.800  1.00 55.63      A    C
ATOM    183  CD  LYS    83     -59.854  14.109 -41.767  1.00 59.44      A    C
ATOM    184  CE  LYS    83     -60.934  14.899 -41.036  1.00 62.03      A    C
ATOM    185  NZ  LYS    83     -61.717  15.751 -41.986  1.00 64.41      A    N
ATOM    186  C   LYS    83     -55.560  11.562 -41.418  1.00 51.74      A    C
ATOM    187  O   LYS    83     -54.770  11.901 -42.301  1.00 50.75      A    O
ATOM    188  N   GLU    84     -55.663  10.309 -40.997  1.00 55.37      A    N
ATOM    189  CA  GLU    84     -54.787   9.289 -41.549  1.00 60.52      A    C
ATOM    190  CB  GLU    84     -54.910   7.985 -40.756  1.00 63.78      A    C
ATOM    191  CG  GLU    84     -56.292   7.371 -40.753  1.00 69.97      A    C
ATOM    192  CD  GLU    84     -56.295   5.986 -40.129  1.00 74.43      A    C
ATOM    193  OE1 GLU    84     -57.294   5.631 -39.462  1.00 75.98      A    O
ATOM    194  OE2 GLU    84     -55.293   5.254 -40.306  1.00 75.33      A    O
ATOM    195  C   GLU    84     -55.110   9.052 -43.022  1.00 61.14      A    C
ATOM    196  O   GLU    84     -56.248   9.246 -43.458  1.00 60.67      A    O
ATOM    197  N   GLU    85     -54.089   8.649 -43.774  1.00 61.46      A    N
ATOM    198  CA  GLU    85     -54.163   8.519 -45.227  1.00 63.36      A    C
ATOM    199  CB  GLU    85     -55.532   7.980 -45.662  1.00 67.22      A    C
ATOM    200  CG  GLU    85     -55.846   6.597 -45.096  1.00 73.81      A    C
ATOM    201  CD  GLU    85     -56.946   5.875 -45.858  1.00 78.53      A    C
ATOM    202  OE1 GLU    85     -57.733   6.545 -46.567  1.00 80.96      A    O
ATOM    203  OE2 GLU    85     -57.022   4.630 -45.745  1.00 80.38      A    O
ATOM    204  C   GLU    85     -53.865   9.835 -45.944  1.00 61.28      A    C
ATOM    205  O   GLU    85     -53.744   9.869 -47.172  1.00 61.77      A    O
ATOM    206  N   THR    86     -53.735  10.917 -45.181  1.00 57.81      A    N
ATOM    207  CA  THR    86     -53.256  12.173 -45.746  1.00 54.38      A    C
ATOM    208  CB  THR    86     -53.297  13.312 -44.701  1.00 52.95      A    C
ATOM    209  OG1 THR    86     -54.654  13.549 -44.307  1.00 50.34      A    O
ATOM    210  CG2 THR    86     -52.720  14.593 -45.281  1.00 50.02      A    C
ATOM    211  C   THR    86     -51.821  11.989 -46.243  1.00 53.59      A    C
ATOM    212  O   THR    86     -50.973  11.435 -45.539  1.00 52.24      A    O
ATOM    213  N   HIS    87     -51.562  12.442 -47.466  1.00 52.07      A    N
ATOM    214  CA  HIS    87     -50.250  12.287 -48.084  1.00 51.71      A    C
ATOM    215  CB  HIS    87     -50.401  12.183 -49.605  1.00 55.85      A    C
ATOM    216  CG  HIS    87     -51.185  10.986 -50.052  1.00 63.40      A    C
ATOM    217  CD2 HIS    87     -52.429  10.880 -50.579  1.00 64.75      A    C
ATOM    218  ND1 HIS    87     -50.690   9.700 -49.975  1.00 65.63      A    N
ATOM    219  CE1 HIS    87     -51.597   8.854 -50.434  1.00 66.35      A    C
ATOM    220  NE2 HIS    87     -52.660   9.544 -50.807  1.00 65.74      A    N
ATOM    221  C   HIS    87     -49.312  13.445 -47.732  1.00 48.81      A    C
ATOM    222  O   HIS    87     -49.760  14.519 -47.319  1.00 47.37      A    O
ATOM    223  N   LEU    88     -48.011  13.213 -47.896  1.00 46.04      A    N
ATOM    224  CA  LEU    88     -46.992  14.193 -47.536  1.00 44.77      A    C
ATOM    225  CB  LEU    88     -45.601  13.697 -47.944  1.00 43.47      A    C
ATOM    226  CG  LEU    88     -44.448  14.702 -47.814  1.00 44.24      A    C
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ATOM	227	CD1	LEU	88	-44.344	15.214	-46.379	1.00	40.89	A	C
ATOM	228	CD2	LEU	88	-43.149	14.037	-48.245	1.00	42.45	A	C
ATOM	229	C	LEU	88	-47.250	15.548	-48.176	1.00	43.96	A	C
ATOM	230	O	LEU	88	-47.167	16.583	-47.508	1.00	42.78	A	O
ATOM	231	N	SER	89	-47.568	15.543	-49.467	1.00	43.13	A	N
ATOM	232	CA	SER	89	-47.769	16.797	-50.187	1.00	43.67	A	C
ATOM	233	CB	SER	89	-47.949	16.537	-51.691	1.00	43.34	A	C
ATOM	234	OG	SER	89	-48.935	15.547	-51.926	1.00	47.73	A	O
ATOM	235	C	SER	89	-48.968	17.560	-49.634	1.00	41.76	A	C
ATOM	236	O	SER	89	-48.992	18.786	-49.663	1.00	43.20	A	O
ATOM	237	N	GLN	90	-49.984	16.836	-49.116	1.00	40.72	A	N
ATOM	238	CA	GLN	90	-51.101	17.471	-48.471	1.00	41.68	A	C
ATOM	239	CB	GLN	90	-52.241	16.452	-48.289	1.00	45.43	A	C
ATOM	240	CG	GLN	90	-52.820	15.904	-49.606	1.00	51.25	A	C
ATOM	241	CD	GLN	90	-53.850	14.789	-49.394	1.00	54.98	A	C
ATOM	242	OE1	GLN	90	-53.500	13.650	-49.059	1.00	54.79	A	O
ATOM	243	NE2	GLN	90	-55.127	15.117	-49.589	1.00	54.36	A	N
ATOM	244	C	GLN	90	-50.737	18.100	-47.114	1.00	39.35	A	C
ATOM	245	O	GLN	90	-51.158	19.219	-46.820	1.00	38.00	A	O
ATOM	246	N	SER	91	-49.960	17.388	-46.297	1.00	36.66	A	N
ATOM	247	CA	SER	91	-49.456	17.947	-45.037	1.00	36.04	A	C
ATOM	248	CB	SER	91	-48.607	16.919	-44.289	1.00	36.68	A	C
ATOM	249	OG	SER	91	-49.340	15.738	-44.034	1.00	45.07	A	O
ATOM	250	C	SER	91	-48.596	19.173	-45.311	1.00	34.52	A	C
ATOM	251	O	SER	91	-48.730	20.197	-44.648	1.00	31.21	A	O
ATOM	252	N	GLU	92	-47.711	19.056	-46.296	1.00	34.78	A	N
ATOM	253	CA	GLU	92	-46.823	20.147	-46.664	1.00	37.08	A	C
ATOM	254	CB	GLU	92	-45.942	19.724	-47.847	1.00	37.84	A	C
ATOM	255	CG	GLU	92	-44.600	20.442	-47.938	1.00	42.08	A	C
ATOM	256	CD	GLU	92	-44.680	21.789	-48.662	1.00	43.83	A	C
ATOM	257	OE1	GLU	92	-43.855	22.689	-48.358	1.00	38.38	A	O
ATOM	258	OE2	GLU	92	-45.566	21.940	-49.539	1.00	45.37	A	O
ATOM	259	C	GLU	92	-47.649	21.386	-47.028	1.00	37.77	A	C
ATOM	260	O	GLU	92	-47.367	22.486	-46.549	1.00	36.08	A	O
ATOM	261	N	ARG	93	-48.680	21.200	-47.855	1.00	37.72	A	N
ATOM	262	CA	ARG	93	-49.527	22.313	-48.300	1.00	37.76	A	C
ATOM	263	CB	ARG	93	-50.435	21.876	-49.457	1.00	39.31	A	C
ATOM	264	CG	ARG	93	-49.726	21.788	-50.808	1.00	46.60	A	C
ATOM	265	CD	ARG	93	-50.717	21.579	-51.963	1.00	52.39	A	C
ATOM	266	NE	ARG	93	-50.750	20.189	-52.419	1.00	56.69	A	N
ATOM	267	CZ	ARG	93	-51.698	19.309	-52.098	1.00	58.62	A	C
ATOM	268	NH1	ARG	93	-51.630	18.066	-52.563	1.00	58.46	A	N
ATOM	269	NH2	ARG	93	-52.715	19.667	-51.319	1.00	59.41	A	N
ATOM	270	C	ARG	93	-50.387	22.891	-47.181	1.00	35.91	A	C
ATOM	271	O	ARG	93	-50.610	24.097	-47.121	1.00	36.85	A	O
ATOM	272	N	THR	94	-50.869	22.034	-46.292	1.00	34.43	A	N
ATOM	273	CA	THR	94	-51.674	22.500	-45.173	1.00	34.17	A	C
ATOM	274	CB	THR	94	-52.275	21.305	-44.401	1.00	35.39	A	C
ATOM	275	OG1	THR	94	-53.069	20.519	-45.297	1.00	35.45	A	O
ATOM	276	CG2	THR	94	-53.159	21.789	-43.246	1.00	33.36	A	C
ATOM	277	C	THR	94	-50.843	23.370	-44.218	1.00	34.67	A	C
ATOM	278	O	THR	94	-51.341	24.355	-43.673	1.00	34.08	A	O
ATOM	279	N	ALA	95	-49.574	23.019	-44.031	1.00	33.89	A	N
ATOM	280	CA	ALA	95	-48.687	23.832	-43.203	1.00	35.56	A	C
ATOM	281	CB	ALA	95	-47.331	23.137	-43.037	1.00	32.25	A	C
ATOM	282	C	ALA	95	-48.496	25.225	-43.816	1.00	36.52	A	C
ATOM	283	O	ALA	95	-48.536	26.234	-43.107	1.00	35.42	A	O
ATOM	284	N	ARG	96	-48.297	25.280	-45.132	1.00	36.12	A	N
ATOM	285	CA	ARG	96	-48.094	26.559	-45.807	1.00	38.95	A	C
ATOM	286	CB	ARG	96	-47.563	26.342	-47.234	1.00	37.87	A	C
ATOM	287	CG	ARG	96	-46.128	25.808	-47.253	1.00	40.62	A	C
ATOM	288	CD	ARG	96	-45.424	26.077	-48.575	1.00	41.07	A	C
ATOM	289	NE	ARG	96	-44.078	25.501	-48.616	1.00	41.43	A	N
ATOM	290	CZ	ARG	96	-42.998	26.079	-48.093	1.00	41.81	A	C
ATOM	291	NH1	ARG	96	-41.814	25.485	-48.180	1.00	41.20	A	N
ATOM	292	NH2	ARG	96	-43.096	27.252	-47.481	1.00	39.73	A	N
ATOM	293	C	ARG	96	-49.383	27.372	-45.841	1.00	39.48	A	C
ATOM	294	O	ARG	96	-49.357	28.604	-45.812	1.00	39.37	A	O
ATOM	295	N	ARG	97	-50.512	26.676	-45.885	1.00	39.22	A	N
ATOM	296	CA	ARG	97	-51.805	27.334	-45.815	1.00	41.01	A	C
ATOM	297	CB	ARG	97	-52.915	26.305	-46.028	1.00	45.31	A	C
ATOM	298	CG	ARG	97	-54.299	26.890	-46.199	1.00	52.12	A	C
ATOM	299	CD	ARG	97	-55.098	26.064	-47.200	1.00	60.48	A	C
ATOM	300	NE	ARG	97	-56.533	26.107	-46.930	1.00	66.20	A	N
ATOM	301	CZ	ARG	97	-57.321	27.138	-47.226	1.00	68.21	A	C
ATOM	302	NH1	ARG	97	-58.617	27.084	-46.940	1.00	68.65	A	N

ATOM	303	NH2	ARG	97	-56.813	28.223	-47.803	1.00	70.24	A	N
ATOM	304	C	ARG	97	-51.979	28.026	-44.460	1.00	39.37	A	C
ATOM	305	O	ARG	97	-52.377	29.190	-44.393	1.00	38.18	A	O
ATOM	306	N	LEU	98	-51.674	27.311	-43.381	1.00	37.41	A	N
ATOM	307	CA	LEU	98	-51.764	27.897	-42.050	1.00	35.51	A	C
ATOM	308	CB	LEU	98	-51.356	26.884	-40.977	1.00	34.36	A	C
ATOM	309	CG	LEU	98	-51.216	27.460	-39.563	1.00	36.07	A	C
ATOM	310	CD1	LEU	98	-52.558	28.003	-39.099	1.00	35.97	A	C
ATOM	311	CD2	LEU	98	-50.727	26.386	-38.604	1.00	36.03	A	C
ATOM	312	C	LEU	98	-50.862	29.122	-41.961	1.00	35.38	A	C
ATOM	313	O	LEU	98	-51.275	30.158	-41.446	1.00	33.36	A	O
ATOM	314	N	GLN	99	-49.635	29.007	-42.469	1.00	34.49	A	N
ATOM	315	CA	GLN	99	-48.693	30.123	-42.409	1.00	37.42	A	C
ATOM	316	CB	GLN	99	-47.359	29.745	-43.061	1.00	38.96	A	C
ATOM	317	CG	GLN	99	-46.377	29.034	-42.146	1.00	40.19	A	C
ATOM	318	CD	GLN	99	-44.931	29.254	-42.564	1.00	41.88	A	C
ATOM	319	OE1	GLN	99	-44.540	28.924	-43.683	1.00	41.48	A	O
ATOM	320	NE2	GLN	99	-44.129	29.816	-41.661	1.00	40.56	A	N
ATOM	321	C	GLN	99	-49.248	31.371	-43.099	1.00	38.67	A	C
ATOM	322	O	GLN	99	-49.136	32.484	-42.577	1.00	38.83	A	O
ATOM	323	N	ALA	100	-49.844	31.179	-44.273	1.00	39.11	A	N
ATOM	324	CA	ALA	100	-50.344	32.293	-45.074	1.00	40.09	A	C
ATOM	325	CB	ALA	100	-50.704	31.814	-46.483	1.00	38.16	A	C
ATOM	326	C	ALA	100	-51.558	32.935	-44.416	1.00	40.78	A	C
ATOM	327	O	ALA	100	-51.685	34.160	-44.396	1.00	41.97	A	O
ATOM	328	N	GLN	101	-52.447	32.115	-43.870	1.00	39.41	A	N
ATOM	329	CA	GLN	101	-53.600	32.644	-43.160	1.00	39.71	A	C
ATOM	330	CB	GLN	101	-54.533	31.518	-42.724	1.00	38.98	A	C
ATOM	331	CG	GLN	101	-55.204	30.803	-43.877	1.00	41.93	A	C
ATOM	332	CD	GLN	101	-56.129	29.706	-43.405	1.00	44.37	A	C
ATOM	333	OE1	GLN	101	-56.874	29.125	-44.192	1.00	45.91	A	O
ATOM	334	NE2	GLN	101	-56.089	29.417	-42.109	1.00	44.36	A	N
ATOM	335	C	GLN	101	-53.174	33.447	-41.940	1.00	40.25	A	C
ATOM	336	O	GLN	101	-53.731	34.512	-41.671	1.00	42.25	A	O
ATOM	337	N	ALA	102	-52.190	32.942	-41.203	1.00	38.45	A	N
ATOM	338	CA	ALA	102	-51.734	33.620	-39.993	1.00	37.78	A	C
ATOM	339	CB	ALA	102	-50.782	32.710	-39.198	1.00	35.08	A	C
ATOM	340	C	ALA	102	-51.037	34.934	-40.346	1.00	37.94	A	C
ATOM	341	O	ALA	102	-51.148	35.924	-39.615	1.00	37.67	A	O
ATOM	342	N	ALA	103	-50.321	34.940	-41.467	1.00	37.12	A	N
ATOM	343	CA	ALA	103	-49.589	36.127	-41.901	1.00	39.38	A	C
ATOM	344	CB	ALA	103	-48.705	35.787	-43.098	1.00	38.54	A	C
ATOM	345	C	ALA	103	-50.549	37.259	-42.268	1.00	41.97	A	C
ATOM	346	O	ALA	103	-50.268	38.437	-42.012	1.00	41.53	A	O
ATOM	347	N	ARG	104	-51.682	36.897	-42.864	1.00	42.63	A	N
ATOM	348	CA	ARG	104	-52.688	37.879	-43.245	1.00	43.83	A	C
ATOM	349	CB	ARG	104	-53.755	37.233	-44.139	1.00	43.90	A	C
ATOM	350	CG	ARG	104	-53.228	36.835	-45.513	1.00	45.54	A	C
ATOM	351	CD	ARG	104	-54.345	36.651	-46.528	1.00	48.21	A	C
ATOM	352	NE	ARG	104	-55.256	35.566	-46.168	1.00	53.08	A	N
ATOM	353	CZ	ARG	104	-55.105	34.300	-46.558	1.00	54.45	A	C
ATOM	354	NH1	ARG	104	-55.984	33.379	-46.184	1.00	53.80	A	N
ATOM	355	NH2	ARG	104	-54.073	33.952	-47.320	1.00	52.89	A	N
ATOM	356	C	ARG	104	-53.340	38.507	-42.018	1.00	44.46	A	C
ATOM	357	O	ARG	104	-53.924	39.589	-42.105	1.00	46.11	A	O
ATOM	358	N	ARG	105	-53.232	37.834	-40.874	1.00	43.03	A	N
ATOM	359	CA	ARG	105	-53.775	38.365	-39.630	1.00	40.77	A	C
ATOM	360	CB	ARG	105	-54.477	37.259	-38.845	1.00	43.29	A	C
ATOM	361	CG	ARG	105	-55.681	36.688	-39.562	1.00	45.71	A	C
ATOM	362	CD	ARG	105	-56.288	35.538	-38.794	1.00	50.02	A	C
ATOM	363	NE	ARG	105	-57.613	35.199	-39.305	1.00	54.91	A	N
ATOM	364	CZ	ARG	105	-57.854	34.793	-40.549	1.00	59.39	A	C
ATOM	365	NH1	ARG	105	-59.096	34.507	-40.919	1.00	61.37	A	N
ATOM	366	NH2	ARG	105	-56.859	34.672	-41.427	1.00	58.64	A	N
ATOM	367	C	ARG	105	-52.704	39.007	-38.763	1.00	40.45	A	C
ATOM	368	O	ARG	105	-52.958	39.360	-37.610	1.00	41.05	A	O
ATOM	369	N	GLY	106	-51.506	39.156	-39.320	1.00	40.26	A	N
ATOM	370	CA	GLY	106	-50.439	39.843	-38.615	1.00	42.36	A	C
ATOM	371	C	GLY	106	-49.558	38.965	-37.734	1.00	43.39	A	C
ATOM	372	O	GLY	106	-48.806	39.476	-36.896	1.00	41.65	A	O
ATOM	373	N	TYR	107	-49.642	37.648	-37.915	1.00	43.64	A	N
ATOM	374	CA	TYR	107	-48.869	36.719	-37.088	1.00	43.17	A	C
ATOM	375	CB	TYR	107	-49.783	35.631	-36.521	1.00	42.03	A	C
ATOM	376	CG	TYR	107	-50.728	36.113	-35.444	1.00	42.50	A	C
ATOM	377	CD1	TYR	107	-50.404	35.976	-34.097	1.00	41.77	A	C
ATOM	378	CE1	TYR	107	-51.274	36.385	-33.105	1.00	41.27	A	C

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ATOM    379  CD2 TYR   107     -51.955  36.679 -35.770  1.00 41.20      A    C
ATOM    380  CE2 TYR   107     -52.835  37.092 -34.782  1.00 40.76      A    C
ATOM    381  CZ  TYR   107     -52.490  36.941 -33.452  1.00 42.29      A    C
ATOM    382  OH  TYR   107     -53.364  37.337 -32.463  1.00 42.42      A    O
ATOM    383  C   TYR   107     -47.719  36.059 -37.842  1.00 42.45      A    C
ATOM    384  O   TYR   107     -47.907  35.508 -38.926  1.00 42.84      A    O
ATOM    385  N   LEU   108     -46.528  36.115 -37.259  1.00 42.23      A    N
ATOM    386  CA  LEU   108     -45.416  35.300 -37.728  1.00 43.15      A    C
ATOM    387  CB  LEU   108     -44.090  35.880 -37.237  1.00 46.35      A    C
ATOM    388  CG  LEU   108     -42.840  35.075 -37.603  1.00 50.90      A    C
ATOM    389  CD1 LEU   108     -42.631  35.119 -39.111  1.00 51.51      A    C
ATOM    390  CD2 LEU   108     -41.625  35.650 -36.885  1.00 52.89      A    C
ATOM    391  C   LEU   108     -45.581  33.878 -37.189  1.00 41.70      A    C
ATOM    392  O   LEU   108     -45.960  33.690 -36.031  1.00 42.13      A    O
ATOM    393  N   THR   109     -45.314  32.881 -38.030  1.00 37.66      A    N
ATOM    394  CA  THR   109     -45.293  31.495 -37.574  1.00 35.85      A    C
ATOM    395  CB  THR   109     -46.585  30.736 -37.972  1.00 34.84      A    C
ATOM    396  OG1 THR   109     -46.667  30.637 -39.400  1.00 35.33      A    O
ATOM    397  CG2 THR   109     -47.812  31.465 -37.451  1.00 34.18      A    C
ATOM    398  C   THR   109     -44.095  30.767 -38.174  1.00 35.46      A    C
ATOM    399  O   THR   109     -43.532  31.204 -39.178  1.00 34.99      A    O
ATOM    400  N   LYS   110     -43.699  29.662 -37.551  1.00 34.81      A    N
ATOM    401  CA  LYS   110     -42.624  28.837 -38.086  1.00 34.79      A    C
ATOM    402  CB  LYS   110     -41.374  28.961 -37.217  1.00 38.01      A    C
ATOM    403  CG  LYS   110     -40.739  30.336 -37.252  1.00 44.73      A    C
ATOM    404  CD  LYS   110     -39.687  30.494 -36.164  1.00 51.66      A    C
ATOM    405  CE  LYS   110     -39.100  31.906 -36.163  1.00 57.04      A    C
ATOM    406  NZ  LYS   110     -38.261  32.179 -34.953  1.00 60.67      A    N
ATOM    407  C   LYS   110     -43.044  27.379 -38.169  1.00 33.48      A    C
ATOM    408  O   LYS   110     -43.505  26.798 -37.182  1.00 30.94      A    O
ATOM    409  N   ILE   111     -42.891  26.794 -39.355  1.00 31.38      A    N
ATOM    410  CA  ILE   111     -43.033  25.352 -39.502  1.00 29.98      A    C
ATOM    411  CB  ILE   111     -43.366  24.953 -40.965  1.00 30.23      A    C
ATOM    412  CG2 ILE   111     -43.548  23.447 -41.065  1.00 27.09      A    C
ATOM    413  CG1 ILE   111     -44.646  25.652 -41.423  1.00 30.09      A    C
ATOM    414  CD1 ILE   111     -45.821  25.410 -40.511  1.00 30.43      A    C
ATOM    415  C   ILE   111     -41.689  24.745 -39.118  1.00 28.90      A    C
ATOM    416  O   ILE   111     -40.673  25.036 -39.745  1.00 28.53      A    O
ATOM    417  N   LEU   112     -41.680  23.909 -38.087  1.00 28.45      A    N
ATOM    418  CA  LEU   112     -40.429  23.359 -37.587  1.00 27.85      A    C
ATOM    419  CB  LEU   112     -40.487  23.232 -36.061  1.00 28.31      A    C
ATOM    420  CG  LEU   112     -40.736  24.527 -35.266  1.00 29.90      A    C
ATOM    421  CD1 LEU   112     -40.836  24.200 -33.780  1.00 27.40      A    C
ATOM    422  CD2 LEU   112     -39.608  25.527 -35.517  1.00 26.41      A    C
ATOM    423  C   LEU   112     -40.136  22.001 -38.209  1.00 28.75      A    C
ATOM    424  O   LEU   112     -38.976  21.588 -38.297  1.00 28.71      A    O
ATOM    425  N   HIS   113     -41.192  21.314 -38.645  1.00 28.46      A    N
ATOM    426  CA  HIS   113     -41.075  19.943 -39.129  1.00 29.20      A    C
ATOM    427  CB  HIS   113     -40.758  19.009 -37.954  1.00 29.04      A    C
ATOM    428  CG  HIS   113     -40.325  17.638 -38.367  1.00 29.50      A    C
ATOM    429  CD2 HIS   113     -41.039  16.519 -38.639  1.00 30.05      A    C
ATOM    430  ND1 HIS   113     -38.999  17.296 -38.533  1.00 31.19      A    N
ATOM    431  CE1 HIS   113     -38.915  16.026 -38.890  1.00 29.33      A    C
ATOM    432  NE2 HIS   113     -40.139  15.532 -38.962  1.00 30.67      A    N
ATOM    433  C   HIS   113     -42.377  19.500 -39.796  1.00 29.37      A    C
ATOM    434  O   HIS   113     -43.461  19.850 -39.336  1.00 30.57      A    O
ATOM    435  N   VAL   114     -42.277  18.730 -40.874  1.00 28.34      A    N
ATOM    436  CA  VAL   114     -43.460  18.116 -41.461  1.00 28.24      A    C
ATOM    437  CB  VAL   114     -43.648  18.545 -42.950  1.00 29.58      A    C
ATOM    438  CG1 VAL   114     -44.890  17.869 -43.556  1.00 24.77      A    C
ATOM    439  CG2 VAL   114     -43.802  20.055 -43.027  1.00 25.34      A    C
ATOM    440  C   VAL   114     -43.320  16.606 -41.363  1.00 30.01      A    C
ATOM    441  O   VAL   114     -42.368  16.031 -41.883  1.00 30.39      A    O
ATOM    442  N   PHE   115     -44.267  15.971 -40.679  1.00 31.76      A    N
ATOM    443  CA  PHE   115     -44.191  14.543 -40.424  1.00 35.74      A    C
ATOM    444  CB  PHE   115     -45.024  14.170 -39.198  1.00 33.99      A    C
ATOM    445  CG  PHE   115     -44.490  14.724 -37.909  1.00 34.95      A    C
ATOM    446  CD1 PHE   115     -44.984  15.915 -37.391  1.00 33.81      A    C
ATOM    447  CD2 PHE   115     -43.491  14.054 -37.212  1.00 33.75      A    C
ATOM    448  CE1 PHE   115     -44.490  16.432 -36.195  1.00 33.67      A    C
ATOM    449  CE2 PHE   115     -42.992  14.562 -36.019  1.00 32.89      A    C
ATOM    450  CZ  PHE   115     -43.494  15.757 -35.509  1.00 33.21      A    C
ATOM    451  C   PHE   115     -44.679  13.738 -41.615  1.00 40.69      A    C
ATOM    452  O   PHE   115     -45.672  14.089 -42.257  1.00 42.29      A    O
ATOM    453  N   HIS   116     -43.967  12.656 -41.902  1.00 45.48      A    N
ATOM    454  CA  HIS   116     -44.434  11.637 -42.830  1.00 51.72      A    C
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ATOM    455  CB  HIS   116     -44.084  12.017 -44.279  1.00 55.93      A    C
ATOM    456  CG  HIS   116     -42.638  12.353 -44.495  1.00 62.52      A    C
ATOM    457  CD2 HIS   116     -41.911  13.443 -44.143  1.00 65.77      A    C
ATOM    458  ND1 HIS   116     -41.775  11.528 -45.187  1.00 65.95      A    N
ATOM    459  CE1 HIS   116     -40.582  12.096 -45.255  1.00 67.64      A    C
ATOM    460  NE2 HIS   116     -40.638  13.260 -44.630  1.00 66.85      A    N
ATOM    461  C   HIS   116     -43.776  10.314 -42.461  1.00 53.18      A    C
ATOM    462  O   HIS   116     -42.587  10.270 -42.142  1.00 53.78      A    O
ATOM    463  N   GLY   117     -44.552   9.237 -42.497  1.00 54.01      A    N
ATOM    464  CA  GLY   117     -43.996   7.932 -42.198  1.00 55.08      A    C
ATOM    465  C   GLY   117     -44.538   7.307 -40.927  1.00 55.76      A    C
ATOM    466  O   GLY   117     -44.749   6.093 -40.868  1.00 58.44      A    O
ATOM    467  N   LEU   118     -44.763   8.122 -39.903  1.00 54.93      A    N
ATOM    468  CA  LEU   118     -45.369   7.626 -38.676  1.00 54.61      A    C
ATOM    469  CB  LEU   118     -44.501   7.993 -37.470  1.00 54.38      A    C
ATOM    470  CG  LEU   118     -44.413   6.966 -36.338  1.00 53.70      A    C
ATOM    471  CD1 LEU   118     -43.981   5.612 -36.888  1.00 53.81      A    C
ATOM    472  CD2 LEU   118     -43.423   7.449 -35.300  1.00 54.71      A    C
ATOM    473  C   LEU   118     -46.755   8.238 -38.526  1.00 54.27      A    C
ATOM    474  O   LEU   118     -47.759   7.530 -38.530  1.00 57.04      A    O
ATOM    475  N   LEU   119     -46.802   9.559 -38.397  1.00 52.23      A    N
ATOM    476  CA  LEU   119     -48.061  10.289 -38.399  1.00 49.64      A    C
ATOM    477  CB  LEU   119     -48.289  10.982 -37.050  1.00 52.24      A    C
ATOM    478  CG  LEU   119     -48.274  10.081 -35.808  1.00 57.18      A    C
ATOM    479  CD1 LEU   119     -48.778  10.861 -34.599  1.00 55.90      A    C
ATOM    480  CD2 LEU   119     -49.146   8.844 -36.045  1.00 57.99      A    C
ATOM    481  C   LEU   119     -47.994  11.336 -39.501  1.00 46.53      A    C
ATOM    482  O   LEU   119     -46.925  11.861 -39.808  1.00 46.48      A    O
ATOM    483  N   PRO   120     -49.134  11.634 -40.127  1.00 42.44      A    N
ATOM    484  CD  PRO   120     -50.381  10.850 -40.113  1.00 43.27      A    C
ATOM    485  CA  PRO   120     -49.213  12.766 -41.047  1.00 40.39      A    C
ATOM    486  CB  PRO   120     -50.352  12.380 -41.986  1.00 41.28      A    C
ATOM    487  CG  PRO   120     -51.269  11.576 -41.117  1.00 42.77      A    C
ATOM    488  C   PRO   120     -49.510  14.053 -40.281  1.00 36.94      A    C
ATOM    489  O   PRO   120     -50.454  14.115 -39.494  1.00 35.42      A    O
ATOM    490  N   GLY   121     -48.709  15.080 -40.524  1.00 34.63      A    N
ATOM    491  CA  GLY   121     -48.946  16.355 -39.877  1.00 33.37      A    C
ATOM    492  C   GLY   121     -47.703  17.217 -39.859  1.00 32.35      A    C
ATOM    493  O   GLY   121     -46.770  17.017 -40.649  1.00 31.71      A    O
ATOM    494  N   PHE   122     -47.678  18.189 -38.960  1.00 29.96      A    N
ATOM    495  CA  PHE   122     -46.524  19.062 -38.876  1.00 29.31      A    C
ATOM    496  CB  PHE   122     -46.594  20.142 -39.967  1.00 26.41      A    C
ATOM    497  CG  PHE   122     -47.862  20.952 -39.947  1.00 27.75      A    C
ATOM    498  CD1 PHE   122     -47.950  22.110 -39.188  1.00 26.81      A    C
ATOM    499  CD2 PHE   122     -48.951  20.582 -40.726  1.00 29.77      A    C
ATOM    500  CE1 PHE   122     -49.094  22.893 -39.204  1.00 28.13      A    C
ATOM    501  CE2 PHE   122     -50.104  21.358 -40.750  1.00 29.99      A    C
ATOM    502  CZ  PHE   122     -50.171  22.519 -39.984  1.00 31.24      A    C
ATOM    503  C   PHE   122     -46.397  19.697 -37.506  1.00 29.39      A    C
ATOM    504  O   PHE   122     -47.327  19.652 -36.694  1.00 29.97      A    O
ATOM    505  N   LEU   123     -45.224  20.273 -37.260  1.00 28.83      A    N
ATOM    506  CA  LEU   123     -44.905  20.916 -35.996  1.00 28.28      A    C
ATOM    507  CB  LEU   123     -43.564  20.388 -35.475  1.00 27.66      A    C
ATOM    508  CG  LEU   123     -43.059  20.877 -34.115  1.00 29.62      A    C
ATOM    509  CD1 LEU   123     -44.026  20.443 -33.014  1.00 28.28      A    C
ATOM    510  CD2 LEU   123     -41.672  20.302 -33.859  1.00 27.46      A    C
ATOM    511  C   LEU   123     -44.803  22.405 -36.288  1.00 29.10      A    C
ATOM    512  O   LEU   123     -44.065  22.816 -37.189  1.00 29.03      A    O
ATOM    513  N   VAL   124     -45.550  23.213 -35.542  1.00 27.38      A    N
ATOM    514  CA  VAL   124     -45.576  24.643 -35.802  1.00 26.87      A    C
ATOM    515  CB  VAL   124     -46.916  25.073 -36.456  1.00 26.05      A    C
ATOM    516  CG1 VAL   124     -48.076  24.763 -35.536  1.00 24.54      A    C
ATOM    517  CG2 VAL   124     -46.889  26.561 -36.776  1.00 28.03      A    C
ATOM    518  C   VAL   124     -45.363  25.447 -34.526  1.00 29.30      A    C
ATOM    519  O   VAL   124     -45.985  25.179 -33.486  1.00 28.28      A    O
ATOM    520  N   LYS   125     -44.469  26.425 -34.606  1.00 29.09      A    N
ATOM    521  CA  LYS   125     -44.318  27.401 -33.538  1.00 32.58      A    C
ATOM    522  CB  LYS   125     -42.848  27.803 -33.411  1.00 35.01      A    C
ATOM    523  CG  LYS   125     -42.592  28.896 -32.398  1.00 39.45      A    C
ATOM    524  CD  LYS   125     -41.413  28.552 -31.512  1.00 47.60      A    C
ATOM    525  CE  LYS   125     -40.270  29.542 -31.691  1.00 51.01      A    C
ATOM    526  NZ  LYS   125     -40.658  30.922 -31.270  1.00 53.91      A    N
ATOM    527  C   LYS   125     -45.177  28.624 -33.872  1.00 33.31      A    C
ATOM    528  O   LYS   125     -44.953  29.288 -34.884  1.00 32.62      A    O
ATOM    529  N   MET   126     -46.168  28.908 -33.031  1.00 32.61      A    N
ATOM    530  CA  MET   126     -47.086  30.010 -33.300  1.00 34.07      A    C
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ATOM    531  CB  MET   126      -48.114  29.606 -34.350  1.00 33.99      A    C
ATOM    532  CG  MET   126      -49.105  28.574 -33.839  1.00 34.58      A    C
ATOM    533  SD  MET   126      -50.324  28.127 -35.077  1.00 40.88      A    S
ATOM    534  CE  MET   126      -51.558  27.301 -34.063  1.00 35.19      A    C
ATOM    535  C   MET   126      -47.818  30.398 -32.031  1.00 35.02      A    C
ATOM    536  O   MET   126      -47.805  29.660 -31.050  1.00 34.82      A    O
ATOM    537  N   SER   127      -48.468  31.558 -32.065  1.00 36.79      A    N
ATOM    538  CA  SER   127      -49.278  32.027 -30.950  1.00 37.25      A    C
ATOM    539  CB  SER   127      -49.715  33.476 -31.191  1.00 38.68      A    C
ATOM    540  OG  SER   127      -50.780  33.843 -30.324  1.00 38.32      A    O
ATOM    541  C   SER   127      -50.513  31.154 -30.760  1.00 38.35      A    C
ATOM    542  O   SER   127      -51.135  30.722 -31.735  1.00 37.21      A    O
ATOM    543  N   GLY   128      -50.868  30.911 -29.499  1.00 37.77      A    N
ATOM    544  CA  GLY   128      -52.105  30.216 -29.194  1.00 38.41      A    C
ATOM    545  C   GLY   128      -53.353  30.935 -29.684  1.00 39.05      A    C
ATOM    546  O   GLY   128      -54.423  30.331 -29.770  1.00 39.27      A    O
ATOM    547  N   ASP   129      -53.228  32.218 -30.013  1.00 40.07      A    N
ATOM    548  CA  ASP   129      -54.350  32.965 -30.590  1.00 42.01      A    C
ATOM    549  CB  ASP   129      -53.927  34.389 -30.966  1.00 42.77      A    C
ATOM    550  CG  ASP   129      -53.629  35.254 -29.754  1.00 46.29      A    C
ATOM    551  OD1 ASP   129      -54.080  34.900 -28.640  1.00 44.66      A    O
ATOM    552  OD2 ASP   129      -52.943  36.292 -29.921  1.00 48.24      A    O
ATOM    553  C   ASP   129      -54.878  32.275 -31.842  1.00 42.68      A    C
ATOM    554  O   ASP   129      -56.064  32.371 -32.158  1.00 43.03      A    O
ATOM    555  N   LEU   130      -53.988  31.581 -32.548  1.00 40.92      A    N
ATOM    556  CA  LEU   130      -54.286  31.050 -33.871  1.00 39.89      A    C
ATOM    557  CB  LEU   130      -53.007  31.009 -34.712  1.00 38.83      A    C
ATOM    558  CG  LEU   130      -52.325  32.351 -34.993  1.00 39.26      A    C
ATOM    559  CD1 LEU   130      -50.980  32.102 -35.649  1.00 37.90      A    C
ATOM    560  CD2 LEU   130      -53.211  33.209 -35.892  1.00 38.49      A    C
ATOM    561  C   LEU   130      -54.916  29.658 -33.840  1.00 40.44      A    C
ATOM    562  O   LEU   130      -55.200  29.082 -34.890  1.00 40.55      A    O
ATOM    563  N   LEU   131      -55.134  29.116 -32.646  1.00 41.71      A    N
ATOM    564  CA  LEU   131      -55.621  27.744 -32.522  1.00 43.42      A    C
ATOM    565  CB  LEU   131      -55.691  27.336 -31.048  1.00 44.46      A    C
ATOM    566  CG  LEU   131      -54.345  26.961 -30.421  1.00 47.94      A    C
ATOM    567  CD1 LEU   131      -54.504  26.760 -28.916  1.00 47.71      A    C
ATOM    568  CD2 LEU   131      -53.809  25.692 -31.088  1.00 46.48      A    C
ATOM    569  C   LEU   131      -56.979  27.524 -33.179  1.00 44.30      A    C
ATOM    570  O   LEU   131      -57.199  26.501 -33.827  1.00 43.27      A    O
ATOM    571  N   GLU   132      -57.890  28.479 -33.010  1.00 46.35      A    N
ATOM    572  CA  GLU   132      -59.217  28.363 -33.610  1.00 48.86      A    C
ATOM    573  CB  GLU   132      -60.113  29.525 -33.171  1.00 54.02      A    C
ATOM    574  CG  GLU   132      -60.487  29.509 -31.691  1.00 62.96      A    C
ATOM    575  CD  GLU   132      -61.250  28.252 -31.282  1.00 68.59      A    C
ATOM    576  OE1 GLU   132      -61.985  27.693 -32.128  1.00 71.67      A    O
ATOM    577  OE2 GLU   132      -61.115  27.822 -30.113  1.00 71.20      A    O
ATOM    578  C   GLU   132      -59.084  28.360 -35.126  1.00 46.75      A    C
ATOM    579  O   GLU   132      -59.755  27.596 -35.818  1.00 46.81      A    O
ATOM    580  N   LEU   133      -58.203  29.214 -35.632  1.00 44.76      A    N
ATOM    581  CA  LEU   133      -57.902  29.257 -37.056  1.00 45.11      A    C
ATOM    582  CB  LEU   133      -56.868  30.351 -37.336  1.00 45.98      A    C
ATOM    583  CG  LEU   133      -56.392  30.501 -38.783  1.00 49.58      A    C
ATOM    584  CD1 LEU   133      -57.518  31.052 -39.652  1.00 51.14      A    C
ATOM    585  CD2 LEU   133      -55.197  31.432 -38.826  1.00 49.33      A    C
ATOM    586  C   LEU   133      -57.364  27.910 -37.532  1.00 44.38      A    C
ATOM    587  O   LEU   133      -57.798  27.386 -38.556  1.00 45.21      A    O
ATOM    588  N   ALA   134      -56.424  27.346 -36.779  1.00 42.85      A    N
ATOM    589  CA  ALA   134      -55.745  26.128 -37.203  1.00 41.92      A    C
ATOM    590  CB  ALA   134      -54.524  25.871 -36.316  1.00 39.84      A    C
ATOM    591  C   ALA   134      -56.692  24.937 -37.160  1.00 41.73      A    C
ATOM    592  O   ALA   134      -56.595  24.027 -37.986  1.00 41.34      A    O
ATOM    593  N   LEU   135      -57.611  24.952 -36.198  1.00 42.08      A    N
ATOM    594  CA  LEU   135      -58.586  23.880 -36.049  1.00 43.75      A    C
ATOM    595  CB  LEU   135      -59.368  24.059 -34.746  1.00 41.32      A    C
ATOM    596  CG  LEU   135      -58.641  23.644 -33.463  1.00 40.82      A    C
ATOM    597  CD1 LEU   135      -59.491  23.981 -32.244  1.00 38.47      A    C
ATOM    598  CD2 LEU   135      -58.354  22.152 -33.506  1.00 37.33      A    C
ATOM    599  C   LEU   135      -59.559  23.796 -37.226  1.00 45.90      A    C
ATOM    600  O   LEU   135      -60.250  22.793 -37.394  1.00 46.62      A    O
ATOM    601  N   LYS   136      -59.608  24.844 -38.043  1.00 48.08      A    N
ATOM    602  CA  LYS   136      -60.506  24.866 -39.194  1.00 49.87      A    C
ATOM    603  CB  LYS   136      -61.132  26.257 -39.356  1.00 52.29      A    C
ATOM    604  CG  LYS   136      -62.148  26.597 -38.274  1.00 56.29      A    C
ATOM    605  CD  LYS   136      -62.611  28.041 -38.367  1.00 60.74      A    C
ATOM    606  CE  LYS   136      -63.461  28.418 -37.155  1.00 63.43      A    C
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ATOM    607  NZ  LYS   136     -63.810  29.871 -37.140  1.00 65.11      A    N
ATOM    608  C   LYS   136     -59.831  24.457 -40.500  1.00 49.01      A    C
ATOM    609  O   LYS   136     -60.486  24.373 -41.534  1.00 48.28      A    O
ATOM    610  N   LEU   137     -58.528  24.203 -40.457  1.00 48.46      A    N
ATOM    611  CA  LEU   137     -57.811  23.779 -41.654  1.00 48.15      A    C
ATOM    612  CB  LEU   137     -56.310  23.668 -41.373  1.00 45.68      A    C
ATOM    613  CG  LEU   137     -55.542  24.954 -41.071  1.00 45.92      A    C
ATOM    614  CD1 LEU   137     -54.210  24.602 -40.435  1.00 45.13      A    C
ATOM    615  CD2 LEU   137     -55.333  25.751 -42.347  1.00 45.54      A    C
ATOM    616  C   LEU   137     -58.331  22.434 -42.157  1.00 50.34      A    C
ATOM    617  O   LEU   137     -58.786  21.593 -41.381  1.00 50.05      A    O
ATOM    618  N   PRO   138     -58.259  22.216 -43.475  1.00 51.81      A    N
ATOM    619  CD  PRO   138     -57.640  23.117 -44.464  1.00 52.57      A    C
ATOM    620  CA  PRO   138     -58.681  20.948 -44.072  1.00 51.58      A    C
ATOM    621  CB  PRO   138     -58.532  21.194 -45.570  1.00 53.25      A    C
ATOM    622  CG  PRO   138     -57.444  22.226 -45.664  1.00 54.45      A    C
ATOM    623  C   PRO   138     -57.781  19.819 -43.595  1.00 51.03      A    C
ATOM    624  O   PRO   138     -56.592  20.030 -43.365  1.00 51.14      A    O
ATOM    625  N   HIS   139     -58.357  18.630 -43.445  1.00 49.72      A    N
ATOM    626  CA  HIS   139     -57.612  17.420 -43.104  1.00 49.11      A    C
ATOM    627  CB  HIS   139     -56.319  17.331 -43.922  1.00 53.64      A    C
ATOM    628  CG  HIS   139     -56.525  17.442 -45.400  1.00 58.94      A    C
ATOM    629  CD2 HIS   139     -55.808  18.085 -46.354  1.00 60.61      A    C
ATOM    630  ND1 HIS   139     -57.578  16.838 -46.055  1.00 61.11      A    N
ATOM    631  CE1 HIS   139     -57.500  17.102 -47.347  1.00 61.63      A    C
ATOM    632  NE2 HIS   139     -56.435  17.857 -47.555  1.00 62.34      A    N
ATOM    633  C   HIS   139     -57.263  17.285 -41.622  1.00 45.66      A    C
ATOM    634  O   HIS   139     -56.773  16.242 -41.200  1.00 45.43      A    O
ATOM    635  N   VAL   140     -57.501  18.326 -40.830  1.00 41.49      A    N
ATOM    636  CA  VAL   140     -57.052  18.302 -39.446  1.00 39.22      A    C
ATOM    637  CB  VAL   140     -57.126  19.687 -38.790  1.00 38.25      A    C
ATOM    638  CG1 VAL   140     -56.864  19.562 -37.301  1.00 36.67      A    C
ATOM    639  CG2 VAL   140     -56.103  20.613 -39.421  1.00 36.79      A    C
ATOM    640  C   VAL   140     -57.867  17.343 -38.606  1.00 40.24      A    C
ATOM    641  O   VAL   140     -59.085  17.463 -38.518  1.00 40.95      A    O
ATOM    642  N   ASP   141     -57.180  16.387 -37.990  1.00 40.17      A    N
ATOM    643  CA  ASP   141     -57.803  15.464 -37.048  1.00 39.16      A    C
ATOM    644  CB  ASP   141     -57.006  14.154 -37.017  1.00 40.87      A    C
ATOM    645  CG  ASP   141     -57.716  13.050 -36.253  1.00 42.53      A    C
ATOM    646  OD1 ASP   141     -58.728  13.338 -35.574  1.00 46.58      A    O
ATOM    647  OD2 ASP   141     -57.257  11.890 -36.333  1.00 40.88      A    O
ATOM    648  C   ASP   141     -57.826  16.110 -35.658  1.00 38.98      A    C
ATOM    649  O   ASP   141     -58.887  16.284 -35.058  1.00 39.39      A    O
ATOM    650  N   TYR   142     -56.653  16.476 -35.151  1.00 37.04      A    N
ATOM    651  CA  TYR   142     -56.576  17.204 -33.888  1.00 34.27      A    C
ATOM    652  CB  TYR   142     -56.771  16.246 -32.709  1.00 34.16      A    C
ATOM    653  CG  TYR   142     -55.712  15.171 -32.610  1.00 33.65      A    C
ATOM    654  CD1 TYR   142     -54.583  15.355 -31.824  1.00 33.96      A    C
ATOM    655  CE1 TYR   142     -53.614  14.369 -31.716  1.00 35.51      A    C
ATOM    656  CD2 TYR   142     -55.845  13.968 -33.294  1.00 34.96      A    C
ATOM    657  CE2 TYR   142     -54.879  12.974 -33.194  1.00 35.75      A    C
ATOM    658  CZ  TYR   142     -53.767  13.183 -32.400  1.00 36.08      A    C
ATOM    659  OH  TYR   142     -52.808  12.205 -32.281  1.00 36.47      A    O
ATOM    660  C   TYR   142     -55.245  17.928 -33.753  1.00 32.82      A    C
ATOM    661  O   TYR   142     -54.307  17.675 -34.513  1.00 33.32      A    O
ATOM    662  N   ILE   143     -55.175  18.836 -32.787  1.00 31.85      A    N
ATOM    663  CA  ILE   143     -53.972  19.615 -32.529  1.00 31.41      A    C
ATOM    664  CB  ILE   143     -54.234  21.113 -32.796  1.00 30.29      A    C
ATOM    665  CG2 ILE   143     -53.001  21.943 -32.452  1.00 28.35      A    C
ATOM    666  CG1 ILE   143     -54.622  21.306 -34.262  1.00 30.00      A    C
ATOM    667  CD1 ILE   143     -54.835  22.751 -34.658  1.00 26.92      A    C
ATOM    668  C   ILE   143     -53.538  19.429 -31.074  1.00 32.89      A    C
ATOM    669  O   ILE   143     -54.354  19.544 -30.152  1.00 33.57      A    O
ATOM    670  N   GLU   144     -52.258  19.141 -30.860  1.00 31.53      A    N
ATOM    671  CA  GLU   144     -51.760  18.993 -29.501  1.00 32.32      A    C
ATOM    672  CB  GLU   144     -51.252  17.567 -29.262  1.00 34.26      A    C
ATOM    673  CG  GLU   144     -50.743  17.356 -27.841  1.00 39.57      A    C
ATOM    674  CD  GLU   144     -50.533  15.893 -27.496  1.00 43.13      A    C
ATOM    675  OE1 GLU   144     -51.317  15.052 -27.991  1.00 45.28      A    O
ATOM    676  OE2 GLU   144     -49.584  15.588 -26.731  1.00 41.06      A    O
ATOM    677  C   GLU   144     -50.662  19.990 -29.147  1.00 30.86      A    C
ATOM    678  O   GLU   144     -49.686  20.164 -29.882  1.00 31.43      A    O
ATOM    679  N   GLU   145     -50.831  20.643 -28.005  1.00 30.15      A    N
ATOM    680  CA  GLU   145     -49.824  21.553 -27.476  1.00 29.60      A    C
ATOM    681  CB  GLU   145     -50.415  22.337 -26.310  1.00 29.75      A    C
ATOM    682  CG  GLU   145     -49.471  23.329 -25.667  1.00 33.69      A    C
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ATOM    683  CD  GLU   145     -50.070  23.931 -24.404  1.00 36.75      A    C
ATOM    684  OE1 GLU   145     -49.915  23.325 -23.319  1.00 34.36      A    O
ATOM    685  OE2 GLU   145     -50.705  25.004 -24.499  1.00 39.41      A    O
ATOM    686  C   GLU   145     -48.618  20.749 -26.997  1.00 28.98      A    C
ATOM    687  O   GLU   145     -48.777  19.735 -26.312  1.00 25.84      A    O
ATOM    688  N   ASP   146     -47.417  21.197 -27.350  1.00 27.03      A    N
ATOM    689  CA  ASP   146     -46.210  20.501 -26.925  1.00 28.33      A    C
ATOM    690  CB  ASP   146     -44.974  21.179 -27.513  1.00 30.15      A    C
ATOM    691  CG  ASP   146     -43.820  20.205 -27.738  1.00 34.26      A    C
ATOM    692  OD1 ASP   146     -43.952  19.010 -27.376  1.00 33.09      A    O
ATOM    693  OD2 ASP   146     -42.781  20.640 -28.285  1.00 34.49      A    O
ATOM    694  C   ASP   146     -46.119  20.489 -25.395  1.00 28.56      A    C
ATOM    695  O   ASP   146     -46.792  21.263 -24.712  1.00 27.12      A    O
ATOM    696  N   SER   147     -45.298  19.596 -24.858  1.00 28.76      A    N
ATOM    697  CA  SER   147     -45.097  19.524 -23.415  1.00 29.70      A    C
ATOM    698  CB  SER   147     -46.315  18.887 -22.732  1.00 31.44      A    C
ATOM    699  OG  SER   147     -46.524  17.558 -23.186  1.00 37.75      A    O
ATOM    700  C   SER   147     -43.843  18.713 -23.115  1.00 28.55      A    C
ATOM    701  O   SER   147     -43.271  18.091 -24.015  1.00 27.48      A    O
ATOM    702  N   SER   148     -43.421  18.727 -21.854  1.00 25.24      A    N
ATOM    703  CA  SER   148     -42.114  18.204 -21.478  1.00 26.40      A    C
ATOM    704  CB  SER   148     -41.591  18.950 -20.244  1.00 27.71      A    C
ATOM    705  OG  SER   148     -41.445  20.339 -20.503  1.00 28.90      A    O
ATOM    706  C   SER   148     -42.137  16.710 -21.178  1.00 27.07      A    C
ATOM    707  O   SER   148     -43.147  16.173 -20.707  1.00 27.21      A    O
ATOM    708  N   VAL   149     -41.016  16.046 -21.441  1.00 25.66      A    N
ATOM    709  CA  VAL   149     -40.784  14.704 -20.915  1.00 24.89      A    C
ATOM    710  CB  VAL   149     -40.688  13.659 -22.044  1.00 25.33      A    C
ATOM    711  CG1 VAL   149     -41.989  13.632 -22.833  1.00 24.08      A    C
ATOM    712  CG2 VAL   149     -39.513  13.983 -22.962  1.00 21.20      A    C
ATOM    713  C   VAL   149     -39.485  14.704 -20.111  1.00 26.06      A    C
ATOM    714  O   VAL   149     -38.609  15.546 -20.335  1.00 25.28      A    O
ATOM    715  N   PHE   150     -39.370  13.768 -19.172  1.00 24.42      A    N
ATOM    716  CA  PHE   150     -38.287  13.797 -18.188  1.00 25.00      A    C
ATOM    717  CB  PHE   150     -38.824  14.250 -16.820  1.00 20.88      A    C
ATOM    718  CG  PHE   150     -39.485  15.601 -16.838  1.00 24.00      A    C
ATOM    719  CD1 PHE   150     -40.842  15.721 -17.094  1.00 23.42      A    C
ATOM    720  CD2 PHE   150     -38.752  16.752 -16.574  1.00 22.91      A    C
ATOM    721  CE1 PHE   150     -41.457  16.958 -17.086  1.00 22.83      A    C
ATOM    722  CE2 PHE   150     -39.360  17.998 -16.565  1.00 21.63      A    C
ATOM    723  CZ  PHE   150     -40.712  18.103 -16.820  1.00 23.76      A    C
ATOM    724  C   PHE   150     -37.642  12.421 -18.033  1.00 23.77      A    C
ATOM    725  O   PHE   150     -38.326  11.399 -18.040  1.00 23.53      A    O
ATOM    726  N   ALA   151     -36.326  12.404 -17.874  1.00 24.56      A    N
ATOM    727  CA  ALA   151     -35.626  11.185 -17.508  1.00 25.41      A    C
ATOM    728  CB  ALA   151     -34.193  11.513 -17.136  1.00 24.13      A    C
ATOM    729  C   ALA   151     -36.332  10.537 -16.324  1.00 28.65      A    C
ATOM    730  O   ALA   151     -36.760  11.228 -15.395  1.00 28.24      A    O
ATOM    731  N   GLN   152     -36.467   9.213 -16.349  1.00 30.75      A    N
ATOM    732  CA  GLN   152     -36.936   8.504 -15.160  1.00 32.33      A    C
ATOM    733  CB  GLN   152     -38.119   7.599 -15.515  1.00 30.09      A    C
ATOM    734  CG  GLN   152     -39.322   8.338 -16.062  1.00 26.52      A    C
ATOM    735  CD  GLN   152     -39.871   9.357 -15.078  1.00 29.34      A    C
ATOM    736  OE1 GLN   152     -40.420   8.999 -14.037  1.00 26.44      A    O
ATOM    737  NE2 GLN   152     -39.722  10.637 -15.405  1.00 27.86      A    N
ATOM    738  C   GLN   152     -35.802   7.678 -14.544  1.00 34.54      A    C
ATOM    739  O   GLN   152     -36.084   6.635 -13.913  1.00 35.69      A    O
ATOM    740  OXT GLN   152     -34.632   8.101 -14.686  1.00 36.82      A    O
TER     741      GLN   152                                              A
ATOM    742  CB  SER   153     -18.830 -12.304  -7.860  1.00 80.02      B    C
ATOM    743  OG  SER   153     -19.427 -13.535  -8.246  1.00 82.63      B    O
ATOM    744  C   SER   153     -20.846 -11.007  -8.585  1.00 76.11      B    C
ATOM    745  O   SER   153     -20.475 -10.397  -9.589  1.00 76.49      B    O
ATOM    746  N   SER   153     -20.624 -11.755  -6.227  1.00 78.23      B    N
ATOM    747  CA  SER   153     -19.883 -11.272  -7.430  1.00 78.01      B    C
ATOM    748  N   ILE   154     -22.081 -11.479  -8.439  1.00 72.80      B    N
ATOM    749  CA  ILE   154     -23.127 -11.189  -9.412  1.00 68.74      B    C
ATOM    750  CB  ILE   154     -23.892 -12.470  -9.810  1.00 69.24      B    C
ATOM    751  CG2 ILE   154     -25.019 -12.132 -10.773  1.00 67.60      B    C
ATOM    752  CG1 ILE   154     -22.925 -13.469 -10.450  1.00 70.75      B    C
ATOM    753  CD1 ILE   154     -22.160 -12.910 -11.639  1.00 70.73      B    C
ATOM    754  C   ILE   154     -24.115 -10.187  -8.825  1.00 65.36      B    C
ATOM    755  O   ILE   154     -24.664 -10.406  -7.742  1.00 65.44      B    O
ATOM    756  N   PRO   155     -24.353  -9.072  -9.536  1.00 61.02      B    N
ATOM    757  CD  PRO   155     -23.763  -8.705 -10.836  1.00 60.32      B    C
ATOM    758  CA  PRO   155     -25.287  -8.050  -9.051  1.00 56.72      B    C
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ATOM    759  CB  PRO   155      -25.337  -7.035 -10.194  1.00 57.80      B    C
ATOM    760  CG  PRO   155      -24.044  -7.232 -10.926  1.00 58.94      B    C
ATOM    761  C   PRO   155      -26.648  -8.670  -8.768  1.00 53.19      B    C
ATOM    762  O   PRO   155      -27.121  -9.515  -9.529  1.00 52.31      B    O
ATOM    763  N   TRP   156      -27.273  -8.252  -7.671  1.00 49.27      B    N
ATOM    764  CA  TRP   156      -28.513  -8.872  -7.221  1.00 47.58      B    C
ATOM    765  CB  TRP   156      -29.045  -8.152  -5.978  1.00 45.42      B    C
ATOM    766  CG  TRP   156      -29.708  -6.840  -6.286  1.00 46.00      B    C
ATOM    767  CD2 TRP   156      -31.083  -6.637  -6.635  1.00 44.51      B    C
ATOM    768  CE2 TRP   156      -31.263  -5.253  -6.836  1.00 44.08      B    C
ATOM    769  CE3 TRP   156      -32.178  -7.492  -6.798  1.00 44.17      B    C
ATOM    770  CD1 TRP   156      -29.125  -5.603  -6.290  1.00 45.31      B    C
ATOM    771  NE1 TRP   156      -30.054  -4.645  -6.620  1.00 45.73      B    N
ATOM    772  CZ2 TRP   156      -32.494  -4.709  -7.189  1.00 44.63      B    C
ATOM    773  CZ3 TRP   156      -33.402  -6.945  -7.150  1.00 46.04      B    C
ATOM    774  CH2 TRP   156      -33.550  -5.562  -7.342  1.00 44.87      B    C
ATOM    775  C   TRP   156      -29.572  -8.841  -8.319  1.00 46.81      B    C
ATOM    776  O   TRP   156      -30.349  -9.784  -8.470  1.00 46.89      B    O
ATOM    777  N   ASN   157      -29.592  -7.752  -9.084  1.00 45.77      B    N
ATOM    778  CA  ASN   157      -30.608  -7.547 -10.110  1.00 44.93      B    C
ATOM    779  CB  ASN   157      -30.586  -6.094 -10.592  1.00 43.13      B    C
ATOM    780  CG  ASN   157      -29.204  -5.642 -10.993  1.00 42.32      B    C
ATOM    781  OD1 ASN   157      -28.302  -5.561 -10.159  1.00 43.42      B    O
ATOM    782  ND2 ASN   157      -29.024  -5.346 -12.276  1.00 38.97      B    N
ATOM    783  C   ASN   157      -30.433  -8.478 -11.302  1.00 44.94      B    C
ATOM    784  O   ASN   157      -31.416  -8.900 -11.907  1.00 43.18      B    O
ATOM    785  N   LEU   158      -29.187  -8.793 -11.646  1.00 47.58      B    N
ATOM    786  CA  LEU   158      -28.925  -9.724 -12.742  1.00 50.92      B    C
ATOM    787  CB  LEU   158      -27.458  -9.672 -13.166  1.00 49.39      B    C
ATOM    788  CG  LEU   158      -27.030  -8.392 -13.884  1.00 50.63      B    C
ATOM    789  CD1 LEU   158      -25.669  -8.602 -14.531  1.00 48.36      B    C
ATOM    790  CD2 LEU   158      -28.074  -8.020 -14.933  1.00 48.22      B    C
ATOM    791  C   LEU   158      -29.282 -11.147 -12.338  1.00 52.96      B    C
ATOM    792  O   LEU   158      -29.812 -11.914 -13.139  1.00 54.48      B    O
ATOM    793  N   GLU   159      -28.989 -11.493 -11.091  1.00 55.23      B    N
ATOM    794  CA  GLU   159      -29.383 -12.783 -10.541  1.00 58.07      B    C
ATOM    795  CB  GLU   159      -28.799 -12.949  -9.132  1.00 61.02      B    C
ATOM    796  CG  GLU   159      -29.709 -13.689  -8.161  1.00 66.59      B    C
ATOM    797  CD  GLU   159      -29.956 -12.905  -6.876  1.00 71.01      B    C
ATOM    798  OE1 GLU   159      -31.037 -12.282  -6.747  1.00 71.63      B    O
ATOM    799  OE2 GLU   159      -29.068 -12.914  -5.992  1.00 72.55      B    O
ATOM    800  C   GLU   159      -30.906 -12.907 -10.490  1.00 58.29      B    C
ATOM    801  O   GLU   159      -31.453 -13.990 -10.695  1.00 56.29      B    O
ATOM    802  N   ARG   160      -31.583 -11.790 -10.230  1.00 59.46      B    N
ATOM    803  CA  ARG   160      -33.016 -11.803  -9.944  1.00 60.58      B    C
ATOM    804  CB  ARG   160      -33.447 -10.460  -9.347  1.00 59.26      B    C
ATOM    805  CG  ARG   160      -34.890 -10.429  -8.869  1.00 59.38      B    C
ATOM    806  CD  ARG   160      -35.116 -11.435  -7.746  1.00 60.25      B    C
ATOM    807  NE  ARG   160      -34.131 -11.276  -6.678  1.00 60.39      B    N
ATOM    808  CZ  ARG   160      -34.313 -10.518  -5.601  1.00 61.06      B    C
ATOM    809  NH1 ARG   160      -33.360 -10.429  -4.681  1.00 59.91      B    N
ATOM    810  NH2 ARG   160      -35.450  -9.849  -5.442  1.00 61.11      B    N
ATOM    811  C   ARG   160      -33.874 -12.114 -11.172  1.00 61.99      B    C
ATOM    812  O   ARG   160      -34.907 -12.780 -11.064  1.00 61.34      B    O
ATOM    813  N   ILE   161      -33.449 -11.629 -12.335  1.00 63.71      B    N
ATOM    814  CA  ILE   161      -34.165 -11.912 -13.575  1.00 66.31      B    C
ATOM    815  CB  ILE   161      -33.882 -10.837 -14.652  1.00 64.21      B    C
ATOM    816  CG2 ILE   161      -34.367  -9.476 -14.172  1.00 63.31      B    C
ATOM    817  CG1 ILE   161      -32.387 -10.803 -14.970  1.00 63.10      B    C
ATOM    818  CD1 ILE   161      -32.035  -9.948 -16.159  1.00 62.54      B    C
ATOM    819  C   ILE   161      -33.786 -13.282 -14.144  1.00 68.85      B    C
ATOM    820  O   ILE   161      -34.429 -13.780 -15.068  1.00 68.03      B    O
ATOM    821  N   THR   162      -32.738 -13.885 -13.590  1.00 72.56      B    N
ATOM    822  CA  THR   162      -32.312 -15.215 -14.016  1.00 77.56      B    C
ATOM    823  CB  THR   162      -30.803 -15.420 -13.772  1.00 77.41      B    C
ATOM    824  OG1 THR   162      -30.059 -14.501 -14.582  1.00 76.98      B    O
ATOM    825  CG2 THR   162      -30.395 -16.843 -14.120  1.00 76.95      B    C
ATOM    826  C   THR   162      -33.082 -16.307 -13.277  1.00 81.06      B    C
ATOM    827  O   THR   162      -33.004 -16.420 -12.054  1.00 80.79      B    O
ATOM    828  N   PRO   163      -33.840 -17.125 -14.024  1.00 85.12      B    N
ATOM    829  CD  PRO   163      -33.917 -17.037 -15.493  1.00 86.11      B    C
ATOM    830  CA  PRO   163      -34.684 -18.206 -13.499  1.00 88.61      B    C
ATOM    831  CB  PRO   163      -35.566 -18.574 -14.689  1.00 87.63      B    C
ATOM    832  CG  PRO   163      -34.720 -18.256 -15.875  1.00 86.84      B    C
ATOM    833  C   PRO   163      -33.869 -19.403 -12.999  1.00 92.41      B    C
ATOM    834  O   PRO   163      -32.664 -19.493 -13.242  1.00 91.73      B    O
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ATOM    835  N   PRO   164     -34.527 -20.342 -12.296  1.00 96.24      B    N
ATOM    836  CD  PRO   164     -35.931 -20.261 -11.855  1.00 97.02      B    C
ATOM    837  CA  PRO   164     -33.868 -21.552 -11.788  1.00 99.15      B    C
ATOM    838  CB  PRO   164     -35.004 -22.323 -11.117  1.00 98.26      B    C
ATOM    839  CG  PRO   164     -35.994 -21.271 -10.743  1.00 97.35      B    C
ATOM    840  C   PRO   164     -33.181 -22.377 -12.878  1.00102.09      B    C
ATOM    841  O   PRO   164     -31.965 -22.576 -12.843  1.00101.97      B    O
ATOM    842  N   ARG   165     -33.962 -22.851 -13.844  1.00105.51      B    N
ATOM    843  CA  ARG   165     -33.429 -23.677 -14.924  1.00109.14      B    C
ATOM    844  CB  ARG   165     -34.487 -24.687 -15.379  1.00111.03      B    C
ATOM    845  CG  ARG   165     -33.981 -26.117 -15.510  1.00114.19      B    C
ATOM    846  CD  ARG   165     -32.793 -26.214 -16.458  1.00116.65      B    C
ATOM    847  NE  ARG   165     -32.370 -27.598 -16.661  1.00118.62      B    N
ATOM    848  CZ  ARG   165     -32.623 -28.306 -17.759  1.00119.70      B    C
ATOM    849  NH1 ARG   165     -32.201 -29.560 -17.854  1.00120.20      B    N
ATOM    850  NH2 ARG   165     -33.293 -27.758 -18.764  1.00119.97      B    N
ATOM    851  C   ARG   165     -33.005 -22.809 -16.108  1.00110.22      B    C
ATOM    852  O   ARG   165     -33.766 -22.624 -17.056  1.00110.60      B    O
ATOM    853  N   TYR   166     -31.787 -22.282 -16.052  1.00111.28      B    N
ATOM    854  CA  TYR   166     -31.320 -21.344 -17.066  1.00112.25      B    C
ATOM    855  CB  TYR   166     -30.758 -20.088 -16.391  1.00113.37      B    C
ATOM    856  CG  TYR   166     -30.241 -19.042 -17.354  1.00114.53      B    C
ATOM    857  CD1 TYR   166     -28.954 -18.532 -17.228  1.00114.68      B    C
ATOM    858  CE1 TYR   166     -28.471 -17.583 -18.110  1.00115.01      B    C
ATOM    859  CD2 TYR   166     -31.035 -18.570 -18.393  1.00114.92      B    C
ATOM    860  CE2 TYR   166     -30.561 -17.619 -19.281  1.00114.93      B    C
ATOM    861  CZ  TYR   166     -29.278 -17.130 -19.135  1.00115.16      B    C
ATOM    862  OH  TYR   166     -28.797 -16.189 -20.017  1.00115.03      B    O
ATOM    863  C   TYR   166     -30.265 -21.965 -17.981  1.00112.12      B    C
ATOM    864  O   TYR   166     -29.207 -22.397 -17.525  1.00111.99      B    O
ATOM    865  N   TYR   171     -28.363 -19.940 -25.299  1.00 96.76      B    N
ATOM    866  CA  TYR   171     -29.633 -19.258 -25.518  1.00 96.87      B    C
ATOM    867  CB  TYR   171     -29.623 -17.883 -24.836  1.00 96.75      B    C
ATOM    868  CG  TYR   171     -30.934 -17.536 -24.160  1.00 96.85      B    C
ATOM    869  CD1 TYR   171     -31.135 -17.811 -22.812  1.00 96.51      B    C
ATOM    870  CE1 TYR   171     -32.345 -17.539 -22.198  1.00 96.16      B    C
ATOM    871  CD2 TYR   171     -31.984 -16.972 -24.877  1.00 96.54      B    C
ATOM    872  CE2 TYR   171     -33.199 -16.697 -24.270  1.00 95.77      B    C
ATOM    873  CZ  TYR   171     -33.373 -16.984 -22.932  1.00 96.10      B    C
ATOM    874  OH  TYR   171     -34.584 -16.727 -22.327  1.00 97.20      B    O
ATOM    875  C   TYR   171     -29.890 -19.090 -27.013  1.00 97.20      B    C
ATOM    876  O   TYR   171     -28.992 -19.292 -27.833  1.00 97.67      B    O
ATOM    877  N   LEU   179     -27.947  -9.679 -34.479  1.00 67.11      B    N
ATOM    878  CA  LEU   179     -29.202  -9.880 -35.196  1.00 67.55      B    C
ATOM    879  CB  LEU   179     -29.749 -11.274 -34.885  1.00 68.43      B    C
ATOM    880  CG  LEU   179     -30.345 -12.050 -36.062  1.00 70.66      B    C
ATOM    881  CD1 LEU   179     -30.862 -13.395 -35.557  1.00 70.04      B    C
ATOM    882  CD2 LEU   179     -31.457 -11.237 -36.722  1.00 68.78      B    C
ATOM    883  C   LEU   179     -30.251  -8.816 -34.831  1.00 66.57      B    C
ATOM    884  O   LEU   179     -30.917  -8.255 -35.704  1.00 67.25      B    O
ATOM    885  N   VAL   180     -30.392  -8.542 -33.537  1.00 63.99      B    N
ATOM    886  CA  VAL   180     -31.393  -7.595 -33.052  1.00 61.13      B    C
ATOM    887  CB  VAL   180     -32.163  -8.180 -31.836  1.00 61.02      B    C
ATOM    888  CG1 VAL   180     -31.183  -8.638 -30.773  1.00 60.46      B    C
ATOM    889  CG2 VAL   180     -33.110  -7.133 -31.260  1.00 60.55      B    C
ATOM    890  C   VAL   180     -30.765  -6.260 -32.651  1.00 58.49      B    C
ATOM    891  O   VAL   180     -29.693  -6.223 -32.049  1.00 58.27      B    O
ATOM    892  N   GLU   181     -31.434  -5.162 -32.988  1.00 56.10      B    N
ATOM    893  CA  GLU   181     -30.940  -3.838 -32.618  1.00 54.53      B    C
ATOM    894  CB  GLU   181     -30.896  -2.931 -33.852  1.00 56.53      B    C
ATOM    895  CG  GLU   181     -29.749  -1.928 -33.833  1.00 62.79      B    C
ATOM    896  CD  GLU   181     -28.759  -2.146 -34.968  1.00 66.51      B    C
ATOM    897  OE1 GLU   181     -27.537  -2.078 -34.714  1.00 69.34      B    O
ATOM    898  OE2 GLU   181     -29.201  -2.383 -36.116  1.00 68.31      B    O
ATOM    899  C   GLU   181     -31.810  -3.196 -31.532  1.00 51.03      B    C
ATOM    900  O   GLU   181     -33.041  -3.231 -31.609  1.00 50.12      B    O
ATOM    901  N   VAL   182     -31.166  -2.615 -30.522  1.00 46.36      B    N
ATOM    902  CA  VAL   182     -31.879  -1.871 -29.485  1.00 44.68      B    C
ATOM    903  CB  VAL   182     -31.417  -2.288 -28.073  1.00 44.84      B    C
ATOM    904  CG1 VAL   182     -32.258  -1.584 -27.023  1.00 43.44      B    C
ATOM    905  CG2 VAL   182     -31.521  -3.796 -27.911  1.00 47.01      B    C
ATOM    906  C   VAL   182     -31.643  -0.365 -29.633  1.00 43.25      B    C
ATOM    907  O   VAL   182     -30.504   0.099 -29.566  1.00 42.77      B    O
ATOM    908  N   TYR   183     -32.715   0.397 -29.837  1.00 39.38      B    N
ATOM    909  CA  TYR   183     -32.602   1.851 -29.872  1.00 38.75      B    C
ATOM    910  CB  TYR   183     -33.594   2.448 -30.872  1.00 39.87      B    C
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ATOM    911  CG  TYR   183     -33.217   2.225 -32.319  1.00 42.94      B    C
ATOM    912  CD1 TYR   183     -33.512   1.026 -32.959  1.00 43.91      B    C
ATOM    913  CE1 TYR   183     -33.177   0.824 -34.290  1.00 46.07      B    C
ATOM    914  CD2 TYR   183     -32.575   3.217 -33.049  1.00 43.64      B    C
ATOM    915  CE2 TYR   183     -32.235   3.025 -34.379  1.00 44.65      B    C
ATOM    916  CZ  TYR   183     -32.539   1.828 -34.994  1.00 45.86      B    C
ATOM    917  OH  TYR   183     -32.209   1.635 -36.319  1.00 48.85      B    O
ATOM    918  C   TYR   183     -32.859   2.442 -28.492  1.00 37.29      B    C
ATOM    919  O   TYR   183     -33.803   2.052 -27.804  1.00 34.89      B    O
ATOM    920  N   LEU   184     -32.005   3.380 -28.097  1.00 35.24      B    N
ATOM    921  CA  LEU   184     -32.155   4.093 -26.835  1.00 34.10      B    C
ATOM    922  CB  LEU   184     -30.872   3.962 -26.008  1.00 32.18      B    C
ATOM    923  CG  LEU   184     -30.740   4.832 -24.750  1.00 35.23      B    C
ATOM    924  CD1 LEU   184     -31.760   4.406 -23.708  1.00 32.47      B    C
ATOM    925  CD2 LEU   184     -29.326   4.703 -24.190  1.00 32.96      B    C
ATOM    926  C   LEU   184     -32.438   5.567 -27.122  1.00 33.43      B    C
ATOM    927  O   LEU   184     -31.631   6.245 -27.755  1.00 32.77      B    O
ATOM    928  N   LEU   185     -33.587   6.057 -26.669  1.00 33.29      B    N
ATOM    929  CA  LEU   185     -33.857   7.491 -26.690  1.00 35.12      B    C
ATOM    930  CB  LEU   185     -35.280   7.770 -27.183  1.00 34.68      B    C
ATOM    931  CG  LEU   185     -35.549   7.510 -28.664  1.00 37.37      B    C
ATOM    932  CD1 LEU   185     -35.598   6.010 -28.923  1.00 38.05      B    C
ATOM    933  CD2 LEU   185     -36.866   8.155 -29.057  1.00 39.00      B    C
ATOM    934  C   LEU   185     -33.685   8.070 -25.290  1.00 35.69      B    C
ATOM    935  O   LEU   185     -34.515   7.837 -24.413  1.00 36.10      B    O
ATOM    936  N   ASP   186     -32.610   8.824 -25.086  1.00 35.47      B    N
ATOM    937  CA  ASP   186     -32.280   9.325 -23.759  1.00 38.43      B    C
ATOM    938  CB  ASP   186     -31.663   8.204 -22.920  1.00 44.90      B    C
ATOM    939  CG  ASP   186     -31.865   8.412 -21.424  1.00 53.04      B    C
ATOM    940  OD1 ASP   186     -32.984   8.134 -20.929  1.00 56.81      B    O
ATOM    941  OD2 ASP   186     -30.909   8.853 -20.743  1.00 55.06      B    O
ATOM    942  C   ASP   186     -31.307  10.500 -23.842  1.00 37.51      B    C
ATOM    943  O   ASP   186     -31.315  11.259 -24.816  1.00 36.47      B    O
ATOM    944  N   THR   187     -30.474  10.655 -22.818  1.00 34.82      B    N
ATOM    945  CA  THR   187     -29.470  11.707 -22.830  1.00 35.00      B    C
ATOM    946  CB  THR   187     -28.870  11.957 -21.426  1.00 33.57      B    C
ATOM    947  OG1 THR   187     -28.137  10.800 -21.004  1.00 33.38      B    O
ATOM    948  CG2 THR   187     -29.968  12.267 -20.416  1.00 30.31      B    C
ATOM    949  C   THR   187     -28.341  11.270 -23.756  1.00 36.71      B    C
ATOM    950  O   THR   187     -28.362  10.162 -24.304  1.00 36.59      B    O
ATOM    951  N   SER   188     -27.360  12.144 -23.937  1.00 36.36      B    N
ATOM    952  CA  SER   188     -26.153  11.757 -24.641  1.00 39.03      B    C
ATOM    953  CB  SER   188     -25.196  12.947 -24.728  1.00 39.14      B    C
ATOM    954  OG  SER   188     -24.937  13.475 -23.443  1.00 42.99      B    O
ATOM    955  C   SER   188     -25.525  10.610 -23.851  1.00 39.61      B    C
ATOM    956  O   SER   188     -25.828  10.426 -22.666  1.00 38.53      B    O
ATOM    957  N   ILE   189     -24.680   9.822 -24.509  1.00 39.46      B    N
ATOM    958  CA  ILE   189     -24.001   8.725 -23.833  1.00 40.86      B    C
ATOM    959  CB  ILE   189     -24.496   7.342 -24.326  1.00 41.34      B    C
ATOM    960  CG2 ILE   189     -26.000   7.220 -24.131  1.00 38.20      B    C
ATOM    961  CG1 ILE   189     -24.132   7.148 -25.797  1.00 42.75      B    C
ATOM    962  CD1 ILE   189     -24.517   5.784 -26.342  1.00 41.61      B    C
ATOM    963  C   ILE   189     -22.494   8.796 -24.041  1.00 41.60      B    C
ATOM    964  O   ILE   189     -22.009   9.471 -24.946  1.00 40.48      B    O
ATOM    965  N   GLN   190     -21.763   8.102 -23.177  1.00 43.69      B    N
ATOM    966  CA  GLN   190     -20.313   8.010 -23.273  1.00 44.38      B    C
ATOM    967  CB  GLN   190     -19.717   8.024 -21.861  1.00 46.59      B    C
ATOM    968  CG  GLN   190     -18.210   7.870 -21.793  1.00 52.20      B    C
ATOM    969  CD  GLN   190     -17.488   8.863 -22.678  1.00 57.06      B    C
ATOM    970  OE1 GLN   190     -17.364  10.046 -22.337  1.00 59.79      B    O
ATOM    971  NE2 GLN   190     -17.008   8.391 -23.828  1.00 58.64      B    N
ATOM    972  C   GLN   190     -19.971   6.704 -23.996  1.00 44.38      B    C
ATOM    973  O   GLN   190     -19.828   5.657 -23.367  1.00 43.59      B    O
ATOM    974  N   SER   191     -19.854   6.768 -25.318  1.00 44.25      B    N
ATOM    975  CA  SER   191     -19.731   5.559 -26.131  1.00 46.81      B    C
ATOM    976  CB  SER   191     -20.052   5.872 -27.597  1.00 45.03      B    C
ATOM    977  OG  SER   191     -19.186   6.872 -28.106  1.00 45.93      B    O
ATOM    978  C   SER   191     -18.351   4.902 -26.035  1.00 47.69      B    C
ATOM    979  O   SER   191     -18.156   3.788 -26.520  1.00 48.01      B    O
ATOM    980  N   ASP   192     -17.404   5.595 -25.410  1.00 49.21      B    N
ATOM    981  CA  ASP   192     -16.056   5.073 -25.198  1.00 52.01      B    C
ATOM    982  CB  ASP   192     -15.052   6.223 -25.056  1.00 54.81      B    C
ATOM    983  CG  ASP   192     -14.726   6.885 -26.379  1.00 60.16      B    C
ATOM    984  OD1 ASP   192     -15.012   6.277 -27.436  1.00 62.13      B    O
ATOM    985  OD2 ASP   192     -14.181   8.014 -26.361  1.00 61.90      B    O
ATOM    986  C   ASP   192     -15.944   4.185 -23.959  1.00 51.65      B    C
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ATOM    987  O   ASP   192     -14.901   3.577 -23.721  1.00 52.13      B    O
ATOM    988  N   HIS   193     -17.001   4.124 -23.159  1.00 50.05      B    N
ATOM    989  CA  HIS   193     -16.932   3.376 -21.914  1.00 48.15      B    C
ATOM    990  CB  HIS   193     -18.204   3.569 -21.091  1.00 44.37      B    C
ATOM    991  CG  HIS   193     -18.091   3.048 -19.693  1.00 43.27      B    C
ATOM    992  CD2 HIS   193     -17.873   3.686 -18.518  1.00 40.34      B    C
ATOM    993  ND1 HIS   193     -18.178   1.705 -19.389  1.00 42.52      B    N
ATOM    994  CE1 HIS   193     -18.020   1.538 -18.088  1.00 40.12      B    C
ATOM    995  NE2 HIS   193     -17.833   2.725 -17.536  1.00 40.42      B    N
ATOM    996  C   HIS   193     -16.725   1.891 -22.188  1.00 48.80      B    C
ATOM    997  O   HIS   193     -17.291   1.333 -23.131  1.00 48.40      B    O
ATOM    998  N   ARG   194     -15.918   1.256 -21.347  1.00 49.72      B    N
ATOM    999  CA  ARG   194     -15.475  -0.106 -21.594  1.00 51.21      B    C
ATOM   1000  CB  ARG   194     -14.469  -0.527 -20.520  1.00 55.14      B    C
ATOM   1001  CG  ARG   194     -13.596  -1.708 -20.912  1.00 62.40      B    C
ATOM   1002  CD  ARG   194     -14.042  -2.993 -20.227  1.00 68.22      B    C
ATOM   1003  NE  ARG   194     -13.251  -4.143 -20.660  1.00 73.86      B    N
ATOM   1004  CZ  ARG   194     -13.497  -5.402 -20.306  1.00 76.07      B    C
ATOM   1005  NH1 ARG   194     -12.721  -6.383 -20.754  1.00 77.25      B    N
ATOM   1006  NH2 ARG   194     -14.516  -5.685 -19.504  1.00 76.69      B    N
ATOM   1007  C   ARG   194     -16.650  -1.077 -21.620  1.00 49.97      B    C
ATOM   1008  O   ARG   194     -16.568  -2.141 -22.227  1.00 49.91      B    O
ATOM   1009  N   GLU   195     -17.748  -0.710 -20.969  1.00 48.05      B    N
ATOM   1010  CA  GLU   195     -18.907  -1.591 -20.911  1.00 46.66      B    C
ATOM   1011  CB  GLU   195     -19.884  -1.116 -19.833  1.00 47.09      B    C
ATOM   1012  CG  GLU   195     -19.582  -1.665 -18.442  1.00 48.87      B    C
ATOM   1013  CD  GLU   195     -19.912  -3.146 -18.320  1.00 51.11      B    C
ATOM   1014  OE1 GLU   195     -18.981  -3.979 -18.447  1.00 52.26      B    O
ATOM   1015  OE2 GLU   195     -21.100  -3.478 -18.101  1.00 48.64      B    O
ATOM   1016  C   GLU   195     -19.634  -1.698 -22.247  1.00 46.38      B    C
ATOM   1017  O   GLU   195     -20.254  -2.718 -22.542  1.00 44.77      B    O
ATOM   1018  N   ILE   196     -19.557  -0.648 -23.057  1.00 46.54      B    N
ATOM   1019  CA  ILE   196     -20.347  -0.599 -24.280  1.00 48.16      B    C
ATOM   1020  CB  ILE   196     -21.486   0.446 -24.164  1.00 46.92      B    C
ATOM   1021  CG2 ILE   196     -22.476   0.015 -23.102  1.00 45.09      B    C
ATOM   1022  CG1 ILE   196     -20.904   1.822 -23.830  1.00 47.02      B    C
ATOM   1023  CD1 ILE   196     -21.951   2.902 -23.616  1.00 47.07      B    C
ATOM   1024  C   ILE   196     -19.533  -0.296 -25.537  1.00 50.05      B    C
ATOM   1025  O   ILE   196     -20.061  -0.348 -26.647  1.00 50.50      B    O
ATOM   1026  N   GLU   197     -18.253   0.015 -25.371  1.00 52.60      B    N
ATOM   1027  CA  GLU   197     -17.453   0.482 -26.497  1.00 55.84      B    C
ATOM   1028  CB  GLU   197     -16.007   0.709 -26.068  1.00 59.17      B    C
ATOM   1029  CG  GLU   197     -15.164   1.378 -27.140  1.00 66.13      B    C
ATOM   1030  CD  GLU   197     -13.691   1.374 -26.800  1.00 71.16      B    C
ATOM   1031  OE1 GLU   197     -13.222   0.370 -26.214  1.00 74.15      B    O
ATOM   1032  OE2 GLU   197     -13.003   2.371 -27.116  1.00 72.99      B    O
ATOM   1033  C   GLU   197     -17.478  -0.479 -27.685  1.00 55.61      B    C
ATOM   1034  O   GLU   197     -17.271  -1.684 -27.536  1.00 55.13      B    O
ATOM   1035  N   GLY   198     -17.740   0.071 -28.866  1.00 56.15      B    N
ATOM   1036  CA  GLY   198     -17.715  -0.727 -30.077  1.00 56.83      B    C
ATOM   1037  C   GLY   198     -18.987  -1.514 -30.321  1.00 57.44      B    C
ATOM   1038  O   GLY   198     -19.150  -2.122 -31.377  1.00 56.60      B    O
ATOM   1039  N   ARG   199     -19.892  -1.513 -29.349  1.00 57.74      B    N
ATOM   1040  CA  ARG   199     -21.160  -2.213 -29.500  1.00 57.82      B    C
ATOM   1041  CB  ARG   199     -21.349  -3.211 -28.355  1.00 58.86      B    C
ATOM   1042  CG  ARG   199     -20.312  -4.333 -28.336  1.00 63.05      B    C
ATOM   1043  CD  ARG   199     -20.287  -5.097 -29.659  1.00 67.78      B    C
ATOM   1044  NE  ARG   199     -21.503  -5.885 -29.868  1.00 71.51      B    N
ATOM   1045  CZ  ARG   199     -21.995  -6.214 -31.060  1.00 72.46      B    C
ATOM   1046  NH1 ARG   199     -23.109  -6.933 -31.145  1.00 72.69      B    N
ATOM   1047  NH2 ARG   199     -21.376  -5.821 -32.168  1.00 73.16      B    N
ATOM   1048  C   ARG   199     -22.353  -1.256 -29.568  1.00 57.64      B    C
ATOM   1049  O   ARG   199     -23.429  -1.633 -30.030  1.00 58.42      B    O
ATOM   1050  N   VAL   200     -22.169  -0.023 -29.105  1.00 56.67      B    N
ATOM   1051  CA  VAL   200     -23.198   1.001 -29.266  1.00 56.34      B    C
ATOM   1052  CB  VAL   200     -23.575   1.661 -27.921  1.00 56.89      B    C
ATOM   1053  CG1 VAL   200     -24.037   0.602 -26.933  1.00 59.60      B    C
ATOM   1054  CG2 VAL   200     -22.390   2.424 -27.369  1.00 58.38      B    C
ATOM   1055  C   VAL   200     -22.735   2.095 -30.219  1.00 55.25      B    C
ATOM   1056  O   VAL   200     -21.632   2.629 -30.089  1.00 55.38      B    O
ATOM   1057  N   MET   201     -23.588   2.421 -31.182  1.00 53.57      B    N
ATOM   1058  CA  MET   201     -23.303   3.491 -32.122  1.00 53.33      B    C
ATOM   1059  CB  MET   201     -23.629   3.037 -33.542  1.00 57.41      B    C
ATOM   1060  CG  MET   201     -23.473   4.127 -34.587  1.00 65.55      B    C
ATOM   1061  SD  MET   201     -24.977   4.367 -35.566  1.00 76.08      B    S
ATOM   1062  CE  MET   201     -24.388   5.567 -36.805  1.00 73.70      B    C
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ATOM   1063  C   MET   201     -24.129   4.725 -31.773  1.00 50.12      B    C
ATOM   1064  O   MET   201     -25.322   4.627 -31.491  1.00 49.66      B    O
ATOM   1065  N   VAL   202     -23.486   5.885 -31.786  1.00 46.03      B    N
ATOM   1066  CA  VAL   202     -24.199   7.143 -31.616  1.00 43.65      B    C
ATOM   1067  CB  VAL   202     -23.269   8.238 -31.051  1.00 42.67      B    C
ATOM   1068  CG1 VAL   202     -24.025   9.549 -30.928  1.00 39.96      B    C
ATOM   1069  CG2 VAL   202     -22.721   7.809 -29.699  1.00 42.01      B    C
ATOM   1070  C   VAL   202     -24.724   7.609 -32.970  1.00 42.11      B    C
ATOM   1071  O   VAL   202     -23.941   7.963 -33.847  1.00 41.82      B    O
ATOM   1072  N   THR   203     -26.042   7.607 -33.146  1.00 39.83      B    N
ATOM   1073  CA  THR   203     -26.621   8.158 -34.367  1.00 39.51      B    C
ATOM   1074  CB  THR   203     -28.127   7.878 -34.458  1.00 37.45      B    C
ATOM   1075  OG1 THR   203     -28.818   8.690 -33.499  1.00 38.09      B    O
ATOM   1076  CG2 THR   203     -28.411   6.412 -34.179  1.00 33.14      B    C
ATOM   1077  C   THR   203     -26.413   9.668 -34.327  1.00 41.01      B    C
ATOM   1078  O   THR   203     -26.080  10.233 -33.284  1.00 41.22      B    O
ATOM   1079  N   ASP   204     -26.595  10.346 -35.446  1.00 41.79      B    N
ATOM   1080  CA  ASP   204     -26.440  11.788 -35.378  1.00 44.36      B    C
ATOM   1081  CB  ASP   204     -25.818  12.331 -36.671  1.00 49.04      B    C
ATOM   1082  CG  ASP   204     -26.486  11.799 -37.913  1.00 55.58      B    C
ATOM   1083  OD1 ASP   204     -27.729  11.633 -37.895  1.00 58.79      B    O
ATOM   1084  OD2 ASP   204     -25.761  11.549 -38.907  1.00 57.72      B    O
ATOM   1085  C   ASP   204     -27.752  12.495 -35.059  1.00 41.84      B    C
ATOM   1086  O   ASP   204     -27.864  13.706 -35.232  1.00 42.04      B    O
ATOM   1087  N   PHE   205     -28.737  11.745 -34.571  1.00 38.16      B    N
ATOM   1088  CA  PHE   205     -29.985  12.363 -34.147  1.00 36.96      B    C
ATOM   1089  CB  PHE   205     -31.130  11.351 -34.078  1.00 35.73      B    C
ATOM   1090  CG  PHE   205     -32.449  11.984 -33.744  1.00 36.34      B    C
ATOM   1091  CD1 PHE   205     -32.800  12.238 -32.428  1.00 34.81      B    C
ATOM   1092  CD2 PHE   205     -33.292  12.424 -34.750  1.00 36.67      B    C
ATOM   1093  CE1 PHE   205     -33.958  12.926 -32.122  1.00 34.86      B    C
ATOM   1094  CE2 PHE   205     -34.455  13.117 -34.450  1.00 36.51      B    C
ATOM   1095  CZ  PHE   205     -34.786  13.368 -33.136  1.00 36.57      B    C
ATOM   1096  C   PHE   205     -29.882  13.048 -32.787  1.00 37.17      B    C
ATOM   1097  O   PHE   205     -29.454  12.449 -31.799  1.00 35.61      B    O
ATOM   1098  N   GLU   206     -30.287  14.308 -32.741  1.00 36.47      B    N
ATOM   1099  CA  GLU   206     -30.548  14.937 -31.469  1.00 39.25      B    C
ATOM   1100  CB  GLU   206     -29.245  15.413 -30.821  1.00 43.55      B    C
ATOM   1101  CG  GLU   206     -28.695  16.714 -31.344  1.00 51.60      B    C
ATOM   1102  CD  GLU   206     -27.488  17.186 -30.544  1.00 57.35      B    C
ATOM   1103  OE1 GLU   206     -26.372  17.216 -31.118  1.00 58.84      B    O
ATOM   1104  OE2 GLU   206     -27.655  17.524 -29.344  1.00 56.01      B    O
ATOM   1105  C   GLU   206     -31.520  16.087 -31.617  1.00 37.64      B    C
ATOM   1106  O   GLU   206     -31.401  16.916 -32.526  1.00 36.38      B    O
ATOM   1107  N   ASN   207     -32.491  16.112 -30.711  1.00 34.24      B    N
ATOM   1108  CA  ASN   207     -33.505  17.155 -30.673  1.00 32.56      B    C
ATOM   1109  CB  ASN   207     -34.736  16.702 -31.466  1.00 30.25      B    C
ATOM   1110  CG  ASN   207     -35.636  17.855 -31.868  1.00 31.68      B    C
ATOM   1111  OD1 ASN   207     -35.748  18.189 -33.054  1.00 32.01      B    O
ATOM   1112  ND2 ASN   207     -36.291  18.465 -30.888  1.00 28.37      B    N
ATOM   1113  C   ASN   207     -33.864  17.364 -29.196  1.00 32.44      B    C
ATOM   1114  O   ASN   207     -34.658  16.611 -28.624  1.00 30.19      B    O
ATOM   1115  N   VAL   208     -33.257  18.374 -28.579  1.00 30.82      B    N
ATOM   1116  CA  VAL   208     -33.467  18.638 -27.163  1.00 30.29      B    C
ATOM   1117  CB  VAL   208     -32.274  18.127 -26.309  1.00 28.56      B    C
ATOM   1118  CG1 VAL   208     -32.051  16.638 -26.560  1.00 27.68      B    C
ATOM   1119  CG2 VAL   208     -31.016  18.917 -26.641  1.00 25.16      B    C
ATOM   1120  C   VAL   208     -33.628  20.134 -26.924  1.00 31.06      B    C
ATOM   1121  O   VAL   208     -33.063  20.956 -27.651  1.00 30.60      B    O
ATOM   1122  N   PRO   209     -34.408  20.503 -25.897  1.00 30.47      B    N
ATOM   1123  CD  PRO   209     -35.207  19.578 -25.072  1.00 30.10      B    C
ATOM   1124  CA  PRO   209     -34.536  21.894 -25.447  1.00 31.51      B    C
ATOM   1125  CB  PRO   209     -35.713  21.848 -24.474  1.00 31.36      B    C
ATOM   1126  CG  PRO   209     -35.683  20.448 -23.932  1.00 29.45      B    C
ATOM   1127  C   PRO   209     -33.256  22.396 -24.774  1.00 31.01      B    C
ATOM   1128  O   PRO   209     -32.478  21.612 -24.234  1.00 28.87      B    O
ATOM   1129  N   GLU   210     -33.045  23.704 -24.838  1.00 33.74      B    N
ATOM   1130  CA  GLU   210     -32.010  24.400 -24.068  1.00 36.85      B    C
ATOM   1131  CB  GLU   210     -32.237  25.908 -24.171  1.00 41.10      B    C
ATOM   1132  CG  GLU   210     -31.186  26.678 -24.925  1.00 52.07      B    C
ATOM   1133  CD  GLU   210     -31.540  28.153 -25.021  1.00 56.74      B    C
ATOM   1134  OE1 GLU   210     -32.743  28.464 -25.188  1.00 57.58      B    O
ATOM   1135  OE2 GLU   210     -30.622  29.000 -24.924  1.00 61.67      B    O
ATOM   1136  C   GLU   210     -32.079  24.017 -22.586  1.00 33.62      B    C
ATOM   1137  O   GLU   210     -33.165  23.878 -22.037  1.00 31.67      B    O
ATOM   1138  N   GLU   211     -30.927  23.878 -21.940  1.00 34.04      B    N
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ATOM	1139	CA	GLU	211	-30.877	23.780	-20.478	1.00	35.09	B	C
ATOM	1140	CB	GLU	211	-29.440	23.595	-19.993	1.00	37.07	B	C
ATOM	1141	CG	GLU	211	-28.520	22.943	-21.003	1.00	45.78	B	C
ATOM	1142	CD	GLU	211	-28.779	21.466	-21.132	1.00	49.35	B	C
ATOM	1143	OE1	GLU	211	-29.395	20.908	-20.197	1.00	53.87	B	O
ATOM	1144	OE2	GLU	211	-28.372	20.867	-22.156	1.00	49.08	B	O
ATOM	1145	C	GLU	211	-31.414	25.075	-19.887	1.00	34.57	B	C
ATOM	1146	O	GLU	211	-31.403	26.112	-20.552	1.00	33.61	B	O
ATOM	1147	N	ASP	212	-31.876	25.020	-18.640	1.00	34.88	B	N
ATOM	1148	CA	ASP	212	-32.155	26.237	-17.880	1.00	33.98	B	C
ATOM	1149	CB	ASP	212	-33.019	25.913	-16.654	1.00	36.73	B	C
ATOM	1150	CG	ASP	212	-33.263	27.131	-15.764	1.00	39.80	B	C
ATOM	1151	OD1	ASP	212	-32.845	28.252	-16.132	1.00	43.49	B	O
ATOM	1152	OD2	ASP	212	-33.873	26.965	-14.692	1.00	43.52	B	O
ATOM	1153	C	ASP	212	-30.828	26.858	-17.438	1.00	34.59	B	C
ATOM	1154	O	ASP	212	-30.185	26.385	-16.492	1.00	34.03	B	O
ATOM	1155	N	GLY	213	-30.425	27.920	-18.127	1.00	34.74	B	N
ATOM	1156	CA	GLY	213	-29.103	28.488	-17.926	1.00	35.64	B	C
ATOM	1157	C	GLY	213	-28.796	29.033	-16.535	1.00	37.75	B	C
ATOM	1158	O	GLY	213	-27.629	29.101	-16.151	1.00	38.98	B	O
ATOM	1159	N	THR	214	-29.818	29.422	-15.776	1.00	38.08	B	N
ATOM	1160	CA	THR	214	-29.594	29.989	-14.444	1.00	39.22	B	C
ATOM	1161	CB	THR	214	-30.863	30.667	-13.889	1.00	39.16	B	C
ATOM	1162	OG1	THR	214	-31.853	29.671	-13.609	1.00	42.43	B	O
ATOM	1163	CG2	THR	214	-31.427	31.653	-14.900	1.00	40.99	B	C
ATOM	1164	C	THR	214	-29.145	28.934	-13.429	1.00	38.48	B	C
ATOM	1165	O	THR	214	-28.695	29.273	-12.334	1.00	41.23	B	O
ATOM	1166	N	ARG	215	-29.278	27.661	-13.790	1.00	36.13	B	N
ATOM	1167	CA	ARG	215	-28.842	26.571	-12.926	1.00	33.09	B	C
ATOM	1168	CB	ARG	215	-30.044	25.753	-12.458	1.00	35.26	B	C
ATOM	1169	CG	ARG	215	-30.978	26.507	-11.525	1.00	39.25	B	C
ATOM	1170	CD	ARG	215	-31.976	25.562	-10.860	1.00	38.84	B	C
ATOM	1171	NE	ARG	215	-33.095	25.240	-11.741	1.00	41.79	B	N
ATOM	1172	CZ	ARG	215	-34.023	24.325	-11.469	1.00	42.83	B	C
ATOM	1173	NH1	ARG	215	-35.005	24.102	-12.330	1.00	42.89	B	N
ATOM	1174	NH2	ARG	215	-33.965	23.631	-10.341	1.00	40.76	B	N
ATOM	1175	C	ARG	215	-27.850	25.649	-13.621	1.00	31.31	B	C
ATOM	1176	O	ARG	215	-27.465	24.620	-13.072	1.00	30.45	B	O
ATOM	1177	N	PHE	216	-27.440	26.007	-14.831	1.00	29.98	B	N
ATOM	1178	CA	PHE	216	-26.577	25.126	-15.606	1.00	31.33	B	C
ATOM	1179	CB	PHE	216	-27.099	25.006	-17.041	1.00	30.11	B	C
ATOM	1180	CG	PHE	216	-26.373	23.979	-17.865	1.00	30.31	B	C
ATOM	1181	CD1	PHE	216	-25.816	24.317	-19.090	1.00	30.81	B	C
ATOM	1182	CD2	PHE	216	-26.258	22.671	-17.419	1.00	29.75	B	C
ATOM	1183	CE1	PHE	216	-25.156	23.363	-19.861	1.00	32.24	B	C
ATOM	1184	CE2	PHE	216	-25.602	21.712	-18.178	1.00	30.04	B	C
ATOM	1185	CZ	PHE	216	-25.049	22.056	-19.402	1.00	31.01	B	C
ATOM	1186	C	PHE	216	-25.122	25.604	-15.618	1.00	33.88	B	C
ATOM	1187	O	PHE	216	-24.824	26.718	-16.051	1.00	33.72	B	O
ATOM	1188	N	HIS	217	-24.226	24.750	-15.128	1.00	34.66	B	N
ATOM	1189	CA	HIS	217	-22.795	25.014	-15.165	1.00	36.02	B	C
ATOM	1190	CB	HIS	217	-22.209	25.037	-13.748	1.00	39.37	B	C
ATOM	1191	CG	HIS	217	-22.800	26.089	-12.860	1.00	44.56	B	C
ATOM	1192	CD2	HIS	217	-24.077	26.516	-12.707	1.00	46.84	B	C
ATOM	1193	ND1	HIS	217	-22.040	26.842	-11.990	1.00	47.01	B	N
ATOM	1194	CE1	HIS	217	-22.822	27.688	-11.341	1.00	46.30	B	C
ATOM	1195	NE2	HIS	217	-24.062	27.510	-11.757	1.00	48.13	B	N
ATOM	1196	C	HIS	217	-22.147	23.897	-15.962	1.00	36.30	B	C
ATOM	1197	O	HIS	217	-21.915	22.804	-15.445	1.00	35.57	B	O
ATOM	1198	N	ARG	218	-21.855	24.183	-17.223	1.00	37.33	B	N
ATOM	1199	CA	ARG	218	-21.393	23.173	-18.161	1.00	40.43	B	C
ATOM	1200	CB	ARG	218	-21.112	23.836	-19.514	1.00	43.99	B	C
ATOM	1201	CG	ARG	218	-20.457	22.930	-20.535	1.00	52.67	B	C
ATOM	1202	CD	ARG	218	-21.475	22.263	-21.444	1.00	60.20	B	C
ATOM	1203	NE	ARG	218	-21.024	20.935	-21.860	1.00	66.00	B	N
ATOM	1204	CZ	ARG	218	-21.834	19.961	-22.267	1.00	67.40	B	C
ATOM	1205	NH1	ARG	218	-21.333	18.783	-22.622	1.00	67.52	B	N
ATOM	1206	NH2	ARG	218	-23.145	20.168	-22.323	1.00	68.80	B	N
ATOM	1207	C	ARG	218	-20.150	22.423	-17.666	1.00	39.21	B	C
ATOM	1208	O	ARG	218	-20.039	21.212	-17.841	1.00	38.82	B	O
ATOM	1209	N	GLN	219	-19.220	23.136	-17.042	1.00	39.79	B	N
ATOM	1210	CA	GLN	219	-17.972	22.517	-16.608	1.00	41.38	B	C
ATOM	1211	CB	GLN	219	-16.984	23.584	-16.120	1.00	44.07	B	C
ATOM	1212	CG	GLN	219	-16.482	24.518	-17.227	1.00	53.87	B	C
ATOM	1213	CD	GLN	219	-14.958	24.512	-17.374	1.00	59.95	B	C
ATOM	1214	OE1	GLN	219	-14.421	24.216	-18.452	1.00	62.01	B	O

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ATOM   1215  NE2 GLN   219     -14.256  24.839 -16.289  1.00 62.25      B    N
ATOM   1216  C   GLN   219     -18.180  21.467 -15.519  1.00 37.67      B    C
ATOM   1217  O   GLN   219     -17.400  20.527 -15.406  1.00 37.10      B    O
ATOM   1218  N   ALA   220     -19.234  21.619 -14.725  1.00 35.89      B    N
ATOM   1219  CA  ALA   220     -19.503  20.680 -13.638  1.00 35.91      B    C
ATOM   1220  CB  ALA   220     -20.027  21.428 -12.427  1.00 33.00      B    C
ATOM   1221  C   ALA   220     -20.489  19.581 -14.037  1.00 35.70      B    C
ATOM   1222  O   ALA   220     -20.497  18.507 -13.453  1.00 36.94      B    O
ATOM   1223  N   SER   221     -21.312  19.856 -15.039  1.00 37.81      B    N
ATOM   1224  CA  SER   221     -22.373  18.941 -15.444  1.00 39.11      B    C
ATOM   1225  CB  SER   221     -23.316  19.659 -16.411  1.00 40.05      B    C
ATOM   1226  OG  SER   221     -24.304  18.777 -16.916  1.00 44.84      B    O
ATOM   1227  C   SER   221     -21.854  17.652 -16.087  1.00 38.84      B    C
ATOM   1228  O   SER   221     -20.932  17.674 -16.900  1.00 38.31      B    O
ATOM   1229  N   LYS   222     -22.449  16.525 -15.715  1.00 39.04      B    N
ATOM   1230  CA  LYS   222     -22.186  15.259 -16.396  1.00 41.16      B    C
ATOM   1231  CB  LYS   222     -22.026  14.127 -15.377  1.00 42.57      B    C
ATOM   1232  CG  LYS   222     -20.878  14.311 -14.402  1.00 46.43      B    C
ATOM   1233  CD  LYS   222     -21.183  13.589 -13.100  1.00 53.68      B    C
ATOM   1234  CE  LYS   222     -19.918  13.140 -12.377  1.00 57.05      B    C
ATOM   1235  NZ  LYS   222     -20.212  12.021 -11.418  1.00 57.68      B    N
ATOM   1236  C   LYS   222     -23.353  14.939 -17.321  1.00 41.01      B    C
ATOM   1237  O   LYS   222     -24.327  14.316 -16.904  1.00 42.43      B    O
ATOM   1238  N   CYS   223     -23.260  15.361 -18.576  1.00 41.97      B    N
ATOM   1239  CA  CYS   223     -24.396  15.259 -19.484  1.00 44.73      B    C
ATOM   1240  C   CYS   223     -24.721  13.819 -19.876  1.00 42.78      B    C
ATOM   1241  O   CYS   223     -25.844  13.528 -20.280  1.00 42.75      B    O
ATOM   1242  CB  CYS   223     -24.148  16.104 -20.745  1.00 48.24      B    C
ATOM   1243  SG  CYS   223     -24.066  17.911 -20.440  1.00 61.09      B    S
ATOM   1244  N   ASP   224     -23.736  12.931 -19.735  1.00 42.89      B    N
ATOM   1245  CA  ASP   224     -23.820  11.549 -20.215  1.00 41.94      B    C
ATOM   1246  CB  ASP   224     -22.448  11.060 -20.677  1.00 45.97      B    C
ATOM   1247  CG  ASP   224     -21.896  11.860 -21.828  1.00 51.61      B    C
ATOM   1248  OD1 ASP   224     -22.698  12.450 -22.585  1.00 55.30      B    O
ATOM   1249  OD2 ASP   224     -20.653  11.896 -21.978  1.00 55.41      B    O
ATOM   1250  C   ASP   224     -24.318  10.570 -19.163  1.00 40.44      B    C
ATOM   1251  O   ASP   224     -24.500   9.391 -19.446  1.00 40.53      B    O
ATOM   1252  N   SER   225     -24.514  11.043 -17.943  1.00 39.18      B    N
ATOM   1253  CA  SER   225     -24.670  10.136 -16.817  1.00 38.21      B    C
ATOM   1254  CB  SER   225     -24.814  10.936 -15.527  1.00 38.32      B    C
ATOM   1255  OG  SER   225     -25.257  10.088 -14.488  1.00 45.34      B    O
ATOM   1256  C   SER   225     -25.838   9.155 -16.939  1.00 35.49      B    C
ATOM   1257  O   SER   225     -25.671   7.957 -16.727  1.00 35.90      B    O
ATOM   1258  N   HIS   226     -27.018   9.664 -17.275  1.00 34.83      B    N
ATOM   1259  CA  HIS   226     -28.235   8.852 -17.285  1.00 34.19      B    C
ATOM   1260  CB  HIS   226     -29.458   9.771 -17.415  1.00 33.13      B    C
ATOM   1261  CG  HIS   226     -30.774   9.083 -17.207  1.00 34.07      B    C
ATOM   1262  CD2 HIS   226     -31.387   8.650 -16.078  1.00 33.89      B    C
ATOM   1263  ND1 HIS   226     -31.659   8.841 -18.238  1.00 33.86      B    N
ATOM   1264  CE1 HIS   226     -32.760   8.295 -17.753  1.00 34.25      B    C
ATOM   1265  NE2 HIS   226     -32.621   8.168 -16.445  1.00 35.48      B    N
ATOM   1266  C   HIS   226     -28.206   7.844 -18.440  1.00 34.61      B    C
ATOM   1267  O   HIS   226     -28.456   6.656 -18.248  1.00 34.03      B    O
ATOM   1268  N   GLY   227     -27.887   8.321 -19.639  1.00 33.03      B    N
ATOM   1269  CA  GLY   227     -27.905   7.449 -20.795  1.00 33.67      B    C
ATOM   1270  C   GLY   227     -26.827   6.378 -20.768  1.00 33.99      B    C
ATOM   1271  O   GLY   227     -27.082   5.231 -21.140  1.00 32.66      B    O
ATOM   1272  N   THR   228     -25.623   6.742 -20.332  1.00 33.72      B    N
ATOM   1273  CA  THR   228     -24.514   5.793 -20.305  1.00 35.16      B    C
ATOM   1274  CB  THR   228     -23.218   6.436 -19.759  1.00 36.04      B    C
ATOM   1275  OG1 THR   228     -22.857   7.564 -20.564  1.00 36.59      B    O
ATOM   1276  CG2 THR   228     -22.085   5.427 -19.778  1.00 35.96      B    C
ATOM   1277  C   THR   228     -24.866   4.620 -19.399  1.00 35.86      B    C
ATOM   1278  O   THR   228     -24.596   3.459 -19.716  1.00 34.91      B    O
ATOM   1279  N   HIS   229     -25.473   4.933 -18.262  1.00 36.05      B    N
ATOM   1280  CA  HIS   229     -25.755   3.920 -17.264  1.00 35.40      B    C
ATOM   1281  CB  HIS   229     -26.279   4.582 -15.990  1.00 35.21      B    C
ATOM   1282  CG  HIS   229     -26.547   3.619 -14.879  1.00 34.42      B    C
ATOM   1283  CD2 HIS   229     -25.754   3.166 -13.879  1.00 35.18      B    C
ATOM   1284  ND1 HIS   229     -27.758   2.981 -14.726  1.00 36.17      B    N
ATOM   1285  CE1 HIS   229     -27.700   2.173 -13.681  1.00 38.58      B    C
ATOM   1286  NE2 HIS   229     -26.494   2.267 -13.150  1.00 37.21      B    N
ATOM   1287  C   HIS   229     -26.787   2.947 -17.817  1.00 36.81      B    C
ATOM   1288  O   HIS   229     -26.699   1.736 -17.603  1.00 37.62      B    O
ATOM   1289  N   LEU   230     -27.764   3.486 -18.540  1.00 37.41      B    N
ATOM   1290  CA  LEU   230     -28.840   2.674 -19.092  1.00 36.89      B    C
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ATOM	1291	CB	LEU	230	-29.971	3.580	-19.591	1.00	35.14	B	C
ATOM	1292	CG	LEU	230	-30.761	4.295	-18.489	1.00	36.53	B	C
ATOM	1293	CD1	LEU	230	-31.805	5.216	-19.118	1.00	34.66	B	C
ATOM	1294	CD2	LEU	230	-31.430	3.263	-17.585	1.00	31.80	B	C
ATOM	1295	C	LEU	230	-28.345	1.773	-20.224	1.00	37.42	B	C
ATOM	1296	O	LEU	230	-28.719	0.602	-20.300	1.00	37.18	B	O
ATOM	1297	N	ALA	231	-27.505	2.317	-21.099	1.00	36.72	B	N
ATOM	1298	CA	ALA	231	-26.895	1.514	-22.154	1.00	37.67	B	C
ATOM	1299	CB	ALA	231	-25.934	2.365	-22.967	1.00	38.58	B	C
ATOM	1300	C	ALA	231	-26.145	0.353	-21.506	1.00	38.18	B	C
ATOM	1301	O	ALA	231	-26.195	-0.781	-21.985	1.00	38.02	B	O
ATOM	1302	N	GLY	232	-25.466	0.650	-20.403	1.00	37.63	B	N
ATOM	1303	CA	GLY	232	-24.740	-0.371	-19.674	1.00	37.42	B	C
ATOM	1304	C	GLY	232	-25.634	-1.433	-19.062	1.00	39.21	B	C
ATOM	1305	O	GLY	232	-25.294	-2.620	-19.096	1.00	39.54	B	O
ATOM	1306	N	VAL	233	-26.774	-1.025	-18.504	1.00	37.80	B	N
ATOM	1307	CA	VAL	233	-27.706	-1.991	-17.932	1.00	36.04	B	C
ATOM	1308	CB	VAL	233	-28.907	-1.305	-17.247	1.00	34.34	B	C
ATOM	1309	CG1	VAL	233	-29.925	-2.348	-16.842	1.00	32.45	B	C
ATOM	1310	CG2	VAL	233	-28.446	-0.528	-16.024	1.00	32.92	B	C
ATOM	1311	C	VAL	233	-28.246	-2.917	-19.014	1.00	37.82	B	C
ATOM	1312	O	VAL	233	-28.417	-4.113	-18.788	1.00	38.19	B	O
ATOM	1313	N	VAL	234	-28.514	-2.363	-20.191	1.00	38.91	B	N
ATOM	1314	CA	VAL	234	-29.036	-3.164	-21.291	1.00	39.62	B	C
ATOM	1315	CB	VAL	234	-29.552	-2.271	-22.453	1.00	39.19	B	C
ATOM	1316	CG1	VAL	234	-29.920	-3.138	-23.640	1.00	37.07	B	C
ATOM	1317	CG2	VAL	234	-30.775	-1.470	-22.005	1.00	38.06	B	C
ATOM	1318	C	VAL	234	-27.980	-4.127	-21.849	1.00	41.00	B	C
ATOM	1319	O	VAL	234	-28.241	-5.323	-22.000	1.00	38.81	B	O
ATOM	1320	N	SER	235	-26.788	-3.609	-22.141	1.00	41.61	B	N
ATOM	1321	CA	SER	235	-25.837	-4.343	-22.969	1.00	43.40	B	C
ATOM	1322	CB	SER	235	-25.898	-3.804	-24.395	1.00	43.13	B	C
ATOM	1323	OG	SER	235	-25.378	-2.486	-24.430	1.00	43.09	B	O
ATOM	1324	C	SER	235	-24.384	-4.320	-22.493	1.00	44.31	B	C
ATOM	1325	O	SER	235	-23.488	-4.670	-23.256	1.00	44.63	B	O
ATOM	1326	N	GLY	236	-24.141	-3.903	-21.255	1.00	44.68	B	N
ATOM	1327	CA	GLY	236	-22.771	-3.852	-20.768	1.00	47.17	B	C
ATOM	1328	C	GLY	236	-22.139	-5.234	-20.731	1.00	48.76	B	C
ATOM	1329	O	GLY	236	-22.823	-6.220	-20.447	1.00	47.27	B	O
ATOM	1330	N	ARG	237	-20.840	-5.327	-21.011	1.00	50.91	B	N
ATOM	1331	CA	ARG	237	-20.216	-6.644	-21.113	1.00	54.98	B	C
ATOM	1332	CB	ARG	237	-18.866	-6.565	-21.839	1.00	57.32	B	C
ATOM	1333	CG	ARG	237	-17.921	-5.481	-21.369	1.00	62.69	B	C
ATOM	1334	CD	ARG	237	-16.673	-5.455	-22.254	1.00	67.42	B	C
ATOM	1335	NE	ARG	237	-16.997	-5.685	-23.664	1.00	72.57	B	N
ATOM	1336	CZ	ARG	237	-16.678	-4.861	-24.663	1.00	74.70	B	C
ATOM	1337	NH1	ARG	237	-17.021	-5.164	-25.911	1.00	74.47	B	N
ATOM	1338	NH2	ARG	237	-16.013	-3.737	-24.420	1.00	75.30	B	N
ATOM	1339	C	ARG	237	-20.047	-7.351	-19.772	1.00	54.54	B	C
ATOM	1340	O	ARG	237	-20.056	-8.578	-19.716	1.00	56.23	B	O
ATOM	1341	N	ASP	238	-19.911	-6.584	-18.694	1.00	54.25	B	N
ATOM	1342	CA	ASP	238	-19.780	-7.167	-17.362	1.00	54.09	B	C
ATOM	1343	CB	ASP	238	-18.663	-6.463	-16.580	1.00	56.58	B	C
ATOM	1344	CG	ASP	238	-17.288	-6.673	-17.199	1.00	59.27	B	C
ATOM	1345	OD1	ASP	238	-17.167	-7.481	-18.144	1.00	60.55	B	O
ATOM	1346	OD2	ASP	238	-16.323	-6.026	-16.738	1.00	61.66	B	O
ATOM	1347	C	ASP	238	-21.077	-7.101	-16.556	1.00	53.15	B	C
ATOM	1348	O	ASP	238	-21.436	-8.060	-15.875	1.00	52.85	B	O
ATOM	1349	N	ALA	239	-21.775	-5.970	-16.627	1.00	52.70	B	N
ATOM	1350	CA	ALA	239	-22.913	-5.727	-15.742	1.00	50.29	B	C
ATOM	1351	CB	ALA	239	-22.663	-4.476	-14.915	1.00	49.27	B	C
ATOM	1352	C	ALA	239	-24.243	-5.607	-16.483	1.00	49.76	B	C
ATOM	1353	O	ALA	239	-25.267	-5.280	-15.881	1.00	48.97	B	O
ATOM	1354	N	GLY	240	-24.229	-5.881	-17.783	1.00	48.12	B	N
ATOM	1355	CA	GLY	240	-25.448	-5.783	-18.566	1.00	47.47	B	C
ATOM	1356	C	GLY	240	-26.327	-7.023	-18.552	1.00	47.44	B	C
ATOM	1357	O	GLY	240	-25.912	-8.098	-18.113	1.00	48.79	B	O
ATOM	1358	N	VAL	241	-27.553	-6.872	-19.042	1.00	46.01	B	N
ATOM	1359	CA	VAL	241	-28.499	-7.977	-19.115	1.00	45.03	B	C
ATOM	1360	CB	VAL	241	-29.949	-7.465	-19.065	1.00	43.05	B	C
ATOM	1361	CG1	VAL	241	-30.907	-8.578	-19.431	1.00	39.38	B	C
ATOM	1362	CG2	VAL	241	-30.261	-6.939	-17.668	1.00	42.38	B	C
ATOM	1363	C	VAL	241	-28.307	-8.797	-20.388	1.00	47.14	B	C
ATOM	1364	O	VAL	241	-28.223	-10.025	-20.334	1.00	47.85	B	O
ATOM	1365	N	ALA	242	-28.243	-8.114	-21.528	1.00	46.60	B	N
ATOM	1366	CA	ALA	242	-27.917	-8.759	-22.797	1.00	48.43	B	C

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ATOM   1367  CB  ALA  242     -28.913  -8.342 -23.873  1.00 45.85      B    C
ATOM   1368  C   ALA  242     -26.504  -8.350 -23.202  1.00 50.23      B    C
ATOM   1369  O   ALA  242     -26.312  -7.414 -23.983  1.00 50.74      B    O
ATOM   1370  N   LYS  243     -25.517  -9.058 -22.663  1.00 51.90      B    N
ATOM   1371  CA  LYS  243     -24.134  -8.615 -22.748  1.00 53.47      B    C
ATOM   1372  CB  LYS  243     -23.239  -9.546 -21.923  1.00 54.37      B    C
ATOM   1373  CG  LYS  243     -23.605  -9.581 -20.437  1.00 56.48      B    C
ATOM   1374  CD  LYS  243     -22.714 -10.528 -19.649  1.00 57.08      B    C
ATOM   1375  CE  LYS  243     -23.346 -10.907 -18.314  1.00 58.87      B    C
ATOM   1376  NZ  LYS  243     -23.227  -9.834 -17.291  1.00 59.43      B    N
ATOM   1377  C   LYS  243     -23.658  -8.552 -24.196  1.00 53.57      B    C
ATOM   1378  O   LYS  243     -23.857  -9.491 -24.972  1.00 53.48      B    O
ATOM   1379  N   GLY  244     -23.047  -7.425 -24.555  1.00 53.12      B    N
ATOM   1380  CA  GLY  244     -22.519  -7.255 -25.896  1.00 51.90      B    C
ATOM   1381  C   GLY  244     -23.555  -6.909 -26.952  1.00 52.11      B    C
ATOM   1382  O   GLY  244     -23.209  -6.729 -28.121  1.00 51.13      B    O
ATOM   1383  N   ALA  245     -24.823  -6.810 -26.557  1.00 51.54      B    N
ATOM   1384  CA  ALA  245     -25.892  -6.538 -27.517  1.00 49.79      B    C
ATOM   1385  CB  ALA  245     -27.240  -6.513 -26.814  1.00 49.57      B    C
ATOM   1386  C   ALA  245     -25.663  -5.221 -28.248  1.00 48.94      B    C
ATOM   1387  O   ALA  245     -25.013  -4.308 -27.732  1.00 50.04      B    O
ATOM   1388  N   SER  246     -26.206  -5.135 -29.455  1.00 47.10      B    N
ATOM   1389  CA  SER  246     -25.985  -3.992 -30.326  1.00 46.40      B    C
ATOM   1390  CB  SER  246     -26.152  -4.432 -31.785  1.00 46.20      B    C
ATOM   1391  OG  SER  246     -26.115  -3.332 -32.676  1.00 49.27      B    O
ATOM   1392  C   SER  246     -26.971  -2.866 -29.992  1.00 46.30      B    C
ATOM   1393  O   SER  246     -28.166  -3.109 -29.813  1.00 46.52      B    O
ATOM   1394  N   MET  247     -26.475  -1.635 -29.909  1.00 44.47      B    N
ATOM   1395  CA  MET  247     -27.341  -0.506 -29.591  1.00 43.77      B    C
ATOM   1396  CB  MET  247     -27.184  -0.116 -28.120  1.00 44.42      B    C
ATOM   1397  CG  MET  247     -27.610  -1.193 -27.145  1.00 47.82      B    C
ATOM   1398  SD  MET  247     -27.687  -0.576 -25.455  1.00 51.27      B    S
ATOM   1399  CE  MET  247     -29.166   0.438 -25.535  1.00 48.79      B    C
ATOM   1400  C   MET  247     -27.092   0.718 -30.464  1.00 42.00      B    C
ATOM   1401  O   MET  247     -25.972   0.962 -30.916  1.00 39.97      B    O
ATOM   1402  N   ARG  248     -28.155   1.480 -30.694  1.00 39.53      B    N
ATOM   1403  CA  ARG  248     -28.057   2.770 -31.359  1.00 39.90      B    C
ATOM   1404  CB  ARG  248     -28.716   2.699 -32.739  1.00 42.19      B    C
ATOM   1405  CG  ARG  248     -28.111   1.636 -33.646  1.00 47.22      B    C
ATOM   1406  CD  ARG  248     -28.800   1.575 -35.001  1.00 52.45      B    C
ATOM   1407  NE  ARG  248     -28.003   2.220 -36.040  1.00 59.77      B    N
ATOM   1408  CZ  ARG  248     -28.404   3.269 -36.754  1.00 64.16      B    C
ATOM   1409  NH1 ARG  248     -27.599   3.789 -37.674  1.00 65.74      B    N
ATOM   1410  NH2 ARG  248     -29.609   3.798 -36.560  1.00 65.15      B    N
ATOM   1411  C   ARG  248     -28.751   3.816 -30.492  1.00 39.62      B    C
ATOM   1412  O   ARG  248     -29.907   3.641 -30.096  1.00 39.17      B    O
ATOM   1413  N   SER  249     -28.047   4.901 -30.185  1.00 37.99      B    N
ATOM   1414  CA  SER  249     -28.588   5.898 -29.280  1.00 37.24      B    C
ATOM   1415  CB  SER  249     -27.549   6.265 -28.214  1.00 36.64      B    C
ATOM   1416  OG  SER  249     -26.336   6.689 -28.805  1.00 42.82      B    O
ATOM   1417  C   SER  249     -29.048   7.147 -30.020  1.00 36.18      B    C
ATOM   1418  O   SER  249     -28.395   7.608 -30.960  1.00 36.34      B    O
ATOM   1419  N   LEU  250     -30.192   7.674 -29.595  1.00 33.93      B    N
ATOM   1420  CA  LEU  250     -30.690   8.965 -30.062  1.00 33.58      B    C
ATOM   1421  CB  LEU  250     -32.085   8.810 -30.683  1.00 31.97      B    C
ATOM   1422  CG  LEU  250     -32.225   8.090 -32.028  1.00 35.15      B    C
ATOM   1423  CD1 LEU  250     -31.875   6.607 -31.885  1.00 32.63      B    C
ATOM   1424  CD2 LEU  250     -33.649   8.262 -32.537  1.00 29.57      B    C
ATOM   1425  C   LEU  250     -30.784   9.893 -28.853  1.00 32.36      B    C
ATOM   1426  O   LEU  250     -31.214   9.472 -27.778  1.00 32.19      B    O
ATOM   1427  N   ARG  251     -30.403  11.153 -29.024  1.00 31.24      B    N
ATOM   1428  CA  ARG  251     -30.497  12.105 -27.924  1.00 31.28      B    C
ATOM   1429  CB  ARG  251     -29.299  13.049 -27.937  1.00 32.49      B    C
ATOM   1430  CG  ARG  251     -29.320  14.019 -26.778  1.00 34.71      B    C
ATOM   1431  CD  ARG  251     -28.075  14.868 -26.739  1.00 35.70      B    C
ATOM   1432  NE  ARG  251     -27.962  15.534 -25.447  1.00 39.37      B    N
ATOM   1433  CZ  ARG  251     -27.585  16.796 -25.294  1.00 38.08      B    C
ATOM   1434  NH1 ARG  251     -27.282  17.528 -26.356  1.00 38.81      B    N
ATOM   1435  NH2 ARG  251     -27.518  17.322 -24.082  1.00 38.72      B    N
ATOM   1436  C   ARG  251     -31.785  12.929 -27.951  1.00 30.86      B    C
ATOM   1437  O   ARG  251     -31.947  13.813 -28.800  1.00 31.68      B    O
ATOM   1438  N   VAL  252     -32.697  12.643 -27.021  1.00 28.82      B    N
ATOM   1439  CA  VAL  252     -33.927  13.423 -26.892  1.00 28.33      B    C
ATOM   1440  CB  VAL  252     -35.187  12.564 -27.173  1.00 28.02      B    C
ATOM   1441  CG1 VAL  252     -35.122  12.000 -28.589  1.00 27.63      B    C
ATOM   1442  CG2 VAL  252     -35.295  11.437 -26.162  1.00 25.95      B    C
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ATOM   1443  C   VAL   252     -34.084  14.077 -25.521  1.00 27.82      B    C
ATOM   1444  O   VAL   252     -35.091  14.727 -25.259  1.00 28.30      B    O
ATOM   1445  N   LEU   253     -33.092  13.897 -24.652  1.00 26.78      B    N
ATOM   1446  CA  LEU   253     -33.075  14.563 -23.349  1.00 27.02      B    C
ATOM   1447  CB  LEU   253     -33.128  13.527 -22.222  1.00 23.74      B    C
ATOM   1448  CG  LEU   253     -34.374  12.643 -22.136  1.00 24.22      B    C
ATOM   1449  CD1 LEU   253     -34.152  11.582 -21.061  1.00 23.90      B    C
ATOM   1450  CD2 LEU   253     -35.602  13.489 -21.807  1.00 21.35      B    C
ATOM   1451  C   LEU   253     -31.798  15.387 -23.223  1.00 25.77      B    C
ATOM   1452  O   LEU   253     -30.723  14.910 -23.570  1.00 26.91      B    O
ATOM   1453  N   ASN   254     -31.908  16.619 -22.732  1.00 26.13      B    N
ATOM   1454  CA  ASN   254     -30.723  17.458 -22.575  1.00 26.64      B    C
ATOM   1455  CB  ASN   254     -31.111  18.945 -22.552  1.00 24.80      B    C
ATOM   1456  CG  ASN   254     -31.999  19.309 -21.380  1.00 25.34      B    C
ATOM   1457  OD1 ASN   254     -32.052  18.598 -20.374  1.00 22.46      B    O
ATOM   1458  ND2 ASN   254     -32.704  20.430 -21.505  1.00 22.21      B    N
ATOM   1459  C   ASN   254     -29.940  17.088 -21.314  1.00 27.37      B    C
ATOM   1460  O   ASN   254     -30.229  16.079 -20.670  1.00 25.26      B    O
ATOM   1461  N   CYS   255     -28.941  17.898 -20.974  1.00 28.60      B    N
ATOM   1462  CA  CYS   255     -28.069  17.609 -19.837  1.00 31.55      B    C
ATOM   1463  C   CYS   255     -28.845  17.611 -18.526  1.00 29.84      B    C
ATOM   1464  O   CYS   255     -28.449  16.956 -17.561  1.00 29.79      B    O
ATOM   1465  CB  CYS   255     -26.944  18.643 -19.756  1.00 39.06      B    C
ATOM   1466  SG  CYS   255     -25.816  18.660 -21.190  1.00 50.83      B    S
ATOM   1467  N   GLN   256     -29.946  18.350 -18.499  1.00 25.96      B    N
ATOM   1468  CA  GLN   256     -30.795  18.413 -17.319  1.00 25.56      B    C
ATOM   1469  CB  GLN   256     -31.304  19.855 -17.118  1.00 26.50      B    C
ATOM   1470  CG  GLN   256     -30.203  20.805 -16.621  1.00 27.07      B    C
ATOM   1471  CD  GLN   256     -30.618  22.269 -16.546  1.00 29.99      B    C
ATOM   1472  OE1 GLN   256     -31.569  22.698 -17.202  1.00 29.18      B    O
ATOM   1473  NE2 GLN   256     -29.889  23.051 -15.740  1.00 28.68      B    N
ATOM   1474  C   GLN   256     -31.955  17.422 -17.400  1.00 25.12      B    C
ATOM   1475  O   GLN   256     -32.960  17.577 -16.714  1.00 25.75      B    O
ATOM   1476  N   GLY   257     -31.806  16.393 -18.231  1.00 25.10      B    N
ATOM   1477  CA  GLY   257     -32.761  15.291 -18.221  1.00 24.84      B    C
ATOM   1478  C   GLY   257     -34.124  15.640 -18.805  1.00 26.13      B    C
ATOM   1479  O   GLY   257     -35.115  14.940 -18.570  1.00 24.93      B    O
ATOM   1480  N   LYS   258     -34.183  16.722 -19.571  1.00 24.07      B    N
ATOM   1481  CA  LYS   258     -35.462  17.231 -20.040  1.00 28.15      B    C
ATOM   1482  CB  LYS   258     -35.657  18.660 -19.522  1.00 32.37      B    C
ATOM   1483  CG  LYS   258     -37.070  19.183 -19.644  1.00 39.14      B    C
ATOM   1484  CD  LYS   258     -37.359  20.253 -18.588  1.00 41.81      B    C
ATOM   1485  CE  LYS   258     -37.363  21.648 -19.204  1.00 45.11      B    C
ATOM   1486  NZ  LYS   258     -38.425  22.526 -18.618  1.00 47.02      B    N
ATOM   1487  C   LYS   258     -35.557  17.191 -21.568  1.00 27.03      B    C
ATOM   1488  O   LYS   258     -34.595  17.513 -22.274  1.00 25.30      B    O
ATOM   1489  N   GLY   259     -36.715  16.771 -22.072  1.00 26.32      B    N
ATOM   1490  CA  GLY   259     -36.963  16.800 -23.507  1.00 25.87      B    C
ATOM   1491  C   GLY   259     -38.363  17.309 -23.804  1.00 26.15      B    C
ATOM   1492  O   GLY   259     -39.028  17.847 -22.920  1.00 25.50      B    O
ATOM   1493  N   THR   260     -38.822  17.143 -25.041  1.00 25.47      B    N
ATOM   1494  CA  THR   260     -40.191  17.523 -25.394  1.00 25.25      B    C
ATOM   1495  CB  THR   260     -40.210  18.742 -26.335  1.00 25.76      B    C
ATOM   1496  OG1 THR   260     -39.565  18.402 -27.572  1.00 24.26      B    O
ATOM   1497  CG2 THR   260     -39.479  19.921 -25.689  1.00 22.58      B    C
ATOM   1498  C   THR   260     -40.920  16.377 -26.082  1.00 24.53      B    C
ATOM   1499  O   THR   260     -40.289  15.486 -26.647  1.00 25.79      B    O
ATOM   1500  N   VAL   261     -42.248  16.393 -26.027  1.00 25.73      B    N
ATOM   1501  CA  VAL   261     -43.038  15.432 -26.789  1.00 25.75      B    C
ATOM   1502  CB  VAL   261     -44.558  15.676 -26.619  1.00 25.68      B    C
ATOM   1503  CG1 VAL   261     -45.328  14.955 -27.717  1.00 23.94      B    C
ATOM   1504  CG2 VAL   261     -45.019  15.164 -25.262  1.00 24.05      B    C
ATOM   1505  C   VAL   261     -42.682  15.543 -28.271  1.00 27.72      B    C
ATOM   1506  O   VAL   261     -42.517  14.534 -28.953  1.00 28.50      B    O
ATOM   1507  N   SER   262     -42.542  16.768 -28.766  1.00 26.86      B    N
ATOM   1508  CA  SER   262     -42.305  16.961 -30.193  1.00 28.93      B    C
ATOM   1509  CB  SER   262     -42.371  18.457 -30.557  1.00 28.78      B    C
ATOM   1510  OG  SER   262     -41.371  19.209 -29.887  1.00 30.58      B    O
ATOM   1511  C   SER   262     -40.955  16.370 -30.599  1.00 29.25      B    C
ATOM   1512  O   SER   262     -40.839  15.735 -31.655  1.00 30.56      B    O
ATOM   1513  N   GLY   263     -39.943  16.561 -29.756  1.00 27.34      B    N
ATOM   1514  CA  GLY   263     -38.631  16.013 -30.051  1.00 25.52      B    C
ATOM   1515  C   GLY   263     -38.647  14.496 -30.038  1.00 27.31      B    C
ATOM   1516  O   GLY   263     -37.978  13.841 -30.836  1.00 27.93      B    O
ATOM   1517  N   THR   264     -39.429  13.930 -29.130  1.00 27.25      B    N
ATOM   1518  CA  THR   264     -39.541  12.487 -29.020  1.00 27.72      B    C
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ATOM   1519  CB  THR   264     -40.297  12.112 -27.718  1.00 30.16      B    C
ATOM   1520  OG1 THR   264     -39.571  12.628 -26.588  1.00 30.13      B    O
ATOM   1521  CG2 THR   264     -40.431  10.597 -27.581  1.00 29.89      B    C
ATOM   1522  C   THR   264     -40.250  11.899 -30.253  1.00 28.07      B    C
ATOM   1523  O   THR   264     -39.831  10.866 -30.775  1.00 28.06      B    O
ATOM   1524  N   LEU   265     -41.305  12.562 -30.724  1.00 26.93      B    N
ATOM   1525  CA  LEU   265     -41.970  12.162 -31.962  1.00 29.22      B    C
ATOM   1526  CB  LEU   265     -43.076  13.152 -32.324  1.00 32.15      B    C
ATOM   1527  CG  LEU   265     -44.282  13.225 -31.394  1.00 36.81      B    C
ATOM   1528  CD1 LEU   265     -45.239  14.303 -31.906  1.00 37.93      B    C
ATOM   1529  CD2 LEU   265     -44.970  11.868 -31.331  1.00 35.61      B    C
ATOM   1530  C   LEU   265     -40.984  12.092 -33.126  1.00 28.75      B    C
ATOM   1531  O   LEU   265     -40.900  11.089 -33.824  1.00 30.75      B    O
ATOM   1532  N   ILE   266     -40.238  13.168 -33.325  1.00 28.71      B    N
ATOM   1533  CA  ILE   266     -39.268  13.236 -34.400  1.00 28.41      B    C
ATOM   1534  CB  ILE   266     -38.591  14.620 -34.408  1.00 27.07      B    C
ATOM   1535  CG2 ILE   266     -37.539  14.694 -35.506  1.00 24.79      B    C
ATOM   1536  CG1 ILE   266     -39.657  15.701 -34.618  1.00 28.34      B    C
ATOM   1537  CD1 ILE   266     -39.111  17.116 -34.626  1.00 27.25      B    C
ATOM   1538  C   ILE   266     -38.219  12.128 -34.261  1.00 30.28      B    C
ATOM   1539  O   ILE   266     -37.689  11.636 -35.252  1.00 31.29      B    O
ATOM   1540  N   GLY   267     -37.928  11.726 -33.030  1.00 31.04      B    N
ATOM   1541  CA  GLY   267     -36.995  10.633 -32.829  1.00 30.24      B    C
ATOM   1542  C   GLY   267     -37.617   9.293 -33.185  1.00 31.88      B    C
ATOM   1543  O   GLY   267     -36.952   8.420 -33.746  1.00 30.74      B    O
ATOM   1544  N   LEU   268     -38.897   9.124 -32.865  1.00 31.46      B    N
ATOM   1545  CA  LEU   268     -39.597   7.900 -33.224  1.00 33.36      B    C
ATOM   1546  CB  LEU   268     -40.980   7.857 -32.566  1.00 29.79      B    C
ATOM   1547  CG  LEU   268     -40.963   7.777 -31.039  1.00 30.96      B    C
ATOM   1548  CD1 LEU   268     -42.378   7.836 -30.496  1.00 28.19      B    C
ATOM   1549  CD2 LEU   268     -40.279   6.495 -30.593  1.00 30.02      B    C
ATOM   1550  C   LEU   268     -39.732   7.805 -34.744  1.00 34.79      B    C
ATOM   1551  O   LEU   268     -39.605   6.726 -35.316  1.00 38.35      B    O
ATOM   1552  N   GLU   269     -39.970   8.934 -35.399  1.00 35.31      B    N
ATOM   1553  CA  GLU   269     -40.034   8.948 -36.857  1.00 37.68      B    C
ATOM   1554  CB  GLU   269     -40.449  10.332 -37.356  1.00 38.65      B    C
ATOM   1555  CG  GLU   269     -40.399  10.460 -38.862  1.00 45.31      B    C
ATOM   1556  CD  GLU   269     -40.836  11.789 -39.363  1.00 49.12      B    C
ATOM   1557  OE1 GLU   269     -40.121  12.707 -39.599  1.00 50.11      B    O
ATOM   1558  OE2 GLU   269     -42.171  11.908 -39.523  1.00 51.18      B    O
ATOM   1559  C   GLU   269     -38.687   8.555 -37.479  1.00 36.64      B    C
ATOM   1560  O   GLU   269     -38.640   7.837 -38.473  1.00 35.50      B    O
ATOM   1561  N   PHE   270     -37.599   9.029 -36.882  1.00 35.93      B    N
ATOM   1562  CA  PHE   270     -36.251   8.670 -37.310  1.00 35.58      B    C
ATOM   1563  CB  PHE   270     -35.221   9.347 -36.399  1.00 35.25      B    C
ATOM   1564  CG  PHE   270     -33.793   8.989 -36.712  1.00 33.73      B    C
ATOM   1565  CD1 PHE   270     -33.013   9.819 -37.505  1.00 35.52      B    C
ATOM   1566  CD2 PHE   270     -33.219   7.841 -36.190  1.00 34.10      B    C
ATOM   1567  CE1 PHE   270     -31.679   9.511 -37.770  1.00 35.84      B    C
ATOM   1568  CE2 PHE   270     -31.890   7.527 -36.451  1.00 36.68      B    C
ATOM   1569  CZ  PHE   270     -31.119   8.365 -37.243  1.00 34.03      B    C
ATOM   1570  C   PHE   270     -36.050   7.159 -37.272  1.00 35.72      B    C
ATOM   1571  O   PHE   270     -35.440   6.587 -38.167  1.00 34.84      B    O
ATOM   1572  N   ILE   271     -36.556   6.522 -36.222  1.00 37.31      B    N
ATOM   1573  CA  ILE   271     -36.427   5.083 -36.064  1.00 37.51      B    C
ATOM   1574  CB  ILE   271     -36.898   4.639 -34.662  1.00 36.29      B    C
ATOM   1575  CG2 ILE   271     -37.081   3.123 -34.616  1.00 35.12      B    C
ATOM   1576  CG1 ILE   271     -35.873   5.088 -33.615  1.00 37.36      B    C
ATOM   1577  CD1 ILE   271     -36.342   4.928 -32.182  1.00 35.75      B    C
ATOM   1578  C   ILE   271     -37.243   4.359 -37.126  1.00 39.47      B    C
ATOM   1579  O   ILE   271     -36.798   3.364 -37.689  1.00 39.37      B    O
ATOM   1580  N   ARG   272     -38.439   4.864 -37.395  1.00 41.04      B    N
ATOM   1581  CA  ARG   272     -39.271   4.311 -38.448  1.00 44.21      B    C
ATOM   1582  CB  ARG   272     -40.590   5.078 -38.516  1.00 44.70      B    C
ATOM   1583  CG  ARG   272     -41.536   4.599 -39.595  1.00 47.39      B    C
ATOM   1584  CD  ARG   272     -41.632   3.085 -39.624  1.00 49.12      B    C
ATOM   1585  NE  ARG   272     -42.740   2.650 -40.470  1.00 54.01      B    N
ATOM   1586  CZ  ARG   272     -42.873   1.425 -40.970  1.00 53.66      B    C
ATOM   1587  NH1 ARG   272     -43.923   1.138 -41.727  1.00 53.00      B    N
ATOM   1588  NH2 ARG   272     -41.961   0.492 -40.723  1.00 51.99      B    N
ATOM   1589  C   ARG   272     -38.532   4.409 -39.781  1.00 46.71      B    C
ATOM   1590  O   ARG   272     -38.598   3.501 -40.607  1.00 45.87      B    O
ATOM   1591  N   LYS   273     -37.811   5.509 -39.969  1.00 49.33      B    N
ATOM   1592  CA  LYS   273     -37.071   5.752 -41.200  1.00 52.95      B    C
ATOM   1593  CB  LYS   273     -36.497   7.171 -41.189  1.00 55.26      B    C
ATOM   1594  CG  LYS   273     -36.154   7.732 -42.556  1.00 58.76      B    C
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ATOM   1595  CD  LYS   273     -37.012   8.960 -42.878  1.00 63.12      B    C
ATOM   1596  CE  LYS   273     -36.916  10.023 -41.780  1.00 66.31      B    C
ATOM   1597  NZ  LYS   273     -37.681  11.264 -42.104  1.00 67.27      B    N
ATOM   1598  C   LYS   273     -35.937   4.740 -41.351  1.00 54.29      B    C
ATOM   1599  O   LYS   273     -35.729   4.192 -42.428  1.00 54.42      B    O
ATOM   1600  N   SER   274     -35.212   4.492 -40.265  1.00 56.43      B    N
ATOM   1601  CA  SER   274     -34.101   3.547 -40.283  1.00 59.57      B    C
ATOM   1602  CB  SER   274     -33.378   3.548 -38.936  1.00 59.27      B    C
ATOM   1603  OG  SER   274     -32.785   4.806 -38.679  1.00 63.53      B    O
ATOM   1604  C   SER   274     -34.588   2.137 -40.578  1.00 61.72      B    C
ATOM   1605  O   SER   274     -33.957   1.398 -41.327  1.00 62.46      B    O
ATOM   1606  N   GLN   275     -35.717   1.774 -39.980  1.00 63.45      B    N
ATOM   1607  CA  GLN   275     -36.247   0.423 -40.080  1.00 65.81      B    C
ATOM   1608  CB  GLN   275     -37.531   0.308 -39.258  1.00 63.90      B    C
ATOM   1609  CG  GLN   275     -38.092  -1.096 -39.169  1.00 62.86      B    C
ATOM   1610  CD  GLN   275     -39.504  -1.118 -38.619  1.00 61.93      B    C
ATOM   1611  OE1 GLN   275     -40.202  -0.105 -38.636  1.00 62.80      B    O
ATOM   1612  NE2 GLN   275     -39.932  -2.275 -38.129  1.00 60.07      B    N
ATOM   1613  C   GLN   275     -36.530   0.032 -41.528  1.00 68.65      B    C
ATOM   1614  O   GLN   275     -36.466  -1.146 -41.886  1.00 69.70      B    O
ATOM   1615  N   LEU   276     -36.838   1.021 -42.359  1.00 71.49      B    N
ATOM   1616  CA  LEU   276     -37.200   0.763 -43.748  1.00 75.33      B    C
ATOM   1617  CB  LEU   276     -37.977   1.950 -44.316  1.00 73.74      B    C
ATOM   1618  CG  LEU   276     -39.275   2.267 -43.577  1.00 73.60      B    C
ATOM   1619  CD1 LEU   276     -39.941   3.488 -44.189  1.00 72.59      B    C
ATOM   1620  CD2 LEU   276     -40.191   1.060 -43.637  1.00 72.21      B    C
ATOM   1621  C   LEU   276     -35.981   0.489 -44.621  1.00 78.31      B    C
ATOM   1622  O   LEU   276     -35.921  -0.525 -45.315  1.00 80.06      B    O
ATOM   1623  N   VAL   277     -35.008   1.392 -44.581  1.00 81.41      B    N
ATOM   1624  CA  VAL   277     -33.861   1.310 -45.475  1.00 84.34      B    C
ATOM   1625  CB  VAL   277     -33.217   2.702 -45.684  1.00 84.33      B    C
ATOM   1626  CG1 VAL   277     -32.431   3.098 -44.444  1.00 83.58      B    C
ATOM   1627  CG2 VAL   277     -32.328   2.690 -46.925  1.00 84.75      B    C
ATOM   1628  C   VAL   277     -32.803   0.357 -44.921  1.00 86.12      B    C
ATOM   1629  O   VAL   277     -31.731   0.196 -45.505  1.00 87.28      B    O
ATOM   1630  N   GLN   278     -33.103  -0.271 -43.790  1.00 87.17      B    N
ATOM   1631  CA  GLN   278     -32.162  -1.200 -43.173  1.00 88.06      B    C
ATOM   1632  CB  GLN   278     -31.904  -0.802 -41.716  1.00 89.74      B    C
ATOM   1633  CG  GLN   278     -31.033   0.439 -41.567  1.00 92.15      B    C
ATOM   1634  CD  GLN   278     -30.362   0.529 -40.207  1.00 93.65      B    C
ATOM   1635  OE1 GLN   278     -29.199   0.927 -40.102  1.00 93.33      B    O
ATOM   1636  NE2 GLN   278     -31.092   0.161 -39.158  1.00 93.86      B    N
ATOM   1637  C   GLN   278     -32.653  -2.642 -43.247  1.00 87.50      B    C
ATOM   1638  O   GLN   278     -33.829  -2.897 -43.517  1.00 86.93      B    O
ATOM   1639  N   PRO   279     -31.747  -3.606 -43.013  1.00 87.19      B    N
ATOM   1640  CD  PRO   279     -30.371  -3.404 -42.522  1.00 87.00      B    C
ATOM   1641  CA  PRO   279     -32.081  -5.029 -43.144  1.00 86.05      B    C
ATOM   1642  CB  PRO   279     -30.748  -5.735 -42.902  1.00 86.93      B    C
ATOM   1643  CG  PRO   279     -29.961  -4.772 -42.064  1.00 86.83      B    C
ATOM   1644  C   PRO   279     -33.141  -5.447 -42.134  1.00 84.78      B    C
ATOM   1645  O   PRO   279     -32.885  -5.457 -40.927  1.00 84.29      B    O
ATOM   1646  N   VAL   280     -34.328  -5.791 -42.629  1.00 82.85      B    N
ATOM   1647  CA  VAL   280     -35.447  -6.087 -41.744  1.00 81.21      B    C
ATOM   1648  CB  VAL   280     -36.705  -6.529 -42.531  1.00 81.67      B    C
ATOM   1649  CG1 VAL   280     -37.219  -5.370 -43.376  1.00 82.27      B    C
ATOM   1650  CG2 VAL   280     -36.384  -7.730 -43.407  1.00 82.13      B    C
ATOM   1651  C   VAL   280     -35.077  -7.170 -40.739  1.00 78.95      B    C
ATOM   1652  O   VAL   280     -34.685  -8.280 -41.109  1.00 79.20      B    O
ATOM   1653  N   GLY   281     -35.190  -6.817 -39.463  1.00 75.44      B    N
ATOM   1654  CA  GLY   281     -34.881  -7.733 -38.382  1.00 70.08      B    C
ATOM   1655  C   GLY   281     -35.525  -7.216 -37.112  1.00 65.88      B    C
ATOM   1656  O   GLY   281     -36.125  -6.140 -37.118  1.00 65.69      B    O
ATOM   1657  N   PRO   282     -35.426  -7.955 -36.000  1.00 61.85      B    N
ATOM   1658  CD  PRO   282     -34.668  -9.198 -35.779  1.00 60.16      B    C
ATOM   1659  CA  PRO   282     -36.086  -7.478 -34.782  1.00 58.35      B    C
ATOM   1660  CB  PRO   282     -35.892  -8.629 -33.797  1.00 58.52      B    C
ATOM   1661  CG  PRO   282     -34.665  -9.334 -34.282  1.00 59.82      B    C
ATOM   1662  C   PRO   282     -35.450  -6.182 -34.288  1.00 54.89      B    C
ATOM   1663  O   PRO   282     -34.226  -6.045 -34.290  1.00 53.58      B    O
ATOM   1664  N   LEU   283     -36.271  -5.223 -33.875  1.00 51.11      B    N
ATOM   1665  CA  LEU   283     -35.724  -4.084 -33.152  1.00 49.71      B    C
ATOM   1666  CB  LEU   283     -35.573  -2.863 -34.070  1.00 51.30      B    C
ATOM   1667  CG  LEU   283     -36.798  -2.124 -34.592  1.00 53.34      B    C
ATOM   1668  CD1 LEU   283     -36.331  -1.003 -35.513  1.00 55.23      B    C
ATOM   1669  CD2 LEU   283     -37.712  -3.082 -35.342  1.00 55.78      B    C
ATOM   1670  C   LEU   283     -36.509  -3.712 -31.904  1.00 46.72      B    C
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ATOM	1671	O	LEU	283	-37.742	-3.770	-31.872	1.00	44.82	B	O
ATOM	1672	N	VAL	284	-35.756	-3.356	-30.870	1.00	43.49	B	N
ATOM	1673	CA	VAL	284	-36.301	-2.949	-29.589	1.00	39.58	B	C
ATOM	1674	CB	VAL	284	-35.610	-3.713	-28.433	1.00	40.26	B	C
ATOM	1675	CG1	VAL	284	-36.194	-3.289	-27.093	1.00	37.10	B	C
ATOM	1676	CG2	VAL	284	-35.765	-5.210	-28.634	1.00	36.99	B	C
ATOM	1677	C	VAL	284	-36.040	-1.454	-29.427	1.00	38.56	B	C
ATOM	1678	O	VAL	284	-34.955	-0.964	-29.752	1.00	36.89	B	O
ATOM	1679	N	VAL	285	-37.041	-0.731	-28.935	1.00	35.58	B	N
ATOM	1680	CA	VAL	285	-36.884	0.687	-28.646	1.00	32.71	B	C
ATOM	1681	CB	VAL	285	-37.877	1.540	-29.464	1.00	32.36	B	C
ATOM	1682	CG1	VAL	285	-37.805	2.988	-29.019	1.00	32.21	B	C
ATOM	1683	CG2	VAL	285	-37.555	1.436	-30.951	1.00	32.10	B	C
ATOM	1684	C	VAL	285	-37.113	0.942	-27.161	1.00	31.36	B	C
ATOM	1685	O	VAL	285	-38.186	0.664	-26.629	1.00	29.48	B	O
ATOM	1686	N	LEU	286	-36.093	1.471	-26.497	1.00	31.80	B	N
ATOM	1687	CA	LEU	286	-36.168	1.762	-25.074	1.00	30.07	B	C
ATOM	1688	CB	LEU	286	-34.864	1.331	-24.395	1.00	29.18	B	C
ATOM	1689	CG	LEU	286	-34.687	1.718	-22.920	1.00	30.76	B	C
ATOM	1690	CD1	LEU	286	-35.823	1.180	-22.088	1.00	28.26	B	C
ATOM	1691	CD2	LEU	286	-33.365	1.174	-22.422	1.00	30.08	B	C
ATOM	1692	C	LEU	286	-36.430	3.252	-24.842	1.00	31.06	B	C
ATOM	1693	O	LEU	286	-35.642	4.106	-25.265	1.00	30.24	B	O
ATOM	1694	N	LEU	287	-37.546	3.545	-24.174	1.00	32.05	B	N
ATOM	1695	CA	LEU	287	-37.976	4.911	-23.859	1.00	33.54	B	C
ATOM	1696	CB	LEU	287	-39.387	5.146	-24.398	1.00	35.72	B	C
ATOM	1697	CG	LEU	287	-39.574	5.223	-25.911	1.00	39.35	B	C
ATOM	1698	CD1	LEU	287	-41.043	5.030	-26.265	1.00	39.79	B	C
ATOM	1699	CD2	LEU	287	-39.074	6.571	-26.409	1.00	40.77	B	C
ATOM	1700	C	LEU	287	-37.980	5.171	-22.345	1.00	33.23	B	C
ATOM	1701	O	LEU	287	-39.026	5.134	-21.697	1.00	34.79	B	O
ATOM	1702	N	PRO	288	-36.811	5.445	-21.766	1.00	32.54	B	N
ATOM	1703	CD	PRO	288	-35.532	5.725	-22.440	1.00	31.44	B	C
ATOM	1704	CA	PRO	288	-36.699	5.567	-20.310	1.00	32.08	B	C
ATOM	1705	CB	PRO	288	-35.239	5.233	-20.048	1.00	31.66	B	C
ATOM	1706	CG	PRO	288	-34.536	5.750	-21.298	1.00	32.64	B	C
ATOM	1707	C	PRO	288	-37.060	6.968	-19.835	1.00	31.22	B	C
ATOM	1708	O	PRO	288	-36.271	7.622	-19.162	1.00	31.60	B	O
ATOM	1709	N	LEU	289	-38.247	7.431	-20.203	1.00	30.48	B	N
ATOM	1710	CA	LEU	289	-38.643	8.804	-19.929	1.00	29.51	B	C
ATOM	1711	CB	LEU	289	-38.173	9.725	-21.062	1.00	28.77	B	C
ATOM	1712	CG	LEU	289	-38.586	9.309	-22.485	1.00	30.49	B	C
ATOM	1713	CD1	LEU	289	-40.055	9.665	-22.733	1.00	28.81	B	C
ATOM	1714	CD2	LEU	289	-37.695	10.011	-23.505	1.00	27.89	B	C
ATOM	1715	C	LEU	289	-40.151	8.842	-19.808	1.00	28.98	B	C
ATOM	1716	O	LEU	289	-40.824	7.888	-20.185	1.00	30.19	B	O
ATOM	1717	N	ALA	290	-40.685	9.933	-19.274	1.00	27.61	B	N
ATOM	1718	CA	ALA	290	-42.134	10.075	-19.168	1.00	26.79	B	C
ATOM	1719	CB	ALA	290	-42.656	9.340	-17.925	1.00	25.41	B	C
ATOM	1720	C	ALA	290	-42.546	11.535	-19.118	1.00	26.19	B	C
ATOM	1721	O	ALA	290	-41.813	12.384	-18.613	1.00	25.46	B	O
ATOM	1722	N	GLY	291	-43.725	11.809	-19.665	1.00	25.85	B	N
ATOM	1723	CA	GLY	291	-44.355	13.105	-19.521	1.00	24.71	B	C
ATOM	1724	C	GLY	291	-45.814	12.869	-19.196	1.00	25.09	B	C
ATOM	1725	O	GLY	291	-46.195	11.747	-18.858	1.00	25.42	B	O
ATOM	1726	N	GLY	292	-46.632	13.912	-19.293	1.00	25.44	B	N
ATOM	1727	CA	GLY	292	-48.053	13.746	-19.062	1.00	25.43	B	C
ATOM	1728	C	GLY	292	-48.682	12.967	-20.201	1.00	29.85	B	C
ATOM	1729	O	GLY	292	-48.090	12.849	-21.281	1.00	28.22	B	O
ATOM	1730	N	TYR	293	-49.877	12.425	-19.967	1.00	30.01	B	N
ATOM	1731	CA	TYR	293	-50.562	11.671	-20.999	1.00	30.44	B	C
ATOM	1732	CB	TYR	293	-51.984	11.317	-20.549	1.00	34.81	B	C
ATOM	1733	CG	TYR	293	-52.862	10.849	-21.689	1.00	36.25	B	C
ATOM	1734	CD1	TYR	293	-52.816	9.534	-22.131	1.00	39.72	B	C
ATOM	1735	CE1	TYR	293	-53.555	9.122	-23.230	1.00	42.49	B	C
ATOM	1736	CD2	TYR	293	-53.679	11.744	-22.372	1.00	37.69	B	C
ATOM	1737	CE2	TYR	293	-54.418	11.347	-23.467	1.00	40.77	B	C
ATOM	1738	CZ	TYR	293	-54.351	10.036	-23.895	1.00	42.73	B	C
ATOM	1739	OH	TYR	293	-55.069	9.645	-25.001	1.00	46.40	B	O
ATOM	1740	C	TYR	293	-50.616	12.501	-22.277	1.00	30.43	B	C
ATOM	1741	O	TYR	293	-51.099	13.633	-22.268	1.00	28.65	B	O
ATOM	1742	N	SER	294	-50.117	11.936	-23.373	1.00	29.63	B	N
ATOM	1743	CA	SER	294	-50.139	12.616	-24.663	1.00	30.31	B	C
ATOM	1744	CB	SER	294	-48.714	13.001	-25.067	1.00	30.52	B	C
ATOM	1745	OG	SER	294	-48.628	13.246	-26.461	1.00	32.50	B	O
ATOM	1746	C	SER	294	-50.766	11.740	-25.753	1.00	31.59	B	C

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ATOM   1747  O   SER   294     -50.261  10.657 -26.070  1.00 33.33       B     O
ATOM   1748  N   ARG   295     -51.863  12.213 -26.328  1.00 32.46       B     N
ATOM   1749  CA  ARG   295     -52.540  11.471 -27.382  1.00 34.59       B     C
ATOM   1750  CB  ARG   295     -53.791  12.231 -27.839  1.00 36.79       B     C
ATOM   1751  CG  ARG   295     -54.144  11.990 -29.293  1.00 42.69       B     C
ATOM   1752  CD  ARG   295     -55.529  11.403 -29.477  1.00 43.11       B     C
ATOM   1753  NE  ARG   295     -56.529  12.421 -29.789  1.00 44.13       B     N
ATOM   1754  CZ  ARG   295     -57.453  12.301 -30.741  1.00 43.05       B     C
ATOM   1755  NH1 ARG   295     -57.506  11.207 -31.488  1.00 40.61       B     N
ATOM   1756  NH2 ARG   295     -58.340  13.269 -30.932  1.00 41.43       B     N
ATOM   1757  C   ARG   295     -51.620  11.207 -28.580  1.00 33.96       B     C
ATOM   1758  O   ARG   295     -51.573  10.084 -29.097  1.00 34.38       B     O
ATOM   1759  N   VAL   296     -50.878  12.225 -29.012  1.00 31.60       B     N
ATOM   1760  CA  VAL   296     -50.047  12.080 -30.200  1.00 30.10       B     C
ATOM   1761  CB  VAL   296     -49.564  13.476 -30.748  1.00 29.55       B     C
ATOM   1762  CG1 VAL   296     -48.665  14.165 -29.744  1.00 26.60       B     C
ATOM   1763  CG2 VAL   296     -48.836  13.296 -32.081  1.00 27.87       B     C
ATOM   1764  C   VAL   296     -48.845  11.173 -29.929  1.00 31.23       B     C
ATOM   1765  O   VAL   296     -48.451  10.380 -30.788  1.00 32.00       B     O
ATOM   1766  N   LEU   297     -48.268  11.259 -28.734  1.00 29.72       B     N
ATOM   1767  CA  LEU   297     -47.151  10.381 -28.405  1.00 29.56       B     C
ATOM   1768  CB  LEU   297     -46.496  10.813 -27.092  1.00 30.51       B     C
ATOM   1769  CG  LEU   297     -45.183  10.098 -26.782  1.00 30.69       B     C
ATOM   1770  CD1 LEU   297     -44.244  10.249 -27.966  1.00 30.90       B     C
ATOM   1771  CD2 LEU   297     -44.554  10.680 -25.526  1.00 30.10       B     C
ATOM   1772  C   LEU   297     -47.602   8.919 -28.301  1.00 30.80       B     C
ATOM   1773  O   LEU   297     -46.916   8.015 -28.786  1.00 30.98       B     O
ATOM   1774  N   ASN   298     -48.751   8.682 -27.673  1.00 28.12       B     N
ATOM   1775  CA  ASN   298     -49.312   7.333 -27.628  1.00 28.96       B     C
ATOM   1776  CB  ASN   298     -50.579   7.291 -26.758  1.00 25.34       B     C
ATOM   1777  CG  ASN   298     -50.267   7.390 -25.270  1.00 30.20       B     C
ATOM   1778  OD1 ASN   298     -49.120   7.635 -24.877  1.00 29.99       B     O
ATOM   1779  ND2 ASN   298     -51.286   7.203 -24.435  1.00 28.93       B     N
ATOM   1780  C   ASN   298     -49.631   6.815 -29.032  1.00 30.19       B     C
ATOM   1781  O   ASN   298     -49.391   5.644 -29.323  1.00 30.64       B     O
ATOM   1782  N   ALA   299     -50.155   7.684 -29.899  1.00 29.73       B     N
ATOM   1783  CA  ALA   299     -50.489   7.284 -31.271  1.00 31.32       B     C
ATOM   1784  CB  ALA   299     -51.300   8.386 -31.974  1.00 26.66       B     C
ATOM   1785  C   ALA   299     -49.244   6.957 -32.094  1.00 32.43       B     C
ATOM   1786  O   ALA   299     -49.247   6.012 -32.883  1.00 34.42       B     O
ATOM   1787  N   ALA   300     -48.177   7.728 -31.913  1.00 32.41       B     N
ATOM   1788  CA  ALA   300     -46.930   7.441 -32.615  1.00 34.19       B     C
ATOM   1789  CB  ALA   300     -45.915   8.553 -32.362  1.00 32.82       B     C
ATOM   1790  C   ALA   300     -46.362   6.091 -32.161  1.00 35.95       B     C
ATOM   1791  O   ALA   300     -45.826   5.326 -32.968  1.00 35.23       B     O
ATOM   1792  N   CYS   301     -46.483   5.798 -30.870  1.00 35.94       B     N
ATOM   1793  CA  CYS   301     -45.982   4.537 -30.341  1.00 38.41       B     C
ATOM   1794  CB  CYS   301     -46.000   4.550 -28.809  1.00 35.59       B     C
ATOM   1795  SG  CYS   301     -44.722   5.609 -28.072  1.00 39.73       B     S
ATOM   1796  C   CYS   301     -46.810   3.364 -30.857  1.00 39.27       B     C
ATOM   1797  O   CYS   301     -46.260   2.319 -31.207  1.00 38.08       B     O
ATOM   1798  N   GLN   302     -48.128   3.535 -30.913  1.00 40.20       B     N
ATOM   1799  CA  GLN   302     -48.982   2.474 -31.431  1.00 43.44       B     C
ATOM   1800  CB  GLN   302     -50.454   2.857 -31.318  1.00 44.90       B     C
ATOM   1801  CG  GLN   302     -51.371   1.816 -31.935  1.00 49.59       B     C
ATOM   1802  CD  GLN   302     -52.832   2.201 -31.856  1.00 53.42       B     C
ATOM   1803  OE1 GLN   302     -53.351   2.884 -32.741  1.00 56.88       B     O
ATOM   1804  NE2 GLN   302     -53.509   1.760 -30.798  1.00 52.21       B     N
ATOM   1805  C   GLN   302     -48.649   2.178 -32.892  1.00 43.80       B     C
ATOM   1806  O   GLN   302     -48.551   1.021 -33.298  1.00 44.02       B     O
ATOM   1807  N   ARG   303     -48.467   3.230 -33.677  1.00 43.42       B     N
ATOM   1808  CA  ARG   303     -48.146   3.068 -35.083  1.00 45.57       B     C
ATOM   1809  CB  ARG   303     -48.065   4.440 -35.747  1.00 49.32       B     C
ATOM   1810  CG  ARG   303     -48.758   4.523 -37.097  1.00 58.51       B     C
ATOM   1811  CD  ARG   303     -47.880   3.939 -38.194  1.00 66.09       B     C
ATOM   1812  NE  ARG   303     -48.459   4.098 -39.527  1.00 72.79       B     N
ATOM   1813  CZ  ARG   303     -47.840   3.755 -40.655  1.00 75.84       B     C
ATOM   1814  NH1 ARG   303     -48.439   3.933 -41.828  1.00 78.15       B     N
ATOM   1815  NH2 ARG   303     -46.617   3.236 -40.611  1.00 75.76       B     N
ATOM   1816  C   ARG   303     -46.828   2.304 -35.266  1.00 45.17       B     C
ATOM   1817  O   ARG   303     -46.701   1.481 -36.169  1.00 45.58       B     O
ATOM   1818  N   LEU   304     -45.850   2.564 -34.405  1.00 44.81       B     N
ATOM   1819  CA  LEU   304     -44.571   1.860 -34.489  1.00 44.87       B     C
ATOM   1820  CB  LEU   304     -43.534   2.512 -33.572  1.00 44.25       B     C
ATOM   1821  CG  LEU   304     -42.401   3.336 -34.182  1.00 45.66       B     C
ATOM   1822  CD1 LEU   304     -41.558   3.899 -33.052  1.00 46.67       B     C
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ATOM   1823  CD2 LEU   304     -41.537   2.485 -35.105  1.00 44.90      B    C
ATOM   1824  C   LEU   304     -44.741   0.398 -34.086  1.00 44.94      B    C
ATOM   1825  O   LEU   304     -44.161  -0.495 -34.698  1.00 45.02      B    O
ATOM   1826  N   ALA   305     -45.536   0.163 -33.047  1.00 45.15      B    N
ATOM   1827  CA  ALA   305     -45.766  -1.186 -32.548  1.00 46.08      B    C
ATOM   1828  CB  ALA   305     -46.657  -1.144 -31.310  1.00 42.15      B    C
ATOM   1829  C   ALA   305     -46.406  -2.054 -33.629  1.00 47.90      B    C
ATOM   1830  O   ALA   305     -46.089  -3.236 -33.754  1.00 48.31      B    O
ATOM   1831  N   ARG   306     -47.300  -1.461 -34.414  1.00 49.18      B    N
ATOM   1832  CA  ARG   306     -47.972  -2.189 -35.482  1.00 51.26      B    C
ATOM   1833  CB  ARG   306     -49.248  -1.449 -35.895  1.00 53.10      B    C
ATOM   1834  CG  ARG   306     -50.258  -1.368 -34.759  1.00 59.04      B    C
ATOM   1835  CD  ARG   306     -51.626  -0.912 -35.228  1.00 63.57      B    C
ATOM   1836  NE  ARG   306     -52.625  -1.044 -34.168  1.00 68.39      B    N
ATOM   1837  CZ  ARG   306     -53.925  -0.806 -34.332  1.00 71.24      B    C
ATOM   1838  NH1 ARG   306     -54.765  -0.952 -33.313  1.00 71.47      B    N
ATOM   1839  NH2 ARG   306     -54.385  -0.421 -35.516  1.00 72.89      B    N
ATOM   1840  C   ARG   306     -47.070  -2.422 -36.691  1.00 50.78      B    C
ATOM   1841  O   ARG   306     -47.418  -3.184 -37.590  1.00 51.65      B    O
ATOM   1842  N   ALA   307     -45.910  -1.773 -36.706  1.00 48.97      B    N
ATOM   1843  CA  ALA   307     -44.895  -2.049 -37.719  1.00 47.77      B    C
ATOM   1844  CB  ALA   307     -44.162  -0.772 -38.088  1.00 47.29      B    C
ATOM   1845  C   ALA   307     -43.897  -3.093 -37.221  1.00 48.22      B    C
ATOM   1846  O   ALA   307     -42.846  -3.301 -37.833  1.00 48.90      B    O
ATOM   1847  N   GLY   308     -44.218  -3.734 -36.100  1.00 47.59      B    N
ATOM   1848  CA  GLY   308     -43.385  -4.819 -35.604  1.00 47.86      B    C
ATOM   1849  C   GLY   308     -42.267  -4.428 -34.648  1.00 47.93      B    C
ATOM   1850  O   GLY   308     -41.354  -5.219 -34.408  1.00 50.24      B    O
ATOM   1851  N   VAL   309     -42.324  -3.218 -34.099  1.00 45.49      B    N
ATOM   1852  CA  VAL   309     -41.312  -2.776 -33.145  1.00 41.95      B    C
ATOM   1853  CB  VAL   309     -41.060  -1.255 -33.264  1.00 42.45      B    C
ATOM   1854  CG1 VAL   309     -40.007  -0.818 -32.256  1.00 41.14      B    C
ATOM   1855  CG2 VAL   309     -40.616  -0.912 -34.674  1.00 42.26      B    C
ATOM   1856  C   VAL   309     -41.739  -3.092 -31.714  1.00 40.15      B    C
ATOM   1857  O   VAL   309     -42.892  -2.892 -31.343  1.00 40.60      B    O
ATOM   1858  N   VAL   310     -40.804  -3.594 -30.917  1.00 38.32      B    N
ATOM   1859  CA  VAL   310     -41.023  -3.780 -29.488  1.00 35.74      B    C
ATOM   1860  CB  VAL   310     -40.163  -4.936 -28.942  1.00 36.40      B    C
ATOM   1861  CG1 VAL   310     -40.404  -5.099 -27.449  1.00 35.15      B    C
ATOM   1862  CG2 VAL   310     -40.483  -6.226 -29.689  1.00 35.82      B    C
ATOM   1863  C   VAL   310     -40.625  -2.509 -28.743  1.00 35.88      B    C
ATOM   1864  O   VAL   310     -39.440  -2.173 -28.665  1.00 35.37      B    O
ATOM   1865  N   LEU   311     -41.608  -1.798 -28.199  1.00 34.34      B    N
ATOM   1866  CA  LEU   311     -41.322  -0.609 -27.407  1.00 34.02      B    C
ATOM   1867  CB  LEU   311     -42.283   0.526 -27.768  1.00 35.30      B    C
ATOM   1868  CG  LEU   311     -42.024   1.154 -29.138  1.00 41.31      B    C
ATOM   1869  CD1 LEU   311     -42.701   0.315 -30.213  1.00 42.98      B    C
ATOM   1870  CD2 LEU   311     -42.552   2.580 -29.168  1.00 41.58      B    C
ATOM   1871  C   LEU   311     -41.410  -0.898 -25.915  1.00 33.33      B    C
ATOM   1872  O   LEU   311     -42.376  -1.502 -25.439  1.00 32.23      B    O
ATOM   1873  N   VAL   312     -40.394  -0.454 -25.184  1.00 31.89      B    N
ATOM   1874  CA  VAL   312     -40.347  -0.631 -23.739  1.00 31.45      B    C
ATOM   1875  CB  VAL   312     -39.169  -1.529 -23.335  1.00 30.23      B    C
ATOM   1876  CG1 VAL   312     -39.200  -1.774 -21.836  1.00 29.35      B    C
ATOM   1877  CG2 VAL   312     -39.225  -2.842 -24.109  1.00 29.71      B    C
ATOM   1878  C   VAL   312     -40.165   0.729 -23.079  1.00 32.28      B    C
ATOM   1879  O   VAL   312     -39.319   1.521 -23.507  1.00 30.28      B    O
ATOM   1880  N   THR   313     -40.950   1.003 -22.041  1.00 31.04      B    N
ATOM   1881  CA  THR   313     -40.906   2.317 -21.411  1.00 32.69      B    C
ATOM   1882  CB  THR   313     -42.075   3.223 -21.908  1.00 34.74      B    C
ATOM   1883  OG1 THR   313     -41.654   4.571 -21.471  1.00 40.86      B    O
ATOM   1884  CG2 THR   313     -43.405   2.750 -21.339  1.00 34.22      B    C
ATOM   1885  C   THR   313     -40.971   2.221 -19.890  1.00 32.73      B    C
ATOM   1886  O   THR   313     -41.479   1.243 -19.341  1.00 31.28      B    O
ATOM   1887  N   ALA   314     -40.456   3.246 -19.216  1.00 31.17      B    N
ATOM   1888  CA  ALA   314     -40.574   3.344 -17.767  1.00 29.71      B    C
ATOM   1889  CB  ALA   314     -39.636   4.439 -17.235  1.00 27.09      B    C
ATOM   1890  C   ALA   314     -42.017   3.664 -17.383  1.00 30.01      B    C
ATOM   1891  O   ALA   314     -42.696   4.430 -18.071  1.00 31.46      B    O
ATOM   1892  N   ALA   315     -42.480   3.087 -16.278  1.00 29.03      B    N
ATOM   1893  CA  ALA   315     -43.813   3.390 -15.767  1.00 28.17      B    C
ATOM   1894  CB  ALA   315     -44.191   2.398 -14.663  1.00 28.01      B    C
ATOM   1895  C   ALA   315     -43.904   4.816 -15.227  1.00 27.59      B    C
ATOM   1896  O   ALA   315     -44.992   5.405 -15.194  1.00 26.65      B    O
ATOM   1897  N   GLY   316     -42.766   5.361 -14.795  1.00 26.83      B    N
ATOM   1898  CA  GLY   316     -42.761   6.652 -14.123  1.00 27.76      B    C
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ATOM   1899  C   GLY   316     -42.447   6.543 -12.635  1.00 29.58      B    C
ATOM   1900  O   GLY   316     -42.688   5.505 -12.019  1.00 29.69      B    O
ATOM   1901  N   ASN   317     -41.924   7.621 -12.053  1.00 28.12      B    N
ATOM   1902  CA  ASN   317     -41.375   7.586 -10.702  1.00 26.22      B    C
ATOM   1903  CB  ASN   317     -39.962   8.172 -10.705  1.00 25.39      B    C
ATOM   1904  CG  ASN   317     -38.976   7.290 -11.441  1.00 27.85      B    C
ATOM   1905  OD1 ASN   317     -39.265   6.123 -11.712  1.00 27.83      B    O
ATOM   1906  ND2 ASN   317     -37.807   7.839 -11.772  1.00 24.72      B    N
ATOM   1907  C   ASN   317     -42.214   8.332  -9.675  1.00 27.33      B    C
ATOM   1908  O   ASN   317     -41.697   8.759  -8.637  1.00 26.61      B    O
ATOM   1909  N   PHE   318     -43.504   8.494  -9.948  1.00 26.04      B    N
ATOM   1910  CA  PHE   318     -44.293   9.446  -9.180  1.00 27.75      B    C
ATOM   1911  CB  PHE   318     -45.017  10.392 -10.146  1.00 28.25      B    C
ATOM   1912  CG  PHE   318     -44.086  11.087 -11.107  1.00 29.82      B    C
ATOM   1913  CD1 PHE   318     -43.300  12.146 -10.684  1.00 29.15      B    C
ATOM   1914  CD2 PHE   318     -43.966  10.653 -12.419  1.00 26.83      B    C
ATOM   1915  CE1 PHE   318     -42.405  12.760 -11.553  1.00 29.49      B    C
ATOM   1916  CE2 PHE   318     -43.076  11.261 -13.288  1.00 29.07      B    C
ATOM   1917  CZ  PHE   318     -42.293  12.316 -12.855  1.00 28.36      B    C
ATOM   1918  C   PHE   318     -45.284   8.806  -8.208  1.00 28.02      B    C
ATOM   1919  O   PHE   318     -46.189   9.477  -7.708  1.00 25.42      B    O
ATOM   1920  N   ARG   319     -45.102   7.514  -7.945  1.00 28.99      B    N
ATOM   1921  CA  ARG   319     -46.019   6.760  -7.097  1.00 31.68      B    C
ATOM   1922  CB  ARG   319     -45.726   7.047  -5.624  1.00 35.03      B    C
ATOM   1923  CG  ARG   319     -46.361   6.058  -4.674  1.00 40.29      B    C
ATOM   1924  CD  ARG   319     -45.680   6.070  -3.317  1.00 43.31      B    C
ATOM   1925  NE  ARG   319     -46.115   4.937  -2.505  1.00 50.86      B    N
ATOM   1926  CZ  ARG   319     -45.568   4.592  -1.340  1.00 55.39      B    C
ATOM   1927  NH1 ARG   319     -44.555   5.299  -0.845  1.00 56.32      B    N
ATOM   1928  NH2 ARG   319     -46.034   3.537  -0.673  1.00 54.10      B    N
ATOM   1929  C   ARG   319     -47.462   7.131  -7.428  1.00 30.53      B    C
ATOM   1930  O   ARG   319     -48.235   7.551  -6.566  1.00 30.48      B    O
ATOM   1931  N   ASP   320     -47.809   6.969  -8.696  1.00 30.77      B    N
ATOM   1932  CA  ASP   320     -49.078   7.444  -9.226  1.00 31.58      B    C
ATOM   1933  CB  ASP   320     -48.857   8.803  -9.906  1.00 33.04      B    C
ATOM   1934  CG  ASP   320     -50.140   9.599 -10.080  1.00 35.04      B    C
ATOM   1935  OD1 ASP   320     -51.190   9.185  -9.540  1.00 37.48      B    O
ATOM   1936  OD2 ASP   320     -50.095  10.649 -10.766  1.00 36.23      B    O
ATOM   1937  C   ASP   320     -49.594   6.412 -10.240  1.00 31.73      B    C
ATOM   1938  O   ASP   320     -48.915   5.423 -10.546  1.00 29.24      B    O
ATOM   1939  N   ASP   321     -50.795   6.649 -10.751  1.00 31.26      B    N
ATOM   1940  CA  ASP   321     -51.382   5.803 -11.782  1.00 32.35      B    C
ATOM   1941  CB  ASP   321     -52.880   6.109 -11.894  1.00 35.85      B    C
ATOM   1942  CG  ASP   321     -53.610   5.160 -12.831  1.00 40.53      B    C
ATOM   1943  OD1 ASP   321     -52.947   4.460 -13.623  1.00 41.07      B    O
ATOM   1944  OD2 ASP   321     -54.859   5.116 -12.772  1.00 46.57      B    O
ATOM   1945  C   ASP   321     -50.691   6.067 -13.123  1.00 31.57      B    C
ATOM   1946  O   ASP   321     -50.791   7.163 -13.671  1.00 30.24      B    O
ATOM   1947  N   ALA   322     -50.007   5.052 -13.648  1.00 30.47      B    N
ATOM   1948  CA  ALA   322     -49.207   5.181 -14.862  1.00 30.58      B    C
ATOM   1949  CB  ALA   322     -48.459   3.876 -15.132  1.00 29.62      B    C
ATOM   1950  C   ALA   322     -50.035   5.562 -16.085  1.00 32.45      B    C
ATOM   1951  O   ALA   322     -49.496   5.993 -17.108  1.00 32.37      B    O
ATOM   1952  N   CYS   323     -51.347   5.410 -15.989  1.00 33.17      B    N
ATOM   1953  CA  CYS   323     -52.199   5.763 -17.114  1.00 36.70      B    C
ATOM   1954  C   CYS   323     -52.321   7.285 -17.294  1.00 34.79      B    C
ATOM   1955  O   CYS   323     -52.950   7.757 -18.244  1.00 35.34      B    O
ATOM   1956  CB  CYS   323     -53.595   5.156 -16.942  1.00 41.21      B    C
ATOM   1957  SG  CYS   323     -53.747   3.326 -16.881  1.00 51.68      B    S
ATOM   1958  N   LEU   324     -51.723   8.053 -16.388  1.00 32.69      B    N
ATOM   1959  CA  LEU   324     -51.751   9.516 -16.496  1.00 31.24      B    C
ATOM   1960  CB  LEU   324     -51.868  10.146 -15.108  1.00 31.16      B    C
ATOM   1961  CG  LEU   324     -53.119   9.756 -14.313  1.00 36.04      B    C
ATOM   1962  CD1 LEU   324     -53.026  10.321 -12.893  1.00 34.08      B    C
ATOM   1963  CD2 LEU   324     -54.363  10.281 -15.024  1.00 33.82      B    C
ATOM   1964  C   LEU   324     -50.497  10.047 -17.192  1.00 29.73      B    C
ATOM   1965  O   LEU   324     -50.338  11.258 -17.366  1.00 28.10      B    O
ATOM   1966  N   TYR   325     -49.619   9.126 -17.586  1.00 28.37      B    N
ATOM   1967  CA  TYR   325     -48.319   9.457 -18.161  1.00 27.58      B    C
ATOM   1968  CB  TYR   325     -47.211   8.953 -17.225  1.00 24.47      B    C
ATOM   1969  CG  TYR   325     -47.331   9.551 -15.838  1.00 26.90      B    C
ATOM   1970  CD1 TYR   325     -46.692  10.742 -15.519  1.00 25.31      B    C
ATOM   1971  CE1 TYR   325     -46.917  11.372 -14.301  1.00 27.10      B    C
ATOM   1972  CD2 TYR   325     -48.187   8.992 -14.890  1.00 24.93      B    C
ATOM   1973  CE2 TYR   325     -48.413   9.614 -13.672  1.00 25.78      B    C
ATOM   1974  CZ  TYR   325     -47.779  10.807 -13.389  1.00 26.84      B    C
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ATOM   1975  OH  TYR   325     -48.035  11.452 -12.203  1.00 29.69      B    O
ATOM   1976  C   TYR   325     -48.167   8.829 -19.550  1.00 28.62      B    C
ATOM   1977  O   TYR   325     -48.836   7.852 -19.872  1.00 28.99      B    O
ATOM   1978  N   SER   326     -47.293   9.401 -20.372  1.00 28.16      B    N
ATOM   1979  CA  SER   326     -46.932   8.797 -21.649  1.00 27.92      B    C
ATOM   1980  CB  SER   326     -47.539   9.583 -22.813  1.00 27.53      B    C
ATOM   1981  OG  SER   326     -48.955   9.520 -22.795  1.00 29.55      B    O
ATOM   1982  C   SER   326     -45.417   8.790 -21.786  1.00 29.65      B    C
ATOM   1983  O   SER   326     -44.732   9.679 -21.264  1.00 28.75      B    O
ATOM   1984  N   PRO   327     -44.872   7.801 -22.514  1.00 30.05      B    N
ATOM   1985  CD  PRO   327     -43.446   7.791 -22.891  1.00 29.85      B    C
ATOM   1986  CA  PRO   327     -45.635   6.756 -23.212  1.00 30.77      B    C
ATOM   1987  CB  PRO   327     -44.677   6.278 -24.297  1.00 30.62      B    C
ATOM   1988  CG  PRO   327     -43.319   6.561 -23.736  1.00 32.87      B    C
ATOM   1989  C   PRO   327     -46.133   5.602 -22.333  1.00 32.88      B    C
ATOM   1990  O   PRO   327     -46.680   4.615 -22.844  1.00 34.26      B    O
ATOM   1991  N   ALA   328     -45.954   5.723 -21.020  1.00 30.65      B    N
ATOM   1992  CA  ALA   328     -46.395   4.682 -20.099  1.00 31.33      B    C
ATOM   1993  CB  ALA   328     -46.348   5.200 -18.650  1.00 29.11      B    C
ATOM   1994  C   ALA   328     -47.805   4.180 -20.427  1.00 31.48      B    C
ATOM   1995  O   ALA   328     -48.063   2.982 -20.381  1.00 30.12      B    O
ATOM   1996  N   SER   329     -48.712   5.095 -20.760  1.00 32.07      B    N
ATOM   1997  CA  SER   329     -50.122   4.743 -20.930  1.00 32.34      B    C
ATOM   1998  CB  SER   329     -51.012   5.962 -20.652  1.00 31.27      B    C
ATOM   1999  OG  SER   329     -50.757   7.013 -21.575  1.00 29.75      B    O
ATOM   2000  C   SER   329     -50.467   4.182 -22.313  1.00 34.16      B    C
ATOM   2001  O   SER   329     -51.615   3.818 -22.560  1.00 35.28      B    O
ATOM   2002  N   ALA   330     -49.494   4.125 -23.218  1.00 35.08      B    N
ATOM   2003  CA  ALA   330     -49.737   3.524 -24.531  1.00 38.27      B    C
ATOM   2004  CB  ALA   330     -48.570   3.812 -25.467  1.00 34.75      B    C
ATOM   2005  C   ALA   330     -49.923   2.014 -24.365  1.00 40.98      B    C
ATOM   2006  O   ALA   330     -49.071   1.325 -23.797  1.00 39.23      B    O
ATOM   2007  N   PRO   331     -51.054   1.479 -24.846  1.00 45.25      B    N
ATOM   2008  CD  PRO   331     -52.211   2.171 -25.439  1.00 46.82      B    C
ATOM   2009  CA  PRO   331     -51.352   0.072 -24.530  1.00 48.31      B    C
ATOM   2010  CB  PRO   331     -52.786  -0.120 -25.039  1.00 48.39      B    C
ATOM   2011  CG  PRO   331     -53.367   1.275 -25.047  1.00 49.16      B    C
ATOM   2012  C   PRO   331     -50.373  -0.922 -25.156  1.00 49.41      B    C
ATOM   2013  O   PRO   331     -50.012  -1.924 -24.532  1.00 51.91      B    O
ATOM   2014  N   GLU   332     -49.933  -0.636 -26.377  1.00 46.64      B    N
ATOM   2015  CA  GLU   332     -49.056  -1.547 -27.097  1.00 46.33      B    C
ATOM   2016  CB  GLU   332     -49.176  -1.311 -28.605  1.00 49.01      B    C
ATOM   2017  CG  GLU   332     -49.292   0.153 -29.001  1.00 57.16      B    C
ATOM   2018  CD  GLU   332     -50.640   0.770 -28.631  1.00 60.17      B    C
ATOM   2019  OE1 GLU   332     -51.647   0.490 -29.323  1.00 59.75      B    O
ATOM   2020  OE2 GLU   332     -50.688   1.538 -27.646  1.00 61.46      B    O
ATOM   2021  C   GLU   332     -47.597  -1.445 -26.665  1.00 44.15      B    C
ATOM   2022  O   GLU   332     -46.767  -2.243 -27.091  1.00 45.37      B    O
ATOM   2023  N   VAL   333     -47.286  -0.470 -25.816  1.00 41.02      B    N
ATOM   2024  CA  VAL   333     -45.942  -0.334 -25.261  1.00 37.77      B    C
ATOM   2025  CB  VAL   333     -45.622   1.142 -24.918  1.00 38.27      B    C
ATOM   2026  CG1 VAL   333     -44.275   1.235 -24.232  1.00 37.40      B    C
ATOM   2027  CG2 VAL   333     -45.625   1.989 -26.188  1.00 35.51      B    C
ATOM   2028  C   VAL   333     -45.821  -1.168 -23.986  1.00 36.92      B    C
ATOM   2029  O   VAL   333     -46.752  -1.229 -23.182  1.00 39.57      B    O
ATOM   2030  N   ILE   334     -44.679  -1.816 -23.800  1.00 33.69      B    N
ATOM   2031  CA  ILE   334     -44.452  -2.578 -22.582  1.00 32.34      B    C
ATOM   2032  CB  ILE   334     -43.411  -3.694 -22.825  1.00 33.09      B    C
ATOM   2033  CG2 ILE   334     -43.133  -4.462 -21.531  1.00 29.80      B    C
ATOM   2034  CG1 ILE   334     -43.944  -4.661 -23.889  1.00 34.39      B    C
ATOM   2035  CD1 ILE   334     -42.924  -5.688 -24.336  1.00 36.56      B    C
ATOM   2036  C   ILE   334     -43.974  -1.629 -21.487  1.00 31.39      B    C
ATOM   2037  O   ILE   334     -42.896  -1.038 -21.581  1.00 31.20      B    O
ATOM   2038  N   THR   335     -44.802  -1.476 -20.459  1.00 29.87      B    N
ATOM   2039  CA  THR   335     -44.595  -0.468 -19.428  1.00 27.70      B    C
ATOM   2040  CB  THR   335     -45.916   0.263 -19.129  1.00 28.65      B    C
ATOM   2041  OG1 THR   335     -46.402   0.865 -20.335  1.00 28.99      B    O
ATOM   2042  CG2 THR   335     -45.717   1.341 -18.064  1.00 28.22      B    C
ATOM   2043  C   THR   335     -44.086  -1.147 -18.162  1.00 28.59      B    C
ATOM   2044  O   THR   335     -44.718  -2.075 -17.658  1.00 26.97      B    O
ATOM   2045  N   VAL   336     -42.945  -0.687 -17.652  1.00 27.73      B    N
ATOM   2046  CA  VAL   336     -42.243  -1.412 -16.599  1.00 27.54      B    C
ATOM   2047  CB  VAL   336     -40.832  -1.821 -17.063  1.00 28.07      B    C
ATOM   2048  CG1 VAL   336     -40.221  -2.801 -16.070  1.00 26.47      B    C
ATOM   2049  CG2 VAL   336     -40.893  -2.421 -18.460  1.00 27.88      B    C
ATOM   2050  C   VAL   336     -42.101  -0.604 -15.310  1.00 28.83      B    C
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ATOM   2051  O   VAL   336     -41.441   0.435 -15.294  1.00 28.91      B    O
ATOM   2052  N   GLY   337     -42.716  -1.092 -14.233  1.00 28.31      B    N
ATOM   2053  CA  GLY   337     -42.482  -0.526 -12.915  1.00 27.90      B    C
ATOM   2054  C   GLY   337     -41.205  -1.068 -12.289  1.00 29.62      B    C
ATOM   2055  O   GLY   337     -40.568  -1.968 -12.846  1.00 29.27      B    O
ATOM   2056  N   ALA   338     -40.825  -0.522 -11.135  1.00 29.78      B    N
ATOM   2057  CA  ALA   338     -39.560  -0.879 -10.490  1.00 32.25      B    C
ATOM   2058  CB  ALA   338     -38.677   0.360 -10.348  1.00 29.13      B    C
ATOM   2059  C   ALA   338     -39.764  -1.526  -9.118  1.00 32.86      B    C
ATOM   2060  O   ALA   338     -40.581  -1.062  -8.317  1.00 32.98      B    O
ATOM   2061  N   THR   339     -39.010  -2.593  -8.861  1.00 33.23      B    N
ATOM   2062  CA  THR   339     -39.001  -3.263  -7.561  1.00 33.80      B    C
ATOM   2063  CB  THR   339     -39.556  -4.709  -7.667  1.00 34.64      B    C
ATOM   2064  OG1 THR   339     -38.991  -5.349  -8.819  1.00 35.54      B    O
ATOM   2065  CG2 THR   339     -41.079  -4.706  -7.774  1.00 32.31      B    C
ATOM   2066  C   THR   339     -37.577  -3.324  -7.006  1.00 34.89      B    C
ATOM   2067  O   THR   339     -36.603  -3.162  -7.749  1.00 34.08      B    O
ATOM   2068  N   ASN   340     -37.456  -3.554  -5.702  1.00 34.56      B    N
ATOM   2069  CA  ASN   340     -36.147  -3.582  -5.069  1.00 37.60      B    C
ATOM   2070  CB  ASN   340     -36.175  -2.780  -3.765  1.00 38.22      B    C
ATOM   2071  CG  ASN   340     -37.163  -3.339  -2.753  1.00 40.64      B    C
ATOM   2072  OD1 ASN   340     -37.492  -4.529  -2.771  1.00 37.96      B    O
ATOM   2073  ND2 ASN   340     -37.639  -2.478  -1.859  1.00 40.07      B    N
ATOM   2074  C   ASN   340     -35.680  -5.010  -4.799  1.00 38.46      B    C
ATOM   2075  O   ASN   340     -36.291  -5.972  -5.262  1.00 38.14      B    O
ATOM   2076  N   ALA   341     -34.593  -5.140  -4.046  1.00 39.59      B    N
ATOM   2077  CA  ALA   341     -33.991  -6.443  -3.783  1.00 42.44      B    C
ATOM   2078  CB  ALA   341     -32.656  -6.258  -3.059  1.00 41.05      B    C
ATOM   2079  C   ALA   341     -34.914  -7.361  -2.971  1.00 43.87      B    C
ATOM   2080  O   ALA   341     -34.759  -8.580  -2.988  1.00 43.82      B    O
ATOM   2081  N   GLN   342     -35.875  -6.778  -2.263  1.00 45.12      B    N
ATOM   2082  CA  GLN   342     -36.859  -7.577  -1.546  1.00 47.53      B    C
ATOM   2083  CB  GLN   342     -37.226  -6.902  -0.222  1.00 49.70      B    C
ATOM   2084  CG  GLN   342     -36.050  -6.727   0.734  1.00 55.38      B    C
ATOM   2085  CD  GLN   342     -35.164  -5.536   0.378  1.00 61.16      B    C
ATOM   2086  OE1 GLN   342     -35.658  -4.444   0.073  1.00 63.32      B    O
ATOM   2087  NE2 GLN   342     -33.848  -5.742   0.415  1.00 61.47      B    N
ATOM   2088  C   GLN   342     -38.118  -7.803  -2.386  1.00 47.92      B    C
ATOM   2089  O   GLN   342     -39.140  -8.270  -1.876  1.00 48.27      B    O
ATOM   2090  N   ASP   343     -38.037  -7.468  -3.673  1.00 47.35      B    N
ATOM   2091  CA  ASP   343     -39.170  -7.602  -4.589  1.00 47.37      B    C
ATOM   2092  CB  ASP   343     -39.621  -9.068  -4.669  1.00 49.86      B    C
ATOM   2093  CG  ASP   343     -38.718  -9.919  -5.560  1.00 52.80      B    C
ATOM   2094  OD1 ASP   343     -38.041  -9.368  -6.457  1.00 52.60      B    O
ATOM   2095  OD2 ASP   343     -38.691 -11.154  -5.366  1.00 56.09      B    O
ATOM   2096  C   ASP   343     -40.371  -6.721  -4.227  1.00 45.38      B    C
ATOM   2097  O   ASP   343     -41.501  -7.023  -4.607  1.00 46.25      B    O
ATOM   2098  N   GLN   344     -40.132  -5.634  -3.498  1.00 42.74      B    N
ATOM   2099  CA  GLN   344     -41.194  -4.674  -3.199  1.00 42.66      B    C
ATOM   2100  CB  GLN   344     -41.142  -4.269  -1.720  1.00 44.37      B    C
ATOM   2101  CG  GLN   344     -41.297  -5.432  -0.745  1.00 46.69      B    C
ATOM   2102  CD  GLN   344     -42.386  -6.418  -1.167  1.00 49.58      B    C
ATOM   2103  OE1 GLN   344     -43.575  -6.081  -1.194  1.00 50.25      B    O
ATOM   2104  NE2 GLN   344     -41.979  -7.644  -1.498  1.00 47.71      B    N
ATOM   2105  C   GLN   344     -41.068  -3.426  -4.081  1.00 40.29      B    C
ATOM   2106  O   GLN   344     -39.986  -3.105  -4.570  1.00 39.64      B    O
ATOM   2107  N   PRO   345     -42.176  -2.703  -4.289  1.00 38.37      B    N
ATOM   2108  CD  PRO   345     -43.521  -2.949  -3.745  1.00 38.04      B    C
ATOM   2109  CA  PRO   345     -42.139  -1.503  -5.136  1.00 37.61      B    C
ATOM   2110  CB  PRO   345     -43.569  -0.966  -5.068  1.00 36.21      B    C
ATOM   2111  CG  PRO   345     -44.400  -2.140  -4.664  1.00 37.92      B    C
ATOM   2112  C   PRO   345     -41.123  -0.490  -4.604  1.00 36.80      B    C
ATOM   2113  O   PRO   345     -41.083  -0.218  -3.406  1.00 36.00      B    O
ATOM   2114  N   VAL   346     -40.304   0.062  -5.493  1.00 34.88      B    N
ATOM   2115  CA  VAL   346     -39.250   0.981  -5.082  1.00 35.68      B    C
ATOM   2116  CB  VAL   346     -38.282   1.265  -6.251  1.00 37.41      B    C
ATOM   2117  CG1 VAL   346     -37.162   2.189  -5.790  1.00 41.37      B    C
ATOM   2118  CG2 VAL   346     -37.701  -0.037  -6.769  1.00 39.72      B    C
ATOM   2119  C   VAL   346     -39.826   2.310  -4.593  1.00 35.58      B    C
ATOM   2120  O   VAL   346     -40.746   2.856  -5.199  1.00 36.23      B    O
ATOM   2121  N   THR   347     -39.292   2.833  -3.494  1.00 35.01      B    N
ATOM   2122  CA  THR   347     -39.591   4.210  -3.118  1.00 35.51      B    C
ATOM   2123  CB  THR   347     -39.701   4.390  -1.582  1.00 37.60      B    C
ATOM   2124  OG1 THR   347     -38.467   4.006  -0.967  1.00 39.20      B    O
ATOM   2125  CG2 THR   347     -40.832   3.548  -1.021  1.00 34.55      B    C
ATOM   2126  C   THR   347     -38.464   5.100  -3.635  1.00 33.43      B    C
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ATOM	2127	O	THR	347	-37.300	4.716	-3.611	1.00	31.83	B	O
ATOM	2128	N	LEU	348	-38.820	6.283	-4.122	1.00	32.22	B	N
ATOM	2129	CA	LEU	348	-37.833	7.231	-4.627	1.00	30.40	B	C
ATOM	2130	CB	LEU	348	-37.946	7.326	-6.158	1.00	29.53	B	C
ATOM	2131	CG	LEU	348	-37.908	5.965	-6.886	1.00	30.64	B	C
ATOM	2132	CD1	LEU	348	-38.457	6.091	-8.307	1.00	27.70	B	C
ATOM	2133	CD2	LEU	348	-36.477	5.436	-6.912	1.00	26.20	B	C
ATOM	2134	C	LEU	348	-38.141	8.573	-3.974	1.00	28.73	B	C
ATOM	2135	O	LEU	348	-39.136	9.214	-4.303	1.00	27.79	B	O
ATOM	2136	N	GLY	349	-37.304	8.997	-3.035	1.00	28.70	B	N
ATOM	2137	CA	GLY	349	-37.703	10.112	-2.193	1.00	27.69	B	C
ATOM	2138	C	GLY	349	-38.999	9.720	-1.508	1.00	28.80	B	C
ATOM	2139	O	GLY	349	-39.113	8.603	-0.998	1.00	28.55	B	O
ATOM	2140	N	THR	350	-39.988	10.612	-1.509	1.00	28.59	B	N
ATOM	2141	CA	THR	350	-41.280	10.309	-0.900	1.00	28.89	B	C
ATOM	2142	CB	THR	350	-41.941	11.574	-0.331	1.00	30.10	B	C
ATOM	2143	OG1	THR	350	-42.291	12.450	-1.411	1.00	26.35	B	O
ATOM	2144	CG2	THR	350	-40.985	12.295	0.634	1.00	27.56	B	C
ATOM	2145	C	THR	350	-42.253	9.686	-1.906	1.00	30.35	B	C
ATOM	2146	O	THR	350	-43.407	9.403	-1.576	1.00	29.28	B	O
ATOM	2147	N	LEU	351	-41.790	9.496	-3.137	1.00	29.53	B	N
ATOM	2148	CA	LEU	351	-42.617	8.896	-4.178	1.00	30.47	B	C
ATOM	2149	CB	LEU	351	-42.669	9.820	-5.409	1.00	28.80	B	C
ATOM	2150	CG	LEU	351	-43.186	11.246	-5.126	1.00	30.54	B	C
ATOM	2151	CD1	LEU	351	-43.285	12.048	-6.426	1.00	30.77	B	C
ATOM	2152	CD2	LEU	351	-44.552	11.184	-4.448	1.00	25.93	B	C
ATOM	2153	C	LEU	351	-42.038	7.525	-4.539	1.00	30.64	B	C
ATOM	2154	O	LEU	351	-41.513	6.825	-3.677	1.00	31.17	B	O
ATOM	2155	N	GLY	352	-42.129	7.136	-5.804	1.00	30.87	B	N
ATOM	2156	CA	GLY	352	-41.659	5.817	-6.181	1.00	30.18	B	C
ATOM	2157	C	GLY	352	-42.283	5.320	-7.466	1.00	30.83	B	C
ATOM	2158	O	GLY	352	-42.842	6.104	-8.233	1.00	30.98	B	O
ATOM	2159	N	THR	353	-42.188	4.018	-7.715	1.00	29.91	B	N
ATOM	2160	CA	THR	353	-42.672	3.480	-8.974	1.00	29.17	B	C
ATOM	2161	CB	THR	353	-42.424	1.964	-9.074	1.00	32.00	B	C
ATOM	2162	OG1	THR	353	-42.923	1.490	-10.333	1.00	28.63	B	O
ATOM	2163	CG2	THR	353	-43.124	1.223	-7.924	1.00	28.13	B	C
ATOM	2164	C	THR	353	-44.164	3.744	-9.122	1.00	28.69	B	C
ATOM	2165	O	THR	353	-44.906	3.718	-8.140	1.00	26.24	B	O
ATOM	2166	N	ASN	354	-44.588	4.022	-10.352	1.00	27.36	B	N
ATOM	2167	CA	ASN	354	-46.005	4.117	-10.680	1.00	28.00	B	C
ATOM	2168	CB	ASN	354	-46.194	4.770	-12.055	1.00	30.00	B	C
ATOM	2169	CG	ASN	354	-46.090	6.295	-12.010	1.00	31.69	B	C
ATOM	2170	OD1	ASN	354	-46.028	6.902	-10.938	1.00	29.46	B	O
ATOM	2171	ND2	ASN	354	-46.079	6.916	-13.184	1.00	30.77	B	N
ATOM	2172	C	ASN	354	-46.625	2.716	-10.692	1.00	29.07	B	C
ATOM	2173	O	ASN	354	-45.915	1.705	-10.658	1.00	28.69	B	O
ATOM	2174	N	PHE	355	-47.948	2.661	-10.741	1.00	28.04	B	N
ATOM	2175	CA	PHE	355	-48.665	1.397	-10.658	1.00	29.51	B	C
ATOM	2176	CB	PHE	355	-49.041	1.118	-9.201	1.00	29.13	B	C
ATOM	2177	CG	PHE	355	-49.523	2.333	-8.465	1.00	29.69	B	C
ATOM	2178	CD1	PHE	355	-48.712	2.959	-7.522	1.00	30.05	B	C
ATOM	2179	CD2	PHE	355	-50.759	2.885	-8.752	1.00	27.67	B	C
ATOM	2180	CE1	PHE	355	-49.128	4.115	-6.887	1.00	28.38	B	C
ATOM	2181	CE2	PHE	355	-51.185	4.043	-8.121	1.00	27.82	B	C
ATOM	2182	CZ	PHE	355	-50.370	4.659	-7.189	1.00	28.92	B	C
ATOM	2183	C	PHE	355	-49.920	1.472	-11.527	1.00	30.09	B	C
ATOM	2184	O	PHE	355	-50.086	2.418	-12.309	1.00	30.27	B	O
ATOM	2185	N	GLY	356	-50.792	0.474	-11.398	1.00	28.41	B	N
ATOM	2186	CA	GLY	356	-52.069	0.519	-12.091	1.00	28.33	B	C
ATOM	2187	C	GLY	356	-52.135	-0.379	-13.314	1.00	30.65	B	C
ATOM	2188	O	GLY	356	-51.204	-1.129	-13.603	1.00	29.89	B	O
ATOM	2189	N	ARG	357	-53.242	-0.295	-14.041	1.00	32.66	B	N
ATOM	2190	CA	ARG	357	-53.536	-1.246	-15.109	1.00	34.90	B	C
ATOM	2191	CB	ARG	357	-55.011	-1.151	-15.493	1.00	34.24	B	C
ATOM	2192	CG	ARG	357	-55.381	0.225	-15.994	1.00	34.62	B	C
ATOM	2193	CD	ARG	357	-56.832	0.346	-16.383	1.00	36.26	B	C
ATOM	2194	NE	ARG	357	-57.105	1.695	-16.856	1.00	40.35	B	N
ATOM	2195	CZ	ARG	357	-56.884	2.102	-18.102	1.00	43.01	B	C
ATOM	2196	NH1	ARG	357	-57.153	3.355	-18.448	1.00	42.62	B	N
ATOM	2197	NH2	ARG	357	-56.406	1.252	-19.005	1.00	41.68	B	N
ATOM	2198	C	ARG	357	-52.683	-1.024	-16.358	1.00	36.92	B	C
ATOM	2199	O	ARG	357	-52.661	-1.881	-17.242	1.00	38.02	B	O
ATOM	2200	N	CYS	358	-51.997	0.119	-16.441	1.00	36.61	B	N
ATOM	2201	CA	CYS	358	-51.170	0.438	-17.612	1.00	35.72	B	C
ATOM	2202	C	CYS	358	-49.745	-0.097	-17.458	1.00	35.12	B	C

ATOM	2203	O	CYS	358	-48.966	-0.093	-18.410	1.00	34.79	B	O
ATOM	2204	CB	CYS	358	-51.153	1.961	-17.870	1.00	36.69	B	C
ATOM	2205	SG	CYS	358	-52.749	2.584	-18.519	1.00	43.14	B	S
ATOM	2206	N	VAL	359	-49.413	-0.558	-16.255	1.00	33.77	B	N
ATOM	2207	CA	VAL	359	-48.133	-1.222	-16.000	1.00	34.99	B	C
ATOM	2208	CB	VAL	359	-47.739	-1.110	-14.506	1.00	34.05	B	C
ATOM	2209	CG1	VAL	359	-46.472	-1.904	-14.236	1.00	33.57	B	C
ATOM	2210	CG2	VAL	359	-47.546	0.350	-14.127	1.00	35.28	B	C
ATOM	2211	C	VAL	359	-48.233	-2.709	-16.357	1.00	36.56	B	C
ATOM	2212	O	VAL	359	-49.117	-3.410	-15.874	1.00	36.77	B	O
ATOM	2213	N	ASP	360	-47.325	-3.194	-17.193	1.00	37.85	B	N
ATOM	2214	CA	ASP	360	-47.394	-4.581	-17.641	1.00	39.04	B	C
ATOM	2215	CB	ASP	360	-46.818	-4.710	-19.054	1.00	40.07	B	C
ATOM	2216	CG	ASP	360	-47.665	-3.983	-20.093	1.00	45.74	B	C
ATOM	2217	OD1	ASP	360	-48.754	-4.500	-20.443	1.00	46.61	B	O
ATOM	2218	OD2	ASP	360	-47.253	-2.891	-20.553	1.00	46.26	B	O
ATOM	2219	C	ASP	360	-46.669	-5.516	-16.688	1.00	38.35	B	C
ATOM	2220	O	ASP	360	-47.138	-6.618	-16.416	1.00	38.52	B	O
ATOM	2221	N	LEU	361	-45.529	-5.073	-16.178	1.00	37.53	B	N
ATOM	2222	CA	LEU	361	-44.819	-5.839	-15.160	1.00	36.62	B	C
ATOM	2223	CB	LEU	361	-44.105	-7.040	-15.797	1.00	36.73	B	C
ATOM	2224	CG	LEU	361	-42.822	-6.803	-16.601	1.00	38.67	B	C
ATOM	2225	CD1	LEU	361	-42.187	-8.145	-16.944	1.00	39.40	B	C
ATOM	2226	CD2	LEU	361	-43.123	-6.017	-17.870	1.00	36.31	B	C
ATOM	2227	C	LEU	361	-43.809	-4.941	-14.461	1.00	35.32	B	C
ATOM	2228	O	LEU	361	-43.605	-3.803	-14.870	1.00	36.05	B	O
ATOM	2229	N	PHE	362	-43.188	-5.457	-13.405	1.00	33.99	B	N
ATOM	2230	CA	PHE	362	-42.132	-4.743	-12.694	1.00	33.08	B	C
ATOM	2231	CB	PHE	362	-42.420	-4.724	-11.190	1.00	30.37	B	C
ATOM	2232	CG	PHE	362	-43.654	-3.967	-10.831	1.00	30.89	B	C
ATOM	2233	CD1	PHE	362	-43.569	-2.670	-10.348	1.00	30.96	B	C
ATOM	2234	CD2	PHE	362	-44.906	-4.530	-11.021	1.00	29.23	B	C
ATOM	2235	CE1	PHE	362	-44.715	-1.942	-10.065	1.00	30.60	B	C
ATOM	2236	CE2	PHE	362	-46.055	-3.809	-10.741	1.00	31.39	B	C
ATOM	2237	CZ	PHE	362	-45.960	-2.512	-10.264	1.00	31.01	B	C
ATOM	2238	C	PHE	362	-40.791	-5.415	-12.943	1.00	33.60	B	C
ATOM	2239	O	PHE	362	-40.741	-6.566	-13.376	1.00	34.30	B	O
ATOM	2240	N	ALA	363	-39.710	-4.691	-12.676	1.00	31.44	B	N
ATOM	2241	CA	ALA	363	-38.369	-5.240	-12.817	1.00	33.53	B	C
ATOM	2242	CB	ALA	363	-37.893	-5.107	-14.276	1.00	28.46	B	C
ATOM	2243	C	ALA	363	-37.427	-4.497	-11.869	1.00	34.75	B	C
ATOM	2244	O	ALA	363	-37.762	-3.423	-11.364	1.00	36.92	B	O
ATOM	2245	N	PRO	364	-36.241	-5.064	-11.606	1.00	35.95	B	N
ATOM	2246	CD	PRO	364	-35.706	-6.328	-12.149	1.00	36.33	B	C
ATOM	2247	CA	PRO	364	-35.321	-4.446	-10.645	1.00	36.66	B	C
ATOM	2248	CB	PRO	364	-34.068	-5.323	-10.733	1.00	36.17	B	C
ATOM	2249	CG	PRO	364	-34.587	-6.654	-11.198	1.00	36.16	B	C
ATOM	2250	C	PRO	364	-35.032	-2.988	-11.003	1.00	38.49	B	C
ATOM	2251	O	PRO	364	-34.633	-2.685	-12.129	1.00	38.54	B	O
ATOM	2252	N	GLY	365	-35.234	-2.089	-10.043	1.00	38.32	B	N
ATOM	2253	CA	GLY	365	-35.031	-0.679	-10.315	1.00	39.76	B	C
ATOM	2254	C	GLY	365	-34.405	0.111	-9.179	1.00	40.07	B	C
ATOM	2255	O	GLY	365	-34.394	1.339	-9.212	1.00	41.59	B	O
ATOM	2256	N	GLU	366	-33.889	-0.583	-8.172	1.00	40.60	B	N
ATOM	2257	CA	GLU	366	-33.183	0.077	-7.080	1.00	40.85	B	C
ATOM	2258	CB	GLU	366	-33.917	-0.150	-5.752	1.00	42.58	B	C
ATOM	2259	CG	GLU	366	-33.248	0.521	-4.559	1.00	48.40	B	C
ATOM	2260	CD	GLU	366	-34.106	0.485	-3.304	1.00	54.29	B	C
ATOM	2261	OE1	GLU	366	-34.917	1.418	-3.111	1.00	57.21	B	O
ATOM	2262	OE2	GLU	366	-33.970	-0.473	-2.509	1.00	57.42	B	O
ATOM	2263	C	GLU	366	-31.758	-0.446	-6.965	1.00	40.03	B	C
ATOM	2264	O	GLU	366	-31.525	-1.650	-7.067	1.00	40.58	B	O
ATOM	2265	N	ASP	367	-30.805	0.457	-6.750	1.00	39.89	B	N
ATOM	2266	CA	ASP	367	-29.431	0.049	-6.463	1.00	41.10	B	C
ATOM	2267	CB	ASP	367	-29.395	-0.733	-5.141	1.00	42.64	B	C
ATOM	2268	CG	ASP	367	-28.009	-1.250	-4.795	1.00	45.84	B	C
ATOM	2269	OD1	ASP	367	-27.011	-0.548	-5.074	1.00	44.71	B	O
ATOM	2270	OD2	ASP	367	-27.924	-2.370	-4.241	1.00	48.26	B	O
ATOM	2271	C	ASP	367	-28.901	-0.811	-7.609	1.00	40.40	B	C
ATOM	2272	O	ASP	367	-28.403	-1.919	-7.396	1.00	40.34	B	O
ATOM	2273	N	ILE	368	-29.028	-0.286	-8.827	1.00	39.65	B	N
ATOM	2274	CA	ILE	368	-28.660	-1.006	-10.041	1.00	37.49	B	C
ATOM	2275	CB	ILE	368	-29.666	-0.712	-11.182	1.00	36.39	B	C
ATOM	2276	CG2	ILE	368	-29.280	-1.479	-12.429	1.00	34.37	B	C
ATOM	2277	CG1	ILE	368	-31.077	-1.111	-10.751	1.00	36.39	B	C
ATOM	2278	CD1	ILE	368	-31.243	-2.594	-10.488	1.00	34.86	B	C

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ATOM   2279  C   ILE   368     -27.273  -0.570 -10.488  1.00 38.13      B    C
ATOM   2280  O   ILE   368     -27.066   0.586 -10.850  1.00 39.56      B    O
ATOM   2281  N   ILE   369     -26.321  -1.496 -10.464  1.00 38.77      B    N
ATOM   2282  CA  ILE   369     -24.942  -1.164 -10.793  1.00 38.05      B    C
ATOM   2283  CB  ILE   369     -23.963  -2.254 -10.257  1.00 40.93      B    C
ATOM   2284  CG2 ILE   369     -24.175  -3.561 -10.994  1.00 40.09      B    C
ATOM   2285  CG1 ILE   369     -22.513  -1.796 -10.431  1.00 41.68      B    C
ATOM   2286  CD1 ILE   369     -22.120  -0.661  -9.512  1.00 43.24      B    C
ATOM   2287  C   ILE   369     -24.832  -1.059 -12.310  1.00 36.91      B    C
ATOM   2288  O   ILE   369     -25.458  -1.832 -13.032  1.00 36.69      B    O
ATOM   2289  N   GLY   370     -24.059  -0.089 -12.793  1.00 35.97      B    N
ATOM   2290  CA  GLY   370     -23.903   0.086 -14.227  1.00 33.72      B    C
ATOM   2291  C   GLY   370     -22.820   1.086 -14.582  1.00 35.47      B    C
ATOM   2292  O   GLY   370     -22.236   1.723 -13.704  1.00 35.95      B    O
ATOM   2293  N   ALA   371     -22.554   1.235 -15.876  1.00 35.98      B    N
ATOM   2294  CA  ALA   371     -21.494   2.122 -16.342  1.00 36.57      B    C
ATOM   2295  CB  ALA   371     -21.434   2.105 -17.872  1.00 35.87      B    C
ATOM   2296  C   ALA   371     -21.666   3.559 -15.846  1.00 37.33      B    C
ATOM   2297  O   ALA   371     -22.748   4.140 -15.947  1.00 37.39      B    O
ATOM   2298  N   SER   372     -20.587   4.124 -15.314  1.00 37.76      B    N
ATOM   2299  CA  SER   372     -20.548   5.536 -14.962  1.00 39.26      B    C
ATOM   2300  CB  SER   372     -19.956   5.725 -13.568  1.00 38.89      B    C
ATOM   2301  OG  SER   372     -19.422   7.029 -13.430  1.00 38.76      B    O
ATOM   2302  C   SER   372     -19.698   6.290 -15.970  1.00 41.19      B    C
ATOM   2303  O   SER   372     -18.614   5.839 -16.325  1.00 41.88      B    O
ATOM   2304  N   SER   373     -20.184   7.438 -16.429  1.00 41.46      B    N
ATOM   2305  CA  SER   373     -19.452   8.216 -17.417  1.00 44.85      B    C
ATOM   2306  CB  SER   373     -20.398   9.175 -18.148  1.00 44.46      B    C
ATOM   2307  OG  SER   373     -21.089  10.016 -17.241  1.00 45.05      B    O
ATOM   2308  C   SER   373     -18.294   8.998 -16.792  1.00 46.64      B    C
ATOM   2309  O   SER   373     -17.564   9.697 -17.499  1.00 45.89      B    O
ATOM   2310  N   ASP   374     -18.133   8.878 -15.474  1.00 48.09      B    N
ATOM   2311  CA  ASP   374     -16.982   9.461 -14.777  1.00 51.70      B    C
ATOM   2312  CB  ASP   374     -16.901   8.951 -13.334  1.00 53.53      B    C
ATOM   2313  CG  ASP   374     -17.947   9.568 -12.428  1.00 57.35      B    C
ATOM   2314  OD1 ASP   374     -18.587  10.561 -12.841  1.00 56.93      B    O
ATOM   2315  OD2 ASP   374     -18.124   9.057 -11.296  1.00 59.88      B    O
ATOM   2316  C   ASP   374     -15.689   9.083 -15.489  1.00 52.87      B    C
ATOM   2317  O   ASP   374     -14.824   9.926 -15.718  1.00 52.95      B    O
ATOM   2318  N   CYS   375     -15.563   7.804 -15.826  1.00 53.58      B    N
ATOM   2319  CA  CYS   375     -14.369   7.301 -16.484  1.00 54.62      B    C
ATOM   2320  C   CYS   375     -14.696   5.974 -17.154  1.00 53.54      B    C
ATOM   2321  O   CYS   375     -15.668   5.314 -16.787  1.00 52.45      B    O
ATOM   2322  CB  CYS   375     -13.242   7.123 -15.460  1.00 58.13      B    C
ATOM   2323  SG  CYS   375     -13.124   5.485 -14.667  1.00 62.82      B    S
ATOM   2324  N   SER   376     -13.883   5.583 -18.130  1.00 52.69      B    N
ATOM   2325  CA  SER   376     -14.237   4.485 -19.023  1.00 52.99      B    C
ATOM   2326  CB  SER   376     -13.244   4.417 -20.186  1.00 55.67      B    C
ATOM   2327  OG  SER   376     -11.912   4.305 -19.715  1.00 60.26      B    O
ATOM   2328  C   SER   376     -14.334   3.113 -18.358  1.00 51.31      B    C
ATOM   2329  O   SER   376     -14.848   2.170 -18.956  1.00 51.62      B    O
ATOM   2330  N   THR   377     -13.847   2.994 -17.129  1.00 50.35      B    N
ATOM   2331  CA  THR   377     -13.944   1.729 -16.406  1.00 50.61      B    C
ATOM   2332  CB  THR   377     -12.543   1.174 -16.063  1.00 50.74      B    C
ATOM   2333  OG1 THR   377     -11.827   2.134 -15.275  1.00 50.85      B    O
ATOM   2334  CG2 THR   377     -11.754   0.887 -17.343  1.00 49.43      B    C
ATOM   2335  C   THR   377     -14.745   1.877 -15.115  1.00 51.23      B    C
ATOM   2336  O   THR   377     -14.872   0.932 -14.336  1.00 51.68      B    O
ATOM   2337  N   CYS   378     -15.289   3.068 -14.898  1.00 50.52      B    N
ATOM   2338  CA  CYS   378     -15.970   3.396 -13.654  1.00 50.70      B    C
ATOM   2339  C   CYS   378     -17.405   2.861 -13.638  1.00 48.67      B    C
ATOM   2340  O   CYS   378     -18.055   2.764 -14.676  1.00 47.78      B    O
ATOM   2341  CB  CYS   378     -15.942   4.919 -13.456  1.00 55.40      B    C
ATOM   2342  SG  CYS   378     -14.291   5.579 -12.988  1.00 65.66      B    S
ATOM   2343  N   PHE   379     -17.883   2.487 -12.456  1.00 47.08      B    N
ATOM   2344  CA  PHE   379     -19.251   1.997 -12.291  1.00 45.71      B    C
ATOM   2345  CB  PHE   379     -19.256   0.512 -11.908  1.00 45.48      B    C
ATOM   2346  CG  PHE   379     -18.985  -0.414 -13.056  1.00 47.99      B    C
ATOM   2347  CD1 PHE   379     -17.686  -0.645 -13.490  1.00 47.70      B    C
ATOM   2348  CD2 PHE   379     -20.030  -1.060 -13.703  1.00 48.07      B    C
ATOM   2349  CE1 PHE   379     -17.435  -1.504 -14.549  1.00 48.26      B    C
ATOM   2350  CE2 PHE   379     -19.784  -1.921 -14.765  1.00 48.64      B    C
ATOM   2351  CZ  PHE   379     -18.485  -2.143 -15.187  1.00 47.52      B    C
ATOM   2352  C   PHE   379     -19.993   2.782 -11.211  1.00 44.80      B    C
ATOM   2353  O   PHE   379     -19.377   3.334 -10.296  1.00 43.54      B    O
ATOM   2354  N   VAL   380     -21.317   2.820 -11.316  1.00 42.05      B    N
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ATOM   2355  CA  VAL   380     -22.127   3.521 -10.332  1.00 40.25       B    C
ATOM   2356  CB  VAL   380     -22.259   5.027 -10.681  1.00 40.74       B    C
ATOM   2357  CG1 VAL   380     -23.084   5.213 -11.954  1.00 41.04       B    C
ATOM   2358  CG2 VAL   380     -22.875   5.770  -9.522  1.00 40.29       B    C
ATOM   2359  C   VAL   380     -23.510   2.899 -10.228  1.00 40.11       B    C
ATOM   2360  O   VAL   380     -23.989   2.262 -11.164  1.00 41.19       B    O
ATOM   2361  N   SER   381     -24.143   3.085  -9.078  1.00 40.08       B    N
ATOM   2362  CA  SER   381     -25.440   2.485  -8.795  1.00 40.60       B    C
ATOM   2363  CB  SER   381     -25.420   1.908  -7.377  1.00 42.23       B    C
ATOM   2364  OG  SER   381     -26.290   0.800  -7.255  1.00 50.37       B    O
ATOM   2365  C   SER   381     -26.535   3.555  -8.916  1.00 38.85       B    C
ATOM   2366  O   SER   381     -26.363   4.673  -8.436  1.00 37.49       B    O
ATOM   2367  N   GLN   382     -27.652   3.218  -9.556  1.00 38.13       B    N
ATOM   2368  CA  GLN   382     -28.749   4.177  -9.728  1.00 38.91       B    C
ATOM   2369  CB  GLN   382     -28.698   4.803 -11.135  1.00 40.19       B    C
ATOM   2370  CG  GLN   382     -27.519   5.759 -11.344  1.00 43.50       B    C
ATOM   2371  CD  GLN   382     -27.521   6.441 -12.706  1.00 46.54       B    C
ATOM   2372  OE1 GLN   382     -26.583   7.166 -13.047  1.00 48.47       B    O
ATOM   2373  NE2 GLN   382     -28.573   6.214 -13.491  1.00 46.84       B    N
ATOM   2374  C   GLN   382     -30.110   3.525  -9.492  1.00 37.66       B    C
ATOM   2375  O   GLN   382     -30.231   2.301  -9.541  1.00 36.85       B    O
ATOM   2376  N   SER   383     -31.129   4.345  -9.232  1.00 36.81       B    N
ATOM   2377  CA  SER   383     -32.485   3.844  -8.985  1.00 35.64       B    C
ATOM   2378  CB  SER   383     -32.823   3.934  -7.496  1.00 33.25       B    C
ATOM   2379  OG  SER   383     -31.868   3.238  -6.722  1.00 36.82       B    O
ATOM   2380  C   SER   383     -33.545   4.608  -9.776  1.00 34.96       B    C
ATOM   2381  O   SER   383     -33.435   5.824  -9.952  1.00 36.04       B    O
ATOM   2382  N   GLY   384     -34.574   3.890 -10.227  1.00 32.10       B    N
ATOM   2383  CA  GLY   384     -35.681   4.510 -10.940  1.00 31.63       B    C
ATOM   2384  C   GLY   384     -36.335   3.539 -11.913  1.00 31.40       B    C
ATOM   2385  O   GLY   384     -35.765   2.487 -12.214  1.00 30.25       B    O
ATOM   2386  N   THR   385     -37.523   3.868 -12.414  1.00 28.45       B    N
ATOM   2387  CA  THR   385     -38.168   2.968 -13.363  1.00 30.65       B    C
ATOM   2388  CB  THR   385     -39.672   3.309 -13.567  1.00 29.80       B    C
ATOM   2389  OG1 THR   385     -39.818   4.667 -13.991  1.00 29.28       B    O
ATOM   2390  CG2 THR   385     -40.441   3.099 -12.277  1.00 28.97       B    C
ATOM   2391  C   THR   385     -37.457   2.947 -14.726  1.00 31.82       B    C
ATOM   2392  O   THR   385     -37.699   2.057 -15.537  1.00 32.77       B    O
ATOM   2393  N   SER   386     -36.571   3.911 -14.974  1.00 31.74       B    N
ATOM   2394  CA  SER   386     -35.708   3.850 -16.157  1.00 33.93       B    C
ATOM   2395  CB  SER   386     -34.783   5.069 -16.234  1.00 33.38       B    C
ATOM   2396  OG  SER   386     -35.443   6.164 -16.832  1.00 36.51       B    O
ATOM   2397  C   SER   386     -34.850   2.591 -16.139  1.00 34.63       B    C
ATOM   2398  O   SER   386     -34.772   1.872 -17.141  1.00 34.01       B    O
ATOM   2399  N   GLN   387     -34.208   2.333 -14.999  1.00 33.67       B    N
ATOM   2400  CA  GLN   387     -33.372   1.145 -14.838  1.00 33.04       B    C
ATOM   2401  CB  GLN   387     -32.711   1.129 -13.457  1.00 34.28       B    C
ATOM   2402  CG  GLN   387     -31.445   1.968 -13.356  1.00 37.41       B    C
ATOM   2403  CD  GLN   387     -31.704   3.443 -13.583  1.00 40.32       B    C
ATOM   2404  OE1 GLN   387     -30.868   4.154 -14.147  1.00 40.63       B    O
ATOM   2405  NE2 GLN   387     -32.869   3.912 -13.148  1.00 39.93       B    N
ATOM   2406  C   GLN   387     -34.190  -0.126 -15.022  1.00 33.03       B    C
ATOM   2407  O   GLN   387     -33.733  -1.074 -15.660  1.00 34.05       B    O
ATOM   2408  N   ALA   388     -35.401  -0.144 -14.477  1.00 31.18       B    N
ATOM   2409  CA  ALA   388     -36.252  -1.320 -14.606  1.00 31.67       B    C
ATOM   2410  CB  ALA   388     -37.524  -1.156 -13.771  1.00 28.74       B    C
ATOM   2411  C   ALA   388     -36.607  -1.551 -16.073  1.00 32.05       B    C
ATOM   2412  O   ALA   388     -36.515  -2.678 -16.565  1.00 31.61       B    O
ATOM   2413  N   ALA   389     -37.004  -0.490 -16.775  1.00 31.17       B    N
ATOM   2414  CA  ALA   389     -37.349  -0.628 -18.190  1.00 32.08       B    C
ATOM   2415  CB  ALA   389     -37.793   0.711 -18.771  1.00 30.72       B    C
ATOM   2416  C   ALA   389     -36.150  -1.164 -18.974  1.00 32.23       B    C
ATOM   2417  O   ALA   389     -36.309  -1.978 -19.880  1.00 31.54       B    O
ATOM   2418  N   ALA   390     -34.954  -0.708 -18.611  1.00 32.68       B    N
ATOM   2419  CA  ALA   390     -33.736  -1.155 -19.274  1.00 34.67       B    C
ATOM   2420  CB  ALA   390     -32.518  -0.409 -18.713  1.00 31.96       B    C
ATOM   2421  C   ALA   390     -33.546  -2.663 -19.113  1.00 35.82       B    C
ATOM   2422  O   ALA   390     -33.020  -3.320 -20.009  1.00 37.57       B    O
ATOM   2423  N   HIS   391     -33.975  -3.211 -17.980  1.00 35.70       B    N
ATOM   2424  CA  HIS   391     -33.864  -4.649 -17.754  1.00 36.87       B    C
ATOM   2425  CB  HIS   391     -34.258  -5.010 -16.317  1.00 37.29       B    C
ATOM   2426  CG  HIS   391     -33.151  -4.835 -15.331  1.00 40.99       B    C
ATOM   2427  CD2 HIS   391     -32.270  -5.724 -14.813  1.00 42.03       B    C
ATOM   2428  ND1 HIS   391     -32.831  -3.611 -14.781  1.00 42.60       B    N
ATOM   2429  CE1 HIS   391     -31.798  -3.754 -13.968  1.00 43.92       B    C
ATOM   2430  NE2 HIS   391     -31.439  -5.026 -13.970  1.00 43.61       B    N
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ATOM   2431  C   HIS   391     -34.754  -5.411 -18.717  1.00 35.90      B    C
ATOM   2432  O   HIS   391     -34.342  -6.416 -19.292  1.00 36.51      B    O
ATOM   2433  N   VAL   392     -35.981  -4.935 -18.886  1.00 34.43      B    N
ATOM   2434  CA  VAL   392     -36.923  -5.612 -19.755  1.00 33.29      B    C
ATOM   2435  CB  VAL   392     -38.351  -5.076 -19.546  1.00 32.77      B    C
ATOM   2436  CG1 VAL   392     -39.298  -5.720 -20.546  1.00 27.80      B    C
ATOM   2437  CG2 VAL   392     -38.811  -5.369 -18.108  1.00 29.54      B    C
ATOM   2438  C   VAL   392     -36.527  -5.460 -21.221  1.00 35.56      B    C
ATOM   2439  O   VAL   392     -36.869  -6.308 -22.043  1.00 35.03      B    O
ATOM   2440  N   ALA   393     -35.799  -4.392 -21.549  1.00 34.87      B    N
ATOM   2441  CA  ALA   393     -35.324  -4.205 -22.918  1.00 36.30      B    C
ATOM   2442  CB  ALA   393     -34.815  -2.771 -23.126  1.00 33.77      B    C
ATOM   2443  C   ALA   393     -34.207  -5.200 -23.204  1.00 36.76      B    C
ATOM   2444  O   ALA   393     -34.095  -5.712 -24.316  1.00 36.97      B    O
ATOM   2445  N   GLY   394     -33.387  -5.467 -22.191  1.00 37.87      B    N
ATOM   2446  CA  GLY   394     -32.357  -6.481 -22.312  1.00 39.01      B    C
ATOM   2447  C   GLY   394     -32.949  -7.871 -22.474  1.00 40.10      B    C
ATOM   2448  O   GLY   394     -32.509  -8.648 -23.324  1.00 39.59      B    O
ATOM   2449  N   ILE   395     -33.956  -8.181 -21.664  1.00 39.13      B    N
ATOM   2450  CA  ILE   395     -34.650  -9.455 -21.757  1.00 39.13      B    C
ATOM   2451  CB  ILE   395     -35.748  -9.560 -20.674  1.00 38.01      B    C
ATOM   2452  CG2 ILE   395     -36.645 -10.753 -20.945  1.00 36.45      B    C
ATOM   2453  CG1 ILE   395     -35.098  -9.659 -19.294  1.00 36.97      B    C
ATOM   2454  CD1 ILE   395     -36.095  -9.694 -18.144  1.00 37.21      B    C
ATOM   2455  C   ILE   395     -35.273  -9.623 -23.138  1.00 40.54      B    C
ATOM   2456  O   ILE   395     -35.123 -10.667 -23.769  1.00 41.76      B    O
ATOM   2457  N   ALA   396     -35.961  -8.591 -23.615  1.00 40.97      B    N
ATOM   2458  CA  ALA   396     -36.552  -8.636 -24.946  1.00 41.51      B    C
ATOM   2459  CB  ALA   396     -37.257  -7.324 -25.256  1.00 40.19      B    C
ATOM   2460  C   ALA   396     -35.480  -8.907 -25.994  1.00 43.15      B    C
ATOM   2461  O   ALA   396     -35.696  -9.683 -26.925  1.00 42.21      B    O
ATOM   2462  N   ALA   397     -34.323  -8.265 -25.838  1.00 43.28      B    N
ATOM   2463  CA  ALA   397     -33.256  -8.384 -26.824  1.00 44.93      B    C
ATOM   2464  CB  ALA   397     -32.144  -7.388 -26.517  1.00 41.15      B    C
ATOM   2465  C   ALA   397     -32.691  -9.805 -26.871  1.00 46.63      B    C
ATOM   2466  O   ALA   397     -32.350 -10.308 -27.941  1.00 46.80      B    O
ATOM   2467  N   MET   398     -32.594 -10.450 -25.714  1.00 48.13      B    N
ATOM   2468  CA  MET   398     -32.121 -11.827 -25.656  1.00 50.03      B    C
ATOM   2469  CB  MET   398     -31.804 -12.228 -24.218  1.00 52.81      B    C
ATOM   2470  CG  MET   398     -30.565 -11.560 -23.665  1.00 57.71      B    C
ATOM   2471  SD  MET   398     -29.966 -12.387 -22.193  1.00 67.28      B    S
ATOM   2472  CE  MET   398     -28.585 -13.313 -22.886  1.00 65.86      B    C
ATOM   2473  C   MET   398     -33.148 -12.788 -26.231  1.00 49.80      B    C
ATOM   2474  O   MET   398     -32.789 -13.787 -26.851  1.00 50.58      B    O
ATOM   2475  N   MET   399     -34.425 -12.485 -26.028  1.00 47.98      B    N
ATOM   2476  CA  MET   399     -35.484 -13.327 -26.562  1.00 47.27      B    C
ATOM   2477  CB  MET   399     -36.840 -12.911 -25.990  1.00 45.22      B    C
ATOM   2478  CG  MET   399     -37.017 -13.255 -24.527  1.00 45.38      B    C
ATOM   2479  SD  MET   399     -38.534 -12.577 -23.818  1.00 48.51      B    S
ATOM   2480  CE  MET   399     -39.806 -13.462 -24.743  1.00 46.63      B    C
ATOM   2481  C   MET   399     -35.530 -13.262 -28.081  1.00 47.45      B    C
ATOM   2482  O   MET   399     -35.728 -14.277 -28.746  1.00 49.42      B    O
ATOM   2483  N   LEU   400     -35.351 -12.069 -28.632  1.00 47.24      B    N
ATOM   2484  CA  LEU   400     -35.459 -11.880 -30.073  1.00 47.13      B    C
ATOM   2485  CB  LEU   400     -35.760 -10.413 -30.387  1.00 44.02      B    C
ATOM   2486  CG  LEU   400     -37.187 -10.011 -30.011  1.00 44.58      B    C
ATOM   2487  CD1 LEU   400     -37.354  -8.501 -30.076  1.00 43.41      B    C
ATOM   2488  CD2 LEU   400     -38.164 -10.711 -30.952  1.00 42.10      B    C
ATOM   2489  C   LEU   400     -34.183 -12.319 -30.775  1.00 48.10      B    C
ATOM   2490  O   LEU   400     -34.180 -12.610 -31.971  1.00 47.33      B    O
ATOM   2491  N   SER   401     -33.094 -12.366 -30.023  1.00 50.32      B    N
ATOM   2492  CA  SER   401     -31.834 -12.838 -30.567  1.00 53.70      B    C
ATOM   2493  CB  SER   401     -30.697 -12.527 -29.594  1.00 54.32      B    C
ATOM   2494  OG  SER   401     -29.444 -12.848 -30.170  1.00 57.99      B    O
ATOM   2495  C   SER   401     -31.918 -14.344 -30.813  1.00 54.86      B    C
ATOM   2496  O   SER   401     -31.319 -14.864 -31.754  1.00 54.93      B    O
ATOM   2497  N   ALA   402     -32.675 -15.036 -29.968  1.00 55.61      B    N
ATOM   2498  CA  ALA   402     -32.846 -16.477 -30.100  1.00 57.04      B    C
ATOM   2499  CB  ALA   402     -33.043 -17.113 -28.726  1.00 55.75      B    C
ATOM   2500  C   ALA   402     -34.024 -16.815 -31.006  1.00 58.15      B    C
ATOM   2501  O   ALA   402     -34.006 -17.836 -31.688  1.00 60.14      B    O
ATOM   2502  N   GLU   403     -35.045 -15.964 -31.011  1.00 58.70      B    N
ATOM   2503  CA  GLU   403     -36.194 -16.160 -31.889  1.00 59.99      B    C
ATOM   2504  CB  GLU   403     -37.413 -16.601 -31.081  1.00 62.05      B    C
ATOM   2505  CG  GLU   403     -37.226 -17.901 -30.323  1.00 66.30      B    C
ATOM   2506  CD  GLU   403     -38.523 -18.402 -29.719  1.00 69.10      B    C
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ATOM   2507  OE1 GLU   403     -38.640 -18.412 -28.474  1.00 70.24      B    O
ATOM   2508  OE2 GLU   403     -39.429 -18.761 -30.494  1.00 71.43      B    O
ATOM   2509  C   GLU   403     -36.539 -14.887 -32.654  1.00 60.03      B    C
ATOM   2510  O   GLU   403     -37.484 -14.178 -32.304  1.00 59.11      B    O
ATOM   2511  N   PRO   404     -35.783 -14.594 -33.723  1.00 60.25      B    N
ATOM   2512  CD  PRO   404     -34.707 -15.459 -34.235  1.00 60.12      B    C
ATOM   2513  CA  PRO   404     -35.905 -13.356 -34.502  1.00 60.18      B    C
ATOM   2514  CB  PRO   404     -34.887 -13.536 -35.629  1.00 59.91      B    C
ATOM   2515  CG  PRO   404     -33.905 -14.525 -35.090  1.00 60.66      B    C
ATOM   2516  C   PRO   404     -37.311 -13.140 -35.044  1.00 60.30      B    C
ATOM   2517  O   PRO   404     -37.687 -12.021 -35.397  1.00 60.57      B    O
ATOM   2518  N   GLU   405     -38.085 -14.215 -35.106  1.00 60.27      B    N
ATOM   2519  CA  GLU   405     -39.386 -14.164 -35.752  1.00 61.10      B    C
ATOM   2520  CB  GLU   405     -39.648 -15.476 -36.500  1.00 65.84      B    C
ATOM   2521  CG  GLU   405     -40.630 -15.348 -37.662  1.00 73.21      B    C
ATOM   2522  CD  GLU   405     -40.025 -14.645 -38.874  1.00 77.69      B    C
ATOM   2523  OE1 GLU   405     -39.331 -13.618 -38.689  1.00 79.87      B    O
ATOM   2524  OE2 GLU   405     -40.244 -15.121 -40.012  1.00 79.14      B    O
ATOM   2525  C   GLU   405     -40.511 -13.892 -34.754  1.00 58.30      B    C
ATOM   2526  O   GLU   405     -41.654 -13.658 -35.149  1.00 57.76      B    O
ATOM   2527  N   LEU   406     -40.184 -13.922 -33.463  1.00 55.23      B    N
ATOM   2528  CA  LEU   406     -41.152 -13.604 -32.414  1.00 52.90      B    C
ATOM   2529  CB  LEU   406     -40.458 -13.511 -31.057  1.00 53.28      B    C
ATOM   2530  CG  LEU   406     -40.619 -14.650 -30.053  1.00 53.34      B    C
ATOM   2531  CD1 LEU   406     -39.995 -14.228 -28.738  1.00 53.19      B    C
ATOM   2532  CD2 LEU   406     -42.085 -14.973 -29.858  1.00 52.98      B    C
ATOM   2533  C   LEU   406     -41.835 -12.275 -32.696  1.00 51.51      B    C
ATOM   2534  O   LEU   406     -41.177 -11.300 -33.058  1.00 52.21      B    O
ATOM   2535  N   THR   407     -43.152 -12.233 -32.521  1.00 49.89      B    N
ATOM   2536  CA  THR   407     -43.900 -10.987 -32.661  1.00 48.26      B    C
ATOM   2537  CB  THR   407     -45.320 -11.235 -33.201  1.00 48.96      B    C
ATOM   2538  OG1 THR   407     -46.078 -11.974 -32.234  1.00 49.44      B    O
ATOM   2539  CG2 THR   407     -45.264 -12.023 -34.499  1.00 49.09      B    C
ATOM   2540  C   THR   407     -44.029 -10.299 -31.306  1.00 47.33      B    C
ATOM   2541  O   THR   407     -43.786 -10.911 -30.263  1.00 47.26      B    O
ATOM   2542  N   LEU   408     -44.424  -9.029 -31.328  1.00 45.56      B    N
ATOM   2543  CA  LEU   408     -44.622  -8.271 -30.100  1.00 43.80      B    C
ATOM   2544  CB  LEU   408     -45.075  -6.843 -30.426  1.00 42.62      B    C
ATOM   2545  CG  LEU   408     -45.567  -5.998 -29.245  1.00 41.47      B    C
ATOM   2546  CD1 LEU   408     -44.511  -5.948 -28.163  1.00 40.64      B    C
ATOM   2547  CD2 LEU   408     -45.904  -4.605 -29.731  1.00 42.36      B    C
ATOM   2548  C   LEU   408     -45.651  -8.948 -29.197  1.00 43.17      B    C
ATOM   2549  O   LEU   408     -45.408  -9.139 -28.007  1.00 42.28      B    O
ATOM   2550  N   ALA   409     -46.791  -9.322 -29.771  1.00 43.40      B    N
ATOM   2551  CA  ALA   409     -47.849  -9.978 -29.007  1.00 44.35      B    C
ATOM   2552  CB  ALA   409     -49.023 -10.307 -29.922  1.00 44.12      B    C
ATOM   2553  C   ALA   409     -47.339 -11.251 -28.322  1.00 44.96      B    C
ATOM   2554  O   ALA   409     -47.599 -11.475 -27.137  1.00 43.20      B    O
ATOM   2555  N   GLU   410     -46.602 -12.074 -29.066  1.00 44.33      B    N
ATOM   2556  CA  GLU   410     -46.036 -13.294 -28.506  1.00 46.70      B    C
ATOM   2557  CB  GLU   410     -45.371 -14.130 -29.608  1.00 50.37      B    C
ATOM   2558  CG  GLU   410     -46.344 -14.648 -30.661  1.00 57.24      B    C
ATOM   2559  CD  GLU   410     -45.650 -15.186 -31.909  1.00 61.50      B    C
ATOM   2560  OE1 GLU   410     -46.351 -15.766 -32.769  1.00 65.56      B    O
ATOM   2561  OE2 GLU   410     -44.413 -15.031 -32.035  1.00 61.64      B    O
ATOM   2562  C   GLU   410     -45.015 -12.977 -27.419  1.00 44.90      B    C
ATOM   2563  O   GLU   410     -44.977 -13.631 -26.374  1.00 44.41      B    O
ATOM   2564  N   LEU   411     -44.186 -11.971 -27.662  1.00 43.46      B    N
ATOM   2565  CA  LEU   411     -43.159 -11.610 -26.700  1.00 41.39      B    C
ATOM   2566  CB  LEU   411     -42.219 -10.568 -27.303  1.00 41.35      B    C
ATOM   2567  CG  LEU   411     -41.034 -10.165 -26.428  1.00 41.21      B    C
ATOM   2568  CD1 LEU   411     -39.814  -9.918 -27.296  1.00 42.34      B    C
ATOM   2569  CD2 LEU   411     -41.398  -8.924 -25.626  1.00 43.75      B    C
ATOM   2570  C   LEU   411     -43.780 -11.082 -25.408  1.00 40.36      B    C
ATOM   2571  O   LEU   411     -43.311 -11.404 -24.313  1.00 39.22      B    O
ATOM   2572  N   ARG   412     -44.839 -10.284 -25.531  1.00 38.70      B    N
ATOM   2573  CA  ARG   412     -45.516  -9.746 -24.353  1.00 39.79      B    C
ATOM   2574  CB  ARG   412     -46.621  -8.759 -24.763  1.00 39.43      B    C
ATOM   2575  CG  ARG   412     -47.375  -8.151 -23.573  1.00 40.81      B    C
ATOM   2576  CD  ARG   412     -48.584  -7.329 -24.013  1.00 42.83      B    C
ATOM   2577  NE  ARG   412     -48.208  -6.224 -24.891  1.00 46.65      B    N
ATOM   2578  CZ  ARG   412     -48.520  -6.140 -26.181  1.00 49.92      B    C
ATOM   2579  NH1 ARG   412     -48.126  -5.090 -26.891  1.00 51.57      B    N
ATOM   2580  NH2 ARG   412     -49.230  -7.099 -26.766  1.00 50.17      B    N
ATOM   2581  C   ARG   412     -46.120 -10.868 -23.501  1.00 40.47      B    C
ATOM   2582  O   ARG   412     -45.977 -10.870 -22.271  1.00 39.56      B    O
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ATOM   2583  N   GLN   413     -46.791 -11.817 -24.156  1.00 40.04       B    N
ATOM   2584  CA  GLN   413     -47.414 -12.944 -23.456  1.00 38.98       B    C
ATOM   2585  CB  GLN   413     -48.226 -13.807 -24.432  1.00 37.49       B    C
ATOM   2586  CG  GLN   413     -49.486 -13.134 -24.982  1.00 35.37       B    C
ATOM   2587  CD  GLN   413     -50.730 -13.433 -24.154  1.00 38.44       B    C
ATOM   2588  OE1 GLN   413     -50.652 -14.037 -23.079  1.00 37.99       B    O
ATOM   2589  NE2 GLN   413     -51.888 -13.012 -24.655  1.00 38.63       B    N
ATOM   2590  C   GLN   413     -46.362 -13.804 -22.765  1.00 38.99       B    C
ATOM   2591  O   GLN   413     -46.598 -14.322 -21.676  1.00 39.28       B    O
ATOM   2592  N   ARG   414     -45.197 -13.943 -23.387  1.00 40.24       B    N
ATOM   2593  CA  ARG   414     -44.119 -14.719 -22.787  1.00 43.34       B    C
ATOM   2594  CB  ARG   414     -43.034 -15.020 -23.825  1.00 46.55       B    C
ATOM   2595  CG  ARG   414     -43.365 -16.199 -24.712  1.00 52.72       B    C
ATOM   2596  CD  ARG   414     -42.260 -16.497 -25.713  1.00 58.62       B    C
ATOM   2597  NE  ARG   414     -42.618 -17.605 -26.599  1.00 65.06       B    N
ATOM   2598  CZ  ARG   414     -43.771 -17.696 -27.262  1.00 68.06       B    C
ATOM   2599  NH1 ARG   414     -44.692 -16.745 -27.146  1.00 68.20       B    N
ATOM   2600  NH2 ARG   414     -44.005 -18.741 -28.049  1.00 69.29       B    N
ATOM   2601  C   ARG   414     -43.502 -14.018 -21.583  1.00 44.39       B    C
ATOM   2602  O   ARG   414     -43.229 -14.654 -20.564  1.00 44.57       B    O
ATOM   2603  N   LEU   415     -43.283 -12.710 -21.693  1.00 44.56       B    N
ATOM   2604  CA  LEU   415     -42.792 -11.938 -20.555  1.00 45.03       B    C
ATOM   2605  CB  LEU   415     -42.684 -10.453 -20.906  1.00 46.00       B    C
ATOM   2606  CG  LEU   415     -41.424 -10.032 -21.661  1.00 46.93       B    C
ATOM   2607  CD1 LEU   415     -41.516  -8.555 -22.037  1.00 46.93       B    C
ATOM   2608  CD2 LEU   415     -40.210 -10.296 -20.789  1.00 44.97       B    C
ATOM   2609  C   LEU   415     -43.726 -12.103 -19.368  1.00 44.14       B    C
ATOM   2610  O   LEU   415     -43.279 -12.333 -18.247  1.00 43.74       B    O
ATOM   2611  N   ILE   416     -45.026 -11.988 -19.620  1.00 43.81       B    N
ATOM   2612  CA  ILE   416     -46.013 -12.126 -18.556  1.00 44.62       B    C
ATOM   2613  CB  ILE   416     -47.445 -11.843 -19.063  1.00 43.70       B    C
ATOM   2614  CG2 ILE   416     -48.453 -12.190 -17.978  1.00 41.87       B    C
ATOM   2615  CG1 ILE   416     -47.576 -10.375 -19.475  1.00 43.04       B    C
ATOM   2616  CD1 ILE   416     -48.959  -9.995 -19.974  1.00 40.89       B    C
ATOM   2617  C   ILE   416     -45.985 -13.534 -17.982  1.00 46.84       B    C
ATOM   2618  O   ILE   416     -46.015 -13.719 -16.765  1.00 47.38       B    O
ATOM   2619  N   HIS   417     -45.920 -14.526 -18.865  1.00 48.11       B    N
ATOM   2620  CA  HIS   417     -46.011 -15.916 -18.452  1.00 48.47       B    C
ATOM   2621  CB  HIS   417     -46.122 -16.829 -19.675  1.00 50.72       B    C
ATOM   2622  CG  HIS   417     -46.218 -18.280 -19.327  1.00 52.76       B    C
ATOM   2623  CD2 HIS   417     -45.281 -19.259 -19.319  1.00 52.85       B    C
ATOM   2624  ND1 HIS   417     -47.385 -18.862 -18.879  1.00 54.15       B    N
ATOM   2625  CE1 HIS   417     -47.162 -20.135 -18.607  1.00 54.24       B    C
ATOM   2626  NE2 HIS   417     -45.894 -20.401 -18.865  1.00 53.33       B    N
ATOM   2627  C   HIS   417     -44.836 -16.367 -17.593  1.00 48.85       B    C
ATOM   2628  O   HIS   417     -45.018 -17.142 -16.654  1.00 49.47       B    O
ATOM   2629  N   PHE   418     -43.638 -15.885 -17.906  1.00 48.81       B    N
ATOM   2630  CA  PHE   418     -42.432 -16.358 -17.229  1.00 50.61       B    C
ATOM   2631  CB  PHE   418     -41.310 -16.566 -18.247  1.00 53.20       B    C
ATOM   2632  CG  PHE   418     -41.549 -17.719 -19.181  1.00 56.54       B    C
ATOM   2633  CD1 PHE   418     -42.039 -17.505 -20.460  1.00 56.42       B    C
ATOM   2634  CD2 PHE   418     -41.291 -19.021 -18.775  1.00 58.19       B    C
ATOM   2635  CE1 PHE   418     -42.269 -18.565 -21.320  1.00 57.36       B    C
ATOM   2636  CE2 PHE   418     -41.519 -20.089 -19.632  1.00 58.59       B    C
ATOM   2637  CZ  PHE   418     -42.009 -19.859 -20.906  1.00 57.63       B    C
ATOM   2638  C   PHE   418     -41.931 -15.465 -16.089  1.00 50.45       B    C
ATOM   2639  O   PHE   418     -40.865 -15.715 -15.523  1.00 50.20       B    O
ATOM   2640  N   SER   419     -42.694 -14.431 -15.751  1.00 49.81       B    N
ATOM   2641  CA  SER   419     -42.356 -13.573 -14.615  1.00 49.94       B    C
ATOM   2642  CB  SER   419     -43.176 -12.285 -14.664  1.00 50.42       B    C
ATOM   2643  OG  SER   419     -42.857 -11.527 -15.811  1.00 53.88       B    O
ATOM   2644  C   SER   419     -42.638 -14.284 -13.299  1.00 48.56       B    C
ATOM   2645  O   SER   419     -43.483 -15.173 -13.238  1.00 49.17       B    O
ATOM   2646  N   ALA   420     -41.941 -13.884 -12.242  1.00 47.18       B    N
ATOM   2647  CA  ALA   420     -42.274 -14.368 -10.909  1.00 47.78       B    C
ATOM   2648  CB  ALA   420     -41.183 -13.978  -9.926  1.00 45.27       B    C
ATOM   2649  C   ALA   420     -43.615 -13.777 -10.475  1.00 48.76       B    C
ATOM   2650  O   ALA   420     -43.809 -12.565 -10.530  1.00 50.00       B    O
ATOM   2651  N   LYS   421     -44.536 -14.639 -10.054  1.00 48.99       B    N
ATOM   2652  CA  LYS   421     -45.865 -14.210  -9.635  1.00 50.27       B    C
ATOM   2653  CB  LYS   421     -46.913 -15.265 -10.019  1.00 52.20       B    C
ATOM   2654  CG  LYS   421     -47.365 -15.245 -11.477  1.00 53.70       B    C
ATOM   2655  CD  LYS   421     -46.221 -15.548 -12.432  1.00 58.05       B    C
ATOM   2656  CE  LYS   421     -46.722 -16.031 -13.797  1.00 59.76       B    C
ATOM   2657  NZ  LYS   421     -47.727 -15.111 -14.413  1.00 60.22       B    N
ATOM   2658  C   LYS   421     -45.935 -13.964  -8.125  1.00 51.72       B    C
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ATOM   2659  O   LYS   421      -45.293 -14.664  -7.340  1.00 51.06      B    O
ATOM   2660  N   ASP   422      -46.711 -12.959  -7.731  1.00 51.17      B    N
ATOM   2661  CA  ASP   422      -47.116 -12.783  -6.343  1.00 52.74      B    C
ATOM   2662  CB  ASP   422      -47.980 -13.967  -5.899  1.00 55.89      B    C
ATOM   2663  CG  ASP   422      -49.341 -13.978  -6.571  1.00 60.38      B    C
ATOM   2664  OD1 ASP   422      -50.008 -12.919  -6.588  1.00 62.29      B    O
ATOM   2665  OD2 ASP   422      -49.744 -15.045  -7.085  1.00 63.14      B    O
ATOM   2666  C   ASP   422      -45.987 -12.586  -5.339  1.00 51.77      B    C
ATOM   2667  O   ASP   422      -46.155 -12.874  -4.157  1.00 51.33      B    O
ATOM   2668  N   VAL   423      -44.843 -12.091  -5.791  1.00 51.65      B    N
ATOM   2669  CA  VAL   423      -43.753 -11.799  -4.867  1.00 50.55      B    C
ATOM   2670  CB  VAL   423      -42.385 -11.905  -5.564  1.00 52.04      B    C
ATOM   2671  CG1 VAL   423      -42.114 -13.351  -5.942  1.00 51.21      B    C
ATOM   2672  CG2 VAL   423      -42.361 -11.022  -6.803  1.00 51.94      B    C
ATOM   2673  C   VAL   423      -43.878 -10.414  -4.238  1.00 50.83      B    C
ATOM   2674  O   VAL   423      -43.196 -10.111  -3.260  1.00 50.92      B    O
ATOM   2675  N   ILE   424      -44.756  -9.581  -4.797  1.00 49.90      B    N
ATOM   2676  CA  ILE   424      -44.927  -8.203  -4.335  1.00 48.89      B    C
ATOM   2677  CB  ILE   424      -45.264  -7.261  -5.511  1.00 47.42      B    C
ATOM   2678  CG2 ILE   424      -45.468  -5.845  -5.006  1.00 46.42      B    C
ATOM   2679  CG1 ILE   424      -44.148  -7.295  -6.551  1.00 46.55      B    C
ATOM   2680  CD1 ILE   424      -44.528  -6.612  -7.846  1.00 44.25      B    C
ATOM   2681  C   ILE   424      -46.059  -8.081  -3.316  1.00 49.43      B    C
ATOM   2682  O   ILE   424      -47.160  -8.578  -3.544  1.00 49.10      B    O
ATOM   2683  N   ASN   425      -45.802  -7.400  -2.205  1.00 51.29      B    N
ATOM   2684  CA  ASN   425      -46.873  -7.103  -1.256  1.00 53.93      B    C
ATOM   2685  CB  ASN   425      -46.306  -6.833   0.143  1.00 53.60      B    C
ATOM   2686  CG  ASN   425      -47.398  -6.613   1.179  1.00 55.99      B    C
ATOM   2687  OD1 ASN   425      -48.587  -6.600   0.856  1.00 56.43      B    O
ATOM   2688  ND2 ASN   425      -46.997  -6.443   2.433  1.00 57.69      B    N
ATOM   2689  C   ASN   425      -47.668  -5.889  -1.728  1.00 54.29      B    C
ATOM   2690  O   ASN   425      -47.196  -4.754  -1.654  1.00 55.32      B    O
ATOM   2691  N   GLU   426      -48.883  -6.134  -2.200  1.00 54.68      B    N
ATOM   2692  CA  GLU   426      -49.667  -5.102  -2.863  1.00 56.26      B    C
ATOM   2693  CB  GLU   426      -50.764  -5.753  -3.708  1.00 59.36      B    C
ATOM   2694  CG  GLU   426      -51.692  -6.660  -2.922  1.00 65.60      B    C
ATOM   2695  CD  GLU   426      -52.426  -7.656  -3.807  1.00 70.48      B    C
ATOM   2696  OE1 GLU   426      -52.050  -7.798  -4.999  1.00 71.64      B    O
ATOM   2697  OE2 GLU   426      -53.377  -8.298  -3.303  1.00 71.69      B    O
ATOM   2698  C   GLU   426      -50.286  -4.088  -1.901  1.00 55.20      B    C
ATOM   2699  O   GLU   426      -51.016  -3.191  -2.327  1.00 54.27      B    O
ATOM   2700  N   ALA   427      -49.997  -4.227  -0.611  1.00 53.77      B    N
ATOM   2701  CA  ALA   427      -50.452  -3.248   0.375  1.00 52.69      B    C
ATOM   2702  CB  ALA   427      -50.142  -3.741   1.788  1.00 51.14      B    C
ATOM   2703  C   ALA   427      -49.772  -1.900   0.125  1.00 51.48      B    C
ATOM   2704  O   ALA   427      -50.293  -0.850   0.505  1.00 50.75      B    O
ATOM   2705  N   TRP   428      -48.610  -1.939  -0.524  1.00 49.80      B    N
ATOM   2706  CA  TRP   428      -47.888  -0.727  -0.896  1.00 49.29      B    C
ATOM   2707  CB  TRP   428      -46.564  -1.105  -1.566  1.00 51.42      B    C
ATOM   2708  CG  TRP   428      -45.580   0.034  -1.742  1.00 55.20      B    C
ATOM   2709  CD2 TRP   428      -45.451   0.901  -2.883  1.00 55.07      B    C
ATOM   2710  CE2 TRP   428      -44.347   1.749  -2.642  1.00 55.79      B    C
ATOM   2711  CE3 TRP   428      -46.157   1.039  -4.085  1.00 54.96      B    C
ATOM   2712  CD1 TRP   428      -44.500   0.393  -0.879  1.00 55.62      B    C
ATOM   2713  NE1 TRP   428      -43.835   1.419  -1.414  1.00 55.96      B    N
ATOM   2714  CZ2 TRP   428      -43.932   2.722  -3.561  1.00 54.75      B    C
ATOM   2715  CZ3 TRP   428      -45.742   2.009  -5.000  1.00 53.56      B    C
ATOM   2716  CH2 TRP   428      -44.640   2.834  -4.730  1.00 52.93      B    C
ATOM   2717  C   TRP   428      -48.724   0.152  -1.838  1.00 48.43      B    C
ATOM   2718  O   TRP   428      -48.662   1.383  -1.767  1.00 47.88      B    O
ATOM   2719  N   PHE   429      -49.511  -0.478  -2.709  1.00 44.61      B    N
ATOM   2720  CA  PHE   429      -50.352   0.262  -3.645  1.00 43.45      B    C
ATOM   2721  CB  PHE   429      -50.741  -0.622  -4.832  1.00 41.72      B    C
ATOM   2722  CG  PHE   429      -49.569  -1.259  -5.525  1.00 43.10      B    C
ATOM   2723  CD1 PHE   429      -49.537  -2.630  -5.745  1.00 42.99      B    C
ATOM   2724  CD2 PHE   429      -48.493  -0.492  -5.943  1.00 41.43      B    C
ATOM   2725  CE1 PHE   429      -48.448  -3.224  -6.368  1.00 43.54      B    C
ATOM   2726  CE2 PHE   429      -47.403  -1.079  -6.567  1.00 42.19      B    C
ATOM   2727  CZ  PHE   429      -47.380  -2.446  -6.780  1.00 42.81      B    C
ATOM   2728  C   PHE   429      -51.616   0.750  -2.951  1.00 43.79      B    C
ATOM   2729  O   PHE   429      -52.095   0.126  -2.008  1.00 44.37      B    O
ATOM   2730  N   PRO   430      -52.180   1.874  -3.414  1.00 43.28      B    N
ATOM   2731  CD  PRO   430      -51.705   2.808  -4.448  1.00 43.12      B    C
ATOM   2732  CA  PRO   430      -53.488   2.253  -2.877  1.00 44.00      B    C
ATOM   2733  CB  PRO   430      -53.822   3.562  -3.600  1.00 41.64      B    C
ATOM   2734  CG  PRO   430      -52.943   3.592  -4.791  1.00 42.89      B    C
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ATOM	2735	C	PRO	430	-54.508	1.154	-3.147	1.00	46.37	B	C
ATOM	2736	O	PRO	430	-54.406	0.428	-4.141	1.00	45.21	B	O
ATOM	2737	N	GLU	431	-55.488	1.037	-2.257	1.00	48.35	B	N
ATOM	2738	CA	GLU	431	-56.412	-0.092	-2.274	1.00	51.07	B	C
ATOM	2739	CB	GLU	431	-57.493	0.099	-1.209	1.00	55.08	B	C
ATOM	2740	CG	GLU	431	-58.513	-1.030	-1.154	1.00	61.74	B	C
ATOM	2741	CD	GLU	431	-59.551	-0.832	-0.057	1.00	67.18	B	C
ATOM	2742	OE1	GLU	431	-59.472	0.186	0.670	1.00	67.29	B	O
ATOM	2743	OE2	GLU	431	-60.446	-1.699	0.078	1.00	69.23	B	O
ATOM	2744	C	GLU	431	-57.073	-0.291	-3.633	1.00	50.09	B	C
ATOM	2745	O	GLU	431	-57.153	-1.413	-4.127	1.00	49.89	B	O
ATOM	2746	N	ASP	432	-57.543	0.798	-4.232	1.00	48.97	B	N
ATOM	2747	CA	ASP	432	-58.306	0.715	-5.470	1.00	49.55	B	C
ATOM	2748	CB	ASP	432	-59.094	2.015	-5.696	1.00	53.81	B	C
ATOM	2749	CG	ASP	432	-58.192	3.233	-5.900	1.00	58.64	B	C
ATOM	2750	OD1	ASP	432	-56.988	3.178	-5.558	1.00	61.37	B	O
ATOM	2751	OD2	ASP	432	-58.699	4.256	-6.407	1.00	61.51	B	O
ATOM	2752	C	ASP	432	-57.440	0.412	-6.697	1.00	48.17	B	C
ATOM	2753	O	ASP	432	-57.948	0.348	-7.815	1.00	48.22	B	O
ATOM	2754	N	GLN	433	-56.141	0.222	-6.488	1.00	45.67	B	N
ATOM	2755	CA	GLN	433	-55.230	-0.089	-7.588	1.00	44.78	B	C
ATOM	2756	CB	GLN	433	-54.055	0.901	-7.605	1.00	43.98	B	C
ATOM	2757	CG	GLN	433	-54.442	2.340	-7.909	1.00	44.11	B	C
ATOM	2758	CD	GLN	433	-54.838	2.543	-9.360	1.00	45.54	B	C
ATOM	2759	OE1	GLN	433	-54.530	1.723	-10.222	1.00	45.75	B	O
ATOM	2760	NE2	GLN	433	-55.523	3.641	-9.636	1.00	46.92	B	N
ATOM	2761	C	GLN	433	-54.685	-1.511	-7.468	1.00	44.14	B	C
ATOM	2762	O	GLN	433	-54.026	-2.017	-8.376	1.00	42.38	B	O
ATOM	2763	N	ARG	434	-54.950	-2.153	-6.339	1.00	44.08	B	N
ATOM	2764	CA	ARG	434	-54.354	-3.453	-6.068	1.00	45.47	B	C
ATOM	2765	CB	ARG	434	-54.660	-3.868	-4.635	1.00	45.94	B	C
ATOM	2766	CG	ARG	434	-53.833	-3.115	-3.623	1.00	48.40	B	C
ATOM	2767	CD	ARG	434	-54.429	-3.182	-2.233	1.00	48.11	B	C
ATOM	2768	NE	ARG	434	-53.674	-2.337	-1.314	1.00	49.63	B	N
ATOM	2769	CZ	ARG	434	-54.127	-1.911	-0.141	1.00	50.56	B	C
ATOM	2770	NH1	ARG	434	-53.360	-1.144	0.624	1.00	50.03	B	N
ATOM	2771	NH2	ARG	434	-55.343	-2.252	0.267	1.00	49.58	B	N
ATOM	2772	C	ARG	434	-54.827	-4.530	-7.045	1.00	45.92	B	C
ATOM	2773	O	ARG	434	-54.049	-5.396	-7.460	1.00	46.72	B	O
ATOM	2774	N	VAL	435	-56.093	-4.472	-7.428	1.00	44.65	B	N
ATOM	2775	CA	VAL	435	-56.615	-5.460	-8.353	1.00	46.78	B	C
ATOM	2776	CB	VAL	435	-58.163	-5.521	-8.286	1.00	47.93	B	C
ATOM	2777	CG1	VAL	435	-58.772	-4.227	-8.817	1.00	46.92	B	C
ATOM	2778	CG2	VAL	435	-58.660	-6.720	-9.061	1.00	48.93	B	C
ATOM	2779	C	VAL	435	-56.167	-5.141	-9.780	1.00	46.07	B	C
ATOM	2780	O	VAL	435	-56.031	-6.035	-10.614	1.00	47.62	B	O
ATOM	2781	N	LEU	436	-55.917	-3.865	-10.051	1.00	44.26	B	N
ATOM	2782	CA	LEU	436	-55.553	-3.424	-11.394	1.00	41.98	B	C
ATOM	2783	CB	LEU	436	-55.896	-1.946	-11.568	1.00	42.62	B	C
ATOM	2784	CG	LEU	436	-57.362	-1.584	-11.362	1.00	43.63	B	C
ATOM	2785	CD1	LEU	436	-57.563	-0.083	-11.533	1.00	44.20	B	C
ATOM	2786	CD2	LEU	436	-58.195	-2.346	-12.366	1.00	44.78	B	C
ATOM	2787	C	LEU	436	-54.079	-3.631	-11.725	1.00	40.11	B	C
ATOM	2788	O	LEU	436	-53.700	-3.603	-12.891	1.00	40.58	B	O
ATOM	2789	N	THR	437	-53.246	-3.825	-10.708	1.00	38.98	B	N
ATOM	2790	CA	THR	437	-51.802	-3.821	-10.918	1.00	38.27	B	C
ATOM	2791	CB	THR	437	-51.070	-3.044	-9.797	1.00	36.73	B	C
ATOM	2792	OG1	THR	437	-51.579	-1.708	-9.721	1.00	38.09	B	O
ATOM	2793	CG2	THR	437	-49.576	-2.989	-10.077	1.00	35.50	B	C
ATOM	2794	C	THR	437	-51.238	-5.238	-10.959	1.00	38.94	B	C
ATOM	2795	O	THR	437	-51.376	-5.995	-10.003	1.00	39.17	B	O
ATOM	2796	N	PRO	438	-50.580	-5.610	-12.068	1.00	39.47	B	N
ATOM	2797	CD	PRO	438	-50.366	-4.833	-13.299	1.00	39.57	B	C
ATOM	2798	CA	PRO	438	-50.051	-6.972	-12.182	1.00	38.85	B	C
ATOM	2799	CB	PRO	438	-49.477	-7.026	-13.601	1.00	39.01	B	C
ATOM	2800	CG	PRO	438	-50.153	-5.911	-14.333	1.00	41.00	B	C
ATOM	2801	C	PRO	438	-48.983	-7.238	-11.129	1.00	39.53	B	C
ATOM	2802	O	PRO	438	-48.046	-6.454	-10.967	1.00	39.03	B	O
ATOM	2803	N	ASN	439	-49.124	-8.349	-10.418	1.00	39.63	B	N
ATOM	2804	CA	ASN	439	-48.128	-8.746	-9.437	1.00	40.67	B	C
ATOM	2805	CB	ASN	439	-48.802	-9.512	-8.293	1.00	38.48	B	C
ATOM	2806	CG	ASN	439	-47.993	-9.469	-7.011	1.00	39.26	B	C
ATOM	2807	OD1	ASN	439	-46.780	-9.700	-7.017	1.00	37.92	B	O
ATOM	2808	ND2	ASN	439	-48.659	-9.164	-5.902	1.00	38.27	B	N
ATOM	2809	C	ASN	439	-47.074	-9.617	-10.119	1.00	40.97	B	C
ATOM	2810	O	ASN	439	-47.017	-10.827	-9.904	1.00	42.65	B	O

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ATOM   2811  N   LEU   440     -46.245  -8.990 -10.949  1.00 42.11      B    N
ATOM   2812  CA  LEU   440     -45.272  -9.705 -11.774  1.00 41.16      B    C
ATOM   2813  CB  LEU   440     -45.709  -9.701 -13.239  1.00 40.54      B    C
ATOM   2814  CG  LEU   440     -47.068 -10.285 -13.613  1.00 42.30      B    C
ATOM   2815  CD1 LEU   440     -47.350  -9.994 -15.085  1.00 40.42      B    C
ATOM   2816  CD2 LEU   440     -47.070 -11.791 -13.350  1.00 43.28      B    C
ATOM   2817  C   LEU   440     -43.903  -9.043 -11.689  1.00 41.93      B    C
ATOM   2818  O   LEU   440     -43.789  -7.821 -11.825  1.00 41.74      B    O
ATOM   2819  N   VAL   441     -42.868  -9.848 -11.477  1.00 40.59      B    N
ATOM   2820  CA  VAL   441     -41.505  -9.363 -11.592  1.00 41.54      B    C
ATOM   2821  CB  VAL   441     -40.738  -9.535 -10.271  1.00 41.16      B    C
ATOM   2822  CG1 VAL   441     -39.296  -9.066 -10.442  1.00 38.34      B    C
ATOM   2823  CG2 VAL   441     -41.427  -8.732  -9.173  1.00 38.68      B    C
ATOM   2824  C   VAL   441     -40.778 -10.104 -12.710  1.00 43.69      B    C
ATOM   2825  O   VAL   441     -40.721 -11.335 -12.724  1.00 44.56      B    O
ATOM   2826  N   ALA   442     -40.224  -9.339 -13.644  1.00 43.06      B    N
ATOM   2827  CA  ALA   442     -39.686  -9.891 -14.878  1.00 43.65      B    C
ATOM   2828  CB  ALA   442     -39.105  -8.771 -15.740  1.00 41.79      B    C
ATOM   2829  C   ALA   442     -38.621 -10.945 -14.617  1.00 44.49      B    C
ATOM   2830  O   ALA   442     -37.872 -10.867 -13.645  1.00 44.26      B    O
ATOM   2831  N   ALA   443     -38.565 -11.933 -15.498  1.00 46.18      B    N
ATOM   2832  CA  ALA   443     -37.482 -12.903 -15.495  1.00 50.11      B    C
ATOM   2833  CB  ALA   443     -37.823 -14.070 -14.570  1.00 48.90      B    C
ATOM   2834  C   ALA   443     -37.282 -13.397 -16.921  1.00 52.32      B    C
ATOM   2835  O   ALA   443     -38.227 -13.430 -17.709  1.00 51.83      B    O
ATOM   2836  N   LEU   444     -36.049 -13.761 -17.253  1.00 56.33      B    N
ATOM   2837  CA  LEU   444     -35.764 -14.429 -18.520  1.00 61.28      B    C
ATOM   2838  CB  LEU   444     -34.263 -14.678 -18.656  1.00 61.00      B    C
ATOM   2839  CG  LEU   444     -33.435 -13.506 -19.170  1.00 63.12      B    C
ATOM   2840  CD1 LEU   444     -32.002 -13.614 -18.669  1.00 63.64      B    C
ATOM   2841  CD2 LEU   444     -33.493 -13.496 -20.692  1.00 64.35      B    C
ATOM   2842  C   LEU   444     -36.495 -15.761 -18.584  1.00 64.69      B    C
ATOM   2843  O   LEU   444     -36.498 -16.525 -17.619  1.00 64.31      B    O
ATOM   2844  N   PRO   445     -37.133 -16.057 -19.723  1.00 69.05      B    N
ATOM   2845  CD  PRO   445     -37.575 -15.168 -20.811  1.00 69.76      B    C
ATOM   2846  CA  PRO   445     -37.611 -17.430 -19.905  1.00 73.71      B    C
ATOM   2847  CB  PRO   445     -38.227 -17.410 -21.302  1.00 72.46      B    C
ATOM   2848  CG  PRO   445     -38.647 -15.982 -21.487  1.00 71.36      B    C
ATOM   2849  C   PRO   445     -36.445 -18.408 -19.793  1.00 78.03      B    C
ATOM   2850  O   PRO   445     -35.362 -18.161 -20.325  1.00 78.13      B    O
ATOM   2851  N   PRO   446     -36.651 -19.527 -19.080  1.00 82.32      B    N
ATOM   2852  CD  PRO   446     -37.943 -19.951 -18.510  1.00 83.90      B    C
ATOM   2853  CA  PRO   446     -35.597 -20.525 -18.854  1.00 85.07      B    C
ATOM   2854  CB  PRO   446     -36.232 -21.496 -17.861  1.00 84.72      B    C
ATOM   2855  CG  PRO   446     -37.698 -21.392 -18.128  1.00 84.66      B    C
ATOM   2856  C   PRO   446     -35.212 -21.212 -20.162  1.00 87.03      B    C
ATOM   2857  O   PRO   446     -34.532 -22.240 -20.165  1.00 87.51      B    O
ATOM   2858  N   SER   447     -35.651 -20.625 -21.271  1.00 89.12      B    N
ATOM   2859  CA  SER   447     -35.758 -21.351 -22.525  1.00 90.87      B    C
ATOM   2860  CB  SER   447     -37.050 -22.160 -22.518  1.00 91.88      B    C
ATOM   2861  OG  SER   447     -38.112 -21.370 -21.998  1.00 92.23      B    O
ATOM   2862  C   SER   447     -35.741 -20.452 -23.756  1.00 90.74      B    C
ATOM   2863  O   SER   447     -36.541 -19.521 -23.874  1.00 90.17      B    O
ATOM   2864  N   THR   448     -34.832 -20.753 -24.679  1.00 91.65      B    N
ATOM   2865  CA  THR   448     -34.918 -20.222 -26.032  1.00 94.19      B    C
ATOM   2866  CB  THR   448     -33.801 -20.786 -26.906  1.00 93.55      B    C
ATOM   2867  OG1 THR   448     -32.534 -20.327 -26.417  1.00 93.33      B    O
ATOM   2868  CG2 THR   448     -33.977 -20.355 -28.356  1.00 94.55      B    C
ATOM   2869  C   THR   448     -36.260 -20.634 -26.617  1.00 93.02      B    C
ATOM   2870  O   THR   448     -36.811 -19.945 -27.471  1.00 92.63      B    O
ATOM   2871  N   HIS   449     -36.766 -21.772 -26.150  1.00 95.06      B    N
ATOM   2872  CA  HIS   449     -38.137 -22.187 -26.413  1.00 97.82      B    C
ATOM   2873  CB  HIS   449     -38.453 -22.138 -27.911  1.00 99.76      B    C
ATOM   2874  CG  HIS   449     -39.916 -22.245 -28.218  1.00102.61      B    C
ATOM   2875  CD2 HIS   449     -41.006 -22.021 -27.445  1.00103.41      B    C
ATOM   2876  ND1 HIS   449     -40.394 -22.630 -29.452  1.00103.87      B    N
ATOM   2877  CE1 HIS   449     -41.715 -22.640 -29.427  1.00104.64      B    C
ATOM   2878  NE2 HIS   449     -42.112 -22.275 -28.221  1.00104.60      B    N
ATOM   2879  C   HIS   449     -38.379 -23.600 -25.893  1.00 97.73      B    C
ATOM   2880  O   HIS   449     -38.521 -23.812 -24.688  1.00 98.20      B    O
ATOM   2881  N   TRP   453     -45.204 -26.487 -27.136  1.00 95.56      B    N
ATOM   2882  CA  TRP   453     -46.152 -25.433 -27.487  1.00 95.17      B    C
ATOM   2883  CB  TRP   453     -46.794 -25.739 -28.844  1.00 99.16      B    C
ATOM   2884  CG  TRP   453     -47.498 -24.563 -29.442  1.00103.55      B    C
ATOM   2885  CD2 TRP   453     -48.830 -24.117 -29.151  1.00105.47      B    C
ATOM   2886  CE2 TRP   453     -49.061 -22.960 -29.923  1.00106.65      B    C
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ATOM   2887  CE3 TRP   453     -49.848 -24.582 -28.313  1.00106.05      B    C
ATOM   2888  CD1 TRP   453     -46.992 -23.683 -30.355  1.00105.12      B    C
ATOM   2889  NE1 TRP   453     -47.924 -22.716 -30.649  1.00106.63      B    N
ATOM   2890  CZ2 TRP   453     -50.268 -22.262 -29.881  1.00107.38      B    C
ATOM   2891  CZ3 TRP   453     -51.046 -23.887 -28.273  1.00107.05      B    C
ATOM   2892  CH2 TRP   453     -51.246 -22.740 -29.052  1.00107.25      B    C
ATOM   2893  C   TRP   453     -47.238 -25.302 -26.416  1.00 92.26      B    C
ATOM   2894  O   TRP   453     -47.614 -26.289 -25.779  1.00 92.72      B    O
ATOM   2895  N   GLN   454     -47.738 -24.083 -26.218  1.00 87.21      B    N
ATOM   2896  CA  GLN   454     -48.759 -23.836 -25.203  1.00 81.58      B    C
ATOM   2897  CB  GLN   454     -48.099 -23.699 -23.827  1.00 83.76      B    C
ATOM   2898  CG  GLN   454     -46.758 -22.984 -23.846  1.00 86.32      B    C
ATOM   2899  CD  GLN   454     -46.034 -23.072 -22.515  1.00 88.32      B    C
ATOM   2900  OE1 GLN   454     -44.803 -23.013 -22.458  1.00 88.83      B    O
ATOM   2901  NE2 GLN   454     -46.796 -23.215 -21.434  1.00 88.87      B    N
ATOM   2902  C   GLN   454     -49.641 -22.617 -25.486  1.00 76.01      B    C
ATOM   2903  O   GLN   454     -49.328 -21.790 -26.343  1.00 74.63      B    O
ATOM   2904  N   LEU   455     -50.752 -22.527 -24.759  1.00 69.41      B    N
ATOM   2905  CA  LEU   455     -51.741 -21.472 -24.956  1.00 63.14      B    C
ATOM   2906  CB  LEU   455     -53.148 -22.057 -24.850  1.00 60.88      B    C
ATOM   2907  CG  LEU   455     -54.309 -21.066 -24.893  1.00 60.03      B    C
ATOM   2908  CD1 LEU   455     -54.324 -20.347 -26.226  1.00 60.46      B    C
ATOM   2909  CD2 LEU   455     -55.611 -21.803 -24.681  1.00 59.51      B    C
ATOM   2910  C   LEU   455     -51.576 -20.354 -23.928  1.00 59.87      B    C
ATOM   2911  O   LEU   455     -51.792 -20.560 -22.735  1.00 58.28      B    O
ATOM   2912  N   PHE   456     -51.200 -19.169 -24.399  1.00 56.42      B    N
ATOM   2913  CA  PHE   456     -50.964 -18.031 -23.517  1.00 53.64      B    C
ATOM   2914  CB  PHE   456     -49.730 -17.251 -23.971  1.00 55.54      B    C
ATOM   2915  CG  PHE   456     -48.446 -18.011 -23.830  1.00 57.27      B    C
ATOM   2916  CD1 PHE   456     -48.030 -18.468 -22.594  1.00 58.30      B    C
ATOM   2917  CD2 PHE   456     -47.649 -18.260 -24.934  1.00 59.52      B    C
ATOM   2918  CE1 PHE   456     -46.839 -19.163 -22.456  1.00 60.51      B    C
ATOM   2919  CE2 PHE   456     -46.457 -18.954 -24.805  1.00 61.44      B    C
ATOM   2920  CZ  PHE   456     -46.052 -19.406 -23.561  1.00 60.76      B    C
ATOM   2921  C   PHE   456     -52.156 -17.093 -23.496  1.00 50.46      B    C
ATOM   2922  O   PHE   456     -52.654 -16.692 -24.547  1.00 50.85      B    O
ATOM   2923  N   CYS   457     -52.611 -16.746 -22.298  1.00 47.18      B    N
ATOM   2924  CA  CYS   457     -53.642 -15.726 -22.136  1.00 45.36      B    C
ATOM   2925  C   CYS   457     -53.203 -14.696 -21.102  1.00 44.05      B    C
ATOM   2926  O   CYS   457     -52.406 -15.003 -20.219  1.00 43.28      B    O
ATOM   2927  CB  CYS   457     -54.951 -16.347 -21.660  1.00 45.94      B    C
ATOM   2928  SG  CYS   457     -55.799 -17.516 -22.771  1.00 45.03      B    S
ATOM   2929  N   ARG   458     -53.741 -13.483 -21.209  1.00 42.97      B    N
ATOM   2930  CA  ARG   458     -53.467 -12.411 -20.249  1.00 42.03      B    C
ATOM   2931  CB  ARG   458     -52.447 -11.420 -20.827  1.00 40.47      B    C
ATOM   2932  CG  ARG   458     -52.907 -10.782 -22.127  1.00 42.86      B    C
ATOM   2933  CD  ARG   458     -51.781 -10.065 -22.866  1.00 44.13      B    C
ATOM   2934  NE  ARG   458     -51.841  -8.632 -22.622  1.00 49.28      B    N
ATOM   2935  CZ  ARG   458     -52.288  -7.728 -23.485  1.00 47.66      B    C
ATOM   2936  NH1 ARG   458     -52.301  -6.450 -23.135  1.00 50.19      B    N
ATOM   2937  NH2 ARG   458     -52.706  -8.086 -24.690  1.00 47.08      B    N
ATOM   2938  C   ARG   458     -54.770 -11.677 -19.950  1.00 41.18      B    C
ATOM   2939  O   ARG   458     -55.714 -11.721 -20.746  1.00 39.62      B    O
ATOM   2940  N   THR   459     -54.822 -11.004 -18.805  1.00 39.42      B    N
ATOM   2941  CA  THR   459     -55.998 -10.222 -18.453  1.00 39.74      B    C
ATOM   2942  CB  THR   459     -56.336 -10.376 -16.964  1.00 38.66      B    C
ATOM   2943  OG1 THR   459     -56.615 -11.750 -16.683  1.00 41.49      B    O
ATOM   2944  CG2 THR   459     -57.555  -9.544 -16.603  1.00 38.15      B    C
ATOM   2945  C   THR   459     -55.787  -8.743 -18.764  1.00 40.32      B    C
ATOM   2946  O   THR   459     -54.721  -8.189 -18.491  1.00 41.35      B    O
ATOM   2947  N   VAL   460     -56.806  -8.111 -19.336  1.00 39.71      B    N
ATOM   2948  CA  VAL   460     -56.747  -6.691 -19.670  1.00 40.13      B    C
ATOM   2949  CB  VAL   460     -56.849  -6.476 -21.193  1.00 40.81      B    C
ATOM   2950  CG1 VAL   460     -56.918  -4.986 -21.512  1.00 39.94      B    C
ATOM   2951  CG2 VAL   460     -55.657  -7.121 -21.885  1.00 40.01      B    C
ATOM   2952  C   VAL   460     -57.896  -5.940 -19.007  1.00 41.10      B    C
ATOM   2953  O   VAL   460     -59.059  -6.148 -19.350  1.00 41.34      B    O
ATOM   2954  N   TRP   461     -57.577  -5.066 -18.058  1.00 40.15      B    N
ATOM   2955  CA  TRP   461     -58.606  -4.243 -17.438  1.00 41.32      B    C
ATOM   2956  CB  TRP   461     -58.210  -3.864 -16.013  1.00 40.29      B    C
ATOM   2957  CG  TRP   461     -58.327  -4.982 -15.025  1.00 39.72      B    C
ATOM   2958  CD2 TRP   461     -59.316  -5.113 -13.997  1.00 38.69      B    C
ATOM   2959  CE2 TRP   461     -59.013  -6.284 -13.272  1.00 40.64      B    C
ATOM   2960  CE3 TRP   461     -60.426  -4.353 -13.618  1.00 37.91      B    C
ATOM   2961  CD1 TRP   461     -57.490  -6.053 -14.891  1.00 39.19      B    C
ATOM   2962  NE1 TRP   461     -57.894  -6.840 -13.836  1.00 40.77      B    N
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ATOM   2963  CZ2 TRP   461     -59.783  -6.713 -12.191  1.00 39.89      B    C
ATOM   2964  CZ3 TRP   461     -61.189  -4.779 -12.543  1.00 39.15      B    C
ATOM   2965  CH2 TRP   461     -60.863  -5.949 -11.841  1.00 39.06      B    C
ATOM   2966  C   TRP   461     -58.835  -2.974 -18.246  1.00 42.82      B    C
ATOM   2967  O   TRP   461     -57.895  -2.381 -18.774  1.00 42.86      B    O
ATOM   2968  N   SER   462     -60.091  -2.558 -18.339  1.00 44.55      B    N
ATOM   2969  CA  SER   462     -60.421  -1.295 -18.981  1.00 45.47      B    C
ATOM   2970  CB  SER   462     -61.849  -1.328 -19.521  1.00 45.66      B    C
ATOM   2971  OG  SER   462     -62.779  -1.265 -18.452  1.00 43.37      B    O
ATOM   2972  C   SER   462     -60.310  -0.175 -17.956  1.00 48.34      B    C
ATOM   2973  O   SER   462     -60.104  -0.422 -16.764  1.00 46.67      B    O
ATOM   2974  N   ALA   463     -60.460   1.057 -18.429  1.00 50.60      B    N
ATOM   2975  CA  ALA   463     -60.634   2.192 -17.537  1.00 54.46      B    C
ATOM   2976  CB  ALA   463     -60.531   3.493 -18.325  1.00 52.98      B    C
ATOM   2977  C   ALA   463     -61.998   2.095 -16.860  1.00 56.97      B    C
ATOM   2978  O   ALA   463     -62.918   1.478 -17.396  1.00 57.30      B    O
ATOM   2979  N   HIS   464     -62.118   2.704 -15.683  1.00 61.61      B    N
ATOM   2980  CA  HIS   464     -63.404   2.863 -15.000  1.00 65.67      B    C
ATOM   2981  CB  HIS   464     -63.204   3.685 -13.725  1.00 68.34      B    C
ATOM   2982  CG  HIS   464     -64.295   3.520 -12.715  1.00 71.76      B    C
ATOM   2983  CD2 HIS   464     -65.503   4.121 -12.599  1.00 73.40      B    C
ATOM   2984  ND1 HIS   464     -64.191   2.658 -11.644  1.00 73.71      B    N
ATOM   2985  CE1 HIS   464     -65.287   2.736 -10.910  1.00 74.59      B    C
ATOM   2986  NE2 HIS   464     -66.100   3.616 -11.468  1.00 75.04      B    N
ATOM   2987  C   HIS   464     -64.389   3.581 -15.926  1.00 67.55      B    C
ATOM   2988  O   HIS   464     -63.996   4.455 -16.697  1.00 67.46      B    O
ATOM   2989  N   SER   465     -65.667   3.220 -15.847  1.00 70.08      B    N
ATOM   2990  CA  SER   465     -66.655   3.714 -16.806  1.00 72.95      B    C
ATOM   2991  CB  SER   465     -67.866   2.783 -16.852  1.00 71.92      B    C
ATOM   2992  OG  SER   465     -68.608   2.866 -15.648  1.00 70.23      B    O
ATOM   2993  C   SER   465     -67.141   5.122 -16.487  1.00 75.78      B    C
ATOM   2994  O   SER   465     -67.431   5.905 -17.393  1.00 75.65      B    O
ATOM   2995  N   GLY   466     -67.240   5.436 -15.200  1.00 78.79      B    N
ATOM   2996  CA  GLY   466     -67.851   6.689 -14.794  1.00 83.44      B    C
ATOM   2997  C   GLY   466     -69.263   6.487 -14.272  1.00 86.66      B    C
ATOM   2998  O   GLY   466     -69.931   5.523 -14.649  1.00 87.19      B    O
ATOM   2999  N   PRO   467     -69.744   7.388 -13.400  1.00 89.00      B    N
ATOM   3000  CD  PRO   467     -68.954   8.535 -12.924  1.00 89.56      B    C
ATOM   3001  CA  PRO   467     -71.005   7.256 -12.655  1.00 90.53      B    C
ATOM   3002  CB  PRO   467     -70.953   8.409 -11.649  1.00 90.49      B    C
ATOM   3003  CG  PRO   467     -69.503   8.764 -11.549  1.00 90.14      B    C
ATOM   3004  C   PRO   467     -72.283   7.299 -13.496  1.00 91.90      B    C
ATOM   3005  O   PRO   467     -73.355   6.926 -13.016  1.00 92.02      B    O
ATOM   3006  N   THR   468     -72.171   7.755 -14.741  1.00 93.44      B    N
ATOM   3007  CA  THR   468     -73.337   7.899 -15.612  1.00 95.40      B    C
ATOM   3008  CB  THR   468     -72.914   8.237 -17.059  1.00 95.23      B    C
ATOM   3009  OG1 THR   468     -72.106   9.420 -17.060  1.00 95.12      B    O
ATOM   3010  CG2 THR   468     -74.140   8.468 -17.934  1.00 94.69      B    C
ATOM   3011  C   THR   468     -74.178   6.621 -15.635  1.00 97.16      B    C
ATOM   3012  O   THR   468     -73.651   5.514 -15.509  1.00 97.49      B    O
ATOM   3013  N   ARG   469     -75.488   6.780 -15.798  1.00 98.73      B    N
ATOM   3014  CA  ARG   469     -76.407   5.646 -15.781  1.00 99.90      B    C
ATOM   3015  CB  ARG   469     -77.854   6.143 -15.688  1.00102.98      B    C
ATOM   3016  CG  ARG   469     -78.326   6.922 -16.911  1.00106.29      B    C
ATOM   3017  CD  ARG   469     -79.787   7.337 -16.781  1.00108.83      B    C
ATOM   3018  NE  ARG   469     -80.366   7.714 -18.069  1.00111.09      B    N
ATOM   3019  CZ  ARG   469     -81.164   6.930 -18.789  1.00112.01      B    C
ATOM   3020  NH1 ARG   469     -81.644   7.356 -19.951  1.00112.06      B    N
ATOM   3021  NH2 ARG   469     -81.486   5.722 -18.345  1.00112.44      B    N
ATOM   3022  C   ARG   469     -76.243   4.786 -17.032  1.00 98.96      B    C
ATOM   3023  O   ARG   469     -76.478   3.578 -17.002  1.00 98.89      B    O
ATOM   3024  N   MET   470     -75.841   5.422 -18.128  1.00 97.91      B    N
ATOM   3025  CA  MET   470     -75.657   4.732 -19.401  1.00 96.48      B    C
ATOM   3026  CB  MET   470     -76.405   5.475 -20.515  1.00 98.76      B    C
ATOM   3027  CG  MET   470     -77.923   5.370 -20.432  1.00101.49      B    C
ATOM   3028  SD  MET   470     -78.555   3.783 -21.034  1.00104.46      B    S
ATOM   3029  CE  MET   470     -78.069   3.857 -22.774  1.00102.90      B    C
ATOM   3030  C   MET   470     -74.179   4.616 -19.770  1.00 94.05      B    C
ATOM   3031  O   MET   470     -73.837   4.427 -20.938  1.00 93.54      B    O
ATOM   3032  N   ALA   471     -73.308   4.732 -18.771  1.00 90.93      B    N
ATOM   3033  CA  ALA   471     -71.867   4.671 -18.998  1.00 87.61      B    C
ATOM   3034  CB  ALA   471     -71.123   5.251 -17.800  1.00 87.53      B    C
ATOM   3035  C   ALA   471     -71.406   3.239 -19.251  1.00 85.13      B    C
ATOM   3036  O   ALA   471     -71.870   2.300 -18.598  1.00 85.32      B    O
ATOM   3037  N   THR   472     -70.491   3.080 -20.203  1.00 81.39      B    N
ATOM   3038  CA  THR   472     -69.941   1.769 -20.528  1.00 77.96      B    C
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ATOM   3039  CB  THR   472     -70.443   1.275 -21.911  1.00 78.06      B    C
ATOM   3040  OG1 THR   472     -70.026   2.188 -22.933  1.00 77.52      B    O
ATOM   3041  CG2 THR   472     -71.962   1.176 -21.921  1.00 77.75      B    C
ATOM   3042  C   THR   472     -68.412   1.784 -20.539  1.00 75.20      B    C
ATOM   3043  O   THR   472     -67.789   2.776 -20.922  1.00 74.74      B    O
ATOM   3044  N   ALA   473     -67.813   0.676 -20.113  1.00 71.92      B    N
ATOM   3045  CA  ALA   473     -66.366   0.503 -20.194  1.00 68.57      B    C
ATOM   3046  CB  ALA   473     -65.827   0.020 -18.858  1.00 67.75      B    C
ATOM   3047  C   ALA   473     -66.020  -0.497 -21.297  1.00 66.77      B    C
ATOM   3048  O   ALA   473     -66.775  -1.433 -21.554  1.00 66.07      B    O
ATOM   3049  N   ILE   474     -64.879  -0.293 -21.947  1.00 64.89      B    N
ATOM   3050  CA  ILE   474     -64.443  -1.170 -23.025  1.00 63.32      B    C
ATOM   3051  CB  ILE   474     -64.477  -0.437 -24.369  1.00 64.64      B    C
ATOM   3052  CG2 ILE   474     -64.014  -1.361 -25.482  1.00 63.94      B    C
ATOM   3053  CG1 ILE   474     -65.896   0.056 -24.652  1.00 67.76      B    C
ATOM   3054  CD1 ILE   474     -65.983   0.995 -25.843  1.00 71.25      B    C
ATOM   3055  C   ILE   474     -63.026  -1.685 -22.801  1.00 61.20      B    C
ATOM   3056  O   ILE   474     -62.100  -0.907 -22.575  1.00 62.29      B    O
ATOM   3057  N   ALA   475     -62.864  -3.002 -22.866  1.00 57.61      B    N
ATOM   3058  CA  ALA   475     -61.548  -3.619 -22.783  1.00 54.56      B    C
ATOM   3059  CB  ALA   475     -61.525  -4.641 -21.662  1.00 53.14      B    C
ATOM   3060  C   ALA   475     -61.222  -4.284 -24.113  1.00 53.19      B    C
ATOM   3061  O   ALA   475     -62.025  -5.050 -24.643  1.00 52.48      B    O
ATOM   3062  N   ARG   476     -60.046  -3.984 -24.655  1.00 52.78      B    N
ATOM   3063  CA  ARG   476     -59.651  -4.501 -25.960  1.00 52.83      B    C
ATOM   3064  CB  ARG   476     -59.378  -3.352 -26.936  1.00 55.93      B    C
ATOM   3065  CG  ARG   476     -60.608  -2.550 -27.326  1.00 61.00      B    C
ATOM   3066  CD  ARG   476     -60.279  -1.499 -28.382  1.00 64.35      B    C
ATOM   3067  NE  ARG   476     -61.444  -0.677 -28.700  1.00 67.87      B    N
ATOM   3068  CZ  ARG   476     -61.840   0.369 -27.978  1.00 70.61      B    C
ATOM   3069  NH1 ARG   476     -61.160   0.726 -26.894  1.00 70.19      B    N
ATOM   3070  NH2 ARG   476     -62.925   1.048 -28.331  1.00 71.97      B    N
ATOM   3071  C   ARG   476     -58.411  -5.376 -25.873  1.00 51.34      B    C
ATOM   3072  O   ARG   476     -57.620  -5.264 -24.938  1.00 51.28      B    O
ATOM   3073  N   CYS   477     -58.247  -6.244 -26.863  1.00 50.36      B    N
ATOM   3074  CA  CYS   477     -57.048  -7.057 -26.983  1.00 50.00      B    C
ATOM   3075  C   CYS   477     -56.135  -6.474 -28.053  1.00 49.37      B    C
ATOM   3076  O   CYS   477     -56.556  -5.638 -28.852  1.00 48.36      B    O
ATOM   3077  CB  CYS   477     -57.416  -8.495 -27.358  1.00 51.30      B    C
ATOM   3078  SG  CYS   477     -58.397  -9.398 -26.112  1.00 54.92      B    S
ATOM   3079  N   ALA   478     -54.883  -6.917 -28.065  1.00 49.69      B    N
ATOM   3080  CA  ALA   478     -53.979  -6.597 -29.163  1.00 52.29      B    C
ATOM   3081  CB  ALA   478     -52.579  -7.101 -28.852  1.00 50.02      B    C
ATOM   3082  C   ALA   478     -54.502  -7.248 -30.443  1.00 54.47      B    C
ATOM   3083  O   ALA   478     -55.233  -8.239 -30.394  1.00 54.91      B    O
ATOM   3084  N   PRO   479     -54.131  -6.695 -31.608  1.00 56.47      B    N
ATOM   3085  CD  PRO   479     -53.268  -5.506 -31.723  1.00 56.66      B    C
ATOM   3086  CA  PRO   479     -54.627  -7.142 -32.919  1.00 56.71      B    C
ATOM   3087  CB  PRO   479     -53.914  -6.215 -33.905  1.00 57.17      B    C
ATOM   3088  CG  PRO   479     -53.584  -4.994 -33.097  1.00 57.72      B    C
ATOM   3089  C   PRO   479     -54.370  -8.621 -33.235  1.00 56.70      B    C
ATOM   3090  O   PRO   479     -55.207  -9.283 -33.849  1.00 56.36      B    O
ATOM   3091  N   ASP   480     -53.217  -9.137 -32.822  1.00 56.51      B    N
ATOM   3092  CA  ASP   480     -52.884 -10.536 -33.077  1.00 56.90      B    C
ATOM   3093  CB  ASP   480     -51.368 -10.737 -33.009  1.00 60.77      B    C
ATOM   3094  CG  ASP   480     -50.655 -10.216 -34.244  1.00 65.51      B    C
ATOM   3095  OD1 ASP   480     -51.324  -9.580 -35.090  1.00 67.40      B    O
ATOM   3096  OD2 ASP   480     -49.430 -10.448 -34.372  1.00 67.85      B    O
ATOM   3097  C   ASP   480     -53.566 -11.511 -32.113  1.00 55.43      B    C
ATOM   3098  O   ASP   480     -53.440 -12.727 -32.262  1.00 55.84      B    O
ATOM   3099  N   GLU   481     -54.281 -10.977 -31.125  1.00 52.35      B    N
ATOM   3100  CA  GLU   481     -54.880 -11.795 -30.074  1.00 50.24      B    C
ATOM   3101  CB  GLU   481     -54.600 -11.134 -28.704  1.00 47.61      B    C
ATOM   3102  CG  GLU   481     -53.231 -11.099 -28.221  1.00 46.45      B    C
ATOM   3103  CD  GLU   481     -53.058 -10.302 -26.934  1.00 44.57      B    C
ATOM   3104  OE1 GLU   481     -52.173 -10.651 -26.123  1.00 42.50      B    O
ATOM   3105  OE2 GLU   481     -53.811  -9.326 -26.737  1.00 43.51      B    O
ATOM   3106  C   GLU   481     -56.370 -12.031 -30.306  1.00 49.74      B    C
ATOM   3107  O   GLU   481     -57.041 -11.239 -30.967  1.00 49.49      B    O
ATOM   3108  N   GLU   482     -56.879 -13.130 -29.759  1.00 48.52      B    N
ATOM   3109  CA  GLU   482     -58.315 -13.385 -29.740  1.00 46.82      B    C
ATOM   3110  CB  GLU   482     -58.594 -14.859 -30.040  1.00 46.77      B    C
ATOM   3111  CG  GLU   482     -57.890 -15.387 -31.271  1.00 49.47      B    C
ATOM   3112  CD  GLU   482     -58.568 -14.957 -32.554  1.00 51.58      B    C
ATOM   3113  OE1 GLU   482     -59.695 -14.418 -32.472  1.00 50.06      B    O
ATOM   3114  OE2 GLU   482     -57.975 -15.162 -33.638  1.00 54.21      B    O
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ATOM	3115	C	GLU	482	-58.851	-13.050	-28.356	1.00	45.01	B	C
ATOM	3116	O	GLU	482	-58.211	-13.351	-27.352	1.00	44.86	B	O
ATOM	3117	N	LEU	483	-60.024	-12.433	-28.301	1.00	44.07	B	N
ATOM	3118	CA	LEU	483	-60.709	-12.256	-27.030	1.00	45.33	B	C
ATOM	3119	CB	LEU	483	-61.583	-11.003	-27.075	1.00	46.54	B	C
ATOM	3120	CG	LEU	483	-62.243	-10.640	-25.742	1.00	49.38	B	C
ATOM	3121	CD1	LEU	483	-62.160	-9.140	-25.513	1.00	52.67	B	C
ATOM	3122	CD2	LEU	483	-63.684	-11.100	-25.745	1.00	51.11	B	C
ATOM	3123	C	LEU	483	-61.570	-13.484	-26.731	1.00	44.81	B	C
ATOM	3124	O	LEU	483	-62.583	-13.710	-27.390	1.00	44.85	B	O
ATOM	3125	N	LEU	484	-61.165	-14.277	-25.741	1.00	44.25	B	N
ATOM	3126	CA	LEU	484	-61.836	-15.545	-25.459	1.00	43.90	B	C
ATOM	3127	CB	LEU	484	-60.820	-16.618	-25.064	1.00	41.25	B	C
ATOM	3128	CG	LEU	484	-59.820	-17.008	-26.151	1.00	40.18	B	C
ATOM	3129	CD1	LEU	484	-59.149	-18.324	-25.797	1.00	38.70	B	C
ATOM	3130	CD2	LEU	484	-60.544	-17.131	-27.465	1.00	40.84	B	C
ATOM	3131	C	LEU	484	-62.908	-15.444	-24.380	1.00	44.90	B	C
ATOM	3132	O	LEU	484	-63.754	-16.326	-24.262	1.00	46.30	B	O
ATOM	3133	N	SER	485	-62.873	-14.380	-23.587	1.00	46.22	B	N
ATOM	3134	CA	SER	485	-63.958	-14.123	-22.647	1.00	46.47	B	C
ATOM	3135	CB	SER	485	-63.866	-15.065	-21.450	1.00	46.34	B	C
ATOM	3136	OG	SER	485	-62.827	-14.667	-20.580	1.00	45.49	B	O
ATOM	3137	C	SER	485	-63.972	-12.684	-22.152	1.00	46.84	B	C
ATOM	3138	O	SER	485	-63.101	-11.886	-22.486	1.00	47.34	B	O
ATOM	3139	N	CYS	486	-64.968	-12.373	-21.333	1.00	48.22	B	N
ATOM	3140	CA	CYS	486	-65.299	-10.997	-21.000	1.00	49.92	B	C
ATOM	3141	C	CYS	486	-66.118	-10.992	-19.719	1.00	49.99	B	C
ATOM	3142	O	CYS	486	-67.213	-11.557	-19.675	1.00	50.31	B	O
ATOM	3143	CB	CYS	486	-66.103	-10.386	-22.151	1.00	52.24	B	C
ATOM	3144	SG	CYS	486	-66.813	-8.726	-21.890	1.00	59.49	B	S
ATOM	3145	N	SER	487	-65.581	-10.372	-18.672	1.00	49.69	B	N
ATOM	3146	CA	SER	487	-66.331	-10.198	-17.431	1.00	49.46	B	C
ATOM	3147	CB	SER	487	-65.750	-11.083	-16.328	1.00	48.38	B	C
ATOM	3148	OG	SER	487	-64.402	-10.754	-16.063	1.00	49.39	B	O
ATOM	3149	C	SER	487	-66.328	-8.736	-16.989	1.00	49.59	B	C
ATOM	3150	O	SER	487	-65.852	-7.863	-17.713	1.00	49.27	B	O
ATOM	3151	N	SER	488	-66.872	-8.470	-15.807	1.00	49.56	B	N
ATOM	3152	CA	SER	488	-66.998	-7.100	-15.333	1.00	51.42	B	C
ATOM	3153	CB	SER	488	-68.295	-6.485	-15.853	1.00	50.06	B	C
ATOM	3154	OG	SER	488	-69.420	-7.186	-15.358	1.00	50.85	B	O
ATOM	3155	C	SER	488	-66.963	-7.035	-13.814	1.00	53.00	B	C
ATOM	3156	O	SER	488	-67.233	-8.022	-13.135	1.00	54.35	B	O
ATOM	3157	N	PHE	489	-66.614	-5.869	-13.284	1.00	55.50	B	N
ATOM	3158	CA	PHE	489	-66.478	-5.700	-11.842	1.00	58.01	B	C
ATOM	3159	CB	PHE	489	-65.049	-6.037	-11.405	1.00	56.05	B	C
ATOM	3160	CG	PHE	489	-64.765	-5.733	-9.959	1.00	56.36	B	C
ATOM	3161	CD1	PHE	489	-65.412	-6.426	-8.949	1.00	56.87	B	C
ATOM	3162	CD2	PHE	489	-63.842	-4.759	-9.611	1.00	55.96	B	C
ATOM	3163	CE1	PHE	489	-65.145	-6.156	-7.615	1.00	56.29	B	C
ATOM	3164	CE2	PHE	489	-63.570	-4.483	-8.281	1.00	55.63	B	C
ATOM	3165	CZ	PHE	489	-64.223	-5.183	-7.281	1.00	55.87	B	C
ATOM	3166	C	PHE	489	-66.829	-4.280	-11.405	1.00	60.88	B	C
ATOM	3167	O	PHE	489	-66.417	-3.298	-12.028	1.00	60.67	B	O
ATOM	3168	N	SER	490	-67.604	-4.188	-10.330	1.00	64.50	B	N
ATOM	3169	CA	SER	490	-67.931	-2.913	-9.704	1.00	68.24	B	C
ATOM	3170	CB	SER	490	-69.370	-2.513	-10.038	1.00	68.91	B	C
ATOM	3171	OG	SER	490	-69.721	-1.289	-9.418	1.00	72.44	B	O
ATOM	3172	C	SER	490	-67.779	-3.094	-8.200	1.00	70.33	B	C
ATOM	3173	O	SER	490	-68.275	-4.068	-7.636	1.00	70.86	B	O
ATOM	3174	N	ARG	491	-67.085	-2.167	-7.549	1.00	72.98	B	N
ATOM	3175	CA	ARG	491	-66.842	-2.300	-6.118	1.00	75.53	B	C
ATOM	3176	CB	ARG	491	-65.779	-1.295	-5.666	1.00	77.17	B	C
ATOM	3177	CG	ARG	491	-64.422	-1.536	-6.308	1.00	80.24	B	C
ATOM	3178	CD	ARG	491	-63.325	-0.685	-5.687	1.00	82.34	B	C
ATOM	3179	NE	ARG	491	-62.027	-0.962	-6.299	1.00	83.86	B	N
ATOM	3180	CZ	ARG	491	-61.603	-0.412	-7.434	1.00	84.91	B	C
ATOM	3181	NH1	ARG	491	-60.408	-0.725	-7.917	1.00	84.85	B	N
ATOM	3182	NH2	ARG	491	-62.373	0.453	-8.086	1.00	84.91	B	N
ATOM	3183	C	ARG	491	-68.133	-2.099	-5.334	1.00	76.00	B	C
ATOM	3184	O	ARG	491	-68.302	-2.649	-4.248	1.00	75.57	B	O
ATOM	3185	N	SER	492	-69.047	-1.317	-5.901	1.00	77.48	B	N
ATOM	3186	CA	SER	492	-70.354	-1.096	-5.290	1.00	79.37	B	C
ATOM	3187	CB	SER	492	-71.028	0.129	-5.911	1.00	79.14	B	C
ATOM	3188	OG	SER	492	-71.317	-0.090	-7.282	1.00	79.09	B	O
ATOM	3189	C	SER	492	-71.240	-2.321	-5.494	1.00	80.75	B	C
ATOM	3190	O	SER	492	-72.065	-2.652	-4.642	1.00	81.44	B	O

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ATOM   3191  N   GLY   493     -71.062  -2.988  -6.631  1.00 81.13      B    N
ATOM   3192  CA  GLY   493     -71.864  -4.157  -6.939  1.00 80.72      B    C
ATOM   3193  C   GLY   493     -73.023  -3.825  -7.858  1.00 80.78      B    C
ATOM   3194  O   GLY   493     -73.689  -4.719  -8.378  1.00 80.89      B    O
ATOM   3195  N   LYS   494     -73.263  -2.533  -8.060  1.00 80.45      B    N
ATOM   3196  CA  LYS   494     -74.352  -2.082  -8.917  1.00 79.64      B    C
ATOM   3197  CB  LYS   494     -74.821  -0.690  -8.484  1.00 81.16      B    C
ATOM   3198  CG  LYS   494     -75.389  -0.633  -7.071  1.00 83.07      B    C
ATOM   3199  CD  LYS   494     -75.838   0.779  -6.712  1.00 84.54      B    C
ATOM   3200  CE  LYS   494     -76.744   0.789  -5.483  1.00 85.65      B    C
ATOM   3201  NZ  LYS   494     -76.058   0.292  -4.255  1.00 85.21      B    N
ATOM   3202  C   LYS   494     -73.900  -2.040 -10.369  1.00 78.44      B    C
ATOM   3203  O   LYS   494     -73.266  -1.081 -10.805  1.00 78.70      B    O
ATOM   3204  N   ARG   495     -74.232  -3.087 -11.115  1.00 77.53      B    N
ATOM   3205  CA  ARG   495     -73.848  -3.179 -12.519  1.00 76.32      B    C
ATOM   3206  CB  ARG   495     -72.525  -3.933 -12.655  1.00 76.14      B    C
ATOM   3207  CG  ARG   495     -72.587  -5.346 -12.128  1.00 75.82      B    C
ATOM   3208  CD  ARG   495     -71.211  -5.965 -12.011  1.00 76.68      B    C
ATOM   3209  NE  ARG   495     -71.292  -7.320 -11.477  1.00 77.74      B    N
ATOM   3210  CZ  ARG   495     -71.266  -8.420 -12.223  1.00 78.22      B    C
ATOM   3211  NH1 ARG   495     -71.350  -9.613 -11.649  1.00 78.62      B    N
ATOM   3212  NH2 ARG   495     -71.150  -8.330 -13.541  1.00 78.31      B    N
ATOM   3213  C   ARG   495     -74.930  -3.909 -13.297  1.00 75.33      B    C
ATOM   3214  O   ARG   495     -75.787  -4.564 -12.708  1.00 75.07      B    O
ATOM   3215  N   ARG   496     -74.890  -3.795 -14.619  1.00 74.53      B    N
ATOM   3216  CA  ARG   496     -75.827  -4.522 -15.464  1.00 74.26      B    C
ATOM   3217  CB  ARG   496     -76.653  -3.540 -16.299  1.00 75.26      B    C
ATOM   3218  CG  ARG   496     -77.599  -2.676 -15.472  1.00 77.17      B    C
ATOM   3219  CD  ARG   496     -78.601  -1.945 -16.354  1.00 78.98      B    C
ATOM   3220  NE  ARG   496     -77.937  -1.117 -17.358  1.00 82.03      B    N
ATOM   3221  CZ  ARG   496     -78.553  -0.564 -18.398  1.00 83.71      B    C
ATOM   3222  NH1 ARG   496     -79.855  -0.747 -18.580  1.00 84.51      B    N
ATOM   3223  NH2 ARG   496     -77.864   0.174 -19.260  1.00 84.84      B    N
ATOM   3224  C   ARG   496     -75.124  -5.533 -16.374  1.00 73.26      B    C
ATOM   3225  O   ARG   496     -75.604  -5.837 -17.467  1.00 72.51      B    O
ATOM   3226  N   GLY   497     -73.985  -6.049 -15.917  1.00 71.89      B    N
ATOM   3227  CA  GLY   497     -73.320  -7.124 -16.632  1.00 70.45      B    C
ATOM   3228  C   GLY   497     -72.398  -6.668 -17.747  1.00 69.59      B    C
ATOM   3229  O   GLY   497     -71.964  -5.517 -17.776  1.00 69.36      B    O
ATOM   3230  N   GLU   498     -72.098  -7.578 -18.670  1.00 68.77      B    N
ATOM   3231  CA  GLU   498     -71.173  -7.288 -19.760  1.00 68.48      B    C
ATOM   3232  CB  GLU   498     -69.744  -7.653 -19.350  1.00 67.95      B    C
ATOM   3233  CG  GLU   498     -69.474  -9.150 -19.254  1.00 67.56      B    C
ATOM   3234  CD  GLU   498     -70.106  -9.795 -18.030  1.00 68.08      B    C
ATOM   3235  OE1 GLU   498     -69.782  -9.383 -16.895  1.00 67.57      B    O
ATOM   3236  OE2 GLU   498     -70.927 -10.720 -18.202  1.00 69.43      B    O
ATOM   3237  C   GLU   498     -71.550  -8.052 -21.024  1.00 68.67      B    C
ATOM   3238  O   GLU   498     -72.284  -9.037 -20.968  1.00 67.70      B    O
ATOM   3239  N   ARG   499     -71.046  -7.590 -22.162  1.00 69.43      B    N
ATOM   3240  CA  ARG   499     -71.295  -8.254 -23.434  1.00 72.09      B    C
ATOM   3241  CB  ARG   499     -72.483  -7.606 -24.149  1.00 74.63      B    C
ATOM   3242  CG  ARG   499     -72.297  -6.129 -24.460  1.00 79.98      B    C
ATOM   3243  CD  ARG   499     -73.461  -5.591 -25.277  1.00 83.74      B    C
ATOM   3244  NE  ARG   499     -73.573  -6.288 -26.557  1.00 88.28      B    N
ATOM   3245  CZ  ARG   499     -74.571  -6.120 -27.420  1.00 90.00      B    C
ATOM   3246  NH1 ARG   499     -74.582  -6.802 -28.559  1.00 90.05      B    N
ATOM   3247  NH2 ARG   499     -75.558  -5.275 -27.142  1.00 90.84      B    N
ATOM   3248  C   ARG   499     -70.065  -8.186 -24.329  1.00 72.19      B    C
ATOM   3249  O   ARG   499     -69.243  -7.284 -24.200  1.00 71.77      B    O
ATOM   3250  N   MET   500     -69.939  -9.151 -25.232  1.00 73.12      B    N
ATOM   3251  CA  MET   500     -68.895  -9.109 -26.245  1.00 74.86      B    C
ATOM   3252  CB  MET   500     -68.290 -10.498 -26.439  1.00 75.04      B    C
ATOM   3253  CG  MET   500     -67.541 -11.022 -25.231  1.00 75.98      B    C
ATOM   3254  SD  MET   500     -67.019 -12.735 -25.437  1.00 77.64      B    S
ATOM   3255  CE  MET   500     -68.472 -13.608 -24.783  1.00 75.19      B    C
ATOM   3256  C   MET   500     -69.471  -8.611 -27.565  1.00 75.89      B    C
ATOM   3257  O   MET   500     -70.349  -9.244 -28.147  1.00 75.73      B    O
ATOM   3258  N   GLU   501     -68.971  -7.471 -28.029  1.00 77.75      B    N
ATOM   3259  CA  GLU   501     -69.475  -6.838 -29.242  1.00 79.49      B    C
ATOM   3260  CB  GLU   501     -69.823  -5.372 -28.975  1.00 80.04      B    C
ATOM   3261  CG  GLU   501     -70.879  -5.151 -27.908  1.00 82.37      B    C
ATOM   3262  CD  GLU   501     -71.193  -3.677 -27.702  1.00 83.60      B    C
ATOM   3263  OE1 GLU   501     -70.530  -2.832 -28.343  1.00 83.92      B    O
ATOM   3264  OE2 GLU   501     -72.102  -3.367 -26.901  1.00 84.18      B    O
ATOM   3265  C   GLU   501     -68.432  -6.897 -30.343  1.00 80.62      B    C
ATOM   3266  O   GLU   501     -67.253  -7.122 -30.080  1.00 81.37      B    O
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ATOM	3267	N	ALA	502	-68.871	-6.685	-31.577	1.00	81.89	B N
ATOM	3268	CA	ALA	502	-67.950	-6.483	-32.684	1.00	83.20	B C
ATOM	3269	CB	ALA	502	-68.497	-7.137	-33.942	1.00	83.17	B C
ATOM	3270	C	ALA	502	-67.739	-4.987	-32.918	1.00	84.38	B C
ATOM	3271	O	ALA	502	-68.689	-4.202	-32.889	1.00	84.50	B O
ATOM	3272	N	GLN	503	-66.485	-4.601	-33.133	1.00	85.34	B N
ATOM	3273	CA	GLN	503	-66.143	-3.237	-33.521	1.00	85.73	B C
ATOM	3274	CB	GLN	503	-65.807	-2.394	-32.291	1.00	85.79	B C
ATOM	3275	CG	GLN	503	-66.989	-2.127	-31.379	1.00	87.34	B C
ATOM	3276	CD	GLN	503	-66.707	-1.020	-30.381	1.00	88.20	B C
ATOM	3277	OE1	GLN	503	-65.587	-0.508	-30.304	1.00	88.07	B O
ATOM	3278	NE2	GLN	503	-67.723	-0.644	-29.610	1.00	88.12	B N
ATOM	3279	C	GLN	503	-64.941	-3.268	-34.456	1.00	86.14	B C
ATOM	3280	O	GLN	503	-63.865	-3.733	-34.081	1.00	86.49	B O
ATOM	3281	N	GLY	504	-65.127	-2.775	-35.675	1.00	85.95	B N
ATOM	3282	CA	GLY	504	-64.066	-2.858	-36.659	1.00	84.92	B C
ATOM	3283	C	GLY	504	-63.668	-4.298	-36.918	1.00	84.27	B C
ATOM	3284	O	GLY	504	-62.487	-4.601	-37.096	1.00	84.30	B O
ATOM	3285	N	GLY	505	-64.652	-5.192	-36.929	1.00	83.46	B N
ATOM	3286	CA	GLY	505	-64.376	-6.587	-37.221	1.00	82.44	B C
ATOM	3287	C	GLY	505	-63.505	-7.254	-36.172	1.00	81.46	B C
ATOM	3288	O	GLY	505	-62.900	-8.296	-36.426	1.00	81.89	B O
ATOM	3289	N	LYS	506	-63.434	-6.645	-34.992	1.00	79.89	B N
ATOM	3290	CA	LYS	506	-62.694	-7.214	-33.870	1.00	77.91	B C
ATOM	3291	CB	LYS	506	-61.499	-6.320	-33.519	1.00	79.66	B C
ATOM	3292	CG	LYS	506	-60.246	-7.075	-33.110	1.00	81.29	B C
ATOM	3293	CD	LYS	506	-59.493	-7.584	-34.332	1.00	83.87	B C
ATOM	3294	CE	LYS	506	-58.221	-8.327	-33.940	1.00	85.75	B C
ATOM	3295	NZ	LYS	506	-57.438	-8.754	-35.136	1.00	86.14	B N
ATOM	3296	C	LYS	506	-63.631	-7.314	-32.669	1.00	75.15	B C
ATOM	3297	O	LYS	506	-64.476	-6.442	-32.465	1.00	74.81	B O
ATOM	3298	N	LEU	507	-63.487	-8.371	-31.877	1.00	71.57	B N
ATOM	3299	CA	LEU	507	-64.313	-8.524	-30.684	1.00	68.58	B C
ATOM	3300	CB	LEU	507	-64.419	-10.000	-30.294	1.00	70.01	B C
ATOM	3301	CG	LEU	507	-65.504	-10.799	-31.017	1.00	71.09	B C
ATOM	3302	CD1	LEU	507	-65.466	-12.256	-30.572	1.00	72.08	B C
ATOM	3303	CD2	LEU	507	-66.861	-10.184	-30.716	1.00	70.93	B C
ATOM	3304	C	LEU	507	-63.774	-7.719	-29.505	1.00	65.55	B C
ATOM	3305	O	LEU	507	-62.595	-7.808	-29.162	1.00	65.40	B O
ATOM	3306	N	VAL	508	-64.647	-6.929	-28.891	1.00	62.61	B N
ATOM	3307	CA	VAL	508	-64.284	-6.166	-27.706	1.00	60.31	B C
ATOM	3308	CB	VAL	508	-64.402	-4.648	-27.958	1.00	59.66	B C
ATOM	3309	CG1	VAL	508	-63.534	-4.250	-29.141	1.00	58.42	B C
ATOM	3310	CG2	VAL	508	-65.850	-4.271	-28.199	1.00	58.70	B C
ATOM	3311	C	VAL	508	-65.181	-6.544	-26.534	1.00	59.23	B C
ATOM	3312	O	VAL	508	-66.215	-7.183	-26.713	1.00	58.09	B O
ATOM	3313	N	CYS	509	-64.773	-6.149	-25.335	1.00	59.26	B N
ATOM	3314	CA	CYS	509	-65.501	-6.483	-24.119	1.00	60.27	B C
ATOM	3315	C	CYS	509	-66.128	-5.225	-23.534	1.00	61.97	B C
ATOM	3316	O	CYS	509	-65.421	-4.335	-23.065	1.00	62.31	B O
ATOM	3317	CB	CYS	509	-64.533	-7.109	-23.113	1.00	59.87	B C
ATOM	3318	SG	CYS	509	-65.198	-7.558	-21.476	1.00	60.83	B S
ATOM	3319	N	ARG	510	-67.455	-5.155	-23.565	1.00	63.76	B N
ATOM	3320	CA	ARG	510	-68.173	-3.970	-23.107	1.00	65.70	B C
ATOM	3321	CB	ARG	510	-69.146	-3.499	-24.191	1.00	67.66	B C
ATOM	3322	CG	ARG	510	-69.977	-2.286	-23.801	1.00	71.78	B C
ATOM	3323	CD	ARG	510	-69.457	-1.026	-24.473	1.00	76.06	B C
ATOM	3324	NE	ARG	510	-70.306	-0.602	-25.585	1.00	79.19	B N
ATOM	3325	CZ	ARG	510	-69.871	0.077	-26.643	1.00	81.45	B C
ATOM	3326	NH1	ARG	510	-70.715	0.424	-27.605	1.00	82.26	B N
ATOM	3327	NH2	ARG	510	-68.590	0.407	-26.746	1.00	81.82	B N
ATOM	3328	C	ARG	510	-68.940	-4.240	-21.817	1.00	66.34	B C
ATOM	3329	O	ARG	510	-69.755	-5.157	-21.751	1.00	66.22	B O
ATOM	3330	N	ALA	511	-68.674	-3.435	-20.794	1.00	67.79	B N
ATOM	3331	CA	ALA	511	-69.380	-3.557	-19.525	1.00	69.49	B C
ATOM	3332	CB	ALA	511	-68.390	-3.565	-18.375	1.00	69.05	B C
ATOM	3333	C	ALA	511	-70.366	-2.408	-19.359	1.00	72.13	B C
ATOM	3334	O	ALA	511	-70.060	-1.260	-19.685	1.00	71.53	B O
ATOM	3335	N	HIS	512	-71.552	-2.725	-18.852	1.00	75.10	B N
ATOM	3336	CA	HIS	512	-72.610	-1.734	-18.709	1.00	77.76	B C
ATOM	3337	CB	HIS	512	-73.915	-2.276	-19.305	1.00	79.16	B C
ATOM	3338	CG	HIS	512	-73.832	-2.568	-20.773	1.00	80.99	B C
ATOM	3339	CD2	HIS	512	-73.653	-3.732	-21.442	1.00	81.47	B C
ATOM	3340	ND1	HIS	512	-73.926	-1.584	-21.735	1.00	81.65	B N
ATOM	3341	CE1	HIS	512	-73.807	-2.130	-22.933	1.00	81.85	B C
ATOM	3342	NE2	HIS	512	-73.640	-3.431	-22.783	1.00	81.72	B N

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ATOM   3343  C   HIS   512     -72.814  -1.370 -17.244  1.00 78.56      B    C
ATOM   3344  O   HIS   512     -72.797  -2.238 -16.371  1.00 77.69      B    O
ATOM   3345  N   ASN   513     -73.006  -0.081 -16.981  1.00 80.56      B    N
ATOM   3346  CA  ASN   513     -73.163   0.401 -15.613  1.00 83.31      B    C
ATOM   3347  CB  ASN   513     -72.626   1.831 -15.495  1.00 82.08      B    C
ATOM   3348  CG  ASN   513     -72.534   2.302 -14.054  1.00 81.68      B    C
ATOM   3349  OD1 ASN   513     -72.666   1.510 -13.120  1.00 81.74      B    O
ATOM   3350  ND2 ASN   513     -72.304   3.597 -13.867  1.00 80.52      B    N
ATOM   3351  C   ASN   513     -74.626   0.363 -15.172  1.00 85.41      B    C
ATOM   3352  O   ASN   513     -75.534   0.530 -15.987  1.00 85.81      B    O
ATOM   3353  N   ALA   514     -74.847   0.140 -13.880  1.00 87.99      B    N
ATOM   3354  CA  ALA   514     -76.188   0.207 -13.311  1.00 91.44      B    C
ATOM   3355  CB  ALA   514     -76.222  -0.503 -11.962  1.00 90.86      B    C
ATOM   3356  C   ALA   514     -76.608   1.666 -13.149  1.00 94.10      B    C
ATOM   3357  O   ALA   514     -75.762   2.559 -13.074  1.00 94.52      B    O
ATOM   3358  N   PHE   515     -77.916   1.905 -13.094  1.00 96.76      B    N
ATOM   3359  CA  PHE   515     -78.440   3.265 -13.033  1.00 98.60      B    C
ATOM   3360  CB  PHE   515     -79.951   3.262 -13.296  1.00101.08      B    C
ATOM   3361  CG  PHE   515     -80.327   2.774 -14.671  1.00103.86      B    C
ATOM   3362  CD1 PHE   515     -81.351   1.853 -14.840  1.00104.87      B    C
ATOM   3363  CD2 PHE   515     -79.658   3.237 -15.795  1.00104.65      B    C
ATOM   3364  CE1 PHE   515     -81.701   1.400 -16.103  1.00105.30      B    C
ATOM   3365  CE2 PHE   515     -80.003   2.789 -17.061  1.00105.48      B    C
ATOM   3366  CZ  PHE   515     -81.026   1.869 -17.215  1.00105.53      B    C
ATOM   3367  C   PHE   515     -78.147   3.936 -11.692  1.00 98.36      B    C
ATOM   3368  O   PHE   515     -78.217   5.160 -11.576  1.00 98.94      B    O
ATOM   3369  N   GLY   516     -77.816   3.135 -10.684  1.00 97.52      B    N
ATOM   3370  CA  GLY   516     -77.447   3.691  -9.394  1.00 96.59      B    C
ATOM   3371  C   GLY   516     -75.972   3.529  -9.061  1.00 96.24      B    C
ATOM   3372  O   GLY   516     -75.559   3.767  -7.925  1.00 96.40      B    O
ATOM   3373  N   GLY   517     -75.175   3.130 -10.049  1.00 95.19      B    N
ATOM   3374  CA  GLY   517     -73.790   2.774  -9.787  1.00 92.79      B    C
ATOM   3375  C   GLY   517     -72.776   3.864 -10.079  1.00 91.22      B    C
ATOM   3376  O   GLY   517     -73.064   4.826 -10.794  1.00 91.57      B    O
ATOM   3377  N   GLU   518     -71.577   3.701  -9.525  1.00 89.22      B    N
ATOM   3378  CA  GLU   518     -70.512   4.693  -9.645  1.00 86.83      B    C
ATOM   3379  CB  GLU   518     -69.724   4.773  -8.334  1.00 89.07      B    C
ATOM   3380  CG  GLU   518     -69.853   3.538  -7.443  1.00 92.53      B    C
ATOM   3381  CD  GLU   518     -69.338   2.270  -8.105  1.00 94.95      B    C
ATOM   3382  OE1 GLU   518     -68.149   1.936  -7.912  1.00 96.07      B    O
ATOM   3383  OE2 GLU   518     -70.124   1.603  -8.813  1.00 96.67      B    O
ATOM   3384  C   GLU   518     -69.557   4.398 -10.798  1.00 83.45      B    C
ATOM   3385  O   GLU   518     -68.691   5.213 -11.122  1.00 83.02      B    O
ATOM   3386  N   GLY   519     -69.717   3.231 -11.413  1.00 79.54      B    N
ATOM   3387  CA  GLY   519     -68.872   2.864 -12.536  1.00 73.49      B    C
ATOM   3388  C   GLY   519     -68.469   1.401 -12.522  1.00 69.03      B    C
ATOM   3389  O   GLY   519     -68.432   0.761 -11.468  1.00 68.16      B    O
ATOM   3390  N   VAL   520     -68.164   0.868 -13.701  1.00 64.49      B    N
ATOM   3391  CA  VAL   520     -67.812  -0.540 -13.838  1.00 60.24      B    C
ATOM   3392  CB  VAL   520     -68.924  -1.328 -14.570  1.00 59.70      B    C
ATOM   3393  CG1 VAL   520     -70.216  -1.262 -13.780  1.00 60.31      B    C
ATOM   3394  CG2 VAL   520     -69.128  -0.761 -15.968  1.00 58.45      B    C
ATOM   3395  C   VAL   520     -66.525  -0.699 -14.632  1.00 57.65      B    C
ATOM   3396  O   VAL   520     -66.190   0.146 -15.462  1.00 55.74      B    O
ATOM   3397  N   TYR   521     -65.806  -1.786 -14.376  1.00 55.08      B    N
ATOM   3398  CA  TYR   521     -64.711  -2.180 -15.252  1.00 53.22      B    C
ATOM   3399  CB  TYR   521     -63.522  -2.693 -14.438  1.00 52.94      B    C
ATOM   3400  CG  TYR   521     -62.891  -1.659 -13.537  1.00 53.88      B    C
ATOM   3401  CD1 TYR   521     -63.254  -1.566 -12.200  1.00 54.58      B    C
ATOM   3402  CE1 TYR   521     -62.667  -0.639 -11.362  1.00 55.31      B    C
ATOM   3403  CD2 TYR   521     -61.919  -0.790 -14.016  1.00 53.24      B    C
ATOM   3404  CE2 TYR   521     -61.326   0.140 -13.185  1.00 54.89      B    C
ATOM   3405  CZ  TYR   521     -61.704   0.210 -11.859  1.00 55.53      B    C
ATOM   3406  OH  TYR   521     -61.120   1.134 -11.021  1.00 57.45      B    O
ATOM   3407  C   TYR   521     -65.161  -3.272 -16.215  1.00 51.73      B    C
ATOM   3408  O   TYR   521     -65.941  -4.155 -15.853  1.00 51.92      B    O
ATOM   3409  N   ALA   522     -64.671  -3.201 -17.446  1.00 49.79      B    N
ATOM   3410  CA  ALA   522     -64.726  -4.338 -18.350  1.00 48.15      B    C
ATOM   3411  CB  ALA   522     -64.927  -3.866 -19.782  1.00 47.08      B    C
ATOM   3412  C   ALA   522     -63.401  -5.076 -18.220  1.00 48.00      B    C
ATOM   3413  O   ALA   522     -62.348  -4.451 -18.090  1.00 48.65      B    O
ATOM   3414  N   ILE   523     -63.456  -6.403 -18.240  1.00 45.80      B    N
ATOM   3415  CA  ILE   523     -62.257  -7.208 -18.080  1.00 43.99      B    C
ATOM   3416  CB  ILE   523     -62.240  -7.891 -16.715  1.00 43.33      B    C
ATOM   3417  CG2 ILE   523     -60.937  -8.642 -16.529  1.00 42.17      B    C
ATOM   3418  CG1 ILE   523     -62.406  -6.837 -15.618  1.00 43.33      B    C
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ATOM   3419  CD1 ILE   523     -62.845  -7.401 -14.295  1.00 41.92      B    C
ATOM   3420  C   ILE   523     -62.187  -8.267 -19.165  1.00 44.59      B    C
ATOM   3421  O   ILE   523     -63.005  -9.190 -19.209  1.00 46.06      B    O
ATOM   3422  N   ALA   524     -61.207  -8.127 -20.048  1.00 42.53      B    N
ATOM   3423  CA  ALA   524     -61.078  -9.025 -21.179  1.00 41.89      B    C
ATOM   3424  CB  ALA   524     -60.713  -8.238 -22.431  1.00 41.27      B    C
ATOM   3425  C   ALA   524     -60.024 -10.080 -20.898  1.00 41.28      B    C
ATOM   3426  O   ALA   524     -59.078  -9.851 -20.150  1.00 41.18      B    O
ATOM   3427  N   ARG   525     -60.200 -11.247 -21.497  1.00 41.36      B    N
ATOM   3428  CA  ARG   525     -59.173 -12.270 -21.451  1.00 41.15      B    C
ATOM   3429  CB  ARG   525     -59.781 -13.570 -20.924  1.00 40.18      B    C
ATOM   3430  CG  ARG   525     -58.773 -14.635 -20.566  1.00 39.48      B    C
ATOM   3431  CD  ARG   525     -58.041 -14.338 -19.268  1.00 37.58      B    C
ATOM   3432  NE  ARG   525     -57.191 -15.473 -18.924  1.00 37.63      B    N
ATOM   3433  CZ  ARG   525     -56.069 -15.401 -18.221  1.00 38.00      B    C
ATOM   3434  NH1 ARG   525     -55.376 -16.508 -17.975  1.00 35.82      B    N
ATOM   3435  NH2 ARG   525     -55.643 -14.230 -17.761  1.00 37.77      B    N
ATOM   3436  C   ARG   525     -58.628 -12.439 -22.869  1.00 40.98      B    C
ATOM   3437  O   ARG   525     -59.364 -12.790 -23.787  1.00 41.72      B    O
ATOM   3438  N   CYS   526     -57.342 -12.163 -23.050  1.00 41.85      B    N
ATOM   3439  CA  CYS   526     -56.764 -12.090 -24.387  1.00 42.79      B    C
ATOM   3440  C   CYS   526     -55.720 -13.167 -24.581  1.00 42.97      B    C
ATOM   3441  O   CYS   526     -54.816 -13.322 -23.761  1.00 44.19      B    O
ATOM   3442  CB  CYS   526     -56.121 -10.723 -24.615  1.00 45.96      B    C
ATOM   3443  SG  CYS   526     -57.256  -9.311 -24.431  1.00 49.70      B    S
ATOM   3444  N   CYS   527     -55.837 -13.912 -25.674  1.00 43.71      B    N
ATOM   3445  CA  CYS   527     -55.014 -15.099 -25.851  1.00 43.31      B    C
ATOM   3446  C   CYS   527     -54.445 -15.210 -27.257  1.00 43.72      B    C
ATOM   3447  O   CYS   527     -55.045 -14.739 -28.226  1.00 41.42      B    O
ATOM   3448  CB  CYS   527     -55.830 -16.351 -25.559  1.00 42.73      B    C
ATOM   3449  SG  CYS   527     -56.857 -16.347 -24.051  1.00 43.67      B    S
ATOM   3450  N   LEU   528     -53.287 -15.852 -27.355  1.00 46.11      B    N
ATOM   3451  CA  LEU   528     -52.702 -16.187 -28.644  1.00 49.34      B    C
ATOM   3452  CB  LEU   528     -51.176 -16.118 -28.565  1.00 49.97      B    C
ATOM   3453  CG  LEU   528     -50.603 -14.699 -28.532  1.00 51.33      B    C
ATOM   3454  CD1 LEU   528     -49.103 -14.751 -28.316  1.00 51.51      B    C
ATOM   3455  CD2 LEU   528     -50.934 -13.991 -29.840  1.00 50.91      B    C
ATOM   3456  C   LEU   528     -53.145 -17.583 -29.064  1.00 51.31      B    C
ATOM   3457  O   LEU   528     -52.695 -18.589 -28.511  1.00 51.03      B    O
ATOM   3458  N   LEU   529     -54.042 -17.622 -30.043  1.00 52.37      B    N
ATOM   3459  CA  LEU   529     -54.606 -18.865 -30.548  1.00 53.39      B    C
ATOM   3460  CB  LEU   529     -56.102 -18.934 -30.230  1.00 53.20      B    C
ATOM   3461  CG  LEU   529     -56.625 -20.120 -29.423  1.00 52.77      B    C
ATOM   3462  CD1 LEU   529     -58.118 -20.235 -29.646  1.00 52.41      B    C
ATOM   3463  CD2 LEU   529     -55.929 -21.402 -29.849  1.00 53.54      B    C
ATOM   3464  C   LEU   529     -54.418 -18.876 -32.057  1.00 54.80      B    C
ATOM   3465  O   LEU   529     -55.236 -18.328 -32.799  1.00 53.85      B    O
ATOM   3466  N   PRO   530     -53.332 -19.495 -32.532  1.00 56.84      B    N
ATOM   3467  CD  PRO   530     -52.292 -20.195 -31.760  1.00 57.17      B    C
ATOM   3468  CA  PRO   530     -53.078 -19.522 -33.976  1.00 58.95      B    C
ATOM   3469  CB  PRO   530     -51.726 -20.233 -34.096  1.00 58.96      B    C
ATOM   3470  CG  PRO   530     -51.557 -20.980 -32.809  1.00 58.65      B    C
ATOM   3471  C   PRO   530     -54.184 -20.219 -34.772  1.00 60.63      B    C
ATOM   3472  O   PRO   530     -54.712 -21.253 -34.353  1.00 59.30      B    O
ATOM   3473  N   GLN   531     -54.538 -19.623 -35.909  1.00 62.90      B    N
ATOM   3474  CA  GLN   531     -55.518 -20.193 -36.832  1.00 65.75      B    C
ATOM   3475  CB  GLN   531     -54.988 -21.495 -37.431  1.00 67.57      B    C
ATOM   3476  CG  GLN   531     -53.804 -21.301 -38.356  1.00 72.19      B    C
ATOM   3477  CD  GLN   531     -52.752 -22.372 -38.173  1.00 75.19      B    C
ATOM   3478  OE1 GLN   531     -53.063 -23.506 -37.801  1.00 76.85      B    O
ATOM   3479  NE2 GLN   531     -51.496 -22.019 -38.429  1.00 76.74      B    N
ATOM   3480  C   GLN   531     -56.863 -20.449 -36.175  1.00 65.84      B    C
ATOM   3481  O   GLN   531     -57.506 -21.465 -36.427  1.00 66.30      B    O
ATOM   3482  N   ALA   532     -57.288 -19.521 -35.330  1.00 66.37      B    N
ATOM   3483  CA  ALA   532     -58.562 -19.657 -34.652  1.00 66.44      B    C
ATOM   3484  CB  ALA   532     -58.439 -19.188 -33.211  1.00 66.93      B    C
ATOM   3485  C   ALA   532     -59.632 -18.856 -35.373  1.00 66.62      B    C
ATOM   3486  O   ALA   532     -59.369 -17.771 -35.891  1.00 67.32      B    O
ATOM   3487  N   ALA   533     -60.840 -19.404 -35.407  1.00 66.62      B    N
ATOM   3488  CA  ALA   533     -61.998 -18.674 -35.893  1.00 66.37      B    C
ATOM   3489  CB  ALA   533     -62.599 -19.385 -37.101  1.00 65.70      B    C
ATOM   3490  C   ALA   533     -63.017 -18.593 -34.763  1.00 66.18      B    C
ATOM   3491  O   ALA   533     -63.818 -19.510 -34.563  1.00 66.80      B    O
ATOM   3492  N   CYS   534     -62.975 -17.495 -34.020  1.00 65.19      B    N
ATOM   3493  CA  CYS   534     -63.886 -17.309 -32.903  1.00 64.70      B    C
ATOM   3494  C   CYS   534     -65.066 -16.439 -33.307  1.00 64.08      B    C
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ATOM   3495  O   CYS   534     -65.030 -15.764 -34.333  1.00 63.92      B    O
ATOM   3496  CB  CYS   534     -63.164 -16.654 -31.733  1.00 64.32      B    C
ATOM   3497  SG  CYS   534     -61.664 -17.496 -31.140  1.00 66.91      B    S
ATOM   3498  N   SER   535     -66.107 -16.458 -32.482  1.00 64.06      B    N
ATOM   3499  CA  SER   535     -67.332 -15.722 -32.756  1.00 64.37      B    C
ATOM   3500  CB  SER   535     -68.104 -16.394 -33.894  1.00 65.10      B    C
ATOM   3501  OG  SER   535     -68.461 -17.724 -33.551  1.00 67.16      B    O
ATOM   3502  C   SER   535     -68.192 -15.695 -31.500  1.00 64.34      B    C
ATOM   3503  O   SER   535     -67.978 -16.480 -30.579  1.00 64.67      B    O
ATOM   3504  N   VAL   536     -69.162 -14.789 -31.467  1.00 64.90      B    N
ATOM   3505  CA  VAL   536     -70.068 -14.687 -30.331  1.00 65.96      B    C
ATOM   3506  CB  VAL   536     -70.133 -13.241 -29.791  1.00 65.53      B    C
ATOM   3507  CG1 VAL   536     -71.068 -13.175 -28.594  1.00 64.39      B    C
ATOM   3508  CG2 VAL   536     -68.743 -12.769 -29.405  1.00 65.88      B    C
ATOM   3509  C   VAL   536     -71.471 -15.116 -30.738  1.00 67.18      B    C
ATOM   3510  O   VAL   536     -71.923 -14.820 -31.843  1.00 65.81      B    O
ATOM   3511  N   HIS   537     -72.153 -15.818 -29.840  1.00 69.33      B    N
ATOM   3512  CA  HIS   537     -73.527 -16.238 -30.076  1.00 72.07      B    C
ATOM   3513  CB  HIS   537     -73.586 -17.754 -30.277  1.00 73.31      B    C
ATOM   3514  CG  HIS   537     -72.754 -18.240 -31.423  1.00 75.71      B    C
ATOM   3515  CD2 HIS   537     -71.437 -18.549 -31.502  1.00 76.61      B    C
ATOM   3516  ND1 HIS   537     -73.269 -18.438 -32.686  1.00 76.54      B    N
ATOM   3517  CE1 HIS   537     -72.306 -18.847 -33.494  1.00 77.41      B    C
ATOM   3518  NE2 HIS   537     -71.184 -18.922 -32.800  1.00 78.04      B    N
ATOM   3519  C   HIS   537     -74.395 -15.831 -28.894  1.00 73.31      B    C
ATOM   3520  O   HIS   537     -74.074 -16.135 -27.745  1.00 73.15      B    O
ATOM   3521  N   THR   538     -75.493 -15.140 -29.182  1.00 75.42      B    N
ATOM   3522  CA  THR   538     -76.318 -14.538 -28.141  1.00 77.29      B    C
ATOM   3523  CB  THR   538     -76.397 -13.005 -28.311  1.00 77.11      B    C
ATOM   3524  OG1 THR   538     -75.077 -12.447 -28.274  1.00 76.83      B    O
ATOM   3525  CG2 THR   538     -77.238 -12.389 -27.203  1.00 76.60      B    C
ATOM   3526  C   THR   538     -77.740 -15.086 -28.151  1.00 79.06      B    C
ATOM   3527  O   THR   538     -78.284 -15.411 -29.207  1.00 78.65      B    O
ATOM   3528  N   ALA   539     -78.335 -15.186 -26.968  1.00 80.99      B    N
ATOM   3529  CA  ALA   539     -79.752 -15.495 -26.845  1.00 83.77      B    C
ATOM   3530  CB  ALA   539     -79.933 -16.889 -26.265  1.00 83.09      B    C
ATOM   3531  C   ALA   539     -80.416 -14.461 -25.944  1.00 86.16      B    C
ATOM   3532  O   ALA   539     -79.932 -14.177 -24.850  1.00 86.73      B    O
ATOM   3533  N   PRO   540     -81.537 -13.880 -26.396  1.00 88.31      B    N
ATOM   3534  CD  PRO   540     -82.164 -14.122 -27.707  1.00 88.43      B    C
ATOM   3535  CA  PRO   540     -82.293 -12.912 -25.592  1.00 90.15      B    C
ATOM   3536  CB  PRO   540     -83.360 -12.399 -26.556  1.00 89.67      B    C
ATOM   3537  CG  PRO   540     -83.517 -13.493 -27.559  1.00 89.11      B    C
ATOM   3538  C   PRO   540     -82.898 -13.553 -24.344  1.00 92.40      B    C
ATOM   3539  O   PRO   540     -82.934 -14.776 -24.219  1.00 91.96      B    O
ATOM   3540  N   PRO   541     -83.380 -12.729 -23.400  1.00 94.77      B    N
ATOM   3541  CD  PRO   541     -83.368 -11.257 -23.433  1.00 94.86      B    C
ATOM   3542  CA  PRO   541     -83.932 -13.243 -22.141  1.00 97.44      B    C
ATOM   3543  CB  PRO   541     -84.361 -11.980 -21.393  1.00 96.46      B    C
ATOM   3544  CG  PRO   541     -83.528 -10.892 -21.986  1.00 95.58      B    C
ATOM   3545  C   PRO   541     -85.103 -14.200 -22.362  1.00100.50      B    C
ATOM   3546  O   PRO   541     -85.968 -13.950 -23.201  1.00100.57      B    O
ATOM   3547  N   ALA   542     -85.120 -15.293 -21.604  1.00104.26      B    N
ATOM   3548  CA  ALA   542     -86.190 -16.280 -21.700  1.00107.63      B    C
ATOM   3549  CB  ALA   542     -85.685 -17.646 -21.250  1.00107.30      B    C
ATOM   3550  C   ALA   542     -87.377 -15.855 -20.843  1.00110.11      B    C
ATOM   3551  O   ALA   542     -88.513 -15.808 -21.316  1.00110.36      B    O
ATOM   3552  N   GLU   543     -87.100 -15.550 -19.578  1.00113.18      B    N
ATOM   3553  CA  GLU   543     -88.102 -15.008 -18.666  1.00116.02      B    C
ATOM   3554  CB  GLU   543     -88.709 -13.731 -19.254  1.00116.65      B    C
ATOM   3555  CG  GLU   543     -89.515 -12.911 -18.263  1.00118.13      B    C
ATOM   3556  CD  GLU   543     -89.438 -11.425 -18.547  1.00118.89      B    C
ATOM   3557  OE1 GLU   543     -90.443 -10.716 -18.316  1.00119.23      B    O
ATOM   3558  OE2 GLU   543     -88.369 -10.961 -19.002  1.00118.98      B    O
ATOM   3559  C   GLU   543     -89.202 -16.025 -18.376  1.00117.48      B    C
ATOM   3560  O   GLU   543     -89.605 -16.788 -19.258  1.00117.72      B    O
ATOM   3561  N   ALA   544     -89.682 -16.033 -17.134  1.00118.85      B    N
ATOM   3562  CA  ALA   544     -90.647 -17.034 -16.688  1.00120.01      B    C
ATOM   3563  CB  ALA   544     -91.991 -16.816 -17.382  1.00119.80      B    C
ATOM   3564  C   ALA   544     -90.118 -18.437 -16.987  1.00120.71      B    C
ATOM   3565  O   ALA   544     -90.876 -19.410 -17.014  1.00121.08      B    O
ATOM   3566  N   SER   545     -88.810 -18.526 -17.212  1.00120.84      B    N
ATOM   3567  CA  SER   545     -88.169 -19.770 -17.617  1.00120.61      B    C
ATOM   3568  CB  SER   545     -87.431 -19.561 -18.942  1.00120.62      B    C
ATOM   3569  OG  SER   545     -86.831 -20.762 -19.392  1.00121.09      B    O
ATOM   3570  C   SER   545     -87.189 -20.247 -16.548  1.00120.42      B    C
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ATOM   3571  O   SER   545     -87.011 -19.591 -15.522  1.00120.46      B     O
ATOM   3572  N   MET   546     -86.559 -21.394 -16.789  1.00120.05      B     N
ATOM   3573  CA  MET   546     -85.549 -21.918 -15.875  1.00119.53      B     C
ATOM   3574  CB  MET   546     -85.530 -23.451 -15.921  1.00120.98      B     C
ATOM   3575  CG  MET   546     -84.814 -24.046 -17.132  1.00122.54      B     C
ATOM   3576  SD  MET   546     -85.634 -23.714 -18.707  1.00124.63      B     S
ATOM   3577  CE  MET   546     -84.540 -24.571 -19.851  1.00124.00      B     C
ATOM   3578  C   MET   546     -84.161 -21.374 -16.225  1.00118.19      B     C
ATOM   3579  O   MET   546     -83.141 -21.923 -15.803  1.00118.60      B     O
ATOM   3580  N   GLY   547     -84.131 -20.294 -17.001  1.00115.77      B     N
ATOM   3581  CA  GLY   547     -82.868 -19.668 -17.351  1.00112.18      B     C
ATOM   3582  C   GLY   547     -82.687 -19.464 -18.842  1.00109.64      B     C
ATOM   3583  O   GLY   547     -83.268 -20.186 -19.654  1.00109.67      B     O
ATOM   3584  N   THR   548     -81.879 -18.473 -19.205  1.00106.80      B     N
ATOM   3585  CA  THR   548     -81.561 -18.211 -20.604  1.00103.45      B     C
ATOM   3586  CB  THR   548     -81.332 -16.709 -20.848  1.00102.95      B     C
ATOM   3587  OG1 THR   548     -82.486 -15.973 -20.422  1.00102.38      B     O
ATOM   3588  CG2 THR   548     -81.089 -16.441 -22.322  1.00102.67      B     C
ATOM   3589  C   THR   548     -80.296 -18.974 -20.989  1.00101.35      B     C
ATOM   3590  O   THR   548     -79.298 -18.932 -20.270  1.00101.40      B     O
ATOM   3591  N   ARG   549     -80.342 -19.671 -22.120  1.00 98.30      B     N
ATOM   3592  CA  ARG   549     -79.255 -20.565 -22.504  1.00 95.33      B     C
ATOM   3593  CB  ARG   549     -79.680 -22.022 -22.300  1.00 97.27      B     C
ATOM   3594  CG  ARG   549     -80.069 -22.352 -20.871  1.00 99.72      B     C
ATOM   3595  CD  ARG   549     -80.915 -23.613 -20.793  1.00101.92      B     C
ATOM   3596  NE  ARG   549     -81.482 -23.787 -19.459  1.00104.40      B     N
ATOM   3597  CZ  ARG   549     -80.888 -24.457 -18.475  1.00105.40      B     C
ATOM   3598  NH1 ARG   549     -81.478 -24.562 -17.289  1.00105.68      B     N
ATOM   3599  NH2 ARG   549     -79.707 -25.026 -18.677  1.00105.41      B     N
ATOM   3600  C   ARG   549     -78.808 -20.370 -23.946  1.00 92.19      B     C
ATOM   3601  O   ARG   549     -79.616 -20.090 -24.828  1.00 91.75      B     O
ATOM   3602  N   VAL   550     -77.509 -20.522 -24.174  1.00 88.72      B     N
ATOM   3603  CA  VAL   550     -76.959 -20.534 -25.520  1.00 85.99      B     C
ATOM   3604  CB  VAL   550     -76.520 -19.120 -25.963  1.00 85.94      B     C
ATOM   3605  CG1 VAL   550     -75.387 -18.624 -25.080  1.00 85.93      B     C
ATOM   3606  CG2 VAL   550     -76.093 -19.138 -27.420  1.00 85.60      B     C
ATOM   3607  C   VAL   550     -75.751 -21.461 -25.532  1.00 84.16      B     C
ATOM   3608  O   VAL   550     -75.046 -21.584 -24.532  1.00 83.76      B     O
ATOM   3609  N   HIS   551     -75.522 -22.127 -26.657  1.00 82.33      B     N
ATOM   3610  CA  HIS   551     -74.368 -23.001 -26.788  1.00 81.02      B     C
ATOM   3611  CB  HIS   551     -74.732 -24.438 -26.390  1.00 83.78      B     C
ATOM   3612  CG  HIS   551     -75.882 -25.015 -27.158  1.00 86.60      B     C
ATOM   3613  CD2 HIS   551     -75.968 -25.468 -28.431  1.00 87.05      B     C
ATOM   3614  ND1 HIS   551     -77.129 -25.203 -26.600  1.00 87.85      B     N
ATOM   3615  CE1 HIS   551     -77.933 -25.747 -27.496  1.00 88.26      B     C
ATOM   3616  NE2 HIS   551     -77.253 -25.919 -28.615  1.00 88.32      B     N
ATOM   3617  C   HIS   551     -73.800 -22.976 -28.198  1.00 78.87      B     C
ATOM   3618  O   HIS   551     -74.450 -22.512 -29.135  1.00 78.95      B     O
ATOM   3619  N   CYS   552     -72.575 -23.469 -28.334  1.00 76.26      B     N
ATOM   3620  CA  CYS   552     -71.885 -23.474 -29.612  1.00 74.68      B     C
ATOM   3621  C   CYS   552     -72.362 -24.649 -30.456  1.00 76.00      B     C
ATOM   3622  O   CYS   552     -71.934 -25.786 -30.253  1.00 75.80      B     O
ATOM   3623  CB  CYS   552     -70.375 -23.572 -29.387  1.00 71.57      B     C
ATOM   3624  SG  CYS   552     -69.677 -22.320 -28.252  1.00 67.04      B     S
ATOM   3625  N   HIS   553     -73.247 -24.363 -31.406  1.00 77.76      B     N
ATOM   3626  CA  HIS   553     -73.889 -25.399 -32.212  1.00 79.66      B     C
ATOM   3627  CB  HIS   553     -75.055 -24.803 -33.013  1.00 84.15      B     C
ATOM   3628  CG  HIS   553     -76.263 -24.472 -32.188  1.00 88.61      B     C
ATOM   3629  CD2 HIS   553     -76.626 -23.332 -31.553  1.00 90.41      B     C
ATOM   3630  ND1 HIS   553     -77.290 -25.369 -31.981  1.00 90.68      B     N
ATOM   3631  CE1 HIS   553     -78.234 -24.795 -31.256  1.00 91.59      B     C
ATOM   3632  NE2 HIS   553     -77.857 -23.558 -30.983  1.00 91.84      B     N
ATOM   3633  C   HIS   553     -72.910 -26.066 -33.177  1.00 78.29      B     C
ATOM   3634  O   HIS   553     -72.901 -27.289 -33.317  1.00 78.79      B     O
ATOM   3635  N   GLN   554     -72.091 -25.257 -33.841  1.00 75.87      B     N
ATOM   3636  CA  GLN   554     -71.197 -25.755 -34.878  1.00 73.89      B     C
ATOM   3637  CB  GLN   554     -70.435 -24.592 -35.516  1.00 76.44      B     C
ATOM   3638  CG  GLN   554     -71.323 -23.528 -36.140  1.00 80.50      B     C
ATOM   3639  CD  GLN   554     -70.578 -22.230 -36.395  1.00 83.46      B     C
ATOM   3640  OE1 GLN   554     -70.183 -21.937 -37.527  1.00 83.91      B     O
ATOM   3641  NE2 GLN   554     -70.376 -21.445 -35.337  1.00 84.48      B     N
ATOM   3642  C   GLN   554     -70.204 -26.781 -34.340  1.00 71.10      B     C
ATOM   3643  O   GLN   554     -69.724 -26.668 -33.212  1.00 70.66      B     O
ATOM   3644  N   GLN   555     -69.900 -27.781 -35.161  1.00 67.93      B     N
ATOM   3645  CA  GLN   555     -68.960 -28.829 -34.792  1.00 65.48      B     C
ATOM   3646  CB  GLN   555     -69.080 -30.006 -35.762  1.00 66.77      B     C
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ATOM   3647  CG  GLN   555     -70.449 -30.677 -35.766  1.00 68.17      B    C
ATOM   3648  CD  GLN   555     -70.635 -31.626 -34.596  1.00 69.56      B    C
ATOM   3649  OE1 GLN   555     -70.909 -31.202 -33.472  1.00 68.69      B    O
ATOM   3650  NE2 GLN   555     -70.483 -32.919 -34.857  1.00 70.20      B    N
ATOM   3651  C   GLN   555     -67.535 -28.290 -34.818  1.00 63.77      B    C
ATOM   3652  O   GLN   555     -67.166 -27.537 -35.718  1.00 63.08      B    O
ATOM   3653  N   GLY   556     -66.737 -28.678 -33.828  1.00 61.44      B    N
ATOM   3654  CA  GLY   556     -65.364 -28.211 -33.765  1.00 58.50      B    C
ATOM   3655  C   GLY   556     -65.237 -26.820 -33.169  1.00 56.70      B    C
ATOM   3656  O   GLY   556     -64.138 -26.277 -33.072  1.00 56.59      B    O
ATOM   3657  N   HIS   557     -66.363 -26.238 -32.775  1.00 54.74      B    N
ATOM   3658  CA  HIS   557     -66.359 -24.945 -32.104  1.00 54.39      B    C
ATOM   3659  CB  HIS   557     -67.488 -24.066 -32.654  1.00 56.06      B    C
ATOM   3660  CG  HIS   557     -67.257 -23.603 -34.059  1.00 59.40      B    C
ATOM   3661  CD2 HIS   557     -67.399 -22.383 -34.631  1.00 60.00      B    C
ATOM   3662  ND1 HIS   557     -66.794 -24.440 -35.052  1.00 60.87      B    N
ATOM   3663  CE1 HIS   557     -66.658 -23.756 -36.175  1.00 60.41      B    C
ATOM   3664  NE2 HIS   557     -67.018 -22.506 -35.946  1.00 60.64      B    N
ATOM   3665  C   HIS   557     -66.517 -25.126 -30.597  1.00 52.41      B    C
ATOM   3666  O   HIS   557     -67.488 -25.718 -30.129  1.00 52.00      B    O
ATOM   3667  N   VAL   558     -65.551 -24.610 -29.843  1.00 50.33      B    N
ATOM   3668  CA  VAL   558     -65.494 -24.841 -28.408  1.00 48.46      B    C
ATOM   3669  CB  VAL   558     -64.078 -25.271 -27.989  1.00 48.10      B    C
ATOM   3670  CG1 VAL   558     -64.079 -25.737 -26.546  1.00 48.76      B    C
ATOM   3671  CG2 VAL   558     -63.581 -26.372 -28.913  1.00 47.38      B    C
ATOM   3672  C   VAL   558     -65.884 -23.582 -27.642  1.00 47.50      B    C
ATOM   3673  O   VAL   558     -65.607 -22.468 -28.080  1.00 48.38      B    O
ATOM   3674  N   LEU   559     -66.536 -23.761 -26.499  1.00 46.06      B    N
ATOM   3675  CA  LEU   559     -66.947 -22.632 -25.674  1.00 45.93      B    C
ATOM   3676  CB  LEU   559     -68.119 -23.041 -24.782  1.00 45.03      B    C
ATOM   3677  CG  LEU   559     -68.664 -21.975 -23.833  1.00 47.58      B    C
ATOM   3678  CD1 LEU   559     -69.353 -20.881 -24.635  1.00 45.95      B    C
ATOM   3679  CD2 LEU   559     -69.639 -22.617 -22.847  1.00 46.31      B    C
ATOM   3680  C   LEU   559     -65.777 -22.167 -24.810  1.00 45.32      B    C
ATOM   3681  O   LEU   559     -65.271 -22.923 -23.988  1.00 46.56      B    O
ATOM   3682  N   THR   560     -65.342 -20.926 -24.999  1.00 44.72      B    N
ATOM   3683  CA  THR   560     -64.186 -20.413 -24.267  1.00 43.84      B    C
ATOM   3684  CB  THR   560     -63.195 -19.709 -25.204  1.00 42.82      B    C
ATOM   3685  OG1 THR   560     -63.826 -18.565 -25.792  1.00 42.05      B    O
ATOM   3686  CG2 THR   560     -62.746 -20.654 -26.302  1.00 42.95      B    C
ATOM   3687  C   THR   560     -64.574 -19.433 -23.167  1.00 43.79      B    C
ATOM   3688  O   THR   560     -63.807 -19.219 -22.231  1.00 44.66      B    O
ATOM   3689  N   GLY   561     -65.757 -18.834 -23.281  1.00 44.15      B    N
ATOM   3690  CA  GLY   561     -66.217 -17.908 -22.260  1.00 45.65      B    C
ATOM   3691  C   GLY   561     -67.688 -17.546 -22.363  1.00 48.91      B    C
ATOM   3692  O   GLY   561     -68.264 -17.559 -23.452  1.00 48.94      B    O
ATOM   3693  N   CYS   562     -68.296 -17.223 -21.223  1.00 52.04      B    N
ATOM   3694  CA  CYS   562     -69.694 -16.790 -21.173  1.00 56.94      B    C
ATOM   3695  C   CYS   562     -69.821 -15.372 -20.622  1.00 57.36      B    C
ATOM   3696  O   CYS   562     -69.176 -15.019 -19.638  1.00 57.14      B    O
ATOM   3697  CB  CYS   562     -70.525 -17.726 -20.286  1.00 60.25      B    C
ATOM   3698  SG  CYS   562     -70.658 -19.459 -20.842  1.00 68.25      B    S
ATOM   3699  N   SER   563     -70.665 -14.566 -21.254  1.00 59.16      B    N
ATOM   3700  CA  SER   563     -70.980 -13.238 -20.743  1.00 60.40      B    C
ATOM   3701  CB  SER   563     -70.453 -12.167 -21.697  1.00 59.60      B    C
ATOM   3702  OG  SER   563     -69.037 -12.162 -21.714  1.00 60.14      B    O
ATOM   3703  C   SER   563     -72.484 -13.073 -20.570  1.00 61.68      B    C
ATOM   3704  O   SER   563     -73.270 -13.626 -21.140  1.00 61.85      B    O
ATOM   3705  N   SER   564     -72.883 -12.085 -19.780  1.00 63.51      B    N
ATOM   3706  CA  SER   564     -74.296 -11.841 -19.540  1.00 65.75      B    C
ATOM   3707  CB  SER   564     -74.798 -12.760 -18.424  1.00 65.21      B    C
ATOM   3708  OG  SER   564     -76.187 -12.593 -18.216  1.00 64.95      B    O
ATOM   3709  C   SER   564     -74.542 -10.389 -19.156  1.00 67.58      B    C
ATOM   3710  O   SER   564     -73.912  -9.870 -18.236  1.00 67.06      B    O
ATOM   3711  N   HIS   565     -75.458  -9.735 -19.862  1.00 70.45      B    N
ATOM   3712  CA  HIS   565     -75.896  -8.401 -19.475  1.00 74.26      B    C
ATOM   3713  CB  HIS   565     -75.342  -7.352 -20.444  1.00 74.10      B    C
ATOM   3714  CG  HIS   565     -76.129  -7.216 -21.710  1.00 75.03      B    C
ATOM   3715  CD2 HIS   565     -77.193  -6.437 -22.019  1.00 75.03      B    C
ATOM   3716  ND1 HIS   565     -75.824  -7.918 -22.857  1.00 75.79      B    N
ATOM   3717  CE1 HIS   565     -76.664  -7.576 -23.817  1.00 75.02      B    C
ATOM   3718  NE2 HIS   565     -77.505  -6.679 -23.335  1.00 74.96      B    N
ATOM   3719  C   HIS   565     -77.417  -8.327 -19.447  1.00 76.76      B    C
ATOM   3720  O   HIS   565     -78.093  -9.033 -20.191  1.00 76.35      B    O
ATOM   3721  N   TRP   566     -77.952  -7.471 -18.582  1.00 80.24      B    N
ATOM   3722  CA  TRP   566     -79.395  -7.319 -18.457  1.00 83.92      B    C
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ATOM   3723  CB  TRP   566     -79.885  -7.999 -17.181  1.00 82.87      B    C
ATOM   3724  CG  TRP   566     -79.070  -7.676 -15.975  1.00 81.41      B    C
ATOM   3725  CD2 TRP   566     -77.859  -8.319 -15.564  1.00 80.85      B    C
ATOM   3726  CE2 TRP   566     -77.461  -7.724 -14.351  1.00 80.81      B    C
ATOM   3727  CE3 TRP   566     -77.073  -9.342 -16.103  1.00 80.36      B    C
ATOM   3728  CD1 TRP   566     -79.348  -6.740 -15.025  1.00 81.33      B    C
ATOM   3729  NE1 TRP   566     -78.387  -6.762 -14.043  1.00 81.30      B    N
ATOM   3730  CZ2 TRP   566     -76.314  -8.118 -13.667  1.00 80.57      B    C
ATOM   3731  CZ3 TRP   566     -75.934  -9.732 -15.423  1.00 80.05      B    C
ATOM   3732  CH2 TRP   566     -75.565  -9.122 -14.218  1.00 80.06      B    C
ATOM   3733  C   TRP   566     -79.835  -5.861 -18.459  1.00 87.50      B    C
ATOM   3734  O   TRP   566     -79.009  -4.951 -18.373  1.00 87.72      B    O
ATOM   3735  N   GLU   567     -81.146  -5.651 -18.555  1.00 91.79      B    N
ATOM   3736  CA  GLU   567     -81.711  -4.311 -18.649  1.00 95.79      B    C
ATOM   3737  CB  GLU   567     -82.770  -4.270 -19.749  1.00 96.18      B    C
ATOM   3738  CG  GLU   567     -82.799  -2.967 -20.519  1.00 98.36      B    C
ATOM   3739  CD  GLU   567     -81.568  -2.786 -21.384  1.00 99.33      B    C
ATOM   3740  OE1 GLU   567     -80.732  -3.715 -21.431  1.00 98.92      B    O
ATOM   3741  OE2 GLU   567     -81.436  -1.717 -22.019  1.00100.24      B    O
ATOM   3742  C   GLU   567     -82.333  -3.866 -17.328  1.00 98.41      B    C
ATOM   3743  O   GLU   567     -82.113  -2.742 -16.876  1.00 98.57      B    O
ATOM   3744  N   VAL   568     -83.111  -4.751 -16.714  1.00101.90      B    N
ATOM   3745  CA  VAL   568     -83.733  -4.454 -15.428  1.00105.56      B    C
ATOM   3746  CB  VAL   568     -84.660  -5.600 -14.971  1.00105.81      B    C
ATOM   3747  CG1 VAL   568     -85.738  -5.844 -16.016  1.00105.97      B    C
ATOM   3748  CG2 VAL   568     -83.847  -6.864 -14.728  1.00105.81      B    C
ATOM   3749  C   VAL   568     -82.657  -4.251 -14.371  1.00107.91      B    C
ATOM   3750  O   VAL   568     -81.478  -4.485 -14.625  1.00108.57      B    O
ATOM   3751  N   GLU   569     -83.063  -3.812 -13.187  1.00110.90      B    N
ATOM   3752  CA  GLU   569     -82.113  -3.566 -12.111  1.00113.78      B    C
ATOM   3753  CB  GLU   569     -82.451  -2.250 -11.407  1.00114.54      B    C
ATOM   3754  CG  GLU   569     -82.319  -1.029 -12.303  1.00116.01      B    C
ATOM   3755  CD  GLU   569     -80.930  -0.895 -12.904  1.00116.56      B    C
ATOM   3756  OE1 GLU   569     -79.972  -0.649 -12.140  1.00116.54      B    O
ATOM   3757  OE2 GLU   569     -80.796  -1.036 -14.139  1.00116.94      B    O
ATOM   3758  C   GLU   569     -82.088  -4.706 -11.099  1.00115.27      B    C
ATOM   3759  O   GLU   569     -81.021  -5.120 -10.646  1.00115.42      B    O
ATOM   3760  N   ASP   570     -83.267  -5.215 -10.753  1.00116.99      B    N
ATOM   3761  CA  ASP   570     -83.379  -6.268  -9.751  1.00118.36      B    C
ATOM   3762  CB  ASP   570     -84.494  -5.930  -8.758  1.00119.50      B    C
ATOM   3763  CG  ASP   570     -84.260  -4.610  -8.044  1.00120.50      B    C
ATOM   3764  OD1 ASP   570     -85.090  -4.241  -7.185  1.00120.98      B    O
ATOM   3765  OD2 ASP   570     -83.247  -3.941  -8.343  1.00120.72      B    O
ATOM   3766  C   ASP   570     -83.652  -7.627 -10.388  1.00118.64      B    C
ATOM   3767  O   ASP   570     -84.496  -7.752 -11.277  1.00118.66      B    O
ATOM   3768  N   LEU   571     -82.927  -8.641  -9.928  1.00118.81      B    N
ATOM   3769  CA  LEU   571     -83.060  -9.990 -10.463  1.00118.86      B    C
ATOM   3770  CB  LEU   571     -81.836 -10.344 -11.317  1.00118.99      B    C
ATOM   3771  CG  LEU   571     -81.609  -9.554 -12.613  1.00118.90      B    C
ATOM   3772  CD1 LEU   571     -81.205  -8.124 -12.294  1.00119.28      B    C
ATOM   3773  CD2 LEU   571     -80.523 -10.230 -13.434  1.00118.88      B    C
ATOM   3774  C   LEU   571     -83.215 -11.009  -9.336  1.00118.75      B    C
ATOM   3775  O   LEU   571     -84.241 -11.682  -9.227  1.00118.42      B    O
ATOM   3776  N   PRO   585     -72.292 -29.792 -21.993  1.00 83.79      B    N
ATOM   3777  CD  PRO   585     -72.479 -31.194 -22.411  1.00 84.05      B    C
ATOM   3778  CA  PRO   585     -70.992 -29.260 -22.422  1.00 83.07      B    C
ATOM   3779  CB  PRO   585     -70.252 -30.493 -22.939  1.00 83.40      B    C
ATOM   3780  CG  PRO   585     -71.337 -31.427 -23.365  1.00 83.43      B    C
ATOM   3781  C   PRO   585     -71.129 -28.186 -23.496  1.00 81.48      B    C
ATOM   3782  O   PRO   585     -71.931 -28.321 -24.418  1.00 80.88      B    O
ATOM   3783  N   ASN   586     -70.340 -27.124 -23.365  1.00 80.01      B    N
ATOM   3784  CA  ASN   586     -70.363 -26.010 -24.311  1.00 78.83      B    C
ATOM   3785  CB  ASN   586     -70.162 -26.507 -25.749  1.00 79.79      B    C
ATOM   3786  CG  ASN   586     -68.793 -27.124 -25.974  1.00 80.52      B    C
ATOM   3787  OD1 ASN   586     -67.783 -26.421 -26.031  1.00 79.27      B    O
ATOM   3788  ND2 ASN   586     -68.754 -28.447 -26.108  1.00 81.46      B    N
ATOM   3789  C   ASN   586     -71.659 -25.214 -24.243  1.00 77.08      B    C
ATOM   3790  O   ASN   586     -72.162 -24.768 -25.271  1.00 76.87      B    O
ATOM   3791  N   GLN   587     -72.197 -25.034 -23.040  1.00 75.05      B    N
ATOM   3792  CA  GLN   587     -73.408 -24.238 -22.875  1.00 74.57      B    C
ATOM   3793  CB  GLN   587     -74.588 -25.142 -22.512  1.00 74.98      B    C
ATOM   3794  CG  GLN   587     -75.876 -24.385 -22.224  1.00 75.59      B    C
ATOM   3795  CD  GLN   587     -77.085 -25.298 -22.114  1.00 76.09      B    C
ATOM   3796  OE1 GLN   587     -77.886 -25.396 -23.044  1.00 76.13      B    O
ATOM   3797  NE2 GLN   587     -77.223 -25.970 -20.973  1.00 75.12      B    N
ATOM   3798  C   GLN   587     -73.263 -23.146 -21.819  1.00 73.56      B    C
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ATOM   3799  O   GLN   587     -72.681 -23.372 -20.759  1.00 73.19      B    O
ATOM   3800  N   CYS   588     -73.796 -21.964 -22.118  1.00 72.91      B    N
ATOM   3801  CA  CYS   588     -73.872 -20.875 -21.144  1.00 72.44      B    C
ATOM   3802  C   CYS   588     -75.271 -20.786 -20.551  1.00 73.33      B    C
ATOM   3803  O   CYS   588     -76.266 -20.833 -21.276  1.00 73.45      B    O
ATOM   3804  CB  CYS   588     -73.541 -19.530 -21.796  1.00 70.56      B    C
ATOM   3805  SG  CYS   588     -71.864 -19.371 -22.479  1.00 68.54      B    S
ATOM   3806  N   VAL   589     -75.348 -20.645 -19.234  1.00 74.61      B    N
ATOM   3807  CA  VAL   589     -76.629 -20.427 -18.579  1.00 76.86      B    C
ATOM   3808  CB  VAL   589     -76.974 -21.577 -17.618  1.00 76.35      B    C
ATOM   3809  CG1 VAL   589     -78.348 -21.346 -17.016  1.00 76.62      B    C
ATOM   3810  CG2 VAL   589     -76.926 -22.905 -18.355  1.00 76.25      B    C
ATOM   3811  C   VAL   589     -76.612 -19.128 -17.790  1.00 78.39      B    C
ATOM   3812  O   VAL   589     -75.810 -18.956 -16.874  1.00 78.41      B    O
ATOM   3813  N   GLY   590     -77.503 -18.214 -18.152  1.00 80.86      B    N
ATOM   3814  CA  GLY   590     -77.608 -16.965 -17.427  1.00 83.83      B    C
ATOM   3815  C   GLY   590     -78.956 -16.817 -16.753  1.00 85.87      B    C
ATOM   3816  O   GLY   590     -79.812 -17.695 -16.854  1.00 86.28      B    O
ATOM   3817  N   HIS   591     -79.143 -15.700 -16.061  1.00 87.95      B    N
ATOM   3818  CA  HIS   591     -80.426 -15.381 -15.452  1.00 89.48      B    C
ATOM   3819  CB  HIS   591     -80.329 -14.043 -14.716  1.00 90.53      B    C
ATOM   3820  CG  HIS   591     -81.482 -13.768 -13.803  1.00 91.97      B    C
ATOM   3821  CD2 HIS   591     -81.585 -13.837 -12.454  1.00 92.54      B    C
ATOM   3822  ND1 HIS   591     -82.714 -13.355 -14.262  1.00 92.28      B    N
ATOM   3823  CE1 HIS   591     -83.527 -13.181 -13.235  1.00 92.99      B    C
ATOM   3824  NE2 HIS   591     -82.867 -13.466 -12.126  1.00 92.92      B    N
ATOM   3825  C   HIS   591     -81.503 -15.308 -16.534  1.00 89.95      B    C
ATOM   3826  O   HIS   591     -81.210 -15.038 -17.700  1.00 89.23      B    O
ATOM   3827  N   ARG   592     -82.749 -15.554 -16.143  1.00 90.86      B    N
ATOM   3828  CA  ARG   592     -83.861 -15.556 -17.088  1.00 91.76      B    C
ATOM   3829  CB  ARG   592     -85.141 -16.024 -16.390  1.00 94.45      B    C
ATOM   3830  CG  ARG   592     -85.501 -15.201 -15.163  1.00 98.10      B    C
ATOM   3831  CD  ARG   592     -86.947 -15.407 -14.740  1.00101.13      B    C
ATOM   3832  NE  ARG   592     -87.177 -16.735 -14.177  1.00103.71      B    N
ATOM   3833  CZ  ARG   592     -88.312 -17.112 -13.596  1.00105.10      B    C
ATOM   3834  NH1 ARG   592     -88.435 -18.342 -13.112  1.00105.47      B    N
ATOM   3835  NH2 ARG   592     -89.325 -16.259 -13.497  1.00105.79      B    N
ATOM   3836  C   ARG   592     -84.085 -14.169 -17.684  1.00 90.52      B    C
ATOM   3837  O   ARG   592     -84.473 -14.038 -18.845  1.00 89.74      B    O
ATOM   3838  N   GLU   593     -83.834 -13.136 -16.886  1.00 89.53      B    N
ATOM   3839  CA  GLU   593     -84.083 -11.762 -17.310  1.00 88.86      B    C
ATOM   3840  CB  GLU   593     -84.361 -10.877 -16.092  1.00 90.90      B    C
ATOM   3841  CG  GLU   593     -85.593 -11.278 -15.292  1.00 93.06      B    C
ATOM   3842  CD  GLU   593     -85.834 -10.371 -14.095  1.00 94.88      B    C
ATOM   3843  OE1 GLU   593     -86.475  -9.311 -14.270  1.00 95.59      B    O
ATOM   3844  OE2 GLU   593     -85.383 -10.717 -12.980  1.00 95.43      B    O
ATOM   3845  C   GLU   593     -82.917 -11.181 -18.102  1.00 86.91      B    C
ATOM   3846  O   GLU   593     -83.001 -10.063 -18.611  1.00 86.77      B    O
ATOM   3847  N   ALA   594     -81.831 -11.941 -18.206  1.00 84.52      B    N
ATOM   3848  CA  ALA   594     -80.616 -11.452 -18.852  1.00 81.65      B    C
ATOM   3849  CB  ALA   594     -79.410 -11.736 -17.961  1.00 81.53      B    C
ATOM   3850  C   ALA   594     -80.394 -12.061 -20.234  1.00 79.44      B    C
ATOM   3851  O   ALA   594     -80.829 -13.179 -20.508  1.00 79.69      B    O
ATOM   3852  N   SER   595     -79.718 -11.316 -21.105  1.00 76.43      B    N
ATOM   3853  CA  SER   595     -79.199 -11.875 -22.345  1.00 73.80      B    C
ATOM   3854  CB  SER   595     -78.882 -10.764 -23.343  1.00 73.82      B    C
ATOM   3855  OG  SER   595     -80.030  -9.986 -23.621  1.00 74.77      B    O
ATOM   3856  C   SER   595     -77.925 -12.644 -22.024  1.00 72.54      B    C
ATOM   3857  O   SER   595     -77.189 -12.282 -21.108  1.00 72.16      B    O
ATOM   3858  N   ILE   596     -77.669 -13.708 -22.776  1.00 71.00      B    N
ATOM   3859  CA  ILE   596     -76.505 -14.546 -22.527  1.00 68.81      B    C
ATOM   3860  CB  ILE   596     -76.932 -15.985 -22.162  1.00 69.08      B    C
ATOM   3861  CG2 ILE   596     -77.525 -16.682 -23.376  1.00 69.30      B    C
ATOM   3862  CG1 ILE   596     -75.731 -16.768 -21.634  1.00 68.96      B    C
ATOM   3863  CD1 ILE   596     -75.266 -16.303 -20.276  1.00 68.21      B    C
ATOM   3864  C   ILE   596     -75.613 -14.572 -23.765  1.00 67.78      B    C
ATOM   3865  O   ILE   596     -76.101 -14.666 -24.892  1.00 67.52      B    O
ATOM   3866  N   HIS   597     -74.304 -14.478 -23.551  1.00 66.74      B    N
ATOM   3867  CA  HIS   597     -73.350 -14.407 -24.653  1.00 65.51      B    C
ATOM   3868  CB  HIS   597     -72.680 -13.033 -24.678  1.00 67.23      B    C
ATOM   3869  CG  HIS   597     -73.646 -11.893 -24.586  1.00 69.06      B    C
ATOM   3870  CD2 HIS   597     -74.065 -11.162 -23.525  1.00 68.78      B    C
ATOM   3871  ND1 HIS   597     -74.330 -11.405 -25.680  1.00 69.46      B    N
ATOM   3872  CE1 HIS   597     -75.128 -10.425 -25.297  1.00 69.61      B    C
ATOM   3873  NE2 HIS   597     -74.987 -10.258 -23.993  1.00 69.86      B    N
ATOM   3874  C   HIS   597     -72.292 -15.492 -24.519  1.00 63.90      B    C
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ATOM   3875  O   HIS   597     -71.750 -15.713 -23.440  1.00 64.34      B    O
ATOM   3876  N   ALA   598     -72.000 -16.169 -25.622  1.00 61.69      B    N
ATOM   3877  CA  ALA   598     -71.035 -17.257 -25.606  1.00 59.24      B    C
ATOM   3878  CB  ALA   598     -71.733 -18.580 -25.919  1.00 59.25      B    C
ATOM   3879  C   ALA   598     -69.922 -17.002 -26.610  1.00 57.16      B    C
ATOM   3880  O   ALA   598     -70.179 -16.648 -27.759  1.00 57.27      B    O
ATOM   3881  N   SER   599     -68.684 -17.182 -26.172  1.00 55.41      B    N
ATOM   3882  CA  SER   599     -67.548 -17.103 -27.077  1.00 54.67      B    C
ATOM   3883  CB  SER   599     -66.342 -16.495 -26.349  1.00 54.38      B    C
ATOM   3884  OG  SER   599     -65.192 -16.463 -27.181  1.00 53.69      B    O
ATOM   3885  C   SER   599     -67.212 -18.506 -27.586  1.00 54.57      B    C
ATOM   3886  O   SER   599     -66.904 -19.401 -26.800  1.00 53.49      B    O
ATOM   3887  N   CYS   600     -67.282 -18.698 -28.901  1.00 55.25      B    N
ATOM   3888  CA  CYS   600     -66.994 -19.999 -29.498  1.00 58.00      B    C
ATOM   3889  C   CYS   600     -65.816 -19.885 -30.457  1.00 58.00      B    C
ATOM   3890  O   CYS   600     -65.784 -18.994 -31.300  1.00 58.78      B    O
ATOM   3891  CB  CYS   600     -68.218 -20.525 -30.259  1.00 59.36      B    C
ATOM   3892  SG  CYS   600     -69.778 -20.583 -29.309  1.00 63.79      B    S
ATOM   3893  N   CYS   601     -64.848 -20.784 -30.330  1.00 57.59      B    N
ATOM   3894  CA  CYS   601     -63.696 -20.762 -31.220  1.00 59.73      B    C
ATOM   3895  C   CYS   601     -63.475 -22.110 -31.892  1.00 59.37      B    C
ATOM   3896  O   CYS   601     -63.589 -23.160 -31.260  1.00 58.29      B    O
ATOM   3897  CB  CYS   601     -62.420 -20.395 -30.461  1.00 61.31      B    C
ATOM   3898  SG  CYS   601     -62.321 -18.753 -29.670  1.00 67.30      B    S
ATOM   3899  N   HIS   602     -63.147 -22.065 -33.177  1.00 59.49      B    N
ATOM   3900  CA  HIS   602     -62.668 -23.232 -33.895  1.00 60.04      B    C
ATOM   3901  CB  HIS   602     -63.299 -23.278 -35.293  1.00 62.97      B    C
ATOM   3902  CG  HIS   602     -62.908 -24.479 -36.099  1.00 66.97      B    C
ATOM   3903  CD2 HIS   602     -63.643 -25.517 -36.565  1.00 68.22      B    C
ATOM   3904  ND1 HIS   602     -61.615 -24.706 -36.521  1.00 68.55      B    N
ATOM   3905  CE1 HIS   602     -61.570 -25.832 -37.211  1.00 68.47      B    C
ATOM   3906  NE2 HIS   602     -62.788 -26.344 -37.253  1.00 68.91      B    N
ATOM   3907  C   HIS   602     -61.156 -23.094 -34.005  1.00 59.13      B    C
ATOM   3908  O   HIS   602     -60.658 -22.117 -34.560  1.00 58.95      B    O
ATOM   3909  N   ALA   603     -60.425 -24.059 -33.462  1.00 57.63      B    N
ATOM   3910  CA  ALA   603     -58.972 -24.045 -33.551  1.00 57.29      B    C
ATOM   3911  CB  ALA   603     -58.392 -23.137 -32.478  1.00 57.06      B    C
ATOM   3912  C   ALA   603     -58.414 -25.455 -33.405  1.00 58.27      B    C
ATOM   3913  O   ALA   603     -58.907 -26.254 -32.611  1.00 57.05      B    O
ATOM   3914  N   PRO   604     -57.363 -25.770 -34.170  1.00 59.55      B    N
ATOM   3915  CD  PRO   604     -56.540 -24.798 -34.914  1.00 60.80      B    C
ATOM   3916  CA  PRO   604     -56.843 -27.137 -34.279  1.00 59.96      B    C
ATOM   3917  CB  PRO   604     -55.719 -27.016 -35.308  1.00 60.62      B    C
ATOM   3918  CG  PRO   604     -55.294 -25.580 -35.237  1.00 61.97      B    C
ATOM   3919  C   PRO   604     -56.344 -27.701 -32.954  1.00 59.56      B    C
ATOM   3920  O   PRO   604     -55.343 -27.238 -32.409  1.00 60.04      B    O
ATOM   3921  N   GLY   605     -57.046 -28.708 -32.445  1.00 58.14      B    N
ATOM   3922  CA  GLY   605     -56.616 -29.359 -31.223  1.00 55.88      B    C
ATOM   3923  C   GLY   605     -57.024 -28.610 -29.969  1.00 54.81      B    C
ATOM   3924  O   GLY   605     -56.578 -28.938 -28.870  1.00 55.62      B    O
ATOM   3925  N   LEU   606     -57.871 -27.600 -30.126  1.00 52.76      B    N
ATOM   3926  CA  LEU   606     -58.380 -26.859 -28.980  1.00 50.04      B    C
ATOM   3927  CB  LEU   606     -59.100 -25.593 -29.445  1.00 48.66      B    C
ATOM   3928  CG  LEU   606     -59.707 -24.770 -28.305  1.00 47.30      B    C
ATOM   3929  CD1 LEU   606     -58.587 -24.229 -27.437  1.00 47.38      B    C
ATOM   3930  CD2 LEU   606     -60.546 -23.640 -28.862  1.00 46.45      B    C
ATOM   3931  C   LEU   606     -59.352 -27.715 -28.179  1.00 50.49      B    C
ATOM   3932  O   LEU   606     -60.373 -28.170 -28.705  1.00 50.29      B    O
ATOM   3933  N   GLU   607     -59.044 -27.932 -26.905  1.00 49.74      B    N
ATOM   3934  CA  GLU   607     -60.027 -28.511 -26.004  1.00 49.92      B    C
ATOM   3935  CB  GLU   607     -59.688 -29.975 -25.689  1.00 51.37      B    C
ATOM   3936  CG  GLU   607     -58.375 -30.208 -24.972  1.00 54.60      B    C
ATOM   3937  CD  GLU   607     -58.294 -31.599 -24.346  1.00 56.01      B    C
ATOM   3938  OE1 GLU   607     -57.200 -32.202 -24.360  1.00 56.47      B    O
ATOM   3939  OE2 GLU   607     -59.325 -32.089 -23.836  1.00 57.69      B    O
ATOM   3940  C   GLU   607     -60.151 -27.707 -24.720  1.00 49.55      B    C
ATOM   3941  O   GLU   607     -59.161 -27.218 -24.179  1.00 49.44      B    O
ATOM   3942  N   CYS   608     -61.384 -27.565 -24.249  1.00 49.05      B    N
ATOM   3943  CA  CYS   608     -61.677 -26.794 -23.050  1.00 49.93      B    C
ATOM   3944  C   CYS   608     -62.611 -27.598 -22.155  1.00 51.10      B    C
ATOM   3945  O   CYS   608     -63.400 -28.410 -22.641  1.00 51.03      B    O
ATOM   3946  CB  CYS   608     -62.374 -25.482 -23.406  1.00 49.04      B    C
ATOM   3947  SG  CYS   608     -61.522 -24.352 -24.553  1.00 51.08      B    S
ATOM   3948  N   LYS   609     -62.526 -27.357 -20.852  1.00 51.23      B    N
ATOM   3949  CA  LYS   609     -63.435 -27.976 -19.901  1.00 52.84      B    C
ATOM   3950  CB  LYS   609     -62.791 -29.219 -19.286  1.00 53.14      B    C
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ATOM   3951  CG  LYS   609     -61.523 -28.955 -18.501  1.00 56.59      B    C
ATOM   3952  CD  LYS   609     -60.931 -30.262 -17.996  1.00 59.65      B    C
ATOM   3953  CE  LYS   609     -59.776 -30.034 -17.035  1.00 62.05      B    C
ATOM   3954  NZ  LYS   609     -59.273 -31.333 -16.491  1.00 64.13      B    N
ATOM   3955  C   LYS   609     -63.807 -26.985 -18.809  1.00 53.60      B    C
ATOM   3956  O   LYS   609     -63.155 -25.953 -18.652  1.00 54.60      B    O
ATOM   3957  N   VAL   610     -64.859 -27.297 -18.059  1.00 53.79      B    N
ATOM   3958  CA  VAL   610     -65.302 -26.448 -16.961  1.00 54.51      B    C
ATOM   3959  CB  VAL   610     -66.830 -26.320 -16.947  1.00 53.63      B    C
ATOM   3960  CG1 VAL   610     -67.271 -25.529 -15.731  1.00 53.41      B    C
ATOM   3961  CG2 VAL   610     -67.302 -25.645 -18.218  1.00 53.06      B    C
ATOM   3962  C   VAL   610     -64.856 -26.978 -15.603  1.00 56.27      B    C
ATOM   3963  O   VAL   610     -64.852 -28.186 -15.371  1.00 56.36      B    O
ATOM   3964  N   LYS   611     -64.472 -26.064 -14.714  1.00 58.50      B    N
ATOM   3965  CA  LYS   611     -64.239 -26.387 -13.308  1.00 60.09      B    C
ATOM   3966  CB  LYS   611     -62.755 -26.257 -12.963  1.00 60.15      B    C
ATOM   3967  CG  LYS   611     -61.822 -26.853 -14.002  1.00 62.60      B    C
ATOM   3968  CD  LYS   611     -60.861 -27.869 -13.396  1.00 64.20      B    C
ATOM   3969  CE  LYS   611     -59.980 -27.252 -12.319  1.00 63.58      B    C
ATOM   3970  NZ  LYS   611     -60.714 -27.067 -11.039  1.00 62.59      B    N
ATOM   3971  C   LYS   611     -65.042 -25.421 -12.449  1.00 61.71      B    C
ATOM   3972  O   LYS   611     -65.139 -24.235 -12.764  1.00 62.06      B    O
ATOM   3973  N   GLU   612     -65.622 -25.929 -11.369  1.00 63.68      B    N
ATOM   3974  CA  GLU   612     -66.417 -25.101 -10.474  1.00 66.24      B    C
ATOM   3975  CB  GLU   612     -67.877 -25.550 -10.484  1.00 67.02      B    C
ATOM   3976  CG  GLU   612     -68.589 -25.366 -11.802  1.00 70.34      B    C
ATOM   3977  CD  GLU   612     -70.077 -25.645 -11.685  1.00 72.40      B    C
ATOM   3978  OE1 GLU   612     -70.810 -25.417 -12.671  1.00 73.78      B    O
ATOM   3979  OE2 GLU   612     -70.512 -26.091 -10.600  1.00 73.79      B    O
ATOM   3980  C   GLU   612     -65.901 -25.158  -9.042  1.00 68.03      B    C
ATOM   3981  O   GLU   612     -65.141 -26.056  -8.675  1.00 66.96      B    O
ATOM   3982  N   HIS   613     -66.327 -24.185  -8.241  1.00 70.17      B    N
ATOM   3983  CA  HIS   613     -66.140 -24.226  -6.798  1.00 72.95      B    C
ATOM   3984  CB  HIS   613     -64.726 -23.777  -6.426  1.00 73.69      B    C
ATOM   3985  CG  HIS   613     -64.399 -23.948  -4.975  1.00 74.96      B    C
ATOM   3986  CD2 HIS   613     -64.782 -23.243  -3.885  1.00 75.11      B    C
ATOM   3987  ND1 HIS   613     -63.574 -24.951  -4.512  1.00 76.02      B    N
ATOM   3988  CE1 HIS   613     -63.462 -24.856  -3.199  1.00 75.34      B    C
ATOM   3989  NE2 HIS   613     -64.185 -23.827  -2.793  1.00 75.59      B    N
ATOM   3990  C   HIS   613     -67.162 -23.310  -6.142  1.00 75.10      B    C
ATOM   3991  O   HIS   613     -67.342 -22.165  -6.557  1.00 74.72      B    O
ATOM   3992  N   GLY   614     -67.840 -23.824  -5.124  1.00 78.12      B    N
ATOM   3993  CA  GLY   614     -68.806 -23.017  -4.404  1.00 81.74      B    C
ATOM   3994  C   GLY   614     -68.482 -22.968  -2.926  1.00 84.51      B    C
ATOM   3995  O   GLY   614     -67.916 -23.913  -2.376  1.00 84.08      B    O
ATOM   3996  N   ILE   615     -68.829 -21.860  -2.281  1.00 87.55      B    N
ATOM   3997  CA  ILE   615     -68.696 -21.749  -0.836  1.00 90.95      B    C
ATOM   3998  CB  ILE   615     -67.545 -20.795  -0.445  1.00 90.68      B    C
ATOM   3999  CG2 ILE   615     -67.478 -20.648   1.070  1.00 90.91      B    C
ATOM   4000  CG1 ILE   615     -66.218 -21.342  -0.970  1.00 90.52      B    C
ATOM   4001  CD1 ILE   615     -65.019 -20.498  -0.596  1.00 91.21      B    C
ATOM   4002  C   ILE   615     -70.001 -21.236  -0.240  1.00 93.64      B    C
ATOM   4003  O   ILE   615     -70.584 -20.268  -0.734  1.00 93.43      B    O
ATOM   4004  N   PRO   616     -70.480 -21.892   0.829  1.00 96.49      B    N
ATOM   4005  CD  PRO   616     -69.870 -23.095   1.420  1.00 97.34      B    C
ATOM   4006  CA  PRO   616     -71.741 -21.540   1.490  1.00 98.31      B    C
ATOM   4007  CB  PRO   616     -71.878 -22.590   2.592  1.00 98.23      B    C
ATOM   4008  CG  PRO   616     -71.022 -23.732   2.137  1.00 97.91      B    C
ATOM   4009  C   PRO   616     -71.695 -20.127   2.056  1.00100.07      B    C
ATOM   4010  O   PRO   616     -72.555 -19.296   1.757  1.00100.25      B    O
ATOM   4011  N   ALA   617     -70.681 -19.863   2.875  1.00101.81      B    N
ATOM   4012  CA  ALA   617     -70.478 -18.536   3.440  1.00103.72      B    C
ATOM   4013  CB  ALA   617     -70.126 -18.644   4.920  1.00103.96      B    C
ATOM   4014  C   ALA   617     -69.369 -17.810   2.690  1.00104.59      B    C
ATOM   4015  O   ALA   617     -68.188 -17.958   3.006  1.00104.51      B    O
ATOM   4016  N   PRO   618     -69.740 -17.009   1.681  1.00105.43      B    N
ATOM   4017  CD  PRO   618     -71.117 -16.737   1.232  1.00105.79      B    C
ATOM   4018  CA  PRO   618     -68.748 -16.235   0.929  1.00105.78      B    C
ATOM   4019  CB  PRO   618     -69.568 -15.579  -0.181  1.00106.12      B    C
ATOM   4020  CG  PRO   618     -70.960 -15.521   0.363  1.00106.24      B    C
ATOM   4021  C   PRO   618     -68.052 -15.211   1.821  1.00105.78      B    C
ATOM   4022  O   PRO   618     -68.610 -14.157   2.131  1.00105.85      B    O
ATOM   4023  N   GLN   619     -66.829 -15.536   2.228  1.00105.22      B    N
ATOM   4024  CA  GLN   619     -66.071 -14.709   3.158  1.00104.16      B    C
ATOM   4025  CB  GLN   619     -64.819 -15.465   3.610  1.00106.08      B    C
ATOM   4026  CG  GLN   619     -63.990 -14.740   4.651  1.00109.04      B    C
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ATOM	4027	CD	GLN	619	-62.910	-15.623	5.243	1.00	110.77	B	C
ATOM	4028	OE1	GLN	619	-62.885	-15.867	6.451	1.00	111.68	B	O
ATOM	4029	NE2	GLN	619	-62.010	-16.111	4.394	1.00	111.21	B	N
ATOM	4030	C	GLN	619	-65.675	-13.369	2.532	1.00	101.86	B	C
ATOM	4031	O	GLN	619	-65.431	-12.389	3.237	1.00	102.25	B	O
ATOM	4032	N	GLY	620	-65.619	-13.337	1.204	1.00	98.47	B	N
ATOM	4033	CA	GLY	620	-65.264	-12.120	0.497	1.00	93.13	B	C
ATOM	4034	C	GLY	620	-64.971	-12.411	-0.964	1.00	89.75	B	C
ATOM	4035	O	GLY	620	-65.153	-11.556	-1.832	1.00	89.35	B	O
ATOM	4036	N	GLN	621	-64.518	-13.630	-1.236	1.00	86.41	B	N
ATOM	4037	CA	GLN	621	-64.285	-14.065	-2.604	1.00	81.21	B	C
ATOM	4038	CB	GLN	621	-62.963	-13.499	-3.122	1.00	80.76	B	C
ATOM	4039	CG	GLN	621	-61.738	-14.039	-2.412	1.00	79.63	B	C
ATOM	4040	CD	GLN	621	-60.449	-13.607	-3.080	1.00	80.09	B	C
ATOM	4041	OE1	GLN	621	-60.360	-12.509	-3.631	1.00	79.55	B	O
ATOM	4042	NE2	GLN	621	-59.441	-14.471	-3.036	1.00	79.53	B	N
ATOM	4043	C	GLN	621	-64.261	-15.585	-2.717	1.00	78.31	B	C
ATOM	4044	O	GLN	621	-63.759	-16.280	-1.833	1.00	77.75	B	O
ATOM	4045	N	VAL	622	-64.810	-16.094	-3.813	1.00	74.56	B	N
ATOM	4046	CA	VAL	622	-64.715	-17.511	-4.135	1.00	70.71	B	C
ATOM	4047	CB	VAL	622	-66.096	-18.084	-4.517	1.00	70.57	B	C
ATOM	4048	CG1	VAL	622	-65.987	-19.574	-4.772	1.00	70.29	B	C
ATOM	4049	CG2	VAL	622	-67.099	-17.800	-3.413	1.00	70.77	B	C
ATOM	4050	C	VAL	622	-63.773	-17.661	-5.323	1.00	68.02	B	C
ATOM	4051	O	VAL	622	-63.844	-16.883	-6.273	1.00	67.71	B	O
ATOM	4052	N	THR	623	-62.889	-18.652	-5.272	1.00	65.11	B	N
ATOM	4053	CA	THR	623	-61.955	-18.871	-6.371	1.00	62.26	B	C
ATOM	4054	CB	THR	623	-60.535	-18.421	-6.005	1.00	62.27	B	C
ATOM	4055	OG1	THR	623	-60.016	-19.274	-4.979	1.00	62.80	B	O
ATOM	4056	CG2	THR	623	-60.539	-16.982	-5.515	1.00	61.70	B	C
ATOM	4057	C	THR	623	-61.868	-20.329	-6.808	1.00	61.06	B	C
ATOM	4058	O	THR	623	-62.154	-21.243	-6.037	1.00	60.46	B	O
ATOM	4059	N	VAL	624	-61.463	-20.528	-8.059	1.00	58.95	B	N
ATOM	4060	CA	VAL	624	-61.115	-21.845	-8.570	1.00	56.86	B	C
ATOM	4061	CB	VAL	624	-62.357	-22.553	-9.183	1.00	56.41	B	C
ATOM	4062	CG1	VAL	624	-62.927	-21.727	-10.326	1.00	56.31	B	C
ATOM	4063	CG2	VAL	624	-61.979	-23.939	-9.672	1.00	55.46	B	C
ATOM	4064	C	VAL	624	-60.026	-21.684	-9.634	1.00	55.91	B	C
ATOM	4065	O	VAL	624	-60.034	-20.723	-10.399	1.00	54.97	B	O
ATOM	4066	N	ALA	625	-59.085	-22.620	-9.674	1.00	55.07	B	N
ATOM	4067	CA	ALA	625	-57.945	-22.504	-10.573	1.00	56.43	B	C
ATOM	4068	CB	ALA	625	-56.656	-22.419	-9.766	1.00	55.50	B	C
ATOM	4069	C	ALA	625	-57.869	-23.673	-11.548	1.00	57.84	B	C
ATOM	4070	O	ALA	625	-58.319	-24.776	-11.241	1.00	58.31	B	O
ATOM	4071	N	CYS	626	-57.294	-23.424	-12.722	1.00	57.81	B	N
ATOM	4072	CA	CYS	626	-57.054	-24.479	-13.692	1.00	59.13	B	C
ATOM	4073	C	CYS	626	-55.829	-25.288	-13.308	1.00	61.75	B	C
ATOM	4074	O	CYS	626	-54.831	-24.741	-12.845	1.00	62.27	B	O
ATOM	4075	CB	CYS	626	-56.840	-23.896	-15.088	1.00	56.64	B	C
ATOM	4076	SG	CYS	626	-58.258	-22.975	-15.751	1.00	57.44	B	S
ATOM	4077	N	GLU	627	-55.911	-26.596	-13.525	1.00	64.64	B	N
ATOM	4078	CA	GLU	627	-54.810	-27.501	-13.241	1.00	67.51	B	C
ATOM	4079	CB	GLU	627	-55.229	-28.938	-13.553	1.00	70.84	B	C
ATOM	4080	CG	GLU	627	-56.560	-29.333	-12.944	1.00	76.92	B	C
ATOM	4081	CD	GLU	627	-57.452	-30.066	-13.929	1.00	81.54	B	C
ATOM	4082	OE1	GLU	627	-57.100	-31.203	-14.322	1.00	83.33	B	O
ATOM	4083	OE2	GLU	627	-58.503	-29.502	-14.311	1.00	83.28	B	O
ATOM	4084	C	GLU	627	-53.576	-27.147	-14.061	1.00	67.30	B	C
ATOM	4085	O	GLU	627	-53.669	-26.488	-15.097	1.00	67.09	B	O
ATOM	4086	N	GLU	628	-52.420	-27.595	-13.587	1.00	67.21	B	N
ATOM	4087	CA	GLU	628	-51.178	-27.462	-14.333	1.00	67.26	B	C
ATOM	4088	CB	GLU	628	-50.071	-28.260	-13.638	1.00	71.29	B	C
ATOM	4089	CG	GLU	628	-48.693	-28.097	-14.259	1.00	76.58	B	C
ATOM	4090	CD	GLU	628	-48.155	-26.687	-14.105	1.00	80.28	B	C
ATOM	4091	OE1	GLU	628	-47.423	-26.433	-13.121	1.00	82.51	B	O
ATOM	4092	OE2	GLU	628	-48.466	-25.833	-14.965	1.00	81.67	B	O
ATOM	4093	C	GLU	628	-51.377	-27.986	-15.751	1.00	65.26	B	C
ATOM	4094	O	GLU	628	-51.897	-29.084	-15.943	1.00	65.44	B	O
ATOM	4095	N	GLY	629	-50.963	-27.198	-16.738	1.00	62.46	B	N
ATOM	4096	CA	GLY	629	-51.086	-27.618	-18.123	1.00	58.56	B	C
ATOM	4097	C	GLY	629	-52.260	-26.986	-18.853	1.00	56.59	B	C
ATOM	4098	O	GLY	629	-52.297	-26.971	-20.085	1.00	56.28	B	O
ATOM	4099	N	TRP	630	-53.220	-26.466	-18.094	1.00	54.07	B	N
ATOM	4100	CA	TRP	630	-54.389	-25.811	-18.670	1.00	52.31	B	C
ATOM	4101	CB	TRP	630	-55.667	-26.350	-18.020	1.00	53.37	B	C
ATOM	4102	CG	TRP	630	-55.912	-27.798	-18.298	1.00	56.24	B	C

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ATOM   4103  CD2 TRP   630     -56.828 -28.348 -19.253  1.00 56.84      B    C
ATOM   4104  CE2 TRP   630     -56.728 -29.752 -19.168  1.00 57.44      B    C
ATOM   4105  CE3 TRP   630     -57.724 -27.791 -20.170  1.00 56.63      B    C
ATOM   4106  CD1 TRP   630     -55.310 -28.861 -17.691  1.00 56.81      B    C
ATOM   4107  NE1 TRP   630     -55.794 -30.037 -18.206  1.00 57.15      B    N
ATOM   4108  CZ2 TRP   630     -57.489 -30.608 -19.964  1.00 56.00      B    C
ATOM   4109  CZ3 TRP   630     -58.482 -28.644 -20.962  1.00 58.10      B    C
ATOM   4110  CH2 TRP   630     -58.357 -30.037 -20.852  1.00 56.56      B    C
ATOM   4111  C   TRP   630     -54.325 -24.295 -18.487  1.00 50.00      B    C
ATOM   4112  O   TRP   630     -53.696 -23.802 -17.555  1.00 49.79      B    O
ATOM   4113  N   THR   631     -54.987 -23.563 -19.378  1.00 46.86      B    N
ATOM   4114  CA  THR   631     -55.000 -22.106 -19.313  1.00 42.14      B    C
ATOM   4115  CB  THR   631     -54.472 -21.490 -20.628  1.00 41.03      B    C
ATOM   4116  OG1 THR   631     -53.126 -21.924 -20.850  1.00 38.22      B    O
ATOM   4117  CG2 THR   631     -54.510 -19.966 -20.568  1.00 39.04      B    C
ATOM   4118  C   THR   631     -56.415 -21.601 -19.069  1.00 41.59      B    C
ATOM   4119  O   THR   631     -57.349 -21.977 -19.777  1.00 41.97      B    O
ATOM   4120  N   LEU   632     -56.573 -20.743 -18.068  1.00 40.02      B    N
ATOM   4121  CA  LEU   632     -57.866 -20.126 -17.804  1.00 39.61      B    C
ATOM   4122  CB  LEU   632     -57.799 -19.316 -16.509  1.00 38.76      B    C
ATOM   4123  CG  LEU   632     -59.077 -18.608 -16.057  1.00 42.02      B    C
ATOM   4124  CD1 LEU   632     -60.205 -19.619 -15.901  1.00 41.47      B    C
ATOM   4125  CD2 LEU   632     -58.816 -17.899 -14.731  1.00 41.91      B    C
ATOM   4126  C   LEU   632     -58.232 -19.212 -18.973  1.00 40.31      B    C
ATOM   4127  O   LEU   632     -57.476 -18.301 -19.311  1.00 39.62      B    O
ATOM   4128  N   THR   633     -59.377 -19.465 -19.601  1.00 40.05      B    N
ATOM   4129  CA  THR   633     -59.849 -18.597 -20.676  1.00 40.37      B    C
ATOM   4130  CB  THR   633     -60.166 -19.385 -21.974  1.00 40.11      B    C
ATOM   4131  OG1 THR   633     -61.267 -20.270 -21.740  1.00 42.19      B    O
ATOM   4132  CG2 THR   633     -58.958 -20.187 -22.430  1.00 39.36      B    C
ATOM   4133  C   THR   633     -61.113 -17.855 -20.267  1.00 40.80      B    C
ATOM   4134  O   THR   633     -61.408 -16.790 -20.797  1.00 42.82      B    O
ATOM   4135  N   GLY   634     -61.868 -18.423 -19.335  1.00 41.84      B    N
ATOM   4136  CA  GLY   634     -63.129 -17.816 -18.949  1.00 43.05      B    C
ATOM   4137  C   GLY   634     -63.344 -17.877 -17.454  1.00 44.20      B    C
ATOM   4138  O   GLY   634     -62.980 -18.859 -16.813  1.00 44.52      B    O
ATOM   4139  N   CYS   635     -63.927 -16.823 -16.892  1.00 46.22      B    N
ATOM   4140  CA  CYS   635     -64.151 -16.751 -15.452  1.00 49.45      B    C
ATOM   4141  C   CYS   635     -65.479 -16.072 -15.176  1.00 50.61      B    C
ATOM   4142  O   CYS   635     -65.802 -15.046 -15.774  1.00 51.08      B    O
ATOM   4143  CB  CYS   635     -63.019 -15.979 -14.770  1.00 49.68      B    C
ATOM   4144  SG  CYS   635     -63.191 -15.778 -12.960  1.00 54.65      B    S
ATOM   4145  N   SER   636     -66.245 -16.647 -14.261  1.00 52.89      B    N
ATOM   4146  CA  SER   636     -67.661 -16.339 -14.186  1.00 56.15      B    C
ATOM   4147  CB  SER   636     -68.390 -17.073 -15.317  1.00 56.84      B    C
ATOM   4148  OG  SER   636     -69.709 -16.597 -15.483  1.00 59.24      B    O
ATOM   4149  C   SER   636     -68.215 -16.773 -12.839  1.00 57.63      B    C
ATOM   4150  O   SER   636     -67.656 -17.654 -12.187  1.00 57.25      B    O
ATOM   4151  N   ALA   637     -69.310 -16.148 -12.422  1.00 60.91      B    N
ATOM   4152  CA  ALA   637     -70.052 -16.611 -11.253  1.00 64.91      B    C
ATOM   4153  CB  ALA   637     -70.318 -15.446 -10.305  1.00 62.85      B    C
ATOM   4154  C   ALA   637     -71.373 -17.231 -11.704  1.00 68.14      B    C
ATOM   4155  O   ALA   637     -71.992 -16.760 -12.664  1.00 66.65      B    O
ATOM   4156  N   LEU   638     -71.803 -18.286 -11.016  1.00 72.27      B    N
ATOM   4157  CA  LEU   638     -73.124 -18.855 -11.268  1.00 76.39      B    C
ATOM   4158  CB  LEU   638     -73.332 -20.127 -10.447  1.00 76.34      B    C
ATOM   4159  CG  LEU   638     -72.627 -21.379 -10.965  1.00 77.63      B    C
ATOM   4160  CD1 LEU   638     -73.027 -22.581 -10.120  1.00 77.21      B    C
ATOM   4161  CD2 LEU   638     -73.002 -21.604 -12.420  1.00 76.50      B    C
ATOM   4162  C   LEU   638     -74.195 -17.836 -10.907  1.00 79.19      B    C
ATOM   4163  O   LEU   638     -74.146 -17.221  -9.840  1.00 79.59      B    O
ATOM   4164  N   PRO   639     -75.180 -17.645 -11.797  1.00 81.71      B    N
ATOM   4165  CD  PRO   639     -75.359 -18.404 -13.047  1.00 81.80      B    C
ATOM   4166  CA  PRO   639     -76.228 -16.636 -11.609  1.00 84.07      B    C
ATOM   4167  CB  PRO   639     -76.986 -16.652 -12.936  1.00 83.38      B    C
ATOM   4168  CG  PRO   639     -76.738 -18.011 -13.494  1.00 82.31      B    C
ATOM   4169  C   PRO   639     -77.139 -16.934 -10.421  1.00 86.54      B    C
ATOM   4170  O   PRO   639     -77.156 -18.053  -9.904  1.00 86.12      B    O
ATOM   4171  N   GLY   640     -77.887 -15.921  -9.989  1.00 89.30      B    N
ATOM   4172  CA  GLY   640     -78.889 -16.126  -8.958  1.00 92.85      B    C
ATOM   4173  C   GLY   640     -78.395 -15.864  -7.549  1.00 95.28      B    C
ATOM   4174  O   GLY   640     -78.954 -16.386  -6.583  1.00 95.56      B    O
ATOM   4175  N   THR   641     -77.345 -15.059  -7.426  1.00 97.49      B    N
ATOM   4176  CA  THR   641     -76.803 -14.714  -6.115  1.00 99.35      B    C
ATOM   4177  CB  THR   641     -75.305 -15.069  -6.020  1.00100.06      B    C
ATOM   4178  OG1 THR   641     -75.106 -16.425  -6.444  1.00100.46      B    O
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ATOM   4179  CG2 THR   641     -74.817 -14.922  -4.584  1.00100.75      B    C
ATOM   4180  C   THR   641     -76.969 -13.219  -5.859  1.00 99.93      B    C
ATOM   4181  O   THR   641     -77.280 -12.800  -4.741  1.00100.09      B    O
ATOM   4182  N   SER   642     -76.758 -12.426  -6.907  1.00100.54      B    N
ATOM   4183  CA  SER   642     -76.919 -10.976  -6.847  1.00100.83      B    C
ATOM   4184  CB  SER   642     -78.311 -10.616  -6.311  1.00101.56      B    C
ATOM   4185  OG  SER   642     -79.322 -10.989  -7.233  1.00102.34      B    O
ATOM   4186  C   SER   642     -75.846 -10.293  -6.000  1.00100.09      B    C
ATOM   4187  O   SER   642     -75.262  -9.292  -6.421  1.00100.47      B    O
ATOM   4188  N   HIS   643     -75.587 -10.835  -4.813  1.00 98.31      B    N
ATOM   4189  CA  HIS   643     -74.555 -10.297  -3.931  1.00 95.91      B    C
ATOM   4190  CB  HIS   643     -74.722 -10.862  -2.519  1.00 98.80      B    C
ATOM   4191  CG  HIS   643     -75.936 -10.348  -1.808  1.00102.15      B    C
ATOM   4192  CD2 HIS   643     -76.772 -10.941  -0.923  1.00103.19      B    C
ATOM   4193  ND1 HIS   643     -76.411  -9.064  -1.982  1.00103.40      B    N
ATOM   4194  CE1 HIS   643     -77.485  -8.889  -1.233  1.00103.99      B    C
ATOM   4195  NE2 HIS   643     -77.726 -10.012  -0.581  1.00104.35      B    N
ATOM   4196  C   HIS   643     -73.158 -10.606  -4.463  1.00 92.18      B    C
ATOM   4197  O   HIS   643     -72.247 -10.950  -3.707  1.00 92.08      B    O
ATOM   4198  N   VAL   644     -73.012 -10.481  -5.779  1.00 86.97      B    N
ATOM   4199  CA  VAL   644     -71.728 -10.628  -6.450  1.00 81.49      B    C
ATOM   4200  CB  VAL   644     -71.815 -11.675  -7.583  1.00 81.51      B    C
ATOM   4201  CG1 VAL   644     -70.522 -11.693  -8.383  1.00 80.44      B    C
ATOM   4202  CG2 VAL   644     -72.090 -13.049  -6.993  1.00 81.28      B    C
ATOM   4203  C   VAL   644     -71.298  -9.289  -7.042  1.00 77.33      B    C
ATOM   4204  O   VAL   644     -72.070  -8.629  -7.739  1.00 75.98      B    O
ATOM   4205  N   LEU   645     -70.061  -8.895  -6.756  1.00 72.69      B    N
ATOM   4206  CA  LEU   645     -69.520  -7.632  -7.242  1.00 68.29      B    C
ATOM   4207  CB  LEU   645     -68.409  -7.148  -6.307  1.00 68.02      B    C
ATOM   4208  CG  LEU   645     -68.789  -7.055  -4.827  1.00 67.91      B    C
ATOM   4209  CD1 LEU   645     -67.603  -6.554  -4.015  1.00 67.93      B    C
ATOM   4210  CD2 LEU   645     -69.979  -6.125  -4.668  1.00 67.53      B    C
ATOM   4211  C   LEU   645     -68.971  -7.801  -8.657  1.00 65.15      B    C
ATOM   4212  O   LEU   645     -68.985  -6.864  -9.457  1.00 64.28      B    O
ATOM   4213  N   GLY   646     -68.493  -9.006  -8.957  1.00 61.10      B    N
ATOM   4214  CA  GLY   646     -67.971  -9.289 -10.280  1.00 56.54      B    C
ATOM   4215  C   GLY   646     -66.980 -10.434 -10.276  1.00 53.72      B    C
ATOM   4216  O   GLY   646     -66.789 -11.101  -9.259  1.00 53.37      B    O
ATOM   4217  N   ALA   647     -66.342 -10.665 -11.417  1.00 50.95      B    N
ATOM   4218  CA  ALA   647     -65.407 -11.776 -11.555  1.00 47.43      B    C
ATOM   4219  CB  ALA   647     -66.143 -13.011 -12.073  1.00 46.90      B    C
ATOM   4220  C   ALA   647     -64.276 -11.404 -12.510  1.00 45.07      B    C
ATOM   4221  O   ALA   647     -64.484 -10.664 -13.471  1.00 45.31      B    O
ATOM   4222  N   TYR   648     -63.081 -11.918 -12.243  1.00 41.18      B    N
ATOM   4223  CA  TYR   648     -61.950 -11.701 -13.132  1.00 40.43      B    C
ATOM   4224  CB  TYR   648     -61.315 -10.324 -12.880  1.00 38.89      B    C
ATOM   4225  CG  TYR   648     -61.108  -9.998 -11.418  1.00 37.05      B    C
ATOM   4226  CD1 TYR   648     -59.993 -10.455 -10.733  1.00 37.52      B    C
ATOM   4227  CE1 TYR   648     -59.825 -10.189  -9.381  1.00 37.90      B    C
ATOM   4228  CD2 TYR   648     -62.051  -9.262 -10.714  1.00 37.16      B    C
ATOM   4229  CE2 TYR   648     -61.891  -8.992  -9.370  1.00 38.07      B    C
ATOM   4230  CZ  TYR   648     -60.778  -9.460  -8.708  1.00 37.64      B    C
ATOM   4231  OH  TYR   648     -60.630  -9.197  -7.362  1.00 40.14      B    O
ATOM   4232  C   TYR   648     -60.905 -12.786 -12.953  1.00 40.67      B    C
ATOM   4233  O   TYR   648     -60.817 -13.412 -11.896  1.00 40.70      B    O
ATOM   4234  N   ALA   649     -60.112 -13.001 -13.995  1.00 40.21      B    N
ATOM   4235  CA  ALA   649     -59.012 -13.949 -13.936  1.00 40.94      B    C
ATOM   4236  CB  ALA   649     -58.743 -14.513 -15.329  1.00 40.44      B    C
ATOM   4237  C   ALA   649     -57.755 -13.273 -13.395  1.00 41.98      B    C
ATOM   4238  O   ALA   649     -57.403 -12.169 -13.813  1.00 42.47      B    O
ATOM   4239  N   VAL   650     -57.089 -13.942 -12.460  1.00 41.96      B    N
ATOM   4240  CA  VAL   650     -55.743 -13.576 -12.056  1.00 41.03      B    C
ATOM   4241  CB  VAL   650     -55.659 -13.331 -10.528  1.00 41.39      B    C
ATOM   4242  CG1 VAL   650     -54.239 -12.904 -10.141  1.00 36.26      B    C
ATOM   4243  CG2 VAL   650     -56.675 -12.274 -10.115  1.00 38.03      B    C
ATOM   4244  C   VAL   650     -54.814 -14.728 -12.429  1.00 42.88      B    C
ATOM   4245  O   VAL   650     -54.775 -15.751 -11.746  1.00 43.57      B    O
ATOM   4246  N   ASP   651     -54.068 -14.555 -13.514  1.00 45.00      B    N
ATOM   4247  CA  ASP   651     -53.278 -15.641 -14.084  1.00 47.26      B    C
ATOM   4248  CB  ASP   651     -52.219 -16.108 -13.086  1.00 50.04      B    C
ATOM   4249  CG  ASP   651     -51.177 -17.014 -13.721  1.00 55.64      B    C
ATOM   4250  OD1 ASP   651     -51.113 -17.076 -14.973  1.00 57.69      B    O
ATOM   4251  OD2 ASP   651     -50.420 -17.664 -12.965  1.00 57.35      B    O
ATOM   4252  C   ASP   651     -54.206 -16.803 -14.451  1.00 47.02      B    C
ATOM   4253  O   ASP   651     -55.093 -16.645 -15.288  1.00 45.87      B    O
ATOM   4254  N   ASN   652     -54.010 -17.959 -13.821  1.00 46.56      B    N
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ATOM   4255  CA  ASN   652      -54.874 -19.113 -14.064  1.00 47.62      B    C
ATOM   4256  CB  ASN   652      -54.041 -20.381 -14.269  1.00 46.87      B    C
ATOM   4257  CG  ASN   652      -53.552 -20.527 -15.695  1.00 48.35      B    C
ATOM   4258  OD1 ASN   652      -54.179 -20.029 -16.633  1.00 47.24      B    O
ATOM   4259  ND2 ASN   652      -52.426 -21.207 -15.869  1.00 49.09      B    N
ATOM   4260  C   ASN   652      -55.900 -19.366 -12.969  1.00 47.56      B    C
ATOM   4261  O   ASN   652      -56.470 -20.453 -12.895  1.00 48.79      B    O
ATOM   4262  N   THR   653      -56.140 -18.374 -12.118  1.00 46.14      B    N
ATOM   4263  CA  THR   653      -57.173 -18.506 -11.102  1.00 44.77      B    C
ATOM   4264  CB  THR   653      -56.648 -18.136  -9.704  1.00 44.11      B    C
ATOM   4265  OG1 THR   653      -55.569 -19.011  -9.353  1.00 44.54      B    O
ATOM   4266  CG2 THR   653      -57.760 -18.266  -8.670  1.00 42.08      B    C
ATOM   4267  C   THR   653      -58.366 -17.622 -11.416  1.00 46.06      B    C
ATOM   4268  O   THR   653      -58.213 -16.455 -11.769  1.00 46.42      B    O
ATOM   4269  N   CYS   654      -59.556 -18.191 -11.288  1.00 46.34      B    N
ATOM   4270  CA  CYS   654      -60.790 -17.449 -11.481  1.00 47.21      B    C
ATOM   4271  C   CYS   654      -61.240 -16.909 -10.132  1.00 47.79      B    C
ATOM   4272  O   CYS   654      -61.341 -17.657  -9.161  1.00 48.70      B    O
ATOM   4273  CB  CYS   654      -61.857 -18.375 -12.065  1.00 47.10      B    C
ATOM   4274  SG  CYS   654      -63.519 -17.664 -12.264  1.00 49.57      B    S
ATOM   4275  N   VAL   655      -61.502 -15.609 -10.074  1.00 47.25      B    N
ATOM   4276  CA  VAL   655      -61.840 -14.952  -8.821  1.00 46.56      B    C
ATOM   4277  CB  VAL   655      -60.806 -13.854  -8.478  1.00 46.21      B    C
ATOM   4278  CG1 VAL   655      -61.252 -13.070  -7.252  1.00 45.65      B    C
ATOM   4279  CG2 VAL   655      -59.449 -14.483  -8.231  1.00 43.82      B    C
ATOM   4280  C   VAL   655      -63.221 -14.326  -8.914  1.00 48.25      B    C
ATOM   4281  O   VAL   655      -63.492 -13.520  -9.804  1.00 48.62      B    O
ATOM   4282  N   VAL   656      -64.098 -14.710  -7.995  1.00 50.22      B    N
ATOM   4283  CA  VAL   656      -65.419 -14.105  -7.906  1.00 52.83      B    C
ATOM   4284  CB  VAL   656      -66.527 -15.182  -7.912  1.00 52.93      B    C
ATOM   4285  CG1 VAL   656      -67.889 -14.530  -7.732  1.00 51.33      B    C
ATOM   4286  CG2 VAL   656      -66.485 -15.959  -9.227  1.00 50.98      B    C
ATOM   4287  C   VAL   656      -65.512 -13.288  -6.621  1.00 55.73      B    C
ATOM   4288  O   VAL   656      -65.155 -13.767  -5.544  1.00 55.45      B    O
ATOM   4289  N   ARG   657      -65.984 -12.051  -6.742  1.00 58.13      B    N
ATOM   4290  CA  ARG   657      -66.028 -11.146  -5.602  1.00 62.52      B    C
ATOM   4291  CB  ARG   657      -65.441  -9.786  -5.988  1.00 61.88      B    C
ATOM   4292  CG  ARG   657      -63.976  -9.841  -6.376  1.00 61.18      B    C
ATOM   4293  CD  ARG   657      -63.099 -10.001  -5.152  1.00 61.37      B    C
ATOM   4294  NE  ARG   657      -63.292  -8.886  -4.232  1.00 61.93      B    N
ATOM   4295  CZ  ARG   657      -62.779  -7.674  -4.415  1.00 63.19      B    C
ATOM   4296  NH1 ARG   657      -63.010  -6.715  -3.527  1.00 63.51      B    N
ATOM   4297  NH2 ARG   657      -62.032  -7.419  -5.484  1.00 62.31      B    N
ATOM   4298  C   ARG   657      -67.446 -10.964  -5.082  1.00 65.59      B    C
ATOM   4299  O   ARG   657      -68.333 -10.523  -5.811  1.00 65.58      B    O
ATOM   4300  N   SER   658      -67.647 -11.305  -3.813  1.00 70.15      B    N
ATOM   4301  CA  SER   658      -68.945 -11.149  -3.162  1.00 75.05      B    C
ATOM   4302  CB  SER   658      -69.352 -12.461  -2.488  1.00 75.03      B    C
ATOM   4303  OG  SER   658      -68.362 -12.879  -1.561  1.00 76.50      B    O
ATOM   4304  C   SER   658      -68.876 -10.036  -2.118  1.00 77.90      B    C
ATOM   4305  O   SER   658      -67.807  -9.754  -1.574  1.00 77.97      B    O
ATOM   4306  N   ARG   659      -70.014  -9.409  -1.837  1.00 81.25      B    N
ATOM   4307  CA  ARG   659      -70.058  -8.343  -0.843  1.00 84.24      B    C
ATOM   4308  CB  ARG   659      -71.156  -7.333  -1.185  1.00 86.82      B    C
ATOM   4309  CG  ARG   659      -71.021  -6.010  -0.441  1.00 90.54      B    C
ATOM   4310  CD  ARG   659      -72.248  -5.131  -0.625  1.00 93.57      B    C
ATOM   4311  NE  ARG   659      -73.441  -5.738  -0.040  1.00 96.92      B    N
ATOM   4312  CZ  ARG   659      -74.593  -5.100   0.150  1.00 98.47      B    C
ATOM   4313  NH1 ARG   659      -74.716  -3.824  -0.199  1.00 98.24      B    N
ATOM   4314  NH2 ARG   659      -75.624  -5.739   0.691  1.00 99.04      B    N
ATOM   4315  C   ARG   659      -70.303  -8.910   0.551  1.00 84.43      B    C
ATOM   4316  O   ARG   659      -69.628  -8.534   1.510  1.00 85.18      B    O
ATOM   4317  N   ALA   671      -74.537 -18.439  -0.545  1.00 74.52      B    N
ATOM   4318  CA  ALA   671      -73.500 -19.241  -1.186  1.00 75.31      B    C
ATOM   4319  CB  ALA   671      -73.891 -20.711  -1.157  1.00 75.29      B    C
ATOM   4320  C   ALA   671      -73.234 -18.801  -2.628  1.00 75.10      B    C
ATOM   4321  O   ALA   671      -74.154 -18.416  -3.357  1.00 76.03      B    O
ATOM   4322  N   VAL   672      -71.970 -18.867  -3.035  1.00 73.34      B    N
ATOM   4323  CA  VAL   672      -71.565 -18.405  -4.359  1.00 70.84      B    C
ATOM   4324  CB  VAL   672      -70.845 -17.046  -4.262  1.00 71.44      B    C
ATOM   4325  CG1 VAL   672      -70.430 -16.578  -5.642  1.00 71.15      B    C
ATOM   4326  CG2 VAL   672      -71.759 -16.022  -3.605  1.00 70.69      B    C
ATOM   4327  C   VAL   672      -70.637 -19.410  -5.044  1.00 68.72      B    C
ATOM   4328  O   VAL   672      -69.829 -20.069  -4.385  1.00 68.45      B    O
ATOM   4329  N   THR   673      -70.760 -19.527  -6.364  1.00 65.32      B    N
ATOM   4330  CA  THR   673      -69.944 -20.476  -7.114  1.00 62.53      B    C
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ATOM   4331  CB  THR   673     -70.817 -21.586  -7.725  1.00 63.63      B    C
ATOM   4332  OG1 THR   673     -71.524 -22.264  -6.679  1.00 64.64      B    O
ATOM   4333  CG2 THR   673     -69.952 -22.590  -8.473  1.00 62.74      B    C
ATOM   4334  C   THR   673     -69.132 -19.821  -8.231  1.00 59.50      B    C
ATOM   4335  O   THR   673     -69.671 -19.105  -9.077  1.00 58.78      B    O
ATOM   4336  N   ALA   674     -67.830 -20.080  -8.221  1.00 56.52      B    N
ATOM   4337  CA  ALA   674     -66.935 -19.595  -9.264  1.00 54.91      B    C
ATOM   4338  CB  ALA   674     -65.567 -19.269  -8.666  1.00 53.50      B    C
ATOM   4339  C   ALA   674     -66.790 -20.634 -10.379  1.00 53.38      B    C
ATOM   4340  O   ALA   674     -66.506 -21.809 -10.121  1.00 52.02      B    O
ATOM   4341  N   VAL   675     -66.974 -20.186 -11.616  1.00 50.64      B    N
ATOM   4342  CA  VAL   675     -66.941 -21.069 -12.776  1.00 49.05      B    C
ATOM   4343  CB  VAL   675     -68.256 -20.961 -13.583  1.00 48.94      B    C
ATOM   4344  CG1 VAL   675     -68.282 -22.007 -14.685  1.00 46.82      B    C
ATOM   4345  CG2 VAL   675     -69.448 -21.114 -12.653  1.00 47.81      B    C
ATOM   4346  C   VAL   675     -65.778 -20.709 -13.701  1.00 47.76      B    C
ATOM   4347  O   VAL   675     -65.758 -19.629 -14.291  1.00 47.79      B    O
ATOM   4348  N   ALA   676     -64.817 -21.619 -13.828  1.00 46.75      B    N
ATOM   4349  CA  ALA   676     -63.688 -21.429 -14.736  1.00 45.42      B    C
ATOM   4350  CB  ALA   676     -62.389 -21.792 -14.037  1.00 43.18      B    C
ATOM   4351  C   ALA   676     -63.840 -22.275 -15.996  1.00 46.23      B    C
ATOM   4352  O   ALA   676     -64.236 -23.440 -15.932  1.00 47.47      B    O
ATOM   4353  N   ILE   677     -63.518 -21.687 -17.142  1.00 45.06      B    N
ATOM   4354  CA  ILE   677     -63.322 -22.462 -18.358  1.00 42.84      B    C
ATOM   4355  CB  ILE   677     -64.044 -21.805 -19.559  1.00 41.75      B    C
ATOM   4356  CG2 ILE   677     -63.786 -22.593 -20.834  1.00 38.64      B    C
ATOM   4357  CG1 ILE   677     -65.548 -21.746 -19.287  1.00 41.12      B    C
ATOM   4358  CD1 ILE   677     -66.349 -21.146 -20.430  1.00 42.32      B    C
ATOM   4359  C   ILE   677     -61.824 -22.534 -18.622  1.00 42.86      B    C
ATOM   4360  O   ILE   677     -61.159 -21.508 -18.733  1.00 43.27      B    O
ATOM   4361  N   CYS   678     -61.303 -23.754 -18.701  1.00 44.51      B    N
ATOM   4362  CA  CYS   678     -59.877 -24.001 -18.896  1.00 46.16      B    C
ATOM   4363  C   CYS   678     -59.656 -24.664 -20.248  1.00 46.08      B    C
ATOM   4364  O   CYS   678     -60.392 -25.574 -20.615  1.00 46.07      B    O
ATOM   4365  CB  CYS   678     -59.346 -24.930 -17.800  1.00 48.42      B    C
ATOM   4366  SG  CYS   678     -59.691 -24.383 -16.096  1.00 55.45      B    S
ATOM   4367  N   CYS   679     -58.641 -24.221 -20.981  1.00 46.30      B    N
ATOM   4368  CA  CYS   679     -58.335 -24.816 -22.273  1.00 48.35      B    C
ATOM   4369  C   CYS   679     -56.863 -25.140 -22.441  1.00 50.02      B    C
ATOM   4370  O   CYS   679     -56.021 -24.723 -21.646  1.00 49.93      B    O
ATOM   4371  CB  CYS   679     -58.737 -23.893 -23.415  1.00 47.56      B    C
ATOM   4372  SG  CYS   679     -60.399 -23.172 -23.340  1.00 48.97      B    S
ATOM   4373  N   ARG   680     -56.574 -25.885 -23.501  1.00 52.89      B    N
ATOM   4374  CA  ARG   680     -55.214 -26.136 -23.944  1.00 57.75      B    C
ATOM   4375  CB  ARG   680     -54.573 -27.236 -23.100  1.00 58.25      B    C
ATOM   4376  CG  ARG   680     -55.435 -28.470 -22.933  1.00 61.86      B    C
ATOM   4377  CD  ARG   680     -54.577 -29.715 -22.813  1.00 64.43      B    C
ATOM   4378  NE  ARG   680     -55.302 -30.826 -22.205  1.00 68.80      B    N
ATOM   4379  CZ  ARG   680     -54.806 -32.052 -22.065  1.00 70.93      B    C
ATOM   4380  NH1 ARG   680     -55.532 -33.006 -21.497  1.00 71.39      B    N
ATOM   4381  NH2 ARG   680     -53.583 -32.327 -22.501  1.00 73.17      B    N
ATOM   4382  C   ARG   680     -55.261 -26.561 -25.404  1.00 60.89      B    C
ATOM   4383  O   ARG   680     -56.339 -26.741 -25.970  1.00 59.44      B    O
ATOM   4384  N   SER   681     -54.093 -26.704 -26.017  1.00 66.22      B    N
ATOM   4385  CA  SER   681     -54.003 -27.243 -27.367  1.00 72.84      B    C
ATOM   4386  CB  SER   681     -53.377 -26.219 -28.317  1.00 73.28      B    C
ATOM   4387  OG  SER   681     -54.177 -25.056 -28.429  1.00 75.41      B    O
ATOM   4388  C   SER   681     -53.148 -28.501 -27.357  1.00 76.77      B    C
ATOM   4389  O   SER   681     -51.946 -28.435 -27.096  1.00 77.43      B    O
ATOM   4390  N   ARG   682     -53.764 -29.645 -27.644  1.00 81.26      B    N
ATOM   4391  CA  ARG   682     -53.031 -30.905 -27.694  1.00 85.36      B    C
ATOM   4392  CB  ARG   682     -54.005 -32.087 -27.634  1.00 88.32      B    C
ATOM   4393  CG  ARG   682     -54.534 -32.394 -26.237  1.00 92.63      B    C
ATOM   4394  CD  ARG   682     -55.114 -33.804 -26.167  1.00 96.94      B    C
ATOM   4395  NE  ARG   682     -54.231 -34.789 -26.789  1.00100.46      B    N
ATOM   4396  CZ  ARG   682     -54.568 -36.052 -27.043  1.00102.13      B    C
ATOM   4397  NH1 ARG   682     -53.691 -36.869 -27.616  1.00102.64      B    N
ATOM   4398  NH2 ARG   682     -55.777 -36.500 -26.724  1.00102.84      B    N
ATOM   4399  C   ARG   682     -52.165 -31.008 -28.953  1.00 86.02      B    C
ATOM   4400  O   ARG   682     -52.319 -30.158 -29.860  1.00 85.88      B    O
ATOM   4401  OXT ARG   682     -51.337 -31.944 -29.016  1.00 86.99      B    O
TER    4402       ARG   682                                             B
ATOM   4403  CB  GLU     1     -36.231  38.731   6.129  1.00 73.54      L    C
ATOM   4404  CG  GLU     1     -35.182  39.159   5.116  1.00 77.83      L    C
ATOM   4405  CD  GLU     1     -33.792  39.221   5.713  1.00 80.96      L    C
ATOM   4406  OE1 GLU     1     -33.555  38.539   6.736  1.00 81.61      L    O
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ATOM	4407	OE2	GLU	1	-32.939	39.951	5.159	1.00	81.99	L	O
ATOM	4408	C	GLU	1	-37.606	37.144	4.782	1.00	68.14	L	C
ATOM	4409	O	GLU	1	-36.842	36.943	3.836	1.00	68.19	L	O
ATOM	4410	N	GLU	1	-38.659	38.513	6.573	1.00	71.43	L	N
ATOM	4411	CA	GLU	1	-37.610	38.484	5.515	1.00	70.62	L	C
ATOM	4412	N	SER	2	-38.458	36.226	5.230	1.00	64.55	L	N
ATOM	4413	CA	SER	2	-38.645	34.953	4.540	1.00	60.66	L	C
ATOM	4414	CB	SER	2	-39.040	33.862	5.545	1.00	60.92	L	C
ATOM	4415	OG	SER	2	-40.188	34.231	6.293	1.00	59.32	L	O
ATOM	4416	C	SER	2	-39.713	35.079	3.450	1.00	57.06	L	C
ATOM	4417	O	SER	2	-40.693	35.810	3.607	1.00	55.88	L	O
ATOM	4418	N	VAL	3	-39.514	34.364	2.346	1.00	52.88	L	N
ATOM	4419	CA	VAL	3	-40.437	34.408	1.212	1.00	48.89	L	C
ATOM	4420	CB	VAL	3	-39.850	33.648	-0.013	1.00	49.36	L	C
ATOM	4421	CG1	VAL	3	-40.818	33.703	-1.170	1.00	49.74	L	C
ATOM	4422	CG2	VAL	3	-38.519	34.253	-0.420	1.00	48.94	L	C
ATOM	4423	C	VAL	3	-41.798	33.801	1.562	1.00	45.34	L	C
ATOM	4424	O	VAL	3	-42.831	34.242	1.057	1.00	46.42	L	O
ATOM	4425	N	LEU	4	-41.796	32.783	2.419	1.00	40.20	L	N
ATOM	4426	CA	LEU	4	-43.039	32.204	2.919	1.00	36.80	L	C
ATOM	4427	CB	LEU	4	-42.943	30.675	2.932	1.00	35.15	L	C
ATOM	4428	CG	LEU	4	-42.440	30.024	1.639	1.00	35.77	L	C
ATOM	4429	CD1	LEU	4	-42.428	28.516	1.799	1.00	33.82	L	C
ATOM	4430	CD2	LEU	4	-43.339	30.424	0.473	1.00	33.23	L	C
ATOM	4431	C	LEU	4	-43.312	32.723	4.330	1.00	34.95	L	C
ATOM	4432	O	LEU	4	-42.414	32.790	5.167	1.00	34.99	L	O
ATOM	4433	N	THR	5	-44.554	33.096	4.599	1.00	34.09	L	N
ATOM	4434	CA	THR	5	-44.870	33.701	5.880	1.00	33.56	L	C
ATOM	4435	CB	THR	5	-45.693	34.989	5.686	1.00	34.89	L	C
ATOM	4436	OG1	THR	5	-44.944	35.912	4.883	1.00	36.65	L	O
ATOM	4437	CG2	THR	5	-45.998	35.636	7.037	1.00	33.40	L	C
ATOM	4438	C	THR	5	-45.637	32.749	6.790	1.00	33.35	L	C
ATOM	4439	O	THR	5	-46.701	32.250	6.421	1.00	33.73	L	O
ATOM	4440	N	GLN	6	-45.088	32.508	7.979	1.00	32.27	L	N
ATOM	4441	CA	GLN	6	-45.755	31.719	9.014	1.00	32.42	L	C
ATOM	4442	CB	GLN	6	-44.886	30.527	9.440	1.00	31.42	L	C
ATOM	4443	CG	GLN	6	-44.498	29.545	8.353	1.00	29.42	L	C
ATOM	4444	CD	GLN	6	-43.610	28.432	8.891	1.00	29.99	L	C
ATOM	4445	OE1	GLN	6	-42.530	28.172	8.362	1.00	30.78	L	O
ATOM	4446	NE2	GLN	6	-44.061	27.775	9.956	1.00	26.07	L	N
ATOM	4447	C	GLN	6	-45.959	32.603	10.241	1.00	33.06	L	C
ATOM	4448	O	GLN	6	-45.203	33.544	10.462	1.00	33.60	L	O
ATOM	4449	N	PRO	7	-46.963	32.293	11.074	1.00	32.74	L	N
ATOM	4450	CD	PRO	7	-48.008	31.265	10.922	1.00	32.27	L	C
ATOM	4451	CA	PRO	7	-47.036	32.979	12.369	1.00	33.15	L	C
ATOM	4452	CB	PRO	7	-48.356	32.492	12.963	1.00	32.34	L	C
ATOM	4453	CG	PRO	7	-48.581	31.147	12.312	1.00	33.16	L	C
ATOM	4454	C	PRO	7	-45.835	32.600	13.241	1.00	33.91	L	C
ATOM	4455	O	PRO	7	-45.385	31.457	13.237	1.00	34.46	L	O
ATOM	4456	N	PRO	8	-45.295	33.564	13.994	1.00	35.09	L	N
ATOM	4457	CD	PRO	8	-45.722	34.973	14.069	1.00	34.70	L	C
ATOM	4458	CA	PRO	8	-44.112	33.292	14.825	1.00	34.71	L	C
ATOM	4459	CB	PRO	8	-43.814	34.640	15.494	1.00	36.39	L	C
ATOM	4460	CG	PRO	8	-44.501	35.661	14.617	1.00	38.46	L	C
ATOM	4461	C	PRO	8	-44.371	32.196	15.858	1.00	33.60	L	C
ATOM	4462	O	PRO	8	-43.480	31.418	16.188	1.00	33.87	L	O
ATOM	4463	N	SER	9	-45.591	32.131	16.373	1.00	32.78	L	N
ATOM	4464	CA	SER	9	-45.886	31.153	17.409	1.00	34.68	L	C
ATOM	4465	CB	SER	9	-45.491	31.713	18.776	1.00	36.06	L	C
ATOM	4466	OG	SER	9	-46.262	32.861	19.072	1.00	40.67	L	O
ATOM	4467	C	SER	9	-47.349	30.732	17.439	1.00	32.15	L	C
ATOM	4468	O	SER	9	-48.225	31.463	16.988	1.00	33.49	L	O
ATOM	4469	N	VAL	10	-47.598	29.540	17.967	1.00	29.27	L	N
ATOM	4470	CA	VAL	10	-48.952	29.078	18.237	1.00	28.93	L	C
ATOM	4471	CB	VAL	10	-49.503	28.157	17.101	1.00	29.21	L	C
ATOM	4472	CG1	VAL	10	-49.559	28.921	15.784	1.00	32.11	L	C
ATOM	4473	CG2	VAL	10	-48.621	26.931	16.952	1.00	29.17	L	C
ATOM	4474	C	VAL	10	-48.888	28.271	19.513	1.00	28.40	L	C
ATOM	4475	O	VAL	10	-47.809	27.826	19.925	1.00	29.78	L	O
ATOM	4476	N	SER	11	-50.035	28.067	20.143	1.00	27.75	L	N
ATOM	4477	CA	SER	11	-50.049	27.307	21.377	1.00	30.00	L	C
ATOM	4478	CB	SER	11	-49.688	28.225	22.550	1.00	30.66	L	C
ATOM	4479	OG	SER	11	-50.592	29.314	22.626	1.00	33.85	L	O
ATOM	4480	C	SER	11	-51.397	26.652	21.621	1.00	27.91	L	C
ATOM	4481	O	SER	11	-52.418	27.101	21.105	1.00	27.57	L	O
ATOM	4482	N	GLY	12	-51.381	25.578	22.404	1.00	27.56	L	N

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ATOM   4483  CA  GLY  12     -52.605  24.933  22.836  1.00 25.58      L    C
ATOM   4484  C   GLY  12     -52.322  24.003  24.004  1.00 27.45      L    C
ATOM   4485  O   GLY  12     -51.165  23.668  24.271  1.00 26.91      L    O
ATOM   4486  N   ALA  13     -53.375  23.593  24.706  1.00 26.20      L    N
ATOM   4487  CA  ALA  13     -53.259  22.596  25.762  1.00 27.69      L    C
ATOM   4488  CB  ALA  13     -54.372  22.804  26.804  1.00 27.24      L    C
ATOM   4489  C   ALA  13     -53.345  21.186  25.171  1.00 26.76      L    C
ATOM   4490  O   ALA  13     -53.861  20.992  24.071  1.00 27.16      L    O
ATOM   4491  N   PRO  14     -52.839  20.182  25.900  1.00 26.62      L    N
ATOM   4492  CD  PRO  14     -52.181  20.255  27.219  1.00 26.21      L    C
ATOM   4493  CA  PRO  14     -52.938  18.803  25.408  1.00 25.50      L    C
ATOM   4494  CB  PRO  14     -52.425  17.965  26.587  1.00 27.17      L    C
ATOM   4495  CG  PRO  14     -51.520  18.906  27.343  1.00 24.03      L    C
ATOM   4496  C   PRO  14     -54.382  18.467  25.046  1.00 26.24      L    C
ATOM   4497  O   PRO  14     -55.303  18.750  25.816  1.00 25.55      L    O
ATOM   4498  N   GLY  15     -54.575  17.885  23.864  1.00 25.21      L    N
ATOM   4499  CA  GLY  15     -55.907  17.508  23.432  1.00 25.03      L    C
ATOM   4500  C   GLY  15     -56.561  18.482  22.464  1.00 26.75      L    C
ATOM   4501  O   GLY  15     -57.538  18.137  21.794  1.00 28.23      L    O
ATOM   4502  N   GLN  16     -56.037  19.700  22.382  1.00 25.53      L    N
ATOM   4503  CA  GLN  16     -56.621  20.711  21.502  1.00 25.79      L    C
ATOM   4504  CB  GLN  16     -56.273  22.122  22.001  1.00 25.68      L    C
ATOM   4505  CG  GLN  16     -56.942  22.481  23.327  1.00 27.59      L    C
ATOM   4506  CD  GLN  16     -56.922  23.976  23.606  1.00 30.78      L    C
ATOM   4507  OE1 GLN  16     -55.867  24.552  23.878  1.00 30.95      L    O
ATOM   4508  NE2 GLN  16     -58.094  24.613  23.540  1.00 27.50      L    N
ATOM   4509  C   GLN  16     -56.155  20.548  20.064  1.00 25.94      L    C
ATOM   4510  O   GLN  16     -55.245  19.772  19.772  1.00 24.98      L    O
ATOM   4511  N   ARG  17     -56.798  21.276  19.161  1.00 28.78      L    N
ATOM   4512  CA  ARG  17     -56.407  21.283  17.756  1.00 29.05      L    C
ATOM   4513  CB  ARG  17     -57.628  20.990  16.876  1.00 31.19      L    C
ATOM   4514  CG  ARG  17     -57.375  21.121  15.385  1.00 32.39      L    C
ATOM   4515  CD  ARG  17     -58.629  20.818  14.577  1.00 32.91      L    C
ATOM   4516  NE  ARG  17     -58.384  20.933  13.140  1.00 36.49      L    N
ATOM   4517  CZ  ARG  17     -58.443  22.079  12.466  1.00 39.15      L    C
ATOM   4518  NH1 ARG  17     -58.737  23.210  13.098  1.00 40.18      L    N
ATOM   4519  NH2 ARG  17     -58.214  22.101  11.160  1.00 39.44      L    N
ATOM   4520  C   ARG  17     -55.853  22.661  17.430  1.00 29.77      L    C
ATOM   4521  O   ARG  17     -56.492  23.674  17.716  1.00 30.44      L    O
ATOM   4522  N   VAL  18     -54.664  22.708  16.845  1.00 28.97      L    N
ATOM   4523  CA  VAL  18     -54.095  23.984  16.439  1.00 30.60      L    C
ATOM   4524  CB  VAL  18     -52.847  24.339  17.288  1.00 32.96      L    C
ATOM   4525  CG1 VAL  18     -53.234  24.451  18.756  1.00 32.49      L    C
ATOM   4526  CG2 VAL  18     -51.771  23.274  17.108  1.00 34.57      L    C
ATOM   4527  C   VAL  18     -53.706  23.936  14.967  1.00 30.86      L    C
ATOM   4528  O   VAL  18     -53.442  22.869  14.418  1.00 31.31      L    O
ATOM   4529  N   THR  19     -53.680  25.093  14.322  1.00 30.68      L    N
ATOM   4530  CA  THR  19     -53.253  25.153  12.937  1.00 30.55      L    C
ATOM   4531  CB  THR  19     -54.419  25.572  12.003  1.00 33.05      L    C
ATOM   4532  OG1 THR  19     -54.922  26.852  12.412  1.00 37.51      L    O
ATOM   4533  CG2 THR  19     -55.549  24.544  12.057  1.00 30.05      L    C
ATOM   4534  C   THR  19     -52.109  26.140  12.772  1.00 30.41      L    C
ATOM   4535  O   THR  19     -51.977  27.094  13.535  1.00 31.46      L    O
ATOM   4536  N   ILE  20     -51.282  25.893  11.766  1.00 30.36      L    N
ATOM   4537  CA  ILE  20     -50.173  26.767  11.430  1.00 28.18      L    C
ATOM   4538  CB  ILE  20     -48.827  26.063  11.700  1.00 28.17      L    C
ATOM   4539  CG2 ILE  20     -47.670  26.922  11.207  1.00 24.45      L    C
ATOM   4540  CG1 ILE  20     -48.706  25.749  13.199  1.00 30.05      L    C
ATOM   4541  CD1 ILE  20     -47.458  24.962  13.567  1.00 27.37      L    C
ATOM   4542  C   ILE  20     -50.295  27.064   9.944  1.00 31.43      L    C
ATOM   4543  O   ILE  20     -50.325  26.142   9.119  1.00 31.91      L    O
ATOM   4544  N   SER  21     -50.375  28.344   9.600  1.00 31.04      L    N
ATOM   4545  CA  SER  21     -50.526  28.735   8.206  1.00 31.99      L    C
ATOM   4546  CB  SER  21     -51.455  29.946   8.086  1.00 32.54      L    C
ATOM   4547  OG  SER  21     -50.847  31.097   8.642  1.00 34.59      L    O
ATOM   4548  C   SER  21     -49.187  29.064   7.567  1.00 32.68      L    C
ATOM   4549  O   SER  21     -48.208  29.381   8.248  1.00 32.74      L    O
ATOM   4550  N   CYS  22     -49.159  28.979   6.245  1.00 33.22      L    N
ATOM   4551  CA  CYS  22     -47.970  29.269   5.456  1.00 35.47      L    C
ATOM   4552  C   CYS  22     -48.488  29.994   4.222  1.00 36.21      L    C
ATOM   4553  O   CYS  22     -49.257  29.423   3.447  1.00 37.64      L    O
ATOM   4554  CB  CYS  22     -47.295  27.956   5.053  1.00 33.85      L    C
ATOM   4555  SG  CYS  22     -45.777  28.046   4.037  1.00 40.59      L    S
ATOM   4556  N   THR  23     -48.099  31.248   4.033  1.00 35.52      L    N
ATOM   4557  CA  THR  23     -48.563  31.957   2.849  1.00 36.01      L    C
ATOM   4558  CB  THR  23     -49.375  33.215   3.218  1.00 36.77      L    C
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ATOM	4559	OG1	THR	23	-48.493	34.218	3.725	1.00	43.95	L	O
ATOM	4560	CG2	THR	23	-50.408	32.868	4.285	1.00	35.35	L	C
ATOM	4561	C	THR	23	-47.399	32.358	1.953	1.00	35.25	L	C
ATOM	4562	O	THR	23	-46.386	32.893	2.418	1.00	33.54	L	O
ATOM	4563	N	GLY	24	-47.549	32.077	0.663	1.00	33.91	L	N
ATOM	4564	CA	GLY	24	-46.526	32.434	-0.295	1.00	34.67	L	C
ATOM	4565	C	GLY	24	-47.058	33.355	-1.373	1.00	35.72	L	C
ATOM	4566	O	GLY	24	-47.847	34.259	-1.100	1.00	35.18	L	O
ATOM	4567	N	SER	25	-46.622	33.120	-2.604	1.00	36.22	L	N
ATOM	4568	CA	SER	25	-46.957	33.996	-3.717	1.00	38.10	L	C
ATOM	4569	CB	SER	25	-45.879	35.067	-3.880	1.00	36.60	L	C
ATOM	4570	OG	SER	25	-44.723	34.513	-4.495	1.00	38.31	L	O
ATOM	4571	C	SER	25	-47.065	33.195	-5.012	1.00	38.80	L	C
ATOM	4572	O	SER	25	-46.973	31.967	-5.011	1.00	37.63	L	O
ATOM	4573	N	SER	26	-47.240	33.909	-6.118	1.00	39.46	L	N
ATOM	4574	CA	SER	26	-47.413	33.284	-7.425	1.00	39.78	L	C
ATOM	4575	CB	SER	26	-47.955	34.309	-8.423	1.00	40.90	L	C
ATOM	4576	OG	SER	26	-47.059	35.407	-8.537	1.00	42.25	L	O
ATOM	4577	C	SER	26	-46.113	32.699	-7.966	1.00	38.37	L	C
ATOM	4578	O	SER	26	-46.125	31.988	-8.965	1.00	40.32	L	O
ATOM	4579	N	SER	27	-44.988	33.000	-7.325	1.00	37.56	L	N
ATOM	4580	CA	SER	27	-43.725	32.405	-7.756	1.00	36.90	L	C
ATOM	4581	CB	SER	27	-42.586	33.423	-7.636	1.00	37.67	L	C
ATOM	4582	OG	SER	27	-42.376	33.797	-6.290	1.00	46.23	L	O
ATOM	4583	C	SER	27	-43.359	31.119	-7.003	1.00	34.65	L	C
ATOM	4584	O	SER	27	-42.387	30.450	-7.356	1.00	34.64	L	O
ATOM	4585	N	ASN	28	-44.126	30.766	-5.972	1.00	32.35	L	N
ATOM	4586	CA	ASN	28	-43.899	29.493	-5.288	1.00	31.77	L	C
ATOM	4587	CB	ASN	28	-43.209	29.711	-3.930	1.00	29.14	L	C
ATOM	4588	CG	ASN	28	-43.767	30.891	-3.162	1.00	29.35	L	C
ATOM	4589	OD1	ASN	28	-44.919	30.878	-2.717	1.00	30.17	L	O
ATOM	4590	ND2	ASN	28	-42.943	31.918	-2.986	1.00	27.56	L	N
ATOM	4591	C	ASN	28	-45.150	28.649	-5.101	1.00	32.18	L	C
ATOM	4592	O	ASN	28	-45.499	27.846	-5.971	1.00	33.39	L	O
ATOM	4593	N	ILE	29	-45.830	28.818	-3.974	1.00	31.68	L	N
ATOM	4594	CA	ILE	29	-47.009	28.012	-3.703	1.00	34.34	L	C
ATOM	4595	CB	ILE	29	-47.614	28.348	-2.323	1.00	34.09	L	C
ATOM	4596	CG2	ILE	29	-48.921	27.592	-2.126	1.00	33.22	L	C
ATOM	4597	CG1	ILE	29	-46.613	27.993	-1.221	1.00	32.59	L	C
ATOM	4598	CD1	ILE	29	-47.137	28.233	0.186	1.00	33.16	L	C
ATOM	4599	C	ILE	29	-48.063	28.232	-4.786	1.00	36.40	L	C
ATOM	4600	O	ILE	29	-48.703	27.281	-5.241	1.00	37.01	L	O
ATOM	4601	N	GLY	30	-48.230	29.485	-5.202	1.00	36.65	L	N
ATOM	4602	CA	GLY	30	-49.224	29.801	-6.212	1.00	37.76	L	C
ATOM	4603	C	GLY	30	-48.821	29.425	-7.630	1.00	39.98	L	C
ATOM	4604	O	GLY	30	-49.600	29.610	-8.562	1.00	41.54	L	O
ATOM	4605	N	ALA	31	-47.612	28.896	-7.800	1.00	39.82	L	N
ATOM	4606	CA	ALA	31	-47.131	28.495	-9.119	1.00	38.00	L	C
ATOM	4607	CB	ALA	31	-45.687	28.942	-9.313	1.00	34.50	L	C
ATOM	4608	C	ALA	31	-47.233	26.991	-9.305	1.00	39.19	L	C
ATOM	4609	O	ALA	31	-46.697	26.441	-10.268	1.00	40.14	L	O
ATOM	4610	N	GLY	32	-47.907	26.319	-8.377	1.00	39.51	L	N
ATOM	4611	CA	GLY	32	-48.112	24.889	-8.526	1.00	39.02	L	C
ATOM	4612	C	GLY	32	-47.097	23.998	-7.825	1.00	39.80	L	C
ATOM	4613	O	GLY	32	-47.106	22.779	-8.004	1.00	42.17	L	O
ATOM	4614	N	TYR	33	-46.213	24.586	-7.029	1.00	36.85	L	N
ATOM	4615	CA	TYR	33	-45.240	23.785	-6.297	1.00	35.51	L	C
ATOM	4616	CB	TYR	33	-43.951	24.592	-6.092	1.00	34.69	L	C
ATOM	4617	CG	TYR	33	-43.300	24.951	-7.410	1.00	34.74	L	C
ATOM	4618	CD1	TYR	33	-42.695	23.974	-8.198	1.00	33.78	L	C
ATOM	4619	CE1	TYR	33	-42.170	24.279	-9.443	1.00	34.40	L	C
ATOM	4620	CD2	TYR	33	-43.356	26.250	-7.903	1.00	35.64	L	C
ATOM	4621	CE2	TYR	33	-42.834	26.566	-9.149	1.00	34.16	L	C
ATOM	4622	CZ	TYR	33	-42.247	25.579	-9.913	1.00	35.08	L	C
ATOM	4623	OH	TYR	33	-41.747	25.893	-11.157	1.00	38.62	L	O
ATOM	4624	C	TYR	33	-45.835	23.345	-4.962	1.00	32.94	L	C
ATOM	4625	O	TYR	33	-46.581	24.095	-4.340	1.00	33.35	L	O
ATOM	4626	N	ASP	34	-45.526	22.120	-4.544	1.00	29.96	L	N
ATOM	4627	CA	ASP	34	-46.046	21.580	-3.288	1.00	28.49	L	C
ATOM	4628	CB	ASP	34	-45.809	20.071	-3.221	1.00	30.81	L	C
ATOM	4629	CG	ASP	34	-46.733	19.280	-4.136	1.00	34.47	L	C
ATOM	4630	OD1	ASP	34	-46.567	18.041	-4.189	1.00	36.39	L	O
ATOM	4631	OD2	ASP	34	-47.618	19.879	-4.793	1.00	32.29	L	O
ATOM	4632	C	ASP	34	-45.377	22.229	-2.073	1.00	27.51	L	C
ATOM	4633	O	ASP	34	-44.238	22.701	-2.151	1.00	25.64	L	O
ATOM	4634	N	VAL	35	-46.090	22.241	-0.954	1.00	26.98	L	N

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ATOM   4635  CA  VAL   35     -45.523  22.652   0.326  1.00 28.45      L    C
ATOM   4636  CB  VAL   35     -46.543  23.485   1.139  1.00 27.84      L    C
ATOM   4637  CG1 VAL   35     -45.970  23.840   2.497  1.00 25.77      L    C
ATOM   4638  CG2 VAL   35     -46.897  24.756   0.375  1.00 29.67      L    C
ATOM   4639  C   VAL   35     -45.143  21.413   1.136  1.00 28.31      L    C
ATOM   4640  O   VAL   35     -45.925  20.471   1.237  1.00 28.31      L    O
ATOM   4641  N   HIS   36     -43.943  21.408   1.706  1.00 27.34      L    N
ATOM   4642  CA  HIS   36     -43.556  20.336   2.623  1.00 27.59      L    C
ATOM   4643  CB  HIS   36     -42.284  19.633   2.131  1.00 26.50      L    C
ATOM   4644  CG  HIS   36     -42.256  19.391   0.653  1.00 27.79      L    C
ATOM   4645  CD2 HIS   36     -41.509  19.955  -0.327  1.00 25.62      L    C
ATOM   4646  ND1 HIS   36     -43.056  18.454   0.032  1.00 27.57      L    N
ATOM   4647  CE1 HIS   36     -42.801  18.450  -1.264  1.00 27.14      L    C
ATOM   4648  NE2 HIS   36     -41.866  19.351  -1.508  1.00 27.36      L    N
ATOM   4649  C   HIS   36     -43.309  20.926   4.009  1.00 28.07      L    C
ATOM   4650  O   HIS   36     -42.879  22.082   4.127  1.00 28.94      L    O
ATOM   4651  N   TRP   37     -43.575  20.140   5.051  1.00 25.76      L    N
ATOM   4652  CA  TRP   37     -43.456  20.614   6.426  1.00 24.75      L    C
ATOM   4653  CB  TRP   37     -44.607  20.545   7.145  1.00 22.85      L    C
ATOM   4654  CG  TRP   37     -45.845  21.469   6.588  1.00 25.11      L    C
ATOM   4655  CD2 TRP   37     -46.188  22.769   7.082  1.00 23.55      L    C
ATOM   4656  CE2 TRP   37     -47.265  23.244   6.301  1.00 24.06      L    C
ATOM   4657  CE3 TRP   37     -45.693  23.576   8.111  1.00 25.38      L    C
ATOM   4658  CD1 TRP   37     -46.704  21.215   5.544  1.00 22.14      L    C
ATOM   4659  NE1 TRP   37     -47.559  22.278   5.370  1.00 24.91      L    N
ATOM   4660  CZ2 TRP   37     -47.856  24.488   6.520  1.00 23.30      L    C
ATOM   4661  CZ3 TRP   37     -46.280  24.813   8.330  1.00 24.69      L    C
ATOM   4662  CH2 TRP   37     -47.352  25.257   7.537  1.00 26.88      L    C
ATOM   4663  C   TRP   37     -42.437  19.813   7.220  1.00 26.46      L    C
ATOM   4664  O   TRP   37     -42.348  18.587   7.099  1.00 25.56      L    O
ATOM   4665  N   TYR   38     -41.678  20.528   8.042  1.00 25.83      L    N
ATOM   4666  CA  TYR   38     -40.661  19.928   8.883  1.00 25.04      L    C
ATOM   4667  CB  TYR   38     -39.286  20.427   8.462  1.00 24.09      L    C
ATOM   4668  CG  TYR   38     -39.001  20.144   7.010  1.00 26.55      L    C
ATOM   4669  CD1 TYR   38     -39.377  21.047   6.026  1.00 25.25      L    C
ATOM   4670  CE1 TYR   38     -39.164  20.775   4.686  1.00 26.99      L    C
ATOM   4671  CD2 TYR   38     -38.394  18.953   6.617  1.00 25.37      L    C
ATOM   4672  CE2 TYR   38     -38.174  18.671   5.278  1.00 25.45      L    C
ATOM   4673  CZ  TYR   38     -38.566  19.591   4.317  1.00 26.83      L    C
ATOM   4674  OH  TYR   38     -38.362  19.332   2.981  1.00 27.05      L    O
ATOM   4675  C   TYR   38     -40.911  20.270  10.342  1.00 27.08      L    C
ATOM   4676  O   TYR   38     -41.383  21.365  10.673  1.00 25.71      L    O
ATOM   4677  N   GLN   39     -40.596  19.314  11.204  1.00 27.12      L    N
ATOM   4678  CA  GLN   39     -40.677  19.495  12.644  1.00 26.88      L    C
ATOM   4679  CB  GLN   39     -41.471  18.349  13.262  1.00 26.61      L    C
ATOM   4680  CG  GLN   39     -41.665  18.445  14.767  1.00 27.00      L    C
ATOM   4681  CD  GLN   39     -42.299  17.184  15.326  1.00 28.70      L    C
ATOM   4682  OE1 GLN   39     -41.752  16.087  15.179  1.00 28.98      L    O
ATOM   4683  NE2 GLN   39     -43.461  17.330  15.965  1.00 25.31      L    N
ATOM   4684  C   GLN   39     -39.262  19.475  13.190  1.00 27.65      L    C
ATOM   4685  O   GLN   39     -38.481  18.585  12.855  1.00 27.35      L    O
ATOM   4686  N   GLN   40     -38.921  20.451  14.022  1.00 26.72      L    N
ATOM   4687  CA  GLN   40     -37.609  20.440  14.660  1.00 29.84      L    C
ATOM   4688  CB  GLN   40     -36.766  21.624  14.172  1.00 28.79      L    C
ATOM   4689  CG  GLN   40     -35.331  21.609  14.685  1.00 30.78      L    C
ATOM   4690  CD  GLN   40     -34.505  22.776  14.157  1.00 31.87      L    C
ATOM   4691  OE1 GLN   40     -35.014  23.887  13.984  1.00 30.96      L    O
ATOM   4692  NE2 GLN   40     -33.226  22.527  13.900  1.00 26.76      L    N
ATOM   4693  C   GLN   40     -37.769  20.502  16.175  1.00 31.91      L    C
ATOM   4694  O   GLN   40     -38.186  21.519  16.729  1.00 30.67      L    O
ATOM   4695  N   LEU   41     -37.450  19.398  16.839  1.00 36.61      L    N
ATOM   4696  CA  LEU   41     -37.465  19.353  18.294  1.00 40.30      L    C
ATOM   4697  CB  LEU   41     -37.397  17.906  18.779  1.00 40.69      L    C
ATOM   4698  CG  LEU   41     -38.540  17.014  18.289  1.00 45.02      L    C
ATOM   4699  CD1 LEU   41     -38.322  15.587  18.778  1.00 47.42      L    C
ATOM   4700  CD2 LEU   41     -39.866  17.558  18.790  1.00 45.56      L    C
ATOM   4701  C   LEU   41     -36.268  20.127  18.813  1.00 41.26      L    C
ATOM   4702  O   LEU   41     -35.249  20.244  18.131  1.00 40.76      L    O
ATOM   4703  N   PRO   42     -36.374  20.670  20.031  1.00 44.22      L    N
ATOM   4704  CD  PRO   42     -37.515  20.575  20.961  1.00 44.68      L    C
ATOM   4705  CA  PRO   42     -35.270  21.466  20.584  1.00 45.94      L    C
ATOM   4706  CB  PRO   42     -35.733  21.786  22.006  1.00 45.06      L    C
ATOM   4707  CG  PRO   42     -37.239  21.682  21.942  1.00 45.70      L    C
ATOM   4708  C   PRO   42     -33.955  20.684  20.561  1.00 47.33      L    C
ATOM   4709  O   PRO   42     -33.887  19.550  21.042  1.00 47.47      L    O
ATOM   4710  N   GLY   43     -32.924  21.284  19.976  1.00 48.07      L    N
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ATOM	4711	CA	GLY	43	-31.632	20.626	19.902	1.00	49.97	L	C
ATOM	4712	C	GLY	43	-31.587	19.359	19.058	1.00	51.95	L	C
ATOM	4713	O	GLY	43	-30.963	18.374	19.455	1.00	54.06	L	O
ATOM	4714	N	THR	44	-32.245	19.373	17.899	1.00	49.25	L	N
ATOM	4715	CA	THR	44	-32.135	18.274	16.947	1.00	44.82	L	C
ATOM	4716	CB	THR	44	-33.288	17.256	17.111	1.00	46.65	L	C
ATOM	4717	OG1	THR	44	-34.531	17.869	16.741	1.00	48.05	L	O
ATOM	4718	CG2	THR	44	-33.384	16.781	18.558	1.00	46.96	L	C
ATOM	4719	C	THR	44	-32.171	18.812	15.519	1.00	41.95	L	C
ATOM	4720	O	THR	44	-32.424	19.994	15.294	1.00	41.93	L	O
ATOM	4721	N	ALA	45	-31.912	17.939	14.556	1.00	39.35	L	N
ATOM	4722	CA	ALA	45	-32.058	18.293	13.150	1.00	37.05	L	C
ATOM	4723	CB	ALA	45	-31.294	17.312	12.285	1.00	37.54	L	C
ATOM	4724	C	ALA	45	-33.533	18.269	12.774	1.00	35.57	L	C
ATOM	4725	O	ALA	45	-34.326	17.537	13.369	1.00	34.35	L	O
ATOM	4726	N	PRO	46	-33.918	19.070	11.774	1.00	33.17	L	N
ATOM	4727	CD	PRO	46	-33.124	20.113	11.103	1.00	33.74	L	C
ATOM	4728	CA	PRO	46	-35.280	18.991	11.245	1.00	32.23	L	C
ATOM	4729	CB	PRO	46	-35.279	20.002	10.096	1.00	32.13	L	C
ATOM	4730	CG	PRO	46	-34.178	20.960	10.444	1.00	34.08	L	C
ATOM	4731	C	PRO	46	-35.558	17.582	10.755	1.00	31.89	L	C
ATOM	4732	O	PRO	46	-34.652	16.894	10.283	1.00	32.49	L	O
ATOM	4733	N	LYS	47	-36.803	17.141	10.873	1.00	31.47	L	N
ATOM	4734	CA	LYS	47	-37.210	15.914	10.200	1.00	33.45	L	C
ATOM	4735	CB	LYS	47	-37.460	14.790	11.215	1.00	35.12	L	C
ATOM	4736	CG	LYS	47	-38.870	14.755	11.764	1.00	41.61	L	C
ATOM	4737	CD	LYS	47	-39.162	13.420	12.443	1.00	47.61	L	C
ATOM	4738	CE	LYS	47	-40.662	13.196	12.597	1.00	50.11	L	C
ATOM	4739	NZ	LYS	47	-40.987	11.855	13.161	1.00	52.03	L	N
ATOM	4740	C	LYS	47	-38.475	16.169	9.387	1.00	30.96	L	C
ATOM	4741	O	LYS	47	-39.262	17.059	9.712	1.00	28.92	L	O
ATOM	4742	N	LEU	48	-38.661	15.381	8.332	1.00	30.98	L	N
ATOM	4743	CA	LEU	48	-39.814	15.526	7.448	1.00	29.23	L	C
ATOM	4744	CB	LEU	48	-39.668	14.589	6.245	1.00	29.31	L	C
ATOM	4745	CG	LEU	48	-40.813	14.640	5.230	1.00	30.45	L	C
ATOM	4746	CD1	LEU	48	-40.989	16.074	4.740	1.00	28.18	L	C
ATOM	4747	CD2	LEU	48	-40.515	13.697	4.068	1.00	28.52	L	C
ATOM	4748	C	LEU	48	-41.084	15.182	8.208	1.00	27.62	L	C
ATOM	4749	O	LEU	48	-41.135	14.160	8.882	1.00	29.03	L	O
ATOM	4750	N	LEU	49	-42.105	16.028	8.091	1.00	26.47	L	N
ATOM	4751	CA	LEU	49	-43.375	15.826	8.793	1.00	25.82	L	C
ATOM	4752	CB	LEU	49	-43.688	17.046	9.664	1.00	24.24	L	C
ATOM	4753	CG	LEU	49	-44.941	16.965	10.538	1.00	24.60	L	C
ATOM	4754	CD1	LEU	49	-44.685	15.971	11.675	1.00	18.36	L	C
ATOM	4755	CD2	LEU	49	-45.295	18.353	11.086	1.00	20.80	L	C
ATOM	4756	C	LEU	49	-44.544	15.589	7.825	1.00	28.13	L	C
ATOM	4757	O	LEU	49	-45.342	14.669	8.007	1.00	29.97	L	O
ATOM	4758	N	ILE	50	-44.648	16.438	6.808	1.00	26.87	L	N
ATOM	4759	CA	ILE	50	-45.665	16.312	5.770	1.00	27.06	L	C
ATOM	4760	CB	ILE	50	-46.783	17.382	5.917	1.00	26.55	L	C
ATOM	4761	CG2	ILE	50	-47.716	17.325	4.697	1.00	25.11	L	C
ATOM	4762	CG1	ILE	50	-47.569	17.184	7.215	1.00	24.89	L	C
ATOM	4763	CD1	ILE	50	-48.447	15.944	7.230	1.00	26.04	L	C
ATOM	4764	C	ILE	50	-44.968	16.569	4.432	1.00	28.76	L	C
ATOM	4765	O	ILE	50	-44.224	17.547	4.293	1.00	26.93	L	O
ATOM	4766	N	SER	51	-45.201	15.707	3.449	1.00	26.34	L	N
ATOM	4767	CA	SER	51	-44.674	15.963	2.114	1.00	27.63	L	C
ATOM	4768	CB	SER	51	-43.760	14.821	1.670	1.00	26.45	L	C
ATOM	4769	OG	SER	51	-44.481	13.605	1.599	1.00	28.18	L	O
ATOM	4770	C	SER	51	-45.823	16.112	1.124	1.00	28.06	L	C
ATOM	4771	O	SER	51	-46.910	15.553	1.325	1.00	27.05	L	O
ATOM	4772	N	GLY	52	-45.579	16.868	0.059	1.00	28.20	L	N
ATOM	4773	CA	GLY	52	-46.568	16.999	-0.999	1.00	28.39	L	C
ATOM	4774	C	GLY	52	-47.898	17.525	-0.495	1.00	28.92	L	C
ATOM	4775	O	GLY	52	-48.950	16.985	-0.838	1.00	28.40	L	O
ATOM	4776	N	ASN	53	-47.845	18.571	0.328	1.00	26.76	L	N
ATOM	4777	CA	ASN	53	-49.044	19.222	0.868	1.00	28.99	L	C
ATOM	4778	CB	ASN	53	-50.080	19.496	-0.239	1.00	27.64	L	C
ATOM	4779	CG	ASN	53	-49.514	20.313	-1.391	1.00	31.12	L	C
ATOM	4780	OD1	ASN	53	-48.790	21.288	-1.184	1.00	30.52	L	O
ATOM	4781	ND2	ASN	53	-49.846	19.914	-2.617	1.00	28.74	L	N
ATOM	4782	C	ASN	53	-49.737	18.448	1.986	1.00	28.32	L	C
ATOM	4783	O	ASN	53	-50.223	19.047	2.947	1.00	26.59	L	O
ATOM	4784	N	SER	54	-49.814	17.128	1.862	1.00	30.28	L	N
ATOM	4785	CA	SER	54	-50.744	16.385	2.713	1.00	32.77	L	C
ATOM	4786	CB	SER	54	-52.149	16.428	2.105	1.00	31.85	L	C

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ATOM   4787  OG  SER    54     -52.154  15.785   0.842  1.00 36.17      L    O
ATOM   4788  C   SER    54     -50.386  14.935   3.008  1.00 31.72      L    C
ATOM   4789  O   SER    54     -51.165  14.237   3.659  1.00 32.86      L    O
ATOM   4790  N   ASN    55     -49.232  14.473   2.533  1.00 31.26      L    N
ATOM   4791  CA  ASN    55     -48.818  13.085   2.775  1.00 32.61      L    C
ATOM   4792  CB  ASN    55     -48.020  12.551   1.582  1.00 32.07      L    C
ATOM   4793  CG  ASN    55     -48.839  12.526   0.298  1.00 35.95      L    C
ATOM   4794  OD1 ASN    55     -49.720  11.686   0.134  1.00 36.33      L    O
ATOM   4795  ND2 ASN    55     -48.553  13.453  -0.616  1.00 34.48      L    N
ATOM   4796  C   ASN    55     -47.983  12.937   4.051  1.00 32.40      L    C
ATOM   4797  O   ASN    55     -47.071  13.724   4.305  1.00 31.40      L    O
ATOM   4798  N   ARG    56     -48.300  11.920   4.844  1.00 34.00      L    N
ATOM   4799  CA  ARG    56     -47.554  11.626   6.061  1.00 35.54      L    C
ATOM   4800  CB  ARG    56     -48.506  11.199   7.182  1.00 36.63      L    C
ATOM   4801  CG  ARG    56     -49.371  12.328   7.723  1.00 42.78      L    C
ATOM   4802  CD  ARG    56     -50.289  11.856   8.846  1.00 46.38      L    C
ATOM   4803  NE  ARG    56     -51.374  11.010   8.356  1.00 49.53      L    N
ATOM   4804  CZ  ARG    56     -51.380   9.683   8.449  1.00 54.19      L    C
ATOM   4805  NH1 ARG    56     -52.404   8.991   7.971  1.00 55.69      L    N
ATOM   4806  NH2 ARG    56     -50.363   9.047   9.025  1.00 53.47      L    N
ATOM   4807  C   ARG    56     -46.539  10.523   5.822  1.00 35.53      L    C
ATOM   4808  O   ARG    56     -46.886   9.444   5.353  1.00 35.07      L    O
ATOM   4809  N   PRO    57     -45.267  10.781   6.154  1.00 36.46      L    N
ATOM   4810  CD  PRO    57     -44.712  12.097   6.516  1.00 35.66      L    C
ATOM   4811  CA  PRO    57     -44.244   9.729   6.163  1.00 37.45      L    C
ATOM   4812  CB  PRO    57     -42.968  10.464   6.581  1.00 36.26      L    C
ATOM   4813  CG  PRO    57     -43.240  11.915   6.281  1.00 37.45      L    C
ATOM   4814  C   PRO    57     -44.613   8.632   7.167  1.00 39.29      L    C
ATOM   4815  O   PRO    57     -45.359   8.875   8.121  1.00 37.09      L    O
ATOM   4816  N   SER    58     -44.091   7.430   6.944  1.00 40.87      L    N
ATOM   4817  CA  SER    58     -44.130   6.386   7.965  1.00 43.46      L    C
ATOM   4818  CB  SER    58     -43.252   5.199   7.557  1.00 43.89      L    C
ATOM   4819  OG  SER    58     -43.995   4.260   6.803  1.00 51.42      L    O
ATOM   4820  C   SER    58     -43.621   6.936   9.285  1.00 41.83      L    C
ATOM   4821  O   SER    58     -42.569   7.575   9.338  1.00 43.28      L    O
ATOM   4822  N   GLY    59     -44.363   6.681  10.353  1.00 40.66      L    N
ATOM   4823  CA  GLY    59     -43.914   7.120  11.658  1.00 40.19      L    C
ATOM   4824  C   GLY    59     -44.533   8.423  12.118  1.00 40.38      L    C
ATOM   4825  O   GLY    59     -44.383   8.793  13.278  1.00 44.27      L    O
ATOM   4826  N   VAL    60     -45.225   9.128  11.228  1.00 38.65      L    N
ATOM   4827  CA  VAL    60     -45.922  10.342  11.627  1.00 36.24      L    C
ATOM   4828  CB  VAL    60     -45.788  11.446  10.550  1.00 36.26      L    C
ATOM   4829  CG1 VAL    60     -46.661  12.642  10.917  1.00 33.36      L    C
ATOM   4830  CG2 VAL    60     -44.329  11.874  10.425  1.00 31.61      L    C
ATOM   4831  C   VAL    60     -47.398  10.061  11.882  1.00 36.29      L    C
ATOM   4832  O   VAL    60     -48.119   9.613  10.992  1.00 36.29      L    O
ATOM   4833  N   PRO    61     -47.866  10.327  13.113  1.00 36.76      L    N
ATOM   4834  CD  PRO    61     -47.074  10.985  14.165  1.00 37.16      L    C
ATOM   4835  CA  PRO    61     -49.233  10.023  13.560  1.00 36.64      L    C
ATOM   4836  CB  PRO    61     -49.262  10.509  15.010  1.00 37.12      L    C
ATOM   4837  CG  PRO    61     -47.829  10.647  15.415  1.00 38.26      L    C
ATOM   4838  C   PRO    61     -50.271  10.762  12.714  1.00 38.15      L    C
ATOM   4839  O   PRO    61     -49.997  11.847  12.193  1.00 36.93      L    O
ATOM   4840  N   ASP    62     -51.464  10.189  12.594  1.00 37.72      L    N
ATOM   4841  CA  ASP    62     -52.509  10.802  11.779  1.00 39.64      L    C
ATOM   4842  CB  ASP    62     -53.630   9.790  11.491  1.00 45.58      L    C
ATOM   4843  CG  ASP    62     -54.249   9.209  12.763  1.00 52.82      L    C
ATOM   4844  OD1 ASP    62     -53.929   9.692  13.877  1.00 54.52      L    O
ATOM   4845  OD2 ASP    62     -55.063   8.262  12.642  1.00 57.04      L    O
ATOM   4846  C   ASP    62     -53.085  12.054  12.440  1.00 36.85      L    C
ATOM   4847  O   ASP    62     -53.968  12.709  11.887  1.00 35.85      L    O
ATOM   4848  N   ARG    63     -52.579  12.383  13.623  1.00 33.17      L    N
ATOM   4849  CA  ARG    63     -52.920  13.642  14.275  1.00 33.59      L    C
ATOM   4850  CB  ARG    63     -52.253  13.728  15.658  1.00 34.57      L    C
ATOM   4851  CG  ARG    63     -52.537  12.529  16.546  1.00 40.31      L    C
ATOM   4852  CD  ARG    63     -52.038  12.725  17.983  1.00 41.07      L    C
ATOM   4853  NE  ARG    63     -50.578  12.668  18.119  1.00 39.33      L    N
ATOM   4854  CZ  ARG    63     -49.811  13.746  18.221  1.00 36.57      L    C
ATOM   4855  NH1 ARG    63     -50.373  14.944  18.188  1.00 34.87      L    N
ATOM   4856  NH2 ARG    63     -48.499  13.631  18.394  1.00 34.48      L    N
ATOM   4857  C   ARG    63     -52.455  14.819  13.413  1.00 30.38      L    C
ATOM   4858  O   ARG    63     -53.024  15.908  13.483  1.00 28.85      L    O
ATOM   4859  N   PHE    64     -51.417  14.592  12.613  1.00 27.74      L    N
ATOM   4860  CA  PHE    64     -50.875  15.637  11.736  1.00 29.98      L    C
ATOM   4861  CB  PHE    64     -49.348  15.502  11.621  1.00 26.68      L    C
ATOM   4862  CG  PHE    64     -48.608  15.769  12.911  1.00 28.95      L    C
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ATOM	4863	CD1	PHE	64	-48.214	17.062	13.247	1.00	27.02	L C
ATOM	4864	CD2	PHE	64	-48.295	14.729	13.780	1.00	27.70	L C
ATOM	4865	CE1	PHE	64	-47.518	17.313	14.431	1.00	27.84	L C
ATOM	4866	CE2	PHE	64	-47.599	14.972	14.967	1.00	28.68	L C
ATOM	4867	CZ	PHE	64	-47.211	16.267	15.291	1.00	27.70	L C
ATOM	4868	C	PHE	64	-51.487	15.548	10.340	1.00	30.00	L C
ATOM	4869	O	PHE	64	-51.518	14.475	9.744	1.00	29.23	L O
ATOM	4870	N	SER	65	-51.963	16.673	9.819	1.00	29.23	L N
ATOM	4871	CA	SER	65	-52.463	16.722	8.445	1.00	31.26	L C
ATOM	4872	CB	SER	65	-53.980	16.499	8.413	1.00	30.43	L C
ATOM	4873	OG	SER	65	-54.665	17.575	9.032	1.00	32.55	L O
ATOM	4874	C	SER	65	-52.135	18.061	7.787	1.00	31.18	L C
ATOM	4875	O	SER	65	-51.937	19.066	8.470	1.00	31.63	L O
ATOM	4876	N	GLY	66	-52.077	18.066	6.460	1.00	30.29	L N
ATOM	4877	CA	GLY	66	-51.749	19.283	5.746	1.00	32.02	L C
ATOM	4878	C	GLY	66	-52.688	19.542	4.585	1.00	34.44	L C
ATOM	4879	O	GLY	66	-53.314	18.624	4.064	1.00	35.30	L O
ATOM	4880	N	SER	67	-52.793	20.798	4.174	1.00	35.63	L N
ATOM	4881	CA	SER	67	-53.540	21.126	2.971	1.00	37.94	L C
ATOM	4882	CB	SER	67	-55.018	21.369	3.304	1.00	37.38	L C
ATOM	4883	OG	SER	67	-55.168	22.386	4.280	1.00	39.96	L O
ATOM	4884	C	SER	67	-52.954	22.358	2.300	1.00	39.49	L C
ATOM	4885	O	SER	67	-52.248	23.150	2.936	1.00	39.23	L O
ATOM	4886	N	LYS	68	-53.252	22.507	1.012	1.00	40.52	L N
ATOM	4887	CA	LYS	68	-52.826	23.666	0.236	1.00	42.33	L C
ATOM	4888	CB	LYS	68	-51.729	23.267	-0.755	1.00	42.24	L C
ATOM	4889	CG	LYS	68	-51.463	24.323	-1.831	1.00	44.74	L C
ATOM	4890	CD	LYS	68	-50.479	23.832	-2.890	1.00	46.29	L C
ATOM	4891	CE	LYS	68	-50.427	24.782	-4.086	1.00	47.53	L C
ATOM	4892	NZ	LYS	68	-49.320	24.451	-5.037	1.00	44.81	L N
ATOM	4893	C	LYS	68	-54.004	24.268	-0.531	1.00	42.80	L C
ATOM	4894	O	LYS	68	-54.869	23.549	-1.026	1.00	41.34	L O
ATOM	4895	N	SER	69	-54.026	25.590	-0.640	1.00	43.73	L N
ATOM	4896	CA	SER	69	-55.090	26.268	-1.366	1.00	44.46	L C
ATOM	4897	CB	SER	69	-56.310	26.424	-0.458	1.00	46.75	L C
ATOM	4898	OG	SER	69	-57.267	27.293	-1.036	1.00	51.75	L O
ATOM	4899	C	SER	69	-54.635	27.637	-1.867	1.00	43.71	L C
ATOM	4900	O	SER	69	-54.309	28.523	-1.074	1.00	42.73	L O
ATOM	4901	N	GLY	70	-54.616	27.807	-3.186	1.00	43.48	L N
ATOM	4902	CA	GLY	70	-54.209	29.082	-3.749	1.00	41.87	L C
ATOM	4903	C	GLY	70	-52.744	29.356	-3.470	1.00	42.01	L C
ATOM	4904	O	GLY	70	-51.877	28.578	-3.871	1.00	42.41	L O
ATOM	4905	N	THR	71	-52.460	30.453	-2.775	1.00	40.06	L N
ATOM	4906	CA	THR	71	-51.083	30.799	-2.443	1.00	40.11	L C
ATOM	4907	CB	THR	71	-50.801	32.283	-2.712	1.00	40.96	L C
ATOM	4908	OG1	THR	71	-51.714	33.081	-1.951	1.00	40.94	L O
ATOM	4909	CG2	THR	71	-50.953	32.601	-4.201	1.00	39.96	L C
ATOM	4910	C	THR	71	-50.732	30.511	-0.981	1.00	39.88	L C
ATOM	4911	O	THR	71	-49.700	30.963	-0.487	1.00	40.96	L O
ATOM	4912	N	SER	72	-51.590	29.774	-0.285	1.00	37.34	L N
ATOM	4913	CA	SER	72	-51.303	29.436	1.100	1.00	37.58	L C
ATOM	4914	CB	SER	72	-52.187	30.259	2.041	1.00	38.75	L C
ATOM	4915	OG	SER	72	-53.541	29.863	1.940	1.00	43.69	L O
ATOM	4916	C	SER	72	-51.471	27.941	1.392	1.00	36.49	L C
ATOM	4917	O	SER	72	-52.095	27.201	0.619	1.00	34.73	L O
ATOM	4918	N	ALA	73	-50.880	27.504	2.500	1.00	33.01	L N
ATOM	4919	CA	ALA	73	-50.997	26.124	2.950	1.00	31.45	L C
ATOM	4920	CB	ALA	73	-49.767	25.328	2.543	1.00	30.50	L C
ATOM	4921	C	ALA	73	-51.154	26.114	4.463	1.00	31.13	L C
ATOM	4922	O	ALA	73	-50.896	27.113	5.132	1.00	30.70	L O
ATOM	4923	N	SER	74	-51.583	24.985	5.006	1.00	30.38	L N
ATOM	4924	CA	SER	74	-51.886	24.932	6.421	1.00	32.16	L C
ATOM	4925	CB	SER	74	-53.375	25.219	6.627	1.00	32.93	L C
ATOM	4926	OG	SER	74	-53.697	25.321	8.002	1.00	40.40	L O
ATOM	4927	C	SER	74	-51.510	23.574	7.001	1.00	30.96	L C
ATOM	4928	O	SER	74	-51.747	22.540	6.379	1.00	32.04	L O
ATOM	4929	N	LEU	75	-50.902	23.585	8.184	1.00	31.07	L N
ATOM	4930	CA	LEU	75	-50.638	22.354	8.928	1.00	29.53	L C
ATOM	4931	CB	LEU	75	-49.192	22.336	9.441	1.00	26.68	L C
ATOM	4932	CG	LEU	75	-48.781	21.170	10.362	1.00	28.53	L C
ATOM	4933	CD1	LEU	75	-48.627	19.878	9.553	1.00	23.41	L C
ATOM	4934	CD2	LEU	75	-47.456	21.510	11.059	1.00	24.35	L C
ATOM	4935	C	LEU	75	-51.607	22.306	10.107	1.00	31.37	L C
ATOM	4936	O	LEU	75	-51.767	23.296	10.827	1.00	30.48	L O
ATOM	4937	N	ALA	76	-52.263	21.163	10.300	1.00	30.74	L N
ATOM	4938	CA	ALA	76	-53.173	21.007	11.425	1.00	30.46	L C

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ATOM   4939  CB  ALA    76     -54.600  20.778  10.930  1.00 27.72      L    C
ATOM   4940  C   ALA    76     -52.737  19.857  12.316  1.00 30.96      L    C
ATOM   4941  O   ALA    76     -52.334  18.803  11.833  1.00 31.40      L    O
ATOM   4942  N   ILE    77     -52.822  20.077  13.624  1.00 31.12      L    N
ATOM   4943  CA  ILE    77     -52.447  19.075  14.612  1.00 31.39      L    C
ATOM   4944  CB  ILE    77     -51.234  19.550  15.440  1.00 32.11      L    C
ATOM   4945  CG2 ILE    77     -50.800  18.460  16.412  1.00 31.86      L    C
ATOM   4946  CG1 ILE    77     -50.085  19.930  14.505  1.00 33.01      L    C
ATOM   4947  CD1 ILE    77     -48.913  20.579  15.210  1.00 31.72      L    C
ATOM   4948  C   ILE    77     -53.630  18.887  15.550  1.00 31.92      L    C
ATOM   4949  O   ILE    77     -54.000  19.811  16.271  1.00 33.01      L    O
ATOM   4950  N   THR    78     -54.231  17.701  15.538  1.00 32.75      L    N
ATOM   4951  CA  THR    78     -55.314  17.401  16.474  1.00 34.22      L    C
ATOM   4952  CB  THR    78     -56.425  16.563  15.808  1.00 36.19      L    C
ATOM   4953  OG1 THR    78     -55.877  15.314  15.369  1.00 37.55      L    O
ATOM   4954  CG2 THR    78     -57.014  17.311  14.610  1.00 33.93      L    C
ATOM   4955  C   THR    78     -54.750  16.618  17.652  1.00 33.72      L    C
ATOM   4956  O   THR    78     -53.631  16.110  17.582  1.00 35.50      L    O
ATOM   4957  N   GLY    79     -55.514  16.529  18.735  1.00 33.77      L    N
ATOM   4958  CA  GLY    79     -55.052  15.777  19.891  1.00 31.44      L    C
ATOM   4959  C   GLY    79     -53.649  16.178  20.312  1.00 30.87      L    C
ATOM   4960  O   GLY    79     -52.798  15.327  20.543  1.00 31.97      L    O
ATOM   4961  N   LEU    80     -53.406  17.480  20.410  1.00 29.82      L    N
ATOM   4962  CA  LEU    80     -52.082  18.001  20.731  1.00 30.55      L    C
ATOM   4963  CB  LEU    80     -52.200  19.489  21.088  1.00 31.73      L    C
ATOM   4964  CG  LEU    80     -50.929  20.337  21.084  1.00 31.97      L    C
ATOM   4965  CD1 LEU    80     -50.314  20.326  19.690  1.00 31.53      L    C
ATOM   4966  CD2 LEU    80     -51.271  21.764  21.495  1.00 33.68      L    C
ATOM   4967  C   LEU    80     -51.443  17.231  21.894  1.00 31.41      L    C
ATOM   4968  O   LEU    80     -52.080  17.015  22.924  1.00 28.69      L    O
ATOM   4969  N   GLN    81     -50.187  16.816  21.718  1.00 33.22      L    N
ATOM   4970  CA  GLN    81     -49.425  16.126  22.769  1.00 35.00      L    C
ATOM   4971  CB  GLN    81     -48.930  14.760  22.277  1.00 36.97      L    C
ATOM   4972  CG  GLN    81     -50.027  13.790  21.874  1.00 46.26      L    C
ATOM   4973  CD  GLN    81     -50.816  13.277  23.066  1.00 52.65      L    C
ATOM   4974  OE1 GLN    81     -50.313  12.475  23.858  1.00 56.12      L    O
ATOM   4975  NE2 GLN    81     -52.061  13.739  23.203  1.00 54.96      L    N
ATOM   4976  C   GLN    81     -48.213  16.952  23.190  1.00 34.68      L    C
ATOM   4977  O   GLN    81     -47.676  17.731  22.395  1.00 32.67      L    O
ATOM   4978  N   ALA    82     -47.772  16.754  24.431  1.00 34.64      L    N
ATOM   4979  CA  ALA    82     -46.587  17.427  24.966  1.00 35.20      L    C
ATOM   4980  CB  ALA    82     -46.257  16.872  26.356  1.00 36.44      L    C
ATOM   4981  C   ALA    82     -45.359  17.313  24.069  1.00 34.25      L    C
ATOM   4982  O   ALA    82     -44.597  18.273  23.923  1.00 35.41      L    O
ATOM   4983  N   GLU    83     -45.153  16.145  23.475  1.00 33.73      L    N
ATOM   4984  CA  GLU    83     -43.982  15.936  22.629  1.00 36.46      L    C
ATOM   4985  CB  GLU    83     -43.768  14.438  22.371  1.00 37.92      L    C
ATOM   4986  CG  GLU    83     -44.991  13.694  21.844  1.00 48.23      L    C
ATOM   4987  CD  GLU    83     -45.924  13.198  22.957  1.00 55.83      L    C
ATOM   4988  OE1 GLU    83     -46.852  12.410  22.641  1.00 60.92      L    O
ATOM   4989  OE2 GLU    83     -45.741  13.588  24.140  1.00 54.46      L    O
ATOM   4990  C   GLU    83     -44.070  16.693  21.295  1.00 35.87      L    C
ATOM   4991  O   GLU    83     -43.115  16.696  20.518  1.00 36.46      L    O
ATOM   4992  N   ASP    84     -45.206  17.342  21.040  1.00 33.53      L    N
ATOM   4993  CA  ASP    84     -45.370  18.148  19.830  1.00 32.04      L    C
ATOM   4994  CB  ASP    84     -46.851  18.329  19.482  1.00 29.82      L    C
ATOM   4995  CG  ASP    84     -47.539  17.024  19.156  1.00 33.55      L    C
ATOM   4996  OD1 ASP    84     -46.859  16.092  18.667  1.00 33.27      L    O
ATOM   4997  OD2 ASP    84     -48.763  16.930  19.390  1.00 31.73      L    O
ATOM   4998  C   ASP    84     -44.737  19.528  19.979  1.00 32.06      L    C
ATOM   4999  O   ASP    84     -44.653  20.279  19.003  1.00 30.80      L    O
ATOM   5000  N   GLU    85     -44.309  19.877  21.191  1.00 29.59      L    N
ATOM   5001  CA  GLU    85     -43.660  21.169  21.389  1.00 31.50      L    C
ATOM   5002  CB  GLU    85     -43.317  21.397  22.870  1.00 33.60      L    C
ATOM   5003  CG  GLU    85     -42.876  22.829  23.172  1.00 38.51      L    C
ATOM   5004  CD  GLU    85     -43.034  23.215  24.641  1.00 43.82      L    C
ATOM   5005  OE1 GLU    85     -42.004  23.517  25.290  1.00 47.43      L    O
ATOM   5006  OE2 GLU    85     -44.182  23.229  25.150  1.00 42.38      L    O
ATOM   5007  C   GLU    85     -42.391  21.182  20.549  1.00 29.61      L    C
ATOM   5008  O   GLU    85     -41.542  20.307  20.688  1.00 32.34      L    O
ATOM   5009  N   ALA    86     -42.269  22.164  19.664  1.00 26.98      L    N
ATOM   5010  CA  ALA    86     -41.217  22.135  18.656  1.00 26.63      L    C
ATOM   5011  CB  ALA    86     -41.295  20.824  17.850  1.00 27.34      L    C
ATOM   5012  C   ALA    86     -41.378  23.323  17.725  1.00 27.49      L    C
ATOM   5013  O   ALA    86     -42.335  24.084  17.844  1.00 28.13      L    O
ATOM   5014  N   ASP    87     -40.437  23.480  16.803  1.00 28.02      L    N
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ATOM   5015  CA  ASP    87     -40.565  24.476  15.750  1.00 29.17      L    C
ATOM   5016  CB  ASP    87     -39.227  25.188  15.534  1.00 31.60      L    C
ATOM   5017  CG  ASP    87     -38.845  26.086  16.712  1.00 36.68      L    C
ATOM   5018  OD1 ASP    87     -39.736  26.768  17.260  1.00 40.12      L    O
ATOM   5019  OD2 ASP    87     -37.653  26.111  17.095  1.00 38.12      L    O
ATOM   5020  C   ASP    87     -41.007  23.786  14.458  1.00 28.96      L    C
ATOM   5021  O   ASP    87     -40.577  22.664  14.164  1.00 26.69      L    O
ATOM   5022  N   TYR    88     -41.874  24.452  13.699  1.00 26.76      L    N
ATOM   5023  CA  TYR    88     -42.396  23.884  12.462  1.00 27.47      L    C
ATOM   5024  CB  TYR    88     -43.904  23.641  12.576  1.00 25.61      L    C
ATOM   5025  CG  TYR    88     -44.269  22.601  13.607  1.00 23.78      L    C
ATOM   5026  CD1 TYR    88     -44.328  22.921  14.965  1.00 24.12      L    C
ATOM   5027  CE1 TYR    88     -44.617  21.946  15.920  1.00 22.16      L    C
ATOM   5028  CD2 TYR    88     -44.514  21.288  13.231  1.00 22.59      L    C
ATOM   5029  CE2 TYR    88     -44.802  20.316  14.166  1.00 22.72      L    C
ATOM   5030  CZ  TYR    88     -44.849  20.648  15.511  1.00 22.94      L    C
ATOM   5031  OH  TYR    88     -45.104  19.661  16.437  1.00 21.22      L    O
ATOM   5032  C   TYR    88     -42.117  24.816  11.298  1.00 26.66      L    C
ATOM   5033  O   TYR    88     -42.384  26.008  11.379  1.00 29.21      L    O
ATOM   5034  N   TYR    89     -41.580  24.261  10.219  1.00 27.15      L    N
ATOM   5035  CA  TYR    89     -41.213  25.044   9.044  1.00 25.90      L    C
ATOM   5036  CB  TYR    89     -39.698  24.971   8.814  1.00 26.15      L    C
ATOM   5037  CG  TYR    89     -38.873  25.579   9.931  1.00 28.08      L    C
ATOM   5038  CD1 TYR    89     -38.414  24.801  10.992  1.00 28.94      L    C
ATOM   5039  CE1 TYR    89     -37.671  25.363  12.027  1.00 27.99      L    C
ATOM   5040  CD2 TYR    89     -38.566  26.934   9.934  1.00 27.68      L    C
ATOM   5041  CE2 TYR    89     -37.829  27.503  10.962  1.00 28.70      L    C
ATOM   5042  CZ  TYR    89     -37.384  26.712  12.006  1.00 30.96      L    C
ATOM   5043  OH  TYR    89     -36.651  27.279  13.031  1.00 32.52      L    O
ATOM   5044  C   TYR    89     -41.935  24.510   7.811  1.00 26.61      L    C
ATOM   5045  O   TYR    89     -41.986  23.296   7.584  1.00 24.13      L    O
ATOM   5046  N   CYS    90     -42.493  25.410   7.011  1.00 26.60      L    N
ATOM   5047  CA  CYS    90     -42.966  25.018   5.694  1.00 25.92      L    C
ATOM   5048  C   CYS    90     -41.898  25.294   4.644  1.00 27.13      L    C
ATOM   5049  O   CYS    90     -40.957  26.053   4.877  1.00 27.83      L    O
ATOM   5050  CB  CYS    90     -44.270  25.748   5.342  1.00 29.34      L    C
ATOM   5051  SG  CYS    90     -44.257  27.570   5.342  1.00 38.10      L    S
ATOM   5052  N   GLN    91     -42.036  24.657   3.488  1.00 28.80      L    N
ATOM   5053  CA  GLN    91     -41.044  24.779   2.429  1.00 28.24      L    C
ATOM   5054  CB  GLN    91     -39.966  23.708   2.595  1.00 27.60      L    C
ATOM   5055  CG  GLN    91     -38.890  23.735   1.525  1.00 29.58      L    C
ATOM   5056  CD  GLN    91     -38.225  22.383   1.351  1.00 33.53      L    C
ATOM   5057  OE1 GLN    91     -38.893  21.347   1.360  1.00 30.27      L    O
ATOM   5058  NE2 GLN    91     -36.896  22.382   1.198  1.00 35.14      L    N
ATOM   5059  C   GLN    91     -41.718  24.615   1.078  1.00 29.35      L    C
ATOM   5060  O   GLN    91     -42.679  23.851   0.946  1.00 29.91      L    O
ATOM   5061  N   SER    92     -41.210  25.330   0.076  1.00 29.56      L    N
ATOM   5062  CA  SER    92     -41.704  25.207  -1.293  1.00 31.14      L    C
ATOM   5063  CB  SER    92     -43.013  25.991  -1.451  1.00 32.55      L    C
ATOM   5064  OG  SER    92     -43.443  26.023  -2.802  1.00 34.97      L    O
ATOM   5065  C   SER    92     -40.668  25.731  -2.285  1.00 30.79      L    C
ATOM   5066  O   SER    92     -39.914  26.658  -1.979  1.00 30.97      L    O
ATOM   5067  N   TYR    93     -40.627  25.140  -3.473  1.00 29.05      L    N
ATOM   5068  CA  TYR    93     -39.728  25.633  -4.507  1.00 29.06      L    C
ATOM   5069  CB  TYR    93     -39.677  24.663  -5.698  1.00 28.59      L    C
ATOM   5070  CG  TYR    93     -38.685  25.053  -6.783  1.00 30.48      L    C
ATOM   5071  CD1 TYR    93     -37.317  24.858  -6.609  1.00 31.57      L    C
ATOM   5072  CE1 TYR    93     -36.406  25.209  -7.599  1.00 30.71      L    C
ATOM   5073  CD2 TYR    93     -39.120  25.612  -7.983  1.00 31.67      L    C
ATOM   5074  CE2 TYR    93     -38.220  25.968  -8.984  1.00 30.85      L    C
ATOM   5075  CZ  TYR    93     -36.865  25.765  -8.781  1.00 33.32      L    C
ATOM   5076  OH  TYR    93     -35.969  26.129  -9.760  1.00 32.49      L    O
ATOM   5077  C   TYR    93     -40.233  26.996  -4.958  1.00 29.39      L    C
ATOM   5078  O   TYR    93     -41.441  27.234  -5.017  1.00 27.21      L    O
ATOM   5079  N   ASP    94     -39.303  27.898  -5.248  1.00 29.82      L    N
ATOM   5080  CA  ASP    94     -39.652  29.201  -5.780  1.00 31.94      L    C
ATOM   5081  CB  ASP    94     -39.229  30.295  -4.811  1.00 34.22      L    C
ATOM   5082  CG  ASP    94     -39.679  31.671  -5.256  1.00 39.09      L    C
ATOM   5083  OD1 ASP    94     -40.570  32.253  -4.589  1.00 38.37      L    O
ATOM   5084  OD2 ASP    94     -39.138  32.169  -6.272  1.00 41.24      L    O
ATOM   5085  C   ASP    94     -38.932  29.379  -7.110  1.00 33.39      L    C
ATOM   5086  O   ASP    94     -37.701  29.381  -7.157  1.00 32.83      L    O
ATOM   5087  N   SER    95     -39.694  29.531  -8.189  1.00 32.19      L    N
ATOM   5088  CA  SER    95     -39.098  29.548  -9.519  1.00 35.46      L    C
ATOM   5089  CB  SER    95     -40.176  29.372 -10.598  1.00 33.41      L    C
ATOM   5090  OG  SER    95     -41.217  30.322 -10.442  1.00 37.34      L    O
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ATOM   5091  C   SER    95     -38.274  30.797  -9.811  1.00 35.42      L    C
ATOM   5092  O   SER    95     -37.381  30.757 -10.651  1.00 38.68      L    O
ATOM   5093  N   SER    96     -38.547  31.905  -9.132  1.00 37.35      L    N
ATOM   5094  CA  SER    96     -37.712  33.086  -9.343  1.00 39.64      L    C
ATOM   5095  CB  SER    96     -38.461  34.365  -8.947  1.00 38.48      L    C
ATOM   5096  OG  SER    96     -38.428  34.586  -7.548  1.00 48.03      L    O
ATOM   5097  C   SER    96     -36.378  32.997  -8.589  1.00 40.42      L    C
ATOM   5098  O   SER    96     -35.411  33.664  -8.951  1.00 41.72      L    O
ATOM   5099  N   LEU    97     -36.317  32.164  -7.552  1.00 40.63      L    N
ATOM   5100  CA  LEU    97     -35.085  32.015  -6.776  1.00 39.89      L    C
ATOM   5101  CB  LEU    97     -35.403  32.010  -5.278  1.00 39.69      L    C
ATOM   5102  CG  LEU    97     -36.119  33.256  -4.752  1.00 41.84      L    C
ATOM   5103  CD1 LEU    97     -36.377  33.113  -3.257  1.00 40.69      L    C
ATOM   5104  CD2 LEU    97     -35.271  34.492  -5.034  1.00 39.18      L    C
ATOM   5105  C   LEU    97     -34.330  30.739  -7.142  1.00 40.12      L    C
ATOM   5106  O   LEU    97     -33.166  30.567  -6.775  1.00 38.81      L    O
ATOM   5107  N   SER    98     -35.005  29.847  -7.862  1.00 40.10      L    N
ATOM   5108  CA  SER    98     -34.432  28.559  -8.247  1.00 39.46      L    C
ATOM   5109  CB  SER    98     -33.228  28.768  -9.169  1.00 39.43      L    C
ATOM   5110  OG  SER    98     -33.610  29.450 -10.354  1.00 42.81      L    O
ATOM   5111  C   SER    98     -34.018  27.713  -7.045  1.00 38.54      L    C
ATOM   5112  O   SER    98     -33.073  26.923  -7.123  1.00 38.98      L    O
ATOM   5113  N   GLY    99     -34.728  27.871  -5.934  1.00 37.42      L    N
ATOM   5114  CA  GLY    99     -34.422  27.070  -4.762  1.00 35.97      L    C
ATOM   5115  C   GLY    99     -35.657  26.785  -3.940  1.00 36.31      L    C
ATOM   5116  O   GLY    99     -36.657  27.493  -4.065  1.00 35.39      L    O
ATOM   5117  N   SER   100     -35.588  25.752  -3.100  1.00 36.91      L    N
ATOM   5118  CA  SER   100     -36.707  25.374  -2.246  1.00 37.76      L    C
ATOM   5119  CB  SER   100     -36.765  23.849  -2.076  1.00 39.33      L    C
ATOM   5120  OG  SER   100     -37.335  23.211  -3.212  1.00 39.19      L    O
ATOM   5121  C   SER   100     -36.563  26.035  -0.885  1.00 38.62      L    C
ATOM   5122  O   SER   100     -35.908  25.493   0.012  1.00 40.69      L    O
ATOM   5123  N   VAL   101     -37.197  27.195  -0.735  1.00 35.61      L    N
ATOM   5124  CA  VAL   101     -37.020  28.048   0.428  1.00 34.08      L    C
ATOM   5125  CB  VAL   101     -37.142  29.527   0.033  1.00 34.68      L    C
ATOM   5126  CG1 VAL   101     -36.015  29.890  -0.932  1.00 37.67      L    C
ATOM   5127  CG2 VAL   101     -38.501  29.780  -0.605  1.00 30.50      L    C
ATOM   5128  C   VAL   101     -38.002  27.768   1.563  1.00 33.19      L    C
ATOM   5129  O   VAL   101     -39.047  27.151   1.362  1.00 31.98      L    O
ATOM   5130  N   PHE   102     -37.652  28.246   2.755  1.00 30.75      L    N
ATOM   5131  CA  PHE   102     -38.362  27.915   3.985  1.00 29.44      L    C
ATOM   5132  CB  PHE   102     -37.361  27.473   5.053  1.00 28.91      L    C
ATOM   5133  CG  PHE   102     -36.680  26.179   4.742  1.00 27.81      L    C
ATOM   5134  CD1 PHE   102     -35.472  26.161   4.070  1.00 26.76      L    C
ATOM   5135  CD2 PHE   102     -37.259  24.973   5.110  1.00 26.32      L    C
ATOM   5136  CE1 PHE   102     -34.853  24.960   3.764  1.00 26.61      L    C
ATOM   5137  CE2 PHE   102     -36.645  23.777   4.809  1.00 25.33      L    C
ATOM   5138  CZ  PHE   102     -35.439  23.770   4.133  1.00 25.96      L    C
ATOM   5139  C   PHE   102     -39.158  29.094   4.521  1.00 28.98      L    C
ATOM   5140  O   PHE   102     -38.750  30.244   4.372  1.00 31.15      L    O
ATOM   5141  N   GLY   103     -40.292  28.811   5.154  1.00 28.82      L    N
ATOM   5142  CA  GLY   103     -40.949  29.830   5.956  1.00 27.61      L    C
ATOM   5143  C   GLY   103     -40.094  30.221   7.153  1.00 28.18      L    C
ATOM   5144  O   GLY   103     -39.068  29.593   7.419  1.00 27.21      L    O
ATOM   5145  N   GLY   104     -40.513  31.257   7.877  1.00 29.41      L    N
ATOM   5146  CA  GLY   104     -39.722  31.760   8.987  1.00 30.21      L    C
ATOM   5147  C   GLY   104     -39.801  30.907  10.245  1.00 31.93      L    C
ATOM   5148  O   GLY   104     -39.038  31.114  11.186  1.00 31.09      L    O
ATOM   5149  N   GLY   105     -40.715  29.941  10.262  1.00 31.73      L    N
ATOM   5150  CA  GLY   105     -40.796  29.024  11.385  1.00 30.36      L    C
ATOM   5151  C   GLY   105     -41.890  29.408  12.360  1.00 30.83      L    C
ATOM   5152  O   GLY   105     -42.133  30.591  12.597  1.00 30.41      L    O
ATOM   5153  N   THR   106     -42.557  28.408  12.924  1.00 30.56      L    N
ATOM   5154  CA  THR   106     -43.546  28.648  13.969  1.00 30.39      L    C
ATOM   5155  CB  THR   106     -44.950  28.202  13.538  1.00 30.35      L    C
ATOM   5156  OG1 THR   106     -45.338  28.900  12.354  1.00 32.39      L    O
ATOM   5157  CG2 THR   106     -45.950  28.490  14.635  1.00 30.52      L    C
ATOM   5158  C   THR   106     -43.170  27.865  15.219  1.00 30.74      L    C
ATOM   5159  O   THR   106     -42.953  26.655  15.167  1.00 30.35      L    O
ATOM   5160  N   LYS   107     -43.099  28.568  16.341  1.00 31.45      L    N
ATOM   5161  CA  LYS   107     -42.788  27.954  17.623  1.00 33.01      L    C
ATOM   5162  CB  LYS   107     -42.130  28.999  18.540  1.00 38.62      L    C
ATOM   5163  CG  LYS   107     -41.064  28.452  19.485  1.00 44.72      L    C
ATOM   5164  CD  LYS   107     -41.600  27.310  20.334  1.00 49.04      L    C
ATOM   5165  CE  LYS   107     -40.472  26.415  20.831  1.00 51.13      L    C
ATOM   5166  NZ  LYS   107     -40.985  25.086  21.262  1.00 50.21      L    N
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ATOM   5167  C   LYS   107     -44.107  27.486  18.230  1.00 31.74      L    C
ATOM   5168  O   LYS   107     -44.979  28.307  18.538  1.00 32.77      L    O
ATOM   5169  N   LEU   108     -44.273  26.179  18.393  1.00 28.38      L    N
ATOM   5170  CA  LEU   108     -45.486  25.668  19.037  1.00 30.04      L    C
ATOM   5171  CB  LEU   108     -45.970  24.391  18.351  1.00 26.99      L    C
ATOM   5172  CG  LEU   108     -47.289  23.796  18.861  1.00 29.30      L    C
ATOM   5173  CD1 LEU   108     -47.883  22.884  17.802  1.00 29.60      L    C
ATOM   5174  CD2 LEU   108     -47.058  23.018  20.150  1.00 31.06      L    C
ATOM   5175  C   LEU   108     -45.256  25.376  20.515  1.00 29.02      L    C
ATOM   5176  O   LEU   108     -44.342  24.647  20.871  1.00 31.80      L    O
ATOM   5177  N   THR   109     -46.112  25.921  21.367  1.00 30.44      L    N
ATOM   5178  CA  THR   109     -45.992  25.733  22.812  1.00 32.72      L    C
ATOM   5179  CB  THR   109     -45.942  27.101  23.535  1.00 33.32      L    C
ATOM   5180  OG1 THR   109     -44.728  27.776  23.190  1.00 39.35      L    O
ATOM   5181  CG2 THR   109     -45.991  26.924  25.030  1.00 36.60      L    C
ATOM   5182  C   THR   109     -47.173  24.932  23.352  1.00 31.37      L    C
ATOM   5183  O   THR   109     -48.326  25.247  23.056  1.00 32.44      L    O
ATOM   5184  N   VAL   110     -46.890  23.901  24.141  1.00 29.28      L    N
ATOM   5185  CA  VAL   110     -47.952  23.126  24.759  1.00 30.60      L    C
ATOM   5186  CB  VAL   110     -47.589  21.639  24.843  1.00 31.58      L    C
ATOM   5187  CG1 VAL   110     -48.696  20.885  25.564  1.00 31.21      L    C
ATOM   5188  CG2 VAL   110     -47.391  21.068  23.437  1.00 29.51      L    C
ATOM   5189  C   VAL   110     -48.247  23.642  26.163  1.00 32.83      L    C
ATOM   5190  O   VAL   110     -47.389  23.608  27.041  1.00 32.80      L    O
ATOM   5191  N   LEU   111     -49.470  24.119  26.364  1.00 34.22      L    N
ATOM   5192  CA  LEU   111     -49.831  24.834  27.581  1.00 34.04      L    C
ATOM   5193  CB  LEU   111     -50.785  25.985  27.247  1.00 34.30      L    C
ATOM   5194  CG  LEU   111     -50.233  27.051  26.294  1.00 39.19      L    C
ATOM   5195  CD1 LEU   111     -51.313  28.073  25.964  1.00 37.22      L    C
ATOM   5196  CD2 LEU   111     -49.029  27.728  26.934  1.00 38.67      L    C
ATOM   5197  C   LEU   111     -50.492  23.919  28.598  1.00 34.06      L    C
ATOM   5198  O   LEU   111     -50.631  22.719  28.376  1.00 34.37      L    O
ATOM   5199  N   GLY   112     -50.893  24.498  29.724  1.00 33.66      L    N
ATOM   5200  CA  GLY   112     -51.772  23.797  30.632  1.00 32.43      L    C
ATOM   5201  C   GLY   112     -51.139  23.268  31.903  1.00 32.20      L    C
ATOM   5202  O   GLY   112     -51.853  22.922  32.838  1.00 32.07      L    O
ATOM   5203  N   GLN   113     -49.815  23.199  31.965  1.00 31.98      L    N
ATOM   5204  CA  GLN   113     -49.186  22.612  33.145  1.00 33.88      L    C
ATOM   5205  CB  GLN   113     -47.725  22.251  32.853  1.00 34.74      L    C
ATOM   5206  CG  GLN   113     -46.732  23.380  32.993  1.00 38.41      L    C
ATOM   5207  CD  GLN   113     -45.326  22.922  32.653  1.00 42.73      L    C
ATOM   5208  OE1 GLN   113     -44.513  22.659  33.543  1.00 42.57      L    O
ATOM   5209  NE2 GLN   113     -45.034  22.811  31.354  1.00 42.20      L    N
ATOM   5210  C   GLN   113     -49.282  23.548  34.354  1.00 32.67      L    C
ATOM   5211  O   GLN   113     -49.377  24.770  34.208  1.00 30.96      L    O
ATOM   5212  N   PRO   114     -49.282  22.981  35.568  1.00 32.80      L    N
ATOM   5213  CD  PRO   114     -49.223  21.540  35.868  1.00 33.24      L    C
ATOM   5214  CA  PRO   114     -49.519  23.784  36.777  1.00 32.36      L    C
ATOM   5215  CB  PRO   114     -49.539  22.748  37.902  1.00 32.38      L    C
ATOM   5216  CG  PRO   114     -49.909  21.457  37.213  1.00 34.37      L    C
ATOM   5217  C   PRO   114     -48.458  24.863  37.007  1.00 30.82      L    C
ATOM   5218  O   PRO   114     -47.285  24.686  36.677  1.00 26.86      L    O
ATOM   5219  N   LYS   115     -48.874  25.989  37.569  1.00 31.80      L    N
ATOM   5220  CA  LYS   115     -47.916  27.022  37.930  1.00 34.33      L    C
ATOM   5221  CB  LYS   115     -48.634  28.202  38.583  1.00 36.99      L    C
ATOM   5222  CG  LYS   115     -47.748  29.426  38.753  1.00 44.60      L    C
ATOM   5223  CD  LYS   115     -47.982  30.122  40.084  1.00 48.66      L    C
ATOM   5224  CE  LYS   115     -49.028  31.218  39.966  1.00 50.81      L    C
ATOM   5225  NZ  LYS   115     -48.997  32.126  41.157  1.00 52.88      L    N
ATOM   5226  C   LYS   115     -46.879  26.437  38.897  1.00 33.33      L    C
ATOM   5227  O   LYS   115     -47.206  25.588  39.736  1.00 30.94      L    O
ATOM   5228  N   ALA   116     -45.630  26.877  38.762  1.00 32.81      L    N
ATOM   5229  CA  ALA   116     -44.570  26.495  39.697  1.00 33.88      L    C
ATOM   5230  CB  ALA   116     -43.738  25.366  39.111  1.00 30.45      L    C
ATOM   5231  C   ALA   116     -43.678  27.697  40.012  1.00 33.75      L    C
ATOM   5232  O   ALA   116     -43.179  28.368  39.105  1.00 33.41      L    O
ATOM   5233  N   ALA   117     -43.487  27.969  41.300  1.00 34.49      L    N
ATOM   5234  CA  ALA   117     -42.664  29.097  41.733  1.00 34.10      L    C
ATOM   5235  CB  ALA   117     -43.008  29.480  43.181  1.00 34.76      L    C
ATOM   5236  C   ALA   117     -41.188  28.724  41.622  1.00 34.26      L    C
ATOM   5237  O   ALA   117     -40.818  27.561  41.781  1.00 34.36      L    O
ATOM   5238  N   PRO   118     -40.329  29.712  41.331  1.00 34.24      L    N
ATOM   5239  CD  PRO   118     -40.679  31.132  41.151  1.00 33.58      L    C
ATOM   5240  CA  PRO   118     -38.902  29.463  41.114  1.00 35.01      L    C
ATOM   5241  CB  PRO   118     -38.394  30.772  40.517  1.00 34.92      L    C
ATOM   5242  CG  PRO   118     -39.339  31.811  41.039  1.00 34.58      L    C
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ATOM	5243	C	PRO	118	-38.172	29.108	42.403	1.00	37.33	L	C
ATOM	5244	O	PRO	118	-38.483	29.643	43.467	1.00	37.79	L	O
ATOM	5245	N	SER	119	-37.212	28.192	42.295	1.00	37.11	L	N
ATOM	5246	CA	SER	119	-36.175	28.046	43.304	1.00	37.74	L	C
ATOM	5247	CB	SER	119	-35.621	26.622	43.312	1.00	36.71	L	C
ATOM	5248	OG	SER	119	-36.607	25.707	43.742	1.00	45.31	L	O
ATOM	5249	C	SER	119	-35.055	29.006	42.951	1.00	37.01	L	C
ATOM	5250	O	SER	119	-34.615	29.064	41.799	1.00	38.07	L	O
ATOM	5251	N	VAL	120	-34.597	29.756	43.944	1.00	35.51	L	N
ATOM	5252	CA	VAL	120	-33.478	30.666	43.758	1.00	35.22	L	C
ATOM	5253	CB	VAL	120	-33.889	32.126	44.059	1.00	34.18	L	C
ATOM	5254	CG1	VAL	120	-32.680	33.048	43.933	1.00	30.75	L	C
ATOM	5255	CG2	VAL	120	-34.995	32.558	43.103	1.00	33.05	L	C
ATOM	5256	C	VAL	120	-32.338	30.278	44.684	1.00	36.13	L	C
ATOM	5257	O	VAL	120	-32.543	30.053	45.879	1.00	35.93	L	O
ATOM	5258	N	THR	121	-31.134	30.187	44.134	1.00	35.67	L	N
ATOM	5259	CA	THR	121	-29.961	30.097	44.975	1.00	36.07	L	C
ATOM	5260	CB	THR	121	-29.413	28.637	45.022	1.00	38.17	L	C
ATOM	5261	OG1	THR	121	-28.028	28.622	44.670	1.00	43.48	L	O
ATOM	5262	CG2	THR	121	-30.191	27.741	44.092	1.00	38.07	L	C
ATOM	5263	C	THR	121	-28.900	31.093	44.518	1.00	35.24	L	C
ATOM	5264	O	THR	121	-28.695	31.310	43.321	1.00	33.07	L	O
ATOM	5265	N	LEU	122	-28.253	31.728	45.490	1.00	34.06	L	N
ATOM	5266	CA	LEU	122	-27.357	32.847	45.220	1.00	34.60	L	C
ATOM	5267	CB	LEU	122	-27.971	34.136	45.762	1.00	31.62	L	C
ATOM	5268	CG	LEU	122	-27.093	35.388	45.764	1.00	33.67	L	C
ATOM	5269	CD1	LEU	122	-26.790	35.813	44.337	1.00	31.18	L	C
ATOM	5270	CD2	LEU	122	-27.815	36.508	46.519	1.00	34.33	L	C
ATOM	5271	C	LEU	122	-25.979	32.625	45.852	1.00	35.18	L	C
ATOM	5272	O	LEU	122	-25.873	32.339	47.048	1.00	35.07	L	O
ATOM	5273	N	PHE	123	-24.934	32.756	45.040	1.00	34.10	L	N
ATOM	5274	CA	PHE	123	-23.563	32.581	45.503	1.00	33.60	L	C
ATOM	5275	CB	PHE	123	-22.803	31.613	44.598	1.00	32.81	L	C
ATOM	5276	CG	PHE	123	-23.292	30.200	44.673	1.00	33.31	L	C
ATOM	5277	CD1	PHE	123	-24.108	29.683	43.680	1.00	31.54	L	C
ATOM	5278	CD2	PHE	123	-22.919	29.380	45.726	1.00	31.90	L	C
ATOM	5279	CE1	PHE	123	-24.545	28.373	43.731	1.00	32.77	L	C
ATOM	5280	CE2	PHE	123	-23.351	28.066	45.783	1.00	31.56	L	C
ATOM	5281	CZ	PHE	123	-24.165	27.562	44.784	1.00	32.86	L	C
ATOM	5282	C	PHE	123	-22.822	33.902	45.507	1.00	34.23	L	C
ATOM	5283	O	PHE	123	-22.887	34.660	44.541	1.00	33.21	L	O
ATOM	5284	N	PRO	124	-22.085	34.183	46.595	1.00	34.60	L	N
ATOM	5285	CD	PRO	124	-22.022	33.339	47.802	1.00	33.97	L	C
ATOM	5286	CA	PRO	124	-21.179	35.334	46.681	1.00	33.22	L	C
ATOM	5287	CB	PRO	124	-20.827	35.396	48.162	1.00	32.92	L	C
ATOM	5288	CG	PRO	124	-20.911	33.971	48.613	1.00	32.54	L	C
ATOM	5289	C	PRO	124	-19.952	35.072	45.815	1.00	33.35	L	C
ATOM	5290	O	PRO	124	-19.764	33.962	45.318	1.00	32.14	L	O
ATOM	5291	N	PRO	125	-19.100	36.089	45.624	1.00	34.18	L	N
ATOM	5292	CD	PRO	125	-19.242	37.482	46.084	1.00	34.13	L	C
ATOM	5293	CA	PRO	125	-17.823	35.865	44.937	1.00	33.64	L	C
ATOM	5294	CB	PRO	125	-17.180	37.253	44.894	1.00	34.39	L	C
ATOM	5295	CG	PRO	125	-18.291	38.220	45.187	1.00	35.53	L	C
ATOM	5296	C	PRO	125	-16.961	34.878	45.732	1.00	35.03	L	C
ATOM	5297	O	PRO	125	-16.917	34.937	46.962	1.00	35.09	L	O
ATOM	5298	N	SER	126	-16.281	33.973	45.035	1.00	34.42	L	N
ATOM	5299	CA	SER	126	-15.335	33.076	45.691	1.00	34.25	L	C
ATOM	5300	CB	SER	126	-14.937	31.950	44.744	1.00	32.73	L	C
ATOM	5301	OG	SER	126	-14.250	32.467	43.621	1.00	32.43	L	O
ATOM	5302	C	SER	126	-14.093	33.870	46.084	1.00	35.04	L	C
ATOM	5303	O	SER	126	-13.777	34.884	45.461	1.00	33.38	L	O
ATOM	5304	N	SER	127	-13.383	33.412	47.110	1.00	35.77	L	N
ATOM	5305	CA	SER	127	-12.176	34.116	47.534	1.00	37.95	L	C
ATOM	5306	CB	SER	127	-11.653	33.549	48.864	1.00	39.02	L	C
ATOM	5307	OG	SER	127	-11.268	32.195	48.731	1.00	45.05	L	O
ATOM	5308	C	SER	127	-11.100	34.022	46.452	1.00	36.52	L	C
ATOM	5309	O	SER	127	-10.301	34.941	46.276	1.00	35.94	L	O
ATOM	5310	N	GLU	128	-11.099	32.928	45.703	1.00	36.21	L	N
ATOM	5311	CA	GLU	128	-10.172	32.797	44.590	1.00	38.71	L	C
ATOM	5312	CB	GLU	128	-10.288	31.401	43.975	1.00	40.30	L	C
ATOM	5313	CG	GLU	128	-9.137	31.038	43.054	1.00	44.77	L	C
ATOM	5314	CD	GLU	128	-9.217	29.605	42.538	1.00	48.56	L	C
ATOM	5315	OE1	GLU	128	-8.448	29.266	41.610	1.00	48.39	L	O
ATOM	5316	OE2	GLU	128	-10.041	28.817	43.061	1.00	49.98	L	O
ATOM	5317	C	GLU	128	-10.396	33.874	43.514	1.00	40.21	L	C
ATOM	5318	O	GLU	128	-9.436	34.479	43.025	1.00	39.54	L	O

ATOM	5319	N	GLU	129	-11.652	34.127	43.147	1.00	40.30	L	N
ATOM	5320	CA	GLU	129	-11.927	35.156	42.142	1.00	38.87	L	C
ATOM	5321	CB	GLU	129	-13.406	35.152	41.720	1.00	39.30	L	C
ATOM	5322	CG	GLU	129	-13.717	36.210	40.653	1.00	40.41	L	C
ATOM	5323	CD	GLU	129	-15.192	36.289	40.261	1.00	42.41	L	C
ATOM	5324	OE1	GLU	129	-15.469	36.599	39.081	1.00	44.06	L	O
ATOM	5325	OE2	GLU	129	-16.072	36.055	41.117	1.00	40.48	L	O
ATOM	5326	C	GLU	129	-11.560	36.538	42.681	1.00	38.41	L	C
ATOM	5327	O	GLU	129	-11.107	37.405	41.933	1.00	36.76	L	O
ATOM	5328	N	LEU	130	-11.759	36.746	43.979	1.00	37.71	L	N
ATOM	5329	CA	LEU	130	-11.361	38.003	44.599	1.00	38.42	L	C
ATOM	5330	CB	LEU	130	-11.837	38.042	46.053	1.00	36.50	L	C
ATOM	5331	CG	LEU	130	-13.358	38.169	46.237	1.00	36.84	L	C
ATOM	5332	CD1	LEU	130	-13.723	38.067	47.713	1.00	34.95	L	C
ATOM	5333	CD2	LEU	130	-13.832	39.497	45.657	1.00	33.05	L	C
ATOM	5334	C	LEU	130	-9.844	38.231	44.515	1.00	40.34	L	C
ATOM	5335	O	LEU	130	-9.401	39.348	44.243	1.00	40.94	L	O
ATOM	5336	N	GLN	131	-9.056	37.176	44.730	1.00	41.55	L	N
ATOM	5337	CA	GLN	131	-7.606	37.236	44.532	1.00	43.80	L	C
ATOM	5338	CB	GLN	131	-6.950	35.883	44.844	1.00	43.76	L	C
ATOM	5339	CG	GLN	131	-6.260	35.816	46.191	1.00	47.33	L	C
ATOM	5340	CD	GLN	131	-5.469	37.072	46.514	1.00	47.15	L	C
ATOM	5341	OE1	GLN	131	-5.756	37.757	47.497	1.00	49.05	L	O
ATOM	5342	NE2	GLN	131	-4.470	37.382	45.692	1.00	45.39	L	N
ATOM	5343	C	GLN	131	-7.248	37.626	43.105	1.00	45.27	L	C
ATOM	5344	O	GLN	131	-6.261	38.333	42.876	1.00	46.89	L	O
ATOM	5345	N	ALA	132	-8.039	37.146	42.147	1.00	45.27	L	N
ATOM	5346	CA	ALA	132	-7.868	37.521	40.743	1.00	43.98	L	C
ATOM	5347	CB	ALA	132	-8.552	36.497	39.841	1.00	43.38	L	C
ATOM	5348	C	ALA	132	-8.435	38.915	40.474	1.00	43.57	L	C
ATOM	5349	O	ALA	132	-8.516	39.351	39.327	1.00	42.33	L	O
ATOM	5350	N	ASN	133	-8.845	39.602	41.537	1.00	43.82	L	N
ATOM	5351	CA	ASN	133	-9.273	40.996	41.433	1.00	46.95	L	C
ATOM	5352	CB	ASN	133	-8.155	41.830	40.792	1.00	48.35	L	C
ATOM	5353	CG	ASN	133	-8.282	43.309	41.101	1.00	50.92	L	C
ATOM	5354	OD1	ASN	133	-8.791	43.694	42.158	1.00	51.72	L	O
ATOM	5355	ND2	ASN	133	-7.821	44.150	40.179	1.00	51.45	L	N
ATOM	5356	C	ASN	133	-10.581	41.179	40.642	1.00	46.81	L	C
ATOM	5357	O	ASN	133	-10.751	42.171	39.929	1.00	47.08	L	O
ATOM	5358	N	LYS	134	-11.494	40.218	40.768	1.00	46.18	L	N
ATOM	5359	CA	LYS	134	-12.830	40.324	40.186	1.00	45.08	L	C
ATOM	5360	CB	LYS	134	-12.955	39.437	38.946	1.00	46.32	L	C
ATOM	5361	CG	LYS	134	-11.819	39.552	37.947	1.00	49.14	L	C
ATOM	5362	CD	LYS	134	-11.935	40.811	37.107	1.00	55.33	L	C
ATOM	5363	CE	LYS	134	-11.460	40.559	35.673	1.00	58.41	L	C
ATOM	5364	NZ	LYS	134	-10.414	39.490	35.603	1.00	59.23	L	N
ATOM	5365	C	LYS	134	-13.854	39.864	41.218	1.00	44.89	L	C
ATOM	5366	O	LYS	134	-13.510	39.179	42.181	1.00	45.02	L	O
ATOM	5367	N	ALA	135	-15.112	40.237	41.013	1.00	43.12	L	N
ATOM	5368	CA	ALA	135	-16.199	39.733	41.845	1.00	42.34	L	C
ATOM	5369	CB	ALA	135	-16.485	40.710	42.986	1.00	40.89	L	C
ATOM	5370	C	ALA	135	-17.465	39.497	41.019	1.00	41.64	L	C
ATOM	5371	O	ALA	135	-17.928	40.383	40.299	1.00	42.40	L	O
ATOM	5372	N	THR	136	-18.019	38.295	41.129	1.00	41.17	L	N
ATOM	5373	CA	THR	136	-19.223	37.928	40.392	1.00	36.95	L	C
ATOM	5374	CB	THR	136	-18.910	36.884	39.301	1.00	36.17	L	C
ATOM	5375	OG1	THR	136	-17.877	37.385	38.448	1.00	34.77	L	O
ATOM	5376	CG2	THR	136	-20.149	36.592	38.460	1.00	35.99	L	C
ATOM	5377	C	THR	136	-20.234	37.326	41.349	1.00	35.86	L	C
ATOM	5378	O	THR	136	-19.952	36.323	42.003	1.00	35.40	L	O
ATOM	5379	N	LEU	137	-21.410	37.940	41.434	1.00	35.37	L	N
ATOM	5380	CA	LEU	137	-22.527	37.333	42.140	1.00	34.20	L	C
ATOM	5381	CB	LEU	137	-23.417	38.412	42.758	1.00	36.75	L	C
ATOM	5382	CG	LEU	137	-22.746	39.314	43.798	1.00	40.79	L	C
ATOM	5383	CD1	LEU	137	-23.790	40.073	44.597	1.00	41.82	L	C
ATOM	5384	CD2	LEU	137	-21.929	38.465	44.725	1.00	42.69	L	C
ATOM	5385	C	LEU	137	-23.332	36.481	41.160	1.00	34.75	L	C
ATOM	5386	O	LEU	137	-23.562	36.877	40.013	1.00	33.76	L	O
ATOM	5387	N	VAL	138	-23.748	35.305	41.616	1.00	33.20	L	N
ATOM	5388	CA	VAL	138	-24.366	34.322	40.741	1.00	33.86	L	C
ATOM	5389	CB	VAL	138	-23.496	33.058	40.627	1.00	33.39	L	C
ATOM	5390	CG1	VAL	138	-24.158	32.065	39.679	1.00	31.94	L	C
ATOM	5391	CG2	VAL	138	-22.096	33.436	40.155	1.00	28.69	L	C
ATOM	5392	C	VAL	138	-25.736	33.916	41.261	1.00	34.92	L	C
ATOM	5393	O	VAL	138	-25.851	33.261	42.303	1.00	35.11	L	O
ATOM	5394	N	CYS	139	-26.772	34.310	40.528	1.00	34.33	L	N

ATOM	5395	CA	CYS	139	-28.145	34.048	40.937	1.00	34.32	L	C
ATOM	5396	C	CYS	139	-28.755	33.014	40.009	1.00	32.61	L	C
ATOM	5397	O	CYS	139	-28.914	33.263	38.816	1.00	32.03	L	O
ATOM	5398	CB	CYS	139	-28.963	35.330	40.868	1.00	36.04	L	C
ATOM	5399	SG	CYS	139	-30.606	35.207	41.630	1.00	39.55	L	S
ATOM	5400	N	LEU	140	-29.092	31.856	40.559	1.00	30.81	L	N
ATOM	5401	CA	LEU	140	-29.656	30.747	39.745	1.00	32.77	L	C
ATOM	5402	CB	LEU	140	-28.722	29.503	40.041	1.00	32.67	L	C
ATOM	5403	CG	LEU	140	-27.225	29.678	39.777	1.00	33.78	L	C
ATOM	5404	CD1	LEU	140	-26.500	28.410	40.172	1.00	31.83	L	C
ATOM	5405	CD2	LEU	140	-26.981	29.995	38.302	1.00	30.83	L	C
ATOM	5406	C	LEU	140	-31.024	30.472	40.018	1.00	33.03	L	C
ATOM	5407	O	LEU	140	-31.432	30.317	41.169	1.00	32.85	L	O
ATOM	5408	N	ILE	141	-31.808	30.409	38.946	1.00	33.23	L	N
ATOM	5409	CA	ILE	141	-33.266	30.360	39.040	1.00	33.94	L	C
ATOM	5410	CB	ILE	141	-33.873	31.630	38.419	1.00	33.58	L	C
ATOM	5411	CG2	ILE	141	-35.333	31.765	38.822	1.00	34.21	L	C
ATOM	5412	CG1	ILE	141	-33.095	32.858	38.899	1.00	32.33	L	C
ATOM	5413	CD1	ILE	141	-33.396	34.114	38.106	1.00	32.35	L	C
ATOM	5414	C	ILE	141	-33.784	29.143	38.277	1.00	33.63	L	C
ATOM	5415	O	ILE	141	-33.506	28.990	37.088	1.00	36.23	L	O
ATOM	5416	N	SER	142	-34.533	28.276	38.947	1.00	32.80	L	N
ATOM	5417	CA	SER	142	-34.925	27.014	38.327	1.00	33.50	L	C
ATOM	5418	CB	SER	142	-33.897	25.918	38.651	1.00	34.71	L	C
ATOM	5419	OG	SER	142	-33.838	25.659	40.040	1.00	39.34	L	O
ATOM	5420	C	SER	142	-36.315	26.537	38.720	1.00	33.10	L	C
ATOM	5421	O	SER	142	-36.918	27.054	39.664	1.00	31.05	L	O
ATOM	5422	N	ASP	143	-36.820	25.562	37.963	1.00	33.73	L	N
ATOM	5423	CA	ASP	143	-38.075	24.877	38.266	1.00	35.27	L	C
ATOM	5424	CB	ASP	143	-37.980	24.159	39.613	1.00	37.99	L	C
ATOM	5425	CG	ASP	143	-36.947	23.050	39.606	1.00	46.04	L	C
ATOM	5426	OD1	ASP	143	-36.195	22.932	40.601	1.00	49.81	L	O
ATOM	5427	OD2	ASP	143	-36.886	22.298	38.604	1.00	46.96	L	O
ATOM	5428	C	ASP	143	-39.301	25.776	38.277	1.00	35.39	L	C
ATOM	5429	O	ASP	143	-40.216	25.564	39.075	1.00	35.57	L	O
ATOM	5430	N	PHE	144	-39.340	26.778	37.408	1.00	32.40	L	N
ATOM	5431	CA	PHE	144	-40.513	27.630	37.392	1.00	33.33	L	C
ATOM	5432	CB	PHE	144	-40.120	29.101	37.632	1.00	32.73	L	C
ATOM	5433	CG	PHE	144	-39.148	29.667	36.624	1.00	32.95	L	C
ATOM	5434	CD1	PHE	144	-39.606	30.394	35.529	1.00	32.15	L	C
ATOM	5435	CD2	PHE	144	-37.776	29.554	36.820	1.00	32.00	L	C
ATOM	5436	CE1	PHE	144	-38.712	31.008	34.649	1.00	32.86	L	C
ATOM	5437	CE2	PHE	144	-36.874	30.164	35.946	1.00	31.42	L	C
ATOM	5438	CZ	PHE	144	-37.345	30.894	34.859	1.00	31.87	L	C
ATOM	5439	C	PHE	144	-41.360	27.491	36.126	1.00	33.15	L	C
ATOM	5440	O	PHE	144	-40.859	27.152	35.051	1.00	31.11	L	O
ATOM	5441	N	TYR	145	-42.658	27.728	36.277	1.00	32.99	L	N
ATOM	5442	CA	TYR	145	-43.570	27.759	35.146	1.00	35.64	L	C
ATOM	5443	CB	TYR	145	-44.099	26.353	34.814	1.00	35.95	L	C
ATOM	5444	CG	TYR	145	-44.958	26.360	33.567	1.00	40.08	L	C
ATOM	5445	CD1	TYR	145	-46.301	26.743	33.623	1.00	40.50	L	C
ATOM	5446	CE1	TYR	145	-47.053	26.893	32.471	1.00	42.12	L	C
ATOM	5447	CD2	TYR	145	-44.399	26.112	32.316	1.00	39.32	L	C
ATOM	5448	CE2	TYR	145	-45.144	26.258	31.158	1.00	41.02	L	C
ATOM	5449	CZ	TYR	145	-46.466	26.653	31.240	1.00	42.27	L	C
ATOM	5450	OH	TYR	145	-47.196	26.833	30.089	1.00	43.82	L	O
ATOM	5451	C	TYR	145	-44.735	28.661	35.499	1.00	35.38	L	C
ATOM	5452	O	TYR	145	-45.261	28.586	36.609	1.00	37.77	L	O
ATOM	5453	N	PRO	146	-45.168	29.521	34.562	1.00	35.04	L	N
ATOM	5454	CD	PRO	146	-46.411	30.290	34.759	1.00	34.49	L	C
ATOM	5455	CA	PRO	146	-44.607	29.714	33.217	1.00	36.14	L	C
ATOM	5456	CB	PRO	146	-45.686	30.514	32.480	1.00	33.39	L	C
ATOM	5457	CG	PRO	146	-46.458	31.194	33.550	1.00	35.09	L	C
ATOM	5458	C	PRO	146	-43.247	30.417	33.179	1.00	38.56	L	C
ATOM	5459	O	PRO	146	-42.752	30.910	34.200	1.00	38.92	L	O
ATOM	5460	N	GLY	147	-42.668	30.477	31.981	1.00	38.91	L	N
ATOM	5461	CA	GLY	147	-41.256	30.772	31.843	1.00	40.73	L	C
ATOM	5462	C	GLY	147	-40.878	32.226	31.670	1.00	41.71	L	C
ATOM	5463	O	GLY	147	-39.990	32.542	30.886	1.00	45.13	L	O
ATOM	5464	N	ALA	148	-41.533	33.117	32.399	1.00	41.10	L	N
ATOM	5465	CA	ALA	148	-41.119	34.510	32.410	1.00	42.96	L	C
ATOM	5466	CB	ALA	148	-42.205	35.384	31.801	1.00	42.15	L	C
ATOM	5467	C	ALA	148	-40.826	34.963	33.838	1.00	43.29	L	C
ATOM	5468	O	ALA	148	-41.658	34.807	34.730	1.00	43.64	L	O
ATOM	5469	N	VAL	149	-39.638	35.517	34.047	1.00	42.89	L	N
ATOM	5470	CA	VAL	149	-39.300	36.155	35.315	1.00	43.55	L	C

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ATOM   5471  CB  VAL   149     -38.333  35.293  36.159  1.00 42.09      L    C
ATOM   5472  CG1 VAL   149     -39.040  34.062  36.679  1.00 40.84      L    C
ATOM   5473  CG2 VAL   149     -37.126  34.903  35.318  1.00 40.82      L    C
ATOM   5474  C   VAL   149     -38.606  37.471  35.027  1.00 44.09      L    C
ATOM   5475  O   VAL   149     -38.039  37.663  33.951  1.00 44.44      L    O
ATOM   5476  N   THR   150     -38.651  38.379  35.990  1.00 44.32      L    N
ATOM   5477  CA  THR   150     -37.788  39.546  35.947  1.00 45.76      L    C
ATOM   5478  CB  THR   150     -38.603  40.854  35.969  1.00 47.68      L    C
ATOM   5479  OG1 THR   150     -39.451  40.866  37.124  1.00 50.74      L    O
ATOM   5480  CG2 THR   150     -39.455  40.971  34.710  1.00 48.08      L    C
ATOM   5481  C   THR   150     -36.871  39.509  37.159  1.00 44.03      L    C
ATOM   5482  O   THR   150     -37.286  39.113  38.251  1.00 42.08      L    O
ATOM   5483  N   VAL   151     -35.623  39.916  36.960  1.00 43.16      L    N
ATOM   5484  CA  VAL   151     -34.645  39.902  38.034  1.00 43.36      L    C
ATOM   5485  CB  VAL   151     -33.424  39.035  37.659  1.00 43.35      L    C
ATOM   5486  CG1 VAL   151     -32.524  38.850  38.880  1.00 42.41      L    C
ATOM   5487  CG2 VAL   151     -33.885  37.694  37.119  1.00 41.25      L    C
ATOM   5488  C   VAL   151     -34.162  41.315  38.343  1.00 43.86      L    C
ATOM   5489  O   VAL   151     -33.789  42.065  37.442  1.00 44.90      L    O
ATOM   5490  N   ALA   152     -34.164  41.671  39.623  1.00 44.46      L    N
ATOM   5491  CA  ALA   152     -33.621  42.953  40.067  1.00 46.25      L    C
ATOM   5492  CB  ALA   152     -34.750  43.846  40.590  1.00 44.32      L    C
ATOM   5493  C   ALA   152     -32.581  42.728  41.167  1.00 46.44      L    C
ATOM   5494  O   ALA   152     -32.771  41.881  42.041  1.00 45.37      L    O
ATOM   5495  N   TRP   153     -31.489  43.488  41.121  1.00 46.96      L    N
ATOM   5496  CA  TRP   153     -30.434  43.375  42.124  1.00 48.34      L    C
ATOM   5497  CB  TRP   153     -29.068  43.258  41.445  1.00 46.62      L    C
ATOM   5498  CG  TRP   153     -28.835  41.977  40.692  1.00 44.35      L    C
ATOM   5499  CD2 TRP   153     -28.153  40.809  41.174  1.00 43.29      L    C
ATOM   5500  CE2 TRP   153     -28.095  39.888  40.106  1.00 42.33      L    C
ATOM   5501  CE3 TRP   153     -27.584  40.454  42.403  1.00 41.90      L    C
ATOM   5502  CD1 TRP   153     -29.158  41.719  39.390  1.00 43.20      L    C
ATOM   5503  NE1 TRP   153     -28.714  40.467  39.030  1.00 41.75      L    N
ATOM   5504  CZ2 TRP   153     -27.487  38.636  40.231  1.00 41.82      L    C
ATOM   5505  CZ3 TRP   153     -26.982  39.210  42.525  1.00 40.30      L    C
ATOM   5506  CH2 TRP   153     -26.938  38.317  41.443  1.00 41.25      L    C
ATOM   5507  C   TRP   153     -30.425  44.586  43.055  1.00 50.76      L    C
ATOM   5508  O   TRP   153     -30.656  45.710  42.619  1.00 51.60      L    O
ATOM   5509  N   LYS   154     -30.158  44.349  44.337  1.00 53.56      L    N
ATOM   5510  CA  LYS   154     -30.028  45.430  45.311  1.00 55.91      L    C
ATOM   5511  CB  LYS   154     -31.100  45.307  46.400  1.00 57.60      L    C
ATOM   5512  CG  LYS   154     -32.529  45.384  45.893  1.00 61.06      L    C
ATOM   5513  CD  LYS   154     -32.750  46.616  45.026  1.00 64.51      L    C
ATOM   5514  CE  LYS   154     -34.157  46.635  44.439  1.00 65.49      L    C
ATOM   5515  NZ  LYS   154     -35.197  46.612  45.509  1.00 67.42      L    N
ATOM   5516  C   LYS   154     -28.649  45.439  45.974  1.00 56.74      L    C
ATOM   5517  O   LYS   154     -28.189  44.416  46.490  1.00 55.15      L    O
ATOM   5518  N   ALA   155     -27.998  46.599  45.952  1.00 57.64      L    N
ATOM   5519  CA  ALA   155     -26.644  46.854  46.809  1.00 59.43      L    C
ATOM   5520  CB  ALA   155     -25.912  47.859  46.147  1.00 58.00      L    C
ATOM   5521  C   ALA   155     -27.352  47.399  48.142  1.00 60.59      L    C
ATOM   5522  O   ALA   155     -27.797  48.546  48.228  1.00 60.44      L    O
ATOM   5523  N   ASP   156     -27.292  46.568  49.176  1.00 62.39      L    N
ATOM   5524  CA  ASP   156     -27.969  46.848  50.426  1.00 65.83      L    C
ATOM   5525  CB  ASP   156     -27.503  48.165  51.032  1.00 67.21      L    C
ATOM   5526  CG  ASP   156     -26.129  48.044  51.654  1.00 69.56      L    C
ATOM   5527  OD1 ASP   156     -25.959  47.193  52.554  1.00 70.11      L    O
ATOM   5528  OD2 ASP   156     -25.220  48.796  51.239  1.00 71.44      L    O
ATOM   5529  C   ASP   156     -29.492  46.918  50.195  1.00 67.34      L    C
ATOM   5530  O   ASP   156     -30.177  45.893  50.179  1.00 68.08      L    O
ATOM   5531  N   SER   157     -30.004  48.131  50.012  1.00 68.19      L    N
ATOM   5532  CA  SER   157     -31.422  48.321  49.739  1.00 69.31      L    C
ATOM   5533  CB  SER   157     -32.119  48.937  50.954  1.00 70.08      L    C
ATOM   5534  OG  SER   157     -32.192  48.008  52.025  1.00 71.57      L    O
ATOM   5535  C   SER   157     -31.650  49.199  48.513  1.00 69.35      L    C
ATOM   5536  O   SER   157     -32.765  49.654  48.265  1.00 69.73      L    O
ATOM   5537  N   SER   158     -30.591  49.433  47.747  1.00 68.83      L    N
ATOM   5538  CA  SER   158     -30.677  50.297  46.580  1.00 69.49      L    C
ATOM   5539  CB  SER   158     -29.621  51.395  46.679  1.00 70.91      L    C
ATOM   5540  OG  SER   158     -29.610  51.946  47.986  1.00 73.20      L    O
ATOM   5541  C   SER   158     -30.488  49.512  45.285  1.00 69.20      L    C
ATOM   5542  O   SER   158     -29.701  48.572  45.225  1.00 69.63      L    O
ATOM   5543  N   PRO   159     -31.215  49.896  44.229  1.00 68.97      L    N
ATOM   5544  CD  PRO   159     -32.254  50.939  44.259  1.00 69.29      L    C
ATOM   5545  CA  PRO   159     -31.157  49.225  42.926  1.00 69.40      L    C
ATOM   5546  CB  PRO   159     -32.184  49.980  42.083  1.00 68.16      L    C
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ATOM	5547	CG	PRO	159	-33.113	50.583	43.082	1.00	69.30	L	C
ATOM	5548	C	PRO	159	-29.771	49.269	42.294	1.00	68.38	L	C
ATOM	5549	O	PRO	159	-29.129	50.320	42.249	1.00	68.62	L	O
ATOM	5550	N	VAL	160	-29.317	48.120	41.804	1.00	67.89	L	N
ATOM	5551	CA	VAL	160	-28.076	48.042	41.047	1.00	67.89	L	C
ATOM	5552	CB	VAL	160	-27.268	46.788	41.426	1.00	67.44	L	C
ATOM	5553	CG1	VAL	160	-25.984	46.736	40.614	1.00	66.24	L	C
ATOM	5554	CG2	VAL	160	-26.970	46.790	42.917	1.00	66.87	L	C
ATOM	5555	C	VAL	160	-28.415	47.962	39.566	1.00	68.75	L	C
ATOM	5556	O	VAL	160	-29.066	47.019	39.128	1.00	70.43	L	O
ATOM	5557	N	LYS	161	-27.968	48.945	38.796	1.00	68.73	L	N
ATOM	5558	CA	LYS	161	-28.304	48.998	37.378	1.00	68.60	L	C
ATOM	5559	CB	LYS	161	-28.436	50.458	36.927	1.00	73.13	L	C
ATOM	5560	CG	LYS	161	-29.469	51.270	37.711	1.00	77.34	L	C
ATOM	5561	CD	LYS	161	-30.885	50.731	37.505	1.00	80.36	L	C
ATOM	5562	CE	LYS	161	-31.912	51.540	38.294	1.00	82.15	L	C
ATOM	5563	NZ	LYS	161	-33.299	51.001	38.144	1.00	82.95	L	N
ATOM	5564	C	LYS	161	-27.254	48.291	36.523	1.00	65.92	L	C
ATOM	5565	O	LYS	161	-27.562	47.346	35.787	1.00	65.61	L	O
ATOM	5566	N	ALA	162	-26.012	48.754	36.634	1.00	61.33	L	N
ATOM	5567	CA	ALA	162	-24.935	48.296	35.763	1.00	57.37	L	C
ATOM	5568	CB	ALA	162	-23.839	49.356	35.699	1.00	57.49	L	C
ATOM	5569	C	ALA	162	-24.344	46.963	36.219	1.00	53.70	L	C
ATOM	5570	O	ALA	162	-24.505	46.561	37.368	1.00	51.79	L	O
ATOM	5571	N	GLY	163	-23.663	46.283	35.303	1.00	50.45	L	N
ATOM	5572	CA	GLY	163	-22.992	45.044	35.646	1.00	49.48	L	C
ATOM	5573	C	GLY	163	-23.885	43.813	35.674	1.00	48.47	L	C
ATOM	5574	O	GLY	163	-23.441	42.737	36.078	1.00	49.78	L	O
ATOM	5575	N	VAL	164	-25.137	43.960	35.252	1.00	44.89	L	N
ATOM	5576	CA	VAL	164	-26.079	42.849	35.282	1.00	42.87	L	C
ATOM	5577	CB	VAL	164	-27.466	43.304	35.757	1.00	41.38	L	C
ATOM	5578	CG1	VAL	164	-28.429	42.123	35.757	1.00	40.11	L	C
ATOM	5579	CG2	VAL	164	-27.365	43.909	37.146	1.00	40.00	L	C
ATOM	5580	C	VAL	164	-26.237	42.197	33.920	1.00	42.51	L	C
ATOM	5581	O	VAL	164	-26.474	42.878	32.926	1.00	43.92	L	O
ATOM	5582	N	GLU	165	-26.106	40.875	33.874	1.00	41.72	L	N
ATOM	5583	CA	GLU	165	-26.428	40.126	32.664	1.00	43.48	L	C
ATOM	5584	CB	GLU	165	-25.143	39.769	31.911	1.00	45.09	L	C
ATOM	5585	CG	GLU	165	-24.375	41.016	31.483	1.00	50.70	L	C
ATOM	5586	CD	GLU	165	-23.153	40.719	30.637	1.00	54.19	L	C
ATOM	5587	OE1	GLU	165	-23.033	41.317	29.542	1.00	54.85	L	O
ATOM	5588	OE2	GLU	165	-22.312	39.899	31.070	1.00	55.72	L	O
ATOM	5589	C	GLU	165	-27.238	38.873	32.984	1.00	42.87	L	C
ATOM	5590	O	GLU	165	-26.797	38.015	33.752	1.00	42.63	L	O
ATOM	5591	N	THR	166	-28.430	38.792	32.394	1.00	41.04	L	N
ATOM	5592	CA	THR	166	-29.392	37.732	32.681	1.00	40.42	L	C
ATOM	5593	CB	THR	166	-30.700	38.326	33.233	1.00	40.61	L	C
ATOM	5594	OG1	THR	166	-30.414	39.093	34.406	1.00	41.13	L	O
ATOM	5595	CG2	THR	166	-31.691	37.226	33.580	1.00	40.33	L	C
ATOM	5596	C	THR	166	-29.719	36.945	31.415	1.00	40.37	L	C
ATOM	5597	O	THR	166	-29.941	37.532	30.360	1.00	42.10	L	O
ATOM	5598	N	THR	167	-29.749	35.620	31.513	1.00	41.03	L	N
ATOM	5599	CA	THR	167	-30.116	34.790	30.365	1.00	40.58	L	C
ATOM	5600	CB	THR	167	-29.722	33.308	30.567	1.00	39.43	L	C
ATOM	5601	OG1	THR	167	-30.455	32.766	31.671	1.00	39.95	L	O
ATOM	5602	CG2	THR	167	-28.233	33.176	30.832	1.00	38.27	L	C
ATOM	5603	C	THR	167	-31.627	34.845	30.149	1.00	41.37	L	C
ATOM	5604	O	THR	167	-32.373	35.320	31.010	1.00	41.43	L	O
ATOM	5605	N	THR	168	-32.078	34.363	28.996	1.00	41.45	L	N
ATOM	5606	CA	THR	168	-33.509	34.207	28.760	1.00	42.31	L	C
ATOM	5607	CB	THR	168	-33.852	34.322	27.256	1.00	44.42	L	C
ATOM	5608	OG1	THR	168	-33.082	33.364	26.522	1.00	48.18	L	O
ATOM	5609	CG2	THR	168	-33.535	35.723	26.732	1.00	44.90	L	C
ATOM	5610	C	THR	168	-33.922	32.831	29.259	1.00	40.42	L	C
ATOM	5611	O	THR	168	-33.178	31.860	29.112	1.00	40.27	L	O
ATOM	5612	N	PRO	169	-35.116	32.729	29.858	1.00	40.18	L	N
ATOM	5613	CD	PRO	169	-36.045	33.834	30.151	1.00	40.79	L	C
ATOM	5614	CA	PRO	169	-35.595	31.453	30.400	1.00	41.42	L	C
ATOM	5615	CB	PRO	169	-37.008	31.776	30.879	1.00	39.84	L	C
ATOM	5616	CG	PRO	169	-36.956	33.241	31.200	1.00	41.04	L	C
ATOM	5617	C	PRO	169	-35.571	30.338	29.364	1.00	42.24	L	C
ATOM	5618	O	PRO	169	-35.869	30.553	28.193	1.00	45.17	L	O
ATOM	5619	N	SER	170	-35.207	29.145	29.805	1.00	42.33	L	N
ATOM	5620	CA	SER	170	-35.041	28.016	28.907	1.00	43.07	L	C
ATOM	5621	CB	SER	170	-33.561	27.872	28.562	1.00	42.97	L	C
ATOM	5622	OG	SER	170	-33.297	26.630	27.950	1.00	50.77	L	O

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ATOM   5623  C   SER   170     -35.566  26.746  29.582  1.00 43.51      L    C
ATOM   5624  O   SER   170     -35.396  26.564  30.790  1.00 43.82      L    O
ATOM   5625  N   LYS   171     -36.211  25.876  28.809  1.00 43.46      L    N
ATOM   5626  CA  LYS   171     -36.836  24.676  29.359  1.00 44.64      L    C
ATOM   5627  CB  LYS   171     -37.741  24.007  28.317  1.00 46.55      L    C
ATOM   5628  CG  LYS   171     -39.099  24.659  28.140  1.00 51.02      L    C
ATOM   5629  CD  LYS   171     -40.179  23.923  28.923  1.00 52.77      L    C
ATOM   5630  CE  LYS   171     -41.558  24.174  28.315  1.00 54.67      L    C
ATOM   5631  NZ  LYS   171     -42.624  23.309  28.902  1.00 54.14      L    N
ATOM   5632  C   LYS   171     -35.805  23.664  29.833  1.00 44.02      L    C
ATOM   5633  O   LYS   171     -34.858  23.350  29.119  1.00 43.67      L    O
ATOM   5634  N   GLN   172     -36.007  23.153  31.041  1.00 44.49      L    N
ATOM   5635  CA  GLN   172     -35.185  22.077  31.583  1.00 46.21      L    C
ATOM   5636  CB  GLN   172     -35.218  22.112  33.116  1.00 45.47      L    C
ATOM   5637  CG  GLN   172     -34.806  23.449  33.714  1.00 49.31      L    C
ATOM   5638  CD  GLN   172     -35.245  23.619  35.167  1.00 49.94      L    C
ATOM   5639  OE1 GLN   172     -34.665  24.403  35.909  1.00 52.52      L    O
ATOM   5640  NE2 GLN   172     -36.273  22.885  35.570  1.00 51.29      L    N
ATOM   5641  C   GLN   172     -35.736  20.743  31.093  1.00 45.91      L    C
ATOM   5642  O   GLN   172     -36.790  20.692  30.461  1.00 46.76      L    O
ATOM   5643  N   SER   173     -35.031  19.662  31.396  1.00 46.60      L    N
ATOM   5644  CA  SER   173     -35.463  18.341  30.961  1.00 48.58      L    C
ATOM   5645  CB  SER   173     -34.335  17.324  31.157  1.00 49.25      L    C
ATOM   5646  OG  SER   173     -34.118  17.066  32.535  1.00 52.49      L    O
ATOM   5647  C   SER   173     -36.715  17.877  31.711  1.00 48.45      L    C
ATOM   5648  O   SER   173     -37.363  16.911  31.307  1.00 48.69      L    O
ATOM   5649  N   ASN   174     -37.058  18.562  32.799  1.00 47.11      L    N
ATOM   5650  CA  ASN   174     -38.284  18.243  33.526  1.00 47.32      L    C
ATOM   5651  CB  ASN   174     -38.067  18.376  35.032  1.00 48.35      L    C
ATOM   5652  CG  ASN   174     -37.959  19.818  35.483  1.00 51.01      L    C
ATOM   5653  OD1 ASN   174     -37.810  20.733  34.666  1.00 50.68      L    O
ATOM   5654  ND2 ASN   174     -38.036  20.032  36.794  1.00 52.49      L    N
ATOM   5655  C   ASN   174     -39.426  19.154  33.093  1.00 47.13      L    C
ATOM   5656  O   ASN   174     -40.484  19.183  33.721  1.00 47.39      L    O
ATOM   5657  N   ASN   175     -39.188  19.912  32.028  1.00 46.42      L    N
ATOM   5658  CA  ASN   175     -40.231  20.697  31.378  1.00 46.60      L    C
ATOM   5659  CB  ASN   175     -41.476  19.840  31.148  1.00 49.67      L    C
ATOM   5660  CG  ASN   175     -42.214  20.236  29.886  1.00 56.34      L    C
ATOM   5661  OD1 ASN   175     -43.406  20.564  29.920  1.00 58.64      L    O
ATOM   5662  ND2 ASN   175     -41.502  20.218  28.757  1.00 57.45      L    N
ATOM   5663  C   ASN   175     -40.621  21.969  32.126  1.00 44.32      L    C
ATOM   5664  O   ASN   175     -41.609  22.620  31.789  1.00 42.56      L    O
ATOM   5665  N   LYS   176     -39.847  22.322  33.144  1.00 42.39      L    N
ATOM   5666  CA  LYS   176     -39.974  23.638  33.753  1.00 40.88      L    C
ATOM   5667  CB  LYS   176     -39.993  23.517  35.281  1.00 41.57      L    C
ATOM   5668  CG  LYS   176     -41.086  22.592  35.806  1.00 43.43      L    C
ATOM   5669  CD  LYS   176     -41.149  22.596  37.330  1.00 46.57      L    C
ATOM   5670  CE  LYS   176     -41.815  21.331  37.867  1.00 48.69      L    C
ATOM   5671  NZ  LYS   176     -42.250  21.472  39.295  1.00 52.64      L    N
ATOM   5672  C   LYS   176     -38.802  24.500  33.290  1.00 39.00      L    C
ATOM   5673  O   LYS   176     -37.914  24.020  32.580  1.00 36.87      L    O
ATOM   5674  N   TYR   177     -38.804  25.768  33.686  1.00 37.05      L    N
ATOM   5675  CA  TYR   177     -37.856  26.731  33.143  1.00 37.79      L    C
ATOM   5676  CB  TYR   177     -38.582  28.028  32.787  1.00 40.04      L    C
ATOM   5677  CG  TYR   177     -39.386  27.955  31.509  1.00 42.10      L    C
ATOM   5678  CD1 TYR   177     -38.803  28.262  30.285  1.00 43.92      L    C
ATOM   5679  CE1 TYR   177     -39.535  28.227  29.114  1.00 46.21      L    C
ATOM   5680  CD2 TYR   177     -40.732  27.604  31.527  1.00 43.08      L    C
ATOM   5681  CE2 TYR   177     -41.477  27.568  30.361  1.00 46.97      L    C
ATOM   5682  CZ  TYR   177     -40.870  27.882  29.153  1.00 47.58      L    C
ATOM   5683  OH  TYR   177     -41.600  27.858  27.987  1.00 48.95      L    O
ATOM   5684  C   TYR   177     -36.682  27.054  34.065  1.00 36.65      L    C
ATOM   5685  O   TYR   177     -36.783  26.952  35.291  1.00 35.52      L    O
ATOM   5686  N   ALA   178     -35.571  27.457  33.458  1.00 34.98      L    N
ATOM   5687  CA  ALA   178     -34.397  27.889  34.205  1.00 34.14      L    C
ATOM   5688  CB  ALA   178     -33.345  26.777  34.220  1.00 33.79      L    C
ATOM   5689  C   ALA   178     -33.804  29.155  33.599  1.00 34.93      L    C
ATOM   5690  O   ALA   178     -33.840  29.362  32.380  1.00 34.79      L    O
ATOM   5691  N   ALA   179     -33.252  30.000  34.461  1.00 32.77      L    N
ATOM   5692  CA  ALA   179     -32.651  31.192  34.019  1.00 31.61      L    C
ATOM   5693  CB  ALA   179     -33.519  32.369  33.943  1.00 30.08      L    C
ATOM   5694  C   ALA   179     -31.450  31.478  34.924  1.00 32.24      L    C
ATOM   5695  O   ALA   179     -31.486  30.976  36.148  1.00 31.98      L    O
ATOM   5696  N   SER   180     -30.468  32.277  34.629  1.00 32.46      L    N
ATOM   5697  CA  SER   180     -29.469  32.730  35.585  1.00 34.62      L    C
ATOM   5698  CB  SER   180     -28.191  31.887  35.475  1.00 34.30      L    C
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ATOM   5699  OG  SER   180     -27.821  31.682  34.130  1.00 36.12      L    O
ATOM   5700  C   SER   180     -29.149  34.201  35.401  1.00 35.73      L    C
ATOM   5701  O   SER   180     -29.371  34.762  34.328  1.00 36.20      L    O
ATOM   5702  N   SER   181     -28.648  34.827  36.464  1.00 35.66      L    N
ATOM   5703  CA  SER   181     -28.306  36.242  36.428  1.00 35.32      L    C
ATOM   5704  CB  SER   181     -29.420  37.066  37.085  1.00 35.39      L    C
ATOM   5705  OG  SER   181     -29.219  38.452  36.863  1.00 35.35      L    O
ATOM   5706  C   SER   181     -26.975  36.503  37.134  1.00 34.93      L    C
ATOM   5707  O   SER   181     -26.740  36.028  38.246  1.00 34.66      L    O
ATOM   5708  N   TYR   182     -26.110  37.263  36.474  1.00 34.88      L    N
ATOM   5709  CA  TYR   182     -24.768  37.537  36.970  1.00 34.05      L    C
ATOM   5710  CB  TYR   182     -23.739  37.082  35.936  1.00 33.06      L    C
ATOM   5711  CG  TYR   182     -23.721  35.591  35.710  1.00 33.67      L    C
ATOM   5712  CD1 TYR   182     -24.573  34.994  34.787  1.00 33.90      L    C
ATOM   5713  CE1 TYR   182     -24.556  33.619  34.580  1.00 32.99      L    C
ATOM   5714  CD2 TYR   182     -22.850  34.775  36.422  1.00 34.60      L    C
ATOM   5715  CE2 TYR   182     -22.823  33.407  36.226  1.00 33.92      L    C
ATOM   5716  CZ  TYR   182     -23.675  32.832  35.304  1.00 34.85      L    C
ATOM   5717  OH  TYR   182     -23.624  31.471  35.101  1.00 35.11      L    O
ATOM   5718  C   TYR   182     -24.576  39.027  37.246  1.00 35.65      L    C
ATOM   5719  O   TYR   182     -24.817  39.863  36.373  1.00 34.81      L    O
ATOM   5720  N   LEU   183     -24.136  39.358  38.456  1.00 35.39      L    N
ATOM   5721  CA  LEU   183     -23.780  40.739  38.778  1.00 36.48      L    C
ATOM   5722  CB  LEU   183     -24.485  41.185  40.062  1.00 36.53      L    C
ATOM   5723  CG  LEU   183     -24.120  42.570  40.611  1.00 38.34      L    C
ATOM   5724  CD1 LEU   183     -24.315  43.637  39.541  1.00 38.29      L    C
ATOM   5725  CD2 LEU   183     -24.983  42.871  41.831  1.00 37.55      L    C
ATOM   5726  C   LEU   183     -22.269  40.871  38.946  1.00 37.69      L    C
ATOM   5727  O   LEU   183     -21.681  40.268  39.847  1.00 38.05      L    O
ATOM   5728  N   SER   184     -21.648  41.650  38.067  1.00 38.46      L    N
ATOM   5729  CA  SER   184     -20.215  41.907  38.142  1.00 40.81      L    C
ATOM   5730  CB  SER   184     -19.639  42.113  36.741  1.00 39.79      L    C
ATOM   5731  OG  SER   184     -19.775  40.942  35.960  1.00 42.53      L    O
ATOM   5732  C   SER   184     -19.913  43.138  38.992  1.00 42.63      L    C
ATOM   5733  O   SER   184     -20.460  44.216  38.759  1.00 44.23      L    O
ATOM   5734  N   LEU   185     -19.036  42.963  39.977  1.00 44.38      L    N
ATOM   5735  CA  LEU   185     -18.597  44.046  40.854  1.00 44.18      L    C
ATOM   5736  CB  LEU   185     -19.093  43.804  42.276  1.00 43.90      L    C
ATOM   5737  CG  LEU   185     -20.592  43.694  42.518  1.00 45.77      L    C
ATOM   5738  CD1 LEU   185     -20.840  43.370  43.987  1.00 43.80      L    C
ATOM   5739  CD2 LEU   185     -21.267  45.005  42.125  1.00 46.33      L    C
ATOM   5740  C   LEU   185     -17.069  44.087  40.879  1.00 45.19      L    C
ATOM   5741  O   LEU   185     -16.414  43.094  40.556  1.00 45.27      L    O
ATOM   5742  N   THR   186     -16.502  45.225  41.271  1.00 44.94      L    N
ATOM   5743  CA  THR   186     -15.103  45.247  41.691  1.00 44.94      L    C
ATOM   5744  CB  THR   186     -14.512  46.671  41.708  1.00 44.97      L    C
ATOM   5745  OG1 THR   186     -15.145  47.439  42.742  1.00 47.27      L    O
ATOM   5746  CG2 THR   186     -14.720  47.353  40.367  1.00 42.76      L    C
ATOM   5747  C   THR   186     -15.067  44.711  43.115  1.00 44.48      L    C
ATOM   5748  O   THR   186     -16.047  44.825  43.859  1.00 43.11      L    O
ATOM   5749  N   PRO   187     -13.937  44.113  43.515  1.00 44.61      L    N
ATOM   5750  CD  PRO   187     -12.753  43.786  42.702  1.00 43.46      L    C
ATOM   5751  CA  PRO   187     -13.813  43.620  44.890  1.00 45.61      L    C
ATOM   5752  CB  PRO   187     -12.374  43.113  44.959  1.00 43.07      L    C
ATOM   5753  CG  PRO   187     -12.048  42.759  43.546  1.00 44.14      L    C
ATOM   5754  C   PRO   187     -14.098  44.714  45.917  1.00 47.37      L    C
ATOM   5755  O   PRO   187     -14.677  44.452  46.973  1.00 47.85      L    O
ATOM   5756  N   GLU   188     -13.707  45.941  45.597  1.00 49.99      L    N
ATOM   5757  CA  GLU   188     -13.931  47.066  46.500  1.00 54.63      L    C
ATOM   5758  CB  GLU   188     -13.236  48.323  45.963  1.00 57.78      L    C
ATOM   5759  CG  GLU   188     -11.728  48.174  45.779  1.00 64.11      L    C
ATOM   5760  CD  GLU   188     -11.354  47.335  44.562  1.00 67.50      L    C
ATOM   5761  OE1 GLU   188     -10.157  47.000  44.411  1.00 70.08      L    O
ATOM   5762  OE2 GLU   188     -12.252  47.012  43.756  1.00 69.12      L    O
ATOM   5763  C   GLU   188     -15.424  47.342  46.695  1.00 54.97      L    C
ATOM   5764  O   GLU   188     -15.875  47.570  47.818  1.00 54.27      L    O
ATOM   5765  N   GLN   189     -16.186  47.318  45.602  1.00 54.96      L    N
ATOM   5766  CA  GLN   189     -17.634  47.478  45.688  1.00 54.99      L    C
ATOM   5767  CB  GLN   189     -18.266  47.452  44.297  1.00 57.33      L    C
ATOM   5768  CG  GLN   189     -17.986  48.677  43.461  1.00 59.74      L    C
ATOM   5769  CD  GLN   189     -18.586  48.570  42.076  1.00 62.10      L    C
ATOM   5770  OE1 GLN   189     -18.126  47.785  41.244  1.00 62.33      L    O
ATOM   5771  NE2 GLN   189     -19.623  49.359  41.819  1.00 63.40      L    N
ATOM   5772  C   GLN   189     -18.245  46.370  46.533  1.00 54.40      L    C
ATOM   5773  O   GLN   189     -19.102  46.623  47.379  1.00 54.55      L    O
ATOM   5774  N   TRP   190     -17.803  45.139  46.298  1.00 53.18      L    N
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ATOM	5775	CA	TRP	190	-18.328	43.996	47.031	1.00	52.69	L	C
ATOM	5776	CB	TRP	190	-17.647	42.713	46.555	1.00	47.71	L	C
ATOM	5777	CG	TRP	190	-17.835	41.541	47.469	1.00	43.39	L	C
ATOM	5778	CD2	TRP	190	-19.078	40.931	47.852	1.00	42.46	L	C
ATOM	5779	CE2	TRP	190	-18.767	39.839	48.690	1.00	41.15	L	C
ATOM	5780	CE3	TRP	190	-20.420	41.201	47.568	1.00	40.32	L	C
ATOM	5781	CD1	TRP	190	-16.853	40.815	48.076	1.00	42.75	L	C
ATOM	5782	NE1	TRP	190	-17.403	39.789	48.810	1.00	40.87	L	N
ATOM	5783	CZ2	TRP	190	-19.749	39.017	49.246	1.00	39.96	L	C
ATOM	5784	CZ3	TRP	190	-21.395	40.381	48.123	1.00	39.31	L	C
ATOM	5785	CH2	TRP	190	-21.054	39.306	48.951	1.00	37.86	L	C
ATOM	5786	C	TRP	190	-18.122	44.161	48.532	1.00	55.45	L	C
ATOM	5787	O	TRP	190	-19.014	43.858	49.325	1.00	56.54	L	O
ATOM	5788	N	LYS	191	-16.949	44.647	48.922	1.00	57.84	L	N
ATOM	5789	CA	LYS	191	-16.606	44.715	50.337	1.00	61.22	L	C
ATOM	5790	CB	LYS	191	-15.087	44.624	50.514	1.00	63.09	L	C
ATOM	5791	CG	LYS	191	-14.517	43.262	50.141	1.00	68.04	L	C
ATOM	5792	CD	LYS	191	-13.238	42.951	50.908	1.00	71.58	L	C
ATOM	5793	CE	LYS	191	-12.009	43.532	50.217	1.00	73.94	L	C
ATOM	5794	NZ	LYS	191	-11.653	42.782	48.975	1.00	75.18	L	N
ATOM	5795	C	LYS	191	-17.142	45.962	51.037	1.00	61.69	L	C
ATOM	5796	O	LYS	191	-17.204	46.008	52.264	1.00	61.30	L	O
ATOM	5797	N	SER	192	-17.545	46.962	50.259	1.00	62.65	L	N
ATOM	5798	CA	SER	192	-17.975	48.237	50.824	1.00	63.95	L	C
ATOM	5799	CB	SER	192	-17.613	49.382	49.874	1.00	64.65	L	C
ATOM	5800	OG	SER	192	-18.392	49.335	48.692	1.00	67.19	L	O
ATOM	5801	C	SER	192	-19.467	48.315	51.161	1.00	64.72	L	C
ATOM	5802	O	SER	192	-19.935	49.335	51.673	1.00	66.00	L	O
ATOM	5803	N	HIS	193	-20.217	47.253	50.874	1.00	63.86	L	N
ATOM	5804	CA	HIS	193	-21.638	47.219	51.216	1.00	61.95	L	C
ATOM	5805	CB	HIS	193	-22.485	46.954	49.970	1.00	61.37	L	C
ATOM	5806	CG	HIS	193	-22.486	48.084	48.988	1.00	61.60	L	C
ATOM	5807	CD2	HIS	193	-23.325	49.136	48.841	1.00	61.56	L	C
ATOM	5808	ND1	HIS	193	-21.535	48.211	47.998	1.00	61.98	L	N
ATOM	5809	CE1	HIS	193	-21.789	49.293	47.283	1.00	62.58	L	C
ATOM	5810	NE2	HIS	193	-22.870	49.872	47.773	1.00	62.31	L	N
ATOM	5811	C	HIS	193	-21.924	46.155	52.260	1.00	62.02	L	C
ATOM	5812	O	HIS	193	-21.148	45.216	52.428	1.00	62.20	L	O
ATOM	5813	N	ARG	194	-23.040	46.305	52.965	1.00	62.45	L	N
ATOM	5814	CA	ARG	194	-23.416	45.343	53.991	1.00	63.53	L	C
ATOM	5815	CB	ARG	194	-24.500	45.926	54.902	1.00	68.20	L	C
ATOM	5816	CG	ARG	194	-23.971	46.895	55.941	1.00	75.45	L	C
ATOM	5817	CD	ARG	194	-23.055	46.173	56.922	1.00	80.85	L	C
ATOM	5818	NE	ARG	194	-22.122	47.082	57.584	1.00	85.64	L	N
ATOM	5819	CZ	ARG	194	-21.136	46.687	58.386	1.00	87.72	L	C
ATOM	5820	NH1	ARG	194	-20.332	47.584	58.947	1.00	88.22	L	N
ATOM	5821	NH2	ARG	194	-20.953	45.394	58.629	1.00	88.28	L	N
ATOM	5822	C	ARG	194	-23.916	44.049	53.369	1.00	61.25	L	C
ATOM	5823	O	ARG	194	-23.696	42.964	53.910	1.00	60.80	L	O
ATOM	5824	N	SER	195	-24.595	44.164	52.232	1.00	57.80	L	N
ATOM	5825	CA	SER	195	-25.120	42.984	51.562	1.00	55.09	L	C
ATOM	5826	CB	SER	195	-26.280	42.393	52.369	1.00	55.10	L	C
ATOM	5827	OG	SER	195	-27.376	43.289	52.411	1.00	57.11	L	O
ATOM	5828	C	SER	195	-25.581	43.272	50.142	1.00	52.03	L	C
ATOM	5829	O	SER	195	-25.760	44.423	49.748	1.00	51.53	L	O
ATOM	5830	N	TYR	196	-25.757	42.205	49.374	1.00	48.88	L	N
ATOM	5831	CA	TYR	196	-26.331	42.298	48.044	1.00	45.63	L	C
ATOM	5832	CB	TYR	196	-25.279	41.968	46.989	1.00	44.45	L	C
ATOM	5833	CG	TYR	196	-24.395	43.139	46.626	1.00	44.49	L	C
ATOM	5834	CD1	TYR	196	-24.664	43.904	45.501	1.00	43.92	L	C
ATOM	5835	CE1	TYR	196	-23.850	44.957	45.140	1.00	46.40	L	C
ATOM	5836	CD2	TYR	196	-23.279	43.464	47.390	1.00	45.08	L	C
ATOM	5837	CE2	TYR	196	-22.451	44.524	47.035	1.00	45.84	L	C
ATOM	5838	CZ	TYR	196	-22.742	45.263	45.907	1.00	47.07	L	C
ATOM	5839	OH	TYR	196	-21.920	46.301	45.516	1.00	48.23	L	O
ATOM	5840	C	TYR	196	-27.493	41.326	47.946	1.00	44.90	L	C
ATOM	5841	O	TYR	196	-27.504	40.290	48.614	1.00	44.93	L	O
ATOM	5842	N	SER	197	-28.480	41.669	47.126	1.00	44.07	L	N
ATOM	5843	CA	SER	197	-29.666	40.838	47.001	1.00	44.08	L	C
ATOM	5844	CB	SER	197	-30.846	41.485	47.727	1.00	43.76	L	C
ATOM	5845	OG	SER	197	-30.752	41.272	49.125	1.00	46.25	L	O
ATOM	5846	C	SER	197	-30.043	40.562	45.556	1.00	43.65	L	C
ATOM	5847	O	SER	197	-29.923	41.432	44.687	1.00	42.68	L	O
ATOM	5848	N	CYS	198	-30.490	39.335	45.311	1.00	42.68	L	N
ATOM	5849	CA	CYS	198	-31.085	38.966	44.037	1.00	43.42	L	C
ATOM	5850	C	CYS	198	-32.590	38.845	44.260	1.00	42.81	L	C

ATOM	5851	O	CYS	198	-33.036	38.049	45.089	1.00	41.88	L	O
ATOM	5852	CB	CYS	198	-30.516	37.624	43.548	1.00	42.55	L	C
ATOM	5853	SG	CYS	198	-31.115	37.164	41.887	1.00	48.59	L	S
ATOM	5854	N	GLN	199	-33.368	39.641	43.532	1.00	42.44	L	N
ATOM	5855	CA	GLN	199	-34.826	39.605	43.650	1.00	43.18	L	C
ATOM	5856	CB	GLN	199	-35.370	41.007	43.922	1.00	45.67	L	C
ATOM	5857	CG	GLN	199	-34.995	41.547	45.290	1.00	53.84	L	C
ATOM	5858	CD	GLN	199	-35.634	42.891	45.584	1.00	56.85	L	C
ATOM	5859	OE1	GLN	199	-35.855	43.699	44.681	1.00	60.04	L	O
ATOM	5860	NE2	GLN	199	-35.935	43.136	46.852	1.00	58.12	L	N
ATOM	5861	C	GLN	199	-35.462	39.057	42.381	1.00	41.40	L	C
ATOM	5862	O	GLN	199	-35.324	39.639	41.306	1.00	40.06	L	O
ATOM	5863	N	VAL	200	-36.160	37.935	42.508	1.00	40.36	L	N
ATOM	5864	CA	VAL	200	-36.772	37.303	41.347	1.00	39.58	L	C
ATOM	5865	CB	VAL	200	-36.363	35.811	41.230	1.00	40.32	L	C
ATOM	5866	CG1	VAL	200	-37.035	35.177	40.010	1.00	37.77	L	C
ATOM	5867	CG2	VAL	200	-34.842	35.699	41.120	1.00	38.20	L	C
ATOM	5868	C	VAL	200	-38.284	37.396	41.433	1.00	39.21	L	C
ATOM	5869	O	VAL	200	-38.897	36.911	42.388	1.00	37.45	L	O
ATOM	5870	N	THR	201	-38.880	38.029	40.429	1.00	39.72	L	N
ATOM	5871	CA	THR	201	-40.323	38.206	40.389	1.00	40.93	L	C
ATOM	5872	CB	THR	201	-40.698	39.680	40.082	1.00	43.40	L	C
ATOM	5873	OG1	THR	201	-40.139	40.532	41.092	1.00	45.08	L	O
ATOM	5874	CG2	THR	201	-42.218	39.857	40.073	1.00	43.33	L	C
ATOM	5875	C	THR	201	-40.935	37.296	39.332	1.00	40.49	L	C
ATOM	5876	O	THR	201	-40.502	37.278	38.176	1.00	39.58	L	O
ATOM	5877	N	HIS	202	-41.942	36.538	39.748	1.00	41.15	L	N
ATOM	5878	CA	HIS	202	-42.586	35.550	38.891	1.00	41.11	L	C
ATOM	5879	CB	HIS	202	-42.032	34.156	39.196	1.00	38.98	L	C
ATOM	5880	CG	HIS	202	-42.664	33.063	38.394	1.00	36.71	L	C
ATOM	5881	CD2	HIS	202	-42.498	32.696	37.101	1.00	35.64	L	C
ATOM	5882	ND1	HIS	202	-43.575	32.179	38.930	1.00	35.58	L	N
ATOM	5883	CE1	HIS	202	-43.941	31.311	38.003	1.00	35.45	L	C
ATOM	5884	NE2	HIS	202	-43.301	31.602	36.883	1.00	35.15	L	N
ATOM	5885	C	HIS	202	-44.081	35.572	39.165	1.00	42.13	L	C
ATOM	5886	O	HIS	202	-44.516	35.276	40.279	1.00	41.36	L	O
ATOM	5887	N	GLU	203	-44.862	35.931	38.151	1.00	44.96	L	N
ATOM	5888	CA	GLU	203	-46.310	36.022	38.298	1.00	47.24	L	C
ATOM	5889	CB	GLU	203	-46.907	34.628	38.489	1.00	49.44	L	C
ATOM	5890	CG	GLU	203	-46.537	33.646	37.394	1.00	53.20	L	C
ATOM	5891	CD	GLU	203	-47.209	33.974	36.076	1.00	57.10	L	C
ATOM	5892	OE1	GLU	203	-46.488	34.219	35.082	1.00	59.14	L	O
ATOM	5893	OE2	GLU	203	-48.460	33.987	36.038	1.00	57.31	L	O
ATOM	5894	C	GLU	203	-46.676	36.900	39.486	1.00	47.57	L	C
ATOM	5895	O	GLU	203	-47.514	36.529	40.306	1.00	47.36	L	O
ATOM	5896	N	GLY	203	-46.031	38.059	39.586	1.00	49.08	L	N
ATOM	5897	CA	GLY	203	-46.380	39.009	40.626	1.00	50.03	L	C
ATOM	5898	C	GLY	203	-45.999	38.603	42.040	1.00	52.56	L	C
ATOM	5899	O	GLY	203	-46.399	39.264	43.000	1.00	54.19	L	O
ATOM	5900	N	SER	205	-45.243	37.519	42.189	1.00	52.78	L	N
ATOM	5901	CA	SER	205	-44.651	37.192	43.484	1.00	53.25	L	C
ATOM	5902	CB	SER	205	-45.060	35.790	43.934	1.00	54.26	L	C
ATOM	5903	OG	SER	205	-46.365	35.790	44.481	1.00	57.58	L	O
ATOM	5904	C	SER	205	-43.135	37.280	43.423	1.00	53.40	L	C
ATOM	5905	O	SER	205	-42.511	36.793	42.476	1.00	53.90	L	O
ATOM	5906	N	THR	206	-42.548	37.902	44.439	1.00	53.20	L	N
ATOM	5907	CA	THR	206	-41.107	38.107	44.486	1.00	54.04	L	C
ATOM	5908	CB	THR	206	-40.781	39.557	44.891	1.00	54.25	L	C
ATOM	5909	OG1	THR	206	-41.343	40.455	43.926	1.00	55.24	L	O
ATOM	5910	CG2	THR	206	-39.279	39.771	44.948	1.00	54.82	L	C
ATOM	5911	C	THR	206	-40.415	37.147	45.457	1.00	53.61	L	C
ATOM	5912	O	THR	206	-40.946	36.985	46.599	1.00	53.73	L	O
ATOM	5913	N	VAL	207	-39.351	36.501	44.987	1.00	52.25	L	N
ATOM	5914	CA	VAL	207	-38.490	35.690	45.848	1.00	51.34	L	C
ATOM	5915	CB	VAL	207	-38.309	34.254	45.282	1.00	51.23	L	C
ATOM	5916	CG1	VAL	207	-37.331	33.472	46.137	1.00	50.68	L	C
ATOM	5917	CG2	VAL	207	-39.642	33.535	45.240	1.00	51.24	L	C
ATOM	5918	C	VAL	207	-37.120	36.365	45.941	1.00	51.56	L	C
ATOM	5919	O	VAL	207	-36.562	36.801	44.930	1.00	50.04	L	O
ATOM	5920	N	GLU	208	-36.582	36.451	47.154	1.00	51.89	L	N
ATOM	5921	CA	GLU	208	-35.353	37.203	47.389	1.00	51.48	L	C
ATOM	5922	CB	GLU	208	-35.661	38.430	48.241	1.00	53.42	L	C
ATOM	5923	CG	GLU	208	-34.455	39.296	48.538	1.00	58.86	L	C
ATOM	5924	CD	GLU	208	-34.835	40.595	49.224	1.00	61.76	L	C
ATOM	5925	OE1	GLU	208	-34.754	41.657	48.569	1.00	63.24	L	O
ATOM	5926	OE2	GLU	208	-35.218	40.554	50.415	1.00	62.32	L	O

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ATOM   5927  C   GLU   208     -34.260  36.375  48.066  1.00 50.33      L    C
ATOM   5928  O   GLU   208     -34.530  35.605  48.989  1.00 49.28      L    O
ATOM   5929  N   LYS   209     -33.026  36.535  47.601  1.00 47.90      L    N
ATOM   5930  CA  LYS   209     -31.877  35.983  48.310  1.00 47.26      L    C
ATOM   5931  CB  LYS   209     -31.271  34.810  47.530  1.00 47.11      L    C
ATOM   5932  CG  LYS   209     -32.148  33.570  47.496  1.00 45.45      L    C
ATOM   5933  CD  LYS   209     -32.621  33.211  48.887  1.00 46.72      L    C
ATOM   5934  CE  LYS   209     -33.750  32.188  48.858  1.00 48.50      L    C
ATOM   5935  NZ  LYS   209     -33.244  30.791  48.874  1.00 49.58      L    N
ATOM   5936  C   LYS   209     -30.825  37.067  48.529  1.00 46.93      L    C
ATOM   5937  O   LYS   209     -30.685  37.986  47.714  1.00 44.32      L    O
ATOM   5938  N   THR   210     -30.096  36.952  49.636  1.00 46.89      L    N
ATOM   5939  CA  THR   210     -29.125  37.968  50.037  1.00 48.05      L    C
ATOM   5940  CB  THR   210     -29.661  38.817  51.222  1.00 49.81      L    C
ATOM   5941  OG1 THR   210     -30.865  39.487  50.831  1.00 50.72      L    O
ATOM   5942  CG2 THR   210     -28.634  39.857  51.643  1.00 50.82      L    C
ATOM   5943  C   THR   210     -27.805  37.330  50.470  1.00 47.03      L    C
ATOM   5944  O   THR   210     -27.796  36.316  51.170  1.00 45.40      L    O
ATOM   5945  N   VAL   211     -26.692  37.926  50.057  1.00 46.63      L    N
ATOM   5946  CA  VAL   211     -25.389  37.516  50.565  1.00 46.45      L    C
ATOM   5947  CB  VAL   211     -24.565  36.770  49.494  1.00 44.37      L    C
ATOM   5948  CG1 VAL   211     -25.186  35.408  49.225  1.00 41.67      L    C
ATOM   5949  CG2 VAL   211     -24.489  37.602  48.217  1.00 40.70      L    C
ATOM   5950  C   VAL   211     -24.580  38.701  51.066  1.00 48.37      L    C
ATOM   5951  O   VAL   211     -24.732  39.823  50.577  1.00 48.59      L    O
ATOM   5952  N   ALA   212     -23.722  38.441  52.048  1.00 50.94      L    N
ATOM   5953  CA  ALA   212     -22.915  39.485  52.676  1.00 52.66      L    C
ATOM   5954  CB  ALA   212     -23.324  39.647  54.135  1.00 51.32      L    C
ATOM   5955  C   ALA   212     -21.431  39.150  52.585  1.00 53.53      L    C
ATOM   5956  O   ALA   212     -21.037  37.991  52.706  1.00 52.26      L    O
ATOM   5957  N   PRO   213     -20.588  40.172  52.383  1.00 55.64      L    N
ATOM   5958  CD  PRO   213     -20.964  41.593  52.338  1.00 56.63      L    C
ATOM   5959  CA  PRO   213     -19.137  39.980  52.293  1.00 59.09      L    C
ATOM   5960  CB  PRO   213     -18.597  41.391  52.069  1.00 57.17      L    C
ATOM   5961  CG  PRO   213     -19.661  42.294  52.577  1.00 57.07      L    C
ATOM   5962  C   PRO   213     -18.579  39.337  53.554  1.00 63.09      L    C
ATOM   5963  O   PRO   213     -17.587  38.605  53.511  1.00 63.69      L    O
ATOM   5964  N   THR   214     -19.236  39.607  54.675  1.00 67.26      L    N
ATOM   5965  CA  THR   214     -18.898  38.972  55.939  1.00 72.25      L    C
ATOM   5966  CB  THR   214     -19.825  39.478  57.069  1.00 73.76      L    C
ATOM   5967  OG1 THR   214     -19.832  40.913  57.072  1.00 74.88      L    O
ATOM   5968  CG2 THR   214     -19.336  38.979  58.429  1.00 74.89      L    C
ATOM   5969  C   THR   214     -19.025  37.447  55.821  1.00 73.85      L    C
ATOM   5970  O   THR   214     -19.975  36.886  56.412  1.00 75.17      L    O
ATOM   5971  OXT THR   214     -18.178  36.832  55.130  1.00 74.44      L    O
TER    5972       THR   214                                             L
ATOM   5973  CB  GLU     1     -30.422   3.027   8.715  1.00 67.00      H    C
ATOM   5974  CG  GLU     1     -29.516   3.883   9.599  1.00 72.68      H    C
ATOM   5975  CD  GLU     1     -30.286   4.614  10.699  1.00 76.76      H    C
ATOM   5976  OE1 GLU     1     -30.875   3.938  11.574  1.00 78.40      H    O
ATOM   5977  OE2 GLU     1     -30.305   5.867  10.687  1.00 78.19      H    O
ATOM   5978  C   GLU     1     -29.301   3.794   6.619  1.00 59.84      H    C
ATOM   5979  O   GLU     1     -28.098   4.037   6.505  1.00 59.47      H    O
ATOM   5980  N   GLU     1     -28.667   1.616   7.662  1.00 63.51      H    N
ATOM   5981  CA  GLU     1     -29.785   2.571   7.395  1.00 63.39      H    C
ATOM   5982  N   VAL     2     -30.249   4.558   6.086  1.00 56.40      H    N
ATOM   5983  CA  VAL     2     -29.937   5.786   5.366  1.00 53.61      H    C
ATOM   5984  CB  VAL     2     -31.204   6.398   4.751  1.00 53.06      H    C
ATOM   5985  CG1 VAL     2     -30.873   7.721   4.084  1.00 53.81      H    C
ATOM   5986  CG2 VAL     2     -31.808   5.430   3.747  1.00 52.96      H    C
ATOM   5987  C   VAL     2     -29.313   6.798   6.314  1.00 52.68      H    C
ATOM   5988  O   VAL     2     -29.805   7.001   7.422  1.00 53.56      H    O
ATOM   5989  N   GLN     3     -28.231   7.438   5.887  1.00 49.43      H    N
ATOM   5990  CA  GLN     3     -27.479   8.274   6.805  1.00 48.34      H    C
ATOM   5991  CB  GLN     3     -26.566   7.385   7.649  1.00 51.60      H    C
ATOM   5992  CG  GLN     3     -25.834   8.110   8.751  1.00 57.66      H    C
ATOM   5993  CD  GLN     3     -25.334   7.163   9.826  1.00 61.96      H    C
ATOM   5994  OE1 GLN     3     -25.022   7.586  10.944  1.00 63.82      H    O
ATOM   5995  NE2 GLN     3     -25.257   5.873   9.496  1.00 62.05      H    N
ATOM   5996  C   GLN     3     -26.663   9.367   6.121  1.00 44.78      H    C
ATOM   5997  O   GLN     3     -26.083   9.154   5.056  1.00 45.81      H    O
ATOM   5998  N   LEU     4     -26.622  10.540   6.745  1.00 41.09      H    N
ATOM   5999  CA  LEU     4     -25.808  11.649   6.257  1.00 39.82      H    C
ATOM   6000  CB  LEU     4     -26.701  12.732   5.635  1.00 36.60      H    C
ATOM   6001  CG  LEU     4     -27.484  12.382   4.368  1.00 36.08      H    C
ATOM   6002  CD1 LEU     4     -28.454  13.513   4.027  1.00 34.06      H    C
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ATOM	6003	CD2	LEU	4	-26.512	12.144	3.226	1.00	32.76	H	C
ATOM	6004	C	LEU	4	-25.009	12.253	7.413	1.00	39.72	H	C
ATOM	6005	O	LEU	4	-25.575	12.613	8.444	1.00	39.93	H	O
ATOM	6006	N	VAL	5	-23.696	12.365	7.241	1.00	39.98	H	N
ATOM	6007	CA	VAL	5	-22.839	12.906	8.293	1.00	39.49	H	C
ATOM	6008	CB	VAL	5	-21.879	11.827	8.851	1.00	40.46	H	C
ATOM	6009	CG1	VAL	5	-21.035	12.426	9.971	1.00	37.54	H	C
ATOM	6010	CG2	VAL	5	-22.672	10.619	9.354	1.00	38.37	H	C
ATOM	6011	C	VAL	5	-21.991	14.059	7.774	1.00	40.34	H	C
ATOM	6012	O	VAL	5	-21.056	13.851	7.004	1.00	40.18	H	O
ATOM	6013	N	GLU	6	-22.308	15.278	8.188	1.00	40.67	H	N
ATOM	6014	CA	GLU	6	-21.511	16.411	7.744	1.00	43.08	H	C
ATOM	6015	CB	GLU	6	-22.383	17.671	7.589	1.00	42.43	H	C
ATOM	6016	CG	GLU	6	-23.333	17.966	8.728	1.00	47.10	H	C
ATOM	6017	CD	GLU	6	-24.611	17.139	8.680	1.00	45.96	H	C
ATOM	6018	OE1	GLU	6	-24.620	16.045	9.272	1.00	46.73	H	O
ATOM	6019	OE2	GLU	6	-25.606	17.582	8.064	1.00	43.63	H	O
ATOM	6020	C	GLU	6	-20.343	16.668	8.689	1.00	42.63	H	C
ATOM	6021	O	GLU	6	-20.410	16.352	9.876	1.00	45.51	H	O
ATOM	6022	N	SER	7	-19.258	17.215	8.151	1.00	41.61	H	N
ATOM	6023	CA	SER	7	-18.100	17.544	8.967	1.00	41.73	H	C
ATOM	6024	CB	SER	7	-17.167	16.334	9.090	1.00	43.65	H	C
ATOM	6025	OG	SER	7	-16.636	15.958	7.832	1.00	47.20	H	O
ATOM	6026	C	SER	7	-17.341	18.722	8.383	1.00	40.62	H	C
ATOM	6027	O	SER	7	-17.662	19.201	7.294	1.00	39.51	H	O
ATOM	6028	N	GLY	8	-16.345	19.195	9.126	1.00	39.23	H	N
ATOM	6029	CA	GLY	8	-15.489	20.261	8.640	1.00	35.58	H	C
ATOM	6030	C	GLY	8	-15.757	21.589	9.309	1.00	36.50	H	C
ATOM	6031	O	GLY	8	-15.050	22.566	9.069	1.00	37.45	H	O
ATOM	6032	N	GLY	9	-16.781	21.642	10.152	1.00	36.88	H	N
ATOM	6033	CA	GLY	9	-17.098	22.895	10.815	1.00	39.47	H	C
ATOM	6034	C	GLY	9	-15.964	23.359	11.717	1.00	40.73	H	C
ATOM	6035	O	GLY	9	-15.073	22.579	12.053	1.00	41.05	H	O
ATOM	6036	N	GLY	10	-15.991	24.628	12.105	1.00	40.13	H	N
ATOM	6037	CA	GLY	10	-14.982	25.132	13.014	1.00	39.76	H	C
ATOM	6038	C	GLY	10	-15.118	26.619	13.266	1.00	41.15	H	C
ATOM	6039	O	GLY	10	-16.118	27.239	12.892	1.00	41.81	H	O
ATOM	6040	N	LEU	11	-14.106	27.189	13.914	1.00	40.63	H	N
ATOM	6041	CA	LEU	11	-14.038	28.628	14.137	1.00	40.05	H	C
ATOM	6042	CB	LEU	11	-13.486	28.926	15.537	1.00	39.42	H	C
ATOM	6043	CG	LEU	11	-13.247	30.405	15.850	1.00	40.30	H	C
ATOM	6044	CD1	LEU	11	-14.555	31.157	15.728	1.00	39.60	H	C
ATOM	6045	CD2	LEU	11	-12.673	30.564	17.258	1.00	42.60	H	C
ATOM	6046	C	LEU	11	-13.109	29.217	13.087	1.00	39.22	H	C
ATOM	6047	O	LEU	11	-12.065	28.642	12.792	1.00	38.39	H	O
ATOM	6048	N	VAL	12	-13.493	30.356	12.521	1.00	39.30	H	N
ATOM	6049	CA	VAL	12	-12.696	31.006	11.489	1.00	41.06	H	C
ATOM	6050	CB	VAL	12	-13.127	30.525	10.062	1.00	43.20	H	C
ATOM	6051	CG1	VAL	12	-14.583	30.861	9.809	1.00	44.60	H	C
ATOM	6052	CG2	VAL	12	-12.266	31.172	9.000	1.00	45.13	H	C
ATOM	6053	C	VAL	12	-12.890	32.511	11.617	1.00	40.57	H	C
ATOM	6054	O	VAL	12	-13.887	32.962	12.179	1.00	40.30	H	O
ATOM	6055	N	LYS	13	-11.934	33.283	11.111	1.00	41.09	H	N
ATOM	6056	CA	LYS	13	-12.029	34.738	11.137	1.00	42.69	H	C
ATOM	6057	CB	LYS	13	-10.633	35.361	11.253	1.00	46.79	H	C
ATOM	6058	CG	LYS	13	-9.867	34.948	12.507	1.00	50.67	H	C
ATOM	6059	CD	LYS	13	-8.787	35.953	12.854	0.50	53.26	H	C
ATOM	6060	CE	LYS	13	-8.152	35.619	14.192	1.00	56.49	H	C
ATOM	6061	NZ	LYS	13	-9.187	35.477	15.251	1.00	58.69	H	N
ATOM	6062	C	LYS	13	-12.712	35.268	9.883	1.00	42.31	H	C
ATOM	6063	O	LYS	13	-12.663	34.645	8.823	1.00	41.76	H	O
ATOM	6064	N	PRO	14	-13.348	36.444	9.987	1.00	41.71	H	N
ATOM	6065	CD	PRO	14	-13.507	37.249	11.210	1.00	40.26	H	C
ATOM	6066	CA	PRO	14	-13.974	37.078	8.822	1.00	40.68	H	C
ATOM	6067	CB	PRO	14	-14.351	38.466	9.331	1.00	40.52	H	C
ATOM	6068	CG	PRO	14	-14.540	38.266	10.813	1.00	41.11	H	C
ATOM	6069	C	PRO	14	-13.008	37.139	7.646	1.00	42.52	H	C
ATOM	6070	O	PRO	14	-11.828	37.437	7.817	1.00	42.90	H	O
ATOM	6071	N	GLY	15	-13.513	36.839	6.452	1.00	42.59	H	N
ATOM	6072	CA	GLY	15	-12.668	36.859	5.275	1.00	40.18	H	C
ATOM	6073	C	GLY	15	-12.003	35.524	5.013	1.00	40.77	H	C
ATOM	6074	O	GLY	15	-11.419	35.320	3.948	1.00	41.52	H	O
ATOM	6075	N	GLY	16	-12.097	34.608	5.974	1.00	40.21	H	N
ATOM	6076	CA	GLY	16	-11.460	33.309	5.822	1.00	42.04	H	C
ATOM	6077	C	GLY	16	-12.291	32.279	5.067	1.00	43.65	H	C
ATOM	6078	O	GLY	16	-13.333	32.611	4.497	1.00	43.64	H	O

ATOM	6079	N	SER	17	-11.829	31.029	5.076	1.00	44.25	H	N
ATOM	6080	CA	SER	17	-12.407	29.961	4.261	1.00	45.74	H	C
ATOM	6081	CB	SER	17	-11.515	29.681	3.050	1.00	46.55	H	C
ATOM	6082	OG	SER	17	-11.340	30.848	2.267	1.00	53.54	H	O
ATOM	6083	C	SER	17	-12.576	28.662	5.047	1.00	45.37	H	C
ATOM	6084	O	SER	17	-11.792	28.372	5.946	1.00	44.90	H	O
ATOM	6085	N	LEU	18	-13.592	27.881	4.681	1.00	44.78	H	N
ATOM	6086	CA	LEU	18	-13.834	26.557	5.255	1.00	44.09	H	C
ATOM	6087	CB	LEU	18	-14.857	26.641	6.391	1.00	44.85	H	C
ATOM	6088	CG	LEU	18	-14.385	26.754	7.838	1.00	47.60	H	C
ATOM	6089	CD1	LEU	18	-15.602	26.870	8.755	1.00	46.16	H	C
ATOM	6090	CD2	LEU	18	-13.554	25.526	8.203	1.00	48.80	H	C
ATOM	6091	C	LEU	18	-14.386	25.624	4.183	1.00	43.53	H	C
ATOM	6092	O	LEU	18	-15.068	26.062	3.262	1.00	44.97	H	O
ATOM	6093	N	ARG	19	-14.106	24.336	4.315	1.00	41.93	H	N
ATOM	6094	CA	ARG	19	-14.658	23.350	3.407	1.00	41.54	H	C
ATOM	6095	CB	ARG	19	-13.538	22.673	2.617	1.00	42.94	H	C
ATOM	6096	CG	ARG	19	-14.030	21.678	1.573	1.00	46.58	H	C
ATOM	6097	CD	ARG	19	-12.969	21.451	0.511	1.00	51.36	H	C
ATOM	6098	NE	ARG	19	-13.459	20.676	-0.627	1.00	56.49	H	N
ATOM	6099	CZ	ARG	19	-13.334	19.357	-0.746	1.00	59.18	H	C
ATOM	6100	NH1	ARG	19	-13.807	18.737	-1.821	1.00	59.16	H	N
ATOM	6101	NH2	ARG	19	-12.743	18.653	0.213	1.00	59.69	H	N
ATOM	6102	C	ARG	19	-15.454	22.300	4.172	1.00	40.18	H	C
ATOM	6103	O	ARG	19	-14.884	21.482	4.898	1.00	39.83	H	O
ATOM	6104	N	LEU	20	-16.773	22.316	4.002	1.00	37.32	H	N
ATOM	6105	CA	LEU	20	-17.617	21.320	4.647	1.00	35.94	H	C
ATOM	6106	CB	LEU	20	-19.007	21.894	4.928	1.00	33.09	H	C
ATOM	6107	CG	LEU	20	-19.027	23.242	5.656	1.00	35.63	H	C
ATOM	6108	CD1	LEU	20	-20.474	23.659	5.917	1.00	34.30	H	C
ATOM	6109	CD2	LEU	20	-18.241	23.140	6.976	1.00	32.60	H	C
ATOM	6110	C	LEU	20	-17.731	20.093	3.763	1.00	35.92	H	C
ATOM	6111	O	LEU	20	-17.699	20.190	2.535	1.00	37.48	H	O
ATOM	6112	N	SER	21	-17.848	18.936	4.396	1.00	35.54	H	N
ATOM	6113	CA	SER	21	-18.112	17.699	3.685	1.00	37.66	H	C
ATOM	6114	CB	SER	21	-16.935	16.730	3.826	1.00	38.97	H	C
ATOM	6115	OG	SER	21	-15.752	17.291	3.290	1.00	45.86	H	O
ATOM	6116	C	SER	21	-19.356	17.061	4.274	1.00	38.28	H	C
ATOM	6117	O	SER	21	-19.740	17.356	5.404	1.00	37.83	H	O
ATOM	6118	N	CYS	22	-19.976	16.184	3.495	1.00	39.37	H	N
ATOM	6119	CA	CYS	22	-21.137	15.428	3.930	1.00	42.63	H	C
ATOM	6120	C	CYS	22	-20.974	14.049	3.305	1.00	42.53	H	C
ATOM	6121	O	CYS	22	-20.881	13.921	2.078	1.00	41.03	H	O
ATOM	6122	CB	CYS	22	-22.413	16.125	3.431	1.00	46.05	H	C
ATOM	6123	SG	CYS	22	-24.012	15.257	3.610	1.00	55.21	H	S
ATOM	6124	N	ALA	23	-20.901	13.019	4.145	1.00	40.59	H	N
ATOM	6125	CA	ALA	23	-20.748	11.658	3.645	1.00	41.14	H	C
ATOM	6126	CB	ALA	23	-19.698	10.905	4.464	1.00	40.28	H	C
ATOM	6127	C	ALA	23	-22.078	10.935	3.712	1.00	41.05	H	C
ATOM	6128	O	ALA	23	-22.754	10.959	4.739	1.00	43.31	H	O
ATOM	6129	N	ALA	24	-22.455	10.292	2.613	1.00	41.38	H	N
ATOM	6130	CA	ALA	24	-23.727	9.581	2.558	1.00	42.11	H	C
ATOM	6131	CB	ALA	24	-24.483	9.956	1.282	1.00	40.56	H	C
ATOM	6132	C	ALA	24	-23.512	8.074	2.611	1.00	42.05	H	C
ATOM	6133	O	ALA	24	-22.507	7.557	2.119	1.00	42.39	H	O
ATOM	6134	N	SER	25	-24.461	7.367	3.207	1.00	40.96	H	N
ATOM	6135	CA	SER	25	-24.384	5.917	3.243	1.00	40.89	H	C
ATOM	6136	CB	SER	25	-23.532	5.462	4.430	1.00	41.08	H	C
ATOM	6137	OG	SER	25	-24.120	5.876	5.651	1.00	44.95	H	O
ATOM	6138	C	SER	25	-25.779	5.347	3.367	1.00	39.60	H	C
ATOM	6139	O	SER	25	-26.697	6.034	3.815	1.00	38.57	H	O
ATOM	6140	N	GLY	26	-25.935	4.089	2.966	1.00	39.26	H	N
ATOM	6141	CA	GLY	26	-27.200	3.406	3.153	1.00	39.31	H	C
ATOM	6142	C	GLY	26	-28.198	3.613	2.031	1.00	40.96	H	C
ATOM	6143	O	GLY	26	-29.320	3.112	2.103	1.00	43.15	H	O
ATOM	6144	N	PHE	27	-27.811	4.348	0.994	1.00	39.72	H	N
ATOM	6145	CA	PHE	27	-28.725	4.576	-0.117	1.00	39.40	H	C
ATOM	6146	CB	PHE	27	-29.741	5.681	0.240	1.00	39.12	H	C
ATOM	6147	CG	PHE	27	-29.157	7.073	0.282	1.00	37.43	H	C
ATOM	6148	CD1	PHE	27	-28.398	7.494	1.369	1.00	34.06	H	C
ATOM	6149	CD2	PHE	27	-29.386	7.967	-0.760	1.00	34.48	H	C
ATOM	6150	CE1	PHE	27	-27.878	8.781	1.421	1.00	33.04	H	C
ATOM	6151	CE2	PHE	27	-28.867	9.262	-0.716	1.00	34.47	H	C
ATOM	6152	CZ	PHE	27	-28.113	9.669	0.376	1.00	34.29	H	C
ATOM	6153	C	PHE	27	-28.014	4.913	-1.422	1.00	38.19	H	C
ATOM	6154	O	PHE	27	-26.808	5.149	-1.449	1.00	38.12	H	O

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ATOM   6155  N   THR  28     -28.779   4.926  -2.506  1.00 38.67      H    N
ATOM   6156  CA  THR  28     -28.236   5.123  -3.842  1.00 37.98      H    C
ATOM   6157  CB  THR  28     -29.221   4.577  -4.889  1.00 39.03      H    C
ATOM   6158  OG1 THR  28     -29.545   3.222  -4.557  1.00 43.37      H    O
ATOM   6159  CG2 THR  28     -28.613   4.603  -6.272  1.00 39.31      H    C
ATOM   6160  C   THR  28     -27.976   6.609  -4.094  1.00 37.20      H    C
ATOM   6161  O   THR  28     -28.784   7.301  -4.715  1.00 37.60      H    O
ATOM   6162  N   PHE  29     -26.829   7.077  -3.613  1.00 35.10      H    N
ATOM   6163  CA  PHE  29     -26.473   8.491  -3.618  1.00 34.39      H    C
ATOM   6164  CB  PHE  29     -25.055   8.640  -3.060  1.00 32.43      H    C
ATOM   6165  CG  PHE  29     -24.583  10.061  -2.929  1.00 31.23      H    C
ATOM   6166  CD1 PHE  29     -25.109  10.898  -1.956  1.00 31.44      H    C
ATOM   6167  CD2 PHE  29     -23.562  10.541  -3.739  1.00 31.88      H    C
ATOM   6168  CE1 PHE  29     -24.623  12.191  -1.784  1.00 30.12      H    C
ATOM   6169  CE2 PHE  29     -23.070  11.828  -3.577  1.00 30.39      H    C
ATOM   6170  CZ  PHE  29     -23.603  12.656  -2.594  1.00 32.44      H    C
ATOM   6171  C   PHE  29     -26.566   9.143  -5.001  1.00 35.36      H    C
ATOM   6172  O   PHE  29     -26.948  10.305  -5.121  1.00 37.04      H    O
ATOM   6173  N   SER  30     -26.226   8.397  -6.044  1.00 35.18      H    N
ATOM   6174  CA  SER  30     -26.089   8.978  -7.373  1.00 36.29      H    C
ATOM   6175  CB  SER  30     -25.270   8.046  -8.271  1.00 36.47      H    C
ATOM   6176  OG  SER  30     -23.971   7.861  -7.726  1.00 41.17      H    O
ATOM   6177  C   SER  30     -27.425   9.297  -8.041  1.00 35.64      H    C
ATOM   6178  O   SER  30     -27.453   9.892  -9.116  1.00 33.90      H    O
ATOM   6179  N   SER  31     -28.529   8.909  -7.405  1.00 34.26      H    N
ATOM   6180  CA  SER  31     -29.853   9.258  -7.912  1.00 32.93      H    C
ATOM   6181  CB  SER  31     -30.753   8.024  -7.953  1.00 32.80      H    C
ATOM   6182  OG  SER  31     -30.280   7.090  -8.908  1.00 35.66      H    O
ATOM   6183  C   SER  31     -30.522  10.351  -7.087  1.00 32.29      H    C
ATOM   6184  O   SER  31     -31.670  10.719  -7.342  1.00 31.47      H    O
ATOM   6185  N   TYR  32     -29.805  10.875  -6.099  1.00 30.85      H    N
ATOM   6186  CA  TYR  32     -30.367  11.926  -5.261  1.00 29.83      H    C
ATOM   6187  CB  TYR  32     -30.301  11.528  -3.783  1.00 28.34      H    C
ATOM   6188  CG  TYR  32     -31.397  10.571  -3.380  1.00 28.23      H    C
ATOM   6189  CD1 TYR  32     -31.313   9.215  -3.680  1.00 28.84      H    C
ATOM   6190  CE1 TYR  32     -32.340   8.338  -3.340  1.00 28.61      H    C
ATOM   6191  CD2 TYR  32     -32.533  11.028  -2.727  1.00 27.49      H    C
ATOM   6192  CE2 TYR  32     -33.555  10.168  -2.383  1.00 27.94      H    C
ATOM   6193  CZ  TYR  32     -33.458   8.826  -2.690  1.00 30.33      H    C
ATOM   6194  OH  TYR  32     -34.487   7.976  -2.339  1.00 32.86      H    O
ATOM   6195  C   TYR  32     -29.676  13.259  -5.465  1.00 30.11      H    C
ATOM   6196  O   TYR  32     -28.450  13.326  -5.575  1.00 30.60      H    O
ATOM   6197  N   SER  33     -30.475  14.319  -5.529  1.00 28.01      H    N
ATOM   6198  CA  SER  33     -29.953  15.673  -5.405  1.00 28.98      H    C
ATOM   6199  CB  SER  33     -31.006  16.688  -5.854  1.00 28.82      H    C
ATOM   6200  OG  SER  33     -31.407  16.426  -7.194  1.00 32.46      H    O
ATOM   6201  C   SER  33     -29.604  15.902  -3.940  1.00 29.47      H    C
ATOM   6202  O   SER  33     -30.146  15.227  -3.058  1.00 28.61      H    O
ATOM   6203  N   MET  34     -28.702  16.849  -3.687  1.00 29.06      H    N
ATOM   6204  CA  MET  34     -28.261  17.168  -2.332  1.00 29.90      H    C
ATOM   6205  CB  MET  34     -26.813  16.705  -2.138  1.00 29.95      H    C
ATOM   6206  CG  MET  34     -26.657  15.197  -2.193  1.00 31.01      H    C
ATOM   6207  SD  MET  34     -27.587  14.381  -0.852  1.00 34.80      H    S
ATOM   6208  CE  MET  34     -26.780  15.106   0.605  1.00 32.82      H    C
ATOM   6209  C   MET  34     -28.379  18.665  -2.061  1.00 29.84      H    C
ATOM   6210  O   MET  34     -28.388  19.474  -2.994  1.00 29.89      H    O
ATOM   6211  N   ASN  35     -28.479  19.030  -0.786  1.00 26.73      H    N
ATOM   6212  CA  ASN  35     -28.740  20.414  -0.411  1.00 27.72      H    C
ATOM   6213  CB  ASN  35     -30.239  20.625  -0.169  1.00 27.88      H    C
ATOM   6214  CG  ASN  35     -31.098  20.079  -1.298  1.00 31.61      H    C
ATOM   6215  OD1 ASN  35     -31.478  20.814  -2.204  1.00 31.06      H    O
ATOM   6216  ND2 ASN  35     -31.410  18.784  -1.244  1.00 27.46      H    N
ATOM   6217  C   ASN  35     -27.985  20.793   0.863  1.00 29.07      H    C
ATOM   6218  O   ASN  35     -27.755  19.951   1.729  1.00 28.59      H    O
ATOM   6219  N   TRP  36     -27.610  22.064   0.967  1.00 27.79      H    N
ATOM   6220  CA  TRP  36     -27.153  22.623   2.224  1.00 26.46      H    C
ATOM   6221  CB  TRP  36     -25.815  23.351   2.044  1.00 26.13      H    C
ATOM   6222  CG  TRP  36     -24.650  22.430   1.747  1.00 30.50      H    C
ATOM   6223  CD2 TRP  36     -23.918  21.632   2.693  1.00 29.81      H    C
ATOM   6224  CE2 TRP  36     -22.939  20.919   1.969  1.00 29.00      H    C
ATOM   6225  CE3 TRP  36     -23.997  21.455   4.080  1.00 29.79      H    C
ATOM   6226  CD1 TRP  36     -24.093  22.176   0.525  1.00 29.38      H    C
ATOM   6227  NE1 TRP  36     -23.065  21.269   0.651  1.00 31.64      H    N
ATOM   6228  CZ2 TRP  36     -22.046  20.040   2.584  1.00 31.48      H    C
ATOM   6229  CZ3 TRP  36     -23.110  20.581   4.692  1.00 29.03      H    C
ATOM   6230  CH2 TRP  36     -22.148  19.884   3.944  1.00 30.42      H    C
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ATOM   6231  C   TRP  36     -28.209  23.599   2.739  1.00 28.02      H    C
ATOM   6232  O   TRP  36     -28.731  24.440   1.990  1.00 26.76      H    O
ATOM   6233  N   VAL  37     -28.517  23.467   4.025  1.00 26.57      H    N
ATOM   6234  CA  VAL  37     -29.437  24.355   4.719  1.00 26.25      H    C
ATOM   6235  CB  VAL  37     -30.750  23.618   5.054  1.00 25.90      H    C
ATOM   6236  CG1 VAL  37     -31.734  24.565   5.733  1.00 21.86      H    C
ATOM   6237  CG2 VAL  37     -31.343  23.019   3.779  1.00 23.23      H    C
ATOM   6238  C   VAL  37     -28.759  24.761   6.023  1.00 28.43      H    C
ATOM   6239  O   VAL  37     -28.183  23.921   6.715  1.00 29.50      H    O
ATOM   6240  N   ARG  38     -28.828  26.039   6.368  1.00 27.29      H    N
ATOM   6241  CA  ARG  38     -28.145  26.503   7.559  1.00 27.16      H    C
ATOM   6242  CB  ARG  38     -27.031  27.481   7.177  1.00 25.84      H    C
ATOM   6243  CG  ARG  38     -27.488  28.872   6.834  1.00 25.35      H    C
ATOM   6244  CD  ARG  38     -26.289  29.696   6.406  1.00 28.59      H    C
ATOM   6245  NE  ARG  38     -26.672  31.009   5.904  1.00 29.47      H    N
ATOM   6246  CZ  ARG  38     -25.815  31.881   5.381  1.00 29.76      H    C
ATOM   6247  NH1 ARG  38     -24.526  31.577   5.291  1.00 29.29      H    N
ATOM   6248  NH2 ARG  38     -26.247  33.058   4.956  1.00 29.33      H    N
ATOM   6249  C   ARG  38     -29.105  27.149   8.545  1.00 28.31      H    C
ATOM   6250  O   ARG  38     -30.212  27.549   8.177  1.00 27.74      H    O
ATOM   6251  N   GLN  39     -28.681  27.234   9.802  1.00 27.51      H    N
ATOM   6252  CA  GLN  39     -29.518  27.796  10.857  1.00 28.91      H    C
ATOM   6253  CB  GLN  39     -30.293  26.674  11.560  1.00 26.80      H    C
ATOM   6254  CG  GLN  39     -31.269  27.149  12.628  1.00 28.63      H    C
ATOM   6255  CD  GLN  39     -32.223  26.054  13.076  1.00 28.81      H    C
ATOM   6256  OE1 GLN  39     -31.833  24.898  13.240  1.00 30.27      H    O
ATOM   6257  NE2 GLN  39     -33.481  26.415  13.274  1.00 28.47      H    N
ATOM   6258  C   GLN  39     -28.644  28.541  11.866  1.00 31.02      H    C
ATOM   6259  O   GLN  39     -27.842  27.932  12.586  1.00 29.71      H    O
ATOM   6260  N   ALA  40     -28.787  29.860  11.900  1.00 33.52      H    N
ATOM   6261  CA  ALA  40     -28.090  30.675  12.886  1.00 37.63      H    C
ATOM   6262  CB  ALA  40     -28.247  32.156  12.547  1.00 37.07      H    C
ATOM   6263  C   ALA  40     -28.708  30.371  14.247  1.00 40.34      H    C
ATOM   6264  O   ALA  40     -29.901  30.098  14.347  1.00 39.58      H    O
ATOM   6265  N   PRO  41     -27.899  30.402  15.315  1.00 44.34      H    N
ATOM   6266  CD  PRO  41     -26.477  30.793  15.353  1.00 44.86      H    C
ATOM   6267  CA  PRO  41     -28.399  29.989  16.638  1.00 43.86      H    C
ATOM   6268  CB  PRO  41     -27.229  30.292  17.572  1.00 45.74      H    C
ATOM   6269  CG  PRO  41     -26.010  30.221  16.665  1.00 47.13      H    C
ATOM   6270  C   PRO  41     -29.668  30.743  17.032  1.00 43.16      H    C
ATOM   6271  O   PRO  41     -29.700  31.973  16.995  1.00 43.04      H    O
ATOM   6272  N   GLY  42     -30.715  29.997  17.383  1.00 42.32      H    N
ATOM   6273  CA  GLY  42     -31.990  30.609  17.720  1.00 40.30      H    C
ATOM   6274  C   GLY  42     -32.770  31.198  16.550  1.00 40.88      H    C
ATOM   6275  O   GLY  42     -33.703  31.977  16.753  1.00 39.91      H    O
ATOM   6276  N   LYS  43     -32.399  30.839  15.324  1.00 40.15      H    N
ATOM   6277  CA  LYS  43     -33.050  31.402  14.135  1.00 39.69      H    C
ATOM   6278  CB  LYS  43     -32.034  32.183  13.297  1.00 44.93      H    C
ATOM   6279  CG  LYS  43     -31.342  33.322  14.031  1.00 49.72      H    C
ATOM   6280  CD  LYS  43     -32.297  34.476  14.299  1.00 55.35      H    C
ATOM   6281  CE  LYS  43     -31.548  35.700  14.832  1.00 59.19      H    C
ATOM   6282  NZ  LYS  43     -30.434  36.120  13.913  1.00 61.48      H    N
ATOM   6283  C   LYS  43     -33.704  30.327  13.258  1.00 36.98      H    C
ATOM   6284  O   LYS  43     -33.728  29.148  13.611  1.00 34.51      H    O
ATOM   6285  N   GLY  44     -34.233  30.748  12.110  1.00 36.40      H    N
ATOM   6286  CA  GLY  44     -34.940  29.831  11.230  1.00 31.67      H    C
ATOM   6287  C   GLY  44     -34.039  29.115  10.240  1.00 30.70      H    C
ATOM   6288  O   GLY  44     -32.887  29.496  10.049  1.00 31.21      H    O
ATOM   6289  N   LEU  45     -34.562  28.069   9.609  1.00 29.44      H    N
ATOM   6290  CA  LEU  45     -33.834  27.377   8.553  1.00 28.81      H    C
ATOM   6291  CB  LEU  45     -34.583  26.103   8.142  1.00 27.99      H    C
ATOM   6292  CG  LEU  45     -34.801  25.086   9.266  1.00 27.14      H    C
ATOM   6293  CD1 LEU  45     -35.685  23.956   8.765  1.00 26.36      H    C
ATOM   6294  CD2 LEU  45     -33.454  24.551   9.750  1.00 24.68      H    C
ATOM   6295  C   LEU  45     -33.674  28.302   7.348  1.00 28.72      H    C
ATOM   6296  O   LEU  45     -34.572  29.085   7.033  1.00 27.69      H    O
ATOM   6297  N   GLU  46     -32.520  28.219   6.690  1.00 28.43      H    N
ATOM   6298  CA  GLU  46     -32.259  29.005   5.489  1.00 28.83      H    C
ATOM   6299  CB  GLU  46     -31.341  30.193   5.798  1.00 31.44      H    C
ATOM   6300  CG  GLU  46     -31.043  31.069   4.571  1.00 39.18      H    C
ATOM   6301  CD  GLU  46     -30.004  32.169   4.826  1.00 43.02      H    C
ATOM   6302  OE1 GLU  46     -30.027  33.182   4.094  1.00 45.99      H    O
ATOM   6303  OE2 GLU  46     -29.163  32.028   5.743  1.00 43.91      H    O
ATOM   6304  C   GLU  46     -31.597  28.124   4.438  1.00 28.50      H    C
ATOM   6305  O   GLU  46     -30.513  27.586   4.664  1.00 28.35      H    O
ATOM   6306  N   TRP  47     -32.256  27.974   3.291  1.00 26.22      H    N
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ATOM   6307  CA  TRP    47     -31.680  27.234   2.174  1.00 23.96      H    C
ATOM   6308  CB  TRP    47     -32.691  27.135   1.020  1.00 25.33      H    C
ATOM   6309  CG  TRP    47     -32.073  26.630  -0.243  1.00 25.29      H    C
ATOM   6310  CD2 TRP    47     -31.798  27.387  -1.431  1.00 24.26      H    C
ATOM   6311  CE2 TRP    47     -31.161  26.517  -2.342  1.00 24.29      H    C
ATOM   6312  CE3 TRP    47     -32.029  28.713  -1.812  1.00 25.96      H    C
ATOM   6313  CD1 TRP    47     -31.608  25.363  -0.478  1.00 24.30      H    C
ATOM   6314  NE1 TRP    47     -31.058  25.292  -1.737  1.00 26.54      H    N
ATOM   6315  CZ2 TRP    47     -30.751  26.931  -3.610  1.00 23.53      H    C
ATOM   6316  CZ3 TRP    47     -31.623  29.124  -3.073  1.00 25.72      H    C
ATOM   6317  CH2 TRP    47     -30.990  28.234  -3.956  1.00 24.04      H    C
ATOM   6318  C   TRP    47     -30.429  27.949   1.684  1.00 23.89      H    C
ATOM   6319  O   TRP    47     -30.433  29.167   1.520  1.00 23.38      H    O
ATOM   6320  N   VAL    48     -29.370  27.186   1.439  1.00 24.56      H    N
ATOM   6321  CA  VAL    48     -28.082  27.748   1.032  1.00 27.13      H    C
ATOM   6322  CB  VAL    48     -26.929  27.214   1.940  1.00 26.57      H    C
ATOM   6323  CG1 VAL    48     -25.589  27.683   1.408  1.00 25.87      H    C
ATOM   6324  CG2 VAL    48     -27.130  27.696   3.373  1.00 25.68      H    C
ATOM   6325  C   VAL    48     -27.727  27.425  -0.424  1.00 27.12      H    C
ATOM   6326  O   VAL    48     -27.359  28.310  -1.200  1.00 28.60      H    O
ATOM   6327  N   SER    49     -27.829  26.153  -0.789  1.00 25.51      H    N
ATOM   6328  CA  SER    49     -27.422  25.723  -2.113  1.00 27.18      H    C
ATOM   6329  CB  SER    49     -25.893  25.755  -2.204  1.00 27.60      H    C
ATOM   6330  OG  SER    49     -25.445  25.455  -3.514  1.00 30.37      H    O
ATOM   6331  C   SER    49     -27.952  24.317  -2.427  1.00 28.05      H    C
ATOM   6332  O   SER    49     -28.228  23.531  -1.518  1.00 26.28      H    O
ATOM   6333  N   SER    50     -28.101  24.015  -3.716  1.00 28.27      H    N
ATOM   6334  CA  SER    50     -28.581  22.704  -4.154  1.00 28.54      H    C
ATOM   6335  CB  SER    50     -30.061  22.782  -4.550  1.00 28.85      H    C
ATOM   6336  OG  SER    50     -30.882  23.050  -3.418  1.00 32.42      H    O
ATOM   6337  C   SER    50     -27.768  22.205  -5.345  1.00 28.12      H    C
ATOM   6338  O   SER    50     -27.315  22.997  -6.169  1.00 28.29      H    O
ATOM   6339  N   ILE    51     -27.595  20.890  -5.441  1.00 26.82      H    N
ATOM   6340  CA  ILE    51     -26.921  20.303  -6.590  1.00 24.36      H    C
ATOM   6341  CB  ILE    51     -25.419  20.114  -6.302  1.00 26.43      H    C
ATOM   6342  CG2 ILE    51     -25.220  19.159  -5.109  1.00 21.83      H    C
ATOM   6343  CG1 ILE    51     -24.709  19.587  -7.552  1.00 25.90      H    C
ATOM   6344  CD1 ILE    51     -23.194  19.662  -7.452  1.00 27.92      H    C
ATOM   6345  C   ILE    51     -27.552  18.958  -6.956  1.00 26.47      H    C
ATOM   6346  O   ILE    51     -27.751  18.097  -6.096  1.00 24.86      H    O
ATOM   6347  N   SER    52     -27.869  18.783  -8.238  1.00 25.97      H    N
ATOM   6348  CA  SER    52     -28.633  17.624  -8.673  1.00 25.28      H    C
ATOM   6349  CB  SER    52     -29.308  17.912 -10.019  1.00 24.40      H    C
ATOM   6350  OG  SER    52     -28.348  18.107 -11.046  1.00 27.24      H    O
ATOM   6351  C   SER    52     -27.713  16.412  -8.782  1.00 26.88      H    C
ATOM   6352  O   SER    52     -26.494  16.536  -8.650  1.00 28.49      H    O
ATOM   6353  N   SER    53     -28.292  15.240  -9.017  1.00 25.41      H    N
ATOM   6354  CA  SER    53     -27.523  14.007  -8.983  1.00 28.14      H    C
ATOM   6355  CB  SER    53     -28.433  12.811  -9.267  1.00 27.87      H    C
ATOM   6356  OG  SER    53     -29.011  12.911 -10.555  1.00 30.31      H    O
ATOM   6357  C   SER    53     -26.344  14.000  -9.959  1.00 30.57      H    C
ATOM   6358  O   SER    53     -25.284  13.447  -9.645  1.00 32.17      H    O
ATOM   6359  N   SER    54     -26.517  14.606 -11.135  1.00 28.22      H    N
ATOM   6360  CA  SER    54     -25.437  14.649 -12.123  1.00 28.16      H    C
ATOM   6361  CB  SER    54     -25.966  14.259 -13.520  1.00 27.82      H    C
ATOM   6362  OG  SER    54     -26.898  15.215 -14.023  1.00 29.46      H    O
ATOM   6363  C   SER    54     -24.766  16.027 -12.185  1.00 27.90      H    C
ATOM   6364  O   SER    54     -23.979  16.302 -13.087  1.00 28.59      H    O
ATOM   6365  N   SER    55     -25.087  16.882 -11.220  1.00 28.14      H    N
ATOM   6366  CA  SER    55     -24.520  18.230 -11.124  1.00 30.31      H    C
ATOM   6367  CB  SER    55     -22.987  18.166 -11.092  1.00 31.88      H    C
ATOM   6368  OG  SER    55     -22.526  17.349 -10.023  1.00 34.17      H    O
ATOM   6369  C   SER    55     -24.968  19.180 -12.244  1.00 30.07      H    C
ATOM   6370  O   SER    55     -24.353  20.230 -12.457  1.00 29.82      H    O
ATOM   6371  N   SER    56     -26.039  18.828 -12.950  1.00 27.46      H    N
ATOM   6372  CA  SER    56     -26.501  19.654 -14.066  1.00 27.70      H    C
ATOM   6373  CB  SER    56     -27.098  18.772 -15.174  1.00 29.06      H    C
ATOM   6374  OG  SER    56     -28.269  18.101 -14.734  1.00 34.97      H    O
ATOM   6375  C   SER    56     -27.505  20.727 -13.637  1.00 27.78      H    C
ATOM   6376  O   SER    56     -27.805  21.642 -14.406  1.00 27.41      H    O
ATOM   6377  N   TYR    57     -28.028  20.611 -12.413  1.00 27.64      H    N
ATOM   6378  CA  TYR    57     -28.685  21.738 -11.742  1.00 27.77      H    C
ATOM   6379  CB  TYR    57     -30.107  21.369 -11.294  1.00 27.25      H    C
ATOM   6380  CG  TYR    57     -31.053  20.945 -12.400  1.00 29.87      H    C
ATOM   6381  CD1 TYR    57     -31.126  19.616 -12.798  1.00 28.73      H    C
ATOM   6382  CE1 TYR    57     -32.050  19.199 -13.735  1.00 30.08      H    C
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ATOM	6383	CD2	TYR	57	-31.933	21.857	-12.984	1.00	28.13	H	C
ATOM	6384	CE2	TYR	57	-32.861	21.448	-13.929	1.00	29.41	H	C
ATOM	6385	CZ	TYR	57	-32.917	20.113	-14.294	1.00	30.49	H	C
ATOM	6386	OH	TYR	57	-33.861	19.669	-15.193	1.00	29.95	H	O
ATOM	6387	C	TYR	57	-27.881	22.156	-10.504	1.00	29.02	H	C
ATOM	6388	O	TYR	57	-27.628	21.344	-9.598	1.00	29.53	H	O
ATOM	6389	N	ILE	58	-27.487	23.424	-10.469	1.00	29.50	H	N
ATOM	6390	CA	ILE	58	-26.789	24.003	-9.325	1.00	29.52	H	C
ATOM	6391	CB	ILE	58	-25.270	24.148	-9.614	1.00	30.72	H	C
ATOM	6392	CG2	ILE	58	-24.589	24.915	-8.498	1.00	29.96	H	C
ATOM	6393	CG1	ILE	58	-24.627	22.765	-9.740	1.00	30.72	H	C
ATOM	6394	CD1	ILE	58	-23.154	22.795	-10.092	1.00	28.84	H	C
ATOM	6395	C	ILE	58	-27.392	25.380	-9.067	1.00	29.74	H	C
ATOM	6396	O	ILE	58	-27.609	26.145	-10.003	1.00	30.44	H	O
ATOM	6397	N	SER	59	-27.697	25.685	-7.809	1.00	27.57	H	N
ATOM	6398	CA	SER	59	-28.161	27.021	-7.458	1.00	29.06	H	C
ATOM	6399	CB	SER	59	-29.680	27.113	-7.616	1.00	28.15	H	C
ATOM	6400	OG	SER	59	-30.321	26.100	-6.874	1.00	34.72	H	O
ATOM	6401	C	SER	59	-27.758	27.454	-6.044	1.00	28.73	H	C
ATOM	6402	O	SER	59	-27.358	26.634	-5.213	1.00	29.00	H	O
ATOM	6403	N	TYR	60	-27.874	28.754	-5.791	1.00	27.30	H	N
ATOM	6404	CA	TYR	60	-27.374	29.366	-4.571	1.00	27.26	H	C
ATOM	6405	CB	TYR	60	-26.039	30.073	-4.834	1.00	25.88	H	C
ATOM	6406	CG	TYR	60	-24.917	29.156	-5.258	1.00	25.19	H	C
ATOM	6407	CD1	TYR	60	-24.210	28.421	-4.315	1.00	23.78	H	C
ATOM	6408	CE1	TYR	60	-23.190	27.572	-4.687	1.00	22.85	H	C
ATOM	6409	CD2	TYR	60	-24.567	29.018	-6.603	1.00	23.48	H	C
ATOM	6410	CE2	TYR	60	-23.543	28.171	-6.988	1.00	21.69	H	C
ATOM	6411	CZ	TYR	60	-22.859	27.447	-6.021	1.00	23.85	H	C
ATOM	6412	OH	TYR	60	-21.854	26.575	-6.377	1.00	24.72	H	O
ATOM	6413	C	TYR	60	-28.379	30.397	-4.087	1.00	28.79	H	C
ATOM	6414	O	TYR	60	-29.032	31.059	-4.894	1.00	27.21	H	O
ATOM	6415	N	ALA	61	-28.499	30.528	-2.770	1.00	28.73	H	N
ATOM	6416	CA	ALA	61	-29.203	31.657	-2.180	1.00	30.11	H	C
ATOM	6417	CB	ALA	61	-29.264	31.491	-0.658	1.00	28.64	H	C
ATOM	6418	C	ALA	61	-28.426	32.926	-2.544	1.00	31.88	H	C
ATOM	6419	O	ALA	61	-27.210	32.883	-2.702	1.00	29.32	H	O
ATOM	6420	N	ASP	62	-29.127	34.046	-2.683	1.00	35.01	H	N
ATOM	6421	CA	ASP	62	-28.475	35.314	-2.995	1.00	40.62	H	C
ATOM	6422	CB	ASP	62	-29.500	36.444	-2.997	1.00	47.18	H	C
ATOM	6423	CG	ASP	62	-29.915	36.842	-4.390	1.00	55.90	H	C
ATOM	6424	OD1	ASP	62	-31.131	36.750	-4.693	1.00	59.73	H	O
ATOM	6425	OD2	ASP	62	-29.023	37.246	-5.177	1.00	57.97	H	O
ATOM	6426	C	ASP	62	-27.366	35.666	-2.007	1.00	41.05	H	C
ATOM	6427	O	ASP	62	-26.320	36.181	-2.394	1.00	41.21	H	O
ATOM	6428	N	SER	63	-27.606	35.387	-0.731	1.00	40.70	H	N
ATOM	6429	CA	SER	63	-26.704	35.804	0.337	1.00	41.03	H	C
ATOM	6430	CB	SER	63	-27.354	35.546	1.695	1.00	41.93	H	C
ATOM	6431	OG	SER	63	-27.781	34.199	1.795	1.00	42.20	H	O
ATOM	6432	C	SER	63	-25.333	35.131	0.302	1.00	40.04	H	C
ATOM	6433	O	SER	63	-24.396	35.606	0.943	1.00	42.70	H	O
ATOM	6434	N	VAL	64	-25.206	34.033	-0.434	1.00	37.58	H	N
ATOM	6435	CA	VAL	64	-23.923	33.342	-0.516	1.00	35.50	H	C
ATOM	6436	CB	VAL	64	-24.023	31.883	-0.010	1.00	34.20	H	C
ATOM	6437	CG1	VAL	64	-24.635	31.851	1.399	1.00	32.45	H	C
ATOM	6438	CG2	VAL	64	-24.844	31.054	-0.987	1.00	32.80	H	C
ATOM	6439	C	VAL	64	-23.386	33.309	-1.941	1.00	36.79	H	C
ATOM	6440	O	VAL	64	-22.329	32.726	-2.203	1.00	35.31	H	O
ATOM	6441	N	LYS	65	-24.117	33.925	-2.862	1.00	37.48	H	N
ATOM	6442	CA	LYS	65	-23.742	33.884	-4.272	1.00	42.50	H	C
ATOM	6443	CB	LYS	65	-24.766	34.661	-5.106	1.00	46.64	H	C
ATOM	6444	CG	LYS	65	-24.604	34.500	-6.605	1.00	52.44	H	C
ATOM	6445	CD	LYS	65	-25.762	33.691	-7.189	1.00	57.69	H	C
ATOM	6446	CE	LYS	65	-27.105	34.302	-6.786	1.00	60.32	H	C
ATOM	6447	NZ	LYS	65	-28.261	33.402	-7.060	1.00	62.20	H	N
ATOM	6448	C	LYS	65	-22.354	34.494	-4.465	1.00	42.87	H	C
ATOM	6449	O	LYS	65	-22.077	35.593	-3.978	1.00	43.54	H	O
ATOM	6450	N	GLY	66	-21.484	33.780	-5.171	1.00	42.33	H	N
ATOM	6451	CA	GLY	66	-20.156	34.301	-5.437	1.00	42.49	H	C
ATOM	6452	C	GLY	66	-19.126	33.988	-4.361	1.00	44.27	H	C
ATOM	6453	O	GLY	66	-17.926	34.142	-4.589	1.00	44.84	H	O
ATOM	6454	N	ARG	67	-19.580	33.549	-3.189	1.00	41.56	H	N
ATOM	6455	CA	ARG	67	-18.663	33.212	-2.108	1.00	39.45	H	C
ATOM	6456	CB	ARG	67	-19.063	33.955	-0.830	1.00	40.38	H	C
ATOM	6457	CG	ARG	67	-19.033	35.478	-0.968	1.00	39.81	H	C
ATOM	6458	CD	ARG	67	-19.250	36.182	0.366	1.00	39.55	H	C

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ATOM   6459  NE  ARG    67     -20.572  35.918   0.933  1.00 39.73      H    N
ATOM   6460  CZ  ARG    67     -20.786  35.246   2.064  1.00 39.95      H    C
ATOM   6461  NH1 ARG    67     -19.760  34.763   2.760  1.00 36.21      H    N
ATOM   6462  NH2 ARG    67     -22.028  35.057   2.498  1.00 37.67      H    N
ATOM   6463  C   ARG    67     -18.616  31.710  -1.849  1.00 38.91      H    C
ATOM   6464  O   ARG    67     -17.549  31.160  -1.566  1.00 38.47      H    O
ATOM   6465  N   PHE    68     -19.769  31.046  -1.951  1.00 37.70      H    N
ATOM   6466  CA  PHE    68     -19.846  29.603  -1.714  1.00 34.66      H    C
ATOM   6467  CB  PHE    68     -21.092  29.247  -0.889  1.00 33.55      H    C
ATOM   6468  CG  PHE    68     -21.064  29.759   0.530  1.00 34.88      H    C
ATOM   6469  CD1 PHE    68     -22.001  29.316   1.453  1.00 34.07      H    C
ATOM   6470  CD2 PHE    68     -20.120  30.692   0.937  1.00 34.38      H    C
ATOM   6471  CE1 PHE    68     -22.004  29.791   2.747  1.00 33.89      H    C
ATOM   6472  CE2 PHE    68     -20.115  31.176   2.238  1.00 34.40      H    C
ATOM   6473  CZ  PHE    68     -21.058  30.725   3.143  1.00 35.06      H    C
ATOM   6474  C   PHE    68     -19.895  28.840  -3.029  1.00 34.39      H    C
ATOM   6475  O   PHE    68     -20.485  29.302  -4.006  1.00 33.33      H    O
ATOM   6476  N   THR    69     -19.275  27.666  -3.045  1.00 33.03      H    N
ATOM   6477  CA  THR    69     -19.317  26.787  -4.204  1.00 33.88      H    C
ATOM   6478  CB  THR    69     -17.954  26.710  -4.911  1.00 34.15      H    C
ATOM   6479  OG1 THR    69     -17.583  28.010  -5.384  1.00 36.58      H    O
ATOM   6480  CG2 THR    69     -18.026  25.739  -6.088  1.00 33.48      H    C
ATOM   6481  C   THR    69     -19.680  25.379  -3.757  1.00 34.59      H    C
ATOM   6482  O   THR    69     -18.985  24.786  -2.930  1.00 34.69      H    O
ATOM   6483  N   ILE    70     -20.758  24.841  -4.315  1.00 32.65      H    N
ATOM   6484  CA  ILE    70     -21.203  23.503  -3.957  1.00 31.01      H    C
ATOM   6485  CB  ILE    70     -22.754  23.432  -3.936  1.00 30.63      H    C
ATOM   6486  CG2 ILE    70     -23.308  23.592  -5.351  1.00 29.46      H    C
ATOM   6487  CG1 ILE    70     -23.219  22.111  -3.329  1.00 29.67      H    C
ATOM   6488  CD1 ILE    70     -24.719  22.090  -3.004  1.00 27.57      H    C
ATOM   6489  C   ILE    70     -20.635  22.534  -4.987  1.00 31.28      H    C
ATOM   6490  O   ILE    70     -20.373  22.914  -6.122  1.00 33.97      H    O
ATOM   6491  N   SER    71     -20.406  21.293  -4.582  1.00 31.56      H    N
ATOM   6492  CA  SER    71     -20.011  20.255  -5.522  1.00 32.33      H    C
ATOM   6493  CB  SER    71     -18.547  20.427  -5.947  1.00 33.86      H    C
ATOM   6494  OG  SER    71     -17.673  20.291  -4.839  1.00 37.91      H    O
ATOM   6495  C   SER    71     -20.195  18.902  -4.861  1.00 32.41      H    C
ATOM   6496  O   SER    71     -20.456  18.821  -3.660  1.00 31.10      H    O
ATOM   6497  N   ARG    72     -20.066  17.843  -5.648  1.00 31.25      H    N
ATOM   6498  CA  ARG    72     -20.285  16.500  -5.140  1.00 32.41      H    C
ATOM   6499  CB  ARG    72     -21.747  16.092  -5.343  1.00 30.85      H    C
ATOM   6500  CG  ARG    72     -22.194  16.055  -6.802  1.00 29.48      H    C
ATOM   6501  CD  ARG    72     -23.691  15.820  -6.912  1.00 28.57      H    C
ATOM   6502  NE  ARG    72     -24.075  14.470  -6.497  1.00 28.84      H    N
ATOM   6503  CZ  ARG    72     -25.264  14.152  -5.989  1.00 27.19      H    C
ATOM   6504  NH1 ARG    72     -26.197  15.086  -5.828  1.00 26.26      H    N
ATOM   6505  NH2 ARG    72     -25.525  12.900  -5.641  1.00 26.57      H    N
ATOM   6506  C   ARG    72     -19.368  15.530  -5.860  1.00 34.09      H    C
ATOM   6507  O   ARG    72     -18.870  15.821  -6.944  1.00 33.62      H    O
ATOM   6508  N   ASP    73     -19.136  14.379  -5.243  1.00 38.28      H    N
ATOM   6509  CA  ASP    73     -18.327  13.325  -5.849  1.00 39.48      H    C
ATOM   6510  CB  ASP    73     -16.947  13.281  -5.179  1.00 41.60      H    C
ATOM   6511  CG  ASP    73     -15.990  12.292  -5.843  1.00 44.95      H    C
ATOM   6512  OD1 ASP    73     -16.426  11.183  -6.225  1.00 45.18      H    O
ATOM   6513  OD2 ASP    73     -14.791  12.630  -5.974  1.00 46.94      H    O
ATOM   6514  C   ASP    73     -19.066  12.013  -5.624  1.00 39.78      H    C
ATOM   6515  O   ASP    73     -18.978  11.423  -4.545  1.00 40.60      H    O
ATOM   6516  N   ASN    74     -19.797  11.560  -6.639  1.00 38.88      H    N
ATOM   6517  CA  ASN    74     -20.653  10.388  -6.486  1.00 39.67      H    C
ATOM   6518  CB  ASN    74     -21.523  10.195  -7.732  1.00 37.88      H    C
ATOM   6519  CG  ASN    74     -22.648  11.215  -7.821  1.00 38.36      H    C
ATOM   6520  OD1 ASN    74     -22.954  11.915  -6.848  1.00 38.42      H    O
ATOM   6521  ND2 ASN    74     -23.271  11.303  -8.989  1.00 35.16      H    N
ATOM   6522  C   ASN    74     -19.865   9.115  -6.212  1.00 41.22      H    C
ATOM   6523  O   ASN    74     -20.336   8.230  -5.499  1.00 42.03      H    O
ATOM   6524  N   ALA    75     -18.667   9.017  -6.780  1.00 41.81      H    N
ATOM   6525  CA  ALA    75     -17.819   7.852  -6.554  1.00 44.00      H    C
ATOM   6526  CB  ALA    75     -16.553   7.948  -7.410  1.00 42.83      H    C
ATOM   6527  C   ALA    75     -17.450   7.748  -5.074  1.00 45.21      H    C
ATOM   6528  O   ALA    75     -17.252   6.654  -4.550  1.00 45.78      H    O
ATOM   6529  N   LYS    76     -17.368   8.892  -4.402  1.00 46.01      H    N
ATOM   6530  CA  LYS    76     -17.043   8.916  -2.982  1.00 47.14      H    C
ATOM   6531  CB  LYS    76     -16.034  10.029  -2.696  1.00 49.80      H    C
ATOM   6532  CG  LYS    76     -14.694   9.846  -3.389  1.00 52.73      H    C
ATOM   6533  CD  LYS    76     -13.726  10.958  -3.012  1.00 56.34      H    C
ATOM   6534  CE  LYS    76     -12.383  10.797  -3.724  1.00 58.48      H    C
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ATOM   6535  NZ  LYS   76     -11.483  11.966  -3.476  1.00 60.20      H    N
ATOM   6536  C   LYS   76     -18.271   9.107  -2.090  1.00 47.47      H    C
ATOM   6537  O   LYS   76     -18.141   9.232  -0.875  1.00 47.19      H    O
ATOM   6538  N   ASN   77     -19.460   9.136  -2.690  1.00 46.65      H    N
ATOM   6539  CA  ASN   77     -20.589   9.376  -1.933  1.00 44.74      H    C
ATOM   6540  CB  ASN   77     -21.023   8.166  -1.052  1.00 45.89      H    C
ATOM   6541  CG  ASN   77     -21.302   6.913  -1.860  1.00 47.96      H    C
ATOM   6542  OD1 ASN   77     -22.410   6.709  -2.352  1.00 49.54      H    O
ATOM   6543  ND2 ASN   77     -20.292   6.065  -2.000  1.00 50.93      H    N
ATOM   6544  C   ASN   77     -20.545  10.617  -1.050  1.00 42.51      H    C
ATOM   6545  O   ASN   77     -20.883  10.588   0.134  1.00 42.35      H    O
ATOM   6546  N   SER   78     -20.039  11.704  -1.621  1.00 39.28      H    N
ATOM   6547  CA  SER   78     -19.787  12.898  -0.836  1.00 38.34      H    C
ATOM   6548  CB  SER   78     -18.291  13.032  -0.565  1.00 40.48      H    C
ATOM   6549  OG  SER   78     -17.830  11.912   0.169  1.00 46.08      H    O
ATOM   6550  C   SER   78     -20.310  14.186  -1.456  1.00 37.03      H    C
ATOM   6551  O   SER   78     -20.320  14.350  -2.678  1.00 37.05      H    O
ATOM   6552  N   LEU   79     -20.733  15.093  -0.580  1.00 33.76      H    N
ATOM   6553  CA  LEU   79     -21.212  16.417  -0.945  1.00 33.07      H    C
ATOM   6554  CB  LEU   79     -22.642  16.599  -0.412  1.00 30.60      H    C
ATOM   6555  CG  LEU   79     -23.219  18.015  -0.350  1.00 29.82      H    C
ATOM   6556  CD1 LEU   79     -23.487  18.523  -1.763  1.00 28.74      H    C
ATOM   6557  CD2 LEU   79     -24.504  18.008   0.470  1.00 26.94      H    C
ATOM   6558  C   LEU   79     -20.263  17.436  -0.294  1.00 33.73      H    C
ATOM   6559  O   LEU   79     -19.834  17.237   0.844  1.00 35.43      H    O
ATOM   6560  N   TYR   80     -19.931  18.512  -1.003  1.00 31.34      H    N
ATOM   6561  CA  TYR   80     -19.001  19.507  -0.479  1.00 30.56      H    C
ATOM   6562  CB  TYR   80     -17.698  19.500  -1.274  1.00 32.57      H    C
ATOM   6563  CG  TYR   80     -17.067  18.136  -1.386  1.00 35.32      H    C
ATOM   6564  CD1 TYR   80     -17.045  17.465  -2.599  1.00 36.45      H    C
ATOM   6565  CE1 TYR   80     -16.486  16.208  -2.711  1.00 38.72      H    C
ATOM   6566  CD2 TYR   80     -16.507  17.510  -0.275  1.00 36.23      H    C
ATOM   6567  CE2 TYR   80     -15.945  16.250  -0.374  1.00 37.75      H    C
ATOM   6568  CZ  TYR   80     -15.938  15.604  -1.598  1.00 40.42      H    C
ATOM   6569  OH  TYR   80     -15.387  14.349  -1.721  1.00 42.98      H    O
ATOM   6570  C   TYR   80     -19.574  20.909  -0.508  1.00 31.38      H    C
ATOM   6571  O   TYR   80     -20.414  21.226  -1.336  1.00 31.59      H    O
ATOM   6572  N   LEU   81     -19.118  21.749   0.413  1.00 32.68      H    N
ATOM   6573  CA  LEU   81     -19.394  23.172   0.341  1.00 33.13      H    C
ATOM   6574  CB  LEU   81     -20.476  23.573   1.350  1.00 30.32      H    C
ATOM   6575  CG  LEU   81     -20.966  25.018   1.180  1.00 31.03      H    C
ATOM   6576  CD1 LEU   81     -21.629  25.162  -0.186  1.00 30.17      H    C
ATOM   6577  CD2 LEU   81     -21.955  25.387   2.279  1.00 29.51      H    C
ATOM   6578  C   LEU   81     -18.106  23.941   0.633  1.00 35.97      H    C
ATOM   6579  O   LEU   81     -17.618  23.955   1.770  1.00 35.94      H    O
ATOM   6580  N   GLN   82     -17.558  24.571  -0.400  1.00 35.80      H    N
ATOM   6581  CA  GLN   82     -16.424  25.463  -0.235  1.00 37.59      H    C
ATOM   6582  CB  GLN   82     -15.592  25.501  -1.521  1.00 38.62      H    C
ATOM   6583  CG  GLN   82     -14.362  26.406  -1.448  1.00 42.10      H    C
ATOM   6584  CD  GLN   82     -13.374  25.975  -0.368  1.00 43.68      H    C
ATOM   6585  OE1 GLN   82     -12.986  24.805  -0.292  1.00 43.76      H    O
ATOM   6586  NE2 GLN   82     -12.968  26.922   0.473  1.00 43.92      H    N
ATOM   6587  C   GLN   82     -16.940  26.854   0.099  1.00 38.25      H    C
ATOM   6588  O   GLN   82     -17.640  27.472  -0.698  1.00 39.65      H    O
ATOM   6589  N   MET   83     -16.600  27.341   1.289  1.00 40.33      H    N
ATOM   6590  CA  MET   83     -17.043  28.657   1.733  1.00 42.26      H    C
ATOM   6591  CB  MET   83     -17.682  28.557   3.120  1.00 42.53      H    C
ATOM   6592  CG  MET   83     -18.815  27.538   3.240  1.00 46.08      H    C
ATOM   6593  SD  MET   83     -19.506  27.440   4.929  1.00 49.67      H    S
ATOM   6594  CE  MET   83     -18.162  26.710   5.786  1.00 50.01      H    C
ATOM   6595  C   MET   83     -15.868  29.634   1.785  1.00 43.58      H    C
ATOM   6596  O   MET   83     -14.917  29.441   2.545  1.00 45.85      H    O
ATOM   6597  N   ASN   84     -15.942  30.686   0.979  1.00 42.66      H    N
ATOM   6598  CA  ASN   84     -14.899  31.699   0.945  1.00 42.04      H    C
ATOM   6599  CB  ASN   84     -14.353  31.857  -0.472  1.00 42.61      H    C
ATOM   6600  CG  ASN   84     -13.545  30.664  -0.917  1.00 45.11      H    C
ATOM   6601  OD1 ASN   84     -13.083  29.866  -0.098  1.00 46.92      H    O
ATOM   6602  ND2 ASN   84     -13.368  30.529  -2.224  1.00 47.46      H    N
ATOM   6603  C   ASN   84     -15.443  33.032   1.416  1.00 42.87      H    C
ATOM   6604  O   ASN   84     -16.657  33.229   1.478  1.00 41.82      H    O
ATOM   6605  N   SER   85     -14.532  33.941   1.752  1.00 42.16      H    N
ATOM   6606  CA  SER   85     -14.898  35.288   2.153  1.00 41.46      H    C
ATOM   6607  CB  SER   85     -15.383  36.077   0.938  1.00 43.66      H    C
ATOM   6608  OG  SER   85     -14.432  36.002  -0.110  1.00 47.73      H    O
ATOM   6609  C   SER   85     -15.981  35.268   3.217  1.00 40.30      H    C
ATOM   6610  O   SER   85     -16.959  36.020   3.135  1.00 39.19      H    O
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ATOM	6611	N	LEU	86	-15.798	34.409	4.217	1.00	39.13	H	N
ATOM	6612	CA	LEU	86	-16.785	34.255	5.281	1.00	39.06	H	C
ATOM	6613	CB	LEU	86	-16.347	33.159	6.245	1.00	37.15	H	C
ATOM	6614	CG	LEU	86	-16.565	31.763	5.669	1.00	38.36	H	C
ATOM	6615	CD1	LEU	86	-15.834	30.724	6.492	1.00	37.27	H	C
ATOM	6616	CD2	LEU	86	-18.054	31.485	5.628	1.00	37.79	H	C
ATOM	6617	C	LEU	86	-17.039	35.540	6.051	1.00	39.98	H	C
ATOM	6618	O	LEU	86	-16.138	36.359	6.235	1.00	41.95	H	O
ATOM	6619	N	ARG	87	-18.280	35.714	6.489	1.00	40.79	H	N
ATOM	6620	CA	ARG	87	-18.662	36.863	7.296	1.00	43.46	H	C
ATOM	6621	CB	ARG	87	-19.635	37.754	6.532	1.00	45.67	H	C
ATOM	6622	CG	ARG	87	-19.140	38.192	5.182	1.00	49.88	H	C
ATOM	6623	CD	ARG	87	-20.293	38.703	4.351	1.00	53.41	H	C
ATOM	6624	NE	ARG	87	-19.920	38.894	2.955	1.00	57.94	H	N
ATOM	6625	CZ	ARG	87	-20.781	39.233	2.001	1.00	59.50	H	C
ATOM	6626	NH1	ARG	87	-20.366	39.389	0.750	1.00	59.64	H	N
ATOM	6627	NH2	ARG	87	-22.062	39.410	2.303	1.00	58.79	H	N
ATOM	6628	C	ARG	87	-19.329	36.387	8.577	1.00	43.92	H	C
ATOM	6629	O	ARG	87	-19.737	35.227	8.681	1.00	42.74	H	O
ATOM	6630	N	ALA	88	-19.444	37.291	9.544	1.00	44.62	H	N
ATOM	6631	CA	ALA	88	-20.030	36.960	10.838	1.00	45.48	H	C
ATOM	6632	CB	ALA	88	-20.133	38.222	11.699	1.00	44.63	H	C
ATOM	6633	C	ALA	88	-21.413	36.335	10.649	1.00	44.68	H	C
ATOM	6634	O	ALA	88	-21.762	35.364	11.320	1.00	43.24	H	O
ATOM	6635	N	GLU	89	-22.180	36.900	9.718	1.00	45.01	H	N
ATOM	6636	CA	GLU	89	-23.546	36.460	9.441	1.00	44.88	H	C
ATOM	6637	CB	GLU	89	-24.219	37.419	8.453	1.00	49.04	H	C
ATOM	6638	CG	GLU	89	-24.223	38.878	8.894	1.00	58.20	H	C
ATOM	6639	CD	GLU	89	-22.918	39.596	8.569	1.00	62.22	H	C
ATOM	6640	OE1	GLU	89	-22.376	40.289	9.461	1.00	64.59	H	O
ATOM	6641	OE2	GLU	89	-22.441	39.466	7.419	1.00	64.36	H	O
ATOM	6642	C	GLU	89	-23.625	35.041	8.883	1.00	41.48	H	C
ATOM	6643	O	GLU	89	-24.703	34.461	8.822	1.00	40.85	H	O
ATOM	6644	N	ASP	90	-22.491	34.487	8.465	1.00	38.97	H	N
ATOM	6645	CA	ASP	90	-22.451	33.104	8.003	1.00	36.93	H	C
ATOM	6646	CB	ASP	90	-21.268	32.884	7.058	1.00	38.32	H	C
ATOM	6647	CG	ASP	90	-21.391	33.687	5.772	1.00	41.49	H	C
ATOM	6648	OD1	ASP	90	-22.526	33.850	5.265	1.00	41.50	H	O
ATOM	6649	OD2	ASP	90	-20.348	34.160	5.272	1.00	42.86	H	O
ATOM	6650	C	ASP	90	-22.357	32.120	9.162	1.00	36.53	H	C
ATOM	6651	O	ASP	90	-22.304	30.907	8.951	1.00	37.19	H	O
ATOM	6652	N	THR	91	-22.334	32.643	10.385	1.00	34.31	H	N
ATOM	6653	CA	THR	91	-22.252	31.797	11.572	1.00	34.08	H	C
ATOM	6654	CB	THR	91	-22.036	32.656	12.849	1.00	34.84	H	C
ATOM	6655	OG1	THR	91	-20.738	33.263	12.799	1.00	36.83	H	O
ATOM	6656	CG2	THR	91	-22.160	31.804	14.109	1.00	31.37	H	C
ATOM	6657	C	THR	91	-23.553	31.019	11.710	1.00	32.40	H	C
ATOM	6658	O	THR	91	-24.628	31.613	11.763	1.00	32.80	H	O
ATOM	6659	N	ALA	92	-23.465	29.695	11.774	1.00	29.43	H	N
ATOM	6660	CA	ALA	92	-24.674	28.884	11.827	1.00	29.39	H	C
ATOM	6661	CB	ALA	92	-25.551	29.177	10.601	1.00	31.93	H	C
ATOM	6662	C	ALA	92	-24.334	27.409	11.867	1.00	28.87	H	C
ATOM	6663	O	ALA	92	-23.197	27.025	11.599	1.00	28.15	H	O
ATOM	6664	N	VAL	93	-25.321	26.581	12.202	1.00	28.51	H	N
ATOM	6665	CA	VAL	93	-25.210	25.145	11.954	1.00	28.45	H	C
ATOM	6666	CB	VAL	93	-26.178	24.339	12.851	1.00	30.44	H	C
ATOM	6667	CG1	VAL	93	-26.090	22.866	12.503	1.00	29.75	H	C
ATOM	6668	CG2	VAL	93	-25.836	24.558	14.344	1.00	31.02	H	C
ATOM	6669	C	VAL	93	-25.562	24.877	10.485	1.00	30.00	H	C
ATOM	6670	O	VAL	93	-26.553	25.403	9.969	1.00	30.05	H	O
ATOM	6671	N	TYR	94	-24.741	24.077	9.812	1.00	29.21	H	N
ATOM	6672	CA	TYR	94	-24.997	23.711	8.424	1.00	29.79	H	C
ATOM	6673	CB	TYR	94	-23.742	23.945	7.570	1.00	29.58	H	C
ATOM	6674	CG	TYR	94	-23.442	25.410	7.332	1.00	32.16	H	C
ATOM	6675	CD1	TYR	94	-23.617	25.981	6.074	1.00	31.93	H	C
ATOM	6676	CE1	TYR	94	-23.410	27.336	5.865	1.00	30.72	H	C
ATOM	6677	CD2	TYR	94	-23.041	26.237	8.379	1.00	31.28	H	C
ATOM	6678	CE2	TYR	94	-22.831	27.592	8.182	1.00	30.20	H	C
ATOM	6679	CZ	TYR	94	-23.022	28.137	6.922	1.00	32.69	H	C
ATOM	6680	OH	TYR	94	-22.863	29.495	6.727	1.00	32.53	H	O
ATOM	6681	C	TYR	94	-25.424	22.249	8.320	1.00	29.99	H	C
ATOM	6682	O	TYR	94	-24.688	21.352	8.750	1.00	30.54	H	O
ATOM	6683	N	PHE	95	-26.610	22.019	7.754	1.00	29.58	H	N
ATOM	6684	CA	PHE	95	-27.123	20.666	7.512	1.00	29.84	H	C
ATOM	6685	CB	PHE	95	-28.601	20.557	7.905	1.00	27.29	H	C
ATOM	6686	CG	PHE	95	-28.900	20.937	9.334	1.00	30.50	H	C

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ATOM   6687  CD1 PHE    95     -28.783  20.004  10.357  1.00 30.20      H    C
ATOM   6688  CD2 PHE    95     -29.336  22.218   9.651  1.00 30.58      H    C
ATOM   6689  CE1 PHE    95     -29.098  20.343  11.671  1.00 27.42      H    C
ATOM   6690  CE2 PHE    95     -29.652  22.560  10.960  1.00 29.43      H    C
ATOM   6691  CZ  PHE    95     -29.531  21.619  11.968  1.00 27.23      H    C
ATOM   6692  C   PHE    95     -27.016  20.347   6.023  1.00 31.72      H    C
ATOM   6693  O   PHE    95     -27.203  21.240   5.185  1.00 30.65      H    O
ATOM   6694  N   CYS    96     -26.734  19.085   5.694  1.00 32.57      H    N
ATOM   6695  CA  CYS    96     -27.051  18.566   4.366  1.00 35.64      H    C
ATOM   6696  C   CYS    96     -28.344  17.753   4.388  1.00 34.02      H    C
ATOM   6697  O   CYS    96     -28.718  17.185   5.414  1.00 33.53      H    O
ATOM   6698  CB  CYS    96     -25.898  17.710   3.784  1.00 41.97      H    C
ATOM   6699  SG  CYS    96     -25.132  16.445   4.858  1.00 55.63      H    S
ATOM   6700  N   ALA    97     -29.028  17.710   3.249  1.00 30.78      H    N
ATOM   6701  CA  ALA    97     -30.273  16.962   3.134  1.00 30.15      H    C
ATOM   6702  CB  ALA    97     -31.450  17.824   3.582  1.00 28.28      H    C
ATOM   6703  C   ALA    97     -30.457  16.540   1.690  1.00 29.71      H    C
ATOM   6704  O   ALA    97     -30.050  17.252   0.768  1.00 31.75      H    O
ATOM   6705  N   ARG    98     -31.067  15.382   1.492  1.00 27.73      H    N
ATOM   6706  CA  ARG    98     -31.226  14.840   0.155  1.00 28.67      H    C
ATOM   6707  CB  ARG    98     -30.979  13.334   0.164  1.00 28.08      H    C
ATOM   6708  CG  ARG    98     -32.090  12.571   0.851  1.00 29.78      H    C
ATOM   6709  CD  ARG    98     -31.942  11.080   0.677  1.00 27.61      H    C
ATOM   6710  NE  ARG    98     -33.217  10.417   0.910  1.00 30.98      H    N
ATOM   6711  CZ  ARG    98     -33.400   9.099   0.859  1.00 31.84      H    C
ATOM   6712  NH1 ARG    98     -32.384   8.292   0.587  1.00 32.85      H    N
ATOM   6713  NH2 ARG    98     -34.605   8.593   1.064  1.00 29.18      H    N
ATOM   6714  C   ARG    98     -32.635  15.105  -0.344  1.00 28.25      H    C
ATOM   6715  O   ARG    98     -33.546  15.321   0.447  1.00 27.82      H    O
ATOM   6716  N   ASP    99     -32.803  15.096  -1.663  1.00 28.56      H    N
ATOM   6717  CA  ASP    99     -34.121  14.950  -2.262  1.00 27.74      H    C
ATOM   6718  CB  ASP    99     -34.795  16.321  -2.466  1.00 28.14      H    C
ATOM   6719  CG  ASP    99     -34.086  17.193  -3.497  1.00 34.77      H    C
ATOM   6720  OD1 ASP    99     -32.927  17.611  -3.252  1.00 34.38      H    O
ATOM   6721  OD2 ASP    99     -34.699  17.474  -4.556  1.00 36.63      H    O
ATOM   6722  C   ASP    99     -33.963  14.219  -3.586  1.00 27.51      H    C
ATOM   6723  O   ASP    99     -33.051  14.518  -4.364  1.00 27.91      H    O
ATOM   6724  N   TYR   100     -34.832  13.244  -3.838  1.00 25.59      H    N
ATOM   6725  CA  TYR   100     -34.683  12.437  -5.036  1.00 25.30      H    C
ATOM   6726  CB  TYR   100     -35.791  11.385  -5.123  1.00 26.94      H    C
ATOM   6727  CG  TYR   100     -35.608  10.449  -6.296  1.00 28.57      H    C
ATOM   6728  CD1 TYR   100     -34.740   9.365  -6.214  1.00 28.82      H    C
ATOM   6729  CE1 TYR   100     -34.511   8.547  -7.310  1.00 28.92      H    C
ATOM   6730  CD2 TYR   100     -36.252  10.686  -7.505  1.00 28.96      H    C
ATOM   6731  CE2 TYR   100     -36.032   9.873  -8.607  1.00 31.42      H    C
ATOM   6732  CZ  TYR   100     -35.157   8.804  -8.504  1.00 32.55      H    C
ATOM   6733  OH  TYR   100     -34.924   7.999  -9.604  1.00 33.58      H    O
ATOM   6734  C   TYR   100     -34.708  13.331  -6.274  1.00 25.64      H    C
ATOM   6735  O   TYR   100     -35.485  14.289  -6.355  1.00 23.00      H    O
ATOM   6736  N   ASP   101     -33.850  13.022  -7.239  1.00 26.57      H    N
ATOM   6737  CA  ASP   101     -33.715  13.877  -8.410  1.00 28.69      H    C
ATOM   6738  CB  ASP   101     -32.296  13.760  -8.978  1.00 28.46      H    C
ATOM   6739  CG  ASP   101     -32.013  14.776 -10.074  1.00 31.11      H    C
ATOM   6740  OD1 ASP   101     -32.949  15.493 -10.483  1.00 30.58      H    O
ATOM   6741  OD2 ASP   101     -30.846  14.860 -10.525  1.00 32.28      H    O
ATOM   6742  C   ASP   101     -34.751  13.461  -9.458  1.00 28.47      H    C
ATOM   6743  O   ASP   101     -34.602  12.433 -10.112  1.00 27.60      H    O
ATOM   6744  N   PHE   102     -35.808  14.258  -9.593  1.00 28.19      H    N
ATOM   6745  CA  PHE   102     -36.838  14.019 -10.604  1.00 28.93      H    C
ATOM   6746  CB  PHE   102     -38.225  14.347 -10.036  1.00 28.62      H    C
ATOM   6747  CG  PHE   102     -38.652  13.455  -8.889  1.00 28.28      H    C
ATOM   6748  CD1 PHE   102     -39.240  12.217  -9.134  1.00 27.61      H    C
ATOM   6749  CD2 PHE   102     -38.482  13.862  -7.573  1.00 28.22      H    C
ATOM   6750  CE1 PHE   102     -39.654  11.399  -8.086  1.00 29.20      H    C
ATOM   6751  CE2 PHE   102     -38.894  13.049  -6.508  1.00 29.50      H    C
ATOM   6752  CZ  PHE   102     -39.481  11.815  -6.767  1.00 28.70      H    C
ATOM   6753  C   PHE   102     -36.578  14.871 -11.859  1.00 28.33      H    C
ATOM   6754  O   PHE   102     -37.447  15.010 -12.725  1.00 29.51      H    O
ATOM   6755  N   TRP   103     -35.384  15.451 -11.939  1.00 25.45      H    N
ATOM   6756  CA  TRP   103     -34.952  16.192 -13.123  1.00 25.17      H    C
ATOM   6757  CB  TRP   103     -34.850  15.245 -14.323  1.00 25.43      H    C
ATOM   6758  CG  TRP   103     -33.937  14.070 -14.091  1.00 26.29      H    C
ATOM   6759  CD2 TRP   103     -32.526  14.010 -14.361  1.00 27.47      H    C
ATOM   6760  CE2 TRP   103     -32.080  12.723 -13.970  1.00 27.34      H    C
ATOM   6761  CE3 TRP   103     -31.598  14.914 -14.898  1.00 26.51      H    C
ATOM   6762  CD1 TRP   103     -34.276  12.853 -13.565  1.00 26.52      H    C
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ATOM	6763	NE1	TRP	103	-33.167	12.040	-13.489	1.00	26.07	H	N
ATOM	6764	CZ2	TRP	103	-30.744	12.321	-14.093	1.00	25.38	H	C
ATOM	6765	CZ3	TRP	103	-30.272	14.514	-15.023	1.00	26.49	H	C
ATOM	6766	CH2	TRP	103	-29.858	13.225	-14.621	1.00	27.17	H	C
ATOM	6767	C	TRP	103	-35.875	17.373	-13.448	1.00	26.58	H	C
ATOM	6768	O	TRP	103	-36.174	17.644	-14.617	1.00	25.55	H	O
ATOM	6769	N	SER	104	-36.310	18.063	-12.393	1.00	26.77	H	N
ATOM	6770	CA	SER	104	-37.202	19.226	-12.456	1.00	27.21	H	C
ATOM	6771	CB	SER	104	-36.686	20.280	-13.463	1.00	28.29	H	C
ATOM	6772	OG	SER	104	-37.065	20.001	-14.806	1.00	28.50	H	O
ATOM	6773	C	SER	104	-38.655	18.875	-12.760	1.00	27.34	H	C
ATOM	6774	O	SER	104	-39.497	19.758	-12.843	1.00	29.98	H	O
ATOM	6775	N	ALA	105	-38.961	17.590	-12.911	1.00	27.05	H	N
ATOM	6776	CA	ALA	105	-40.349	17.178	-13.141	1.00	27.35	H	C
ATOM	6777	CB	ALA	105	-40.406	15.700	-13.521	1.00	25.10	H	C
ATOM	6778	C	ALA	105	-41.196	17.422	-11.893	1.00	27.00	H	C
ATOM	6779	O	ALA	105	-42.404	17.675	-11.979	1.00	26.55	H	O
ATOM	6780	N	TYR	106	-40.554	17.331	-10.733	1.00	26.09	H	N
ATOM	6781	CA	TYR	106	-41.233	17.502	-9.456	1.00	25.70	H	C
ATOM	6782	CB	TYR	106	-41.923	16.190	-9.043	1.00	27.12	H	C
ATOM	6783	CG	TYR	106	-42.597	16.242	-7.682	1.00	27.55	H	C
ATOM	6784	CD1	TYR	106	-42.076	15.545	-6.598	1.00	25.52	H	C
ATOM	6785	CE1	TYR	106	-42.678	15.609	-5.347	1.00	26.89	H	C
ATOM	6786	CD2	TYR	106	-43.745	17.009	-7.482	1.00	26.08	H	C
ATOM	6787	CE2	TYR	106	-44.349	17.085	-6.242	1.00	27.05	H	C
ATOM	6788	CZ	TYR	106	-43.810	16.383	-5.179	1.00	28.47	H	C
ATOM	6789	OH	TYR	106	-44.405	16.471	-3.945	1.00	29.33	H	O
ATOM	6790	C	TYR	106	-40.189	17.884	-8.416	1.00	27.08	H	C
ATOM	6791	O	TYR	106	-39.053	17.414	-8.468	1.00	29.19	H	O
ATOM	6792	N	TYR	107	-40.562	18.734	-7.469	1.00	27.35	H	N
ATOM	6793	CA	TYR	107	-39.632	19.091	-6.403	1.00	29.15	H	C
ATOM	6794	CB	TYR	107	-39.511	20.614	-6.307	1.00	29.00	H	C
ATOM	6795	CG	TYR	107	-38.874	21.203	-7.548	1.00	31.69	H	C
ATOM	6796	CD1	TYR	107	-39.638	21.854	-8.511	1.00	30.78	H	C
ATOM	6797	CE1	TYR	107	-39.059	22.362	-9.664	1.00	31.09	H	C
ATOM	6798	CD2	TYR	107	-37.510	21.075	-7.774	1.00	32.92	H	C
ATOM	6799	CE2	TYR	107	-36.921	21.579	-8.928	1.00	33.30	H	C
ATOM	6800	CZ	TYR	107	-37.702	22.221	-9.868	1.00	31.33	H	C
ATOM	6801	OH	TYR	107	-37.115	22.718	-11.012	1.00	31.72	H	O
ATOM	6802	C	TYR	107	-40.069	18.483	-5.078	1.00	27.80	H	C
ATOM	6803	O	TYR	107	-41.029	18.933	-4.451	1.00	27.16	H	O
ATOM	6804	N	ASP	108	-39.356	17.439	-4.673	1.00	26.30	H	N
ATOM	6805	CA	ASP	108	-39.769	16.524	-3.547	1.00	27.34	H	C
ATOM	6806	CB	ASP	108	-39.196	15.214	-3.692	1.00	28.85	H	C
ATOM	6807	CG	ASP	108	-39.932	14.183	-2.844	1.00	32.64	H	C
ATOM	6808	OD1	ASP	108	-40.918	14.543	-2.149	1.00	28.23	H	O
ATOM	6809	OD2	ASP	108	-39.517	13.002	-2.879	1.00	31.34	H	O
ATOM	6810	C	ASP	108	-39.250	17.262	-2.269	1.00	27.61	H	C
ATOM	6811	O	ASP	108	-38.456	18.203	-2.319	1.00	27.35	H	O
ATOM	6812	N	ALA	109	-39.704	16.742	-1.135	1.00	26.89	H	N
ATOM	6813	CA	ALA	109	-39.248	17.186	0.174	1.00	28.36	H	C
ATOM	6814	CB	ALA	109	-40.160	16.614	1.264	1.00	28.17	H	C
ATOM	6815	C	ALA	109	-37.814	16.734	0.409	1.00	30.74	H	C
ATOM	6816	O	ALA	109	-37.346	15.767	-0.209	1.00	30.34	H	O
ATOM	6817	N	PHE	110	-37.118	17.444	1.299	1.00	30.36	H	N
ATOM	6818	CA	PHE	110	-35.829	16.997	1.815	1.00	29.18	H	C
ATOM	6819	CB	PHE	110	-35.087	18.166	2.481	1.00	27.43	H	C
ATOM	6820	CG	PHE	110	-34.728	19.286	1.537	1.00	26.74	H	C
ATOM	6821	CD1	PHE	110	-34.084	20.420	2.006	1.00	25.08	H	C
ATOM	6822	CD2	PHE	110	-35.044	19.210	0.187	1.00	27.64	H	C
ATOM	6823	CE1	PHE	110	-33.760	21.470	1.146	1.00	29.39	H	C
ATOM	6824	CE2	PHE	110	-34.725	20.253	-0.683	1.00	31.03	H	C
ATOM	6825	CZ	PHE	110	-34.081	21.387	-0.201	1.00	28.21	H	C
ATOM	6826	C	PHE	110	-36.142	15.930	2.856	1.00	30.56	H	C
ATOM	6827	O	PHE	110	-36.469	16.256	3.999	1.00	32.31	H	O
ATOM	6828	N	ASP	111	-36.065	14.657	2.479	1.00	31.23	H	N
ATOM	6829	CA	ASP	111	-36.660	13.634	3.336	1.00	33.06	H	C
ATOM	6830	CB	ASP	111	-37.352	12.546	2.491	1.00	33.67	H	C
ATOM	6831	CG	ASP	111	-36.385	11.710	1.664	1.00	37.65	H	C
ATOM	6832	OD1	ASP	111	-35.276	12.183	1.326	1.00	37.86	H	O
ATOM	6833	OD2	ASP	111	-36.756	10.560	1.344	1.00	38.69	H	O
ATOM	6834	C	ASP	111	-35.717	13.009	4.355	1.00	32.70	H	C
ATOM	6835	O	ASP	111	-36.162	12.365	5.305	1.00	33.78	H	O
ATOM	6836	N	VAL	112	-34.418	13.213	4.167	1.00	32.63	H	N
ATOM	6837	CA	VAL	112	-33.425	12.801	5.151	1.00	32.15	H	C
ATOM	6838	CB	VAL	112	-32.689	11.505	4.708	1.00	33.32	H	C

ATOM	6839	CG1	VAL	112	-31.585	11.153	5.718	1.00	30.53	H	C
ATOM	6840	CG2	VAL	112	-33.688	10.356	4.599	1.00	31.40	H	C
ATOM	6841	C	VAL	112	-32.400	13.914	5.343	1.00	32.58	H	C
ATOM	6842	O	VAL	112	-31.889	14.476	4.370	1.00	33.04	H	O
ATOM	6843	N	TRP	113	-32.112	14.228	6.602	1.00	32.15	H	N
ATOM	6844	CA	TRP	113	-31.161	15.277	6.953	1.00	31.65	H	C
ATOM	6845	CB	TRP	113	-31.843	16.338	7.816	1.00	29.50	H	C
ATOM	6846	CG	TRP	113	-32.931	17.105	7.128	1.00	28.47	H	C
ATOM	6847	CD2	TRP	113	-33.006	18.527	6.977	1.00	25.65	H	C
ATOM	6848	CE2	TRP	113	-34.217	18.813	6.315	1.00	26.45	H	C
ATOM	6849	CE3	TRP	113	-32.167	19.586	7.341	1.00	24.52	H	C
ATOM	6850	CD1	TRP	113	-34.069	16.597	6.562	1.00	26.10	H	C
ATOM	6851	NE1	TRP	113	-34.847	17.620	6.074	1.00	28.26	H	N
ATOM	6852	CZ2	TRP	113	-34.611	20.118	6.008	1.00	26.31	H	C
ATOM	6853	CZ3	TRP	113	-32.558	20.886	7.037	1.00	24.40	H	C
ATOM	6854	CH2	TRP	113	-33.771	21.139	6.377	1.00	25.50	H	C
ATOM	6855	C	TRP	113	-29.996	14.687	7.746	1.00	33.12	H	C
ATOM	6856	O	TRP	113	-30.136	13.634	8.371	1.00	31.88	H	O
ATOM	6857	N	GLY	114	-28.855	15.373	7.723	1.00	32.68	H	N
ATOM	6858	CA	GLY	114	-27.776	15.044	8.639	1.00	35.69	H	C
ATOM	6859	C	GLY	114	-27.980	15.743	9.974	1.00	37.38	H	C
ATOM	6860	O	GLY	114	-28.984	16.429	10.170	1.00	38.62	H	O
ATOM	6861	N	GLN	115	-27.036	15.587	10.894	1.00	38.12	H	N
ATOM	6862	CA	GLN	115	-27.189	16.182	12.218	1.00	40.72	H	C
ATOM	6863	CB	GLN	115	-26.590	15.262	13.285	1.00	43.53	H	C
ATOM	6864	CG	GLN	115	-27.488	14.063	13.619	1.00	51.75	H	C
ATOM	6865	CD	GLN	115	-28.955	14.461	13.857	1.00	56.93	H	C
ATOM	6866	OE1	GLN	115	-29.269	15.223	14.778	1.00	58.91	H	O
ATOM	6867	NE2	GLN	115	-29.853	13.944	13.018	1.00	58.11	H	N
ATOM	6868	C	GLN	115	-26.573	17.573	12.303	1.00	39.58	H	C
ATOM	6869	O	GLN	115	-26.820	18.321	13.254	1.00	38.68	H	O
ATOM	6870	N	GLY	116	-25.782	17.928	11.298	1.00	37.78	H	N
ATOM	6871	CA	GLY	116	-25.283	19.285	11.220	1.00	38.42	H	C
ATOM	6872	C	GLY	116	-23.874	19.439	11.752	1.00	39.35	H	C
ATOM	6873	O	GLY	116	-23.407	18.644	12.568	1.00	39.93	H	O
ATOM	6874	N	THR	117	-23.188	20.466	11.267	1.00	39.58	H	N
ATOM	6875	CA	THR	117	-21.862	20.803	11.750	1.00	39.22	H	C
ATOM	6876	CB	THR	117	-20.776	20.415	10.720	1.00	39.77	H	C
ATOM	6877	OG1	THR	117	-19.482	20.645	11.284	1.00	41.61	H	O
ATOM	6878	CG2	THR	117	-20.912	21.241	9.445	1.00	38.56	H	C
ATOM	6879	C	THR	117	-21.849	22.306	11.985	1.00	38.56	H	C
ATOM	6880	O	THR	117	-22.412	23.063	11.194	1.00	39.43	H	O
ATOM	6881	N	MET	118	-21.227	22.733	13.079	1.00	38.05	H	N
ATOM	6882	CA	MET	118	-21.275	24.134	13.498	1.00	36.74	H	C
ATOM	6883	CB	MET	118	-21.189	24.230	15.024	1.00	38.21	H	C
ATOM	6884	CG	MET	118	-21.254	25.649	15.561	1.00	41.58	H	C
ATOM	6885	SD	MET	118	-22.816	26.465	15.157	1.00	50.65	H	S
ATOM	6886	CE	MET	118	-22.441	28.169	15.524	1.00	47.35	H	C
ATOM	6887	C	MET	118	-20.147	24.946	12.871	1.00	34.48	H	C
ATOM	6888	O	MET	118	-19.013	24.487	12.768	1.00	34.26	H	O
ATOM	6889	N	VAL	119	-20.474	26.159	12.449	1.00	34.68	H	N
ATOM	6890	CA	VAL	119	-19.506	27.054	11.833	1.00	33.91	H	C
ATOM	6891	CB	VAL	119	-19.783	27.251	10.311	1.00	33.34	H	C
ATOM	6892	CG1	VAL	119	-19.007	28.445	9.793	1.00	33.26	H	C
ATOM	6893	CG2	VAL	119	-19.376	26.001	9.531	1.00	31.55	H	C
ATOM	6894	C	VAL	119	-19.643	28.389	12.520	1.00	34.59	H	C
ATOM	6895	O	VAL	119	-20.724	28.981	12.517	1.00	37.76	H	O
ATOM	6896	N	THR	120	-18.553	28.863	13.117	1.00	34.91	H	N
ATOM	6897	CA	THR	120	-18.554	30.171	13.762	1.00	35.35	H	C
ATOM	6898	CB	THR	120	-18.134	30.073	15.256	1.00	35.88	H	C
ATOM	6899	OG1	THR	120	-19.037	29.212	15.960	1.00	37.94	H	O
ATOM	6900	CG2	THR	120	-18.154	31.444	15.902	1.00	34.25	H	C
ATOM	6901	C	THR	120	-17.578	31.085	13.038	1.00	35.16	H	C
ATOM	6902	O	THR	120	-16.442	30.705	12.768	1.00	35.56	H	O
ATOM	6903	N	VAL	121	-18.028	32.291	12.721	1.00	36.37	H	N
ATOM	6904	CA	VAL	121	-17.166	33.271	12.088	1.00	37.55	H	C
ATOM	6905	CB	VAL	121	-17.718	33.687	10.703	1.00	38.08	H	C
ATOM	6906	CG1	VAL	121	-16.839	34.765	10.089	1.00	36.30	H	C
ATOM	6907	CG2	VAL	121	-17.774	32.470	9.784	1.00	36.74	H	C
ATOM	6908	C	VAL	121	-17.063	34.491	12.985	1.00	38.28	H	C
ATOM	6909	O	VAL	121	-18.029	35.233	13.156	1.00	39.94	H	O
ATOM	6910	N	SER	122	-15.884	34.694	13.563	1.00	38.45	H	N
ATOM	6911	CA	SER	122	-15.694	35.755	14.550	1.00	39.93	H	C
ATOM	6912	CB	SER	122	-15.947	35.209	15.962	1.00	40.21	H	C
ATOM	6913	OG	SER	122	-15.882	36.242	16.929	1.00	42.15	H	O
ATOM	6914	C	SER	122	-14.284	36.331	14.473	1.00	39.50	H	C

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ATOM   6915  O   SER   122     -13.320  35.615  14.199  1.00 37.65      H    O
ATOM   6916  N   SER   123     -14.165  37.627  14.726  1.00 41.58      H    N
ATOM   6917  CA  SER   123     -12.849  38.253  14.807  1.00 44.16      H    C
ATOM   6918  CB  SER   123     -12.993  39.776  14.777  1.00 44.85      H    C
ATOM   6919  OG  SER   123     -13.875  40.223  15.794  1.00 49.05      H    O
ATOM   6920  C   SER   123     -12.105  37.819  16.079  1.00 44.60      H    C
ATOM   6921  O   SER   123     -10.893  37.995  16.184  1.00 45.68      H    O
ATOM   6922  N   ALA   124     -12.837  37.240  17.031  1.00 43.93      H    N
ATOM   6923  CA  ALA   124     -12.262  36.801  18.301  1.00 42.24      H    C
ATOM   6924  CB  ALA   124     -13.326  36.846  19.390  1.00 40.10      H    C
ATOM   6925  C   ALA   124     -11.675  35.394  18.214  1.00 42.14      H    C
ATOM   6926  O   ALA   124     -12.274  34.500  17.620  1.00 43.33      H    O
ATOM   6927  N   SER   125     -10.503  35.201  18.820  1.00 42.38      H    N
ATOM   6928  CA  SER   125      -9.862  33.887  18.876  1.00 40.59      H    C
ATOM   6929  CB  SER   125      -8.355  34.041  19.109  1.00 42.02      H    C
ATOM   6930  OG  SER   125      -7.752  34.799  18.072  1.00 45.10      H    O
ATOM   6931  C   SER   125     -10.466  33.067  20.010  1.00 38.95      H    C
ATOM   6932  O   SER   125     -11.093  33.619  20.907  1.00 37.48      H    O
ATOM   6933  N   THR   126     -10.271  31.753  19.977  1.00 37.66      H    N
ATOM   6934  CA  THR   126     -10.785  30.905  21.042  1.00 39.23      H    C
ATOM   6935  CB  THR   126     -10.669  29.419  20.674  1.00 38.28      H    C
ATOM   6936  OG1 THR   126     -11.284  28.629  21.697  1.00 41.97      H    O
ATOM   6937  CG2 THR   126      -9.217  29.008  20.533  1.00 37.08      H    C
ATOM   6938  C   THR   126     -10.049  31.147  22.370  1.00 41.01      H    C
ATOM   6939  O   THR   126      -8.841  31.397  22.392  1.00 41.24      H    O
ATOM   6940  N   LYS   127     -10.794  31.074  23.471  1.00 40.95      H    N
ATOM   6941  CA  LYS   127     -10.262  31.339  24.806  1.00 38.98      H    C
ATOM   6942  CB  LYS   127     -10.599  32.771  25.231  1.00 38.46      H    C
ATOM   6943  CG  LYS   127     -10.108  33.139  26.625  1.00 39.85      H    C
ATOM   6944  CD  LYS   127     -10.581  34.529  27.043  1.00 39.69      H    C
ATOM   6945  CE  LYS   127     -10.208  34.833  28.494  1.00 40.59      H    C
ATOM   6946  NZ  LYS   127     -10.600  33.727  29.429  1.00 41.49      H    N
ATOM   6947  C   LYS   127     -10.862  30.359  25.809  1.00 37.64      H    C
ATOM   6948  O   LYS   127     -12.080  30.248  25.916  1.00 37.85      H    O
ATOM   6949  N   GLY   128     -10.005  29.648  26.536  1.00 36.50      H    N
ATOM   6950  CA  GLY   128     -10.476  28.740  27.566  1.00 33.05      H    C
ATOM   6951  C   GLY   128     -11.064  29.484  28.750  1.00 32.53      H    C
ATOM   6952  O   GLY   128     -10.832  30.681  28.923  1.00 32.58      H    O
ATOM   6953  N   PRO   129     -11.856  28.798  29.578  1.00 32.14      H    N
ATOM   6954  CD  PRO   129     -12.255  27.396  29.361  1.00 32.62      H    C
ATOM   6955  CA  PRO   129     -12.553  29.387  30.728  1.00 32.53      H    C
ATOM   6956  CB  PRO   129     -13.695  28.411  30.978  1.00 31.17      H    C
ATOM   6957  CG  PRO   129     -13.119  27.096  30.561  1.00 30.16      H    C
ATOM   6958  C   PRO   129     -11.663  29.522  31.967  1.00 33.22      H    C
ATOM   6959  O   PRO   129     -10.742  28.732  32.161  1.00 31.90      H    O
ATOM   6960  N   SER   130     -11.954  30.513  32.804  1.00 32.62      H    N
ATOM   6961  CA  SER   130     -11.478  30.505  34.185  1.00 35.02      H    C
ATOM   6962  CB  SER   130     -11.275  31.932  34.687  1.00 34.24      H    C
ATOM   6963  OG  SER   130     -10.239  32.566  33.964  1.00 41.41      H    O
ATOM   6964  C   SER   130     -12.522  29.807  35.050  1.00 35.50      H    C
ATOM   6965  O   SER   130     -13.708  30.118  34.958  1.00 37.01      H    O
ATOM   6966  N   VAL   131     -12.091  28.868  35.884  1.00 34.84      H    N
ATOM   6967  CA  VAL   131     -13.026  28.121  36.716  1.00 34.36      H    C
ATOM   6968  CB  VAL   131     -12.840  26.605  36.526  1.00 34.23      H    C
ATOM   6969  CG1 VAL   131     -13.882  25.848  37.333  1.00 33.85      H    C
ATOM   6970  CG2 VAL   131     -12.938  26.251  35.046  1.00 34.73      H    C
ATOM   6971  C   VAL   131     -12.860  28.460  38.198  1.00 35.71      H    C
ATOM   6972  O   VAL   131     -11.760  28.349  38.750  1.00 35.77      H    O
ATOM   6973  N   PHE   132     -13.955  28.878  38.834  1.00 34.57      H    N
ATOM   6974  CA  PHE   132     -13.953  29.184  40.261  1.00 33.58      H    C
ATOM   6975  CB  PHE   132     -14.266  30.664  40.489  1.00 32.57      H    C
ATOM   6976  CG  PHE   132     -13.327  31.593  39.782  1.00 34.21      H    C
ATOM   6977  CD1 PHE   132     -12.103  31.922  40.344  1.00 35.59      H    C
ATOM   6978  CD2 PHE   132     -13.651  32.114  38.538  1.00 33.19      H    C
ATOM   6979  CE1 PHE   132     -11.214  32.750  39.674  1.00 37.01      H    C
ATOM   6980  CE2 PHE   132     -12.769  32.943  37.862  1.00 35.41      H    C
ATOM   6981  CZ  PHE   132     -11.548  33.262  38.430  1.00 36.28      H    C
ATOM   6982  C   PHE   132     -14.963  28.325  41.022  1.00 34.57      H    C
ATOM   6983  O   PHE   132     -16.041  28.012  40.512  1.00 32.06      H    O
ATOM   6984  N   PRO   133     -14.617  27.927  42.258  1.00 34.29      H    N
ATOM   6985  CD  PRO   133     -13.324  28.197  42.918  1.00 34.35      H    C
ATOM   6986  CA  PRO   133     -15.511  27.132  43.108  1.00 35.13      H    C
ATOM   6987  CB  PRO   133     -14.593  26.624  44.220  1.00 33.32      H    C
ATOM   6988  CG  PRO   133     -13.541  27.689  44.336  1.00 34.24      H    C
ATOM   6989  C   PRO   133     -16.645  27.983  43.662  1.00 35.55      H    C
ATOM   6990  O   PRO   133     -16.426  29.128  44.068  1.00 36.64      H    O
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ATOM	6991	N	LEU	134	-17.852	27.424	43.674	1.00	35.32	H	N
ATOM	6992	CA	LEU	134	-18.977	28.048	44.366	1.00	36.08	H	C
ATOM	6993	CB	LEU	134	-20.209	28.094	43.450	1.00	34.28	H	C
ATOM	6994	CG	LEU	134	-19.960	28.860	42.142	1.00	34.73	H	C
ATOM	6995	CD1	LEU	134	-21.140	28.718	41.185	1.00	31.18	H	C
ATOM	6996	CD2	LEU	134	-19.697	30.321	42.475	1.00	33.05	H	C
ATOM	6997	C	LEU	134	-19.259	27.224	45.618	1.00	36.52	H	C
ATOM	6998	O	LEU	134	-19.999	26.234	45.585	1.00	34.85	H	O
ATOM	6999	N	ALA	135	-18.645	27.635	46.720	1.00	37.22	H	N
ATOM	7000	CA	ALA	135	-18.596	26.810	47.917	1.00	40.97	H	C
ATOM	7001	CB	ALA	135	-17.561	27.367	48.891	1.00	38.81	H	C
ATOM	7002	C	ALA	135	-19.957	26.725	48.594	1.00	44.00	H	C
ATOM	7003	O	ALA	135	-20.712	27.697	48.626	1.00	40.78	H	O
ATOM	7004	N	PRO	136	-20.283	25.550	49.148	1.00	48.76	H	N
ATOM	7005	CD	PRO	136	-19.446	24.338	49.152	1.00	49.46	H	C
ATOM	7006	CA	PRO	136	-21.531	25.360	49.890	1.00	54.13	H	C
ATOM	7007	CB	PRO	136	-21.528	23.874	50.232	1.00	52.72	H	C
ATOM	7008	CG	PRO	136	-20.085	23.486	50.205	1.00	51.83	H	C
ATOM	7009	C	PRO	136	-21.586	26.238	51.132	1.00	59.78	H	C
ATOM	7010	O	PRO	136	-20.570	26.460	51.794	1.00	59.54	H	O
ATOM	7011	N	SER	137	-22.782	26.737	51.432	1.00	67.10	H	N
ATOM	7012	CA	SER	137	-22.993	27.638	52.560	1.00	74.14	H	C
ATOM	7013	CB	SER	137	-24.411	28.215	52.507	1.00	75.09	H	C
ATOM	7014	OG	SER	137	-24.664	29.061	53.617	1.00	78.01	H	O
ATOM	7015	C	SER	137	-22.774	26.926	53.892	1.00	77.67	H	C
ATOM	7016	O	SER	137	-22.866	25.699	53.977	1.00	78.44	H	O
ATOM	7017	N	SER	138	-22.492	27.707	54.931	1.00	82.40	H	N
ATOM	7018	CA	SER	138	-22.171	27.163	56.248	1.00	86.84	H	C
ATOM	7019	CB	SER	138	-21.280	28.151	57.011	1.00	87.07	H	C
ATOM	7020	OG	SER	138	-20.112	28.463	56.271	1.00	87.61	H	O
ATOM	7021	C	SER	138	-23.414	26.841	57.087	1.00	89.30	H	C
ATOM	7022	O	SER	138	-23.688	27.512	58.085	1.00	89.91	H	O
ATOM	7023	N	LYS	139	-24.161	25.816	56.678	1.00	91.76	H	N
ATOM	7024	CA	LYS	139	-25.298	25.325	57.457	1.00	93.44	H	C
ATOM	7025	CB	LYS	139	-26.616	25.879	56.899	1.00	93.81	H	C
ATOM	7026	CG	LYS	139	-26.910	27.321	57.291	1.00	94.61	H	C
ATOM	7027	CD	LYS	139	-26.118	28.303	56.442	1.00	95.03	H	C
ATOM	7028	CE	LYS	139	-26.243	29.721	56.976	1.00	95.42	H	C
ATOM	7029	NZ	LYS	139	-25.510	30.700	56.125	1.00	95.24	H	N
ATOM	7030	C	LYS	139	-25.349	23.798	57.464	1.00	94.08	H	C
ATOM	7031	O	LYS	139	-26.414	23.202	57.638	1.00	95.08	H	O
ATOM	7032	N	GLY	143	-30.107	19.527	59.281	1.00	59.11	H	N
ATOM	7033	CA	GLY	143	-29.319	19.470	58.062	1.00	61.11	H	C
ATOM	7034	C	GLY	143	-29.957	18.625	56.968	1.00	60.28	H	C
ATOM	7035	O	GLY	143	-30.100	17.406	57.106	1.00	60.72	H	O
ATOM	7036	N	GLY	144	-30.342	19.280	55.877	1.00	58.29	H	N
ATOM	7037	CA	GLY	144	-30.925	18.578	54.746	1.00	54.60	H	C
ATOM	7038	C	GLY	144	-29.972	18.554	53.562	1.00	52.81	H	C
ATOM	7039	O	GLY	144	-29.023	17.772	53.550	1.00	53.00	H	O
ATOM	7040	N	THR	145	-30.216	19.401	52.565	1.00	49.01	H	N
ATOM	7041	CA	THR	145	-29.334	19.461	51.403	1.00	46.11	H	C
ATOM	7042	CB	THR	145	-30.109	19.311	50.080	1.00	46.35	H	C
ATOM	7043	OG1	THR	145	-31.019	20.410	49.929	1.00	48.50	H	O
ATOM	7044	CG2	THR	145	-30.877	18.001	50.056	1.00	44.98	H	C
ATOM	7045	C	THR	145	-28.562	20.775	51.343	1.00	44.27	H	C
ATOM	7046	O	THR	145	-28.979	21.788	51.912	1.00	42.47	H	O
ATOM	7047	N	ALA	146	-27.428	20.751	50.651	1.00	41.79	H	N
ATOM	7048	CA	ALA	146	-26.691	21.974	50.374	1.00	40.00	H	C
ATOM	7049	CB	ALA	146	-25.407	22.009	51.187	1.00	40.43	H	C
ATOM	7050	C	ALA	146	-26.374	22.057	48.888	1.00	39.46	H	C
ATOM	7051	O	ALA	146	-26.227	21.037	48.212	1.00	38.93	H	O
ATOM	7052	N	ALA	147	-26.278	23.279	48.380	1.00	37.34	H	N
ATOM	7053	CA	ALA	147	-25.943	23.491	46.986	1.00	35.51	H	C
ATOM	7054	CB	ALA	147	-26.894	24.512	46.374	1.00	34.09	H	C
ATOM	7055	C	ALA	147	-24.500	23.977	46.874	1.00	35.59	H	C
ATOM	7056	O	ALA	147	-24.055	24.852	47.620	1.00	35.64	H	O
ATOM	7057	N	LEU	148	-23.765	23.396	45.942	1.00	34.79	H	N
ATOM	7058	CA	LEU	148	-22.421	23.857	45.658	1.00	34.42	H	C
ATOM	7059	CB	LEU	148	-21.402	23.008	46.425	1.00	35.26	H	C
ATOM	7060	CG	LEU	148	-21.395	21.515	46.088	1.00	38.02	H	C
ATOM	7061	CD1	LEU	148	-20.408	21.250	44.961	1.00	38.18	H	C
ATOM	7062	CD2	LEU	148	-21.011	20.713	47.323	1.00	39.69	H	C
ATOM	7063	C	LEU	148	-22.221	23.721	44.159	1.00	33.02	H	C
ATOM	7064	O	LEU	148	-22.934	22.968	43.498	1.00	32.43	H	O
ATOM	7065	N	GLY	149	-21.267	24.457	43.612	1.00	32.54	H	N
ATOM	7066	CA	GLY	149	-21.040	24.349	42.187	1.00	35.20	H	C

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ATOM   7067  C   GLY   149     -19.729  24.945  41.742  1.00 35.35      H    C
ATOM   7068  O   GLY   149     -18.812  25.171  42.537  1.00 34.93      H    O
ATOM   7069  N   CYS   150     -19.626  25.201  40.451  1.00 36.17      H    N
ATOM   7070  CA  CYS   150     -18.464  25.898  39.974  1.00 38.63      H    C
ATOM   7071  C   CYS   150     -18.825  26.843  38.841  1.00 36.98      H    C
ATOM   7072  O   CYS   150     -19.729  26.576  38.048  1.00 37.67      H    O
ATOM   7073  CB  CYS   150     -17.360  24.886  39.601  1.00 42.69      H    C
ATOM   7074  SG  CYS   150     -17.554  23.900  38.089  1.00 55.77      H    S
ATOM   7075  N   LEU   151     -18.141  27.981  38.818  1.00 35.05      H    N
ATOM   7076  CA  LEU   151     -18.431  29.068  37.898  1.00 33.32      H    C
ATOM   7077  CB  LEU   151     -18.334  30.397  38.640  1.00 31.86      H    C
ATOM   7078  CG  LEU   151     -18.258  31.673  37.803  1.00 34.10      H    C
ATOM   7079  CD1 LEU   151     -19.607  31.930  37.141  1.00 31.52      H    C
ATOM   7080  CD2 LEU   151     -17.863  32.841  38.701  1.00 32.35      H    C
ATOM   7081  C   LEU   151     -17.415  29.035  36.764  1.00 35.15      H    C
ATOM   7082  O   LEU   151     -16.204  29.012  37.002  1.00 34.70      H    O
ATOM   7083  N   VAL   152     -17.909  29.034  35.531  1.00 35.15      H    N
ATOM   7084  CA  VAL   152     -17.053  28.860  34.365  1.00 33.97      H    C
ATOM   7085  CB  VAL   152     -17.542  27.662  33.519  1.00 34.25      H    C
ATOM   7086  CG1 VAL   152     -16.662  27.474  32.298  1.00 33.15      H    C
ATOM   7087  CG2 VAL   152     -17.540  26.398  34.373  1.00 33.07      H    C
ATOM   7088  C   VAL   152     -17.090  30.136  33.536  1.00 35.52      H    C
ATOM   7089  O   VAL   152     -18.059  30.388  32.816  1.00 35.70      H    O
ATOM   7090  N   LYS   153     -16.025  30.931  33.644  1.00 35.56      H    N
ATOM   7091  CA  LYS   153     -16.020  32.325  33.194  1.00 36.98      H    C
ATOM   7092  CB  LYS   153     -15.435  33.227  34.288  1.00 37.29      H    C
ATOM   7093  CG  LYS   153     -16.456  33.857  35.205  1.00 44.06      H    C
ATOM   7094  CD  LYS   153     -15.778  34.702  36.286  1.00 45.54      H    C
ATOM   7095  CE  LYS   153     -15.056  35.903  35.687  1.00 48.07      H    C
ATOM   7096  NZ  LYS   153     -14.280  36.672  36.708  1.00 48.70      H    N
ATOM   7097  C   LYS   153     -15.250  32.589  31.896  1.00 35.32      H    C
ATOM   7098  O   LYS   153     -14.178  32.029  31.674  1.00 33.48      H    O
ATOM   7099  N   ASP   154     -15.803  33.466  31.061  1.00 35.67      H    N
ATOM   7100  CA  ASP   154     -15.069  34.081  29.951  1.00 38.13      H    C
ATOM   7101  CB  ASP   154     -13.929  34.957  30.488  1.00 38.72      H    C
ATOM   7102  CG  ASP   154     -14.421  36.122  31.326  1.00 42.66      H    C
ATOM   7103  OD1 ASP   154     -15.540  36.626  31.078  1.00 42.06      H    O
ATOM   7104  OD2 ASP   154     -13.675  36.538  32.241  1.00 47.57      H    O
ATOM   7105  C   ASP   154     -14.473  33.096  28.945  1.00 37.85      H    C
ATOM   7106  O   ASP   154     -13.271  33.134  28.676  1.00 39.38      H    O
ATOM   7107  N   TYR   155     -15.286  32.217  28.377  1.00 36.72      H    N
ATOM   7108  CA  TYR   155     -14.769  31.349  27.326  1.00 35.87      H    C
ATOM   7109  CB  TYR   155     -15.004  29.883  27.690  1.00 35.27      H    C
ATOM   7110  CG  TYR   155     -16.462  29.508  27.807  1.00 37.64      H    C
ATOM   7111  CD1 TYR   155     -17.179  29.084  26.693  1.00 36.63      H    C
ATOM   7112  CE1 TYR   155     -18.510  28.735  26.793  1.00 38.10      H    C
ATOM   7113  CD2 TYR   155     -17.123  29.569  29.032  1.00 36.52      H    C
ATOM   7114  CE2 TYR   155     -18.457  29.217  29.140  1.00 36.48      H    C
ATOM   7115  CZ  TYR   155     -19.144  28.802  28.017  1.00 37.43      H    C
ATOM   7116  OH  TYR   155     -20.475  28.465  28.105  1.00 38.45      H    O
ATOM   7117  C   TYR   155     -15.398  31.670  25.966  1.00 36.12      H    C
ATOM   7118  O   TYR   155     -16.382  32.414  25.881  1.00 35.25      H    O
ATOM   7119  N   PHE   156     -14.814  31.116  24.908  1.00 36.85      H    N
ATOM   7120  CA  PHE   156     -15.321  31.310  23.554  1.00 38.37      H    C
ATOM   7121  CB  PHE   156     -15.072  32.745  23.088  1.00 38.82      H    C
ATOM   7122  CG  PHE   156     -15.588  33.032  21.705  1.00 42.82      H    C
ATOM   7123  CD1 PHE   156     -14.798  32.792  20.588  1.00 43.82      H    C
ATOM   7124  CD2 PHE   156     -16.870  33.526  21.516  1.00 43.40      H    C
ATOM   7125  CE1 PHE   156     -15.280  33.039  19.308  1.00 44.30      H    C
ATOM   7126  CE2 PHE   156     -17.358  33.774  20.235  1.00 43.81      H    C
ATOM   7127  CZ  PHE   156     -16.561  33.530  19.134  1.00 41.90      H    C
ATOM   7128  C   PHE   156     -14.646  30.342  22.593  1.00 39.91      H    C
ATOM   7129  O   PHE   156     -13.452  30.067  22.710  1.00 40.67      H    O
ATOM   7130  N   PRO   157     -15.410  29.793  21.635  1.00 40.59      H    N
ATOM   7131  CD  PRO   157     -14.843  29.059  20.491  1.00 41.29      H    C
ATOM   7132  CA  PRO   157     -16.873  29.867  21.549  1.00 40.24      H    C
ATOM   7133  CB  PRO   157     -17.141  29.658  20.065  1.00 41.39      H    C
ATOM   7134  CG  PRO   157     -16.070  28.698  19.661  1.00 42.40      H    C
ATOM   7135  C   PRO   157     -17.533  28.777  22.402  1.00 38.74      H    C
ATOM   7136  O   PRO   157     -16.863  28.082  23.163  1.00 36.64      H    O
ATOM   7137  N   GLU   158     -18.845  28.619  22.258  1.00 39.77      H    N
ATOM   7138  CA  GLU   158     -19.543  27.476  22.849  1.00 41.74      H    C
ATOM   7139  CB  GLU   158     -21.058  27.665  22.739  1.00 40.97      H    C
ATOM   7140  CG  GLU   158     -21.620  28.752  23.634  1.00 42.62      H    C
ATOM   7141  CD  GLU   158     -22.272  28.194  24.893  1.00 45.45      H    C
ATOM   7142  OE1 GLU   158     -21.641  27.366  25.594  1.00 47.64      H    O
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ATOM   7143  OE2 GLU   158     -23.423  28.583  25.182  1.00 45.92      H    O
ATOM   7144  C   GLU   158     -19.135  26.202  22.117  1.00 41.79      H    C
ATOM   7145  O   GLU   158     -18.668  26.257  20.981  1.00 43.21      H    O
ATOM   7146  N   PRO   159     -19.324  25.031  22.746  1.00 43.12      H    N
ATOM   7147  CD  PRO   159     -19.151  23.773  21.993  1.00 40.80      H    C
ATOM   7148  CA  PRO   159     -19.854  24.763  24.090  1.00 42.75      H    C
ATOM   7149  CB  PRO   159     -20.732  23.545  23.861  1.00 41.21      H    C
ATOM   7150  CG  PRO   159     -19.907  22.746  22.852  1.00 42.04      H    C
ATOM   7151  C   PRO   159     -18.755  24.454  25.116  1.00 43.47      H    C
ATOM   7152  O   PRO   159     -17.610  24.204  24.748  1.00 43.74      H    O
ATOM   7153  N   VAL   160     -19.112  24.440  26.399  1.00 44.59      H    N
ATOM   7154  CA  VAL   160     -18.333  23.682  27.381  1.00 45.26      H    C
ATOM   7155  CB  VAL   160     -17.819  24.563  28.546  1.00 44.95      H    C
ATOM   7156  CG1 VAL   160     -17.135  25.800  28.015  1.00 45.98      H    C
ATOM   7157  CG2 VAL   160     -18.960  24.931  29.455  1.00 47.14      H    C
ATOM   7158  C   VAL   160     -19.223  22.598  27.980  1.00 45.46      H    C
ATOM   7159  O   VAL   160     -20.436  22.770  28.090  1.00 45.76      H    O
ATOM   7160  N   THR   161     -18.621  21.479  28.358  1.00 44.80      H    N
ATOM   7161  CA  THR   161     -19.341  20.457  29.095  1.00 45.47      H    C
ATOM   7162  CB  THR   161     -19.128  19.065  28.470  1.00 47.53      H    C
ATOM   7163  OG1 THR   161     -17.767  18.658  28.658  1.00 50.04      H    O
ATOM   7164  CG2 THR   161     -19.424  19.101  26.973  1.00 47.52      H    C
ATOM   7165  C   THR   161     -18.838  20.440  30.536  1.00 44.68      H    C
ATOM   7166  O   THR   161     -17.665  20.714  30.800  1.00 44.39      H    O
ATOM   7167  N   VAL   162     -19.733  20.131  31.468  1.00 43.69      H    N
ATOM   7168  CA  VAL   162     -19.364  20.021  32.874  1.00 42.46      H    C
ATOM   7169  CB  VAL   162     -19.983  21.165  33.710  1.00 41.02      H    C
ATOM   7170  CG1 VAL   162     -19.557  21.033  35.159  1.00 39.15      H    C
ATOM   7171  CG2 VAL   162     -19.553  22.510  33.154  1.00 39.60      H    C
ATOM   7172  C   VAL   162     -19.853  18.692  33.439  1.00 42.98      H    C
ATOM   7173  O   VAL   162     -21.023  18.344  33.302  1.00 44.77      H    O
ATOM   7174  N   SER   163     -18.955  17.947  34.067  1.00 41.43      H    N
ATOM   7175  CA  SER   163     -19.347  16.736  34.767  1.00 41.82      H    C
ATOM   7176  CB  SER   163     -18.732  15.503  34.095  1.00 43.74      H    C
ATOM   7177  OG  SER   163     -17.328  15.465  34.281  1.00 47.54      H    O
ATOM   7178  C   SER   163     -18.854  16.852  36.199  1.00 40.92      H    C
ATOM   7179  O   SER   163     -18.027  17.710  36.505  1.00 40.33      H    O
ATOM   7180  N   TRP   164     -19.366  16.003  37.081  1.00 39.41      H    N
ATOM   7181  CA  TRP   164     -18.936  16.031  38.469  1.00 39.92      H    C
ATOM   7182  CB  TRP   164     -20.109  16.427  39.367  1.00 38.97      H    C
ATOM   7183  CG  TRP   164     -20.453  17.878  39.251  1.00 36.68      H    C
ATOM   7184  CD2 TRP   164     -19.970  18.936  40.083  1.00 35.35      H    C
ATOM   7185  CE2 TRP   164     -20.531  20.138  39.599  1.00 35.70      H    C
ATOM   7186  CE3 TRP   164     -19.117  18.986  41.189  1.00 35.83      H    C
ATOM   7187  CD1 TRP   164     -21.268  18.463  38.320  1.00 36.06      H    C
ATOM   7188  NE1 TRP   164     -21.319  19.823  38.524  1.00 34.61      H    N
ATOM   7189  CZ2 TRP   164     -20.265  21.372  40.185  1.00 36.66      H    C
ATOM   7190  CZ3 TRP   164     -18.853  20.214  41.770  1.00 38.04      H    C
ATOM   7191  CH2 TRP   164     -19.427  21.392  41.266  1.00 37.96      H    C
ATOM   7192  C   TRP   164     -18.365  14.690  38.899  1.00 41.57      H    C
ATOM   7193  O   TRP   164     -18.923  13.641  38.579  1.00 41.54      H    O
ATOM   7194  N   ASN   165     -17.247  14.734  39.623  1.00 43.99      H    N
ATOM   7195  CA  ASN   165     -16.524  13.531  40.015  1.00 45.28      H    C
ATOM   7196  CB  ASN   165     -17.216  12.870  41.211  1.00 44.38      H    C
ATOM   7197  CG  ASN   165     -17.097  13.698  42.477  1.00 44.74      H    C
ATOM   7198  OD1 ASN   165     -16.354  14.679  42.515  1.00 48.85      H    O
ATOM   7199  ND2 ASN   165     -17.823  13.309  43.520  1.00 43.07      H    N
ATOM   7200  C   ASN   165     -16.426  12.552  38.848  1.00 48.10      H    C
ATOM   7201  O   ASN   165     -16.682  11.358  38.998  1.00 48.86      H    O
ATOM   7202  N   SER   166     -16.071  13.075  37.679  1.00 50.62      H    N
ATOM   7203  CA  SER   166     -15.795  12.251  36.507  1.00 54.00      H    C
ATOM   7204  CB  SER   166     -14.620  11.310  36.787  1.00 54.39      H    C
ATOM   7205  OG  SER   166     -13.465  12.038  37.171  1.00 56.87      H    O
ATOM   7206  C   SER   166     -17.003  11.436  36.065  1.00 55.66      H    C
ATOM   7207  O   SER   166     -16.854  10.353  35.505  1.00 56.97      H    O
ATOM   7208  N   GLY   167     -18.199  11.958  36.317  1.00 56.98      H    N
ATOM   7209  CA  GLY   167     -19.403  11.303  35.835  1.00 56.82      H    C
ATOM   7210  C   GLY   167     -20.035  10.371  36.853  1.00 57.13      H    C
ATOM   7211  O   GLY   167     -21.137   9.865  36.640  1.00 57.48      H    O
ATOM   7212  N   ALA   168     -19.340  10.141  37.962  1.00 55.72      H    N
ATOM   7213  CA  ALA   168     -19.868   9.296  39.024  1.00 56.14      H    C
ATOM   7214  CB  ALA   168     -18.762   8.958  40.018  1.00 54.67      H    C
ATOM   7215  C   ALA   168     -21.030   9.980  39.746  1.00 56.76      H    C
ATOM   7216  O   ALA   168     -21.869   9.316  40.359  1.00 57.75      H    O
ATOM   7217  N   LEU   169     -21.073  11.308  39.676  1.00 55.50      H    N
ATOM   7218  CA  LEU   169     -22.129  12.076  40.325  1.00 53.11      H    C
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ATOM	7219	CB	LEU	169	-21.519	13.117	41.263	1.00	52.46	H	C
ATOM	7220	CG	LEU	169	-22.471	14.079	41.979	1.00	52.36	H	C
ATOM	7221	CD1	LEU	169	-23.459	13.291	42.831	1.00	50.70	H	C
ATOM	7222	CD2	LEU	169	-21.660	15.036	42.847	1.00	50.44	H	C
ATOM	7223	C	LEU	169	-22.989	12.761	39.273	1.00	52.77	H	C
ATOM	7224	O	LEU	169	-22.540	13.678	38.588	1.00	52.53	H	O
ATOM	7225	N	THR	170	-24.230	12.304	39.147	1.00	52.69	H	N
ATOM	7226	CA	THR	170	-25.126	12.796	38.110	1.00	52.29	H	C
ATOM	7227	CB	THR	170	-25.414	11.699	37.077	1.00	52.97	H	C
ATOM	7228	OG1	THR	170	-25.938	10.543	37.744	1.00	52.52	H	O
ATOM	7229	CG2	THR	170	-24.135	11.322	36.339	1.00	51.54	H	C
ATOM	7230	C	THR	170	-26.447	13.274	38.702	1.00	52.12	H	C
ATOM	7231	O	THR	170	-27.083	14.192	38.182	1.00	52.03	H	O
ATOM	7232	N	SER	171	-26.854	12.649	39.798	1.00	51.23	H	N
ATOM	7233	CA	SER	171	-28.080	13.042	40.469	1.00	50.24	H	C
ATOM	7234	CB	SER	171	-28.477	11.974	41.490	1.00	51.95	H	C
ATOM	7235	OG	SER	171	-29.732	12.268	42.075	1.00	55.96	H	O
ATOM	7236	C	SER	171	-27.898	14.392	41.164	1.00	48.92	H	C
ATOM	7237	O	SER	171	-26.913	14.607	41.879	1.00	48.18	H	O
ATOM	7238	N	GLY	172	-28.848	15.298	40.946	1.00	46.35	H	N
ATOM	7239	CA	GLY	172	-28.794	16.600	41.589	1.00	44.63	H	C
ATOM	7240	C	GLY	172	-27.951	17.617	40.839	1.00	43.58	H	C
ATOM	7241	O	GLY	172	-27.815	18.762	41.276	1.00	43.02	H	O
ATOM	7242	N	VAL	173	-27.381	17.206	39.711	1.00	41.66	H	N
ATOM	7243	CA	VAL	173	-26.556	18.099	38.907	1.00	40.61	H	C
ATOM	7244	CB	VAL	173	-25.533	17.313	38.052	1.00	40.49	H	C
ATOM	7245	CG1	VAL	173	-24.782	18.265	37.122	1.00	36.94	H	C
ATOM	7246	CG2	VAL	173	-24.556	16.584	38.959	1.00	38.36	H	C
ATOM	7247	C	VAL	173	-27.414	18.941	37.974	1.00	39.99	H	C
ATOM	7248	O	VAL	173	-28.250	18.415	37.240	1.00	39.43	H	O
ATOM	7249	N	HIS	174	-27.202	20.251	38.008	1.00	39.34	H	N
ATOM	7250	CA	HIS	174	-27.856	21.144	37.065	1.00	39.73	H	C
ATOM	7251	CB	HIS	174	-29.001	21.901	37.751	1.00	41.08	H	C
ATOM	7252	CG	HIS	174	-29.858	22.680	36.800	1.00	45.12	H	C
ATOM	7253	CD2	HIS	174	-29.786	22.828	35.455	1.00	46.87	H	C
ATOM	7254	ND1	HIS	174	-30.944	23.424	37.210	1.00	47.42	H	N
ATOM	7255	CE1	HIS	174	-31.506	23.997	36.159	1.00	46.85	H	C
ATOM	7256	NE2	HIS	174	-30.823	23.652	35.082	1.00	47.87	H	N
ATOM	7257	C	HIS	174	-26.850	22.135	36.485	1.00	38.58	H	C
ATOM	7258	O	HIS	174	-26.371	23.035	37.176	1.00	37.51	H	O
ATOM	7259	N	THR	175	-26.534	21.962	35.208	1.00	38.03	H	N
ATOM	7260	CA	THR	175	-25.664	22.893	34.507	1.00	36.61	H	C
ATOM	7261	CB	THR	175	-24.692	22.135	33.585	1.00	37.00	H	C
ATOM	7262	OG1	THR	175	-23.814	21.331	34.390	1.00	36.64	H	O
ATOM	7263	CG2	THR	175	-23.868	23.115	32.749	1.00	36.37	H	C
ATOM	7264	C	THR	175	-26.504	23.868	33.694	1.00	36.47	H	C
ATOM	7265	O	THR	175	-27.269	23.462	32.822	1.00	38.71	H	O
ATOM	7266	N	PHE	176	-26.370	25.155	33.994	1.00	34.80	H	N
ATOM	7267	CA	PHE	176	-27.245	26.165	33.416	1.00	33.49	H	C
ATOM	7268	CB	PHE	176	-27.366	27.370	34.350	1.00	29.26	H	C
ATOM	7269	CG	PHE	176	-28.218	27.117	35.554	1.00	29.36	H	C
ATOM	7270	CD1	PHE	176	-27.803	26.234	36.543	1.00	28.93	H	C
ATOM	7271	CD2	PHE	176	-29.437	27.761	35.701	1.00	26.76	H	C
ATOM	7272	CE1	PHE	176	-28.597	25.997	37.666	1.00	30.31	H	C
ATOM	7273	CE2	PHE	176	-30.233	27.532	36.814	1.00	28.16	H	C
ATOM	7274	CZ	PHE	176	-29.812	26.648	37.798	1.00	29.02	H	C
ATOM	7275	C	PHE	176	-26.734	26.636	32.068	1.00	34.89	H	C
ATOM	7276	O	PHE	176	-25.530	26.650	31.822	1.00	35.66	H	O
ATOM	7277	N	PRO	177	-27.652	27.033	31.175	1.00	35.42	H	N
ATOM	7278	CD	PRO	177	-29.113	26.855	31.270	1.00	34.38	H	C
ATOM	7279	CA	PRO	177	-27.251	27.589	29.881	1.00	35.34	H	C
ATOM	7280	CB	PRO	177	-28.583	27.873	29.182	1.00	34.43	H	C
ATOM	7281	CG	PRO	177	-29.548	26.907	29.830	1.00	35.56	H	C
ATOM	7282	C	PRO	177	-26.419	28.848	30.082	1.00	35.62	H	C
ATOM	7283	O	PRO	177	-26.689	29.640	30.986	1.00	37.58	H	O
ATOM	7284	N	ALA	178	-25.410	29.022	29.237	1.00	34.61	H	N
ATOM	7285	CA	ALA	178	-24.468	30.122	29.379	1.00	34.64	H	C
ATOM	7286	CB	ALA	178	-23.295	29.930	28.424	1.00	32.42	H	C
ATOM	7287	C	ALA	178	-25.129	31.466	29.121	1.00	34.20	H	C
ATOM	7288	O	ALA	178	-26.116	31.560	28.394	1.00	35.25	H	O
ATOM	7289	N	VAL	179	-24.583	32.507	29.734	1.00	33.24	H	N
ATOM	7290	CA	VAL	179	-24.940	33.864	29.372	1.00	34.36	H	C
ATOM	7291	CB	VAL	179	-24.935	34.784	30.613	1.00	35.45	H	C
ATOM	7292	CG1	VAL	179	-23.532	34.865	31.198	1.00	34.78	H	C
ATOM	7293	CG2	VAL	179	-25.460	36.158	30.246	1.00	33.91	H	C
ATOM	7294	C	VAL	179	-23.904	34.352	28.359	1.00	35.86	H	C

ATOM	7295	O	VAL	179	-22.734	33.962	28.413	1.00	34.73	H	O
ATOM	7296	N	LEU	180	-24.336	35.180	27.416	1.00	39.05	H	N
ATOM	7297	CA	LEU	180	-23.402	35.795	26.476	1.00	41.77	H	C
ATOM	7298	CB	LEU	180	-23.998	36.815	25.065	1.00	41.69	H	C
ATOM	7299	CG	LEU	180	-23.180	36.524	23.982	1.00	43.45	H	C
ATOM	7300	CD1	LEU	180	-21.808	35.879	23.862	1.00	43.12	H	C
ATOM	7301	CD2	LEU	180	-23.925	36.450	22.655	1.00	42.68	H	C
ATOM	7302	C	LEU	180	-23.112	37.213	26.939	1.00	42.76	H	C
ATOM	7303	O	LEU	180	-24.023	38.026	27.071	1.00	44.73	H	O
ATOM	7304	N	GLN	181	-21.845	37.511	27.202	1.00	44.83	H	N
ATOM	7305	CA	GLN	181	-21.494	38.815	27.750	1.00	47.27	H	C
ATOM	7306	CB	GLN	181	-20.189	38.727	28.534	1.00	48.86	H	C
ATOM	7307	CG	GLN	181	-20.135	37.607	29.554	1.00	52.26	H	C
ATOM	7308	CD	GLN	181	-18.767	37.504	30.198	1.00	53.89	H	C
ATOM	7309	OE1	GLN	181	-18.275	38.467	30.794	1.00	56.33	H	O
ATOM	7310	NE2	GLN	181	-18.140	36.341	30.075	1.00	51.83	H	N
ATOM	7311	C	GLN	181	-21.339	39.857	26.648	1.00	47.64	H	C
ATOM	7312	O	GLN	181	-21.008	39.525	25.508	1.00	45.43	H	O
ATOM	7313	N	SER	182	-21.569	41.117	27.005	1.00	48.50	H	N
ATOM	7314	CA	SER	182	-21.284	42.241	26.117	1.00	50.22	H	C
ATOM	7315	CB	SER	182	-21.346	43.553	26.897	1.00	50.75	H	C
ATOM	7316	OG	SER	182	-22.606	43.704	27.526	1.00	55.71	H	O
ATOM	7317	C	SER	182	-19.910	42.110	25.468	1.00	49.58	H	C
ATOM	7318	O	SER	182	-19.711	42.528	24.327	1.00	50.47	H	O
ATOM	7319	N	SER	183	-18.964	41.531	26.198	1.00	47.76	H	N
ATOM	7320	CA	SER	183	-17.606	41.360	25.693	1.00	46.55	H	C
ATOM	7321	CB	SER	183	-16.691	40.844	26.803	1.00	46.41	H	C
ATOM	7322	OG	SER	183	-16.966	39.479	27.074	1.00	47.06	H	O
ATOM	7323	C	SER	183	-17.558	40.375	24.533	1.00	45.50	H	C
ATOM	7324	O	SER	183	-16.549	40.289	23.830	1.00	47.15	H	O
ATOM	7325	N	GLY	184	-18.634	39.610	24.354	1.00	43.61	H	N
ATOM	7326	CA	GLY	184	-18.630	38.554	23.354	1.00	41.62	H	C
ATOM	7327	C	GLY	184	-18.155	37.206	23.873	1.00	41.63	H	C
ATOM	7328	O	GLY	184	-18.108	36.225	23.127	1.00	42.12	H	O
ATOM	7329	N	LEU	185	-17.797	37.151	25.154	1.00	41.06	H	N
ATOM	7330	CA	LEU	185	-17.391	35.896	25.782	1.00	39.48	H	C
ATOM	7331	CB	LEU	185	-16.222	36.147	26.740	1.00	40.59	H	C
ATOM	7332	CG	LEU	185	-14.954	36.763	26.134	1.00	41.75	H	C
ATOM	7333	CD1	LEU	185	-13.943	37.034	27.238	1.00	41.08	H	C
ATOM	7334	CD2	LEU	185	-14.363	35.829	25.088	1.00	40.63	H	C
ATOM	7335	C	LEU	185	-18.566	35.270	26.541	1.00	37.17	H	C
ATOM	7336	O	LEU	185	-19.534	35.956	26.870	1.00	36.34	H	O
ATOM	7337	N	TYR	186	-18.479	33.969	26.811	1.00	35.21	H	N
ATOM	7338	CA	TYR	186	-19.544	33.257	27.520	1.00	35.82	H	C
ATOM	7339	CB	TYR	186	-19.905	31.958	26.790	1.00	35.22	H	C
ATOM	7340	CG	TYR	186	-20.484	32.132	25.400	1.00	38.74	H	C
ATOM	7341	CD1	TYR	186	-21.859	32.213	25.201	1.00	39.50	H	C
ATOM	7342	CE1	TYR	186	-22.393	32.334	23.930	1.00	41.29	H	C
ATOM	7343	CD2	TYR	186	-19.658	32.181	24.286	1.00	37.63	H	C
ATOM	7344	CE2	TYR	186	-20.178	32.304	23.018	1.00	40.69	H	C
ATOM	7345	CZ	TYR	186	-21.545	32.378	22.840	1.00	43.16	H	C
ATOM	7346	OH	TYR	186	-22.060	32.482	21.565	1.00	45.56	H	O
ATOM	7347	C	TYR	186	-19.137	32.901	28.956	1.00	35.42	H	C
ATOM	7348	O	TYR	186	-17.962	32.669	29.233	1.00	36.05	H	O
ATOM	7349	N	SER	187	-20.118	32.841	29.854	1.00	33.07	H	N
ATOM	7350	CA	SER	187	-19.939	32.219	31.166	1.00	33.69	H	C
ATOM	7351	CB	SER	187	-19.760	33.293	32.249	1.00	30.64	H	C
ATOM	7352	OG	SER	187	-18.614	34.090	31.992	1.00	31.70	H	O
ATOM	7353	C	SER	187	-21.157	31.362	31.508	1.00	33.84	H	C
ATOM	7354	O	SER	187	-22.288	31.709	31.153	1.00	33.92	H	O
ATOM	7355	N	LEU	188	-20.932	30.247	32.198	1.00	33.13	H	N
ATOM	7356	CA	LEU	188	-22.041	29.488	32.762	1.00	32.13	H	C
ATOM	7357	CB	LEU	188	-22.372	28.284	31.880	1.00	30.17	H	C
ATOM	7358	CG	LEU	188	-21.331	27.175	31.701	1.00	32.49	H	C
ATOM	7359	CD1	LEU	188	-21.234	26.317	32.963	1.00	30.00	H	C
ATOM	7360	CD2	LEU	188	-21.747	26.310	30.518	1.00	29.79	H	C
ATOM	7361	C	LEU	188	-21.753	29.023	34.183	1.00	33.34	H	C
ATOM	7362	O	LEU	188	-20.650	29.200	34.700	1.00	33.47	H	O
ATOM	7363	N	SER	189	-22.758	28.430	34.814	1.00	32.89	H	N
ATOM	7364	CA	SER	189	-22.580	27.833	36.125	1.00	32.77	H	C
ATOM	7365	CB	SER	189	-23.270	28.675	37.201	1.00	31.98	H	C
ATOM	7366	OG	SER	189	-22.656	29.947	37.313	1.00	33.26	H	O
ATOM	7367	C	SER	189	-23.159	26.435	36.123	1.00	33.45	H	C
ATOM	7368	O	SER	189	-24.163	26.164	35.458	1.00	32.42	H	O
ATOM	7369	N	SER	190	-22.502	25.546	36.856	1.00	34.16	H	N
ATOM	7370	CA	SER	190	-23.029	24.220	37.108	1.00	34.97	H	C

ATOM	7371	CB	SER	190	-22.098	23.157	36.521	1.00	35.56	H	C
ATOM	7372	OG	SER	190	-22.601	21.852	36.764	1.00	36.81	H	O
ATOM	7373	C	SER	190	-23.130	24.049	38.617	1.00	36.41	H	C
ATOM	7374	O	SER	190	-22.185	24.356	39.350	1.00	35.33	H	O
ATOM	7375	N	VAL	191	-24.281	23.571	39.075	1.00	36.33	H	N
ATOM	7376	CA	VAL	191	-24.522	23.382	40.495	1.00	38.43	H	C
ATOM	7377	CB	VAL	191	-25.583	24.369	41.013	1.00	39.52	H	C
ATOM	7378	CG1	VAL	191	-25.902	24.085	42.471	1.00	41.66	H	C
ATOM	7379	CG2	VAL	191	-25.066	25.774	40.876	1.00	41.62	H	C
ATOM	7380	C	VAL	191	-24.998	21.968	40.761	1.00	39.28	H	C
ATOM	7381	O	VAL	191	-25.663	21.358	39.927	1.00	40.77	H	O
ATOM	7382	N	VAL	192	-24.647	21.437	41.924	1.00	39.29	H	N
ATOM	7383	CA	VAL	192	-25.145	20.134	42.316	1.00	38.27	H	C
ATOM	7384	CB	VAL	192	-24.027	19.066	42.249	1.00	37.49	H	C
ATOM	7385	CG1	VAL	192	-22.875	19.462	43.149	1.00	39.66	H	C
ATOM	7386	CG2	VAL	192	-24.579	17.708	42.650	1.00	38.15	H	C
ATOM	7387	C	VAL	192	-25.700	20.223	43.724	1.00	37.59	H	C
ATOM	7388	O	VAL	192	-25.123	20.882	44.587	1.00	35.77	H	O
ATOM	7389	N	THR	193	-26.838	19.576	43.944	1.00	39.03	H	N
ATOM	7390	CA	THR	193	-27.465	19.546	45.264	1.00	40.19	H	C
ATOM	7391	CB	THR	193	-29.000	19.651	45.142	1.00	42.63	H	C
ATOM	7392	OG1	THR	193	-29.342	20.941	44.618	1.00	45.54	H	O
ATOM	7393	CG2	THR	193	-29.675	19.459	46.504	1.00	41.23	H	C
ATOM	7394	C	THR	193	-27.102	18.246	45.967	1.00	38.72	H	C
ATOM	7395	O	THR	193	-27.343	17.163	45.438	1.00	38.35	H	O
ATOM	7396	N	VAL	194	-26.508	18.357	47.150	1.00	38.96	H	N
ATOM	7397	CA	VAL	194	-25.998	17.183	47.862	1.00	40.70	H	C
ATOM	7398	CB	VAL	194	-24.452	17.105	47.801	1.00	38.53	H	C
ATOM	7399	CG1	VAL	194	-23.986	17.097	46.362	1.00	39.05	H	C
ATOM	7400	CG2	VAL	194	-23.841	18.271	48.566	1.00	34.97	H	C
ATOM	7401	C	VAL	194	-26.402	17.197	49.333	1.00	42.41	H	C
ATOM	7402	O	VAL	194	-26.782	18.239	49.876	1.00	42.66	H	O
ATOM	7403	N	PRO	195	-26.315	16.035	50.003	1.00	44.22	H	N
ATOM	7404	CD	PRO	195	-26.009	14.700	49.463	1.00	45.72	H	C
ATOM	7405	CA	PRO	195	-26.603	15.989	51.439	1.00	46.23	H	C
ATOM	7406	CB	PRO	195	-26.475	14.506	51.786	1.00	47.19	H	C
ATOM	7407	CG	PRO	195	-26.632	13.788	50.476	1.00	47.45	H	C
ATOM	7408	C	PRO	195	-25.585	16.838	52.191	1.00	47.52	H	C
ATOM	7409	O	PRO	195	-24.381	16.671	52.005	1.00	48.34	H	O
ATOM	7410	N	SER	196	-26.054	17.751	53.031	1.00	47.83	H	N
ATOM	7411	CA	SER	196	-25.130	18.582	53.787	1.00	52.32	H	C
ATOM	7412	CB	SER	196	-25.879	19.705	54.514	1.00	52.90	H	C
ATOM	7413	OG	SER	196	-26.802	19.189	55.450	1.00	57.28	H	O
ATOM	7414	C	SER	196	-24.354	17.737	54.789	1.00	53.77	H	C
ATOM	7415	O	SER	196	-23.330	18.170	55.317	1.00	54.68	H	O
ATOM	7416	N	SER	197	-24.838	16.526	55.045	1.00	55.89	H	N
ATOM	7417	CA	SER	197	-24.154	15.615	55.957	1.00	58.39	H	C
ATOM	7418	CB	SER	197	-25.098	14.489	56.397	1.00	57.25	H	C
ATOM	7419	OG	SER	197	-25.329	13.570	55.345	1.00	56.57	H	O
ATOM	7420	C	SER	197	-22.912	15.020	55.292	1.00	59.35	H	C
ATOM	7421	O	SER	197	-21.974	14.607	55.970	1.00	59.77	H	O
ATOM	7422	N	SER	198	-22.909	14.984	53.963	1.00	60.41	H	N
ATOM	7423	CA	SER	198	-21.781	14.441	53.213	1.00	60.89	H	C
ATOM	7424	CB	SER	198	-22.231	14.008	51.818	1.00	59.87	H	C
ATOM	7425	OG	SER	198	-22.374	15.135	50.967	1.00	60.91	H	O
ATOM	7426	C	SER	198	-20.659	15.470	53.080	1.00	61.53	H	C
ATOM	7427	O	SER	198	-19.515	15.116	52.807	1.00	62.27	H	O
ATOM	7428	N	LEU	199	-20.989	16.744	53.264	1.00	62.42	H	N
ATOM	7429	CA	LEU	199	-19.986	17.799	53.170	1.00	63.79	H	C
ATOM	7430	CB	LEU	199	-20.591	19.164	53.518	1.00	60.27	H	C
ATOM	7431	CG	LEU	199	-21.688	19.724	52.607	1.00	58.84	H	C
ATOM	7432	CD1	LEU	199	-22.169	21.058	53.158	1.00	56.44	H	C
ATOM	7433	CD2	LEU	199	-21.159	19.886	51.190	1.00	56.85	H	C
ATOM	7434	C	LEU	199	-18.859	17.490	54.142	1.00	66.35	H	C
ATOM	7435	O	LEU	199	-19.092	17.296	55.337	1.00	68.30	H	O
ATOM	7436	N	GLY	200	-17.635	17.437	53.634	1.00	67.32	H	N
ATOM	7437	CA	GLY	200	-16.505	17.234	54.519	1.00	69.17	H	C
ATOM	7438	C	GLY	200	-16.023	15.799	54.633	1.00	68.88	H	C
ATOM	7439	O	GLY	200	-14.974	15.556	55.224	1.00	70.04	H	O
ATOM	7440	N	THR	201	-16.771	14.846	54.083	1.00	67.65	H	N
ATOM	7441	CA	THR	201	-16.241	13.493	53.930	1.00	67.09	H	C
ATOM	7442	CB	THR	201	-17.079	12.444	54.702	1.00	67.94	H	C
ATOM	7443	OG1	THR	201	-18.284	12.155	53.981	1.00	69.01	H	O
ATOM	7444	CG2	THR	201	-17.430	12.967	56.094	1.00	67.66	H	C
ATOM	7445	C	THR	201	-16.217	13.108	52.456	1.00	65.62	H	C
ATOM	7446	O	THR	201	-15.265	12.486	51.984	1.00	66.99	H	O

ATOM	7447	N	GLN	202	-17.265	13.482	51.731	1.00	63.03	H	N
ATOM	7448	CA	GLN	202	-17.288	13.296	50.286	1.00	60.90	H	C
ATOM	7449	CB	GLN	202	-18.726	13.167	49.795	1.00	61.55	H	C
ATOM	7450	CG	GLN	202	-18.853	13.129	48.269	1.00	64.43	H	C
ATOM	7451	CD	GLN	202	-18.191	11.913	47.677	1.00	66.51	H	C
ATOM	7452	OE1	GLN	202	-18.749	10.814	47.694	1.00	67.79	H	O
ATOM	7453	NE2	GLN	202	-16.994	12.103	47.129	1.00	66.84	H	N
ATOM	7454	C	GLN	202	-16.617	14.487	49.607	1.00	58.78	H	C
ATOM	7455	O	GLN	202	-16.892	15.641	49.936	1.00	58.19	H	O
ATOM	7456	N	THR	203	-15.725	14.214	48.666	1.00	55.82	H	N
ATOM	7457	CA	THR	203	-15.092	15.299	47.934	1.00	53.36	H	C
ATOM	7458	CB	THR	203	-13.615	14.991	47.617	1.00	54.58	H	C
ATOM	7459	OG1	THR	203	-13.537	13.811	46.810	1.00	55.09	H	O
ATOM	7460	CG2	THR	203	-12.826	14.789	48.903	1.00	54.39	H	C
ATOM	7461	C	THR	203	-15.838	15.553	46.631	1.00	49.49	H	C
ATOM	7462	O	THR	203	-16.299	14.617	45.965	1.00	47.29	H	O
ATOM	7463	N	TYR	204	-15.962	16.827	46.278	1.00	45.01	H	N
ATOM	7464	CA	TYR	204	-16.675	17.202	45.068	1.00	42.84	H	C
ATOM	7465	CB	TYR	204	-17.884	18.070	45.427	1.00	40.12	H	C
ATOM	7466	CG	TYR	204	-18.919	17.314	46.224	1.00	38.49	H	C
ATOM	7467	CD1	TYR	204	-19.065	17.527	47.590	1.00	38.48	H	C
ATOM	7468	CE1	TYR	204	-19.994	16.806	48.330	1.00	38.92	H	C
ATOM	7469	CD2	TYR	204	-19.731	16.362	45.615	1.00	38.50	H	C
ATOM	7470	CE2	TYR	204	-20.659	15.639	46.343	1.00	38.51	H	C
ATOM	7471	CZ	TYR	204	-20.785	15.866	47.699	1.00	38.77	H	C
ATOM	7472	OH	TYR	204	-21.706	15.148	48.425	1.00	41.78	H	O
ATOM	7473	C	TYR	204	-15.762	17.924	44.097	1.00	41.23	H	C
ATOM	7474	O	TYR	204	-15.124	18.925	44.436	1.00	40.31	H	O
ATOM	7475	N	ILE	205	-15.694	17.395	42.885	1.00	41.27	H	N
ATOM	7476	CA	ILE	205	-14.812	17.944	41.868	1.00	41.45	H	C
ATOM	7477	CB	ILE	205	-13.640	16.985	41.577	1.00	42.17	H	C
ATOM	7478	CG2	ILE	205	-12.732	17.578	40.499	1.00	40.04	H	C
ATOM	7479	CG1	ILE	205	-12.850	16.734	42.865	1.00	41.38	H	C
ATOM	7480	CD1	ILE	205	-11.695	15.767	42.693	1.00	41.84	H	C
ATOM	7481	C	ILE	205	-15.601	18.158	40.591	1.00	41.22	H	C
ATOM	7482	O	ILE	205	-16.240	17.236	40.083	1.00	41.08	H	O
ATOM	7483	N	CYS	206	-15.563	19.373	40.067	1.00	40.41	H	N
ATOM	7484	CA	CYS	206	-16.252	19.629	38.819	1.00	42.57	H	C
ATOM	7485	C	CYS	206	-15.242	19.546	37.667	1.00	42.00	H	C
ATOM	7486	O	CYS	206	-14.135	20.079	37.752	1.00	41.39	H	O
ATOM	7487	CB	CYS	206	-16.974	20.992	38.895	1.00	45.38	H	C
ATOM	7488	SG	CYS	206	-16.108	22.456	38.247	1.00	54.25	H	S
ATOM	7489	N	ASN	207	-15.613	18.832	36.609	1.00	40.73	H	N
ATOM	7490	CA	ASN	207	-14.714	18.599	35.482	1.00	40.35	H	C
ATOM	7491	CB	ASN	207	-14.680	17.109	35.124	1.00	39.81	H	C
ATOM	7492	CG	ASN	207	-14.565	16.216	36.350	1.00	42.88	H	C
ATOM	7493	OD1	ASN	207	-15.521	15.533	36.725	1.00	42.86	H	O
ATOM	7494	ND2	ASN	207	-13.392	16.217	36.981	1.00	40.06	H	N
ATOM	7495	C	ASN	207	-15.198	19.398	34.281	1.00	39.79	H	C
ATOM	7496	O	ASN	207	-16.219	19.070	33.677	1.00	39.54	H	O
ATOM	7497	N	VAL	208	-14.456	20.447	33.941	1.00	38.89	H	N
ATOM	7498	CA	VAL	208	-14.853	21.353	32.875	1.00	39.31	H	C
ATOM	7499	CB	VAL	208	-14.612	22.825	33.288	1.00	39.03	H	C
ATOM	7500	CG1	VAL	208	-14.979	23.761	32.147	1.00	36.52	H	C
ATOM	7501	CG2	VAL	208	-15.432	23.154	34.533	1.00	36.09	H	C
ATOM	7502	C	VAL	208	-14.069	21.061	31.604	1.00	41.22	H	C
ATOM	7503	O	VAL	208	-12.839	21.053	31.611	1.00	43.41	H	O
ATOM	7504	N	ASN	209	-14.782	20.821	30.511	1.00	43.06	H	N
ATOM	7505	CA	ASN	209	-14.136	20.546	29.234	1.00	44.34	H	C
ATOM	7506	CB	ASN	209	-14.500	19.136	28.756	1.00	47.58	H	C
ATOM	7507	CG	ASN	209	-13.650	18.677	27.578	1.00	54.59	H	C
ATOM	7508	OD1	ASN	209	-12.994	19.486	26.913	1.00	57.77	H	O
ATOM	7509	ND2	ASN	209	-13.657	17.372	27.316	1.00	55.14	H	N
ATOM	7510	C	ASN	209	-14.556	21.583	28.196	1.00	43.67	H	C
ATOM	7511	O	ASN	209	-15.745	21.776	27.942	1.00	44.34	H	O
ATOM	7512	N	HIS	210	-13.573	22.261	27.612	1.00	42.73	H	N
ATOM	7513	CA	HIS	210	-13.823	23.227	26.548	1.00	42.37	H	C
ATOM	7514	CB	HIS	210	-13.448	24.636	27.019	1.00	41.38	H	C
ATOM	7515	CG	HIS	210	-13.788	25.717	26.040	1.00	40.95	H	C
ATOM	7516	CD2	HIS	210	-14.971	26.086	25.494	1.00	40.98	H	C
ATOM	7517	ND1	HIS	210	-12.845	26.584	25.529	1.00	42.03	H	N
ATOM	7518	CE1	HIS	210	-13.432	27.442	24.712	1.00	40.17	H	C
ATOM	7519	NE2	HIS	210	-14.722	27.161	24.673	1.00	41.17	H	N
ATOM	7520	C	HIS	210	-12.987	22.842	25.331	1.00	43.48	H	C
ATOM	7521	O	HIS	210	-11.847	23.286	25.185	1.00	43.46	H	O
ATOM	7522	N	LYS	211	-13.558	22.018	24.455	1.00	44.53	H	N

ATOM	7523	CA	LYS	211	-12.811	21.468	23.328	1.00	45.35	H	C
ATOM	7524	CB	LYS	211	-13.644	20.393	22.623	1.00	48.43	H	C
ATOM	7525	CG	LYS	211	-13.903	19.179	23.508	1.00	54.24	H	C
ATOM	7526	CD	LYS	211	-14.558	18.028	22.751	1.00	59.93	H	C
ATOM	7527	CE	LYS	211	-14.674	16.781	23.638	1.00	63.06	H	C
ATOM	7528	NZ	LYS	211	-15.144	15.575	22.888	1.00	64.57	H	N
ATOM	7529	C	LYS	211	-12.313	22.501	22.317	1.00	41.93	H	C
ATOM	7530	O	LYS	211	-11.200	22.387	21.816	1.00	43.23	H	O
ATOM	7531	N	PRO	212	-13.113	23.534	22.020	1.00	40.22	H	N
ATOM	7532	CD	PRO	212	-14.467	23.852	22.506	1.00	38.83	H	C
ATOM	7533	CA	PRO	212	-12.628	24.526	21.052	1.00	41.43	H	C
ATOM	7534	CB	PRO	212	-13.702	25.618	21.083	1.00	39.54	H	C
ATOM	7535	CG	PRO	212	-14.939	24.906	21.540	1.00	37.50	H	C
ATOM	7536	C	PRO	212	-11.229	25.079	21.373	1.00	44.16	H	C
ATOM	7537	O	PRO	212	-10.457	25.394	20.464	1.00	46.27	H	O
ATOM	7538	N	SER	213	-10.903	25.187	22.661	1.00	44.31	H	N
ATOM	7539	CA	SER	213	-9.593	25.691	23.077	1.00	43.30	H	C
ATOM	7540	CB	SER	213	-9.751	26.781	24.139	1.00	43.36	H	C
ATOM	7541	OG	SER	213	-10.185	26.231	25.375	1.00	42.70	H	O
ATOM	7542	C	SER	213	-8.690	24.594	23.630	1.00	43.68	H	C
ATOM	7543	O	SER	213	-7.572	24.870	24.062	1.00	44.07	H	O
ATOM	7544	N	ASN	214	-9.172	23.357	23.630	1.00	43.44	H	N
ATOM	7545	CA	ASN	214	-8.372	22.234	24.110	1.00	46.08	H	C
ATOM	7546	CB	ASN	214	-7.097	22.093	23.272	1.00	48.48	H	C
ATOM	7547	CG	ASN	214	-7.388	21.690	21.832	1.00	52.65	H	C
ATOM	7548	OD1	ASN	214	-7.140	22.457	20.895	1.00	52.37	H	O
ATOM	7549	ND2	ASN	214	-7.919	20.483	21.652	1.00	52.33	H	N
ATOM	7550	C	ASN	214	-7.999	22.388	25.584	1.00	46.19	H	C
ATOM	7551	O	ASN	214	-6.891	22.046	25.996	1.00	45.93	H	O
ATOM	7552	N	THR	215	-8.935	22.904	26.372	1.00	45.35	H	N
ATOM	7553	CA	THR	215	-8.716	23.108	27.794	1.00	44.99	H	C
ATOM	7554	CB	THR	215	-9.062	24.553	28.198	1.00	44.21	H	C
ATOM	7555	OG1	THR	215	-8.385	25.468	27.329	1.00	44.83	H	O
ATOM	7556	CG2	THR	215	-8.635	24.822	29.632	1.00	43.77	H	C
ATOM	7557	C	THR	215	-9.599	22.162	28.608	1.00	45.50	H	C
ATOM	7558	O	THR	215	-10.815	22.122	28.414	1.00	45.25	H	O
ATOM	7559	N	LYS	216	-8.985	21.402	29.511	1.00	44.26	H	N
ATOM	7560	CA	LYS	216	-9.725	20.736	30.577	1.00	45.60	H	C
ATOM	7561	CB	LYS	216	-9.439	19.234	30.595	1.00	47.96	H	C
ATOM	7562	CG	LYS	216	-10.062	18.437	29.460	1.00	52.62	H	C
ATOM	7563	CD	LYS	216	-9.851	16.937	29.685	1.00	55.74	H	C
ATOM	7564	CE	LYS	216	-10.309	16.114	28.487	1.00	58.57	H	C
ATOM	7565	NZ	LYS	216	-9.548	16.453	27.249	1.00	60.46	H	N
ATOM	7566	C	LYS	216	-9.311	21.332	31.913	1.00	44.05	H	C
ATOM	7567	O	LYS	216	-8.132	21.579	32.144	1.00	43.16	H	O
ATOM	7568	N	VAL	217	-10.282	21.562	32.790	1.00	43.83	H	N
ATOM	7569	CA	VAL	217	-9.998	22.049	34.134	1.00	42.88	H	C
ATOM	7570	CB	VAL	217	-10.428	23.517	34.297	1.00	42.10	H	C
ATOM	7571	CG1	VAL	217	-10.096	24.003	35.702	1.00	40.43	H	C
ATOM	7572	CG2	VAL	217	-9.743	24.379	33.250	1.00	41.13	H	C
ATOM	7573	C	VAL	217	-10.734	21.216	35.178	1.00	44.48	H	C
ATOM	7574	O	VAL	217	-11.927	20.941	35.034	1.00	44.28	H	O
ATOM	7575	N	ASP	218	-10.021	20.812	36.224	1.00	45.43	H	N
ATOM	7576	CA	ASP	218	-10.647	20.153	37.368	1.00	46.54	H	C
ATOM	7577	CB	ASP	218	-9.980	18.810	37.663	1.00	49.28	H	C
ATOM	7578	CG	ASP	218	-10.346	17.743	36.657	1.00	54.06	H	C
ATOM	7579	OD1	ASP	218	-11.245	17.982	35.822	1.00	56.26	H	O
ATOM	7580	OD2	ASP	218	-9.732	16.658	36.701	1.00	59.11	H	O
ATOM	7581	C	ASP	218	-10.534	21.038	38.593	1.00	45.42	H	C
ATOM	7582	O	ASP	218	-9.450	21.516	38.925	1.00	45.41	H	O
ATOM	7583	N	LYS	219	-11.658	21.254	39.265	1.00	43.94	H	N
ATOM	7584	CA	LYS	219	-11.679	22.099	40.444	1.00	43.70	H	C
ATOM	7585	CB	LYS	219	-12.358	23.429	40.119	1.00	43.41	H	C
ATOM	7586	CG	LYS	219	-12.481	24.361	41.302	1.00	45.72	H	C
ATOM	7587	CD	LYS	219	-11.136	24.958	41.689	1.00	46.20	H	C
ATOM	7588	CE	LYS	219	-10.569	25.794	40.560	1.00	45.39	H	C
ATOM	7589	NZ	LYS	219	-9.622	26.809	41.087	1.00	48.37	H	N
ATOM	7590	C	LYS	219	-12.416	21.400	41.578	1.00	44.16	H	C
ATOM	7591	O	LYS	219	-13.581	21.035	41.442	1.00	44.32	H	O
ATOM	7592	N	LYS	220	-11.728	21.210	42.697	1.00	44.00	H	N
ATOM	7593	CA	LYS	220	-12.356	20.635	43.875	1.00	45.01	H	C
ATOM	7594	CB	LYS	220	-11.294	20.017	44.794	1.00	48.33	H	C
ATOM	7595	CG	LYS	220	-11.843	19.277	46.009	1.00	51.01	H	C
ATOM	7596	CD	LYS	220	-10.696	18.744	46.869	1.00	57.34	H	C
ATOM	7597	CE	LYS	220	-11.185	17.829	47.990	1.00	60.75	H	C
ATOM	7598	NZ	LYS	220	-11.920	18.562	49.066	1.00	62.49	H	N

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ATOM   7599  C   LYS   220     -13.092  21.748  44.601  1.00 43.21      H    C
ATOM   7600  O   LYS   220     -12.560  22.846  44.779  1.00 40.91      H    O
ATOM   7601  N   VAL   221     -14.324  21.468  45.007  1.00 42.53      H    N
ATOM   7602  CA  VAL   221     -15.142  22.475  45.666  1.00 43.36      H    C
ATOM   7603  CB  VAL   221     -16.521  22.606  44.975  1.00 40.89      H    C
ATOM   7604  CG1 VAL   221     -17.341  23.702  45.641  1.00 37.25      H    C
ATOM   7605  CG2 VAL   221     -16.330  22.905  43.489  1.00 37.55      H    C
ATOM   7606  C   VAL   221     -15.336  22.092  47.126  1.00 45.76      H    C
ATOM   7607  O   VAL   221     -15.936  21.064  47.428  1.00 45.44      H    O
ATOM   7608  N   GLU   222     -14.814  22.919  48.025  1.00 49.29      H    N
ATOM   7609  CA  GLU   222     -14.822  22.613  49.455  1.00 53.13      H    C
ATOM   7610  CB  GLU   222     -13.394  22.573  50.004  1.00 54.80      H    C
ATOM   7611  CG  GLU   222     -12.501  21.504  49.408  1.00 59.95      H    C
ATOM   7612  CD  GLU   222     -11.109  21.507  50.028  1.00 63.23      H    C
ATOM   7613  OE1 GLU   222     -10.566  20.411  50.209  1.00 65.91      H    O
ATOM   7614  OE2 GLU   222     -10.556  22.607  50.256  1.00 63.80      H    O
ATOM   7615  C   GLU   222     -15.609  23.656  50.234  1.00 53.98      H    C
ATOM   7616  O   GLU   222     -15.745  24.798  49.798  1.00 52.85      H    O
ATOM   7617  N   PRO   223     -16.124  23.271  51.412  1.00 55.55      H    N
ATOM   7618  CD  PRO   223     -16.157  21.876  51.886  1.00 55.58      H    C
ATOM   7619  CA  PRO   223     -16.809  24.179  52.339  1.00 57.83      H    C
ATOM   7620  CB  PRO   223     -17.228  23.267  53.492  1.00 57.45      H    C
ATOM   7621  CG  PRO   223     -17.264  21.894  52.896  1.00 56.88      H    C
ATOM   7622  C   PRO   223     -15.892  25.306  52.807  1.00 61.01      H    C
ATOM   7623  O   PRO   223     -14.674  25.152  52.821  1.00 61.88      H    O
ATOM   7624  N   LYS   224     -16.481  26.434  53.189  1.00 64.76      H    N
ATOM   7625  CA  LYS   224     -15.710  27.608  53.598  1.00 68.89      H    C
ATOM   7626  CB  LYS   224     -16.608  28.848  53.602  1.00 71.49      H    C
ATOM   7627  CG  LYS   224     -17.358  29.095  52.302  1.00 74.50      H    C
ATOM   7628  CD  LYS   224     -18.487  30.095  52.518  1.00 76.86      H    C
ATOM   7629  CE  LYS   224     -19.142  30.504  51.212  1.00 77.87      H    C
ATOM   7630  NZ  LYS   224     -20.308  31.401  51.451  1.00 78.94      H    N
ATOM   7631  C   LYS   224     -15.090  27.431  54.988  1.00 69.98      H    C
ATOM   7632  O   LYS   224     -13.862  27.636  55.125  1.00 70.34      H    O
ATOM   7633  OXT LYS   224     -15.848  27.105  55.928  1.00 70.58      H    O
TER    7634       LYS   224                                             H
ATOM   8057  NA  NA      1     -48.879  -0.173 -21.279  1.00 64.24      ION  N
TER    8058       NA      1                                             ION
END
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Table 35.2

ATOM	1	CB	THR	61	10.449	-40.746	-18.654	1.00	36.37	A	C
ATOM	2	OG1	THR	61	10.788	-42.078	-18.244	1.00	39.29	A	O
ATOM	3	CG2	THR	61	11.631	-39.827	-18.376	1.00	36.88	A	C
ATOM	4	C	THR	61	9.503	-39.926	-16.424	1.00	32.74	A	C
ATOM	5	O	THR	61	10.188	-38.932	-16.133	1.00	32.55	A	O
ATOM	6	N	THR	61	8.558	-39.083	-18.582	1.00	33.91	A	N
ATOM	7	CA	THR	61	9.165	-40.263	-17.892	1.00	34.36	A	C
ATOM	8	N	ALA	62	9.017	-40.768	-15.509	1.00	30.00	A	N
ATOM	9	CA	ALA	62	8.977	-40.452	-14.074	1.00	26.49	A	C
ATOM	10	CB	ALA	62	8.158	-41.501	-13.337	1.00	24.81	A	C
ATOM	11	C	ALA	62	10.345	-40.330	-13.425	1.00	24.06	A	C
ATOM	12	O	ALA	62	11.301	-40.970	-13.849	1.00	26.36	A	O
ATOM	13	N	THR	63	10.427	-39.513	-12.381	1.00	21.77	A	N
ATOM	14	CA	THR	63	11.687	-39.264	-11.691	1.00	18.83	A	C
ATOM	15	CB	THR	63	12.049	-37.757	-11.731	1.00	17.84	A	C
ATOM	16	OG1	THR	63	11.064	-37.000	-11.029	1.00	19.65	A	O
ATOM	17	CG2	THR	63	12.086	-37.256	-13.151	1.00	17.35	A	C
ATOM	18	C	THR	63	11.667	-39.741	-10.234	1.00	17.55	A	C
ATOM	19	O	THR	63	10.611	-39.991	-9.663	1.00	17.43	A	O
ATOM	20	N	PHE	64	12.851	-39.886	-9.649	1.00	18.91	A	N
ATOM	21	CA	PHE	64	12.987	-40.230	-8.239	1.00	18.70	A	C
ATOM	22	CB	PHE	64	13.904	-41.433	-8.066	1.00	18.53	A	C
ATOM	23	CG	PHE	64	14.182	-41.779	-6.631	1.00	19.76	A	C
ATOM	24	CD1	PHE	64	13.146	-42.007	-5.745	1.00	21.07	A	C
ATOM	25	CD2	PHE	64	15.479	-41.893	-6.169	1.00	20.97	A	C
ATOM	26	CE1	PHE	64	13.405	-42.353	-4.418	1.00	22.93	A	C
ATOM	27	CE2	PHE	64	15.745	-42.237	-4.847	1.00	20.76	A	C
ATOM	28	CZ	PHE	64	14.712	-42.465	-3.972	1.00	20.85	A	C
ATOM	29	C	PHE	64	13.562	-39.062	-7.457	1.00	19.32	A	C
ATOM	30	O	PHE	64	14.475	-38.380	-7.932	1.00	19.55	A	O
ATOM	31	N	HIS	65	13.028	-38.846	-6.256	1.00	19.72	A	N
ATOM	32	CA	HIS	65	13.445	-37.744	-5.395	1.00	18.95	A	C
ATOM	33	CB	HIS	65	12.394	-36.638	-5.417	1.00	17.76	A	C
ATOM	34	CG	HIS	65	12.152	-36.074	-6.780	1.00	19.55	A	C
ATOM	35	CD2	HIS	65	11.341	-36.481	-7.783	1.00	20.70	A	C
ATOM	36	ND1	HIS	65	12.813	-34.962	-7.253	1.00	20.16	A	N
ATOM	37	CE1	HIS	65	12.420	-34.709	-8.488	1.00	17.64	A	C
ATOM	38	NE2	HIS	65	11.527	-35.616	-8.834	1.00	17.20	A	N
ATOM	39	C	HIS	65	13.682	-38.196	-3.959	1.00	19.09	A	C
ATOM	40	O	HIS	65	13.039	-39.123	-3.468	1.00	18.05	A	O
ATOM	41	N	ARG	66	14.608	-37.519	-3.289	1.00	19.61	A	N
ATOM	42	CA	ARG	66	15.107	-37.964	-2.000	1.00	20.80	A	C
ATOM	43	CB	ARG	66	16.306	-38.873	-2.238	1.00	20.79	A	C
ATOM	44	CG	ARG	66	16.797	-39.615	-1.039	1.00	23.33	A	C
ATOM	45	CD	ARG	66	18.313	-39.719	-1.112	1.00	25.65	A	C
ATOM	46	NE	ARG	66	18.769	-41.038	-1.533	1.00	25.62	A	N
ATOM	47	CZ	ARG	66	20.031	-41.338	-1.827	1.00	24.86	A	C
ATOM	48	NH1	ARG	66	20.946	-42.573	-2.195	1.00	24.37	A	N
ATOM	49	NH2	ARG	66	20.974	-40.408	-1.760	1.00	23.60	A	N
ATOM	50	C	ARG	66	15.496	-36.737	-1.172	1.00	22.22	A	C
ATOM	51	O	ARG	66	15.836	-35.695	-1.727	1.00	21.06	A	O
ATOM	52	N	CYS	67	15.429	-36.852	0.152	1.00	24.84	A	N
ATOM	53	CA	CYS	67	15.637	-35.694	1.022	1.00	26.62	A	C
ATOM	54	CB	CYS	67	15.124	-35.966	2.437	1.00	26.30	A	C
ATOM	55	SG	CYS	67	15.480	-34.603	3.602	1.00	29.02	A	S
ATOM	56	C	CYS	67	17.105	-35.321	1.107	1.00	28.15	A	C
ATOM	57	O	CYS	67	17.951	-36.170	1.412	1.00	29.15	A	O
ATOM	58	N	ALA	68	17.404	-34.045	0.863	1.00	28.74	A	N
ATOM	59	CA	ALA	68	18.786	-33.593	0.787	1.00	28.43	A	C
ATOM	60	CB	ALA	68	18.856	-32.228	0.139	1.00	27.23	A	C
ATOM	61	C	ALA	68	19.438	-33.564	2.162	1.00	28.95	A	C
ATOM	62	O	ALA	68	20.660	-33.551	2.270	1.00	30.22	A	O
ATOM	63	N	LYS	69	18.623	-33.565	3.212	1.00	29.94	A	N
ATOM	64	CA	LYS	69	19.131	-33.726	4.570	1.00	31.16	A	C
ATOM	65	CB	LYS	69	18.197	-33.047	5.575	1.00	33.82	A	C
ATOM	66	CG	LYS	69	17.843	-31.621	5.209	1.00	36.96	A	C
ATOM	67	CD	LYS	69	18.947	-30.629	5.550	1.00	38.61	A	C
ATOM	68	CE	LYS	69	18.481	-29.195	5.237	1.00	41.46	A	C
ATOM	69	NZ	LYS	69	19.397	-28.113	5.730	1.00	40.68	A	N
ATOM	70	C	LYS	69	19.233	-35.212	4.887	1.00	29.96	A	C
ATOM	71	O	LYS	69	18.252	-35.845	5.273	1.00	30.13	A	O
ATOM	72	N	ASP	70	20.427	-35.764	4.731	1.00	28.75	A	N
ATOM	73	CA	ASP	70	20.580	-37.202	4.782	1.00	27.72	A	C
ATOM	74	CB	ASP	70	22.052	-37.571	4.623	1.00	30.38	A	C

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ATOM     75  CG  ASP   70      22.243 -38.864   3.847  1.00 33.51      A    C
ATOM     76  OD1 ASP   70      21.800 -38.932   2.675  1.00 34.66      A    O
ATOM     77  OD2 ASP   70      22.834 -39.813   4.406  1.00 33.99      A    O
ATOM     78  C   ASP   70      20.008 -37.850   6.045  1.00 25.79      A    C
ATOM     79  O   ASP   70      19.346 -38.878   5.970  1.00 25.05      A    O
ATOM     80  N   PRO   71      20.258 -37.265   7.225  1.00 24.83      A    N
ATOM     81  CD  PRO   71      21.116 -36.107   7.526  1.00 24.43      A    C
ATOM     82  CA  PRO   71      19.729 -37.890   8.445  1.00 24.69      A    C
ATOM     83  CB  PRO   71      20.456 -37.157   9.572  1.00 24.48      A    C
ATOM     84  CG  PRO   71      20.847 -35.854   8.980  1.00 24.31      A    C
ATOM     85  C   PRO   71      18.207 -37.806   8.587  1.00 24.67      A    C
ATOM     86  O   PRO   71      17.603 -38.557   9.353  1.00 25.53      A    O
ATOM     87  N   TRP   72      17.589 -36.892   7.848  1.00 22.88      A    N
ATOM     88  CA  TRP   72      16.145 -36.742   7.892  1.00 21.03      A    C
ATOM     89  CB  TRP   72      15.749 -35.337   7.462  1.00 21.00      A    C
ATOM     90  CG  TRP   72      16.168 -34.308   8.454  1.00 20.77      A    C
ATOM     91  CD2 TRP   72      15.958 -32.894   8.366  1.00 18.78      A    C
ATOM     92  CE2 TRP   72      16.470 -32.326   9.551  1.00 18.39      A    C
ATOM     93  CE3 TRP   72      15.390 -32.056   7.406  1.00 16.68      A    C
ATOM     94  CD1 TRP   72      16.785 -34.531   9.650  1.00 19.91      A    C
ATOM     95  NE1 TRP   72      16.967 -33.345  10.316  1.00 19.30      A    N
ATOM     96  CZ2 TRP   72      16.425 -30.956   9.796  1.00 15.74      A    C
ATOM     97  CZ3 TRP   72      15.348 -30.701   7.654  1.00 14.03      A    C
ATOM     98  CH2 TRP   72      15.861 -30.165   8.838  1.00 14.11      A    C
ATOM     99  C   TRP   72      15.486 -37.762   6.993  1.00 21.80      A    C
ATOM    100  O   TRP   72      14.281 -38.007   7.090  1.00 24.16      A    O
ATOM    101  N   ARG   73      16.288 -38.362   6.120  1.00 21.43      A    N
ATOM    102  CA  ARG   73      15.820 -39.451   5.266  1.00 18.71      A    C
ATOM    103  CB  ARG   73      16.942 -39.936   4.341  1.00 15.20      A    C
ATOM    104  CG  ARG   73      17.321 -38.950   3.272  1.00 13.51      A    C
ATOM    105  CD  ARG   73      18.419 -39.521   2.401  1.00 15.16      A    C
ATOM    106  NE  ARG   73      18.059 -40.830   1.856  1.00 15.36      A    N
ATOM    107  CZ  ARG   73      18.930 -41.794   1.576  1.00 13.61      A    C
ATOM    108  NH1 ARG   73      20.222 -41.601   1.793  1.00 10.92      A    N
ATOM    109  NH2 ARG   73      18.507 -42.950   1.079  1.00 11.59      A    N
ATOM    110  C   ARG   73      15.297 -40.629   6.081  1.00 17.55      A    C
ATOM    111  O   ARG   73      15.826 -40.952   7.147  1.00 16.42      A    O
ATOM    112  N   LEU   74      14.245 -41.248   5.556  1.00 17.09      A    N
ATOM    113  CA  LEU   74      13.646 -42.438   6.131  1.00 17.51      A    C
ATOM    114  CB  LEU   74      12.264 -42.115   6.721  1.00 16.82      A    C
ATOM    115  CG  LEU   74      12.207 -41.101   7.868  1.00 17.36      A    C
ATOM    116  CD1 LEU   74      10.767 -40.742   8.206  1.00 15.71      A    C
ATOM    117  CD2 LEU   74      12.914 -41.689   9.069  1.00 14.95      A    C
ATOM    118  C   LEU   74      13.493 -43.437   4.994  1.00 18.74      A    C
ATOM    119  O   LEU   74      12.401 -43.611   4.451  1.00 20.09      A    O
ATOM    120  N   PRO   75      14.594 -44.102   4.610  1.00 19.33      A    N
ATOM    121  CD  PRO   75      15.932 -43.950   5.198  1.00 17.44      A    C
ATOM    122  CA  PRO   75      14.608 -44.997   3.446  1.00 18.25      A    C
ATOM    123  CB  PRO   75      16.091 -45.274   3.226  1.00 16.29      A    C
ATOM    124  CG  PRO   75      16.714 -45.036   4.531  1.00 15.69      A    C
ATOM    125  C   PRO   75      13.802 -46.280   3.619  1.00 18.01      A    C
ATOM    126  O   PRO   75      13.625 -46.766   4.729  1.00 19.35      A    O
ATOM    127  N   GLY   76      13.319 -46.828   2.510  1.00 17.66      A    N
ATOM    128  CA  GLY   76      12.610 -48.092   2.568  1.00 16.82      A    C
ATOM    129  C   GLY   76      11.119 -47.864   2.556  1.00 16.73      A    C
ATOM    130  O   GLY   76      10.328 -48.792   2.431  1.00 16.37      A    O
ATOM    131  N   THR   77      10.732 -46.610   2.702  1.00 16.94      A    N
ATOM    132  CA  THR   77       9.352 -46.241   2.510  1.00 18.46      A    C
ATOM    133  CB  THR   77       8.753 -45.699   3.799  1.00 20.12      A    C
ATOM    134  OG1 THR   77       8.988 -46.643   4.850  1.00 22.67      A    O
ATOM    135  CG2 THR   77       7.250 -45.499   3.642  1.00 20.19      A    C
ATOM    136  C   THR   77       9.301 -45.189   1.427  1.00 17.73      A    C
ATOM    137  O   THR   77      10.106 -44.259   1.415  1.00 17.66      A    O
ATOM    138  N   TYR   78       8.362 -45.354   0.501  1.00 18.93      A    N
ATOM    139  CA  TYR   78       8.281 -44.474  -0.656  1.00 19.73      A    C
ATOM    140  CB  TYR   78       8.854 -45.173  -1.892  1.00 18.09      A    C
ATOM    141  CG  TYR   78      10.285 -45.575  -1.679  1.00 19.07      A    C
ATOM    142  CD1 TYR   78      10.601 -46.777  -1.050  1.00 19.20      A    C
ATOM    143  CE1 TYR   78      11.909 -47.114  -0.777  1.00 18.84      A    C
ATOM    144  CQ2 TYR   78      11.323 -44.727  -2.035  1.00 18.89      A    C
ATOM    145  CE2 TYR   78      12.633 -45.061  -1.771  1.00 18.34      A    C
ATOM    146  CZ  TYR   78      12.917 -46.252  -1.137  1.00 18.13      A    C
ATOM    147  OH  TYR   78      14.215 -46.565  -0.835  1.00 20.00      A    O
ATOM    148  C   TYR   78       6.861 -44.025  -0.917  1.00 20.34      A    C
ATOM    149  O   TYR   78       5.894 -44.735  -0.617  1.00 21.95      A    O
ATOM    150  N   VAL   79       6.750 -42.826  -1.472  1.00 20.31      A    N
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ATOM    151  CA  VAL    79       5.460 -42.246  -1.791  1.00 19.33      A    C
ATOM    152  CB  VAL    79       5.336 -40.845  -1.171  1.00 19.66      A    C
ATOM    153  CG1 VAL    79       3.957 -40.279  -1.436  1.00 20.37      A    C
ATOM    154  CG2 VAL    79       5.620 -40.916   0.319  1.00 17.38      A    C
ATOM    155  C   VAL    79       5.344 -42.140  -3.302  1.00 18.08      A    C
ATOM    156  O   VAL    79       5.816 -41.188  -3.900  1.00 18.83      A    O
ATOM    157  N   VAL    80       4.716 -43.129  -3.919  1.00 18.93      A    N
ATOM    158  CA  VAL    80       4.508 -43.112  -5.359  1.00 18.97      A    C
ATOM    159  CB  VAL    80       4.142 -44.508  -5.872  1.00 18.62      A    C
ATOM    160  CG1 VAL    80       4.083 -44.504  -7.391  1.00 16.15      A    C
ATOM    161  CG2 VAL    80       5.150 -45.513  -5.356  1.00 16.43      A    C
ATOM    162  C   VAL    80       3.386 -42.157  -5.727  1.00 19.40      A    C
ATOM    163  O   VAL    80       2.241 -42.362  -5.327  1.00 19.75      A    O
ATOM    164  N   VAL    81       3.733 -41.123  -6.492  1.00 20.81      A    N
ATOM    165  CA  VAL    81       2.808 -40.060  -6.889  1.00 22.10      A    C
ATOM    166  CB  VAL    81       3.451 -38.651  -6.730  1.00 22.25      A    C
ATOM    167  CG1 VAL    81       2.536 -37.580  -7.304  1.00 19.18      A    C
ATOM    168  CG2 VAL    81       3.718 -38.366  -5.264  1.00 22.64      A    C
ATOM    169  C   VAL    81       2.398 -40.229  -8.341  1.00 24.17      A    C
ATOM    170  O   VAL    81       3.212 -40.105  -9.259  1.00 24.19      A    O
ATOM    171  N   LEU    82       1.123 -40.509  -8.546  1.00 26.03      A    N
ATOM    172  CA  LEU    82       0.622 -40.717  -9.885  1.00 27.03      A    C
ATOM    173  CB  LEU    82      -0.593 -41.641  -9.828  1.00 27.37      A    C
ATOM    174  CG  LEU    82      -0.289 -42.868  -8.959  1.00 28.39      A    C
ATOM    175  CD1 LEU    82      -1.516 -43.767  -8.828  1.00 28.17      A    C
ATOM    176  CD2 LEU    82       0.889 -43.627  -9.574  1.00 27.31      A    C
ATOM    177  C   LEU    82       0.261 -39.362 -10.448  1.00 27.34      A    C
ATOM    178  O   LEU    82       0.098 -38.404  -9.701  1.00 24.96      A    O
ATOM    179  N   LYS    83       0.162 -39.279 -11.768  1.00 30.55      A    N
ATOM    180  CA  LYS    83      -0.317 -38.067 -12.414  1.00 33.51      A    C
ATOM    181  CB  LYS    83      -0.422 -38.286 -13.915  1.00 31.93      A    C
ATOM    182  CG  LYS    83       0.905 -38.539 -14.576  1.00 34.59      A    C
ATOM    183  CD  LYS    83       0.707 -39.240 -15.898  1.00 37.08      A    C
ATOM    184  CE  LYS    83       1.989 -39.282 -16.705  1.00 38.83      A    C
ATOM    185  NZ  LYS    83       1.749 -39.874 -18.052  1.00 40.48      A    N
ATOM    186  C   LYS    83      -1.682 -37.702 -11.852  1.00 36.22      A    C
ATOM    187  O   LYS    83      -2.568 -38.553 -11.750  1.00 36.20      A    O
ATOM    188  N   GLU    84      -1.855 -36.440 -11.475  1.00 38.84      A    N
ATOM    189  CA  GLU    84      -3.128 -36.023 -10.919  1.00 40.89      A    C
ATOM    190  CB  GLU    84      -3.069 -34.579 -10.428  1.00 42.22      A    C
ATOM    191  CG  GLU    84      -2.933 -33.546 -11.512  1.00 46.22      A    C
ATOM    192  CD  GLU    84      -3.066 -32.138 -10.964  1.00 49.28      A    C
ATOM    193  OE1 GLU    84      -3.370 -31.995  -9.759  1.00 50.03      A    O
ATOM    194  OE2 GLU    84      -2.872 -31.173 -11.734  1.00 51.27      A    O
ATOM    195  C   GLU    84      -4.181 -36.171 -11.996  1.00 41.56      A    C
ATOM    196  O   GLU    84      -3.880 -36.094 -13.190  1.00 40.94      A    O
ATOM    197  N   GLU    85      -5.413 -36.399 -11.558  1.00 42.71      A    N
ATOM    198  CA  GLU    85      -6.483 -36.829 -12.443  1.00 43.97      A    C
ATOM    199  CB  GLU    85      -6.446 -36.052 -13.759  1.00 47.49      A    C
ATOM    200  CG  GLU    85      -6.820 -34.579 -13.615  1.00 53.47      A    C
ATOM    201  CD  GLU    85      -6.879 -33.857 -14.955  1.00 57.25      A    C
ATOM    202  OE1 GLU    85      -7.821 -33.053 -15.160  1.00 58.40      A    O
ATOM    203  OE2 GLU    85      -5.985 -34.096 -15.804  1.00 58.66      A    O
ATOM    204  C   GLU    85      -6.407 -38.323 -12.724  1.00 41.83      A    C
ATOM    205  O   GLU    85      -7.152 -38.839 -13.549  1.00 42.95      A    O
ATOM    206  N   THR    86      -5.503 -39.019 -12.042  1.00 39.72      A    N
ATOM    207  CA  THR    86      -5.578 -40.471 -11.973  1.00 36.25      A    C
ATOM    208  CB  THR    86      -4.213 -41.106 -11.608  1.00 34.62      A    C
ATOM    209  OG1 THR    86      -3.301 -40.957 -12.701  1.00 30.95      A    O
ATOM    210  CG2 THR    86      -4.379 -42.585 -11.295  1.00 33.47      A    C
ATOM    211  C   THR    86      -6.599 -40.825 -10.902  1.00 35.99      A    C
ATOM    212  O   THR    86      -6.593 -40.255  -9.817  1.00 36.01      A    O
ATOM    213  N   HIS    87      -7.482 -41.761 -11.213  1.00 35.71      A    N
ATOM    214  CA  HIS    87      -8.564 -42.089 -10.312  1.00 36.27      A    C
ATOM    215  CB  HIS    87      -9.823 -42.388 -11.112  1.00 38.72      A    C
ATOM    216  CG  HIS    87     -10.446 -41.172 -11.715  1.00 41.65      A    C
ATOM    217  CD2 HIS    87     -11.072 -40.117 -11.143  1.00 41.88      A    C
ATOM    218  ND1 HIS    87     -10.461 -40.937 -13.074  1.00 43.71      A    N
ATOM    219  CE1 HIS    87     -11.070 -39.790 -13.314  1.00 43.53      A    C
ATOM    220  NE2 HIS    87     -11.451 -39.273 -12.159  1.00 43.79      A    N
ATOM    221  C   HIS    87      -8.239 -43.251  -9.393  1.00 35.80      A    C
ATOM    222  O   HIS    87      -7.406 -44.097  -9.705  1.00 36.51      A    O
ATOM    223  N   LEU    88      -8.925 -43.288  -8.258  1.00 34.46      A    N
ATOM    224  CA  LEU    88      -8.567 -44.177  -7.168  1.00 32.12      A    C
ATOM    225  CB  LEU    88      -9.537 -43.994  -6.004  1.00 28.85      A    C
ATOM    226  CG  LEU    88      -9.207 -44.893  -4.817  1.00 26.89      A    C
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ATOM    227  CD1 LEU    88      -7.810 -44.563  -4.298  1.00 25.11      A    C
ATOM    228  CD2 LEU    88     -10.249 -44.724  -3.744  1.00 24.99      A    C
ATOM    229  C   LEU    88      -8.511 -45.652  -7.549  1.00 31.90      A    C
ATOM    230  O   LEU    88      -7.850 -46.438  -6.866  1.00 32.53      A    O
ATOM    231  N   SER    89      -9.184 -46.038  -8.628  1.00 31.60      A    N
ATOM    232  CA  SER    89      -9.173 -47.440  -9.012  1.00 32.76      A    C
ATOM    233  CB  SER    89     -10.412 -47.801  -9.813  1.00 32.88      A    C
ATOM    234  OG  SER    89     -10.215 -47.494 -11.183  1.00 37.17      A    O
ATOM    235  C   SER    89      -7.938 -47.674  -9.875  1.00 32.28      A    C
ATOM    236  O   SER    89      -7.371 -48.771  -9.852  1.00 32.59      A    O
ATOM    237  N   GLN    90      -7.533 -46.652 -10.641  1.00 33.53      A    N
ATOM    238  CA  GLN    90      -6.260 -46.691 -11.384  1.00 35.45      A    C
ATOM    239  CB  GLN    90      -6.130 -45.500 -12.337  1.00 38.30      A    C
ATOM    240  CG  GLN    90      -7.054 -45.578 -13.538  1.00 41.60      A    C
ATOM    241  CD  GLN    90      -7.386 -44.199 -14.107  1.00 44.93      A    C
ATOM    242  OE1 GLN    90      -6.964 -43.179 -13.553  1.00 46.74      A    O
ATOM    243  NE2 GLN    90      -8.148 -44.162 -15.214  1.00 46.21      A    N
ATOM    244  C   GLN    90      -5.081 -46.714 -10.399  1.00 33.56      A    C
ATOM    245  O   GLN    90      -4.123 -47.444 -10.636  1.00 33.85      A    O
ATOM    246  N   SER    91      -5.176 -45.966  -9.289  1.00 32.27      A    N
ATOM    247  CA  SER    91      -4.234 -46.079  -8.168  1.00 31.90      A    C
ATOM    248  CB  SER    91      -4.673 -45.187  -7.010  1.00 32.61      A    C
ATOM    249  OG  SER    91      -4.020 -43.938  -7.044  1.00 35.88      A    O
ATOM    250  C   SER    91      -4.116 -47.509  -7.667  1.00 30.46      A    C
ATOM    251  O   SER    91      -3.053 -48.115  -7.740  1.00 29.82      A    O
ATOM    252  N   GLU    92      -5.219 -48.040  -7.159  1.00 30.78      A    N
ATOM    253  CA  GLU    92      -5.238 -49.385  -6.616  1.00 29.62      A    C
ATOM    254  CB  GLU    92      -6.652 -49.765  -6.200  1.00 30.03      A    C
ATOM    255  CG  GLU    92      -7.297 -48.854  -5.173  1.00 31.12      A    C
ATOM    256  CD  GLU    92      -8.720 -49.289  -4.847  1.00 33.15      A    C
ATOM    257  OE1 GLU    92      -9.229 -50.204  -5.540  1.00 33.55      A    O
ATOM    258  OE2 GLU    92      -9.326 -48.720  -3.908  1.00 32.58      A    O
ATOM    259  C   GLU    92      -4.738 -50.399  -7.632  1.00 28.84      A    C
ATOM    260  O   GLU    92      -4.051 -51.353  -7.275  1.00 27.60      A    O
ATOM    261  N   ARG    93      -5.087 -50.202  -8.899  1.00 28.96      A    N
ATOM    262  CA  ARG    93      -4.744 -51.186  -9.911  1.00 29.05      A    C
ATOM    263  CB  ARG    93      -5.381 -50.820 -11.259  1.00 31.00      A    C
ATOM    264  CG  ARG    93      -5.574 -52.005 -12.218  1.00 34.57      A    C
ATOM    265  CD  ARG    93      -6.296 -51.599 -13.519  1.00 39.82      A    C
ATOM    266  NE  ARG    93      -7.724 -51.277 -13.341  1.00 44.62      A    N
ATOM    267  CZ  ARG    93      -8.288 -50.103 -13.651  1.00 45.59      A    C
ATOM    268  NH1 ARG    93      -9.594 -49.903 -13.456  1.00 44.61      A    N
ATOM    269  NH2 ARG    93      -7.548 -49.120 -14.155  1.00 45.28      A    N
ATOM    270  C   ARG    93      -3.229 -51.201 -10.020  1.00 27.72      A    C
ATOM    271  O   ARG    93      -2.613 -52.261 -10.149  1.00 26.82      A    O
ATOM    272  N   THR    94      -2.640 -50.011  -9.932  1.00 27.00      A    N
ATOM    273  CA  THR    94      -1.197 -49.837 -10.068  1.00 26.48      A    C
ATOM    274  CB  THR    94      -0.822 -48.354 -10.250  1.00 25.41      A    C
ATOM    275  OG1 THR    94      -1.140 -47.942 -11.585  1.00 24.63      A    O
ATOM    276  CG2 THR    94       0.657 -48.145  -9.995  1.00 25.11      A    C
ATOM    277  C   THR    94      -0.407 -50.388  -8.894  1.00 26.50      A    C
ATOM    278  O   THR    94       0.637 -51.007  -9.081  1.00 26.28      A    O
ATOM    279  N   ALA    95      -0.898 -50.155  -7.684  1.00 26.80      A    N
ATOM    280  CA  ALA    95      -0.282 -50.736  -6.508  1.00 26.79      A    C
ATOM    281  CB  ALA    95      -1.066 -50.347  -5.265  1.00 25.54      A    C
ATOM    282  C   ALA    95      -0.241 -52.257  -6.662  1.00 27.87      A    C
ATOM    283  O   ALA    95       0.753 -52.893  -6.322  1.00 28.83      A    O
ATOM    284  N   ARG    96      -1.318 -52.841  -7.180  1.00 28.88      A    N
ATOM    285  CA  ARG    96      -1.386 -54.297  -7.327  1.00 28.84      A    C
ATOM    286  CB  ARG    96      -2.785 -54.731  -7.793  1.00 29.30      A    C
ATOM    287  CG  ARG    96      -3.808 -54.845  -6.680  1.00 29.85      A    C
ATOM    288  CD  ARG    96      -5.163 -55.261  -7.206  1.00 31.96      A    C
ATOM    289  NE  ARG    96      -6.184 -54.262  -6.887  1.00 36.30      A    N
ATOM    290  CZ  ARG    96      -6.973 -53.679  -7.789  1.00 37.39      A    C
ATOM    291  NH1 ARG    96      -7.874 -52.774  -7.410  1.00 37.42      A    N
ATOM    292  NH2 ARG    96      -6.865 -54.005  -9.072  1.00 37.61      A    N
ATOM    293  C   ARG    96      -0.336 -54.783  -8.320  1.00 28.39      A    C
ATOM    294  O   ARG    96       0.303 -55.814  -8.113  1.00 28.76      A    O
ATOM    295  N   ARG    97      -0.163 -54.019  -9.391  1.00 27.77      A    N
ATOM    296  CA  ARG    97       0.755 -54.372 -10.459  1.00 28.70      A    C
ATOM    297  CB  ARG    97       0.576 -53.397 -11.628  1.00 30.62      A    C
ATOM    298  CG  ARG    97       1.419 -53.715 -12.834  1.00 34.83      A    C
ATOM    299  CD  ARG    97       1.054 -52.827 -14.005  1.00 39.80      A    C
ATOM    300  NE  ARG    97       0.580 -53.596 -15.155  1.00 43.80      A    N
ATOM    301  CZ  ARG    97      -0.693 -53.661 -15.537  1.00 45.70      A    C
ATOM    302  NH1 ARG    97      -1.037 -54.385 -16.595  1.00 47.10      A    N
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ATOM    303  NH2 ARG   97      -1.627 -53.000 -14.865  1.00 45.89      A    N
ATOM    304  C   ARG   97       2.202 -54.346  -9.964  1.00 27.88      A    C
ATOM    305  O   ARG   97       2.983 -55.253 -10.248  1.00 28.38      A    O
ATOM    306  N   LEU   98       2.545 -53.296  -9.223  1.00 27.13      A    N
ATOM    307  CA  LEU   98       3.852 -53.161  -8.585  1.00 23.63      A    C
ATOM    308  CB  LEU   98       3.863 -51.911  -7.705  1.00 23.28      A    C
ATOM    309  CG  LEU   98       4.960 -51.732  -6.656  1.00 24.24      A    C
ATOM    310  CD1 LEU   98       6.330 -51.842  -7.301  1.00 25.92      A    C
ATOM    311  CD2 LEU   98       4.798 -50.374  -5.996  1.00 23.55      A    C
ATOM    312  C   LEU   98       4.115 -54.376  -7.732  1.00 22.69      A    C
ATOM    313  O   LEU   98       5.145 -55.031  -7.849  1.00 23.28      A    O
ATOM    314  N   GLN   99       3.147 -54.664  -6.874  1.00 22.76      A    N
ATOM    315  CA  GLN   99       3.208 -55.766  -5.931  1.00 21.72      A    C
ATOM    316  CB  GLN   99       1.848 -55.904  -5.261  1.00 23.87      A    C
ATOM    317  CG  GLN   99       1.895 -56.405  -3.838  1.00 28.10      A    C
ATOM    318  CD  GLN   99       1.583 -55.309  -2.857  1.00 28.27      A    C
ATOM    319  OE1 GLN   99       1.373 -55.555  -1.669  1.00 29.74      A    O
ATOM    320  NE2 GLN   99       1.551 -54.078  -3.351  1.00 29.27      A    N
ATOM    321  C   GLN   99       3.569 -57.066  -6.644  1.00 20.42      A    C
ATOM    322  O   GLN   99       4.434 -57.822  -6.196  1.00 18.61      A    O
ATOM    323  N   ALA  100       2.886 -57.306  -7.759  1.00 19.89      A    N
ATOM    324  CA  ALA  100       3.071 -58.504  -8.567  1.00 19.33      A    C
ATOM    325  CB  ALA  100       2.023 -58.546  -9.668  1.00 17.45      A    C
ATOM    326  C   ALA  100       4.464 -58.551  -9.180  1.00 18.79      A    C
ATOM    327  O   ALA  100       5.216 -59.499  -8.959  1.00 21.98      A    O
ATOM    328  N   GLN  101       4.803 -57.524  -9.950  1.00 16.83      A    N
ATOM    329  CA  GLN  101       6.090 -57.452 -10.617  1.00 14.81      A    C
ATOM    330  CB  GLN  101       6.204 -56.118 -11.339  1.00 14.22      A    C
ATOM    331  CG  GLN  101       5.063 -55.912 -12.293  1.00 17.42      A    C
ATOM    332  CD  GLN  101       5.207 -54.679 -13.146  1.00 20.01      A    C
ATOM    333  OE1 GLN  101       4.281 -54.306 -13.870  1.00 24.17      A    O
ATOM    334  NE2 GLN  101       6.364 -54.033 -13.070  1.00 20.43      A    N
ATOM    335  C   GLN  101       7.217 -57.602  -9.607  1.00 15.13      A    C
ATOM    336  O   GLN  101       8.253 -58.205  -9.893  1.00 14.83      A    O
ATOM    337  N   ALA  102       6.997 -57.054  -8.417  1.00 14.58      A    N
ATOM    338  CA  ALA  102       7.975 -57.129  -7.348  1.00 13.30      A    C
ATOM    339  CB  ALA  102       7.632 -56.117  -6.279  1.00 11.40      A    C
ATOM    340  C   ALA  102       8.047 -58.536  -6.749  1.00 13.44      A    C
ATOM    341  O   ALA  102       9.107 -58.993  -6.339  1.00 12.59      A    O
ATOM    342  N   ALA  103       6.918 -59.226  -6.698  1.00 15.23      A    N
ATOM    343  CA  ALA  103       6.904 -60.566  -6.142  1.00 17.52      A    C
ATOM    344  CB  ALA  103       5.488 -61.014  -5.894  1.00 13.93      A    C
ATOM    345  C   ALA  103       7.587 -61.500  -7.124  1.00 20.68      A    C
ATOM    346  O   ALA  103       8.338 -62.395  -6.731  1.00 20.64      A    O
ATOM    347  N   ARG  104       7.329 -61.291  -8.407  1.00 23.25      A    N
ATOM    348  CA  ARG  104       7.976 -62.104  -9.411  1.00 26.07      A    C
ATOM    349  CB  ARG  104       7.556 -61.665 -10.806  1.00 28.33      A    C
ATOM    350  CG  ARG  104       6.248 -62.282 -11.215  1.00 33.77      A    C
ATOM    351  CD  ARG  104       5.898 -61.942 -12.629  1.00 39.49      A    C
ATOM    352  NE  ARG  104       4.707 -61.099 -12.687  1.00 46.45      A    N
ATOM    353  CZ  ARG  104       4.716 -59.802 -12.988  1.00 48.94      A    C
ATOM    354  NH1 ARG  104       3.572 -59.128 -13.016  1.00 50.90      A    N
ATOM    355  NH2 ARG  104       5.861 -59.179 -13.265  1.00 49.24      A    N
ATOM    356  C   ARG  104       9.481 -62.001  -9.257  1.00 27.46      A    C
ATOM    357  O   ARG  104      10.200 -62.974  -9.467  1.00 29.34      A    O
ATOM    358  N   ARG  105       9.963 -60.828  -8.874  1.00 28.17      A    N
ATOM    359  CA  ARG  105      11.384 -60.669  -8.636  1.00 27.57      A    C
ATOM    360  CB  ARG  105      11.786 -59.203  -8.778  1.00 29.99      A    C
ATOM    361  CG  ARG  105      11.942 -58.762 -10.220  1.00 34.08      A    C
ATOM    362  CD  ARG  105      12.558 -57.373 -10.326  1.00 38.94      A    C
ATOM    363  NE  ARG  105      14.002 -57.376 -10.107  1.00 43.02      A    N
ATOM    364  CZ  ARG  105      14.688 -56.338  -9.635  1.00 46.10      A    C
ATOM    365  NH1 ARG  105      16.004 -56.429  -9.470  1.00 48.21      A    N
ATOM    366  NH2 ARG  105      14.057 -55.210  -9.319  1.00 46.61      A    N
ATOM    367  C   ARG  105      11.775 -61.195  -7.260  1.00 26.22      A    C
ATOM    368  O   ARG  105      12.947 -61.228  -6.917  1.00 25.88      A    O
ATOM    369  N   GLY  106      10.793 -61.609  -6.473  1.00 26.12      A    N
ATOM    370  CA  GLY  106      11.090 -62.194  -5.174  1.00 25.74      A    C
ATOM    371  C   GLY  106      11.029 -61.231  -4.000  1.00 25.62      A    C
ATOM    372  O   GLY  106      11.455 -61.566  -2.892  1.00 25.64      A    O
ATOM    373  N   TYR  107      10.497 -60.035  -4.237  1.00 25.48      A    N
ATOM    374  CA  TYR  107      10.483 -58.990  -3.222  1.00 25.01      A    C
ATOM    375  CB  TYR  107      10.780 -57.627  -3.846  1.00 24.93      A    C
ATOM    376  CG  TYR  107      12.237 -57.386  -4.137  1.00 26.36      A    C
ATOM    377  CD1 TYR  107      13.070 -56.810  -3.183  1.00 25.53      A    C
ATOM    378  CE1 TYR  107      14.406 -56.570  -3.448  1.00 26.30      A    C
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ATOM    379  CD2 TYR   107      12.781 -57.724  -5.369  1.00 26.57      A    C
ATOM    380  CE2 TYR   107      14.117 -57.487  -5.647  1.00 27.76      A    C
ATOM    381  CZ  TYR   107      14.927 -56.909  -4.684  1.00 28.67      A    C
ATOM    382  OH  TYR   107      16.248 -56.640  -4.975  1.00 29.04      A    O
ATOM    383  C   TYR   107       9.173 -58.906  -2.471  1.00 24.19      A    C
ATOM    384  O   TYR   107       8.110 -58.711  -3.054  1.00 25.14      A    O
ATOM    385  N   LEU   108       9.266 -59.045  -1.159  1.00 24.06      A    N
ATOM    386  CA  LEU   108       8.158 -58.736  -0.275  1.00 23.48      A    C
ATOM    387  CB  LEU   108       8.537 -59.153   1.142  1.00 22.61      A    C
ATOM    388  CG  LEU   108       7.498 -58.975   2.234  1.00 24.41      A    C
ATOM    389  CD1 LEU   108       6.261 -59.780   1.881  1.00 27.40      A    C
ATOM    390  CD2 LEU   108       8.078 -59.433   3.562  1.00 25.45      A    C
ATOM    391  C   LEU   108       7.891 -57.223  -0.343  1.00 22.40      A    C
ATOM    392  O   LEU   108       8.830 -56.431  -0.309  1.00 23.50      A    O
ATOM    393  N   THR   109       6.625 -56.822  -0.464  1.00 21.22      A    N
ATOM    394  CA  THR   109       6.268 -55.405  -0.403  1.00 19.07      A    C
ATOM    395  CB  THR   109       6.087 -54.760  -1.794  1.00 19.59      A    C
ATOM    396  OG1 THR   109       4.895 -55.264  -2.398  1.00 19.90      A    O
ATOM    397  CG2 THR   109       7.274 -55.056  -2.694  1.00 20.38      A    C
ATOM    398  C   THR   109       4.956 -55.235   0.309  1.00 17.86      A    C
ATOM    399  O   THR   109       4.152 -56.150   0.351  1.00 17.14      A    O
ATOM    400  N   LYS   110       4.741 -54.044   0.849  1.00 18.44      A    N
ATOM    401  CA  LYS   110       3.512 -53.738   1.552  1.00 19.57      A    C
ATOM    402  CB  LYS   110       3.736 -53.841   3.054  1.00 20.21      A    C
ATOM    403  CG  LYS   110       2.505 -53.554   3.861  1.00 22.66      A    C
ATOM    404  CD  LYS   110       2.449 -54.457   5.061  1.00 24.45      A    C
ATOM    405  CE  LYS   110       1.013 -54.835   5.367  1.00 26.17      A    C
ATOM    406  NZ  LYS   110       0.944 -56.041   6.235  1.00 28.22      A    N
ATOM    407  C   LYS   110       3.001 -52.347   1.219  1.00 19.90      A    C
ATOM    408  O   LYS   110       3.728 -51.359   1.344  1.00 20.91      A    O
ATOM    409  N   ILE   111       1.741 -52.270   0.809  1.00 19.04      A    N
ATOM    410  CA  ILE   111       1.085 -50.982   0.623  1.00 18.90      A    C
ATOM    411  CB  ILE   111      -0.029 -51.093  -0.406  1.00 17.72      A    C
ATOM    412  CG2 ILE   111      -0.751 -49.766  -0.525  1.00 18.36      A    C
ATOM    413  CG1 ILE   111       0.559 -51.524  -1.750  1.00 19.16      A    C
ATOM    414  CD1 ILE   111       1.519 -50.522  -2.347  1.00 17.38      A    C
ATOM    415  C   ILE   111       0.498 -50.431   1.927  1.00 18.80      A    C
ATOM    416  O   ILE   111      -0.408 -51.018   2.512  1.00 17.44      A    O
ATOM    417  N   LEU   112       1.016 -49.291   2.373  1.00 19.52      A    N
ATOM    418  CA  LEU   112       0.667 -48.762   3.687  1.00 19.22      A    C
ATOM    419  CB  LEU   112       1.845 -47.976   4.273  1.00 19.32      A    C
ATOM    420  CG  LEU   112       3.124 -48.811   4.439  1.00 22.22      A    C
ATOM    421  CD1 LEU   112       4.303 -47.914   4.746  1.00 22.17      A    C
ATOM    422  CD2 LEU   112       2.935 -49.838   5.534  1.00 20.39      A    C
ATOM    423  C   LEU   112      -0.565 -47.883   3.622  1.00 18.62      A    C
ATOM    424  O   LEU   112      -1.298 -47.762   4.599  1.00 18.34      A    O
ATOM    425  N   HIS   113      -0.800 -47.284   2.460  1.00 18.84      A    N
ATOM    426  CA  HIS   113      -1.900 -46.344   2.299  1.00 19.02      A    C
ATOM    427  CB  HIS   113      -1.589 -45.050   3.057  1.00 20.19      A    C
ATOM    428  CG  HIS   113      -2.658 -44.007   2.949  1.00 22.51      A    C
ATOM    429  CD2 HIS   113      -2.829 -42.992   2.066  1.00 22.37      A    C
ATOM    430  ND1 HIS   113      -3.716 -43.931   3.830  1.00 21.62      A    N
ATOM    431  CE1 HIS   113      -4.492 -42.916   3.494  1.00 21.55      A    C
ATOM    432  NE2 HIS   113      -3.977 -42.330   2.428  1.00 21.95      A    N
ATOM    433  C   HIS   113      -2.114 -46.036   0.829  1.00 18.98      A    C
ATOM    434  O   HIS   113      -1.163 -45.968   0.059  1.00 19.75      A    O
ATOM    435  N   VAL   114      -3.366 -45.854   0.434  1.00 19.49      A    N
ATOM    436  CA  VAL   114      -3.662 -45.387  -0.910  1.00 20.33      A    C
ATOM    437  CB  VAL   114      -4.528 -46.418  -1.678  1.00 19.03      A    C
ATOM    438  CG1 VAL   114      -4.729 -45.983  -3.119  1.00 16.98      A    C
ATOM    439  CG2 VAL   114      -3.865 -47.777  -1.635  1.00 18.60      A    C
ATOM    440  C   VAL   114      -4.414 -44.071  -0.793  1.00 21.30      A    C
ATOM    441  O   VAL   114      -5.416 -43.998  -0.098  1.00 22.35      A    O
ATOM    442  N   PHE   115      -3.932 -43.035  -1.468  1.00 22.44      A    N
ATOM    443  CA  PHE   115      -4.532 -41.713  -1.351  1.00 25.41      A    C
ATOM    444  CB  PHE   115      -3.513 -40.640  -1.757  1.00 26.99      A    C
ATOM    445  CG  PHE   115      -2.287 -40.613  -0.890  1.00 27.32      A    C
ATOM    446  CD1 PHE   115      -1.182 -41.383  -1.209  1.00 26.01      A    C
ATOM    447  CD2 PHE   115      -2.247 -39.836   0.259  1.00 26.96      A    C
ATOM    448  CE1 PHE   115      -0.068 -41.383  -0.403  1.00 24.13      A    C
ATOM    449  CE2 PHE   115      -1.130 -39.833   1.070  1.00 25.59      A    C
ATOM    450  CZ  PHE   115      -0.043 -40.607   0.736  1.00 24.86      A    C
ATOM    451  C   PHE   115      -5.813 -41.538  -2.171  1.00 26.85      A    C
ATOM    452  O   PHE   115      -5.856 -41.860  -3.371  1.00 27.95      A    O
ATOM    453  N   HIS   116      -6.844 -41.019  -1.503  1.00 26.78      A    N
ATOM    454  CA  HIS   116      -8.118 -40.659  -2.124  1.00 27.32      A    C
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ATOM    455  CB  HIS   116      -9.299 -40.991  -1.192  1.00 24.94      A    C
ATOM    456  CG  HIS   116      -9.464 -42.450  -0.886  1.00 24.70      A    C
ATOM    457  CD2 HIS   116     -10.571 -43.177  -0.597  1.00 24.63      A    C
ATOM    458  ND1 HIS   116      -8.406 -43.330  -0.821  1.00 25.06      A    N
ATOM    459  CE1 HIS   116      -8.852 -44.534  -0.506  1.00 23.26      A    C
ATOM    460  NE2 HIS   116     -10.162 -44.468  -0.365  1.00 22.78      A    N
ATOM    461  C   HIS   116      -8.144 -39.153  -2.418  1.00 28.43      A    C
ATOM    462  O   HIS   116      -9.172 -38.612  -2.819  1.00 29.04      A    O
ATOM    463  N   GLY   117      -7.022 -38.474  -2.215  1.00 29.51      A    N
ATOM    464  CA  GLY   117      -7.061 -37.021  -2.127  1.00 32.31      A    C
ATOM    465  C   GLY   117      -7.039 -36.199  -3.416  1.00 32.25      A    C
ATOM    466  O   GLY   117      -7.454 -36.658  -4.469  1.00 32.39      A    O
ATOM    467  N   LEU   118      -6.568 -34.957  -3.286  1.00 30.44      A    N
ATOM    468  CA  LEU   118      -6.261 -34.099  -4.427  1.00 29.49      A    C
ATOM    469  CB  LEU   118      -6.188 -32.630  -3.991  1.00 27.76      A    C
ATOM    470  CG  LEU   118      -7.442 -31.991  -3.399  1.00 26.47      A    C
ATOM    471  CD1 LEU   118      -7.061 -30.748  -2.612  1.00 26.36      A    C
ATOM    472  CD2 LEU   118      -8.421 -31.662  -4.511  1.00 25.77      A    C
ATOM    473  C   LEU   118      -4.906 -34.520  -4.966  1.00 29.07      A    C
ATOM    474  O   LEU   118      -4.345 -33.876  -5.848  1.00 29.41      A    O
ATOM    475  N   LEU   119      -4.382 -35.602  -4.407  1.00 29.30      A    N
ATOM    476  CA  LEU   119      -3.057 -36.084  -4.747  1.00 29.49      A    C
ATOM    477  CB  LEU   119      -2.093 -35.724  -3.624  1.00 28.72      A    C
ATOM    478  CG  LEU   119      -0.605 -35.959  -3.870  1.00 30.16      A    C
ATOM    479  CD1 LEU   119       0.070 -34.639  -4.268  1.00 29.01      A    C
ATOM    480  CD2 LEU   119       0.025 -36.538  -2.602  1.00 29.14      A    C
ATOM    481  C   LEU   119      -3.090 -37.600  -4.935  1.00 29.79      A    C
ATOM    482  O   LEU   119      -2.874 -38.358  -3.988  1.00 30.11      A    O
ATOM    483  N   PRO   120      -3.369 -38.061  -6.164  1.00 29.03      A    N
ATOM    484  CD  PRO   120      -3.629 -37.259  -7.367  1.00 28.57      A    C
ATOM    485  CA  PRO   120      -3.499 -39.498  -6.422  1.00 28.72      A    C
ATOM    486  CB  PRO   120      -4.078 -39.562  -7.831  1.00 27.70      A    C
ATOM    487  CG  PRO   120      -3.675 -38.291  -8.455  1.00 28.64      A    C
ATOM    488  C   PRO   120      -2.178 -40.242  -6.312  1.00 27.90      A    C
ATOM    489  O   PRO   120      -1.167 -39.838  -6.890  1.00 28.70      A    O
ATOM    490  N   GLY   121      -2.196 -41.334  -5.561  1.00 27.12      A    N
ATOM    491  CA  GLY   121      -0.998 -42.130  -5.392  1.00 25.12      A    C
ATOM    492  C   GLY   121      -1.206 -43.167  -4.315  1.00 24.57      A    C
ATOM    493  O   GLY   121      -2.343 -43.528  -4.009  1.00 25.06      A    O
ATOM    494  N   PHE   122      -0.108 -43.652  -3.746  1.00 23.56      A    N
ATOM    495  CA  PHE   122      -0.166 -44.565  -2.611  1.00 22.66      A    C
ATOM    496  CB  PHE   122      -0.560 -45.971  -3.064  1.00 23.01      A    C
ATOM    497  CG  PHE   122       0.355 -46.552  -4.109  1.00 25.01      A    C
ATOM    498  CD1 PHE   122       1.428 -47.349  -3.747  1.00 24.64      A    C
ATOM    499  CD2 PHE   122       0.128 -46.320  -5.457  1.00 25.34      A    C
ATOM    500  CE1 PHE   122       2.252 -47.903  -4.707  1.00 23.88      A    C
ATOM    501  CE2 PHE   122       0.952 -46.874  -6.421  1.00 25.50      A    C
ATOM    502  CZ  PHE   122       2.015 -47.667  -6.043  1.00 24.61      A    C
ATOM    503  C   PHE   122       1.196 -44.603  -1.945  1.00 22.06      A    C
ATOM    504  O   PHE   122       2.181 -44.162  -2.520  1.00 22.52      A    O
ATOM    505  N   LEU   123       1.237 -45.115  -0.722  1.00 22.50      A    N
ATOM    506  CA  LEU   123       2.466 -45.171   0.064  1.00 23.34      A    C
ATOM    507  CB  LEU   123       2.257 -44.478   1.424  1.00 22.27      A    C
ATOM    508  CG  LEU   123       3.295 -44.748   2.522  1.00 20.76      A    C
ATOM    509  CD1 LEU   123       4.386 -43.683   2.536  1.00 18.72      A    C
ATOM    510  CD2 LEU   123       2.585 -44.776   3.841  1.00 19.62      A    C
ATOM    511  C   LEU   123       2.882 -46.627   0.279  1.00 23.45      A    C
ATOM    512  O   LEU   123       2.122 -47.432   0.828  1.00 23.19      A    O
ATOM    513  N   VAL   124       4.094 -46.958  -0.152  1.00 23.78      A    N
ATOM    514  CA  VAL   124       4.541 -48.342  -0.136  1.00 23.86      A    C
ATOM    515  CB  VAL   124       4.718 -48.873  -1.580  1.00 21.15      A    C
ATOM    516  CG1 VAL   124       5.646 -47.970  -2.357  1.00 19.01      A    C
ATOM    517  CG2 VAL   124       5.263 -50.288  -1.550  1.00 21.68      A    C
ATOM    518  C   VAL   124       5.838 -48.557   0.657  1.00 24.98      A    C
ATOM    519  O   VAL   124       6.769 -47.751   0.596  1.00 23.20      A    O
ATOM    520  N   LYS   125       5.860 -49.648   1.418  1.00 26.11      A    N
ATOM    521  CA  LYS   125       7.057 -50.127   2.098  1.00 27.32      A    C
ATOM    522  CB  LYS   125       6.696 -50.623   3.498  1.00 25.97      A    C
ATOM    523  CG  LYS   125       7.812 -51.373   4.183  1.00 23.58      A    C
ATOM    524  CD  LYS   125       7.383 -51.861   5.549  1.00 25.85      A    C
ATOM    525  CE  LYS   125       8.566 -52.431   6.316  1.00 28.11      A    C
ATOM    526  NZ  LYS   125       8.220 -52.783   7.731  1.00 32.00      A    N
ATOM    527  C   LYS   125       7.673 -51.275   1.289  1.00 29.44      A    C
ATOM    528  O   LYS   125       7.025 -52.302   1.057  1.00 30.59      A    O
ATOM    529  N   MET   126       8.926 -51.097   0.873  1.00 29.55      A    N
ATOM    530  CA  MET   126       9.587 -52.004  -0.066  1.00 27.34      A    C
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ATOM    531  CB  MET   126       8.929 -51.896  -1.435  1.00 25.09      A    C
ATOM    532  CG  MET   126       9.170 -50.536  -2.052  1.00 24.16      A    C
ATOM    533  SD  MET   126       8.824 -50.453  -3.795  1.00 24.67      A    S
ATOM    534  CE  MET   126       9.694 -48.998  -4.214  1.00 24.23      A    C
ATOM    535  C   MET   126      11.054 -51.508  -0.208  1.00 27.55      A    C
ATOM    536  O   MET   126      11.407 -50.438   0.069  1.00 28.53      A    O
ATOM    537  N   SER   127      11.900 -52.514  -0.658  1.00 26.25      A    N
ATOM    538  CA  SER   127      13.272 -52.178  -1.035  1.00 24.19      A    C
ATOM    539  CB  SER   127      13.991 -53.408  -1.590  1.00 23.41      A    C
ATOM    540  OG  SER   127      15.272 -53.060  -2.093  1.00 19.90      A    O
ATOM    541  C   SER   127      13.336 -51.074  -2.081  1.00 23.63      A    C
ATOM    542  O   SER   127      12.560 -51.066  -3.036  1.00 23.04      A    O
ATOM    543  N   GLY   128      14.279 -50.153  -1.907  1.00 23.89      A    N
ATOM    544  CA  GLY   128      14.572 -49.194  -2.958  1.00 25.02      A    C
ATOM    545  C   GLY   128      15.125 -49.844  -4.221  1.00 25.07      A    C
ATOM    546  O   GLY   128      15.369 -49.174  -5.221  1.00 25.81      A    O
ATOM    547  N   ASP   129      15.321 -51.157  -4.182  1.00 24.43      A    N
ATOM    548  CA  ASP   129      15.845 -51.872  -5.327  1.00 23.02      A    C
ATOM    549  CB  ASP   129      16.282 -53.270  -4.923  1.00 25.00      A    C
ATOM    550  CG  ASP   129      17.570 -53.266  -4.157  1.00 26.85      A    C
ATOM    551  OD1 ASP   129      18.089 -52.199  -3.878  1.00 29.00      A    O
ATOM    552  OD2 ASP   129      18.086 -54.365  -3.840  1.00 27.23      A    O
ATOM    553  C   ASP   129      14.833 -51.972  -6.433  1.00 22.97      A    C
ATOM    554  O   ASP   129      15.178 -52.312  -7.556  1.00 24.66      A    O
ATOM    555  N   LEU   130      13.580 -51.677  -6.114  1.00 23.97      A    N
ATOM    556  CA  LEU   130      12.493 -51.807  -7.077  1.00 23.49      A    C
ATOM    557  CB  LEU   130      11.257 -52.375  -6.393  1.00 22.01      A    C
ATOM    558  CG  LEU   130      11.441 -53.659  -5.597  1.00 21.51      A    C
ATOM    559  CD1 LEU   130      10.365 -53.753  -4.524  1.00 19.81      A    C
ATOM    560  CD2 LEU   130      11.390 -54.837  -6.543  1.00 20.89      A    C
ATOM    561  C   LEU   130      12.139 -50.464  -7.700  1.00 23.13      A    C
ATOM    562  O   LEU   130      11.138 -50.344  -8.410  1.00 23.93      A    O
ATOM    563  N   LEU   131      12.954 -49.453  -7.429  1.00 23.29      A    N
ATOM    564  CA  LEU   131      12.666 -48.112  -7.915  1.00 24.47      A    C
ATOM    565  CB  LEU   131      13.665 -47.117  -7.331  1.00 23.09      A    C
ATOM    566  CG  LEU   131      13.505 -46.795  -5.843  1.00 22.95      A    C
ATOM    567  CD1 LEU   131      14.687 -45.956  -5.418  1.00 22.36      A    C
ATOM    568  CD2 LEU   131      12.188 -46.066  -5.568  1.00 20.28      A    C
ATOM    569  C   LEU   131      12.662 -48.002  -9.437  1.00 25.34      A    C
ATOM    570  O   LEU   131      11.704 -47.493 -10.025  1.00 25.04      A    O
ATOM    571  N   GLU   132      13.726 -48.473 -10.077  1.00 27.24      A    N
ATOM    572  CA  GLU   132      13.782 -48.456 -11.534  1.00 28.71      A    C
ATOM    573  CB  GLU   132      15.045 -49.164 -12.034  1.00 30.19      A    C
ATOM    574  CG  GLU   132      16.295 -48.304 -12.008  1.00 32.76      A    C
ATOM    575  CD  GLU   132      17.528 -49.061 -12.454  1.00 34.92      A    C
ATOM    576  OE1 GLU   132      17.661 -50.244 -12.069  1.00 34.12      A    O
ATOM    577  OE2 GLU   132      18.359 -48.476 -13.191  1.00 36.98      A    O
ATOM    578  C   GLU   132      12.547 -49.136 -12.112  1.00 29.50      A    C
ATOM    579  O   GLU   132      12.005 -48.697 -13.128  1.00 30.00      A    O
ATOM    580  N   LEU   133      12.106 -50.207 -11.451  1.00 29.35      A    N
ATOM    581  CA  LEU   133      10.930 -50.960 -11.879  1.00 28.02      A    C
ATOM    582  CB  LEU   133      10.769 -52.229 -11.036  1.00 28.47      A    C
ATOM    583  CG  LEU   133       9.429 -52.959 -11.156  1.00 26.56      A    C
ATOM    584  CD1 LEU   133       9.241 -53.439 -12.578  1.00 26.64      A    C
ATOM    585  CD2 LEU   133       9.397 -54.123 -10.206  1.00 25.35      A    C
ATOM    586  C   LEU   133       9.660 -50.132 -11.769  1.00 27.30      A    C
ATOM    587  O   LEU   133       8.901 -50.022 -12.733  1.00 27.98      A    O
ATOM    588  N   ALA   134       9.435 -49.555 -10.592  1.00 25.72      A    N
ATOM    589  CA  ALA   134       8.198 -48.831 -10.314  1.00 24.04      A    C
ATOM    590  CB  ALA   134       8.147 -48.430  -8.847  1.00 24.07      A    C
ATOM    591  C   ALA   134       8.032 -47.598 -11.195  1.00 21.33      A    C
ATOM    592  O   ALA   134       6.917 -47.263 -11.603  1.00 21.21      A    O
ATOM    593  N   LEU   135       9.141 -46.927 -11.487  1.00 18.76      A    N
ATOM    594  CA  LEU   135       9.103 -45.742 -12.323  1.00 17.09      A    C
ATOM    595  CB  LEU   135      10.491 -45.136 -12.414  1.00 17.50      A    C
ATOM    596  CG  LEU   135      10.969 -44.445 -11.139  1.00 18.25      A    C
ATOM    597  CD1 LEU   135      12.446 -44.197 -11.239  1.00 18.59      A    C
ATOM    598  CD2 LEU   135      10.227 -43.139 -10.937  1.00 19.48      A    C
ATOM    599  C   LEU   135       8.579 -46.054 -13.713  1.00 17.06      A    C
ATOM    600  O   LEU   135       7.968 -45.213 -14.352  1.00 16.31      A    O
ATOM    601  N   LYS   136       8.802 -47.277 -14.178  1.00 18.88      A    N
ATOM    602  CA  LYS   136       8.217 -47.722 -15.438  1.00 18.70      A    C
ATOM    603  CB  LYS   136       8.974 -48.935 -15.967  1.00 19.33      A    C
ATOM    604  CG  LYS   136      10.417 -48.673 -16.319  1.00 21.05      A    C
ATOM    605  CD  LYS   136      11.128 -49.987 -16.527  1.00 25.15      A    C
ATOM    606  CE  LYS   136      12.629 -49.826 -16.488  1.00 29.77      A    C
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ATOM    607  NZ  LYS   136      13.174 -49.445 -17.824  1.00 33.49      A    N
ATOM    608  C   LYS   136       6.718 -48.050 -15.366  1.00 18.94      A    C
ATOM    609  O   LYS   136       6.062 -48.132 -16.403  1.00 19.33      A    O
ATOM    610  N   LEU   137       6.175 -48.236 -14.160  1.00 19.09      A    N
ATOM    611  CA  LEU   137       4.736 -48.494 -13.999  1.00 18.90      A    C
ATOM    612  CB  LEU   137       4.378 -48.748 -12.526  1.00 16.30      A    C
ATOM    613  CG  LEU   137       4.790 -50.075 -11.878  1.00 15.79      A    C
ATOM    614  CD1 LEU   137       4.652 -49.976 -10.379  1.00 14.21      A    C
ATOM    615  CD2 LEU   137       3.936 -51.207 -12.414  1.00 16.03      A    C
ATOM    616  C   LEU   137       3.903 -47.320 -14.522  1.00 20.04      A    C
ATOM    617  O   LEU   137       4.313 -46.163 -14.450  1.00 19.08      A    O
ATOM    618  N   PRO   138       2.713 -47.611 -15.057  1.00 21.62      A    N
ATOM    619  CD  PRO   138       1.982 -48.887 -14.964  1.00 23.18      A    C
ATOM    620  CA  PRO   138       1.918 -46.579 -15.721  1.00 23.00      A    C
ATOM    621  CB  PRO   138       0.862 -47.385 -16.471  1.00 22.04      A    C
ATOM    622  CG  PRO   138       0.653 -48.578 -15.607  1.00 21.35      A    C
ATOM    623  C   PRO   138       1.306 -45.617 -14.705  1.00 22.95      A    C
ATOM    624  O   PRO   138       0.952 -46.032 -13.608  1.00 23.32      A    O
ATOM    625  N   HIS   139       1.195 -44.343 -15.080  1.00 23.11      A    N
ATOM    626  CA  HIS   139       0.543 -43.325 -14.258  1.00 23.62      A    C
ATOM    627  CB  HIS   139      -0.625 -43.920 -13.473  1.00 27.86      A    C
ATOM    628  CG  HIS   139      -1.698 -44.509 -14.334  1.00 31.12      A    C
ATOM    629  CD2 HIS   139      -2.450 -43.963 -15.320  1.00 30.94      A    C
ATOM    630  ND1 HIS   139      -2.103 -45.823 -14.227  1.00 32.87      A    N
ATOM    631  CE1 HIS   139      -3.058 -46.061 -15.108  1.00 32.29      A    C
ATOM    632  NE2 HIS   139      -3.286 -44.949 -15.784  1.00 32.01      A    N
ATOM    633  C   HIS   139       1.476 -42.648 -13.278  1.00 22.78      A    C
ATOM    634  O   HIS   139       1.107 -41.655 -12.658  1.00 23.61      A    O
ATOM    635  N   VAL   140       2.677 -43.195 -13.124  1.00 22.21      A    N
ATOM    636  CA  VAL   140       3.650 -42.649 -12.189  1.00 19.92      A    C
ATOM    637  CB  VAL   140       4.872 -43.576 -12.021  1.00 18.16      A    C
ATOM    638  CG1 VAL   140       5.852 -42.973 -11.029  1.00 16.60      A    C
ATOM    639  CG2 VAL   140       4.427 -44.938 -11.555  1.00 16.58      A    C
ATOM    640  C   VAL   140       4.138 -41.297 -12.677  1.00 19.89      A    C
ATOM    641  O   VAL   140       4.483 -41.124 -13.841  1.00 18.54      A    O
ATOM    642  N   ASP   141       4.158 -40.336 -11.765  1.00 21.38      A    N
ATOM    643  CA  ASP   141       4.647 -39.001 -12.065  1.00 21.70      A    C
ATOM    644  CB  ASP   141       3.736 -37.976 -11.401  1.00 23.53      A    C
ATOM    645  CG  ASP   141       3.853 -36.609 -12.018  1.00 24.88      A    C
ATOM    646  OD1 ASP   141       4.819 -36.375 -12.776  1.00 26.94      A    O
ATOM    647  OD2 ASP   141       2.971 -35.766 -11.741  1.00 27.24      A    O
ATOM    648  C   ASP   141       6.057 -38.889 -11.508  1.00 19.63      A    C
ATOM    649  O   ASP   141       6.989 -38.466 -12.190  1.00 18.59      A    O
ATOM    650  N   TYR   142       6.194 -39.293 -10.256  1.00 18.63      A    N
ATOM    651  CA  TYR   142       7.488 -39.383  -9.609  1.00 19.45      A    C
ATOM    652  CB  TYR   142       8.044 -37.980  -9.335  1.00 18.30      A    C
ATOM    653  CG  TYR   142       7.128 -37.082  -8.527  1.00 19.04      A    C
ATOM    654  CD1 TYR   142       7.309 -36.920  -7.157  1.00 18.23      A    C
ATOM    655  CE1 TYR   142       6.496 -36.077  -6.421  1.00 17.86      A    C
ATOM    656  CD2 TYR   142       6.103 -36.373  -9.141  1.00 18.51      A    C
ATOM    657  CE2 TYR   142       5.285 -35.528  -8.413  1.00 18.75      A    C
ATOM    658  CZ  TYR   142       5.486 -35.382  -7.054  1.00 19.08      A    C
ATOM    659  OH  TYR   142       4.676 -34.537  -6.331  1.00 18.33      A    O
ATOM    660  C   TYR   142       7.333 -40.160  -8.302  1.00 19.33      A    C
ATOM    661  O   TYR   142       6.219 -40.402  -7.843  1.00 19.86      A    O
ATOM    662  N   ILE   143       8.447 -40.564  -7.709  1.00 18.21      A    N
ATOM    663  CA  ILE   143       8.388 -41.236  -6.421  1.00 18.63      A    C
ATOM    664  CB  ILE   143       8.795 -42.710  -6.568  1.00 19.75      A    C
ATOM    665  CG2 ILE   143       8.693 -43.414  -5.224  1.00 19.43      A    C
ATOM    666  CG1 ILE   143       7.907 -43.383  -7.620  1.00 18.58      A    C
ATOM    667  CD1 ILE   143       8.575 -44.520  -8.370  1.00 15.29      A    C
ATOM    668  C   ILE   143       9.348 -40.544  -5.471  1.00 18.03      A    C
ATOM    669  O   ILE   143      10.498 -40.326  -5.819  1.00 20.14      A    O
ATOM    670  N   GLU   144       8.876 -40.182  -4.284  1.00 18.26      A    N
ATOM    671  CA  GLU   144       9.733 -39.529  -3.292  1.00 19.45      A    C
ATOM    672  CB  GLU   144       9.167 -38.157  -2.908  1.00 18.29      A    C
ATOM    673  CG  GLU   144      10.120 -37.303  -2.068  1.00 19.87      A    C
ATOM    674  CD  GLU   144       9.478 -36.013  -1.543  1.00 21.24      A    C
ATOM    675  OE1 GLU   144       8.950 -35.224  -2.362  1.00 22.48      A    O
ATOM    676  OE2 GLU   144       9.503 -35.787  -0.307  1.00 19.26      A    O
ATOM    677  C   GLU   144       9.880 -40.392  -2.041  1.00 19.63      A    C
ATOM    678  O   GLU   144       8.907 -40.944  -1.536  1.00 20.55      A    O
ATOM    679  N   GLU   145      11.100 -40.514  -1.539  1.00 20.79      A    N
ATOM    680  CA  GLU   145      11.344 -41.370  -0.384  1.00 22.29      A    C
ATOM    681  CB  GLU   145      12.780 -41.881  -0.408  1.00 24.00      A    C
ATOM    682  CG  GLU   145      13.257 -42.460   0.904  1.00 25.67      A    C
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ATOM    683  CD  GLU   145      14.657 -41.994   1.229  1.00 27.32      A    C
ATOM    684  OE1 GLU   145      14.909 -40.770   1.121  1.00 29.46      A    O
ATOM    685  OE2 GLU   145      15.501 -42.843   1.578  1.00 25.95      A    O
ATOM    686  C   GLU   145      11.071 -40.648   0.934  1.00 22.03      A    C
ATOM    687  O   GLU   145      11.501 -39.512   1.141  1.00 21.79      A    O
ATOM    688  N   ASP   146      10.361 -41.330   1.826  1.00 21.77      A    N
ATOM    689  CA  ASP   146       9.805 -40.698   3.010  1.00 21.81      A    C
ATOM    690  CB  ASP   146       8.946 -41.693   3.792  1.00 23.57      A    C
ATOM    691  CG  ASP   146       8.061 -41.015   4.831  1.00 26.04      A    C
ATOM    692  OD1 ASP   146       7.680 -39.836   4.626  1.00 25.09      A    O
ATOM    693  OD2 ASP   146       7.735 -41.666   5.851  1.00 26.40      A    O
ATOM    694  C   ASP   146      10.901 -40.164   3.902  1.00 20.29      A    C
ATOM    695  O   ASP   146      11.944 -40.767   4.066  1.00 20.15      A    O
ATOM    696  N   SER   147      10.653 -39.000   4.472  1.00 20.99      A    N
ATOM    697  CA  SER   147      11.635 -38.327   5.320  1.00 21.78      A    C
ATOM    698  CB  SER   147      12.306 -37.175   4.553  1.00 21.45      A    C
ATOM    699  OG  SER   147      11.393 -36.616   3.622  1.00 21.30      A    O
ATOM    700  C   SER   147      10.917 -37.800   6.539  1.00 22.03      A    C
ATOM    701  O   SER   147       9.683 -37.841   6.594  1.00 22.28      A    O
ATOM    702  N   SER   148      11.682 -37.280   7.495  1.00 20.88      A    N
ATOM    703  CA  SER   148      11.097 -36.709   8.695  1.00 18.56      A    C
ATOM    704  CB  SER   148      12.030 -36.971   9.875  1.00 17.76      A    C
ATOM    705  OG  SER   148      11.817 -38.303  10.343  1.00 23.02      A    O
ATOM    706  C   SER   148      10.740 -35.220   8.591  1.00 17.96      A    C
ATOM    707  O   SER   148      11.243 -34.500   7.705  1.00 18.68      A    O
ATOM    708  N   VAL   149       9.817 -34.797   9.464  1.00 17.56      A    N
ATOM    709  CA  VAL   149       9.528 -33.382   9.732  1.00 14.60      A    C
ATOM    710  CB  VAL   149       8.123 -32.931   9.188  1.00 10.11      A    C
ATOM    711  CG1 VAL   149       8.098 -32.979   7.681  1.00  7.29      A    C
ATOM    712  CG2 VAL   149       7.022 -33.790   9.768  1.00  4.64      A    C
ATOM    713  C   VAL   149       9.565 -33.145  11.243  1.00 14.75      A    C
ATOM    714  O   VAL   149       9.483 -34.090  12.034  1.00 13.75      A    O
ATOM    715  N   PHE   150       9.687 -31.882  11.635  1.00 14.76      A    N
ATOM    716  CA  PHE   150       9.958 -31.564  13.020  1.00 14.98      A    C
ATOM    717  CB  PHE   150      11.455 -31.301  13.209  1.00 13.78      A    C
ATOM    718  CG  PHE   150      12.340 -32.416  12.706  1.00 13.73      A    C
ATOM    719  CD1 PHE   150      12.825 -32.408  11.399  1.00 13.23      A    C
ATOM    720  CD2 PHE   150      12.699 -33.464  13.544  1.00 12.73      A    C
ATOM    721  CE1 PHE   150      13.651 -33.424  10.940  1.00 13.20      A    C
ATOM    722  CE2 PHE   150      13.528 -34.488  13.093  1.00 12.62      A    C
ATOM    723  CZ  PHE   150      14.007 -34.468  11.789  1.00 13.49      A    C
ATOM    724  C   PHE   150       9.161 -30.358  13.493  1.00 15.59      A    C
ATOM    725  O   PHE   150       9.022 -29.371  12.769  1.00 15.48      A    O
ATOM    726  N   ALA   151       8.652 -30.462  14.721  1.00 16.70      A    N
ATOM    727  CA  ALA   151       7.941 -29.391  15.405  1.00 17.59      A    C
ATOM    728  CB  ALA   151       7.546 -29.861  16.800  1.00 14.66      A    C
ATOM    729  C   ALA   151       8.807 -28.134  15.499  1.00 20.08      A    C
ATOM    730  O   ALA   151       9.987 -28.214  15.863  1.00 21.40      A    O
ATOM    731  N   GLN   152       8.224 -26.974  15.185  1.00 21.45      A    N
ATOM    732  CA  GLN   152       8.999 -25.734  15.123  1.00 22.94      A    C
ATOM    733  CB  GLN   152       8.816 -25.094  13.751  1.00 22.61      A    C
ATOM    734  CG  GLN   152       9.251 -25.981  12.600  1.00 24.12      A    C
ATOM    735  CD  GLN   152      10.729 -26.298  12.639  1.00 24.23      A    C
ATOM    736  OE1 GLN   152      11.555 -25.426  12.900  1.00 25.38      A    O
ATOM    737  NE2 GLN   152      11.073 -27.553  12.384  1.00 25.68      A    N
ATOM    738  C   GLN   152       8.690 -24.693  16.212  1.00 24.95      A    C
ATOM    739  O   GLN   152       8.769 -23.483  15.892  1.00 25.70      A    O
ATOM    740  OXT GLN   152       8.403 -25.079  17.372  1.00 25.96      A    O
TER     741       GLN   152                                              A
ATOM    742  CB  SER   153      -2.118  -5.652  31.558  1.00 64.33      B    C
ATOM    743  OG  SER   153      -3.502  -5.729  31.243  1.00 65.73      B    O
ATOM    744  C   SER   153      -1.938  -7.936  30.526  1.00 60.48      B    C
ATOM    745  O   SER   153      -2.486  -9.027  30.714  1.00 61.28      B    O
ATOM    746  N   SER   153      -0.031  -6.953  31.819  1.00 62.19      B    N
ATOM    747  CA  SER   153      -1.516  -7.054  31.716  1.00 62.27      B    C
ATOM    748  N   ILE   154      -1.691  -7.442  29.312  1.00 56.98      B    N
ATOM    749  CA  ILE   154      -1.789  -8.241  28.096  1.00 53.64      B    C
ATOM    750  CB  ILE   154      -2.590  -7.497  26.997  1.00 54.07      B    C
ATOM    751  CG2 ILE   154      -2.566  -8.298  25.696  1.00 53.46      B    C
ATOM    752  CG1 ILE   154      -4.033  -7.284  27.459  1.00 54.70      B    C
ATOM    753  CD1 ILE   154      -4.702  -8.539  27.999  1.00 55.43      B    C
ATOM    754  C   ILE   154      -0.387  -8.536  27.576  1.00 50.51      B    C
ATOM    755  O   ILE   154       0.459  -7.638  27.493  1.00 51.05      B    O
ATOM    756  N   PRO   155      -0.124  -9.807  27.215  1.00 46.28      B    N
ATOM    757  CD  PRO   155      -1.077 -10.927  27.297  1.00 44.83      B    C
ATOM    758  CA  PRO   155       1.190 -10.235  26.719  1.00 42.26      B    C
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ATOM    759  CB  PRO   155       1.044 -11.747  26.515  1.00 43.54      B    C
ATOM    760  CG  PRO   155      -0.419 -12.012  26.469  1.00 43.77      B    C
ATOM    761  C   PRO   155       1.624  -9.511  25.468  1.00 38.35      B    C
ATOM    762  O   PRO   155       0.810  -9.280  24.581  1.00 37.59      B    O
ATOM    763  N   TRP   156       2.883  -9.073  25.464  1.00 35.05      B    N
ATOM    764  CA  TRP   156       3.404  -8.189  24.434  1.00 31.77      B    C
ATOM    765  CB  TRP   156       4.931  -8.188  24.492  1.00 30.82      B    C
ATOM    766  CG  TRP   156       5.560  -9.409  23.877  1.00 29.15      B    C
ATOM    767  CD2 TRP   156       5.935  -9.573  22.504  1.00 28.01      B    C
ATOM    768  CE2 TRP   156       6.429 -10.887  22.365  1.00 28.46      B    C
ATOM    769  CE3 TRP   156       5.893  -8.741  21.383  1.00 26.98      B    C
ATOM    770  CD1 TRP   156       5.845 -10.597  24.498  1.00 28.63      B    C
ATOM    771  NE1 TRP   156       6.367 -11.489  23.592  1.00 27.53      B    N
ATOM    772  CZ2 TRP   156       6.878 -11.380  21.145  1.00 27.62      B    C
ATOM    773  CZ3 TRP   156       6.337  -9.230  20.183  1.00 27.01      B    C
ATOM    774  CH2 TRP   156       6.821 -10.538  20.068  1.00 27.44      B    C
ATOM    775  C   TRP   156       2.945  -8.565  23.034  1.00 30.37      B    C
ATOM    776  O   TRP   156       2.745  -7.698  22.188  1.00 29.94      B    O
ATOM    777  N   ASN   157       2.790  -9.860  22.792  1.00 30.00      B    N
ATOM    778  CA  ASN   157       2.496 -10.358  21.461  1.00 30.04      B    C
ATOM    779  CB  ASN   157       2.841 -11.844  21.387  1.00 30.90      B    C
ATOM    780  CG  ASN   157       2.297 -12.631  22.566  1.00 32.76      B    C
ATOM    781  OD1 ASN   157       2.678 -12.393  23.720  1.00 32.26      B    O
ATOM    782  ND2 ASN   157       1.406 -13.582  22.282  1.00 32.40      B    N
ATOM    783  C   ASN   157       1.042 -10.132  21.050  1.00 29.16      B    C
ATOM    784  O   ASN   157       0.759  -9.778  19.905  1.00 28.18      B    O
ATOM    785  N   LEU   158       0.118 -10.340  21.979  1.00 28.71      B    N
ATOM    786  CA  LEU   158      -1.275 -10.005  21.720  1.00 29.10      B    C
ATOM    787  CB  LEU   158      -2.159 -10.499  22.869  1.00 26.66      B    C
ATOM    788  CG  LEU   158      -2.299 -12.012  23.001  1.00 23.99      B    C
ATOM    789  CD1 LEU   158      -3.369 -12.316  24.011  1.00 22.11      B    C
ATOM    790  CD2 LEU   158      -2.656 -12.624  21.661  1.00 22.63      B    C
ATOM    791  C   LEU   158      -1.380  -8.493  21.595  1.00 30.36      B    C
ATOM    792  O   LEU   158      -2.015  -7.956  20.690  1.00 27.77      B    O
ATOM    793  N   GLU   159      -0.734  -7.818  22.531  1.00 34.34      B    N
ATOM    794  CA  GLU   159      -0.599  -6.377  22.499  1.00 38.03      B    C
ATOM    795  CB  GLU   159       0.373  -5.958  23.596  1.00 42.36      B    C
ATOM    796  CG  GLU   159       0.570  -4.472  23.754  1.00 49.60      B    C
ATOM    797  CD  GLU   159       1.497  -4.146  24.917  1.00 54.93      B    C
ATOM    798  OE1 GLU   159       1.838  -5.078  25.690  1.00 56.11      B    O
ATOM    799  OE2 GLU   159       1.884  -2.960  25.053  1.00 57.72      B    O
ATOM    800  C   GLU   159      -0.081  -5.946  21.133  1.00 37.92      B    C
ATOM    801  O   GLU   159      -0.657  -5.065  20.497  1.00 38.35      B    O
ATOM    802  N   ARG   160       0.993  -6.595  20.685  1.00 37.80      B    N
ATOM    803  CA  ARG   160       1.690  -6.234  19.451  1.00 37.18      B    C
ATOM    804  CB  ARG   160       2.871  -7.182  19.217  1.00 38.09      B    C
ATOM    805  CG  ARG   160       3.727  -6.822  18.011  1.00 40.19      B    C
ATOM    806  CD  ARG   160       4.297  -5.430  18.181  1.00 43.31      B    C
ATOM    807  NE  ARG   160       4.594  -5.181  19.590  1.00 47.57      B    N
ATOM    808  CZ  ARG   160       5.820  -5.112  20.106  1.00 48.38      B    C
ATOM    809  NH1 ARG   160       5.986  -4.890  21.409  1.00 48.57      B    N
ATOM    810  NH2 ARG   160       6.879  -5.250  19.321  1.00 47.72      B    N
ATOM    811  C   ARG   160       0.789  -6.255  18.220  1.00 36.61      B    C
ATOM    812  O   ARG   160       0.748  -5.290  17.457  1.00 35.07      B    O
ATOM    813  N   ILE   161       0.071  -7.361  18.034  1.00 37.09      B    N
ATOM    814  CA  ILE   161      -0.735  -7.569  16.832  1.00 36.70      B    C
ATOM    815  CB  ILE   161      -1.083  -9.056  16.649  1.00 34.95      B    C
ATOM    816  CG2 ILE   161       0.167  -9.895  16.787  1.00 34.16      B    C
ATOM    817  CG1 ILE   161      -2.096  -9.495  17.698  1.00 31.99      B    C
ATOM    818  CD1 ILE   161      -2.663 -10.865  17.425  1.00 31.11      B    C
ATOM    819  C   ILE   161      -2.037  -6.760  16.842  1.00 37.39      B    C
ATOM    820  O   ILE   161      -2.845  -6.851  15.922  1.00 36.70      B    O
ATOM    821  N   THR   162      -2.228  -5.972  17.893  1.00 38.96      B    N
ATOM    822  CA  THR   162      -3.346  -5.046  17.977  1.00 39.63      B    C
ATOM    823  CB  THR   162      -3.960  -5.062  19.366  1.00 37.84      B    C
ATOM    824  OG1 THR   162      -4.768  -6.230  19.511  1.00 36.55      B    O
ATOM    825  CG2 THR   162      -4.795  -3.828  19.588  1.00 37.94      B    C
ATOM    826  C   THR   162      -2.896  -3.629  17.696  1.00 42.17      B    C
ATOM    827  O   THR   162      -2.007  -3.116  18.373  1.00 43.46      B    O
ATOM    828  N   PRO   163      -3.507  -2.973  16.695  1.00 44.52      B    N
ATOM    829  CD  PRO   163      -4.456  -3.541  15.722  1.00 44.90      B    C
ATOM    830  CA  PRO   163      -3.231  -1.557  16.428  1.00 46.29      B    C
ATOM    831  CB  PRO   163      -3.865  -1.316  15.062  1.00 45.47      B    C
ATOM    832  CG  PRO   163      -4.953  -2.330  14.977  1.00 45.09      B    C
ATOM    833  C   PRO   163      -3.824  -0.652  17.508  1.00 48.53      B    C
ATOM    834  O   PRO   163      -4.773  -1.027  18.198  1.00 47.74      B    O
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ATOM    835  N   PRO   164      -3.266   0.559  17.657  1.00 50.33      B    N
ATOM    836  CD  PRO   164      -2.151   1.049  16.825  1.00 51.64      B    C
ATOM    837  CA  PRO   164      -3.672   1.546  18.666  1.00 50.37      B    C
ATOM    838  CB  PRO   164      -2.839   2.779  18.314  1.00 50.83      B    C
ATOM    839  CG  PRO   164      -1.638   2.229  17.606  1.00 51.59      B    C
ATOM    840  C   PRO   164      -5.170   1.845  18.648  1.00 49.88      B    C
ATOM    841  O   PRO   164      -5.784   2.055  19.696  1.00 49.10      B    O
ATOM    842  N   GLY   176     -14.144 -13.280  18.361  1.00 32.04      B    N
ATOM    843  CA  GLY   176     -12.995 -13.701  17.570  1.00 33.60      B    C
ATOM    844  C   GLY   176     -13.330 -14.191  16.169  1.00 33.76      B    C
ATOM    845  O   GLY   176     -14.095 -13.548  15.451  1.00 36.77      B    O
ATOM    846  N   GLY   177     -12.770 -15.327  15.766  1.00 32.58      B    N
ATOM    847  CA  GLY   177     -12.976 -15.794  14.401  1.00 31.46      B    C
ATOM    848  C   GLY   177     -14.045 -16.865  14.275  1.00 30.95      B    C
ATOM    849  O   GLY   177     -13.760 -18.006  13.924  1.00 30.29      B    O
ATOM    850  N   SER   178     -15.288 -16.490  14.552  1.00 30.81      B    N
ATOM    851  CA  SER   178     -16.381 -17.452  14.695  1.00 30.89      B    C
ATOM    852  CB  SER   178     -17.619 -16.729  15.221  1.00 30.25      B    C
ATOM    853  OG  SER   178     -17.669 -15.393  14.738  1.00 32.95      B    O
ATOM    854  C   SER   178     -16.747 -18.256  13.437  1.00 30.60      B    C
ATOM    855  O   SER   178     -17.146 -19.414  13.539  1.00 31.32      B    O
ATOM    856  N   LEU   179     -16.613 -17.655  12.258  1.00 29.16      B    N
ATOM    857  CA  LEU   179     -16.991 -18.327  11.022  1.00 28.05      B    C
ATOM    858  CB  LEU   179     -17.207 -17.306   9.900  1.00 27.27      B    C
ATOM    859  CG  LEU   179     -18.273 -16.239  10.143  1.00 26.59      B    C
ATOM    860  CD1 LEU   179     -18.583 -15.514   8.855  1.00 22.84      B    C
ATOM    861  CD2 LEU   179     -19.514 -16.895  10.699  1.00 26.13      B    C
ATOM    862  C   LEU   179     -15.968 -19.359  10.554  1.00 28.95      B    C
ATOM    863  O   LEU   179     -16.306 -20.296   9.832  1.00 29.75      B    O
ATOM    864  N   VAL   180     -14.709 -19.194  10.939  1.00 29.57      B    N
ATOM    865  CA  VAL   180     -13.690 -20.076  10.392  1.00 28.73      B    C
ATOM    866  CB  VAL   180     -12.380 -19.321  10.040  1.00 29.20      B    C
ATOM    867  CG1 VAL   180     -12.692 -17.928   9.531  1.00 28.98      B    C
ATOM    868  CG2 VAL   180     -11.467 -19.283  11.232  1.00 30.37      B    C
ATOM    869  C   VAL   180     -13.350 -21.229  11.329  1.00 28.24      B    C
ATOM    870  O   VAL   180     -13.657 -21.211  12.525  1.00 26.19      B    O
ATOM    871  N   GLU   181     -12.718 -22.240  10.746  1.00 29.33      B    N
ATOM    872  CA  GLU   181     -12.366 -23.461  11.439  1.00 30.53      B    C
ATOM    873  CB  GLU   181     -13.206 -24.615  10.894  1.00 32.40      B    C
ATOM    874  CG  GLU   181     -13.309 -25.810  11.806  1.00 37.22      B    C
ATOM    875  CD  GLU   181     -14.641 -26.513  11.661  1.00 40.66      B    C
ATOM    876  OE1 GLU   181     -14.661 -27.688  11.235  1.00 44.08      B    O
ATOM    877  OE2 GLU   181     -15.675 -25.885  11.971  1.00 42.45      B    O
ATOM    878  C   GLU   181     -10.881 -23.710  11.186  1.00 29.82      B    C
ATOM    879  O   GLU   181     -10.433 -23.763  10.036  1.00 28.64      B    O
ATOM    880  N   VAL   182     -10.120 -23.844  12.267  1.00 29.16      B    N
ATOM    881  CA  VAL   182      -8.683 -24.032  12.162  1.00 27.69      B    C
ATOM    882  CB  VAL   182      -7.935 -23.145  13.172  1.00 26.22      B    C
ATOM    883  CG1 VAL   182      -6.457 -23.419  13.117  1.00 25.33      B    C
ATOM    884  CG2 VAL   182      -8.186 -21.694  12.847  1.00 26.31      B    C
ATOM    885  C   VAL   182      -8.309 -25.483  12.384  1.00 27.38      B    C
ATOM    886  O   VAL   182      -8.629 -26.073  13.415  1.00 26.94      B    O
ATOM    887  N   TYR   183      -7.644 -26.058  11.391  1.00 28.09      B    N
ATOM    888  CA  TYR   183      -7.113 -27.402  11.515  1.00 29.37      B    C
ATOM    889  CB  TYR   183      -7.221 -28.135  10.183  1.00 29.70      B    C
ATOM    890  CG  TYR   183      -8.562 -28.795   9.965  1.00 31.89      B    C
ATOM    891  CD1 TYR   183      -9.592 -28.132   9.300  1.00 31.43      B    C
ATOM    892  CE1 TYR   183     -10.818 -28.743   9.094  1.00 31.92      B    C
ATOM    893  CD2 TYR   183      -8.797 -30.087  10.416  1.00 32.07      B    C
ATOM    894  CE2 TYR   183     -10.014 -30.704  10.213  1.00 32.69      B    C
ATOM    895  CZ  TYR   183     -11.024 -30.032   9.554  1.00 33.05      B    C
ATOM    896  OH  TYR   183     -12.237 -30.666   9.360  1.00 34.31      B    O
ATOM    897  C   TYR   183      -5.661 -27.315  11.944  1.00 30.17      B    C
ATOM    898  O   TYR   183      -4.868 -26.596  11.339  1.00 31.22      B    O
ATOM    899  N   LEU   184      -5.326 -28.045  12.999  1.00 30.04      B    N
ATOM    900  CA  LEU   184      -3.963 -28.099  13.502  1.00 30.29      B    C
ATOM    901  CB  LEU   184      -3.959 -27.740  14.991  1.00 30.85      B    C
ATOM    902  CG  LEU   184      -2.652 -27.812  15.786  1.00 32.45      B    C
ATOM    903  CD1 LEU   184      -1.684 -26.740  15.303  1.00 32.56      B    C
ATOM    904  CD2 LEU   184      -2.056 -27.627  17.278  1.00 31.96      B    C
ATOM    905  C   LEU   184      -3.300 -29.499  13.294  1.00 30.49      B    C
ATOM    906  O   LEU   184      -3.897 -30.484  13.819  1.00 30.53      B    O
ATOM    907  N   LEU   185      -2.312 -29.594  12.515  1.00 30.52      B    N
ATOM    908  CA  LEU   185      -1.579 -30.847  12.432  1.00 32.04      B    C
ATOM    909  CB  LEU   185      -1.178 -31.162  10.991  1.00 32.70      B    C
ATOM    910  CG  LEU   185      -2.326 -31.422  10.023  1.00 33.68      B    C
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ATOM    911  CD1 LEU   185      -2.960 -30.097   9.644  1.00 34.74      B    C
ATOM    912  CD2 LEU   185      -1.813 -32.130   8.788  1.00 33.81      B    C
ATOM    913  C   LEU   185      -0.340 -30.715  13.284  1.00 32.10      B    C
ATOM    914  O   LEU   185       0.518 -29.875  13.016  1.00 33.06      B    O
ATOM    915  N   ASP   186      -0.244 -31.547  14.312  1.00 32.16      B    N
ATOM    916  CA  ASP   186       0.716 -31.296  15.365  1.00 33.03      B    C
ATOM    917  CB  ASP   186       0.299 -30.036  16.118  1.00 35.85      B    C
ATOM    918  CG  ASP   186       1.439 -29.410  16.876  1.00 40.10      B    C
ATOM    919  OD1 ASP   186       2.197 -28.616  16.272  1.00 41.27      B    O
ATOM    920  OD2 ASP   186       1.575 -29.714  18.082  1.00 44.20      B    O
ATOM    921  C   ASP   186       0.797 -32.485  16.315  1.00 32.10      B    C
ATOM    922  O   ASP   186      -0.362 -33.583  15.981  1.00 33.15      B    O
ATOM    923  N   THR   187       1.365 -32.265  17.494  1.00 30.56      B    N
ATOM    924  CA  THR   187       1.426 -33.291  18.525  1.00 29.30      B    C
ATOM    925  CB  THR   187       2.265 -32.840  19.706  1.00 29.76      B    C
ATOM    926  OG1 THR   187       1.568 -31.790  20.387  1.00 28.42      B    O
ATOM    927  CG2 THR   187       3.625 -32.325  19.235  1.00 28.84      B    C
ATOM    928  C   THR   187       0.021 -33.501  19.042  1.00 29.16      B    C
ATOM    929  O   THR   187      -0.934 -32.861  18.496  1.00 30.34      B    O
ATOM    930  N   SER   188      -0.101 -34.279  20.109  1.00 29.21      B    N
ATOM    931  CA  SER   188      -1.387 -34.471  20.756  1.00 28.11      B    C
ATOM    932  CB  SER   188      -1.352 -35.725  21.637  1.00 25.78      B    C
ATOM    933  OG  SER   188      -0.424 -35.577  22.688  1.00 23.67      B    O
ATOM    934  C   SER   188      -1.703 -33.237  21.594  1.00 29.15      B    C
ATOM    935  O   SER   188      -0.801 -32.520  22.020  1.00 29.20      B    O
ATOM    936  N   ILE   189      -2.991 -32.993  21.815  1.00 31.00      B    N
ATOM    937  CA  ILE   189      -3.458 -31.824  22.560  1.00 31.96      B    C
ATOM    938  CB  ILE   189      -4.547 -31.091  21.767  1.00 32.72      B    C
ATOM    939  CG2 ILE   189      -5.002 -29.863  22.512  1.00 34.17      B    C
ATOM    940  CG1 ILE   189      -4.000 -30.715  20.393  1.00 34.80      B    C
ATOM    941  CD1 ILE   189      -2.550 -30.250  20.422  1.00 34.94      B    C
ATOM    942  C   ILE   189      -4.033 -32.223  23.915  1.00 31.62      B    C
ATOM    943  O   ILE   189      -4.439 -33.365  24.105  1.00 32.27      B    O
ATOM    944  N   GLN   190      -4.063 -31.292  24.860  1.00 32.36      B    N
ATOM    945  CA  GLN   190      -4.861 -31.489  26.064  1.00 33.98      B    C
ATOM    946  CB  GLN   190      -4.083 -31.063  27.305  1.00 32.89      B    C
ATOM    947  CG  GLN   190      -4.502 -31.817  28.548  1.00 36.37      B    C
ATOM    948  CD  GLN   190      -5.107 -30.914  29.599  1.00 39.51      B    C
ATOM    949  OE1 GLN   190      -6.187 -30.364  29.403  1.00 42.37      B    O
ATOM    950  NE2 GLN   190      -4.410 -30.750  30.726  1.00 40.58      B    N
ATOM    951  C   GLN   190      -6.135 -30.666  25.936  1.00 34.78      B    C
ATOM    952  O   GLN   190      -6.184 -29.509  26.356  1.00 35.39      B    O
ATOM    953  N   SER   191      -7.162 -31.280  25.353  1.00 35.46      B    N
ATOM    954  CA  SER   191      -8.255 -30.546  24.721  1.00 36.57      B    C
ATOM    955  CB  SER   191      -9.034 -31.458  23.787  1.00 36.91      B    C
ATOM    956  OG  SER   191      -9.937 -32.247  24.540  1.00 37.76      B    O
ATOM    957  C   SER   191      -9.231 -29.962  25.718  1.00 37.07      B    C
ATOM    958  O   SER   191     -10.183 -29.285  25.333  1.00 36.92      B    O
ATOM    959  N   ASP   192      -9.010 -30.245  26.995  1.00 37.93      B    N
ATOM    960  CA  ASP   192      -9.853 -29.681  28.032  1.00 38.96      B    C
ATOM    961  CB  ASP   192     -10.593 -30.797  28.784  1.00 43.84      B    C
ATOM    962  CG  ASP   192      -9.655 -31.797  29.420  1.00 47.60      B    C
ATOM    963  OD1 ASP   192      -8.434 -31.704  29.155  1.00 50.20      B    O
ATOM    964  OD2 ASP   192     -10.142 -32.669  30.182  1.00 48.68      B    O
ATOM    965  C   ASP   192      -9.078 -28.793  29.001  1.00 36.46      B    C
ATOM    966  O   ASP   192      -9.563 -28.451  30.078  1.00 35.83      B    O
ATOM    967  N   HIS   193      -7.872 -28.411  28.611  1.00 34.81      B    N
ATOM    968  CA  HIS   193      -7.174 -27.356  29.319  1.00 34.03      B    C
ATOM    969  CB  HIS   193      -5.859 -27.022  28.619  1.00 33.96      B    C
ATOM    970  CG  HIS   193      -5.066 -25.959  29.308  1.00 33.51      B    C
ATOM    971  CD2 HIS   193      -3.978 -26.038  30.108  1.00 33.22      B    C
ATOM    972  ND1 HIS   193      -5.379 -24.621  29.215  1.00 33.43      B    N
ATOM    973  CE1 HIS   193      -4.517 -23.920  29.930  1.00 33.67      B    C
ATOM    974  NE2 HIS   193      -3.658 -24.756  30.483  1.00 33.30      B    N
ATOM    975  C   HIS   193      -8.089 -26.138  29.309  1.00 33.86      B    C
ATOM    976  O   HIS   193      -8.669 -25.795  28.274  1.00 34.18      B    O
ATOM    977  N   ARG   194      -8.214 -25.488  30.461  1.00 32.52      B    N
ATOM    978  CA  ARG   194      -9.264 -24.507  30.662  1.00 31.10      B    C
ATOM    979  CB  ARG   194      -9.101 -23.830  32.026  1.00 29.16      B    C
ATOM    980  CG  ARG   194      -8.795 -22.340  31.959  1.00 28.99      B    C
ATOM    981  CD  ARG   194      -9.657 -21.522  32.934  1.00 26.01      B    C
ATOM    982  NE  ARG   194      -8.938 -21.075  34.132  1.00 23.97      B    N
ATOM    983  CZ  ARG   194      -7.921 -20.212  34.126  1.00 23.47      B    C
ATOM    984  NH1 ARG   194      -7.334 -19.864  35.262  1.00 23.18      B    N
ATOM    985  NH2 ARG   194      -7.479 -19.699  32.984  1.00 23.23      B    N
ATOM    986  C   ARG   194      -9.235 -23.471  29.552  1.00 31.87      B    C
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ATOM    987  O   ARG   194     -10.245 -22.835  29.249  1.00 32.31      B    O
ATOM    988  N   GLU   195      -8.073 -23.327  28.930  1.00 31.93      B    N
ATOM    989  CA  GLU   195      -7.834 -22.250  27.981  1.00 31.02      B    C
ATOM    990  CB  GLU   195      -6.324 -22.075  27.807  1.00 31.22      B    C
ATOM    991  CG  GLU   195      -5.917 -21.036  26.791  1.00 32.56      B    C
ATOM    992  CD  GLU   195      -5.879 -19.625  27.358  1.00 33.36      B    C
ATOM    993  OE1 GLU   195      -6.529 -19.363  28.404  1.00 31.05      B    O
ATOM    994  OE2 GLU   195      -5.191 -18.782  26.736  1.00 32.52      B    O
ATOM    995  C   GLU   195      -8.506 -22.462  26.617  1.00 30.14      B    C
ATOM    996  O   GLU   195      -8.857 -21.496  25.939  1.00 29.15      B    O
ATOM    997  N   ILE   196      -8.690 -23.720  26.221  1.00 29.91      B    N
ATOM    998  CA  ILE   196      -9.100 -24.037  24.855  1.00 29.25      B    C
ATOM    999  CB  ILE   196      -7.951 -24.687  24.064  1.00 27.84      B    C
ATOM   1000  CG2 ILE   196      -6.854 -23.670  23.801  1.00 26.24      B    C
ATOM   1001  CG1 ILE   196      -7.436 -25.909  24.834  1.00 26.53      B    C
ATOM   1002  CD1 ILE   196      -6.554 -26.819  24.033  1.00 24.05      B    C
ATOM   1003  C   ILE   196     -10.274 -24.999  24.811  1.00 30.60      B    C
ATOM   1004  O   ILE   196     -10.948 -25.135  23.788  1.00 30.51      B    O
ATOM   1005  N   GLU   197     -10.512 -25.686  25.916  1.00 32.24      B    N
ATOM   1006  CA  GLU   197     -11.490 -26.759  25.910  1.00 34.74      B    C
ATOM   1007  CB  GLU   197     -11.610 -27.350  27.309  1.00 35.54      B    C
ATOM   1008  CG  GLU   197     -12.153 -26.413  28.324  1.00 38.11      B    C
ATOM   1009  CD  GLU   197     -13.648 -26.430  28.324  1.00 41.53      B    C
ATOM   1010  OE1 GLU   197     -14.254 -25.395  28.669  1.00 45.11      B    O
ATOM   1011  OE2 GLU   197     -14.219 -27.484  27.971  1.00 43.30      B    O
ATOM   1012  C   GLU   197     -12.855 -26.287  25.413  1.00 34.89      B    C
ATOM   1013  O   GLU   197     -13.359 -25.251  25.844  1.00 34.37      B    O
ATOM   1014  N   GLY   198     -13.446 -27.046  24.497  1.00 34.64      B    N
ATOM   1015  CA  GLY   198     -14.707 -26.631  23.918  1.00 34.42      B    C
ATOM   1016  C   GLY   198     -14.534 -25.954  22.570  1.00 35.86      B    C
ATOM   1017  O   GLY   198     -15.436 -26.002  21.732  1.00 36.41      B    O
ATOM   1018  N   ARG   199     -13.383 -25.318  22.354  1.00 36.46      B    N
ATOM   1019  CA  ARG   199     -13.081 -24.694  21.066  1.00 37.77      B    C
ATOM   1020  CB  ARG   199     -12.258 -23.416  21.257  1.00 39.47      B    C
ATOM   1021  CG  ARG   199     -12.909 -22.372  22.129  1.00 41.61      B    C
ATOM   1022  CD  ARG   199     -11.888 -21.766  23.062  1.00 43.31      B    C
ATOM   1023  NE  ARG   199     -12.472 -20.711  23.877  1.00 45.69      B    N
ATOM   1024  CZ  ARG   199     -12.610 -19.453  23.475  1.00 48.05      B    C
ATOM   1025  NH1 ARG   199     -13.153 -18.552  24.283  1.00 49.22      B    N
ATOM   1026  NH2 ARG   199     -12.208 -19.095  22.262  1.00 48.95      B    N
ATOM   1027  C   ARG   199     -12.275 -25.673  20.242  1.00 37.48      B    C
ATOM   1028  O   ARG   199     -12.100 -25.507  19.042  1.00 37.33      B    O
ATOM   1029  N   VAL   200     -11.768 -26.692  20.916  1.00 38.28      B    N
ATOM   1030  CA  VAL   200     -10.894 -27.662  20.286  1.00 39.41      B    C
ATOM   1031  CB  VAL   200      -9.503 -27.665  20.976  1.00 40.93      B    C
ATOM   1032  CG1 VAL   200      -8.662 -28.836  20.488  1.00 40.39      B    C
ATOM   1033  CG2 VAL   200      -8.791 -26.346  20.688  1.00 40.81      B    C
ATOM   1034  C   VAL   200     -11.516 -29.051  20.361  1.00 39.27      B    C
ATOM   1035  O   VAL   200     -11.652 -29.628  21.443  1.00 39.35      B    O
ATOM   1036  N   MET   201     -11.904 -29.565  19.199  1.00 39.35      B    N
ATOM   1037  CA  MET   201     -12.387 -30.930  19.063  1.00 38.47      B    C
ATOM   1038  CB  MET   201     -13.537 -30.968  18.057  1.00 39.97      B    C
ATOM   1039  CG  MET   201     -14.150 -32.335  17.862  1.00 43.51      B    C
ATOM   1040  SD  MET   201     -13.407 -33.296  16.515  1.00 49.43      B    S
ATOM   1041  CE  MET   201     -13.750 -34.985  17.094  1.00 46.63      B    C
ATOM   1042  C   MET   201     -11.233 -31.801  18.575  1.00 37.52      B    C
ATOM   1043  O   MET   201     -10.835 -31.727  17.414  1.00 37.13      B    O
ATOM   1044  N   VAL   202     -10.680 -32.612  19.467  1.00 36.78      B    N
ATOM   1045  CA  VAL   202      -9.624 -33.533  19.076  1.00 35.76      B    C
ATOM   1046  CB  VAL   202      -8.927 -34.157  20.305  1.00 34.80      B    C
ATOM   1047  CG1 VAL   202      -8.212 -35.428  19.905  1.00 33.07      B    C
ATOM   1048  CG2 VAL   202      -7.934 -33.173  20.888  1.00 33.22      B    C
ATOM   1049  C   VAL   202     -10.226 -34.641  18.230  1.00 36.18      B    C
ATOM   1050  O   VAL   202     -11.025 -35.447  18.716  1.00 35.82      B    O
ATOM   1051  N   THR   203      -9.848 -34.668  16.956  1.00 36.30      B    N
ATOM   1052  CA  THR   203     -10.380 -35.662  16.041  1.00 35.62      B    C
ATOM   1053  CB  THR   203     -10.067 -35.316  14.568  1.00 34.74      B    C
ATOM   1054  OG1 THR   203      -8.719 -35.682  14.257  1.00 34.72      B    O
ATOM   1055  CG2 THR   203     -10.240 -33.821  14.334  1.00 33.32      B    C
ATOM   1056  C   THR   203      -9.745 -36.993  16.389  1.00 36.32      B    C
ATOM   1057  O   THR   203      -9.147 -37.151  17.447  1.00 37.09      B    O
ATOM   1058  N   ASP   204      -9.881 -37.956  15.497  1.00 37.59      B    N
ATOM   1059  CA  ASP   204      -9.461 -39.315  15.786  1.00 38.74      B    C
ATOM   1060  CB  ASP   204     -10.615 -40.264  15.511  1.00 42.30      B    C
ATOM   1061  CG  ASP   204     -11.162 -40.090  14.112  1.00 47.00      B    C
ATOM   1062  OD1 ASP   204     -10.783 -40.892  13.228  1.00 50.09      B    O
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ATOM   1063  OD2 ASP   204     -11.953 -39.137  13.891  1.00 48.97      B    O
ATOM   1064  C   ASP   204      -8.293 -39.671  14.886  1.00 37.76      B    C
ATOM   1065  O   ASP   204      -7.862 -40.823  14.846  1.00 38.21      B    O
ATOM   1066  N   PHE   205      -7.790 -38.684  14.151  1.00 35.84      B    N
ATOM   1067  CA  PHE   205      -6.668 -38.913  13.249  1.00 33.48      B    C
ATOM   1068  CB  PHE   205      -6.603 -37.818  12.190  1.00 31.69      B    C
ATOM   1069  CG  PHE   205      -5.544 -38.049  11.166  1.00 31.35      B    C
ATOM   1070  CD1 PHE   205      -4.288 -37.502  11.317  1.00 31.19      B    C
ATOM   1071  CD2 PHE   205      -5.798 -38.827  10.051  1.00 32.23      B    C
ATOM   1072  CE1 PHE   205      -3.303 -37.726  10.371  1.00 31.97      B    C
ATOM   1073  CE2 PHE   205      -4.813 -39.052   9.102  1.00 32.26      B    C
ATOM   1074  CZ  PHE   205      -3.568 -38.502   9.264  1.00 30.67      B    C
ATOM   1075  C   PHE   205      -5.314 -39.001  13.959  1.00 32.69      B    C
ATOM   1076  O   PHE   205      -4.991 -38.201  14.842  1.00 32.36      B    O
ATOM   1077  N   GLU   206      -4.515 -39.978  13.563  1.00 31.16      B    N
ATOM   1078  CA  GLU   206      -3.160 -40.042  14.060  1.00 30.22      B    C
ATOM   1079  CB  GLU   206      -3.156 -40.639  15.456  1.00 30.05      B    C
ATOM   1080  CG  GLU   206      -1.784 -40.771  16.058  1.00 32.36      B    C
ATOM   1081  CD  GLU   206      -1.797 -41.638  17.292  1.00 35.04      B    C
ATOM   1082  OE1 GLU   206      -1.549 -41.109  18.397  1.00 38.43      B    O
ATOM   1083  OE2 GLU   206      -2.063 -42.850  17.160  1.00 34.75      B    O
ATOM   1084  C   GLU   206      -2.250 -40.843  13.142  1.00 28.87      B    C
ATOM   1085  O   GLU   206      -2.529 -41.991  12.825  1.00 27.95      B    O
ATOM   1086  N   ASN   207      -1.163 -40.213  12.712  1.00 29.34      B    N
ATOM   1087  CA  ASN   207      -0.115 -40.894  11.966  1.00 29.09      B    C
ATOM   1088  CB  ASN   207      -0.283 -40.651  10.462  1.00 29.34      B    C
ATOM   1089  CG  ASN   207       0.661 -41.504   9.620  1.00 28.97      B    C
ATOM   1090  OD1 ASN   207       0.231 -42.440   8.961  1.00 27.55      B    O
ATOM   1091  ND2 ASN   207       1.956 -41.175   9.643  1.00 30.27      B    N
ATOM   1092  C   ASN   207       1.249 -40.382  12.415  1.00 28.09      B    C
ATOM   1093  O   ASN   207       1.760 -39.399  11.878  1.00 29.19      B    O
ATOM   1094  N   VAL   208       1.835 -41.057  13.397  1.00 26.80      B    N
ATOM   1095  CA  VAL   208       3.126 -40.654  13.925  1.00 25.13      B    C
ATOM   1096  CB  VAL   208       2.987 -40.135  15.371  1.00 23.53      B    C
ATOM   1097  CG1 VAL   208       2.211 -38.842  15.388  1.00 22.53      B    C
ATOM   1098  CG2 VAL   208       2.278 -41.159  16.219  1.00 21.79      B    C
ATOM   1099  C   VAL   208       4.127 -41.805  13.892  1.00 26.28      B    C
ATOM   1100  O   VAL   208       3.752 -42.969  13.992  1.00 26.48      B    O
ATOM   1101  N   PRO   209       5.418 -41.482  13.722  1.00 28.18      B    N
ATOM   1102  CD  PRO   209       5.845 -40.119  13.351  1.00 29.06      B    C
ATOM   1103  CA  PRO   209       6.562 -42.394  13.827  1.00 30.05      B    C
ATOM   1104  CB  PRO   209       7.692 -41.608  13.190  1.00 28.79      B    C
ATOM   1105  CG  PRO   209       7.339 -40.177  13.468  1.00 26.64      B    C
ATOM   1106  C   PRO   209       6.872 -42.725  15.280  1.00 33.78      B    C
ATOM   1107  O   PRO   209       6.671 -41.897  16.166  1.00 34.02      B    O
ATOM   1108  N   GLU   210       7.378 -43.924  15.529  1.00 38.83      B    N
ATOM   1109  CA  GLU   210       7.634 -44.337  16.899  1.00 44.60      B    C
ATOM   1110  CB  GLU   210       8.115 -45.777  16.947  1.00 46.99      B    C
ATOM   1111  CG  GLU   210       7.149 -46.753  16.350  1.00 52.51      B    C
ATOM   1112  CD  GLU   210       7.807 -48.078  16.052  1.00 56.68      B    C
ATOM   1113  OE1 GLU   210       8.749 -48.448  16.791  1.00 58.71      B    O
ATOM   1114  OE2 GLU   210       7.390 -48.744  15.077  1.00 58.58      B    O
ATOM   1115  C   GLU   210       8.682 -43.450  17.531  1.00 46.80      B    C
ATOM   1116  O   GLU   210       9.615 -42.998  16.867  1.00 45.67      B    O
ATOM   1117  N   GLU   211       8.532 -43.210  18.823  1.00 50.25      B    N
ATOM   1118  CA  GLU   211       9.508 -42.412  19.545  1.00 53.81      B    C
ATOM   1119  CB  GLU   211       9.122 -42.343  21.020  1.00 54.42      B    C
ATOM   1120  CG  GLU   211       7.641 -42.600  21.242  1.00 56.19      B    C
ATOM   1121  CD  GLU   211       6.791 -41.386  20.950  1.00 56.76      B    C
ATOM   1122  OE1 GLU   211       7.332 -40.269  21.041  1.00 58.23      B    O
ATOM   1123  OE2 GLU   211       5.589 -41.543  20.631  1.00 55.21      B    O
ATOM   1124  C   GLU   211      10.872 -43.060  19.410  1.00 55.25      B    C
ATOM   1125  O   GLU   211      10.981 -44.286  19.400  1.00 55.73      B    O
ATOM   1126  N   ASP   212      11.907 -42.237  19.296  1.00 57.47      B    N
ATOM   1127  CA  ASP   212      13.257 -42.726  19.497  1.00 59.60      B    C
ATOM   1128  CB  ASP   212      13.335 -43.386  20.870  1.00 61.46      B    C
ATOM   1129  CG  ASP   212      14.753 -43.585  21.348  1.00 63.54      B    C
ATOM   1130  OD1 ASP   212      15.701 -43.219  20.613  1.00 65.22      B    O
ATOM   1131  OD2 ASP   212      14.911 -44.119  22.468  1.00 65.41      B    O
ATOM   1132  C   ASP   212      13.595 -43.735  18.407  1.00 61.25      B    C
ATOM   1133  O   ASP   212      14.004 -43.364  17.310  1.00 62.44      B    O
ATOM   1134  N   LYS   222       5.879 -33.583  27.994  1.00 46.08      B    N
ATOM   1135  CA  LYS   222       4.625 -34.186  27.575  1.00 46.76      B    C
ATOM   1136  CB  LYS   222       3.455 -33.569  28.341  1.00 47.87      B    C
ATOM   1137  CG  LYS   222       3.481 -33.867  29.823  1.00 50.38      B    C
ATOM   1138  CD  LYS   222       2.205 -33.438  30.501  1.00 52.96      B    C
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ATOM   1139  CE  LYS   222       2.195 -33.878  31.954  1.00 55.83      B    C
ATOM   1140  NZ  LYS   222       1.005 -33.423  32.731  1.00 59.69      B    N
ATOM   1141  C   LYS   222       4.382 -34.040  26.079  1.00 45.41      B    C
ATOM   1142  O   LYS   222       4.825 -33.080  25.445  1.00 45.49      B    O
ATOM   1143  N   CYS   223       3.646 -35.003  25.540  1.00 43.53      B    N
ATOM   1144  CA  CYS   223       3.327 -35.050  24.125  1.00 41.93      B    C
ATOM   1145  C   CYS   223       2.291 -33.991  23.784  1.00 40.12      B    C
ATOM   1146  O   CYS   223       2.093 -33.666  22.619  1.00 39.88      B    O
ATOM   1147  CB  CYS   223       2.774 -36.424  23.779  1.00 43.01      B    C
ATOM   1148  SG  CYS   223       3.674 -37.758  24.630  1.00 45.72      B    S
ATOM   1149  N   ASP   224       1.613 -33.482  24.809  1.00 38.82      B    N
ATOM   1150  CA  ASP   224       0.589 -32.452  24.646  1.00 36.21      B    C
ATOM   1151  CB  ASP   224      -0.267 -32.332  25.907  1.00 38.96      B    C
ATOM   1152  CG  ASP   224      -0.850 -33.650  26.351  1.00 41.36      B    C
ATOM   1153  OD1 ASP   224      -0.148 -34.679  26.266  1.00 44.99      B    O
ATOM   1154  OD2 ASP   224      -2.017 -33.657  26.792  1.00 42.87      B    O
ATOM   1155  C   ASP   224       1.280 -31.118  24.421  1.00 34.07      B    C
ATOM   1156  O   ASP   224       0.778 -30.248  23.706  1.00 32.46      B    O
ATOM   1157  N   SER   225       2.442 -30.979  25.053  1.00 31.41      B    N
ATOM   1158  CA  SER   225       3.169 -29.721  25.120  1.00 28.08      B    C
ATOM   1159  CB  SER   225       4.651 -30.002  25.351  1.00 26.55      B    C
ATOM   1160  OG  SER   225       5.418 -28.847  25.088  1.00 27.45      B    O
ATOM   1161  C   SER   225       3.004 -28.809  23.906  1.00 26.58      B    C
ATOM   1162  O   SER   225       2.371 -27.760  23.991  1.00 25.91      B    O
ATOM   1163  N   HIS   226       3.576 -29.203  22.778  1.00 25.93      B    N
ATOM   1164  CA  HIS   226       3.721 -28.286  21.651  1.00 25.28      B    C
ATOM   1165  CB  HIS   226       4.530 -28.958  20.541  1.00 25.17      B    C
ATOM   1166  CG  HIS   226       4.905 -28.039  19.419  1.00 25.61      B    C
ATOM   1167  CD2 HIS   226       5.842 -27.065  19.339  1.00 25.96      B    C
ATOM   1168  ND1 HIS   226       4.308 -28.100  18.177  1.00 25.61      B    N
ATOM   1169  CE1 HIS   226       4.864 -27.205  17.380  1.00 26.84      B    C
ATOM   1170  NE2 HIS   226       5.799 -26.563  18.060  1.00 26.81      B    N
ATOM   1171  C   HIS   226       2.382 -27.807  21.095  1.00 24.30      B    C
ATOM   1172  O   HIS   226       2.193 -26.613  20.867  1.00 24.47      B    O
ATOM   1173  N   GLY   227       1.460 -28.742  20.883  1.00 23.30      B    N
ATOM   1174  CA  GLY   227       0.196 -28.419  20.251  1.00 22.28      B    C
ATOM   1175  C   GLY   227      -0.772 -27.718  21.184  1.00 23.20      B    C
ATOM   1176  O   GLY   227      -1.589 -26.907  20.754  1.00 23.88      B    O
ATOM   1177  N   THR   228      -0.684 -28.022  22.471  1.00 23.01      B    N
ATOM   1178  CA  THR   228      -1.561 -27.394  23.447  1.00 22.63      B    C
ATOM   1179  CB  THR   228      -1.394 -28.024  24.842  1.00 23.38      B    C
ATOM   1180  OG1 THR   228      -1.516 -29.450  24.748  1.00 23.12      B    O
ATOM   1181  CG2 THR   228      -2.456 -27.499  25.782  1.00 21.83      B    C
ATOM   1182  C   THR   228      -1.233 -25.913  23.542  1.00 22.38      B    C
ATOM   1183  O   THR   228      -2.121 -25.067  23.614  1.00 23.83      B    O
ATOM   1184  N   HIS   229       0.050 -25.592  23.534  1.00 22.67      B    N
ATOM   1185  CA  HIS   229       0.446 -24.198  23.574  1.00 22.18      B    C
ATOM   1186  CB  HIS   229       1.964 -24.077  23.632  1.00 22.78      B    C
ATOM   1187  CG  HIS   229       2.448 -22.720  24.041  1.00 23.95      B    C
ATOM   1188  CD2 HIS   229       2.598 -22.159  25.264  1.00 23.15      B    C
ATOM   1189  ND1 HIS   229       2.896 -21.783  23.132  1.00 23.61      B    N
ATOM   1190  CE1 HIS   229       3.305 -20.707  23.778  1.00 23.68      B    C
ATOM   1191  NE2 HIS   229       3.136 -20.909  25.074  1.00 23.94      B    N
ATOM   1192  C   HIS   229      -0.080 -23.527  22.321  1.00 21.39      B    C
ATOM   1193  O   HIS   229      -0.646 -22.443  22.381  1.00 20.56      B    O
ATOM   1194  N   LEU   230       0.096 -24.187  21.184  1.00 20.66      B    N
ATOM   1195  CA  LEU   230      -0.314 -23.605  19.919  1.00 21.11      B    C
ATOM   1196  CB  LEU   230       0.134 -24.495  18.764  1.00 21.06      B    C
ATOM   1197  CG  LEU   230       1.647 -24.572  18.592  1.00 22.74      B    C
ATOM   1198  CD1 LEU   230       1.993 -25.388  17.359  1.00 21.39      B    C
ATOM   1199  CD2 LEU   230       2.204 -23.156  18.478  1.00 23.15      B    C
ATOM   1200  C   LEU   230      -1.821 -23.401  19.858  1.00 21.39      B    C
ATOM   1201  O   LEU   230      -2.296 -22.404  19.304  1.00 21.76      B    O
ATOM   1202  N   ALA   231      -2.569 -24.341  20.431  1.00 21.33      B    N
ATOM   1203  CA  ALA   231      -4.026 -24.270  20.417  1.00 21.00      B    C
ATOM   1204  CB  ALA   231      -4.618 -25.474  21.138  1.00 18.46      B    C
ATOM   1205  C   ALA   231      -4.473 -22.979  21.090  1.00 22.12      B    C
ATOM   1206  O   ALA   231      -5.335 -22.264  20.572  1.00 21.92      B    O
ATOM   1207  N   GLY   232      -3.868 -22.685  22.240  1.00 22.96      B    N
ATOM   1208  CA  GLY   232      -4.181 -21.468  22.965  1.00 24.03      B    C
ATOM   1209  C   GLY   232      -3.799 -20.213  22.207  1.00 24.50      B    C
ATOM   1210  O   GLY   232      -4.573 -19.265  22.132  1.00 25.89      B    O
ATOM   1211  N   VAL   233      -2.609 -20.196  21.628  1.00 25.98      B    N
ATOM   1212  CA  VAL   233      -2.169 -19.014  20.908  1.00 26.71      B    C
ATOM   1213  CB  VAL   233      -0.805 -19.224  20.252  1.00 25.92      B    C
ATOM   1214  CG1 VAL   233      -0.390 -17.959  19.531  1.00 25.35      B    C
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ATOM   1215  CG2 VAL   233       0.215 -19.601  21.294  1.00 25.45      B    C
ATOM   1216  C   VAL   233      -3.171 -18.663  19.823  1.00 28.02      B    C
ATOM   1217  O   VAL   233      -3.284 -17.506  19.424  1.00 29.94      B    O
ATOM   1218  N   VAL   234      -3.906 -19.664  19.349  1.00 29.13      B    N
ATOM   1219  CA  VAL   234      -4.922 -19.447  18.320  1.00 28.46      B    C
ATOM   1220  CB  VAL   234      -5.133 -20.693  17.455  1.00 29.28      B    C
ATOM   1221  CG1 VAL   234      -6.540 -20.693  16.905  1.00 29.41      B    C
ATOM   1222  CG2 VAL   234      -4.135 -20.704  16.310  1.00 30.36      B    C
ATOM   1223  C   VAL   234      -6.281 -19.046  18.854  1.00 26.66      B    C
ATOM   1224  O   VAL   234      -6.833 -18.043  18.425  1.00 26.52      B    O
ATOM   1225  N   SER   235      -6.819 -19.832  19.781  1.00 25.95      B    N
ATOM   1226  CA  SER   235      -8.194 -19.648  20.222  1.00 26.62      B    C
ATOM   1227  CB  SER   235      -9.046 -20.814  19.740  1.00 27.81      B    C
ATOM   1228  OG  SER   235      -8.946 -21.905  20.639  1.00 30.51      B    O
ATOM   1229  C   SER   235      -8.354 -19.516  21.735  1.00 26.18      B    C
ATOM   1230  O   SER   235      -9.471 -19.410  22.230  1.00 25.76      B    O
ATOM   1231  N   GLY   236      -7.243 -19.531  22.464  1.00 26.66      B    N
ATOM   1232  CA  GLY   236      -7.295 -19.505  23.917  1.00 26.34      B    C
ATOM   1233  C   GLY   236      -8.147 -18.384  24.488  1.00 27.10      B    C
ATOM   1234  O   GLY   236      -8.256 -17.308  23.899  1.00 26.56      B    O
ATOM   1235  N   ARG   237      -8.749 -18.637  25.646  1.00 27.20      B    N
ATOM   1236  CA  ARG   237      -9.678 -17.693  26.258  1.00 28.66      B    C
ATOM   1237  CB  ARG   237     -10.412 -18.355  27.422  1.00 31.42      B    C
ATOM   1238  CG  ARG   237     -11.558 -19.253  26.990  1.00 36.85      B    C
ATOM   1239  CD  ARG   237     -11.972 -20.233  28.086  1.00 41.17      B    C
ATOM   1240  NE  ARG   237     -13.193 -20.955  27.729  1.00 46.01      B    N
ATOM   1241  CZ  ARG   237     -13.224 -22.136  27.114  1.00 48.46      B    C
ATOM   1242  NH1 ARG   237     -14.395 -22.702  26.833  1.00 49.07      B    N
ATOM   1243  NH2 ARG   237     -12.093 -22.756  26.784  1.00 48.35      B    N
ATOM   1244  C   ARG   237      -9.021 -16.416  26.748  1.00 27.85      B    C
ATOM   1245  O   ARG   237      -9.582 -15.331  26.603  1.00 26.35      B    O
ATOM   1246  N   ASP   238      -7.838 -16.546  27.334  1.00 28.76      B    N
ATOM   1247  CA  ASP   238      -7.118 -15.392  27.861  1.00 30.27      B    C
ATOM   1248  CB  ASP   238      -6.617 -15.685  29.280  1.00 33.62      B    C
ATOM   1249  CG  ASP   238      -7.733 -16.134  30.216  1.00 37.65      B    C
ATOM   1250  OD1 ASP   238      -8.918 -15.968  29.852  1.00 39.82      B    O
ATOM   1251  OD2 ASP   238      -7.429 -16.652  31.316  1.00 37.81      B    O
ATOM   1252  C   ASP   238      -5.935 -15.002  26.978  1.00 29.53      B    C
ATOM   1253  O   ASP   238      -5.566 -13.836  26.902  1.00 29.35      B    O
ATOM   1254  N   ALA   239      -5.342 -15.985  26.311  1.00 30.01      B    N
ATOM   1255  CA  ALA   239      -4.086 -15.779  25.601  1.00 29.41      B    C
ATOM   1256  CB  ALA   239      -3.024 -16.745  26.132  1.00 27.71      B    C
ATOM   1257  C   ALA   239      -4.251 -15.961  24.098  1.00 29.33      B    C
ATOM   1258  O   ALA   239      -3.298 -15.821  23.342  1.00 29.65      B    O
ATOM   1259  N   GLY   240      -5.462 -16.280  23.663  1.00 30.49      B    N
ATOM   1260  CA  GLY   240      -5.670 -16.564  22.258  1.00 29.73      B    C
ATOM   1261  C   GLY   240      -5.746 -15.301  21.434  1.00 29.50      B    C
ATOM   1262  O   GLY   240      -5.981 -14.218  21.967  1.00 28.77      B    O
ATOM   1263  N   VAL   241      -5.539 -15.449  20.129  1.00 29.66      B    N
ATOM   1264  CA  VAL   241      -5.801 -14.389  19.165  1.00 29.00      B    C
ATOM   1265  CB  VAL   241      -4.887 -14.534  17.934  1.00 27.73      B    C
ATOM   1266  CG1 VAL   241      -5.211 -13.462  16.912  1.00 27.00      B    C
ATOM   1267  CG2 VAL   241      -3.441 -14.432  18.359  1.00 27.06      B    C
ATOM   1268  C   VAL   241      -7.261 -14.426  18.703  1.00 29.65      B    C
ATOM   1269  O   VAL   241      -8.006 -13.454  18.869  1.00 30.73      B    O
ATOM   1270  N   ALA   242      -7.670 -15.555  18.130  1.00 29.93      B    N
ATOM   1271  CA  ALA   242      -9.018 -15.693  17.580  1.00 29.53      B    C
ATOM   1272  CB  ALA   242      -8.964 -16.377  16.220  1.00 28.85      B    C
ATOM   1273  C   ALA   242      -9.921 -16.478  18.515  1.00 28.85      B    C
ATOM   1274  O   ALA   242     -10.393 -17.559  18.176  1.00 28.12      B    O
ATOM   1275  N   LYS   243     -10.159 -15.930  19.696  1.00 29.26      B    N
ATOM   1276  CA  LYS   243     -11.157 -16.492  20.582  1.00 29.82      B    C
ATOM   1277  CB  LYS   243     -11.445 -15.500  21.697  1.00 28.15      B    C
ATOM   1278  CG  LYS   243     -10.347 -14.472  21.869  1.00 26.26      B    C
ATOM   1279  CD  LYS   243      -9.612 -14.683  23.175  1.00 26.89      B    C
ATOM   1280  CE  LYS   243      -8.704 -13.506  23.508  1.00 26.93      B    C
ATOM   1281  NZ  LYS   243      -8.473 -13.398  24.973  1.00 26.14      B    N
ATOM   1282  C   LYS   243     -12.415 -16.735  19.747  1.00 31.25      B    C
ATOM   1283  O   LYS   243     -12.645 -16.054  18.750  1.00 33.02      B    O
ATOM   1284  N   GLY   244     -13.220 -17.718  20.123  1.00 31.12      B    N
ATOM   1285  CA  GLY   244     -14.451 -17.936  19.385  1.00 31.63      B    C
ATOM   1286  C   GLY   244     -14.296 -18.595  18.025  1.00 31.24      B    C
ATOM   1287  O   GLY   244     -15.258 -19.141  17.487  1.00 31.27      B    O
ATOM   1288  N   ALA   245     -13.100 -18.542  17.456  1.00 31.32      B    N
ATOM   1289  CA  ALA   245     -12.768 -19.446  16.366  1.00 32.05      B    C
ATOM   1290  CB  ALA   245     -11.446 -19.059  15.757  1.00 31.24      B    C
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ATOM   1291  C   ALA   245     -12.683 -20.847  16.956  1.00 32.80      B    C
ATOM   1292  O   ALA   245     -12.548 -21.012  18.170  1.00 33.98      B    O
ATOM   1293  N   SER   246     -12.764 -21.864  16.111  1.00 33.54      B    N
ATOM   1294  CA  SER   246     -12.791 -23.227  16.625  1.00 34.92      B    C
ATOM   1295  CB  SER   246     -14.207 -23.775  16.536  1.00 36.59      B    C
ATOM   1296  OG  SER   246     -15.062 -23.000  17.353  1.00 39.79      B    O
ATOM   1297  C   SER   246     -11.821 -24.180  15.951  1.00 34.43      B    C
ATOM   1298  O   SER   246     -11.525 -24.051  14.768  1.00 34.27      B    O
ATOM   1299  N   MET   247     -11.334 -25.145  16.719  1.00 34.89      B    N
ATOM   1300  CA  MET   247     -10.205 -25.947  16.292  1.00 35.66      B    C
ATOM   1301  CB  MET   247      -9.026 -25.702  17.237  1.00 38.44      B    C
ATOM   1302  CG  MET   247      -8.439 -24.296  17.103  1.00 41.75      B    C
ATOM   1303  SD  MET   247      -7.185 -23.868  18.326  1.00 45.85      B    S
ATOM   1304  CE  MET   247      -5.832 -24.890  17.796  1.00 44.65      B    C
ATOM   1305  C   MET   247     -10.499 -27.436  16.194  1.00 35.05      B    C
ATOM   1306  O   MET   247     -11.242 -28.001  16.995  1.00 34.54      B    O
ATOM   1307  N   ARG   248      -9.909 -28.066  15.190  1.00 34.28      B    N
ATOM   1308  CA  ARG   248      -9.950 -29.509  15.083  1.00 33.81      B    C
ATOM   1309  CB  ARG   248     -10.815 -29.914  13.897  1.00 36.62      B    C
ATOM   1310  CG  ARG   248     -11.884 -28.894  13.551  1.00 38.31      B    C
ATOM   1311  CD  ARG   248     -12.886 -29.472  12.566  1.00 40.68      B    C
ATOM   1312  NE  ARG   248     -13.589 -30.609  13.149  1.00 44.00      B    N
ATOM   1313  CZ  ARG   248     -14.713 -30.509  13.849  1.00 45.66      B    C
ATOM   1314  NH1 ARG   248     -15.285 -31.596  14.351  1.00 46.06      B    N
ATOM   1315  NH2 ARG   248     -15.268 -29.320  14.041  1.00 46.30      B    N
ATOM   1316  C   ARG   248      -8.526 -30.023  14.905  1.00 32.41      B    C
ATOM   1317  O   ARG   248      -7.850 -29.692  13.928  1.00 32.44      B    O
ATOM   1318  N   SER   249      -8.083 -30.836  15.857  1.00 29.90      B    N
ATOM   1319  CA  SER   249      -6.694 -31.247  15.930  1.00 27.61      B    C
ATOM   1320  CB  SER   249      -6.250 -31.200  17.393  1.00 27.24      B    C
ATOM   1321  OG  SER   249      -5.681 -32.427  17.811  1.00 27.75      B    O
ATOM   1322  C   SER   249      -6.420 -32.638  15.330  1.00 26.69      B    C
ATOM   1323  O   SER   249      -7.102 -33.608  15.649  1.00 27.35      B    O
ATOM   1324  N   LEU   250      -5.415 -32.727  14.462  1.00 25.20      B    N
ATOM   1325  CA  LEU   250      -4.862 -34.016  14.044  1.00 24.40      B    C
ATOM   1326  CB  LEU   250      -4.708 -34.070  12.521  1.00 25.99      B    C
ATOM   1327  CG  LEU   250      -5.964 -34.039  11.643  1.00 25.42      B    C
ATOM   1328  CD1 LEU   250      -6.634 -32.682  11.736  1.00 25.61      B    C
ATOM   1329  CD2 LEU   250      -5.574 -34.329  10.205  1.00 25.09      B    C
ATOM   1330  C   LEU   250      -3.493 -34.224  14.691  1.00 23.33      B    C
ATOM   1331  O   LEU   250      -2.813 -33.257  15.010  1.00 23.67      B    O
ATOM   1332  N   ARG   251      -3.091 -35.477  14.891  1.00 22.89      B    N
ATOM   1333  CA  ARG   251      -1.769 -35.763  15.438  1.00 22.61      B    C
ATOM   1334  CB  ARG   251      -1.859 -36.714  16.630  1.00 21.26      B    C
ATOM   1335  CG  ARG   251      -0.487 -37.028  17.209  1.00 23.82      B    C
ATOM   1336  CD  ARG   251      -0.542 -37.959  18.419  1.00 26.20      B    C
ATOM   1337  NE  ARG   251       0.709 -37.945  19.182  1.00 26.31      B    N
ATOM   1338  CZ  ARG   251       1.351 -39.036  19.585  1.00 25.27      B    C
ATOM   1339  NH1 ARG   251       0.863 -40.233  19.299  1.00 24.93      B    N
ATOM   1340  NH2 ARG   251       2.477 -38.931  20.275  1.00 25.73      B    N
ATOM   1341  C   ARG   251      -0.812 -36.355  14.406  1.00 23.38      B    C
ATOM   1342  O   ARG   251      -0.951 -37.511  13.993  1.00 23.07      B    O
ATOM   1343  N   VAL   252       0.174 -35.559  14.002  1.00 23.27      B    N
ATOM   1344  CA  VAL   252       1.136 -35.992  12.999  1.00 24.29      B    C
ATOM   1345  CB  VAL   252       1.134 -35.048  11.791  1.00 24.61      B    C
ATOM   1346  CG1 VAL   252      -0.182 -35.150  11.053  1.00 24.12      B    C
ATOM   1347  CG2 VAL   252       1.350 -33.633  12.261  1.00 25.97      B    C
ATOM   1348  C   VAL   252       2.546 -36.054  13.570  1.00 24.19      B    C
ATOM   1349  O   VAL   252       3.447 -36.634  12.959  1.00 25.34      B    O
ATOM   1350  N   LEU   253       2.720 -35.454  14.744  1.00 24.84      B    N
ATOM   1351  CA  LEU   253       4.006 -35.411  15.430  1.00 24.76      B    C
ATOM   1352  CB  LEU   253       4.361 -33.967  15.776  1.00 23.72      B    C
ATOM   1353  CG  LEU   253       4.361 -32.960  14.625  1.00 22.29      B    C
ATOM   1354  CD1 LEU   253       4.645 -31.576  15.154  1.00 21.59      B    C
ATOM   1355  CD2 LEU   253       5.408 -33.352  13.607  1.00 22.92      B    C
ATOM   1356  C   LEU   253       3.950 -36.233  16.714  1.00 26.34      B    C
ATOM   1357  O   LEU   253       3.063 -36.031  17.551  1.00 26.64      B    O
ATOM   1358  N   ASN   254       4.907 -37.147  16.870  1.00 27.39      B    N
ATOM   1359  CA  ASN   254       4.999 -37.981  18.066  1.00 27.84      B    C
ATOM   1360  CB  ASN   254       5.988 -39.131  17.832  1.00 28.42      B    C
ATOM   1361  CG  ASN   254       7.439 -38.676  17.836  1.00 28.34      B    C
ATOM   1362  OD1 ASN   254       7.727 -37.486  17.881  1.00 28.47      B    O
ATOM   1363  ND2 ASN   254       8.358 -39.630  17.788  1.00 28.22      B    N
ATOM   1364  C   ASN   254       5.414 -37.177  19.306  1.00 28.27      B    C
ATOM   1365  O   ASN   254       5.418 -35.938  19.295  1.00 28.25      B    O
ATOM   1366  N   CYS   255       5.756 -37.884  20.376  1.00 28.16      B    N
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ATOM   1367  CA  CYS   255       6.024 -37.239  21.655  1.00 28.63      B    C
ATOM   1368  C   CYS   255       7.325 -36.487  21.675  1.00 26.69      B    C
ATOM   1369  O   CYS   255       7.626 -35.806  22.645  1.00 27.25      B    O
ATOM   1370  CB  CYS   255       6.045 -38.258  22.781  1.00 32.30      B    C
ATOM   1371  SG  CYS   255       4.407 -38.942  23.137  1.00 43.56      B    S
ATOM   1372  N   GLN   256       8.111 -36.622  20.617  1.00 24.58      B    N
ATOM   1373  CA  GLN   256       9.348 -35.875  20.520  1.00 21.14      B    C
ATOM   1374  CB  GLN   256      10.503 -36.846  20.406  1.00 21.92      B    C
ATOM   1375  CG  GLN   256      10.613 -37.723  21.623  1.00 24.98      B    C
ATOM   1376  CD  GLN   256      11.095 -39.115  21.295  1.00 28.39      B    C
ATOM   1377  OE1 GLN   256      11.508 -39.398  20.165  1.00 30.69      B    O
ATOM   1378  NE2 GLN   256      11.045 -40.001  22.283  1.00 30.25      B    N
ATOM   1379  C   GLN   256       9.337 -34.893  19.357  1.00 20.02      B    C
ATOM   1380  O   GLN   256      10.385 -34.517  18.843  1.00 19.29      B    O
ATOM   1381  N   GLY   257       8.139 -34.479  18.950  1.00 19.22      B    N
ATOM   1382  CA  GLY   257       8.004 -33.430  17.957  1.00 19.26      B    C
ATOM   1383  C   GLY   257       8.241 -33.906  16.542  1.00 19.45      B    C
ATOM   1384  O   GLY   257       8.408 -33.104  15.619  1.00 18.96      B    O
ATOM   1385  N   LYS   258       8.248 -35.223  16.370  1.00 20.81      B    N
ATOM   1386  CA  LYS   258       8.682 -35.826  15.123  1.00 21.90      B    C
ATOM   1387  CB  LYS   258       9.762 -36.856  15.415  1.00 21.99      B    C
ATOM   1388  CG  LYS   258      10.399 -37.442  14.179  1.00 27.25      B    C
ATOM   1389  CD  LYS   258      11.905 -37.233  14.196  1.00 30.54      B    C
ATOM   1390  CE  LYS   258      12.624 -38.401  13.536  1.00 34.07      B    C
ATOM   1391  NZ  LYS   258      14.066 -38.101  13.269  1.00 37.21      B    N
ATOM   1392  C   LYS   258       7.536 -36.476  14.353  1.00 21.86      B    C
ATOM   1393  O   LYS   258       6.699 -37.167  14.923  1.00 23.60      B    O
ATOM   1394  N   GLY   259       7.504 -36.233  13.050  1.00 22.07      B    N
ATOM   1395  CA  GLY   259       6.571 -36.918  12.178  1.00 22.34      B    C
ATOM   1396  C   GLY   259       7.238 -37.276  10.862  1.00 22.86      B    C
ATOM   1397  O   GLY   259       8.467 -37.230  10.747  1.00 23.37      B    O
ATOM   1398  N   THR   260       6.435 -37.637   9.866  1.00 21.84      B    N
ATOM   1399  CA  THR   260       6.966 -37.958   8.548  1.00 21.86      B    C
ATOM   1400  CB  THR   260       6.869 -39.462   8.253  1.00 20.56      B    C
ATOM   1401  OG1 THR   260       5.496 -39.859   8.248  1.00 19.54      B    O
ATOM   1402  CG2 THR   260       7.622 -40.251   9.309  1.00 19.92      B    C
ATOM   1403  C   THR   260       6.203 -37.201   7.475  1.00 22.65      B    C
ATOM   1404  O   THR   260       5.080 -36.777   7.704  1.00 24.48      B    O
ATOM   1405  N   VAL   261       6.811 -37.024   6.306  1.00 22.69      B    N
ATOM   1406  CA  VAL   261       6.173 -36.262   5.247  1.00 21.49      B    C
ATOM   1407  CB  VAL   261       7.091 -36.153   4.018  1.00 20.59      B    C
ATOM   1408  CG1 VAL   261       6.402 -35.364   2.919  1.00 19.43      B    C
ATOM   1409  CG2 VAL   261       8.388 -35.473   4.409  1.00 19.44      B    C
ATOM   1410  C   VAL   261       4.872 -36.951   4.860  1.00 21.94      B    C
ATOM   1411  O   VAL   261       3.850 -36.301   4.624  1.00 22.12      B    O
ATOM   1412  N   SER   262       4.911 -38.278   4.830  1.00 22.53      B    N
ATOM   1413  CA  SER   262       3.757 -39.076   4.428  1.00 22.26      B    C
ATOM   1414  CB  SER   262       4.165 -40.541   4.284  1.00 22.47      B    C
ATOM   1415  OG  SER   262       4.955 -40.941   5.390  1.00 25.99      B    O
ATOM   1416  C   SER   262       2.605 -38.962   5.414  1.00 21.68      B    C
ATOM   1417  O   SER   262       1.457 -38.779   5.009  1.00 22.09      B    O
ATOM   1418  N   GLY   263       2.916 -39.076   6.702  1.00 21.45      B    N
ATOM   1419  CA  GLY   263       1.909 -38.867   7.724  1.00 21.21      B    C
ATOM   1420  C   GLY   263       1.256 -37.506   7.567  1.00 22.43      B    C
ATOM   1421  O   GLY   263       0.063 -37.328   7.845  1.00 21.36      B    O
ATOM   1422  N   THR   264       2.039 -36.538   7.103  1.00 22.11      B    N
ATOM   1423  CA  THR   264       1.527 -35.202   6.887  1.00 21.72      B    C
ATOM   1424  CB  THR   264       2.644 -34.228   6.597  1.00 21.29      B    C
ATOM   1425  OG1 THR   264       3.532 -34.176   7.715  1.00 20.27      B    O
ATOM   1426  CG2 THR   264       2.074 -32.850   6.348  1.00 21.16      B    C
ATOM   1427  C   THR   264       0.585 -35.208   5.701  1.00 23.10      B    C
ATOM   1428  O   THR   264      -0.547 -34.732   5.796  1.00 25.02      B    O
ATOM   1429  N   LEU   265       1.058 -35.753   4.584  1.00 23.29      B    N
ATOM   1430  CA  LEU   265       0.233 -35.904   3.388  1.00 21.86      B    C
ATOM   1431  CB  LEU   265       0.904 -36.860   2.412  1.00 20.38      B    C
ATOM   1432  CG  LEU   265       2.200 -36.340   1.805  1.00 20.98      B    C
ATOM   1433  CD1 LEU   265       2.879 -37.457   1.015  1.00 20.52      B    C
ATOM   1434  CD2 LEU   265       1.898 -35.139   0.919  1.00 19.35      B    C
ATOM   1435  C   LEU   265      -1.135 -36.455   3.745  1.00 21.27      B    C
ATOM   1436  O   LEU   265      -2.159 -35.900   3.361  1.00 23.43      B    O
ATOM   1437  N   ILE   266      -1.136 -37.555   4.485  1.00 19.76      B    N
ATOM   1438  CA  ILE   266      -2.355 -38.257   4.838  1.00 18.31      B    C
ATOM   1439  CB  ILE   266      -2.002 -39.585   5.543  1.00 16.80      B    C
ATOM   1440  CG2 ILE   266      -3.251 -40.323   5.957  1.00 13.78      B    C
ATOM   1441  CG1 ILE   266      -1.151 -40.436   4.603  1.00 14.04      B    C
ATOM   1442  CD1 ILE   266      -0.945 -41.830   5.071  1.00 14.81      B    C
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ATOM   1443  C   ILE   266      -3.219 -37.386   5.746  1.00 20.40      B    C
ATOM   1444  O   ILE   266      -4.454 -37.447   5.703  1.00 20.07      B    O
ATOM   1445  N   GLY   267      -2.560 -36.566   6.560  1.00 21.94      B    N
ATOM   1446  CA  GLY   267      -3.283 -35.624   7.393  1.00 21.97      B    C
ATOM   1447  C   GLY   267      -4.098 -34.693   6.520  1.00 22.46      B    C
ATOM   1448  O   GLY   267      -5.301 -34.527   6.717  1.00 23.67      B    O
ATOM   1449  N   LEU   268      -3.444 -34.092   5.538  1.00 21.50      B    N
ATOM   1450  CA  LEU   268      -4.124 -33.185   4.639  1.00 21.24      B    C
ATOM   1451  CB  LEU   268      -3.108 -32.532   3.697  1.00 21.08      B    C
ATOM   1452  CG  LEU   268      -1.978 -31.721   4.354  1.00 20.19      B    C
ATOM   1453  CD1 LEU   268      -1.029 -31.208   3.291  1.00 20.01      B    C
ATOM   1454  CD2 LEU   268      -2.550 -30.556   5.130  1.00 19.67      B    C
ATOM   1455  C   LEU   268      -5.214 -33.910   3.841  1.00 22.54      B    C
ATOM   1456  O   LEU   268      -6.269 -33.333   3.558  1.00 24.32      B    O
ATOM   1457  N   GLU   269      -4.983 -35.172   3.488  1.00 21.86      B    N
ATOM   1458  CA  GLU   269      -6.009 -35.930   2.778  1.00 21.18      B    C
ATOM   1459  CB  GLU   269      -5.553 -37.367   2.519  1.00 22.31      B    C
ATOM   1460  CG  GLU   269      -6.368 -38.078   1.431  1.00 24.62      B    C
ATOM   1461  CD  GLU   269      -6.354 -39.593   1.562  1.00 25.91      B    C
ATOM   1462  OE1 GLU   269      -5.320 -40.154   1.966  1.00 27.02      B    O
ATOM   1463  OE2 GLU   269      -7.382 -40.229   1.257  1.00 27.61      B    O
ATOM   1464  C   GLU   269      -7.250 -35.957   3.650  1.00 20.82      B    C
ATOM   1465  O   GLU   269      -8.365 -35.650   3.209  1.00 21.02      B    O
ATOM   1466  N   PHE   270      -7.032 -36.321   4.906  1.00 19.63      B    N
ATOM   1467  CA  PHE   270      -8.100 -36.429   5.881  1.00 17.51      B    C
ATOM   1468  CB  PHE   270      -7.495 -36.822   7.220  1.00 16.07      B    C
ATOM   1469  CG  PHE   270      -8.467 -36.847   8.342  1.00 15.79      B    C
ATOM   1470  CD1 PHE   270      -8.923 -38.046   8.844  1.00 16.82      B    C
ATOM   1471  CD2 PHE   270      -8.884 -35.672   8.940  1.00 16.62      B    C
ATOM   1472  CE1 PHE   270      -9.774 -38.081   9.933  1.00 17.22      B    C
ATOM   1473  CE2 PHE   270      -9.728 -35.697  10.023  1.00 17.59      B    C
ATOM   1474  CZ  PHE   270     -10.176 -36.910  10.523  1.00 18.31      B    C
ATOM   1475  C   PHE   270      -8.860 -35.114   5.997  1.00 16.78      B    C
ATOM   1476  O   PHE   270     -10.089 -35.096   6.065  1.00 17.12      B    O
ATOM   1477  N   ILE   271      -8.125 -34.012   6.008  1.00 15.76      B    N
ATOM   1478  CA  ILE   271      -8.743 -32.713   6.185  1.00 16.00      B    C
ATOM   1479  CB  ILE   271      -7.686 -31.603   6.253  1.00 16.26      B    C
ATOM   1480  CG2 ILE   271      -8.346 -30.272   6.509  1.00 15.52      B    C
ATOM   1481  CG1 ILE   271      -6.715 -31.875   7.390  1.00 16.87      B    C
ATOM   1482  CD1 ILE   271      -5.899 -30.666   7.767  1.00 16.57      B    C
ATOM   1483  C   ILE   271      -9.697 -32.409   5.039  1.00 16.31      B    C
ATOM   1484  O   ILE   271     -10.801 -31.905   5.247  1.00 18.27      B    O
ATOM   1485  N   ARG   272      -9.276 -32.722   3.825  1.00 15.58      B    N
ATOM   1486  CA  ARG   272     -10.058 -32.368   2.657  1.00 14.76      B    C
ATOM   1487  CB  ARG   272      -9.167 -32.521   1.425  1.00 13.53      B    C
ATOM   1488  CG  ARG   272      -9.870 -32.459   0.112  1.00 14.31      B    C
ATOM   1489  CD  ARG   272     -10.647 -31.195  -0.054  1.00 14.45      B    C
ATOM   1490  NE  ARG   272     -10.958 -30.998  -1.464  1.00 18.03      B    N
ATOM   1491  CZ  ARG   272     -11.694 -30.000  -1.939  1.00 17.53      B    C
ATOM   1492  NH1 ARG   272     -11.920 -29.895  -3.237  1.00 18.77      B    N
ATOM   1493  NH2 ARG   272     -12.217 -29.114  -1.117  1.00 17.11      B    N
ATOM   1494  C   ARG   272     -11.332 -33.229   2.569  1.00 14.98      B    C
ATOM   1495  O   ARG   272     -12.411 -32.735   2.240  1.00 13.74      B    O
ATOM   1496  N   LYS   273     -11.213 -34.511   2.898  1.00 16.77      B    N
ATOM   1497  CA  LYS   273     -12.380 -35.393   2.914  1.00 19.15      B    C
ATOM   1498  CB  LYS   273     -11.971 -36.834   3.234  1.00 19.12      B    C
ATOM   1499  CG  LYS   273     -13.115 -37.702   3.711  1.00 21.09      B    C
ATOM   1500  CD  LYS   273     -13.108 -37.797   5.246  1.00 28.83      B    C
ATOM   1501  CE  LYS   273     -14.534 -37.717   5.868  1.00 32.67      B    C
ATOM   1502  NZ  LYS   273     -14.605 -37.111   7.263  1.00 30.13      B    N
ATOM   1503  C   LYS   273     -13.424 -34.937   3.924  1.00 20.09      B    C
ATOM   1504  O   LYS   273     -14.622 -34.957   3.639  1.00 20.19      B    O
ATOM   1505  N   SER   274     -12.970 -34.528   5.104  1.00 20.81      B    N
ATOM   1506  CA  SER   274     -13.879 -34.101   6.166  1.00 21.13      B    C
ATOM   1507  CB  SER   274     -13.089 -33.634   7.385  1.00 20.17      B    C
ATOM   1508  OG  SER   274     -12.088 -34.580   7.707  1.00 21.58      B    O
ATOM   1509  C   SER   274     -14.713 -32.956   5.648  1.00 20.96      B    C
ATOM   1510  O   SER   274     -15.902 -32.838   5.927  1.00 22.15      B    O
ATOM   1511  N   GLN   275     -14.068 -32.112   4.867  1.00 20.71      B    N
ATOM   1512  CA  GLN   275     -14.668 -30.868   4.471  1.00 20.82      B    C
ATOM   1513  CB  GLN   275     -13.566 -29.907   4.085  1.00 19.43      B    C
ATOM   1514  CG  GLN   275     -14.035 -28.600   3.568  1.00 18.84      B    C
ATOM   1515  CD  GLN   275     -13.215 -28.212   2.389  1.00 19.76      B    C
ATOM   1516  OE1 GLN   275     -12.708 -29.082   1.681  1.00 22.34      B    O
ATOM   1517  NE2 GLN   275     -13.057 -26.918   2.163  1.00 18.84      B    N
ATOM   1518  C   GLN   275     -15.632 -31.078   3.320  1.00 21.36      B    C
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ATOM	1519	O	GLN	275	-16.578	-30.317	3.144	1.00	21.85	B	O
ATOM	1520	N	LEU	276	-15.393	-32.124	2.542	1.00	21.35	B	N
ATOM	1521	CA	LEU	276	-16.293	-32.468	1.458	1.00	21.46	B	C
ATOM	1522	CB	LEU	276	-15.569	-33.370	0.468	1.00	20.85	B	C
ATOM	1523	CG	LEU	276	-14.412	-32.655	-0.222	1.00	21.54	B	C
ATOM	1524	CD1	LEU	276	-13.352	-33.659	-0.648	1.00	21.26	B	C
ATOM	1525	CD2	LEU	276	-14.946	-31.877	-1.410	1.00	18.00	B	C
ATOM	1526	C	LEU	276	-17.540	-33.162	1.994	1.00	22.78	B	C
ATOM	1527	O	LEU	276	-18.622	-33.061	1.406	1.00	23.18	B	O
ATOM	1528	N	VAL	277	-17.378	-33.855	3.119	1.00	23.69	B	N
ATOM	1529	CA	VAL	277	-18.449	-34.645	3.719	1.00	23.85	B	C
ATOM	1530	CB	VAL	277	-17.867	-35.768	4.570	1.00	21.82	B	C
ATOM	1531	CG1	VAL	277	-18.977	-36.622	5.132	1.00	21.54	B	C
ATOM	1532	CG2	VAL	277	-16.941	-36.597	3.735	1.00	22.34	B	C
ATOM	1533	C	VAL	277	-19.370	-33.801	4.596	1.00	25.25	B	C
ATOM	1534	O	VAL	277	-20.541	-34.136	4.804	1.00	25.41	B	O
ATOM	1535	N	GLN	278	-18.826	-32.707	5.115	1.00	26.20	B	N
ATOM	1536	CA	GLN	278	-19.583	-31.798	5.960	1.00	26.06	B	C
ATOM	1537	CB	GLN	278	-19.440	-32.202	7.414	1.00	28.61	B	C
ATOM	1538	CG	GLN	278	-20.458	-33.207	7.843	1.00	35.11	B	C
ATOM	1539	CD	GLN	278	-21.228	-32.729	9.047	1.00	39.20	B	C
ATOM	1540	OE1	GLN	278	-20.635	-32.271	10.033	1.00	40.40	B	O
ATOM	1541	NE2	GLN	278	-22.562	-32.821	8.980	1.00	41.21	B	N
ATOM	1542	C	GLN	278	-19.108	-30.372	5.791	1.00	25.07	B	C
ATOM	1543	O	GLN	278	-18.602	-29.760	6.729	1.00	25.51	B	O
ATOM	1544	N	PRO	279	-19.272	-29.817	4.586	1.00	24.43	B	N
ATOM	1545	CD	PRO	279	-20.021	-30.374	3.449	1.00	22.73	B	C
ATOM	1546	CA	PRO	279	-18.773	-28.467	4.320	1.00	23.37	B	C
ATOM	1547	CB	PRO	279	-19.302	-28.148	2.922	1.00	21.77	B	C
ATOM	1548	CG	PRO	279	-20.342	-29.170	2.644	1.00	22.09	B	C
ATOM	1549	C	PRO	279	-19.239	-27.470	5.362	1.00	22.65	B	C
ATOM	1550	O	PRO	279	-20.374	-27.525	5.820	1.00	23.20	B	O
ATOM	1551	N	VAL	280	-18.336	-26.571	5.742	1.00	23.67	B	N
ATOM	1552	CA	VAL	280	-18.598	-25.562	6.768	1.00	22.67	B	C
ATOM	1553	CB	VAL	280	-17.728	-25.800	8.019	1.00	22.17	B	C
ATOM	1554	CG1	VAL	280	-18.175	-27.056	8.729	1.00	20.68	B	C
ATOM	1555	CG2	VAL	280	-16.267	-25.934	7.607	1.00	23.06	B	C
ATOM	1556	C	VAL	280	-18.283	-24.179	6.220	1.00	22.17	B	C
ATOM	1557	O	VAL	280	-18.635	-23.858	5.088	1.00	20.16	B	O
ATOM	1558	N	GLY	281	-17.613	-23.366	7.032	1.00	23.75	B	N
ATOM	1559	CA	GLY	281	-17.163	-22.064	6.570	1.00	24.31	B	C
ATOM	1560	C	GLY	281	-15.768	-22.144	5.984	1.00	24.09	B	C
ATOM	1561	O	GLY	281	-15.317	-23.229	5.636	1.00	24.64	B	O
ATOM	1562	N	PRO	282	-15.059	-21.013	5.865	1.00	24.04	B	N
ATOM	1563	CD	PRO	282	-15.556	-19.651	6.123	1.00	24.55	B	C
ATOM	1564	CA	PRO	282	-13.654	-21.006	5.451	1.00	24.01	B	C
ATOM	1565	CB	PRO	282	-13.274	-19.527	5.493	1.00	24.10	B	C
ATOM	1566	CG	PRO	282	-14.564	-18.795	5.411	1.00	23.06	B	C
ATOM	1567	C	PRO	282	-12.796	-21.839	6.409	1.00	24.43	B	C
ATOM	1568	O	PRO	282	-13.047	-21.856	7.617	1.00	24.10	B	O
ATOM	1569	N	LEU	283	-11.792	-22.528	5.863	1.00	24.32	B	N
ATOM	1570	CA	LEU	283	-10.926	-23.400	6.651	1.00	23.30	B	C
ATOM	1571	CB	LEU	283	-10.884	-24.802	6.057	1.00	22.99	B	C
ATOM	1572	CG	LEU	283	-12.166	-25.615	6.083	1.00	23.79	B	C
ATOM	1573	CD1	LEU	283	-11.845	-27.055	5.708	1.00	24.80	B	C
ATOM	1574	CD2	LEU	283	-12.784	-25.549	7.465	1.00	25.21	B	C
ATOM	1575	C	LEU	283	-9.512	-22.865	6.687	1.00	22.69	B	C
ATOM	1576	O	LEU	283	-8.999	-22.402	5.679	1.00	23.60	B	O
ATOM	1577	N	VAL	284	-8.881	-22.944	7.851	1.00	22.49	B	N
ATOM	1578	CA	VAL	284	-7.471	-22.605	7.973	1.00	21.61	B	C
ATOM	1579	CB	VAL	284	-7.271	-21.405	8.918	1.00	21.87	B	C
ATOM	1580	CG1	VAL	284	-5.794	-21.231	9.247	1.00	20.77	B	C
ATOM	1581	CG2	VAL	284	-7.825	-20.147	8.268	1.00	20.07	B	C
ATOM	1582	C	VAL	284	-6.679	-23.794	8.496	1.00	20.93	B	C
ATOM	1583	O	VAL	284	-6.957	-24.315	9.573	1.00	20.79	B	O
ATOM	1584	N	VAL	285	-5.696	-24.231	7.718	1.00	21.45	B	N
ATOM	1585	CA	VAL	285	-4.803	-25.283	8.175	1.00	21.67	B	C
ATOM	1586	CB	VAL	285	-4.470	-26.269	7.049	1.00	23.02	B	C
ATOM	1587	CG1	VAL	285	-3.716	-27.470	7.613	1.00	22.15	B	C
ATOM	1588	CG2	VAL	285	-5.747	-26.714	6.363	1.00	24.14	B	C
ATOM	1589	C	VAL	285	-3.506	-24.674	8.687	1.00	20.61	B	C
ATOM	1590	O	VAL	285	-2.849	-23.902	7.986	1.00	18.40	B	O
ATOM	1591	N	LEU	286	-3.161	-25.023	9.924	1.00	20.05	B	N
ATOM	1592	CA	LEU	286	-1.912	-24.605	10.535	1.00	20.46	B	C
ATOM	1593	CB	LEU	286	-2.150	-24.169	11.973	1.00	21.88	B	C
ATOM	1594	CG	LEU	286	-0.864	-23.852	12.738	1.00	22.20	B	C

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ATOM   1595  CD1 LEU   286      -0.189 -22.614  12.136  1.00 20.76      B    C
ATOM   1596  CD2 LEU   286      -1.201 -23.638  14.196  1.00 21.62      B    C
ATOM   1597  C   LEU   286      -0.892 -25.734  10.532  1.00 20.34      B    C
ATOM   1598  O   LEU   286      -1.162 -26.824  11.035  1.00 19.43      B    O
ATOM   1599  N   LEU   287       0.286 -25.458   9.978  1.00 21.20      B    N
ATOM   1600  CA  LEU   287       1.346 -26.463   9.881  1.00 21.06      B    C
ATOM   1601  CB  LEU   287       1.646 -26.744   8.406  1.00 20.12      B    C
ATOM   1602  CG  LEU   287       0.453 -27.283   7.608  1.00 17.86      B    C
ATOM   1603  CD1 LEU   287       0.647 -27.013   6.144  1.00 16.35      B    C
ATOM   1604  CD2 LEU   287       0.292 -28.768   7.864  1.00 17.11      B    C
ATOM   1605  C   LEU   287       2.616 -26.007  10.612  1.00 20.61      B    C
ATOM   1606  O   LEU   287       3.545 -25.471  10.001  1.00 20.04      B    O
ATOM   1607  N   PRO   288       2.657 -26.211  11.941  1.00 20.31      B    N
ATOM   1608  CD  PRO   288       1.636 -26.978  12.681  1.00 19.79      B    C
ATOM   1609  CA  PRO   288       3.741 -25.767  12.823  1.00 19.59      B    C
ATOM   1610  CB  PRO   288       3.062 -25.699  14.187  1.00 19.77      B    C
ATOM   1611  CG  PRO   288       2.020 -26.783  14.126  1.00 19.23      B    C
ATOM   1612  C   PRO   288       4.925 -26.736  12.786  1.00 18.73      B    C
ATOM   1613  O   PRO   288       5.436 -27.179  13.829  1.00 17.76      B    O
ATOM   1614  N   LEU   289       5.341 -27.069  11.565  1.00 17.09      B    N
ATOM   1615  CA  LEU   289       6.372 -28.076  11.355  1.00 16.76      B    C
ATOM   1616  CB  LEU   289       5.741 -29.470  11.230  1.00 15.59      B    C
ATOM   1617  CG  LEU   289       4.678 -29.630  10.138  1.00 16.71      B    C
ATOM   1618  CD1 LEU   289       5.301 -30.269   8.897  1.00 17.39      B    C
ATOM   1619  CD2 LEU   289       3.525 -30.474  10.662  1.00 14.29      B    C
ATOM   1620  C   LEU   289       7.200 -27.766  10.120  1.00 16.66      B    C
ATOM   1621  O   LEU   289       6.795 -26.960   9.267  1.00 15.85      B    O
ATOM   1622  N   ALA   290       8.371 -28.399  10.046  1.00 17.02      B    N
ATOM   1623  CA  ALA   290       9.266 -28.245   8.910  1.00 17.79      B    C
ATOM   1624  CB  ALA   290      10.198 -27.063   9.127  1.00 16.37      B    C
ATOM   1625  C   ALA   290      10.080 -29.507   8.714  1.00 19.12      B    C
ATOM   1626  O   ALA   290      10.594 -30.089   9.682  1.00 18.31      B    O
ATOM   1627  N   GLY   291      10.181 -29.920   7.452  1.00 20.09      B    N
ATOM   1628  CA  GLY   291      11.138 -30.934   7.050  1.00 20.48      B    C
ATOM   1629  C   GLY   291      11.980 -30.388   5.915  1.00 20.72      B    C
ATOM   1630  O   GLY   291      11.786 -29.242   5.500  1.00 21.90      B    O
ATOM   1631  N   GLY   292      12.919 -31.186   5.412  1.00 20.93      B    N
ATOM   1632  CA  GLY   292      13.737 -30.734   4.299  1.00 20.49      B    C
ATOM   1633  C   GLY   292      12.862 -30.506   3.088  1.00 20.99      B    C
ATOM   1634  O   GLY   292      11.752 -31.020   3.027  1.00 22.41      B    O
ATOM   1635  N   TYR   293      13.340 -29.744   2.116  1.00 22.10      B    N
ATOM   1636  CA  TYR   293      12.499 -29.422   0.969  1.00 21.97      B    C
ATOM   1637  CB  TYR   293      13.293 -28.690  -0.106  1.00 22.23      B    C
ATOM   1638  CG  TYR   293      12.489 -28.396  -1.344  1.00 22.52      B    C
ATOM   1639  CD1 TYR   293      11.931 -27.137  -1.554  1.00 23.24      B    C
ATOM   1640  CE1 TYR   293      11.215 -26.853  -2.706  1.00 23.07      B    C
ATOM   1641  CD2 TYR   293      12.304 -29.367  -2.318  1.00 22.54      B    C
ATOM   1642  CE2 TYR   293      11.589 -29.095  -3.470  1.00 23.87      B    C
ATOM   1643  CZ  TYR   293      11.051 -27.839  -3.662  1.00 24.37      B    C
ATOM   1644  OH  TYR   293      10.373 -27.571  -4.829  1.00 24.90      B    O
ATOM   1645  C   TYR   293      11.897 -30.681   0.373  1.00 21.19      B    C
ATOM   1646  O   TYR   293      12.579 -31.688   0.186  1.00 21.55      B    O
ATOM   1647  N   SER   294      10.607 -30.608   0.078  1.00 20.40      B    N
ATOM   1648  CA  SER   294       9.865 -31.758  -0.397  1.00 19.74      B    C
ATOM   1649  CB  SER   294       8.982 -32.285   0.724  1.00 18.39      B    C
ATOM   1650  OG  SER   294       8.273 -33.436   0.313  1.00 19.04      B    O
ATOM   1651  C   SER   294       9.012 -31.412  -1.619  1.00 19.99      B    C
ATOM   1652  O   SER   294       8.284 -30.422  -1.625  1.00 19.62      B    O
ATOM   1653  N   ARG   295       9.110 -32.236  -2.658  1.00 20.06      B    N
ATOM   1654  CA  ARG   295       8.328 -32.012  -3.859  1.00 18.83      B    C
ATOM   1655  CB  ARG   295       8.908 -32.790  -5.033  1.00 16.98      B    C
ATOM   1656  CG  ARG   295       7.854 -33.248  -5.999  1.00 18.85      B    C
ATOM   1657  CD  ARG   295       8.040 -32.668  -7.377  1.00 20.56      B    C
ATOM   1658  NE  ARG   295       8.627 -33.644  -8.290  1.00 22.09      B    N
ATOM   1659  CZ  ARG   295       8.278 -33.780  -9.567  1.00 22.80      B    C
ATOM   1660  NH1 ARG   295       7.331 -33.023 -10.103  1.00 20.61      B    N
ATOM   1661  NH2 ARG   295       8.889 -34.707 -10.316  1.00 24.02      B    N
ATOM   1662  C   ARG   295       6.903 -32.457  -3.608  1.00 18.54      B    C
ATOM   1663  O   ARG   295       5.956 -31.735  -3.908  1.00 17.83      B    O
ATOM   1664  N   VAL   296       6.754 -33.652  -3.053  1.00 19.37      B    N
ATOM   1665  CA  VAL   296       5.425 -34.189  -2.812  1.00 19.78      B    C
ATOM   1666  CB  VAL   296       5.459 -35.674  -2.372  1.00 18.32      B    C
ATOM   1667  CG1 VAL   296       6.315 -35.852  -1.154  1.00 16.29      B    C
ATOM   1668  CG2 VAL   296       4.061 -36.142  -2.074  1.00 16.25      B    C
ATOM   1669  C   VAL   296       4.710 -33.375  -1.749  1.00 20.69      B    C
ATOM   1670  O   VAL   296       3.544 -33.021  -1.917  1.00 23.04      B    O
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ATOM   1671  N   LEU   297       5.405 -33.055  -0.664  1.00 20.71      B    N
ATOM   1672  CA  LEU   297       4.784 -32.286   0.408  1.00 20.16      B    C
ATOM   1673  CB  LEU   297       5.766 -32.061   1.561  1.00 19.83      B    C
ATOM   1674  CG  LEU   297       5.115 -31.355   2.748  1.00 19.79      B    C
ATOM   1675  CD1 LEU   297       3.837 -32.103   3.088  1.00 19.32      B    C
ATOM   1676  CD2 LEU   297       6.048 -31.300   3.942  1.00 17.12      B    C
ATOM   1677  C   LEU   297       4.330 -30.945  -0.133  1.00 19.04      B    C
ATOM   1678  O   LEU   297       3.187 -30.549   0.051  1.00 18.61      B    O
ATOM   1679  N   ASN   298       5.237 -30.258  -0.814  1.00 19.03      B    N
ATOM   1680  CA  ASN   298       4.924 -28.982  -1.438  1.00 19.13      B    C
ATOM   1681  CB  ASN   298       6.132 -28.481  -2.233  1.00 19.01      B    C
ATOM   1682  CG  ASN   298       7.070 -27.627  -1.397  1.00 20.79      B    C
ATOM   1683  OD1 ASN   298       7.070 -27.694  -0.161  1.00 19.14      B    O
ATOM   1684  ND2 ASN   298       7.878 -26.813  -2.072  1.00 21.14      B    N
ATOM   1685  C   ASN   298       3.712 -29.055  -2.358  1.00 19.17      B    C
ATOM   1686  O   ASN   298       2.981 -28.080  -2.502  1.00 20.60      B    O
ATOM   1687  N   ALA   299       3.512 -30.209  -2.991  1.00 20.42      B    N
ATOM   1688  CA  ALA   299       2.475 -30.370  -4.010  1.00 19.62      B    C
ATOM   1689  CB  ALA   299       2.808 -31.555  -4.892  1.00 18.75      B    C
ATOM   1690  C   ALA   299       1.113 -30.575  -3.350  1.00 20.32      B    C
ATOM   1691  O   ALA   299       0.110 -29.965  -3.740  1.00 18.83      B    O
ATOM   1692  N   ALA   300       1.090 -31.443  -2.347  1.00 19.46      B    N
ATOM   1693  CA  ALA   300      -0.084 -31.611  -1.521  1.00 19.11      B    C
ATOM   1694  CB  ALA   300       0.283 -32.382  -0.286  1.00 16.69      B    C
ATOM   1695  C   ALA   300      -0.622 -30.239  -1.140  1.00 20.79      B    C
ATOM   1696  O   ALA   300      -1.786 -29.918  -1.387  1.00 21.71      B    O
ATOM   1697  N   CYS   301       0.249 -29.426  -0.551  1.00 22.30      B    N
ATOM   1698  CA  CYS   301      -0.120 -28.099  -0.080  1.00 22.76      B    C
ATOM   1699  CB  CYS   301       1.070 -27.416   0.582  1.00 23.52      B    C
ATOM   1700  SG  CYS   301       1.582 -28.239   2.077  1.00 26.02      B    S
ATOM   1701  C   CYS   301      -0.618 -27.234  -1.200  1.00 22.64      B    C
ATOM   1702  O   CYS   301      -1.618 -26.556  -1.052  1.00 24.93      B    O
ATOM   1703  N   GLN   302       0.077 -27.245  -2.323  1.00 23.10      B    N
ATOM   1704  CA  GLN   302      -0.368 -26.448  -3.442  1.00 24.66      B    C
ATOM   1705  CB  GLN   302       0.572 -26.621  -4.615  1.00 26.73      B    C
ATOM   1706  CG  GLN   302       0.020 -26.012  -5.874  1.00 32.13      B    C
ATOM   1707  CD  GLN   302       0.813 -26.408  -7.091  1.00 36.06      B    C
ATOM   1708  OE1 GLN   302       0.632 -25.850  -8.176  1.00 37.94      B    O
ATOM   1709  NE2 GLN   302       1.708 -27.382  -6.921  1.00 39.51      B    N
ATOM   1710  C   GLN   302      -1.784 -26.816  -3.881  1.00 24.41      B    C
ATOM   1711  O   GLN   302      -2.604 -25.939  -4.146  1.00 25.13      B    O
ATOM   1712  N   ARG   303      -2.075 -28.109  -3.960  1.00 23.65      B    N
ATOM   1713  CA  ARG   303      -3.357 -28.553  -4.490  1.00 21.86      B    C
ATOM   1714  CB  ARG   303      -3.414 -30.069  -4.570  1.00 21.24      B    C
ATOM   1715  CG  ARG   303      -2.200 -30.679  -5.168  1.00 22.97      B    C
ATOM   1716  CD  ARG   303      -2.556 -31.516  -6.363  1.00 23.46      B    C
ATOM   1717  NE  ARG   303      -1.353 -31.932  -7.067  1.00 25.29      B    N
ATOM   1718  CZ  ARG   303      -1.188 -33.120  -7.632  1.00 26.86      B    C
ATOM   1719  NH1 ARG   303      -0.048 -33.399  -8.251  1.00 26.86      B    N
ATOM   1720  NH2 ARG   303      -2.158 -34.025  -7.578  1.00 27.68      B    N
ATOM   1721  C   ARG   303      -4.448 -28.094  -3.561  1.00 21.46      B    C
ATOM   1722  O   ARG   303      -5.502 -27.632  -3.994  1.00 22.14      B    O
ATOM   1723  N   LEU   304      -4.184 -28.247  -2.273  1.00 20.16      B    N
ATOM   1724  CA  LEU   304      -5.166 -27.946  -1.250  1.00 18.95      B    C
ATOM   1725  CB  LEU   304      -4.613 -28.345   0.113  1.00 17.05      B    C
ATOM   1726  CG  LEU   304      -5.650 -28.360   1.225  1.00 17.78      B    C
ATOM   1727  CD1 LEU   304      -6.880 -29.111   0.750  1.00 18.88      B    C
ATOM   1728  CD2 LEU   304      -5.056 -29.006   2.460  1.00 17.39      B    C
ATOM   1729  C   LEU   304      -5.498 -26.458  -1.262  1.00 17.86      B    C
ATOM   1730  O   LEU   304      -6.647 -26.062  -1.060  1.00 16.37      B    O
ATOM   1731  N   ALA   305      -4.474 -25.645  -1.503  1.00 18.14      B    N
ATOM   1732  CA  ALA   305      -4.595 -24.196  -1.499  1.00 18.65      B    C
ATOM   1733  CB  ALA   305      -3.223 -23.568  -1.606  1.00 18.26      B    C
ATOM   1734  C   ALA   305      -5.452 -23.755  -2.664  1.00 19.54      B    C
ATOM   1735  O   ALA   305      -6.277 -22.854  -2.537  1.00 19.94      B    O
ATOM   1736  N   ARG   306      -5.238 -24.399  -3.803  1.00 20.90      B    N
ATOM   1737  CA  ARG   306      -6.015 -24.122  -4.996  1.00 23.22      B    C
ATOM   1738  CB  ARG   306      -5.258 -24.629  -6.217  1.00 26.07      B    C
ATOM   1739  CG  ARG   306      -3.873 -24.036  -6.332  1.00 32.51      B    C
ATOM   1740  CD  ARG   306      -2.949 -24.978  -7.079  1.00 40.27      B    C
ATOM   1741  NE  ARG   306      -2.167 -24.284  -8.100  1.00 46.05      B    N
ATOM   1742  CZ  ARG   306      -1.806 -24.827  -9.261  1.00 48.83      B    C
ATOM   1743  NH1 ARG   306      -1.094 -24.120 -10.133  1.00 50.33      B    N
ATOM   1744  NH2 ARG   306      -2.157 -26.076  -9.552  1.00 49.46      B    N
ATOM   1745  C   ARG   306      -7.400 -24.764  -4.923  1.00 21.55      B    C
ATOM   1746  O   ARG   306      -8.291 -24.445  -5.710  1.00 22.43      B    O
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ATOM   1747  N   ALA   307      -7.581 -25.668  -3.974  1.00 19.75       B    N
ATOM   1748  CA  ALA   307      -8.861 -26.329  -3.807  1.00 18.30       B    C
ATOM   1749  CB  ALA   307      -8.662 -27.684  -3.143  1.00 18.38       B    C
ATOM   1750  C   ALA   307      -9.737 -25.435  -2.946  1.00 18.11       B    C
ATOM   1751  O   ALA   307     -10.925 -25.697  -2.752  1.00 18.26       B    O
ATOM   1752  N   GLY   308      -9.131 -24.373  -2.427  1.00 18.07       B    N
ATOM   1753  CA  GLY   308      -9.883 -23.397  -1.670  1.00 17.27       B    C
ATOM   1754  C   GLY   308      -9.464 -23.190  -0.224  1.00 17.47       B    C
ATOM   1755  O   GLY   308      -9.854 -22.199   0.371  1.00 18.26       B    O
ATOM   1756  N   VAL   309      -8.679 -24.092   0.358  1.00 17.85       B    N
ATOM   1757  CA  VAL   309      -8.288 -23.942   1.762  1.00 17.40       B    C
ATOM   1758  CB  VAL   309      -8.113 -25.327   2.426  1.00 15.52       B    C
ATOM   1759  CG1 VAL   309      -7.427 -25.197   3.750  1.00 16.49       B    C
ATOM   1760  CG2 VAL   309      -9.465 -25.941   2.660  1.00 14.46       B    C
ATOM   1761  C   VAL   309      -7.039 -23.065   2.012  1.00 17.87       B    C
ATOM   1762  O   VAL   309      -6.136 -22.986   1.179  1.00 19.05       B    O
ATOM   1763  N   VAL   310      -7.020 -22.385   3.159  1.00 17.38       B    N
ATOM   1764  CA  VAL   310      -5.918 -21.508   3.550  1.00 18.63       B    C
ATOM   1765  CB  VAL   310      -6.408 -20.376   4.464  1.00 19.64       B    C
ATOM   1766  CG1 VAL   310      -5.227 -19.605   5.018  1.00 18.09       B    C
ATOM   1767  CG2 VAL   310      -7.334 -19.464   3.702  1.00 18.46       B    C
ATOM   1768  C   VAL   310      -4.854 -22.270   4.318  1.00 19.53       B    C
ATOM   1769  O   VAL   310      -5.146 -22.917   5.327  1.00 19.08       B    O
ATOM   1770  N   LEU   311      -3.612 -22.189   3.857  1.00 20.49       B    N
ATOM   1771  CA  LEU   311      -2.539 -22.836   4.601  1.00 20.59       B    C
ATOM   1772  CB  LEU   311      -1.847 -23.898   3.729  1.00 21.10       B    C
ATOM   1773  CG  LEU   311      -2.719 -25.114   3.374  1.00 20.77       B    C
ATOM   1774  CD1 LEU   311      -3.169 -25.035   1.924  1.00 19.01       B    C
ATOM   1775  CD2 LEU   311      -1.929 -26.384   3.595  1.00 19.85       B    C
ATOM   1776  C   LEU   311      -1.523 -21.846   5.170  1.00 19.52       B    C
ATOM   1777  O   LEU   311      -0.989 -20.984   4.465  1.00 18.71       B    O
ATOM   1778  N   VAL   312      -1.287 -21.971   6.469  1.00 19.75       B    N
ATOM   1779  CA  VAL   312      -0.284 -21.166   7.165  1.00 20.13       B    C
ATOM   1780  CB  VAL   312      -0.954 -20.298   8.283  1.00 21.70       B    C
ATOM   1781  CG1 VAL   312       0.103 -19.499   9.048  1.00 20.27       B    C
ATOM   1782  CG2 VAL   312      -1.998 -19.364   7.660  1.00 18.86       B    C
ATOM   1783  C   VAL   312       0.774 -22.086   7.784  1.00 18.09       B    C
ATOM   1784  O   VAL   312       0.442 -23.018   8.528  1.00 15.74       B    O
ATOM   1785  N   THR   313       2.043 -21.833   7.463  1.00 17.81       B    N
ATOM   1786  CA  THR   313       3.130 -22.664   8.001  1.00 17.87       B    C
ATOM   1787  CB  THR   313       3.775 -23.550   6.924  1.00 16.57       B    C
ATOM   1788  OG1 THR   313       4.473 -24.615   7.568  1.00 15.42       B    O
ATOM   1789  CG2 THR   313       4.772 -22.752   6.088  1.00 13.87       B    C
ATOM   1790  C   THR   313       4.265 -21.899   8.670  1.00 17.42       B    C
ATOM   1791  O   THR   313       4.567 -20.752   8.306  1.00 19.31       B    O
ATOM   1792  N   ALA   314       4.893 -22.551   9.645  1.00 15.83       B    N
ATOM   1793  CA  ALA   314       6.148 -22.070  10.220  1.00 14.33       B    C
ATOM   1794  CB  ALA   314       6.611 -23.014  11.316  1.00 12.97       B    C
ATOM   1795  C   ALA   314       7.231 -21.944   9.147  1.00 13.75       B    C
ATOM   1796  O   ALA   314       7.228 -22.653   8.139  1.00 12.56       B    O
ATOM   1797  N   ALA   315       8.152 -21.016   9.366  1.00 14.84       B    N
ATOM   1798  CA  ALA   315       9.267 -20.818   8.455  1.00 15.19       B    C
ATOM   1799  CB  ALA   315       9.764 -19.384   8.563  1.00 12.98       B    C
ATOM   1800  C   ALA   315      10.398 -21.803   8.793  1.00 16.33       B    C
ATOM   1801  O   ALA   315      11.223 -22.147   7.933  1.00 16.84       B    O
ATOM   1802  N   GLY   316      10.416 -22.251  10.049  1.00 15.86       B    N
ATOM   1803  CA  GLY   316      11.460 -23.134  10.522  1.00 15.85       B    C
ATOM   1804  C   GLY   316      12.487 -22.390  11.352  1.00 17.04       B    C
ATOM   1805  O   GLY   316      12.662 -21.181  11.191  1.00 15.26       B    O
ATOM   1806  N   ASN   317      13.173 -23.123  12.230  1.00 18.37       B    N
ATOM   1807  CA  ASN   317      14.084 -22.530  13.205  1.00 20.16       B    C
ATOM   1808  CB  ASN   317      13.729 -23.023  14.600  1.00 17.67       B    C
ATOM   1809  CG  ASN   317      12.270 -22.864  14.894  1.00 17.51       B    C
ATOM   1810  OD1 ASN   317      11.632 -21.947  14.385  1.00 15.76       B    O
ATOM   1811  ND2 ASN   317      11.721 -23.758  15.709  1.00 19.11       B    N
ATOM   1812  C   ASN   317      15.535 -22.853  12.897  1.00 20.80       B    C
ATOM   1813  O   ASN   317      16.302 -23.250  13.779  1.00 20.23       B    O
ATOM   1814  N   PHE   318      15.899 -22.663  11.635  1.00 21.80       B    N
ATOM   1815  CA  PHE   318      17.167 -23.148  11.126  1.00 24.16       B    C
ATOM   1816  CB  PHE   318      16.905 -24.123   9.979  1.00 24.38       B    C
ATOM   1817  CG  PHE   318      16.091 -25.325  10.383  1.00 25.06       B    C
ATOM   1818  CD1 PHE   318      16.578 -26.227  11.319  1.00 23.42       B    C
ATOM   1819  CD2 PHE   318      14.847 -25.562   9.816  1.00 24.55       B    C
ATOM   1820  CE1 PHE   318      15.839 -27.341  11.681  1.00 23.94       B    C
ATOM   1821  CE2 PHE   318      14.104 -26.676  10.175  1.00 22.84       B    C
ATOM   1822  CZ  PHE   318      14.601 -27.565  11.108  1.00 22.93       B    C
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ATOM   1823  C   PHE  318      18.106 -22.038  10.666  1.00 25.02      B    C
ATOM   1824  O   PHE  318      19.162 -22.312  10.109  1.00 26.36      B    O
ATOM   1825  N   ARG  319      17.716 -20.789  10.893  1.00 25.77      B    N
ATOM   1826  CA  ARG  319      18.570 -19.654  10.571  1.00 26.61      B    C
ATOM   1827  CB  ARG  319      19.788 -19.643  11.508  1.00 28.47      B    C
ATOM   1828  CG  ARG  319      20.420 -18.273  11.749  1.00 32.16      B    C
ATOM   1829  CD  ARG  319      21.502 -18.345  12.822  1.00 36.07      B    C
ATOM   1830  NE  ARG  319      21.972 -17.023  13.247  1.00 42.52      B    N
ATOM   1831  CZ  ARG  319      23.231 -16.735  13.597  1.00 45.72      B    C
ATOM   1832  NH1 ARG  319      24.169 -17.674  13.579  1.00 44.82      B    N
ATOM   1833  NH2 ARG  319      23.560 -15.499  13.966  1.00 45.86      B    N
ATOM   1834  C   ARG  319      19.019 -19.766   9.115  1.00 26.48      B    C
ATOM   1835  O   ARG  319      20.203 -19.627   8.808  1.00 26.94      B    O
ATOM   1836  N   ASP  320      18.066 -20.022   8.221  1.00 25.91      B    N
ATOM   1837  CA  ASP  320      18.377 -20.386   6.839  1.00 25.38      B    C
ATOM   1838  CB  ASP  320      18.482 -21.907   6.716  1.00 27.38      B    C
ATOM   1839  CG  ASP  320      19.283 -22.347   5.492  1.00 31.81      B    C
ATOM   1840  OD1 ASP  320      19.625 -21.483   4.646  1.00 30.34      B    O
ATOM   1841  OD2 ASP  320      19.572 -23.567   5.381  1.00 34.03      B    O
ATOM   1842  C   ASP  320      17.310 -19.879   5.882  1.00 23.72      B    C
ATOM   1843  O   ASP  320      16.259 -19.420   6.311  1.00 24.64      B    O
ATOM   1844  N   ASP  321      17.578 -19.959   4.584  1.00 21.64      B    N
ATOM   1845  CA  ASP  321      16.545 -19.676   3.604  1.00 20.23      B    C
ATOM   1846  CB  ASP  321      17.118 -19.655   2.187  1.00 20.01      B    C
ATOM   1847  CG  ASP  321      16.160 -19.032   1.177  1.00 23.00      B    C
ATOM   1848  OD1 ASP  321      15.034 -18.631   1.557  1.00 25.85      B    O
ATOM   1849  OD2 ASP  321      16.529 -18.936  -0.009  1.00 24.40      B    O
ATOM   1850  C   ASP  321      15.500 -20.768   3.715  1.00 20.33      B    C
ATOM   1851  O   ASP  321      15.785 -21.937   3.470  1.00 20.93      B    O
ATOM   1852  N   ALA  322      14.289 -20.376   4.092  1.00 20.81      B    N
ATOM   1853  CA  ALA  322      13.189 -21.311   4.245  1.00 20.82      B    C
ATOM   1854  CB  ALA  322      12.027 -20.619   4.903  1.00 20.87      B    C
ATOM   1855  C   ALA  322      12.773 -21.879   2.896  1.00 22.07      B    C
ATOM   1856  O   ALA  322      11.925 -22.770   2.814  1.00 24.23      B    O
ATOM   1857  N   CYS  323      13.380 -21.362   1.836  1.00 21.62      B    N
ATOM   1858  CA  CYS  323      13.189 -21.906   0.505  1.00 21.78      B    C
ATOM   1859  C   CYS  323      13.745 -23.327   0.420  1.00 20.87      B    C
ATOM   1860  O   CYS  323      13.523 -24.041  -0.562  1.00 20.76      B    O
ATOM   1861  CB  CYS  323      13.883 -21.013  -0.517  1.00 23.38      B    C
ATOM   1862  SG  CYS  323      12.973 -19.497  -0.993  1.00 34.53      B    S
ATOM   1863  N   LEU  324      14.473 -23.729   1.457  1.00 19.41      B    N
ATOM   1864  CA  LEU  324      15.129 -25.034   1.485  1.00 17.45      B    C
ATOM   1865  CB  LEU  324      16.580 -24.895   1.928  1.00 14.89      B    C
ATOM   1866  CG  LEU  324      17.426 -23.921   1.129  1.00 14.62      B    C
ATOM   1867  CD1 LEU  324      18.812 -23.877   1.719  1.00 12.16      B    C
ATOM   1868  CD2 LEU  324      17.470 -24.357  -0.324  1.00 16.39      B    C
ATOM   1869  C   LEU  324      14.436 -26.000   2.426  1.00 17.67      B    C
ATOM   1870  O   LEU  324      14.998 -27.046   2.769  1.00 17.60      B    O
ATOM   1871  N   TYR  325      13.231 -25.643   2.862  1.00 16.35      B    N
ATOM   1872  CA  TYR  325      12.460 -26.512   3.744  1.00 15.57      B    C
ATOM   1873  CB  TYR  325      12.534 -26.011   5.178  1.00 14.14      B    C
ATOM   1874  CG  TYR  325      13.949 -25.807   5.651  1.00 14.56      B    C
ATOM   1875  CD1 TYR  325      14.620 -26.806   6.355  1.00 12.52      B    C
ATOM   1876  CE1 TYR  325      15.915 -26.636   6.755  1.00 11.36      B    C
ATOM   1877  CD2 TYR  325      14.626 -24.631   5.363  1.00 10.83      B    C
ATOM   1878  CE2 TYR  325      15.916 -24.455   5.753  1.00 12.48      B    C
ATOM   1879  CZ  TYR  325      16.563 -25.459   6.449  1.00 13.44      B    C
ATOM   1880  OH  TYR  325      17.869 -25.275   6.839  1.00 14.00      B    O
ATOM   1881  C   TYR  325      11.017 -26.553   3.303  1.00 16.17      B    C
ATOM   1882  O   TYR  325      10.554 -25.661   2.593  1.00 19.07      B    O
ATOM   1883  N   SER  326      10.307 -27.594   3.718  1.00 15.32      B    N
ATOM   1884  CA  SER  326       8.884 -27.706   3.436  1.00 14.45      B    C
ATOM   1885  CB  SER  326       8.643 -28.777   2.365  1.00 13.51      B    C
ATOM   1886  OG  SER  326       9.232 -28.396   1.136  1.00  9.68      B    O
ATOM   1887  C   SER  326       8.098 -28.036   4.702  1.00 14.43      B    C
ATOM   1888  O   SER  326       8.596 -28.721   5.593  1.00 13.96      B    O
ATOM   1889  N   PRO  327       6.846 -27.557   4.791  1.00 16.56      B    N
ATOM   1890  CD  PRO  327       5.970 -27.849   5.938  1.00 16.61      B    C
ATOM   1891  CA  PRO  327       6.131 -26.829   3.729  1.00 17.34      B    C
ATOM   1892  CB  PRO  327       4.669 -26.884   4.171  1.00 15.73      B    C
ATOM   1893  CG  PRO  327       4.735 -27.040   5.641  1.00 16.44      B    C
ATOM   1894  C   PRO  327       6.582 -25.397   3.456  1.00 18.31      B    C
ATOM   1895  O   PRO  327       6.117 -24.779   2.501  1.00 18.41      B    O
ATOM   1896  N   ALA  328       7.478 -24.872   4.286  1.00 20.01      B    N
ATOM   1897  CA  ALA  328       7.811 -23.450   4.231  1.00 22.14      B    C
ATOM   1898  CB  ALA  328       9.031 -23.170   5.107  1.00 21.67      B    C
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ATOM   1899  C   ALA   328       8.056 -22.946   2.799  1.00 23.20      B    C
ATOM   1900  O   ALA   328       7.640 -21.835   2.438  1.00 23.95      B    O
ATOM   1901  N   SER   329       8.711 -23.773   1.986  1.00 22.51      B    N
ATOM   1902  CA  SER   329       9.159 -23.358   0.662  1.00 22.34      B    C
ATOM   1903  CB  SER   329      10.296 -24.261   0.199  1.00 21.26      B    C
ATOM   1904  OG  SER   329       9.830 -25.579  -0.021  1.00 18.71      B    O
ATOM   1905  C   SER   329       8.054 -23.355  -0.399  1.00 23.40      B    C
ATOM   1906  O   SER   329       8.228 -22.787  -1.476  1.00 23.51      B    O
ATOM   1907  N   ALA   330       6.926 -23.993  -0.100  1.00 25.31      B    N
ATOM   1908  CA  ALA   330       5.835 -24.107  -1.064  1.00 26.56      B    C
ATOM   1909  CB  ALA   330       4.822 -25.133  -0.590  1.00 27.06      B    C
ATOM   1910  C   ALA   330       5.160 -22.758  -1.250  1.00 27.53      B    C
ATOM   1911  O   ALA   330       4.727 -22.133  -0.284  1.00 29.15      B    O
ATOM   1912  N   PRO   331       5.065 -22.293  -2.503  1.00 28.24      B    N
ATOM   1913  CD  PRO   331       5.431 -23.095  -3.682  1.00 28.39      B    C
ATOM   1914  CA  PRO   331       4.718 -20.909  -2.866  1.00 29.18      B    C
ATOM   1915  CB  PRO   331       4.844 -20.891  -4.391  1.00 28.66      B    C
ATOM   1916  CG  PRO   331       5.727 -22.051  -4.711  1.00 28.90      B    C
ATOM   1917  C   PRO   331       3.326 -20.464  -2.416  1.00 29.39      B    C
ATOM   1918  O   PRO   331       3.115 -19.296  -2.078  1.00 29.76      B    O
ATOM   1919  N   GLU   332       2.384 -21.400  -2.420  1.00 28.61      B    N
ATOM   1920  CA  GLU   332       0.983 -21.080  -2.211  1.00 28.81      B    C
ATOM   1921  CB  GLU   332       0.121 -22.089  -2.970  1.00 31.96      B    C
ATOM   1922  CG  GLU   332       0.851 -23.396  -3.296  1.00 36.23      B    C
ATOM   1923  CD  GLU   332       1.701 -23.306  -4.566  1.00 39.36      B    C
ATOM   1924  OE1 GLU   332       1.169 -22.910  -5.630  1.00 41.08      B    O
ATOM   1925  OE2 GLU   332       2.904 -23.639  -4.503  1.00 40.19      B    O
ATOM   1926  C   GLU   332       0.637 -21.087  -0.722  1.00 27.73      B    C
ATOM   1927  O   GLU   332      -0.452 -20.683  -0.318  1.00 26.94      B    O
ATOM   1928  N   VAL   333       1.575 -21.554   0.091  1.00 26.86      B    N
ATOM   1929  CA  VAL   333       1.422 -21.524   1.541  1.00 25.32      B    C
ATOM   1930  CB  VAL   333       2.269 -22.633   2.227  1.00 23.61      B    C
ATOM   1931  CG1 VAL   333       2.247 -22.441   3.732  1.00 23.08      B    C
ATOM   1932  CG2 VAL   333       1.727 -24.009   1.873  1.00 23.25      B    C
ATOM   1933  C   VAL   333       1.848 -20.166   2.101  1.00 24.40      B    C
ATOM   1934  O   VAL   333       2.781 -19.529   1.596  1.00 24.78      B    O
ATOM   1935  N   ILE   334       1.156 -19.724   3.145  1.00 23.38      B    N
ATOM   1936  CA  ILE   334       1.588 -18.546   3.878  1.00 22.82      B    C
ATOM   1937  CB  ILE   334       0.413 -17.867   4.582  1.00 23.36      B    C
ATOM   1938  CG2 ILE   334       0.908 -16.682   5.373  1.00 21.37      B    C
ATOM   1939  CG1 ILE   334      -0.617 -17.418   3.546  1.00 22.38      B    C
ATOM   1940  CD1 ILE   334      -1.908 -16.964   4.149  1.00 19.76      B    C
ATOM   1941  C   ILE   334       2.626 -18.950   4.910  1.00 21.94      B    C
ATOM   1942  O   ILE   334       2.337 -19.676   5.863  1.00 20.41      B    O
ATOM   1943  N   THR   335       3.844 -18.471   4.694  1.00 21.71      B    N
ATOM   1944  CA  THR   335       5.010 -18.951   5.419  1.00 22.66      B    C
ATOM   1945  CB  THR   335       6.141 -19.283   4.426  1.00 24.07      B    C
ATOM   1946  OG1 THR   335       5.600 -20.019   3.315  1.00 24.80      B    O
ATOM   1947  CG2 THR   335       7.219 -20.116   5.100  1.00 23.59      B    C
ATOM   1948  C   THR   335       5.463 -17.861   6.394  1.00 22.37      B    C
ATOM   1949  O   THR   335       5.772 -16.741   5.988  1.00 21.19      B    O
ATOM   1950  N   VAL   336       5.486 -18.192   7.682  1.00 22.56      B    N
ATOM   1951  CA  VAL   336       5.552 -17.173   8.731  1.00 22.60      B    C
ATOM   1952  CB  VAL   336       4.355 -17.296   9.692  1.00 22.03      B    C
ATOM   1953  CG1 VAL   336       4.359 -16.147  10.689  1.00 19.46      B    C
ATOM   1954  CG2 VAL   336       3.069 -17.332   8.897  1.00 21.19      B    C
ATOM   1955  C   VAL   336       6.824 -17.242   9.566  1.00 22.61      B    C
ATOM   1956  O   VAL   336       7.119 -18.276  10.160  1.00 22.75      B    O
ATOM   1957  N   GLY   337       7.553 -16.127   9.623  1.00 23.52      B    N
ATOM   1958  CA  GLY   337       8.746 -16.036  10.454  1.00 24.53      B    C
ATOM   1959  C   GLY   337       8.500 -15.342  11.784  1.00 25.08      B    C
ATOM   1960  O   GLY   337       7.451 -14.745  11.996  1.00 26.14      B    O
ATOM   1961  N   ALA   338       9.463 -15.413  12.691  1.00 25.21      B    N
ATOM   1962  CA  ALA   338       9.243 -14.906  14.036  1.00 25.87      B    C
ATOM   1963  CB  ALA   338       9.544 -15.990  15.047  1.00 25.68      B    C
ATOM   1964  C   ALA   338      10.081 -13.674  14.340  1.00 27.22      B    C
ATOM   1965  O   ALA   338      11.271 -13.641  14.033  1.00 28.69      B    O
ATOM   1966  N   THR   339       9.452 -12.665  14.943  1.00 27.73      B    N
ATOM   1967  CA  THR   339      10.165 -11.528  15.529  1.00 27.88      B    C
ATOM   1968  CB  THR   339       9.680 -10.187  14.944  1.00 28.43      B    C
ATOM   1969  OG1 THR   339       8.410  -9.838  15.515  1.00 30.24      B    O
ATOM   1970  CG2 THR   339       9.535 -10.297  13.438  1.00 27.60      B    C
ATOM   1971  C   THR   339       9.912 -11.514  17.038  1.00 28.11      B    C
ATOM   1972  O   THR   339       9.127 -12.313  17.545  1.00 29.32      B    O
ATOM   1973  N   ASN   340      10.576 -10.612  17.754  1.00 28.25      B    N
ATOM   1974  CA  ASN   340      10.390 -10.507  19.197  1.00 28.64      B    C
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ATOM   1975  CB  ASN   340      11.683 -10.900  19.937  1.00 26.79      B    C
ATOM   1976  CG  ASN   340      12.780  -9.837  19.842  1.00 24.82      B    C
ATOM   1977  OD1 ASN   340      12.778  -8.987  18.954  1.00 23.33      B    O
ATOM   1978  ND2 ASN   340      13.718  -9.889  20.768  1.00 21.41      B    N
ATOM   1979  C   ASN   340       9.947  -9.105  19.614  1.00 30.56      B    C
ATOM   1980  O   ASN   340       9.662  -8.253  18.776  1.00 30.36      B    O
ATOM   1981  N   ALA   341       9.886  -8.874  20.917  1.00 32.81      B    N
ATOM   1982  CA  ALA   341       9.383  -7.618  21.435  1.00 34.72      B    C
ATOM   1983  CB  ALA   341       9.352  -7.677  22.943  1.00 33.86      B    C
ATOM   1984  C   ALA   341      10.193  -6.404  20.959  1.00 37.05      B    C
ATOM   1985  O   ALA   341       9.686  -5.281  20.932  1.00 37.22      B    O
ATOM   1986  N   GLN   342      11.449  -6.617  20.582  1.00 39.17      B    N
ATOM   1987  CA  GLN   342      12.237  -5.518  20.037  1.00 41.65      B    C
ATOM   1988  CB  GLN   342      13.682  -5.544  20.558  1.00 43.43      B    C
ATOM   1989  CG  GLN   342      14.069  -6.779  21.353  1.00 47.47      B    C
ATOM   1990  CD  GLN   342      13.394  -6.830  22.709  1.00 49.49      B    C
ATOM   1991  OE1 GLN   342      13.198  -7.907  23.289  1.00 48.68      B    O
ATOM   1992  NE2 GLN   342      13.024  -5.660  23.223  1.00 50.00      B    N
ATOM   1993  C   GLN   342      12.239  -5.576  18.518  1.00 43.22      B    C
ATOM   1994  O   GLN   342      13.174  -5.104  17.869  1.00 43.77      B    O
ATOM   1995  N   ASP   343      11.189  -6.170  17.956  1.00 43.93      B    N
ATOM   1996  CA  ASP   343      10.986  -6.170  16.509  1.00 43.53      B    C
ATOM   1997  CB  ASP   343      10.524  -4.775  16.050  1.00 43.99      B    C
ATOM   1998  CG  ASP   343       9.011  -4.565  16.209  1.00 44.42      B    C
ATOM   1999  OD1 ASP   343       8.245  -5.558  16.276  1.00 42.52      B    O
ATOM   2000  OD2 ASP   343       8.587  -3.393  16.260  1.00 44.78      B    O
ATOM   2001  C   ASP   343      12.244  -6.601  15.737  1.00 42.39      B    C
ATOM   2002  O   ASP   343      12.546  -6.085  14.657  1.00 42.07      B    O
ATOM   2003  N   GLN   344      12.976  -7.552  16.302  1.00 41.78      B    N
ATOM   2004  CA  GLN   344      14.134  -8.127  15.629  1.00 41.56      B    C
ATOM   2005  CB  GLN   344      15.408  -7.797  16.396  1.00 44.09      B    C
ATOM   2006  CG  GLN   344      15.384  -8.247  17.839  1.00 48.25      B    C
ATOM   2007  CD  GLN   344      16.394  -7.499  18.687  1.00 50.65      B    C
ATOM   2008  OE1 GLN   344      16.657  -7.863  19.840  1.00 51.73      B    O
ATOM   2009  NE2 GLN   344      16.968  -6.440  18.118  1.00 51.25      B    N
ATOM   2010  C   GLN   344      13.963  -9.630  15.540  1.00 38.96      B    C
ATOM   2011  O   GLN   344      13.247 -10.227  16.341  1.00 38.19      B    O
ATOM   2012  N   PRO   345      14.614 -10.263  14.556  1.00 37.58      B    N
ATOM   2013  CD  PRO   345      15.492  -9.635  13.555  1.00 36.39      B    C
ATOM   2014  CA  PRO   345      14.416 -11.698  14.298  1.00 37.16      B    C
ATOM   2015  CB  PRO   345      15.331 -11.980  13.101  1.00 35.35      B    C
ATOM   2016  CG  PRO   345      15.514 -10.648  12.446  1.00 35.43      B    C
ATOM   2017  C   PRO   345      14.759 -12.573  15.510  1.00 37.13      B    C
ATOM   2018  O   PRO   345      15.782 -12.364  16.162  1.00 39.08      B    O
ATOM   2019  N   VAL   346      13.910 -13.551  15.811  1.00 36.05      B    N
ATOM   2020  CA  VAL   346      14.149 -14.421  16.957  1.00 34.82      B    C
ATOM   2021  CB  VAL   346      12.915 -15.281  17.287  1.00 33.18      B    C
ATOM   2022  CG1 VAL   346      13.244 -16.221  18.428  1.00 33.07      B    C
ATOM   2023  CG2 VAL   346      11.740 -14.401  17.664  1.00 31.64      B    C
ATOM   2024  C   VAL   346      15.333 -15.359  16.745  1.00 35.10      B    C
ATOM   2025  O   VAL   346      15.385 -16.111  15.770  1.00 34.39      B    O
ATOM   2026  N   THR   347      16.285 -15.311  17.669  1.00 35.85      B    N
ATOM   2027  CA  THR   347      17.388 -16.264  17.671  1.00 37.77      B    C
ATOM   2028  CB  THR   347      18.718 -15.599  18.074  1.00 36.41      B    C
ATOM   2029  OG1 THR   347      18.567 -14.985  19.358  1.00 35.69      B    O
ATOM   2030  CG2 THR   347      19.129 -14.554  17.050  1.00 33.79      B    C
ATOM   2031  C   THR   347      17.084 -17.381  18.661  1.00 39.16      B    C
ATOM   2032  O   THR   347      16.755 -17.124  19.819  1.00 39.34      B    O
ATOM   2033  N   LEU   348      17.188 -18.620  18.196  1.00 40.40      B    N
ATOM   2034  CA  LEU   348      16.839 -19.770  19.014  1.00 42.15      B    C
ATOM   2035  CB  LEU   348      15.727 -20.580  18.334  1.00 42.90      B    C
ATOM   2036  CG  LEU   348      14.364 -19.887  18.207  1.00 44.13      B    C
ATOM   2037  CD1 LEU   348      13.468 -20.686  17.284  1.00 44.95      B    C
ATOM   2038  CD2 LEU   348      13.725 -19.739  19.584  1.00 44.34      B    C
ATOM   2039  C   LEU   348      18.076 -20.631  19.221  1.00 42.53      B    C
ATOM   2040  O   LEU   348      18.481 -21.390  18.333  1.00 44.15      B    O
ATOM   2041  N   GLY   349      18.680 -20.514  20.396  1.00 41.43      B    N
ATOM   2042  CA  GLY   349      19.985 -21.107  20.581  1.00 40.94      B    C
ATOM   2043  C   GLY   349      20.931 -20.405  19.636  1.00 40.49      B    C
ATOM   2044  O   GLY   349      20.898 -19.177  19.536  1.00 40.57      B    O
ATOM   2045  N   THR   350      21.761 -21.172  18.934  1.00 40.43      B    N
ATOM   2046  CA  THR   350      22.649 -20.602  17.923  1.00 40.06      B    C
ATOM   2047  CB  THR   350      23.967 -21.395  17.796  1.00 41.33      B    C
ATOM   2048  OG1 THR   350      23.689 -22.717  17.311  1.00 42.85      B    O
ATOM   2049  CG2 THR   350      24.679 -21.468  19.142  1.00 39.97      B    C
ATOM   2050  C   THR   350      21.976 -20.596  16.557  1.00 38.80      B    C
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ATOM   2051  O   THR  350      22.599 -20.288  15.543  1.00 39.80      B    O
ATOM   2052  N   LEU  351      20.702 -20.960  16.533  1.00 36.87      B    N
ATOM   2053  CA  LEU  351      19.916 -20.858  15.320  1.00 35.16      B    C
ATOM   2054  CB  LEU  351      19.330 -22.236  14.982  1.00 34.71      B    C
ATOM   2055  CG  LEU  351      20.325 -23.192  14.299  1.00 32.23      B    C
ATOM   2056  CD1 LEU  351      20.033 -24.639  14.626  1.00 29.33      B    C
ATOM   2057  CD2 LEU  351      20.247 -22.975  12.807  1.00 32.45      B    C
ATOM   2058  C   LEU  351      18.825 -19.788  15.484  1.00 34.05      B    C
ATOM   2059  O   LEU  351      18.970 -18.858  16.284  1.00 33.71      B    O
ATOM   2060  N   GLY  352      17.747 -19.899  14.717  1.00 33.23      B    N
ATOM   2061  CA  GLY  352      16.666 -18.938  14.839  1.00 30.44      B    C
ATOM   2062  C   GLY  352      15.731 -18.988  13.652  1.00 28.36      B    C
ATOM   2063  O   GLY  352      15.681 -19.984  12.934  1.00 27.74      B    O
ATOM   2064  N   THR  353      14.991 -17.907  13.441  1.00 25.82      B    N
ATOM   2065  CA  THR  353      13.991 -17.868  12.386  1.00 23.39      B    C
ATOM   2066  CB  THR  353      13.177 -16.565  12.446  1.00 23.46      B    C
ATOM   2067  OG1 THR  353      12.050 -16.652  11.559  1.00 22.87      B    O
ATOM   2068  CG2 THR  353      14.054 -15.387  12.031  1.00 23.22      B    C
ATOM   2069  C   THR  353      14.657 -17.954  11.020  1.00 21.85      B    C
ATOM   2070  O   THR  353      15.659 -17.290  10.768  1.00 23.19      B    O
ATOM   2071  N   ASN  354      14.099 -18.776  10.142  1.00 19.78      B    N
ATOM   2072  CA  ASN  354      14.513 -18.776   8.751  1.00 18.88      B    C
ATOM   2073  CB  ASN  354      13.905 -19.981   8.029  1.00 19.41      B    C
ATOM   2074  CG  ASN  354      14.733 -21.232   8.203  1.00 18.91      B    C
ATOM   2075  OD1 ASN  354      15.835 -21.175   8.720  1.00 22.77      B    O
ATOM   2076  ND2 ASN  354      14.211 -22.363   7.765  1.00 20.48      B    N
ATOM   2077  C   ASN  354      14.057 -17.464   8.114  1.00 19.36      B    C
ATOM   2078  O   ASN  354      13.425 -16.641   8.775  1.00 18.72      B    O
ATOM   2079  N   PHE  355      14.380 -17.258   6.842  1.00 20.03      B    N
ATOM   2080  CA  PHE  355      14.126 -15.974   6.191  1.00 22.21      B    C
ATOM   2081  CB  PHE  355      15.174 -14.960   6.638  1.00 19.82      B    C
ATOM   2082  CG  PHE  355      16.550 -15.546   6.737  1.00 20.47      B    C
ATOM   2083  CD1 PHE  355      17.080 -15.893   7.972  1.00 19.70      B    C
ATOM   2084  CD2 PHE  355      17.281 -15.832   5.590  1.00 19.94      B    C
ATOM   2085  CE1 PHE  355      18.305 -16.521   8.064  1.00 18.39      B    C
ATOM   2086  CE2 PHE  355      18.504 -16.457   5.676  1.00 19.14      B    C
ATOM   2087  CZ  PHE  355      19.016 -16.804   6.918  1.00 18.75      B    C
ATOM   2088  C   PHE  355      14.188 -16.140   4.672  1.00 24.66      B    C
ATOM   2089  O   PHE  355      14.063 -17.254   4.154  1.00 26.20      B    O
ATOM   2090  N   GLY  356      14.384 -15.031   3.963  1.00 25.89      B    N
ATOM   2091  CA  GLY  356      14.478 -15.090   2.520  1.00 27.05      B    C
ATOM   2092  C   GLY  356      13.145 -14.824   1.856  1.00 28.45      B    C
ATOM   2093  O   GLY  356      12.172 -14.473   2.509  1.00 28.91      B    O
ATOM   2094  N   ARG  357      13.103 -15.003   0.545  1.00 29.61      B    N
ATOM   2095  CA  ARG  357      11.921 -14.683  -0.235  1.00 30.42      B    C
ATOM   2096  CB  ARG  357      12.312 -14.507  -1.702  1.00 33.04      B    C
ATOM   2097  CG  ARG  357      12.919 -15.749  -2.307  1.00 38.74      B    C
ATOM   2098  CD  ARG  357      12.688 -15.791  -3.794  1.00 43.88      B    C
ATOM   2099  NE  ARG  357      13.524 -14.837  -4.515  1.00 49.58      B    N
ATOM   2100  CZ  ARG  357      14.766 -15.094  -4.921  1.00 52.78      B    C
ATOM   2101  NH1 ARG  357      15.456 -14.164  -5.580  1.00 53.88      B    N
ATOM   2102  NH2 ARG  357      15.322 -16.277  -4.662  1.00 53.07      B    N
ATOM   2103  C   ARG  357      10.802 -15.725  -0.121  1.00 29.64      B    C
ATOM   2104  O   ARG  357       9.708 -15.518  -0.635  1.00 28.09      B    O
ATOM   2105  N   CYS  358      11.064 -16.845   0.543  1.00 29.30      B    N
ATOM   2106  CA  CYS  358       9.998 -17.817   0.778  1.00 28.85      B    C
ATOM   2107  C   CYS  358       9.170 -17.473   2.017  1.00 27.26      B    C
ATOM   2108  O   CYS  358       8.119 -18.067   2.257  1.00 27.00      B    O
ATOM   2109  CB  CYS  358      10.570 -19.239   0.899  1.00 30.23      B    C
ATOM   2110  SG  CYS  358      10.992 -19.934  -0.727  1.00 33.15      B    S
ATOM   2111  N   VAL  359       9.635 -16.508   2.801  1.00 25.41      B    N
ATOM   2112  CA  VAL  359       8.846 -16.037   3.927  1.00 24.19      B    C
ATOM   2113  CB  VAL  359       9.747 -15.506   5.072  1.00 24.35      B    C
ATOM   2114  CG1 VAL  359       8.898 -14.934   6.197  1.00 22.54      B    C
ATOM   2115  CG2 VAL  359      10.602 -16.640   5.615  1.00 25.11      B    C
ATOM   2116  C   VAL  359       7.903 -14.941   3.461  1.00 23.48      B    C
ATOM   2117  O   VAL  359       8.282 -14.058   2.693  1.00 24.15      B    O
ATOM   2118  N   ASP  360       6.660 -15.017   3.914  1.00 22.77      B    N
ATOM   2119  CA  ASP  360       5.648 -14.067   3.504  1.00 20.61      B    C
ATOM   2120  CB  ASP  360       4.286 -14.766   3.445  1.00 23.47      B    C
ATOM   2121  CG  ASP  360       4.162 -15.711   2.237  1.00 26.57      B    C
ATOM   2122  OD1 ASP  360       3.832 -15.224   1.128  1.00 25.68      B    O
ATOM   2123  OD2 ASP  360       4.396 -16.935   2.395  1.00 27.94      B    O
ATOM   2124  C   ASP  360       5.632 -12.899   4.475  1.00 17.89      B    C
ATOM   2125  O   ASP  360       5.638 -11.740   4.069  1.00 16.84      B    O
ATOM   2126  N   LEU  361       5.639 -13.207   5.763  1.00 16.38      B    N
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ATOM	2127	CA	LEU	361	5.770	-12.171	6.777	1.00	16.17	B	C
ATOM	2128	CB	LEU	361	4.447	-11.425	6.946	1.00	17.23	B	C
ATOM	2129	CG	LEU	361	3.252	-12.186	7.523	1.00	16.84	B	C
ATOM	2130	CD1	LEU	361	2.087	-11.229	7.633	1.00	16.25	B	C
ATOM	2131	CD2	LEU	361	2.881	-13.374	6.643	1.00	18.27	B	C
ATOM	2132	C	LEU	361	6.225	-12.719	8.128	1.00	15.83	B	C
ATOM	2133	O	LEU	361	6.477	-13.913	8.278	1.00	16.68	B	O
ATOM	2134	N	PHE	362	6.345	-11.832	9.109	1.00	15.56	B	N
ATOM	2135	CA	PHE	362	6.792	-12.235	10.432	1.00	14.82	B	C
ATOM	2136	CB	PHE	362	8.059	-11.474	10.814	1.00	14.14	B	C
ATOM	2137	CG	PHE	362	9.259	-11.872	10.002	1.00	12.01	B	C
ATOM	2138	CD1	PHE	362	10.157	-12.818	10.486	1.00	9.24	B	C
ATOM	2139	CD2	PHE	362	9.464	-11.332	8.738	1.00	8.63	B	C
ATOM	2140	CE1	PHE	362	11.233	-13.218	9.726	1.00	8.18	B	C
ATOM	2141	CE2	PHE	362	10.534	-11.725	7.970	1.00	7.87	B	C
ATOM	2142	CZ	PHE	362	11.423	-12.671	8.463	1.00	9.03	B	C
ATOM	2143	C	PHE	362	5.710	-11.997	11.450	1.00	14.66	B	C
ATOM	2144	O	PHE	362	4.720	-11.336	11.158	1.00	17.08	B	O
ATOM	2145	N	ALA	363	5.891	-12.547	12.641	1.00	14.32	B	N
ATOM	2146	CA	ALA	363	4.877	-12.453	13.674	1.00	15.69	B	C
ATOM	2147	CB	ALA	363	3.767	-13.462	13.408	1.00	14.68	B	C
ATOM	2148	C	ALA	363	5.509	-12.717	15.025	1.00	17.04	B	C
ATOM	2149	O	ALA	363	6.544	-13.370	15.115	1.00	18.01	B	O
ATOM	2150	N	PRO	364	4.894	-12.207	16.101	1.00	18.24	B	N
ATOM	2151	CD	PRO	364	3.855	-11.163	16.138	1.00	17.59	B	C
ATOM	2152	CA	PRO	364	5.395	-12.521	17.438	1.00	18.94	B	C
ATOM	2153	CB	PRO	364	4.227	-12.146	18.339	1.00	18.02	B	C
ATOM	2154	CG	PRO	364	3.597	-10.978	17.621	1.00	17.86	B	C
ATOM	2155	C	PRO	364	5.753	-13.992	17.540	1.00	20.27	B	C
ATOM	2156	O	PRO	364	4.962	-14.851	17.152	1.00	22.37	B	O
ATOM	2157	N	GLY	365	6.950	-14.279	18.044	1.00	20.34	B	N
ATOM	2158	CA	GLY	365	7.380	-15.658	18.138	1.00	22.18	B	C
ATOM	2159	C	GLY	365	8.373	-15.948	19.243	1.00	23.70	B	C
ATOM	2160	O	GLY	365	8.973	-17.018	19.270	1.00	23.16	B	O
ATOM	2161	N	GLU	366	8.558	-15.002	20.154	1.00	25.89	B	N
ATOM	2162	CA	GLU	366	9.498	-15.193	21.252	1.00	29.11	B	C
ATOM	2163	CB	GLU	366	10.758	-14.359	21.023	1.00	31.40	B	C
ATOM	2164	CG	GLU	366	11.838	-14.585	22.053	1.00	35.84	B	C
ATOM	2165	CD	GLU	366	12.907	-13.508	22.017	1.00	40.32	B	C
ATOM	2166	OE1	GLU	366	13.931	-13.702	21.320	1.00	42.04	B	O
ATOM	2167	OE2	GLU	366	12.723	-12.462	22.685	1.00	43.30	B	O
ATOM	2168	C	GLU	366	8.856	-14.794	22.563	1.00	28.79	B	C
ATOM	2169	O	GLU	366	8.202	-13.753	22.642	1.00	28.23	B	O
ATOM	2170	N	ASP	367	9.048	-15.624	23.586	1.00	29.00	B	N
ATOM	2171	CA	ASP	367	8.400	-15.431	24.884	1.00	29.74	B	C
ATOM	2172	CB	ASP	367	8.868	-14.145	25.549	1.00	33.33	B	C
ATOM	2173	CG	ASP	367	10.172	-14.314	26.270	1.00	38.62	B	C
ATOM	2174	OD1	ASP	367	11.177	-13.708	25.839	1.00	42.20	B	O
ATOM	2175	OD2	ASP	367	10.195	-15.057	27.272	1.00	42.52	B	O
ATOM	2176	C	ASP	367	6.898	-15.371	24.771	1.00	28.66	B	C
ATOM	2177	O	ASP	367	6.273	-14.473	25.327	1.00	29.35	B	O
ATOM	2178	N	ILE	368	6.312	-16.316	24.047	1.00	28.05	B	N
ATOM	2179	CA	ILE	368	4.859	-16.347	23.911	1.00	26.36	B	C
ATOM	2180	CB	ILE	368	4.425	-16.924	22.542	1.00	24.44	B	C
ATOM	2181	CG2	ILE	368	2.921	-16.746	22.359	1.00	22.78	B	C
ATOM	2182	CG1	ILE	368	5.175	-16.208	21.414	1.00	23.14	B	C
ATOM	2183	CD1	ILE	368	4.953	-14.710	21.380	1.00	21.33	B	C
ATOM	2184	C	ILE	368	4.233	-17.175	25.028	1.00	25.21	B	C
ATOM	2185	O	ILE	368	4.365	-18.399	25.076	1.00	25.00	B	O
ATOM	2186	N	ILE	369	3.561	-16.491	25.940	1.00	23.47	B	N
ATOM	2187	CA	ILE	369	2.872	-17.179	27.007	1.00	24.06	B	C
ATOM	2188	CB	ILE	369	2.562	-16.226	28.161	1.00	24.47	B	C
ATOM	2189	CG2	ILE	369	2.407	-14.812	27.645	1.00	24.60	B	C
ATOM	2190	CG1	ILE	369	1.328	-16.723	28.901	1.00	24.42	B	C
ATOM	2191	CD1	ILE	369	0.614	-15.646	29.639	1.00	26.06	B	C
ATOM	2192	C	ILE	369	1.578	-17.780	26.496	1.00	23.71	B	C
ATOM	2193	O	ILE	369	0.827	-17.129	25.783	1.00	24.50	B	O
ATOM	2194	N	GLY	370	1.326	-19.029	26.866	1.00	24.27	B	N
ATOM	2195	CA	GLY	370	0.172	-19.743	26.357	1.00	24.97	B	C
ATOM	2196	C	GLY	370	-0.100	-20.957	27.214	1.00	26.60	B	C
ATOM	2197	O	GLY	370	0.565	-21.153	28.235	1.00	27.10	B	O
ATOM	2198	N	ALA	371	-1.063	-21.778	26.799	1.00	27.73	B	N
ATOM	2199	CA	ALA	371	-1.526	-22.902	27.615	1.00	27.34	B	C
ATOM	2200	CB	ALA	371	-2.705	-23.580	26.946	1.00	26.39	B	C
ATOM	2201	C	ALA	371	-0.431	-23.922	27.880	1.00	27.36	B	C
ATOM	2202	O	ALA	371	0.319	-24.296	26.980	1.00	28.65	B	O

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ATOM   2203  N   SER   372      -0.343 -24.371  29.126  1.00 27.77      B    N
ATOM   2204  CA  SER   372       0.621 -25.397  29.498  1.00 28.83      B    C
ATOM   2205  CB  SER   372       1.439 -24.952  30.706  1.00 29.82      B    C
ATOM   2206  OG  SER   372       2.034 -26.074  31.322  1.00 30.59      B    O
ATOM   2207  C   SER   372      -0.076 -26.700  29.829  1.00 29.05      B    C
ATOM   2208  O   SER   372      -0.969 -26.741  30.665  1.00 29.97      B    O
ATOM   2209  N   SER   373       0.349 -27.770  29.172  1.00 30.11      B    N
ATOM   2210  CA  SER   373      -0.304 -29.060  29.310  1.00 30.04      B    C
ATOM   2211  CB  SER   373       0.107 -29.979  28.159  1.00 30.84      B    C
ATOM   2212  OG  SER   373       1.493 -30.283  28.205  1.00 33.39      B    O
ATOM   2213  C   SER   373       0.062 -29.695  30.641  1.00 30.17      B    C
ATOM   2214  O   SER   373      -0.443 -30.761  30.986  1.00 29.58      B    O
ATOM   2215  N   ASP   374       0.942 -29.030  31.385  1.00 31.28      B    N
ATOM   2216  CA  ASP   374       1.373 -29.532  32.688  1.00 32.37      B    C
ATOM   2217  CB  ASP   374       2.410 -28.602  33.317  1.00 32.98      B    C
ATOM   2218  CG  ASP   374       3.832 -28.993  32.964  1.00 34.09      B    C
ATOM   2219  OD1 ASP   374       4.016 -29.822  32.042  1.00 35.12      B    O
ATOM   2220  OD2 ASP   374       4.763 -28.471  33.615  1.00 34.70      B    O
ATOM   2221  C   ASP   374       0.187 -29.699  33.616  1.00 31.98      B    C
ATOM   2222  O   ASP   374       0.079 -30.578  34.403  1.00 33.02      B    O
ATOM   2223  N   CYS   375      -0.696 -28.654  33.525  1.00 32.06      B    N
ATOM   2224  CA  CYS   375      -1.976 -28.705  34.204  1.00 31.88      B    C
ATOM   2225  C   CYS   375      -2.922 -27.755  33.477  1.00 30.87      B    C
ATOM   2226  O   CYS   375      -2.495 -26.986  32.618  1.00 28.42      B    O
ATOM   2227  CB  CYS   375      -1.809 -28.308  35.674  1.00 31.19      B    C
ATOM   2228  SG  CYS   375      -2.015 -26.538  36.040  1.00 37.01      B    S
ATOM   2229  N   SER   376      -4.207 -27.816  33.807  1.00 32.21      B    N
ATOM   2230  CA  SER   376      -5.221 -27.206  32.954  1.00 33.14      B    C
ATOM   2231  CB  SER   376      -6.529 -27.973  33.071  1.00 32.67      B    C
ATOM   2232  OG  SER   376      -6.680 -28.456  34.388  1.00 36.61      B    O
ATOM   2233  C   SER   376      -5.455 -25.734  33.237  1.00 32.73      B    C
ATOM   2234  O   SER   376      -6.133 -25.052  32.474  1.00 33.18      B    O
ATOM   2235  N   THR   377      -4.892 -25.245  34.335  1.00 33.07      B    N
ATOM   2236  CA  THR   377      -4.816 -23.808  34.571  1.00 32.43      B    C
ATOM   2237  CB  THR   377      -5.466 -23.429  35.916  1.00 32.44      B    C
ATOM   2238  OG1 THR   377      -4.997 -24.310  36.947  1.00 31.10      B    O
ATOM   2239  CG2 THR   377      -6.974 -23.510  35.801  1.00 31.10      B    C
ATOM   2240  C   THR   377      -3.366 -23.319  34.549  1.00 31.50      B    C
ATOM   2241  O   THR   377      -3.047 -22.242  35.061  1.00 30.92      B    O
ATOM   2242  N   CYS   378      -2.497 -24.116  33.936  1.00 29.98      B    N
ATOM   2243  CA  CYS   378      -1.081 -23.786  33.837  1.00 29.05      B    C
ATOM   2244  C   CYS   378      -0.729 -23.067  32.527  1.00 26.98      B    C
ATOM   2245  O   CYS   378      -1.265 -23.382  31.459  1.00 25.69      B    O
ATOM   2246  CB  CYS   378      -0.249 -25.064  33.989  1.00 29.82      B    C
ATOM   2247  SG  CYS   378      -0.190 -25.678  35.704  1.00 33.15      B    S
ATOM   2248  N   PHE   379       0.169 -22.090  32.627  1.00 24.89      B    N
ATOM   2249  CA  PHE   379       0.663 -21.372  31.463  1.00 23.14      B    C
ATOM   2250  CB  PHE   379       0.228 -19.909  31.530  1.00 21.19      B    C
ATOM   2251  CG  PHE   379      -1.233 -19.730  31.289  1.00 22.53      B    C
ATOM   2252  CD1 PHE   379      -2.151 -19.959  32.309  1.00 22.23      B    C
ATOM   2253  CD2 PHE   379      -1.709 -19.409  30.019  1.00 22.19      B    C
ATOM   2254  CE1 PHE   379      -3.520 -19.886  32.059  1.00 22.41      B    C
ATOM   2255  CE2 PHE   379      -3.072 -19.335  29.764  1.00 20.69      B    C
ATOM   2256  CZ  PHE   379      -3.978 -19.571  30.783  1.00 21.30      B    C
ATOM   2257  C   PHE   379       2.172 -21.483  31.357  1.00 23.01      B    C
ATOM   2258  O   PHE   379       2.837 -21.948  32.285  1.00 23.48      B    O
ATOM   2259  N   VAL   380       2.718 -21.073  30.215  1.00 21.60      B    N
ATOM   2260  CA  VAL   380       4.127 -21.259  29.953  1.00 21.47      B    C
ATOM   2261  CB  VAL   380       4.421 -22.730  29.624  1.00 20.54      B    C
ATOM   2262  CG1 VAL   380       3.910 -23.050  28.235  1.00 19.96      B    C
ATOM   2263  CG2 VAL   380       5.895 -23.015  29.760  1.00 19.12      B    C
ATOM   2264  C   VAL   380       4.526 -20.395  28.768  1.00 21.24      B    C
ATOM   2265  O   VAL   380       3.706 -20.143  27.891  1.00 21.49      B    O
ATOM   2266  N   SER   381       5.774 -19.937  28.737  1.00 21.07      B    N
ATOM   2267  CA  SER   381       6.242 -19.163  27.596  1.00 23.27      B    C
ATOM   2268  CB  SER   381       7.095 -17.992  28.073  1.00 24.23      B    C
ATOM   2269  OG  SER   381       7.575 -17.242  26.972  1.00 27.50      B    O
ATOM   2270  C   SER   381       7.038 -20.022  26.603  1.00 24.40      B    C
ATOM   2271  O   SER   381       7.786 -20.921  27.003  1.00 25.03      B    O
ATOM   2272  N   GLN   382       6.866 -19.759  25.310  1.00 22.98      B    N
ATOM   2273  CA  GLN   382       7.558 -20.537  24.294  1.00 24.19      B    C
ATOM   2274  CB  GLN   382       6.699 -21.731  23.851  1.00 25.69      B    C
ATOM   2275  CG  GLN   382       6.625 -22.896  24.858  1.00 28.37      B    C
ATOM   2276  CD  GLN   382       5.873 -24.131  24.310  1.00 31.99      B    C
ATOM   2277  OE1 GLN   382       5.701 -25.134  25.011  1.00 32.02      B    O
ATOM   2278  NE2 GLN   382       5.430 -24.054  23.055  1.00 33.34      B    N
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ATOM   2279  C   GLN   382       7.924 -19.672  23.090  1.00 24.25       B    C
ATOM   2280  O   GLN   382       7.331 -18.621  22.869  1.00 23.92       B    O
ATOM   2281  N   SER   383       8.918 -20.117  22.328  1.00 24.93       B    N
ATOM   2282  CA  SER   383       9.435 -19.366  21.186  1.00 25.23       B    C
ATOM   2283  CB  SER   383      10.741 -18.650  21.569  1.00 24.50       B    C
ATOM   2284  OG  SER   383      10.523 -17.648  22.550  1.00 21.80       B    O
ATOM   2285  C   SER   383       9.686 -20.306  20.003  1.00 25.02       B    C
ATOM   2286  O   SER   383       9.703 -21.521  20.165  1.00 26.23       B    O
ATOM   2287  N   GLY   384       9.892 -19.739  18.816  1.00 25.92       B    N
ATOM   2288  CA  GLY   384       9.975 -20.544  17.611  1.00 25.10       B    C
ATOM   2289  C   GLY   384       9.031 -20.032  16.534  1.00 25.74       B    C
ATOM   2290  O   GLY   384       8.130 -19.234  16.804  1.00 24.41       B    O
ATOM   2291  N   THR   385       9.240 -20.479  15.302  1.00 26.36       B    N
ATOM   2292  CA  THR   385       8.417 -20.030  14.185  1.00 26.63       B    C
ATOM   2293  CB  THR   385       9.056 -20.415  12.841  1.00 25.25       B    C
ATOM   2294  OG1 THR   385       9.504 -21.777  12.890  1.00 23.96       B    O
ATOM   2295  CG2 THR   385      10.221 -19.506  12.541  1.00 25.37       B    C
ATOM   2296  C   THR   385       7.003 -20.615  14.254  1.00 27.62       B    C
ATOM   2297  O   THR   385       6.094 -20.163  13.554  1.00 28.58       B    O
ATOM   2298  N   SER   386       6.820 -21.621  15.099  1.00 26.33       B    N
ATOM   2299  CA  SER   386       5.495 -22.160  15.316  1.00 26.38       B    C
ATOM   2300  CB  SER   386       5.572 -23.414  16.184  1.00 26.41       B    C
ATOM   2301  OG  SER   386       5.940 -24.523  15.395  1.00 27.85       B    O
ATOM   2302  C   SER   386       4.591 -21.125  15.978  1.00 26.58       B    C
ATOM   2303  O   SER   386       3.444 -20.948  15.572  1.00 27.11       B    O
ATOM   2304  N   GLN   387       5.104 -20.443  16.998  1.00 25.64       B    N
ATOM   2305  CA  GLN   387       4.283 -19.532  17.780  1.00 23.63       B    C
ATOM   2306  CB  GLN   387       5.064 -19.006  18.978  1.00 22.61       B    C
ATOM   2307  CG  GLN   387       6.153 -19.938  19.464  1.00 22.42       B    C
ATOM   2308  CD  GLN   387       5.615 -21.198  20.096  1.00 21.76       B    C
ATOM   2309  OE1 GLN   387       4.627 -21.162  20.839  1.00 21.92       B    O
ATOM   2310  NE2 GLN   387       6.263 -22.327  19.809  1.00 20.35       B    N
ATOM   2311  C   GLN   387       3.880 -18.378  16.886  1.00 23.78       B    C
ATOM   2312  O   GLN   387       2.777 -17.846  16.997  1.00 24.91       B    O
ATOM   2313  N   ALA   388       4.787 -18.002  15.992  1.00 23.75       B    N
ATOM   2314  CA  ALA   388       4.497 -17.002  14.972  1.00 23.70       B    C
ATOM   2315  CB  ALA   388       5.740 -16.768  14.094  1.00 23.02       B    C
ATOM   2316  C   ALA   388       3.319 -17.475  14.113  1.00 23.75       B    C
ATOM   2317  O   ALA   388       2.242 -16.873  14.129  1.00 24.11       B    O
ATOM   2318  N   ALA   389       3.529 -18.561  13.374  1.00 22.91       B    N
ATOM   2319  CA  ALA   389       2.503 -19.106  12.498  1.00 21.81       B    C
ATOM   2320  CB  ALA   389       2.887 -20.521  12.086  1.00 19.84       B    C
ATOM   2321  C   ALA   389       1.123 -19.094  13.172  1.00 22.14       B    C
ATOM   2322  O   ALA   389       0.135 -18.644  12.584  1.00 22.22       B    O
ATOM   2323  N   ALA   390       1.057 -19.570  14.410  1.00 22.17       B    N
ATOM   2324  CA  ALA   390      -0.215 -19.627  15.123  1.00 22.04       B    C
ATOM   2325  CB  ALA   390      -0.026 -20.241  16.506  1.00 22.25       B    C
ATOM   2326  C   ALA   390      -0.843 -18.241  15.249  1.00 21.51       B    C
ATOM   2327  O   ALA   390      -2.068 -18.099  15.179  1.00 21.52       B    O
ATOM   2328  N   HIS   391      -0.012 -17.219  15.443  1.00 20.10       B    N
ATOM   2329  CA  HIS   391      -0.526 -15.858  15.538  1.00 18.46       B    C
ATOM   2330  CB  HIS   391       0.615 -14.857  15.771  1.00 19.36       B    C
ATOM   2331  CG  HIS   391       0.912 -14.586  17.214  1.00 20.07       B    C
ATOM   2332  CD2 HIS   391       0.636 -13.513  17.995  1.00 19.81       B    C
ATOM   2333  ND1 HIS   391       1.615 -15.467  18.010  1.00 18.44       B    N
ATOM   2334  CE1 HIS   391       1.761 -14.947  19.216  1.00 18.95       B    C
ATOM   2335  NE2 HIS   391       1.177 -13.762  19.234  1.00 18.77       B    N
ATOM   2336  C   HIS   391      -1.241 -15.527  14.234  1.00 16.67       B    C
ATOM   2337  O   HIS   391      -2.274 -14.852  14.222  1.00 15.39       B    O
ATOM   2338  N   VAL   392      -0.688 -16.027  13.136  1.00 15.23       B    N
ATOM   2339  CA  VAL   392      -1.146 -15.632  11.815  1.00 14.91       B    C
ATOM   2340  CB  VAL   392       0.007 -15.768  10.783  1.00 14.04       B    C
ATOM   2341  CG1 VAL   392      -0.475 -15.429   9.382  1.00 12.46       B    C
ATOM   2342  CG2 VAL   392       1.143 -14.838  11.169  1.00 12.88       B    C
ATOM   2343  C   VAL   392      -2.369 -16.446  11.385  1.00 15.09       B    C
ATOM   2344  O   VAL   392      -3.269 -15.929  10.722  1.00 14.21       B    O
ATOM   2345  N   ALA   393      -2.421 -17.708  11.789  1.00 15.37       B    N
ATOM   2346  CA  ALA   393      -3.658 -18.467  11.655  1.00 17.74       B    C
ATOM   2347  CB  ALA   393      -3.453 -19.888  12.163  1.00 18.52       B    C
ATOM   2348  C   ALA   393      -4.791 -17.776  12.434  1.00 17.78       B    C
ATOM   2349  O   ALA   393      -5.936 -17.716  11.980  1.00 16.51       B    O
ATOM   2350  N   GLY   394      -4.453 -17.248  13.606  1.00 18.67       B    N
ATOM   2351  CA  GLY   394      -5.397 -16.444  14.359  1.00 19.56       B    C
ATOM   2352  C   GLY   394      -5.878 -15.212  13.613  1.00 20.36       B    C
ATOM   2353  O   GLY   394      -7.082 -14.986  13.501  1.00 21.22       B    O
ATOM   2354  N   ILE   395      -4.950 -14.414  13.096  1.00 20.24       B    N
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ATOM   2355  CA  ILE   395      -5.312 -13.195  12.384  1.00 20.11      B    C
ATOM   2356  CB  ILE   395      -4.060 -12.375  12.034  1.00 20.04      B    C
ATOM   2357  CG2 ILE   395      -4.449 -11.094  11.302  1.00 19.78      B    C
ATOM   2358  CG1 ILE   395      -3.301 -12.049  13.316  1.00 20.30      B    C
ATOM   2359  CD1 ILE   395      -2.049 -11.279  13.095  1.00 19.69      B    C
ATOM   2360  C   ILE   395      -6.077 -13.519  11.104  1.00 21.08      B    C
ATOM   2361  O   ILE   395      -6.974 -12.776  10.691  1.00 21.99      B    O
ATOM   2362  N   ALA   396      -5.732 -14.634  10.473  1.00 20.99      B    N
ATOM   2363  CA  ALA   396      -6.436 -15.039   9.267  1.00 20.56      B    C
ATOM   2364  CB  ALA   396      -5.770 -16.267   8.659  1.00 20.18      B    C
ATOM   2365  C   ALA   396      -7.885 -15.338   9.631  1.00 19.85      B    C
ATOM   2366  O   ALA   396      -8.809 -14.845   8.995  1.00 17.95      B    O
ATOM   2367  N   ALA   397      -8.071 -16.136  10.677  1.00 21.02      B    N
ATOM   2368  CA  ALA   397      -9.402 -16.485  11.154  1.00 21.16      B    C
ATOM   2369  CB  ALA   397      -9.305 -17.247  12.464  1.00 17.93      B    C
ATOM   2370  C   ALA   397     -10.244 -15.233  11.342  1.00 22.61      B    C
ATOM   2371  O   ALA   397     -11.378 -15.168  10.875  1.00 25.02      B    O
ATOM   2372  N   MET   398      -9.686 -14.231  12.014  1.00 23.67      B    N
ATOM   2373  CA  MET   398     -10.426 -13.008  12.282  1.00 23.20      B    C
ATOM   2374  CB  MET   398      -9.601 -12.084  13.167  1.00 24.40      B    C
ATOM   2375  CG  MET   398      -9.150 -12.733  14.457  1.00 29.13      B    C
ATOM   2376  SD  MET   398     -10.147 -12.270  15.879  1.00 35.63      B    S
ATOM   2377  CE  MET   398      -9.367 -10.694  16.357  1.00 31.39      B    C
ATOM   2378  C   MET   398     -10.784 -12.298  10.980  1.00 23.27      B    C
ATOM   2379  O   MET   398     -11.925 -11.889  10.782  1.00 22.16      B    O
ATOM   2380  N   MET   399      -9.815 -12.165  10.083  1.00 22.78      B    N
ATOM   2381  CA  MET   399     -10.059 -11.448   8.843  1.00 22.76      B    C
ATOM   2382  CB  MET   399      -8.741 -11.202   8.118  1.00 24.23      B    C
ATOM   2383  CG  MET   399      -7.755 -10.386   8.928  1.00 27.83      B    C
ATOM   2384  SD  MET   399      -6.151 -10.150   8.115  1.00 32.96      B    S
ATOM   2385  CE  MET   399      -6.680  -9.723   6.497  1.00 29.89      B    C
ATOM   2386  C   MET   399     -11.032 -12.195   7.930  1.00 22.15      B    C
ATOM   2387  O   MET   399     -11.828 -11.580   7.218  1.00 21.59      B    O
ATOM   2388  N   LEU   400     -10.975 -13.522   7.957  1.00 21.52      B    N
ATOM   2389  CA  LEU   400     -11.878 -14.320   7.140  1.00 20.75      B    C
ATOM   2390  CB  LEU   400     -11.349 -15.745   6.962  1.00 18.56      B    C
ATOM   2391  CG  LEU   400     -10.109 -15.930   6.085  1.00 16.86      B    C
ATOM   2392  CD1 LEU   400      -9.897 -17.411   5.826  1.00 13.64      B    C
ATOM   2393  CD2 LEU   400     -10.273 -15.166   4.770  1.00 15.30      B    C
ATOM   2394  C   LEU   400     -13.248 -14.379   7.782  1.00 22.24      B    C
ATOM   2395  O   LEU   400     -14.267 -14.449   7.092  1.00 23.47      B    O
ATOM   2396  N   SER   401     -13.284 -14.352   9.108  1.00 22.69      B    N
ATOM   2397  CA  SER   401     -14.566 -14.401   9.786  1.00 23.81      B    C
ATOM   2398  CB  SER   401     -14.375 -14.687  11.269  1.00 25.01      B    C
ATOM   2399  OG  SER   401     -15.532 -15.304  11.804  1.00 25.44      B    O
ATOM   2400  C   SER   401     -15.288 -13.080   9.596  1.00 23.85      B    C
ATOM   2401  O   SER   401     -16.508 -13.006   9.696  1.00 23.69      B    O
ATOM   2402  N   ALA   402     -14.523 -12.035   9.310  1.00 25.13      B    N
ATOM   2403  CA  ALA   402     -15.106 -10.745   9.016  1.00 25.64      B    C
ATOM   2404  CB  ALA   402     -14.124  -9.663   9.291  1.00 23.94      B    C
ATOM   2405  C   ALA   402     -15.531 -10.718   7.560  1.00 28.07      B    C
ATOM   2406  O   ALA   402     -16.724 -10.730   7.279  1.00 31.96      B    O
ATOM   2407  N   GLU   403     -14.576 -10.696   6.631  1.00 28.94      B    N
ATOM   2408  CA  GLU   403     -14.920 -10.756   5.204  1.00 29.12      B    C
ATOM   2409  CB  GLU   403     -14.104  -9.769   4.382  1.00 29.99      B    C
ATOM   2410  CG  GLU   403     -13.762  -8.509   5.092  1.00 33.22      B    C
ATOM   2411  CD  GLU   403     -12.300  -8.220   4.979  1.00 36.14      B    C
ATOM   2412  OE1 GLU   403     -11.906  -7.640   3.941  1.00 37.63      B    O
ATOM   2413  OE2 GLU   403     -11.547  -8.584   5.918  1.00 38.88      B    O
ATOM   2414  C   GLU   403     -14.652 -12.140   4.657  1.00 27.95      B    C
ATOM   2415  O   GLU   403     -13.554 -12.428   4.180  1.00 26.83      B    O
ATOM   2416  N   PRO   404     -15.661 -13.015   4.719  1.00 26.86      B    N
ATOM   2417  CD  PRO   404     -16.921 -12.782   5.442  1.00 25.89      B    C
ATOM   2418  CA  PRO   404     -15.499 -14.423   4.362  1.00 25.99      B    C
ATOM   2419  CB  PRO   404     -16.769 -15.073   4.911  1.00 25.93      B    C
ATOM   2420  CG  PRO   404     -17.237 -14.135   5.983  1.00 25.26      B    C
ATOM   2421  C   PRO   404     -15.296 -14.682   2.868  1.00 25.55      B    C
ATOM   2422  O   PRO   404     -14.748 -15.718   2.479  1.00 26.48      B    O
ATOM   2423  N   GLU   405     -15.727 -13.747   2.029  1.00 25.09      B    N
ATOM   2424  CA  GLU   405     -15.601 -13.931   0.588  1.00 25.06      B    C
ATOM   2425  CB  GLU   405     -16.583 -13.020  -0.153  1.00 27.09      B    C
ATOM   2426  CG  GLU   405     -18.044 -13.438  -0.031  1.00 29.66      B    C
ATOM   2427  CD  GLU   405     -19.001 -12.496  -0.752  1.00 30.19      B    C
ATOM   2428  OE1 GLU   405     -20.212 -12.786  -0.767  1.00 31.61      B    O
ATOM   2429  OE2 GLU   405     -18.553 -11.471  -1.303  1.00 30.56      B    O
ATOM   2430  C   GLU   405     -14.188 -13.693   0.056  1.00 24.86      B    C
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ATOM	2431	O	GLU	405	-13.948	-13.872	-1.132	1.00	24.99	B	O
ATOM	2432	N	LEU	406	-13.256	-13.295	0.922	1.00	24.30	B	N
ATOM	2433	CA	LEU	406	-11.870	-13.043	0.505	1.00	23.76	B	C
ATOM	2434	CB	LEU	406	-11.000	-12.639	1.701	1.00	21.49	B	C
ATOM	2435	CG	LEU	406	-11.111	-11.210	2.222	1.00	20.80	B	C
ATOM	2436	CD1	LEU	406	-10.269	-11.069	3.464	1.00	18.92	B	C
ATOM	2437	CD2	LEU	406	-10.664	-10.233	1.156	1.00	18.73	B	C
ATOM	2438	C	LEU	406	-11.234	-14.263	-0.146	1.00	23.84	B	C
ATOM	2439	O	LEU	406	-11.275	-15.358	0.411	1.00	25.23	B	O
ATOM	2440	N	THR	407	-10.633	-14.067	-1.315	1.00	23.77	B	N
ATOM	2441	CA	THR	407	-9.781	-15.087	-1.918	1.00	23.47	B	C
ATOM	2442	CB	THR	407	-9.533	-14.794	-3.395	1.00	21.81	B	C
ATOM	2443	OG1	THR	407	-8.682	-13.647	-3.517	1.00	23.22	B	O
ATOM	2444	CG2	THR	407	-10.827	-14.507	-4.089	1.00	19.79	B	C
ATOM	2445	C	THR	407	-8.419	-15.138	-1.213	1.00	24.96	B	C
ATOM	2446	O	THR	407	-8.081	-14.268	-0.399	1.00	26.81	B	O
ATOM	2447	N	LEU	408	-7.630	-16.154	-1.535	1.00	24.30	B	N
ATOM	2448	CA	LEU	408	-6.312	-16.292	-0.929	1.00	22.60	B	C
ATOM	2449	CB	LEU	408	-5.678	-17.616	-1.355	1.00	22.93	B	C
ATOM	2450	CG	LEU	408	-4.472	-18.043	-0.525	1.00	24.47	B	C
ATOM	2451	CD1	LEU	408	-4.734	-17.740	0.940	1.00	23.99	B	C
ATOM	2452	CD2	LEU	408	-4.211	-19.529	-0.732	1.00	25.90	B	C
ATOM	2453	C	LEU	408	-5.421	-15.131	-1.344	1.00	20.03	B	C
ATOM	2454	O	LEU	408	-4.552	-14.697	-0.592	1.00	16.95	B	O
ATOM	2455	N	ALA	409	-5.651	-14.636	-2.553	1.00	19.32	B	N
ATOM	2456	CA	ALA	409	-4.906	-13.506	-3.065	1.00	19.43	B	C
ATOM	2457	CB	ALA	409	-5.280	-13.249	-4.499	1.00	20.13	B	C
ATOM	2458	C	ALA	409	-5.248	-12.301	-2.220	1.00	20.61	B	C
ATOM	2459	O	ALA	409	-4.368	-11.579	-1.750	1.00	20.75	B	O
ATOM	2460	N	GLU	410	-6.540	-12.092	-2.012	1.00	21.62	B	N
ATOM	2461	CA	GLU	410	-6.976	-10.967	-1.212	1.00	22.93	B	C
ATOM	2462	CB	GLU	410	-8.480	-10.800	-1.324	1.00	25.38	B	C
ATOM	2463	CG	GLU	410	-8.903	-10.285	-2.661	1.00	30.96	B	C
ATOM	2464	CD	GLU	410	-10.208	-10.888	-3.125	1.00	34.21	B	C
ATOM	2465	OE1	GLU	410	-10.441	-10.877	-4.355	1.00	37.17	B	O
ATOM	2466	OE2	GLU	410	-10.993	-11.371	-2.269	1.00	35.26	B	O
ATOM	2467	C	GLU	410	-6.583	-11.119	0.247	1.00	22.94	B	C
ATOM	2468	O	GLU	410	-6.332	-10.128	0.925	1.00	23.23	B	O
ATOM	2469	N	LEU	411	-6.532	-12.352	0.738	1.00	22.70	B	N
ATOM	2470	CA	LEU	411	-6.175	-12.552	2.129	1.00	22.90	B	C
ATOM	2471	CB	LEU	411	-6.387	-13.998	2.542	1.00	20.63	B	C
ATOM	2472	CG	LEU	411	-5.995	-14.204	4.009	1.00	20.39	B	C
ATOM	2473	CD1	LEU	411	-6.881	-13.338	4.906	1.00	17.47	B	C
ATOM	2474	CD2	LEU	411	-6.123	-15.670	4.375	1.00	18.76	B	C
ATOM	2475	C	LEU	411	-4.722	-12.174	2.373	1.00	24.89	B	C
ATOM	2476	O	LEU	411	-4.369	-11.657	3.438	1.00	24.07	B	O
ATOM	2477	N	ARG	412	-3.877	-12.438	1.383	1.00	27.02	B	N
ATOM	2478	CA	ARG	412	-2.471	-12.099	1.496	1.00	29.77	B	C
ATOM	2479	CB	ARG	412	-1.683	-12.716	0.354	1.00	30.56	B	C
ATOM	2480	CG	ARG	412	-1.384	-14.172	0.527	1.00	32.41	B	C
ATOM	2481	CD	ARG	412	-0.517	-14.632	-0.620	1.00	36.01	B	C
ATOM	2482	NE	ARG	412	0.598	-15.444	-0.152	1.00	38.01	B	N
ATOM	2483	CZ	ARG	412	0.587	-16.768	-0.136	1.00	38.92	B	C
ATOM	2484	NH1	ARG	412	1.643	-17.437	0.308	1.00	38.83	B	N
ATOM	2485	NH2	ARG	412	-0.485	-17.418	-0.569	1.00	38.86	B	N
ATOM	2486	C	ARG	412	-2.327	-10.599	1.436	1.00	30.73	B	C
ATOM	2487	O	ARG	412	-1.628	-9.994	2.248	1.00	30.38	B	O
ATOM	2488	N	GLN	413	-3.004	-10.005	0.461	1.00	33.11	B	N
ATOM	2489	CA	GLN	413	-2.895	-8.581	0.221	1.00	35.39	B	C
ATOM	2490	CB	GLN	413	-3.779	-8.176	-0.948	1.00	38.60	B	C
ATOM	2491	CG	GLN	413	-3.344	-6.875	-1.596	1.00	45.86	B	C
ATOM	2492	CD	GLN	413	-1.860	-6.872	-1.965	1.00	49.77	B	C
ATOM	2493	OE1	GLN	413	-1.485	-7.150	-3.115	1.00	52.54	B	O
ATOM	2494	NE2	GLN	413	-1.008	-6.552	-0.989	1.00	49.77	B	N
ATOM	2495	C	GLN	413	-3.301	-7.805	1.456	1.00	35.24	B	C
ATOM	2496	O	GLN	413	-2.824	-6.696	1.682	1.00	36.49	B	O
ATOM	2497	N	ARG	414	-4.175	-8.405	2.256	1.00	35.12	B	N
ATOM	2498	CA	ARG	414	-4.643	-7.806	3.500	1.00	34.96	B	C
ATOM	2499	CB	ARG	414	-5.975	-8.422	3.901	1.00	36.59	B	C
ATOM	2500	CG	ARG	414	-7.128	-7.989	3.043	1.00	39.95	B	C
ATOM	2501	CD	ARG	414	-8.229	-7.397	3.892	1.00	42.16	B	C
ATOM	2502	NE	ARG	414	-9.264	-6.704	3.125	1.00	44.38	B	N
ATOM	2503	CZ	ARG	414	-9.560	-6.929	1.846	1.00	46.36	B	C
ATOM	2504	NH1	ARG	414	-8.904	-7.835	1.136	1.00	48.71	B	N
ATOM	2505	NH2	ARG	414	-10.541	-6.251	1.274	1.00	47.81	B	N
ATOM	2506	C	ARG	414	-3.657	-7.972	4.646	1.00	34.00	B	C

ATOM	2507	O	ARG	414	-3.252	-6.992	5.270	1.00	34.78	B	O
ATOM	2508	N	LEU	415	-3.278	-9.211	4.931	1.00	32.40	B	N
ATOM	2509	CA	LEU	415	-2.233	-9.466	5.912	1.00	31.47	B	C
ATOM	2510	CB	LEU	415	-1.689	-10.873	5.748	1.00	31.21	B	C
ATOM	2511	CG	LEU	415	-2.574	-11.951	6.346	1.00	31.92	B	C
ATOM	2512	CD1	LEU	415	-2.230	-13.274	5.697	1.00	33.02	B	C
ATOM	2513	CD2	LEU	415	-2.395	-11.990	7.862	1.00	30.21	B	C
ATOM	2514	C	LEU	415	-1.091	-8.479	5.750	1.00	31.04	B	C
ATOM	2515	O	LEU	415	-0.679	-7.823	6.711	1.00	31.86	B	O
ATOM	2516	N	ILE	416	-0.580	-8.380	4.530	1.00	29.96	B	N
ATOM	2517	CA	ILE	416	0.423	-7.378	4.230	1.00	30.82	B	C
ATOM	2518	CB	ILE	416	0.655	-7.241	2.716	1.00	30.83	B	C
ATOM	2519	CG2	ILE	416	1.444	-5.977	2.434	1.00	29.81	B	C
ATOM	2520	CG1	ILE	416	1.374	-8.480	2.178	1.00	29.19	B	C
ATOM	2521	CD1	ILE	416	1.488	-8.503	0.678	1.00	25.78	B	C
ATOM	2522	C	ILE	416	-0.046	-6.031	4.743	1.00	30.89	B	C
ATOM	2523	O	ILE	416	0.545	-5.459	5.663	1.00	31.33	B	O
ATOM	2524	N	HIS	417	-1.130	-5.548	4.145	1.00	31.24	B	N
ATOM	2525	CA	HIS	417	-1.556	-4.166	4.300	1.00	30.14	B	C
ATOM	2526	CB	HIS	417	-2.812	-3.907	3.469	1.00	30.66	B	C
ATOM	2527	CG	HIS	417	-3.393	-2.548	3.679	1.00	32.44	B	C
ATOM	2528	CD2	HIS	417	-2.908	-1.318	3.388	1.00	33.19	B	C
ATOM	2529	ND1	HIS	417	-4.597	-2.345	4.319	1.00	31.59	B	N
ATOM	2530	CE1	HIS	417	-4.828	-1.048	4.417	1.00	32.26	B	C
ATOM	2531	NE2	HIS	417	-3.818	-0.403	3.860	1.00	33.69	B	N
ATOM	2532	C	HIS	417	-1.811	-3.789	5.750	1.00	29.15	B	C
ATOM	2533	O	HIS	417	-1.646	-2.629	6.122	1.00	29.16	B	O
ATOM	2534	N	PHE	418	-2.210	-4.761	6.568	1.00	27.70	B	N
ATOM	2535	CA	PHE	418	-2.394	-4.512	7.992	1.00	27.83	B	C
ATOM	2536	CB	PHE	418	-3.595	-5.285	8.534	1.00	28.70	B	C
ATOM	2537	CG	PHE	418	-4.895	-4.754	8.058	1.00	29.85	B	C
ATOM	2538	CD1	PHE	418	-5.677	-5.485	7.185	1.00	29.88	B	C
ATOM	2539	CD2	PHE	418	-5.285	-3.473	8.393	1.00	30.40	B	C
ATOM	2540	CE1	PHE	418	-6.822	-4.945	6.651	1.00	29.59	B	C
ATOM	2541	CE2	PHE	418	-6.426	-2.929	7.863	1.00	31.18	B	C
ATOM	2542	CZ	PHE	418	-7.195	-3.661	6.986	1.00	30.31	B	C
ATOM	2543	C	PHE	418	-1.164	-4.884	8.777	1.00	27.12	B	C
ATOM	2544	O	PHE	418	-1.177	-4.895	10.005	1.00	26.90	B	O
ATOM	2545	N	SER	419	-0.094	-5.201	8.070	1.00	26.74	B	N
ATOM	2546	CA	SER	419	1.154	-5.463	8.748	1.00	26.69	B	C
ATOM	2547	CB	SER	419	2.006	-6.431	7.931	1.00	25.86	B	C
ATOM	2548	OG	SER	419	1.593	-7.759	8.188	1.00	25.35	B	O
ATOM	2549	C	SER	419	1.905	-4.165	8.983	1.00	26.68	B	C
ATOM	2550	O	SER	419	1.879	-3.262	8.144	1.00	26.77	B	O
ATOM	2551	N	ALA	420	2.544	-4.079	10.147	1.00	25.25	B	N
ATOM	2552	CA	ALA	420	3.554	-3.072	10.412	1.00	24.13	B	C
ATOM	2553	CB	ALA	420	4.069	-3.215	11.816	1.00	24.79	B	C
ATOM	2554	C	ALA	420	4.675	-3.321	9.439	1.00	24.87	B	C
ATOM	2555	O	ALA	420	5.089	-4.466	9.238	1.00	26.34	B	O
ATOM	2556	N	LYS	421	5.172	-2.252	8.838	1.00	24.32	B	N
ATOM	2557	CA	LYS	421	6.132	-2.383	7.760	1.00	25.67	B	C
ATOM	2558	CB	LYS	421	5.557	-1.747	6.497	1.00	23.67	B	C
ATOM	2559	CG	LYS	421	4.124	-2.161	6.249	1.00	23.87	B	C
ATOM	2560	CD	LYS	421	3.716	-2.007	4.797	1.00	23.84	B	C
ATOM	2561	CE	LYS	421	2.331	-2.614	4.550	1.00	23.52	B	C
ATOM	2562	NZ	LYS	421	1.363	-2.280	5.638	1.00	22.85	B	N
ATOM	2563	C	LYS	421	7.468	-1.743	8.125	1.00	27.41	B	C
ATOM	2564	O	LYS	421	7.510	-0.712	8.798	1.00	28.11	B	O
ATOM	2565	N	ASP	422	8.562	-2.361	7.691	1.00	28.06	B	N
ATOM	2566	CA	ASP	422	9.873	-1.745	7.844	1.00	27.60	B	C
ATOM	2567	CB	ASP	422	9.974	-0.503	6.955	1.00	26.92	B	C
ATOM	2568	CG	ASP	422	10.080	-0.856	5.489	1.00	28.56	B	C
ATOM	2569	OD1	ASP	422	10.866	-1.768	5.161	1.00	28.83	B	O
ATOM	2570	OD2	ASP	422	9.381	-0.234	4.663	1.00	29.17	B	O
ATOM	2571	C	ASP	422	10.124	-1.368	9.296	1.00	26.84	B	C
ATOM	2572	O	ASP	422	10.463	-0.227	9.607	1.00	26.44	B	O
ATOM	2573	N	VAL	423	9.944	-2.335	10.182	1.00	25.89	B	N
ATOM	2574	CA	VAL	423	10.146	-2.100	11.599	1.00	26.20	B	C
ATOM	2575	CB	VAL	423	8.831	-2.292	12.396	1.00	27.67	B	C
ATOM	2576	CG1	VAL	423	7.775	-1.320	11.900	1.00	27.74	B	C
ATOM	2577	CG2	VAL	423	8.331	-3.723	12.240	1.00	28.31	B	C
ATOM	2578	C	VAL	423	11.158	-3.110	12.075	1.00	25.50	B	C
ATOM	2579	O	VAL	423	11.697	-3.006	13.168	1.00	25.97	B	O
ATOM	2580	N	ILE	424	11.415	-4.101	11.239	1.00	25.87	B	N
ATOM	2581	CA	ILE	424	12.307	-5.171	11.630	1.00	26.65	B	C
ATOM	2582	CB	ILE	424	11.960	-6.471	10.893	1.00	26.26	B	C

ATOM	2583	CG2	ILE	424	12.495	-7.652	11.674	1.00	27.62	B	C
ATOM	2584	CG1	ILE	424	10.446	-6.635	10.784	1.00	27.06	B	C
ATOM	2585	CD1	ILE	424	10.016	-7.957	10.169	1.00	26.12	B	C
ATOM	2586	C	ILE	424	13.759	-4.800	11.329	1.00	27.01	B	C
ATOM	2587	O	ILE	424	14.072	-4.233	10.274	1.00	26.17	B	O
ATOM	2588	N	ASN	425	14.642	-5.128	12.268	1.00	28.05	B	N
ATOM	2589	CA	ASN	425	16.065	-4.869	12.107	1.00	29.38	B	C
ATOM	2590	CB	ASN	425	16.741	-4.831	13.475	1.00	31.69	B	C
ATOM	2591	CG	ASN	425	18.215	-4.485	13.385	1.00	34.14	B	C
ATOM	2592	OD1	ASN	425	18.782	-4.397	12.292	1.00	35.11	B	O
ATOM	2593	ND2	ASN	425	18.846	-4.287	14.538	1.00	34.72	B	N
ATOM	2594	C	ASN	425	16.729	-5.933	11.241	1.00	28.62	B	C
ATOM	2595	O	ASN	425	16.973	-7.045	11.697	1.00	29.23	B	O
ATOM	2596	N	GLU	426	17.038	-5.576	10.001	1.00	28.49	B	N
ATOM	2597	CA	GLU	426	17.513	-6.536	9.015	1.00	29.63	B	C
ATOM	2598	CB	GLU	426	17.266	-5.985	7.624	1.00	31.63	B	C
ATOM	2599	CG	GLU	426	15.815	-5.841	7.266	1.00	35.09	B	C
ATOM	2600	CD	GLU	426	15.653	-5.209	5.905	1.00	38.74	B	C
ATOM	2601	OE1	GLU	426	14.584	-5.376	5.271	1.00	41.88	B	O
ATOM	2602	OE2	GLU	426	16.615	-4.541	5.466	1.00	40.95	B	O
ATOM	2603	C	GLU	426	18.991	-6.913	9.148	1.00	29.25	B	C
ATOM	2604	O	GLU	426	19.488	-7.779	8.432	1.00	28.10	B	O
ATOM	2605	N	ALA	427	19.688	-6.258	10.065	1.00	30.10	B	N
ATOM	2606	CA	ALA	427	21.111	-6.486	10.263	1.00	29.38	B	C
ATOM	2607	CB	ALA	427	21.607	-5.637	11.417	1.00	28.99	B	C
ATOM	2608	C	ALA	427	21.426	-7.947	10.525	1.00	29.94	B	C
ATOM	2609	O	ALA	427	22.500	-8.422	10.165	1.00	29.35	B	O
ATOM	2610	N	TRP	428	20.497	-8.661	11.153	1.00	31.00	B	N
ATOM	2611	CA	TRP	428	20.745	-10.045	11.540	1.00	31.86	B	C
ATOM	2612	CB	TRP	428	19.724	-10.484	12.597	1.00	34.20	B	C
ATOM	2613	CG	TRP	428	20.016	-11.842	13.223	1.00	37.61	B	C
ATOM	2614	CD2	TRP	428	19.340	-13.084	12.952	1.00	39.01	B	C
ATOM	2615	CE2	TRP	428	19.939	-14.073	13.767	1.00	40.16	B	C
ATOM	2616	CE3	TRP	428	18.287	-13.452	12.106	1.00	38.02	B	C
ATOM	2617	CD1	TRP	428	20.973	-12.129	14.161	1.00	38.36	B	C
ATOM	2618	NE1	TRP	428	20.932	-13.467	14.491	1.00	38.65	B	N
ATOM	2619	CZ2	TRP	428	19.521	-15.409	13.750	1.00	40.19	B	C
ATOM	2620	CZ3	TRP	428	17.875	-14.776	12.092	1.00	38.79	B	C
ATOM	2621	CH2	TRP	428	18.487	-15.737	12.911	1.00	39.60	B	C
ATOM	2622	C	TRP	428	20.731	-11.020	10.353	1.00	31.48	B	C
ATOM	2623	O	TRP	428	21.473	-12.003	10.346	1.00	31.38	B	O
ATOM	2624	N	PHE	429	19.898	-10.763	9.348	1.00	31.09	B	N
ATOM	2625	CA	PHE	429	19.884	-11.621	8.167	1.00	31.73	B	C
ATOM	2626	CB	PHE	429	18.752	-11.254	7.220	1.00	31.93	B	C
ATOM	2627	CG	PHE	429	17.416	-11.186	7.876	1.00	33.61	B	C
ATOM	2628	CD1	PHE	429	16.565	-10.120	7.621	1.00	32.81	B	C
ATOM	2629	CD2	PHE	429	17.011	-12.182	8.753	1.00	31.99	B	C
ATOM	2630	CE1	PHE	429	15.341	-10.047	8.233	1.00	33.79	B	C
ATOM	2631	CE2	PHE	429	15.789	-12.116	9.367	1.00	32.86	B	C
ATOM	2632	CZ	PHE	429	14.946	-11.046	9.108	1.00	33.20	B	C
ATOM	2633	C	PHE	429	21.179	-11.428	7.421	1.00	33.05	B	C
ATOM	2634	O	PHE	429	21.792	-10.366	7.504	1.00	34.48	B	O
ATOM	2635	N	PRO	430	21.610	-12.444	6.662	1.00	33.68	B	N
ATOM	2636	CD	PRO	430	21.130	-13.829	6.543	1.00	34.28	B	C
ATOM	2637	CA	PRO	430	22.700	-12.155	5.736	1.00	34.21	B	C
ATOM	2638	CB	PRO	430	22.977	-13.501	5.066	1.00	33.97	B	C
ATOM	2639	CG	PRO	430	21.737	-14.282	5.249	1.00	34.13	B	C
ATOM	2640	C	PRO	430	22.294	-11.058	4.750	1.00	35.03	B	C
ATOM	2641	O	PRO	430	21.112	-10.832	4.497	1.00	35.24	B	O
ATOM	2642	N	GLU	431	23.293	-10.368	4.214	1.00	36.46	B	N
ATOM	2643	CA	GLU	431	23.079	-9.141	3.475	1.00	36.94	B	C
ATOM	2644	CB	GLU	431	24.409	-8.645	2.928	1.00	38.04	B	C
ATOM	2645	CG	GLU	431	25.515	-8.637	3.950	1.00	39.87	B	C
ATOM	2646	CD	GLU	431	26.828	-8.187	3.358	1.00	42.43	B	C
ATOM	2647	OE1	GLU	431	26.842	-7.845	2.152	1.00	42.43	B	O
ATOM	2648	OE2	GLU	431	27.842	-8.171	4.098	1.00	43.52	B	O
ATOM	2649	C	GLU	431	22.125	-9.374	2.328	1.00	37.62	B	C
ATOM	2650	O	GLU	431	21.094	-8.710	2.205	1.00	38.50	B	O
ATOM	2651	N	ASP	432	22.478	-10.327	1.482	1.00	37.53	B	N
ATOM	2652	CA	ASP	432	21.711	-10.562	0.280	1.00	37.71	B	C
ATOM	2653	CB	ASP	432	22.396	-11.627	-0.560	1.00	41.08	B	C
ATOM	2654	CG	ASP	432	23.813	-11.258	-0.898	1.00	43.17	B	C
ATOM	2655	OD1	ASP	432	24.061	-10.058	-1.146	1.00	45.66	B	O
ATOM	2656	OD2	ASP	432	24.675	-12.159	-0.909	1.00	44.70	B	O
ATOM	2657	C	ASP	432	20.290	-10.982	0.598	1.00	35.78	B	C
ATOM	2658	O	ASP	432	19.410	-10.932	-0.264	1.00	37.15	B	O

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ATOM   2659  N   GLN   433      20.060 -11.395   1.835  1.00 32.85      B    N
ATOM   2660  CA  GLN   433      18.749 -11.887   2.205  1.00 31.54      B    C
ATOM   2661  CB  GLN   433      18.879 -12.932   3.305  1.00 29.75      B    C
ATOM   2662  CG  GLN   433      19.423 -14.245   2.794  1.00 28.29      B    C
ATOM   2663  CD  GLN   433      18.459 -14.955   1.850  1.00 26.47      B    C
ATOM   2664  OE1 GLN   433      18.828 -15.919   1.184  1.00 24.27      B    O
ATOM   2665  NE2 GLN   433      17.221 -14.481   1.794  1.00 26.30      B    N
ATOM   2666  C   GLN   433      17.823 -10.769   2.645  1.00 31.04      B    C
ATOM   2667  O   GLN   433      16.623 -10.974   2.805  1.00 32.65      B    O
ATOM   2668  N   ARG   434      18.373  -9.576   2.819  1.00 30.10      B    N
ATOM   2669  CA  ARG   434      17.588  -8.474   3.343  1.00 28.71      B    C
ATOM   2670  CB  ARG   434      18.518  -7.369   3.845  1.00 29.04      B    C
ATOM   2671  CG  ARG   434      19.390  -7.826   5.012  1.00 28.41      B    C
ATOM   2672  CD  ARG   434      20.174  -6.695   5.646  1.00 26.84      B    C
ATOM   2673  NE  ARG   434      21.266  -7.222   6.459  1.00 27.43      B    N
ATOM   2674  CZ  ARG   434      22.538  -6.869   6.315  1.00 25.70      B    C
ATOM   2675  NH1 ARG   434      23.473  -7.399   7.088  1.00 23.11      B    N
ATOM   2676  NH2 ARG   434      22.869  -5.978   5.398  1.00 25.28      B    N
ATOM   2677  C   ARG   434      16.569  -7.912   2.361  1.00 27.60      B    C
ATOM   2678  O   ARG   434      15.431  -7.664   2.747  1.00 27.25      B    O
ATOM   2679  N   VAL   435      16.951  -7.711   1.101  1.00 27.74      B    N
ATOM   2680  CA  VAL   435      15.975  -7.213   0.128  1.00 27.86      B    C
ATOM   2681  CB  VAL   435      16.599  -6.710  -1.199  1.00 26.45      B    C
ATOM   2682  CG1 VAL   435      17.515  -5.547  -0.943  1.00 27.10      B    C
ATOM   2683  CG2 VAL   435      17.315  -7.832  -1.890  1.00 27.00      B    C
ATOM   2684  C   VAL   435      14.992  -8.302  -0.237  1.00 28.30      B    C
ATOM   2685  O   VAL   435      13.890  -8.011  -0.709  1.00 30.16      B    O
ATOM   2686  N   LEU   436      15.389  -9.554  -0.022  1.00 27.85      B    N
ATOM   2687  CA  LEU   436      14.511 -10.686  -0.307  1.00 27.26      B    C
ATOM   2688  CB  LEU   436      15.312 -11.975  -0.467  1.00 27.16      B    C
ATOM   2689  CG  LEU   436      16.446 -12.004  -1.478  1.00 25.55      B    C
ATOM   2690  CD1 LEU   436      16.995 -13.421  -1.511  1.00 24.68      B    C
ATOM   2691  CD2 LEU   436      15.952 -11.565  -2.849  1.00 22.51      B    C
ATOM   2692  C   LEU   436      13.500 -10.894   0.808  1.00 27.27      B    C
ATOM   2693  O   LEU   436      12.351 -11.262   0.551  1.00 28.79      B    O
ATOM   2694  N   THR   437      13.936 -10.675   2.046  1.00 25.82      B    N
ATOM   2695  CA  THR   437      13.075 -10.904   3.199  1.00 23.86      B    C
ATOM   2696  CB  THR   437      13.887 -11.073   4.494  1.00 23.41      B    C
ATOM   2697  OG1 THR   437      14.782 -12.185   4.362  1.00 23.12      B    O
ATOM   2698  CG2 THR   437      12.949 -11.313   5.668  1.00 22.10      B    C
ATOM   2699  C   THR   437      12.090  -9.759   3.397  1.00 22.78      B    C
ATOM   2700  O   THR   437      12.464  -8.589   3.392  1.00 22.51      B    O
ATOM   2701  N   PRO   438      10.807 -10.095   3.566  1.00 23.25      B    N
ATOM   2702  CD  PRO   438      10.282 -11.469   3.511  1.00 24.11      B    C
ATOM   2703  CA  PRO   438       9.731  -9.118   3.742  1.00 22.75      B    C
ATOM   2704  CB  PRO   438       8.453  -9.957   3.639  1.00 23.28      B    C
ATOM   2705  CG  PRO   438       8.880 -11.258   3.049  1.00 23.27      B    C
ATOM   2706  C   PRO   438       9.847  -8.448   5.099  1.00 21.97      B    C
ATOM   2707  O   PRO   438       9.940  -9.124   6.125  1.00 19.45      B    O
ATOM   2708  N   ASN   439       9.837  -7.121   5.103  1.00 22.81      B    N
ATOM   2709  CA  ASN   439       9.866  -6.388   6.353  1.00 23.40      B    C
ATOM   2710  CB  ASN   439      10.627  -5.085   6.168  1.00 23.33      B    C
ATOM   2711  CG  ASN   439      11.355  -4.657   7.424  1.00 25.22      B    C
ATOM   2712  OD1 ASN   439      10.869  -4.854   8.538  1.00 26.16      B    O
ATOM   2713  ND2 ASN   439      12.533  -4.066   7.252  1.00 24.91      B    N
ATOM   2714  C   ASN   439       8.434  -6.118   6.820  1.00 24.31      B    C
ATOM   2715  O   ASN   439       7.949  -4.983   6.781  1.00 25.14      B    O
ATOM   2716  N   LEU   440       7.771  -7.186   7.256  1.00 24.30      B    N
ATOM   2717  CA  LEU   440       6.378  -7.145   7.690  1.00 25.09      B    C
ATOM   2718  CB  LEU   440       5.470  -7.703   6.589  1.00 23.69      B    C
ATOM   2719  CG  LEU   440       5.503  -6.977   5.247  1.00 24.08      B    C
ATOM   2720  CD1 LEU   440       4.757  -7.796   4.220  1.00 21.26      B    C
ATOM   2721  CD2 LEU   440       4.883  -5.584   5.401  1.00 23.50      B    C
ATOM   2722  C   LEU   440       6.163  -7.965   8.966  1.00 25.32      B    C
ATOM   2723  O   LEU   440       6.447  -9.164   9.007  1.00 24.66      B    O
ATOM   2724  N   VAL   441       5.647  -7.323  10.006  1.00 25.82      B    N
ATOM   2725  CA  VAL   441       5.096  -8.067  11.128  1.00 25.14      B    C
ATOM   2726  CB  VAL   441       5.633  -7.548  12.479  1.00 23.66      B    C
ATOM   2727  CG1 VAL   441       5.249  -8.500  13.580  1.00 21.33      B    C
ATOM   2728  CG2 VAL   441       7.139  -7.385  12.421  1.00 23.17      B    C
ATOM   2729  C   VAL   441       3.581  -7.889  11.083  1.00 26.87      B    C
ATOM   2730  O   VAL   441       3.078  -6.806  10.773  1.00 28.65      B    O
ATOM   2731  N   ALA   442       2.851  -8.953  11.384  1.00 26.88      B    N
ATOM   2732  CA  ALA   442       1.415  -8.952  11.179  1.00 27.49      B    C
ATOM   2733  CB  ALA   442       0.938 -10.369  10.972  1.00 27.84      B    C
ATOM   2734  C   ALA   442       0.648  -8.313  12.329  1.00 28.01      B    C
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ATOM   2735  O   ALA   442       0.863  -8.649  13.490  1.00 27.44       B    O
ATOM   2736  N   ALA   443      -0.260  -7.399  11.996  1.00 29.53       B    N
ATOM   2737  CA  ALA   443      -1.259  -6.929  12.950  1.00 29.76       B    C
ATOM   2738  CB  ALA   443      -1.135  -5.433  13.149  1.00 28.70       B    C
ATOM   2739  C   ALA   443      -2.662  -7.265  12.467  1.00 30.98       B    C
ATOM   2740  O   ALA   443      -2.873  -7.532  11.282  1.00 30.38       B    O
ATOM   2741  N   LEU   444      -3.610  -7.253  13.402  1.00 32.91       B    N
ATOM   2742  CA  LEU   444      -5.037  -7.307  13.095  1.00 33.56       B    C
ATOM   2743  CB  LEU   444      -5.841  -7.636  14.348  1.00 29.03       B    C
ATOM   2744  CG  LEU   444      -5.624  -8.983  15.021  1.00 26.74       B    C
ATOM   2745  CD1 LEU   444      -6.078  -8.902  16.462  1.00 24.31       B    C
ATOM   2746  CD2 LEU   444      -6.383 -10.061  14.269  1.00 25.72       B    C
ATOM   2747  C   LEU   444      -5.503  -5.958  12.565  1.00 37.13       B    C
ATOM   2748  O   LEU   444      -4.915  -4.914  12.869  1.00 36.31       B    O
ATOM   2749  N   PRO   445      -6.578  -5.967  11.766  1.00 40.64       B    N
ATOM   2750  CD  PRO   445      -7.291  -7.151  11.267  1.00 40.97       B    C
ATOM   2751  CA  PRO   445      -7.138  -4.733  11.216  1.00 43.51       B    C
ATOM   2752  CB  PRO   445      -8.215  -5.217  10.252  1.00 41.99       B    C
ATOM   2753  CG  PRO   445      -7.890  -6.645   9.998  1.00 41.69       B    C
ATOM   2754  C   PRO   445      -7.727  -3.918  12.343  1.00 47.02       B    C
ATOM   2755  O   PRO   445      -8.502  -4.429  13.149  1.00 47.30       B    O
ATOM   2756  N   PRO   446      -7.357  -2.637  12.422  1.00 50.39       B    N
ATOM   2757  CD  PRO   446      -6.431  -1.908  11.538  1.00 52.20       B    C
ATOM   2758  CA  PRO   446      -7.984  -1.768  13.418  1.00 52.13       B    C
ATOM   2759  CB  PRO   446      -7.269  -0.429  13.234  1.00 52.77       B    C
ATOM   2760  CG  PRO   446      -6.736  -0.467  11.833  1.00 53.26       B    C
ATOM   2761  C   PRO   446      -9.471  -1.691  13.117  1.00 52.85       B    C
ATOM   2762  O   PRO   446     -10.279  -1.732  14.075  1.00 53.37       B    O
ATOM   2763  OXT PRO   446      -9.793  -1.603  11.911  1.00 53.26       B    O
TER    2764       PRO   446                                               B
ATOM   2765  CB  THR   294      -8.672 -12.634  38.449  1.00 44.50       EGFA C
ATOM   2766  OG1 THR   294      -8.921 -13.914  39.043  1.00 46.09       EGFA O
ATOM   2767  CG2 THR   294      -9.367 -12.555  37.098  1.00 44.95       EGFA C
ATOM   2768  C   THR   294      -6.483 -13.598  37.580  1.00 42.73       EGFA C
ATOM   2769  O   THR   294      -6.465 -14.703  38.105  1.00 42.87       EGFA O
ATOM   2770  N   THR   294      -6.854 -11.126  37.620  1.00 43.75       EGFA N
ATOM   2771  CA  THR   294      -7.135 -12.410  38.303  1.00 43.33       EGFA C
ATOM   2772  N   ASN   295      -5.945 -13.381  36.380  1.00 40.79       EGFA N
ATOM   2773  CA  ASN   295      -5.062 -14.384  35.774  1.00 38.54       EGFA C
ATOM   2774  CB  ASN   295      -5.516 -14.783  34.369  1.00 38.56       EGFA C
ATOM   2775  CG  ASN   295      -4.653 -15.897  33.767  1.00 38.98       EGFA C
ATOM   2776  OD1 ASN   295      -3.455 -15.979  34.023  1.00 39.16       EGFA O
ATOM   2777  ND2 ASN   295      -5.271 -16.757  32.961  1.00 39.03       EGFA N
ATOM   2778  C   ASN   295      -3.656 -13.823  35.711  1.00 37.61       EGFA C
ATOM   2779  O   ASN   295      -3.211 -13.284  34.693  1.00 35.61       EGFA O
ATOM   2780  N   GLU   296      -2.965 -13.993  36.828  1.00 36.47       EGFA N
ATOM   2781  CA  GLU   296      -1.734 -13.293  37.119  1.00 36.71       EGFA C
ATOM   2782  CB  GLU   296      -1.337 -13.587  38.574  1.00 35.44       EGFA C
ATOM   2783  CG  GLU   296      -2.292 -12.988  39.619  1.00 33.45       EGFA C
ATOM   2784  CD  GLU   296      -3.482 -13.882  39.915  1.00 35.35       EGFA C
ATOM   2785  OE1 GLU   296      -3.596 -14.933  39.248  1.00 35.30       EGFA O
ATOM   2786  OE2 GLU   296      -4.300 -13.540  40.804  1.00 33.34       EGFA O
ATOM   2787  C   GLU   296      -0.588 -13.629  36.151  1.00 36.70       EGFA C
ATOM   2788  O   GLU   296       0.499 -13.060  36.244  1.00 36.33       EGFA O
ATOM   2789  N   CYS   297      -0.838 -14.530  35.207  1.00 36.97       EGFA N
ATOM   2790  CA  CYS   297       0.228 -15.025  34.351  1.00 37.51       EGFA C
ATOM   2791  C   CYS   297       0.233 -14.447  32.972  1.00 37.46       EGFA C
ATOM   2792  O   CYS   297       1.110 -14.750  32.177  1.00 36.37       EGFA O
ATOM   2793  CB  CYS   297       0.151 -16.521  34.243  1.00 38.66       EGFA C
ATOM   2794  SG  CYS   297       0.339 -17.282  35.863  1.00 41.23       EGFA S
ATOM   2795  N   LEU   298      -0.759 -13.620  32.688  1.00 38.90       EGFA N
ATOM   2796  CA  LEU   298      -0.761 -12.827  31.478  1.00 38.94       EGFA C
ATOM   2797  CB  LEU   298      -2.174 -12.332  31.205  1.00 38.48       EGFA C
ATOM   2798  CG  LEU   298      -3.084 -13.516  30.877  1.00 39.20       EGFA C
ATOM   2799  CD1 LEU   298      -4.519 -13.178  31.222  1.00 38.76       EGFA C
ATOM   2800  CD2 LEU   298      -2.920 -13.890  29.400  1.00 37.53       EGFA C
ATOM   2801  C   LEU   298       0.202 -11.667  31.660  1.00 40.24       EGFA C
ATOM   2802  O   LEU   298       0.430 -10.878  30.742  1.00 40.91       EGFA O
ATOM   2803  N   ASP   299       0.781 -11.567  32.851  1.00 40.63       EGFA N
ATOM   2804  CA  ASP   299       1.885 -10.648  33.038  1.00 41.29       EGFA C
ATOM   2805  CB  ASP   299       1.615  -9.716  34.207  1.00 45.11       EGFA C
ATOM   2806  CG  ASP   299       2.450  -8.461  34.133  1.00 48.48       EGFA C
ATOM   2807  OD1 ASP   299       2.944  -8.005  35.182  1.00 51.40       EGFA O
ATOM   2808  OD2 ASP   299       2.613  -7.931  33.013  1.00 50.27       EGFA O
ATOM   2809  C   ASP   299       3.199 -11.374  33.265  1.00 40.06       EGFA C
ATOM   2810  O   ASP   299       3.518 -11.759  34.383  1.00 39.62       EGFA O
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ATOM   2811  N   ASN  300       3.952 -11.557  32.187  1.00 38.77      EGFA N
ATOM   2812  CA  ASN  300       5.278 -12.140  32.257  1.00 38.83      EGFA C
ATOM   2813  CB  ASN  300       6.187 -11.246  33.093  1.00 40.88      EGFA C
ATOM   2814  CG  ASN  300       7.613 -11.245  32.593  1.00 43.06      EGFA C
ATOM   2815  OD1 ASN  300       8.562 -11.252  33.378  1.00 43.69      EGFA O
ATOM   2816  ND2 ASN  300       7.775 -11.234  31.273  1.00 44.81      EGFA N
ATOM   2817  C   ASN  300       5.256 -13.547  32.843  1.00 38.88      EGFA C
ATOM   2818  O   ASN  300       6.235 -13.997  33.448  1.00 38.75      EGFA O
ATOM   2819  N   ASN  301       4.136 -14.238  32.661  1.00 38.27      EGFA N
ATOM   2820  CA  ASN  301       3.977 -15.602  33.153  1.00 38.98      EGFA C
ATOM   2821  CB  ASN  301       5.029 -16.514  32.524  1.00 38.60      EGFA C
ATOM   2822  CG  ASN  301       4.603 -17.965  32.508  1.00 39.84      EGFA C
ATOM   2823  OD1 ASN  301       3.431 -18.277  32.280  1.00 39.37      EGFA O
ATOM   2824  ND2 ASN  301       5.554 -18.866  32.748  1.00 39.66      EGFA N
ATOM   2825  C   ASN  301       4.091 -15.648  34.676  1.00 39.19      EGFA C
ATOM   2826  O   ASN  301       4.368 -16.697  35.265  1.00 38.65      EGFA O
ATOM   2827  N   GLY  302       3.874 -14.496  35.305  1.00 38.94      EGFA N
ATOM   2828  CA  GLY  302       3.929 -14.420  36.752  1.00 39.47      EGFA C
ATOM   2829  C   GLY  302       5.356 -14.384  37.255  1.00 39.98      EGFA C
ATOM   2830  O   GLY  302       5.620 -14.710  38.408  1.00 40.36      EGFA O
ATOM   2831  N   GLY  303       6.277 -13.993  36.381  1.00 40.10      EGFA N
ATOM   2832  CA  GLY  303       7.674 -13.946  36.756  1.00 41.13      EGFA C
ATOM   2833  C   GLY  303       8.255 -15.337  36.912  1.00 41.77      EGFA C
ATOM   2834  O   GLY  303       9.411 -15.490  37.318  1.00 43.49      EGFA O
ATOM   2835  N   CYS  304       7.448 -16.347  36.582  1.00 41.53      EGFA N
ATOM   2836  CA  CYS  304       7.819 -17.757  36.718  1.00 39.56      EGFA C
ATOM   2837  C   CYS  304       8.848 -18.175  35.670  1.00 38.81      EGFA C
ATOM   2838  O   CYS  304       8.667 -17.908  34.483  1.00 39.26      EGFA O
ATOM   2839  CB  CYS  304       6.579 -18.636  36.564  1.00 38.73      EGFA C
ATOM   2840  SG  CYS  304       5.347 -18.462  37.890  1.00 37.07      EGFA S
ATOM   2841  N   SER  305       9.917 -18.843  36.093  1.00 37.94      EGFA N
ATOM   2842  CA  SER  305      10.966 -19.202  35.149  1.00 36.26      EGFA C
ATOM   2843  CB  SER  305      12.152 -19.862  35.870  1.00 34.90      EGFA C
ATOM   2844  OG  SER  305      11.790 -21.060  36.529  1.00 33.63      EGFA O
ATOM   2845  C   SER  305      10.398 -20.142  34.094  1.00 36.04      EGFA C
ATOM   2846  O   SER  305      10.667 -19.985  32.904  1.00 35.42      EGFA O
ATOM   2847  N   HIS  306       9.598 -21.105  34.547  1.00 36.85      EGFA N
ATOM   2848  CA  HIS  306       9.060 -22.156  33.689  1.00 36.68      EGFA C
ATOM   2849  CB  HIS  306       9.468 -23.524  34.237  1.00 37.58      EGFA C
ATOM   2850  CG  HIS  306      10.931 -23.805  34.109  1.00 38.79      EGFA C
ATOM   2851  CD2 HIS  306      12.012 -23.054  34.423  1.00 38.86      EGFA C
ATOM   2852  ND1 HIS  306      11.420 -24.974  33.567  1.00 39.47      EGFA N
ATOM   2853  CE1 HIS  306      12.739 -24.929  33.550  1.00 39.63      EGFA C
ATOM   2854  NE2 HIS  306      13.124 -23.774  34.064  1.00 39.05      EGFA N
ATOM   2855  C   HIS  306       7.541 -22.080  33.566  1.00 36.61      EGFA C
ATOM   2856  O   HIS  306       7.014 -21.231  32.842  1.00 36.74      EGFA O
ATOM   2857  N   VAL  307       6.839 -22.967  34.271  1.00 36.53      EGFA N
ATOM   2858  CA  VAL  307       5.381 -22.965  34.252  1.00 36.68      EGFA C
ATOM   2859  CB  VAL  307       4.793 -24.385  34.359  1.00 36.53      EGFA C
ATOM   2860  CG1 VAL  307       3.271 -24.316  34.440  1.00 33.02      EGFA C
ATOM   2861  CG2 VAL  307       5.210 -25.204  33.153  1.00 36.84      EGFA C
ATOM   2862  C   VAL  307       4.785 -22.135  35.370  1.00 37.94      EGFA C
ATOM   2863  O   VAL  307       5.269 -22.143  36.505  1.00 36.79      EGFA O
ATOM   2864  N   CYS  308       3.726 -21.413  35.019  1.00 40.08      EGFA N
ATOM   2865  CA  CYS  308       2.902 -20.690  35.978  1.00 41.42      EGFA C
ATOM   2866  C   CYS  308       1.562 -21.425  36.159  1.00 41.00      EGFA C
ATOM   2867  O   CYS  308       1.026 -22.009  35.211  1.00 40.92      EGFA O
ATOM   2868  CB  CYS  308       2.585 -19.294  35.465  1.00 42.46      EGFA C
ATOM   2869  SG  CYS  308       0.778 -19.184  35.372  1.00 49.21      EGFA S
ATOM   2870  N   ASN  309       1.016 -21.369  37.371  1.00 40.41      EGFA N
ATOM   2871  CA  ASN  309      -0.305 -21.909  37.647  1.00 39.28      EGFA C
ATOM   2872  CB  ASN  309      -0.193 -23.037  38.661  1.00 37.87      EGFA C
ATOM   2873  CG  ASN  309      -1.511 -23.706  38.923  1.00 37.08      EGFA C
ATOM   2874  OD1 ASN  309      -2.383 -23.738  38.061  1.00 37.19      EGFA O
ATOM   2875  ND2 ASN  309      -1.667 -24.249  40.116  1.00 38.27      EGFA N
ATOM   2876  C   ASN  309      -1.244 -20.825  38.181  1.00 40.43      EGFA C
ATOM   2877  O   ASN  309      -1.025 -20.272  39.260  1.00 41.33      EGFA O
ATOM   2878  N   ASP  310      -2.291 -20.529  37.417  1.00 41.20      EGFA N
ATOM   2879  CA  ASP  310      -3.285 -19.531  37.801  1.00 40.90      EGFA C
ATOM   2880  CB  ASP  310      -4.013 -19.017  36.553  1.00 42.26      EGFA C
ATOM   2881  CG  ASP  310      -4.927 -17.844  36.842  1.00 42.34      EGFA C
ATOM   2882  OD1 ASP  310      -4.822 -17.258  37.937  1.00 44.79      EGFA O
ATOM   2883  OD2 ASP  310      -5.749 -17.507  35.968  1.00 41.66      EGFA O
ATOM   2884  C   ASP  310      -4.297 -20.145  38.755  1.00 41.32      EGFA C
ATOM   2885  O   ASP  310      -5.140 -20.944  38.337  1.00 41.03      EGFA O
ATOM   2886  N   LEU  311      -4.206 -19.767  40.031  1.00 41.42      EGFA N
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ATOM   2887  CA  LEU   311      -5.173 -20.183  41.044  1.00 41.31      EGFA C
ATOM   2888  CB  LEU   311      -4.563 -20.085  42.431  1.00 38.52      EGFA C
ATOM   2889  CG  LEU   311      -3.108 -20.509  42.498  1.00 36.88      EGFA C
ATOM   2890  CD1 LEU   311      -2.561 -20.187  43.872  1.00 37.89      EGFA C
ATOM   2891  CD2 LEU   311      -2.991 -21.987  42.182  1.00 34.94      EGFA C
ATOM   2892  C   LEU   311      -6.398 -19.289  40.995  1.00 42.14      EGFA C
ATOM   2893  O   LEU   311      -6.479 -18.376  40.185  1.00 41.25      EGFA O
ATOM   2894  N   LYS   312      -7.360 -19.563  41.862  1.00 43.27      EGFA N
ATOM   2895  CA  LYS   312      -8.478 -18.658  42.005  1.00 45.45      EGFA C
ATOM   2896  CB  LYS   312      -9.624 -19.345  42.749  1.00 48.60      EGFA C
ATOM   2897  CG  LYS   312     -10.251 -20.511  41.979  1.00 53.73      EGFA C
ATOM   2898  CD  LYS   312     -11.599 -20.945  42.575  1.00 56.82      EGFA C
ATOM   2899  CE  LYS   312     -12.280 -22.005  41.700  1.00 57.14      EGFA C
ATOM   2900  NZ  LYS   312     -13.550 -22.527  42.291  1.00 57.94      EGFA N
ATOM   2901  C   LYS   312      -7.973 -17.464  42.793  1.00 44.13      EGFA C
ATOM   2902  O   LYS   312      -8.229 -16.315  42.434  1.00 44.64      EGFA O
ATOM   2903  N   ILE   313      -7.234 -17.754  43.860  1.00 43.02      EGFA N
ATOM   2904  CA  ILE   313      -6.706 -16.727  44.744  1.00 41.44      EGFA C
ATOM   2905  CB  ILE   313      -6.987 -17.073  46.217  1.00 42.03      EGFA C
ATOM   2906  CG2 ILE   313      -7.416 -15.833  46.978  1.00 40.16      EGFA C
ATOM   2907  CG1 ILE   313      -8.083 -18.140  46.283  1.00 43.15      EGFA C
ATOM   2908  CD1 ILE   313      -9.026 -17.992  47.445  1.00 43.53      EGFA C
ATOM   2909  C   ILE   313      -5.207 -16.672  44.521  1.00 40.70      EGFA C
ATOM   2910  O   ILE   313      -4.445 -17.408  45.151  1.00 42.01      EGFA O
ATOM   2911  N   GLY   314      -4.797 -15.805  43.605  1.00 39.34      EGFA N
ATOM   2912  CA  GLY   314      -3.396 -15.694  43.255  1.00 38.80      EGFA C
ATOM   2913  C   GLY   314      -2.928 -16.707  42.230  1.00 38.26      EGFA C
ATOM   2914  O   GLY   314      -3.707 -17.249  41.456  1.00 37.64      EGFA O
ATOM   2915  N   TYR   315      -1.625 -16.948  42.226  1.00 38.97      EGFA N
ATOM   2916  CA  TYR   315      -1.028 -17.987  41.405  1.00 38.95      EGFA C
ATOM   2917  CB  TYR   315      -0.553 -17.405  40.076  1.00 37.90      EGFA C
ATOM   2918  CG  TYR   315       0.616 -16.457  40.208  1.00 38.39      EGFA C
ATOM   2919  CD1 TYR   315       1.874 -16.788  39.706  1.00 38.94      EGFA C
ATOM   2920  CE1 TYR   315       2.942 -15.906  39.803  1.00 38.15      EGFA C
ATOM   2921  CD2 TYR   315       0.459 -15.216  40.807  1.00 38.29      EGFA C
ATOM   2922  CE2 TYR   315       1.517 -14.333  40.910  1.00 38.40      EGFA C
ATOM   2923  CZ  TYR   315       2.756 -14.675  40.405  1.00 38.57      EGFA C
ATOM   2924  OH  TYR   315       3.805 -13.783  40.526  1.00 37.81      EGFA O
ATOM   2925  C   TYR   315       0.153 -18.612  42.123  1.00 39.94      EGFA C
ATOM   2926  O   TYR   315       0.364 -18.401  43.323  1.00 40.28      EGFA O
ATOM   2927  N   GLU   316       0.933 -19.369  41.364  1.00 41.83      EGFA N
ATOM   2928  CA  GLU   316       2.158 -19.964  41.871  1.00 43.58      EGFA C
ATOM   2929  CB  GLU   316       1.819 -21.179  42.721  1.00 44.75      EGFA C
ATOM   2930  CG  GLU   316       1.030 -22.213  41.959  1.00 48.82      EGFA C
ATOM   2931  CD  GLU   316       0.823 -23.461  42.761  1.00 52.94      EGFA C
ATOM   2932  OE1 GLU   316       1.123 -23.438  43.971  1.00 57.20      EGFA O
ATOM   2933  OE2 GLU   316       0.381 -24.475  42.184  1.00 54.04      EGFA O
ATOM   2934  C   GLU   316       3.041 -20.383  40.701  1.00 42.80      EGFA C
ATOM   2935  O   GLU   316       2.568 -20.500  39.575  1.00 44.00      EGFA O
ATOM   2936  N   CYS   317       4.326 -20.592  40.957  1.00 42.14      EGFA N
ATOM   2937  CA  CYS   317       5.195 -21.137  39.927  1.00 41.35      EGFA C
ATOM   2938  C   CYS   317       5.412 -22.630  40.169  1.00 42.56      EGFA C
ATOM   2939  O   CYS   317       5.470 -23.079  41.316  1.00 42.57      EGFA O
ATOM   2940  CB  CYS   317       6.538 -20.413  39.925  1.00 38.97      EGFA C
ATOM   2941  SG  CYS   317       6.442 -18.628  39.586  1.00 35.62      EGFA S
ATOM   2942  N   LEU   318       5.547 -23.397  39.089  1.00 43.52      EGFA N
ATOM   2943  CA  LEU   318       5.832 -24.820  39.210  1.00 43.40      EGFA C
ATOM   2944  CB  LEU   318       4.653 -25.640  38.689  1.00 41.86      EGFA C
ATOM   2945  CG  LEU   318       3.328 -25.268  39.355  1.00 41.42      EGFA C
ATOM   2946  CD1 LEU   318       2.255 -26.234  38.937  1.00 42.17      EGFA C
ATOM   2947  CD2 LEU   318       3.481 -25.286  40.855  1.00 41.53      EGFA C
ATOM   2948  C   LEU   318       7.098 -25.185  38.453  1.00 44.60      EGFA C
ATOM   2949  O   LEU   318       7.489 -24.493  37.510  1.00 44.39      EGFA O
ATOM   2950  N   CYS   319       7.738 -26.273  38.875  1.00 46.62      EGFA N
ATOM   2951  CA  CYS   319       9.002 -26.685  38.276  1.00 48.75      EGFA C
ATOM   2952  C   CYS   319       8.993 -28.073  37.639  1.00 48.42      EGFA C
ATOM   2953  O   CYS   319       8.274 -28.970  38.077  1.00 47.58      EGFA O
ATOM   2954  CB  CYS   319      10.123 -26.615  39.317  1.00 49.62      EGFA C
ATOM   2955  SG  CYS   319      10.467 -24.933  39.923  1.00 50.75      EGFA S
ATOM   2956  N   PRO   320       9.812 -28.263  36.593  1.00 49.01      EGFA N
ATOM   2957  CD  PRO   320      10.541 -27.229  35.842  1.00 48.55      EGFA C
ATOM   2958  CA  PRO   320      10.067 -29.605  36.060  1.00 49.42      EGFA C
ATOM   2959  CB  PRO   320      11.111 -29.369  34.972  1.00 48.61      EGFA C
ATOM   2960  CG  PRO   320      10.927 -27.948  34.582  1.00 48.27      EGFA C
ATOM   2961  C   PRO   320      10.600 -30.500  37.165  1.00 50.82      EGFA C
ATOM   2962  O   PRO   320      11.453 -30.086  37.952  1.00 50.19      EGFA O
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ATOM   2963  N   ASP   321      10.096 -31.724  37.234  1.00 52.17      EGFA N
ATOM   2964  CA  ASP   321      10.479 -32.600  38.325  1.00 52.72      EGFA C
ATOM   2965  CB  ASP   321       9.810 -33.970  38.187  1.00 57.23      EGFA C
ATOM   2966  CG  ASP   321       9.182 -34.447  39.499  1.00 61.91      EGFA C
ATOM   2967  OD1 ASP   321       8.807 -33.586  40.332  1.00 63.09      EGFA O
ATOM   2968  OD2 ASP   321       9.064 -35.680  39.701  1.00 64.11      EGFA O
ATOM   2969  C   ASP   321      11.987 -32.742  38.310  1.00 51.06      EGFA C
ATOM   2970  O   ASP   321      12.599 -32.869  37.246  1.00 49.86      EGFA O
ATOM   2971  N   GLY   322      12.577 -32.695  39.499  1.00 49.84      EGFA N
ATOM   2972  CA  GLY   322      14.023 -32.681  39.614  1.00 48.51      EGFA C
ATOM   2973  C   GLY   322      14.592 -31.285  39.793  1.00 47.34      EGFA C
ATOM   2974  O   GLY   322      15.729 -31.131  40.239  1.00 47.93      EGFA O
ATOM   2975  N   PHE   323      13.807 -30.266  39.445  1.00 45.29      EGFA N
ATOM   2976  CA  PHE   323      14.247 -28.880  39.583  1.00 42.20      EGFA C
ATOM   2977  CB  PHE   323      13.607 -27.999  38.517  1.00 40.51      EGFA C
ATOM   2978  CG  PHE   323      14.157 -28.215  37.141  1.00 38.07      EGFA C
ATOM   2979  CD1 PHE   323      14.250 -29.489  36.610  1.00 36.97      EGFA C
ATOM   2980  CD2 PHE   323      14.550 -27.135  36.364  1.00 37.05      EGFA C
ATOM   2981  CE1 PHE   323      14.723 -29.681  35.330  1.00 37.63      EGFA C
ATOM   2982  CE2 PHE   323      15.024 -27.317  35.082  1.00 36.22      EGFA C
ATOM   2983  CZ  PHE   323      15.110 -28.590  34.561  1.00 37.83      EGFA C
ATOM   2984  C   PHE   323      13.890 -28.332  40.947  1.00 41.10      EGFA C
ATOM   2985  O   PHE   323      13.022 -28.857  41.631  1.00 40.98      EGFA O
ATOM   2986  N   GLN   324      14.567 -27.266  41.339  1.00 41.32      EGFA N
ATOM   2987  CA  GLN   324      14.326 -26.657  42.637  1.00 42.35      EGFA C
ATOM   2988  CB  GLN   324      15.645 -26.527  43.393  1.00 45.11      EGFA C
ATOM   2989  CG  GLN   324      15.543 -26.714  44.897  1.00 48.90      EGFA C
ATOM   2990  CD  GLN   324      16.897 -26.588  45.584  1.00 50.71      EGFA C
ATOM   2991  OE1 GLN   324      16.978 -26.425  46.808  1.00 51.49      EGFA O
ATOM   2992  NE2 GLN   324      17.972 -26.660  44.793  1.00 50.40      EGFA N
ATOM   2993  C   GLN   324      13.729 -25.281  42.408  1.00 41.37      EGFA C
ATOM   2994  O   GLN   324      14.076 -24.607  41.439  1.00 41.39      EGFA O
ATOM   2995  N   LEU   325      12.831 -24.856  43.287  1.00 40.78      EGFA N
ATOM   2996  CA  LEU   325      12.250 -23.527  43.147  1.00 40.37      EGFA C
ATOM   2997  CB  LEU   325      10.740 -23.570  43.373  1.00 38.89      EGFA C
ATOM   2998  CG  LEU   325      10.036 -22.248  43.063  1.00 37.99      EGFA C
ATOM   2999  CD1 LEU   325      10.139 -21.941  41.584  1.00 35.18      EGFA C
ATOM   3000  CD2 LEU   325       8.592 -22.331  43.485  1.00 37.81      EGFA C
ATOM   3001  C   LEU   325      12.880 -22.553  44.131  1.00 40.79      EGFA C
ATOM   3002  O   LEU   325      12.896 -22.806  45.337  1.00 40.04      EGFA O
ATOM   3003  N   VAL   326      13.402 -21.442  43.618  1.00 41.36      EGFA N
ATOM   3004  CA  VAL   326      14.077 -20.477  44.476  1.00 42.53      EGFA C
ATOM   3005  CB  VAL   326      15.592 -20.351  44.122  1.00 41.99      EGFA C
ATOM   3006  CG1 VAL   326      16.010 -21.475  43.201  1.00 41.24      EGFA C
ATOM   3007  CG2 VAL   326      15.883 -19.007  43.506  1.00 41.93      EGFA C
ATOM   3008  C   VAL   326      13.418 -19.109  44.385  1.00 43.44      EGFA C
ATOM   3009  O   VAL   326      12.929 -18.716  43.332  1.00 43.12      EGFA O
ATOM   3010  N   ALA   327      13.394 -18.390  45.501  1.00 45.25      EGFA N
ATOM   3011  CA  ALA   327      12.751 -17.084  45.539  1.00 47.09      EGFA C
ATOM   3012  CB  ALA   327      13.443 -16.139  44.577  1.00 46.26      EGFA C
ATOM   3013  C   ALA   327      11.270 -17.201  45.185  1.00 48.01      EGFA C
ATOM   3014  O   ALA   327      10.567 -16.197  45.078  1.00 48.60      EGFA O
ATOM   3015  N   GLN   328      10.813 -18.435  44.997  1.00 49.94      EGFA N
ATOM   3016  CA  GLN   328       9.413 -18.717  44.692  1.00 51.60      EGFA C
ATOM   3017  CB  GLN   328       8.492 -17.860  45.563  1.00 51.40      EGFA C
ATOM   3018  CG  GLN   328       7.372 -18.643  46.191  1.00 53.05      EGFA C
ATOM   3019  CD  GLN   328       7.288 -18.392  47.666  1.00 56.41      EGFA C
ATOM   3020  OE1 GLN   328       7.462 -19.307  48.477  1.00 58.59      EGFA O
ATOM   3021  NE2 GLN   328       7.022 -17.136  48.036  1.00 57.77      EGFA N
ATOM   3022  C   GLN   328       9.050 -18.516  43.224  1.00 51.40      EGFA C
ATOM   3023  O   GLN   328       7.872 -18.574  42.863  1.00 52.05      EGFA O
ATOM   3024  N   ARG   329      10.045 -18.278  42.376  1.00 52.02      EGFA N
ATOM   3025  CA  ARG   329       9.759 -18.102  40.962  1.00 53.90      EGFA C
ATOM   3026  CB  ARG   329       9.202 -16.710  40.719  1.00 54.36      EGFA C
ATOM   3027  CG  ARG   329      10.176 -15.594  40.973  1.00 55.41      EGFA C
ATOM   3028  CD  ARG   329       9.425 -14.289  40.909  1.00 56.46      EGFA C
ATOM   3029  NE  ARG   329       8.174 -14.397  41.644  1.00 57.62      EGFA N
ATOM   3030  CZ  ARG   329       7.271 -13.427  41.681  1.00 58.17      EGFA C
ATOM   3031  NH1 ARG   329       7.499 -12.294  41.018  1.00 57.94      EGFA N
ATOM   3032  NH2 ARG   329       6.159 -13.570  42.390  1.00 58.09      EGFA N
ATOM   3033  C   ARG   329      10.921 -18.337  40.015  1.00 54.26      EGFA C
ATOM   3034  O   ARG   329      10.899 -17.869  38.881  1.00 54.09      EGFA O
ATOM   3035  N   ARG   330      11.925 -19.079  40.459  1.00 55.91      EGFA N
ATOM   3036  CA  ARG   330      12.957 -19.541  39.545  1.00 56.83      EGFA C
ATOM   3037  CB  ARG   330      14.202 -18.663  39.676  1.00 59.07      EGFA C
ATOM   3038  CG  ARG   330      15.156 -18.777  38.494  1.00 63.56      EGFA C
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ATOM   3039  CD  ARG   330      16.009 -17.526  38.337  1.00 67.44      EGFA C
ATOM   3040  NE  ARG   330      17.205 -17.544  39.183  1.00 71.71      EGFA N
ATOM   3041  CZ  ARG   330      18.101 -16.560  39.232  1.00 73.25      EGFA C
ATOM   3042  NH1 ARG   330      17.931 -15.476  38.483  1.00 74.17      EGFA N
ATOM   3043  NH2 ARG   330      19.165 -16.658  40.026  1.00 73.57      EGFA N
ATOM   3044  C   ARG   330      13.307 -21.007  39.805  1.00 56.14      EGFA C
ATOM   3045  O   ARG   330      13.624 -21.398  40.931  1.00 55.45      EGFA O
ATOM   3046  N   CYS   331      13.237 -21.821  38.757  1.00 55.78      EGFA N
ATOM   3047  CA  CYS   331      13.615 -23.219  38.876  1.00 55.78      EGFA C
ATOM   3048  C   CYS   331      15.046 -23.384  38.407  1.00 56.71      EGFA C
ATOM   3049  O   CYS   331      15.424 -22.886  37.344  1.00 54.16      EGFA O
ATOM   3050  CB  CYS   331      12.707 -24.105  38.024  1.00 54.71      EGFA C
ATOM   3051  SG  CYS   331      10.917 -23.862  38.247  1.00 52.34      EGFA S
ATOM   3052  N   GLU   332      15.843 -24.075  39.210  1.00 59.67      EGFA N
ATOM   3053  CA  GLU   332      17.181 -24.445  38.793  1.00 63.04      EGFA C
ATOM   3054  CB  GLU   332      18.234 -23.765  39.677  1.00 63.37      EGFA C
ATOM   3055  CG  GLU   332      17.872 -23.624  41.151  1.00 64.27      EGFA C
ATOM   3056  CD  GLU   332      18.797 -22.655  41.891  1.00 64.91      EGFA C
ATOM   3057  OE1 GLU   332      19.367 -23.051  42.935  1.00 64.31      EGFA O
ATOM   3058  OE2 GLU   332      18.951 -21.499  41.428  1.00 64.07      EGFA O
ATOM   3059  C   GLU   332      17.350 -25.953  38.811  1.00 65.33      EGFA C
ATOM   3060  O   GLU   332      17.082 -26.620  39.816  1.00 64.39      EGFA O
ATOM   3061  N   ASP   333      17.780 -26.476  37.667  1.00 68.69      EGFA N
ATOM   3062  CA  ASP   333      17.880 -27.910  37.433  1.00 72.25      EGFA C
ATOM   3063  CB  ASP   333      18.483 -28.168  36.056  1.00 74.16      EGFA C
ATOM   3064  CG  ASP   333      19.649 -27.250  35.762  1.00 76.50      EGFA C
ATOM   3065  OD1 ASP   333      19.458 -26.013  35.827  1.00 77.91      EGFA O
ATOM   3066  OD2 ASP   333      20.754 -27.757  35.476  1.00 77.38      EGFA O
ATOM   3067  C   ASP   333      18.756 -28.536  38.488  1.00 73.49      EGFA C
ATOM   3068  O   ASP   333      18.526 -29.673  38.894  1.00 73.58      EGFA O
ATOM   3069  N   ILE   334      19.769 -27.780  38.903  1.00 75.52      EGFA N
ATOM   3070  CA  ILE   334      20.587 -28.098  40.069  1.00 77.98      EGFA C
ATOM   3071  CB  ILE   334      20.070 -27.299  41.320  1.00 78.68      EGFA C
ATOM   3072  CG2 ILE   334      19.986 -28.191  42.551  1.00 78.54      EGFA C
ATOM   3073  CG1 ILE   334      20.989 -26.098  41.585  1.00 78.89      EGFA C
ATOM   3074  CD1 ILE   334      21.191 -25.170  40.391  1.00 77.96      EGFA C
ATOM   3075  C   ILE   334      20.666 -29.597  40.363  1.00 79.11      EGFA C
ATOM   3076  O   ILE   334      21.755 -30.162  40.118  1.00 79.71      EGFA O
ATOM   3077  OXT ILE   334      19.656 -30.196  40.802  1.00 79.97      EGFA O
TER    3078       ILE   334                                              EGFA
ATOM   3079  CA  CA      1      -5.688 -15.911  40.122  1.00 56.81      ION  C
TER    3080       CA      1                                              ION
END
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TABLE 35.3

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ATOM      1  CB  THR    61       7.625 -20.113   2.894  1.00120.20      A    C
ATOM      2  OG1 THR    61       7.048 -19.215   1.936  1.00120.97      A    O
ATOM      3  CG2 THR    61       7.446 -19.537   4.289  1.00121.74      A    C
ATOM      4  C   THR    61       9.869 -19.003   2.758  1.00115.49      A    C
ATOM      5  O   THR    61      10.710 -18.870   3.648  1.00115.58      A    O
ATOM      6  N   THR    61       9.741 -21.360   3.461  1.00119.24      A    N
ATOM      7  CA  THR    61       9.132 -20.323   2.577  1.00118.55      A    C
ATOM      8  N   ALA    62       9.546 -18.032   1.907  1.00111.50      A    N
ATOM      9  CA  ALA    62      10.296 -16.784   1.830  1.00106.53      A    C
ATOM     10  CB  ALA    62      10.124 -16.161   0.455  1.00108.21      A    C
ATOM     11  C   ALA    62       9.882 -15.789   2.902  1.00102.91      A    C
ATOM     12  O   ALA    62       8.701 -15.600   3.174  1.00101.98      A    O
ATOM     13  N   THR    63      10.870 -15.145   3.501  1.00 98.29      A    N
ATOM     14  CA  THR    63      10.624 -14.157   4.534  1.00 93.32      A    C
ATOM     15  CB  THR    63      11.644 -14.326   5.679  1.00 92.70      A    C
ATOM     16  OG1 THR    63      12.973 -14.322   5.147  1.00 92.34      A    O
ATOM     17  CG2 THR    63      11.422 -15.647   6.398  1.00 92.08      A    C
ATOM     18  C   THR    63      10.716 -12.747   3.944  1.00 89.66      A    C
ATOM     19  O   THR    63      11.156 -12.571   2.810  1.00 90.88      A    O
ATOM     20  N   PHE    64      10.280 -11.747   4.705  1.00 83.77      A    N
ATOM     21  CA  PHE    64      10.405 -10.353   4.278  1.00 77.76      A    C
ATOM     22  CB  PHE    64       9.065  -9.825   3.799  1.00 72.88      A    C
ATOM     23  CG  PHE    64       9.028  -8.343   3.648  1.00 67.65      A    C
ATOM     24  CD1 PHE    64       9.305  -7.752   2.429  1.00 66.67      A    C
ATOM     25  CD2 PHE    64       8.697  -7.530   4.726  1.00 66.49      A    C
ATOM     26  CE1 PHE    64       9.251  -6.370   2.285  1.00 66.42      A    C
ATOM     27  CE2 PHE    64       8.640  -6.150   4.594  1.00 64.45      A    C
ATOM     28  CZ  PHE    64       8.917  -5.567   3.374  1.00 64.76      A    C
ATOM     29  C   PHE    64      10.936  -9.426   5.370  1.00 76.67      A    C
ATOM     30  O   PHE    64      10.563  -9.557   6.540  1.00 76.03      A    O
ATOM     31  N   HIS    65      11.787  -8.477   4.975  1.00 75.50      A    N
ATOM     32  CA  HIS    65      12.561  -7.686   5.928  1.00 74.60      A    C
ATOM     33  CB  HIS    65      14.003  -8.192   5.961  1.00 74.07      A    C
ATOM     34  CG  HIS    65      14.117  -9.659   6.235  1.00 74.70      A    C
ATOM     35  CD2 HIS    65      13.881 -10.735   5.448  1.00 74.67      A    C
ATOM     36  ND1 HIS    65      14.517 -10.161   7.456  1.00 75.55      A    N
ATOM     37  CE1 HIS    65      14.523 -11.481   7.409  1.00 74.80      A    C
ATOM     38  NE2 HIS    65      14.141 -11.855   6.201  1.00 74.59      A    N
ATOM     39  C   HIS    65      12.556  -6.187   5.642  1.00 73.68      A    C
ATOM     40  O   HIS    65      12.678  -5.755   4.498  1.00 73.21      A    O
ATOM     41  N   ARG    66      12.425  -5.397   6.700  1.00 73.29      A    N
ATOM     42  CA  ARG    66      12.508  -3.952   6.586  1.00 73.63      A    C
ATOM     43  CB  ARG    66      11.131  -3.320   6.801  1.00 80.94      A    C
ATOM     44  CG  ARG    66      10.664  -3.376   8.239  1.00 91.57      A    C
ATOM     45  CD  ARG    66       9.465  -2.490   8.486  1.00102.14      A    C
ATOM     46  NE  ARG    66       9.120  -2.462   9.905  1.00113.95      A    N
ATOM     47  CZ  ARG    66       9.564  -1.551  10.763  1.00120.16      A    C
ATOM     48  NH1 ARG    66       9.200  -1.602  12.037  1.00124.56      A    N
ATOM     49  NH2 ARG    66      10.368  -0.586  10.342  1.00124.67      A    N
ATOM     50  C   ARG    66      13.482  -3.420   7.630  1.00 68.82      A    C
ATOM     51  O   ARG    66      13.789  -4.088   8.605  1.00 69.03      A    O
ATOM     52  N   CYS    67      13.974  -2.210   7.425  1.00 64.02      A    N
ATOM     53  CA  CYS    67      14.838  -1.590   8.414  1.00 57.74      A    C
ATOM     54  CB  CYS    67      15.397  -0.279   7.865  1.00 55.67      A    C
ATOM     55  SG  CYS    67      16.287   0.702   9.068  1.00 43.75      A    S
ATOM     56  C   CYS    67      14.043  -1.320   9.685  1.00 55.30      A    C
ATOM     57  O   CYS    67      12.919  -0.825   9.630  1.00 53.23      A    O
ATOM     58  N   ALA    68      14.830  -1.644  10.828  1.00 52.57      A    N
ATOM     59  CA  ALA    68      13.953  -1.434  12.097  1.00 51.78      A    C
ATOM     60  CB  ALA    68      14.818  -1.940  13.236  1.00 48.43      A    C
ATOM     61  C   ALA    68      13.624   0.043  12.306  1.00 53.12      A    C
ATOM     62  O   ALA    68      12.511   0.389  12.700  1.00 53.69      A    O
ATOM     63  N   LYS    69      14.596   0.914  12.042  1.00 53.95      A    N
ATOM     64  CA  LYS    69      14.435   2.342  12.306  1.00 53.04      A    C
ATOM     65  CB  LYS    69      15.814   3.015  12.422  1.00 55.99      A    C
ATOM     66  CG  LYS    69      16.931   2.058  12.899  1.00 57.69      A    C
ATOM     67  CD  LYS    69      17.593   2.502  14.208  1.00 57.50      A    C
ATOM     68  CE  LYS    69      17.940   1.307  15.091  1.00 57.22      A    C
ATOM     69  NZ  LYS    69      18.773   1.709  16.260  1.00 57.57      A    N
ATOM     70  C   LYS    69      13.620   2.975  11.188  1.00 51.24      A    C
ATOM     71  O   LYS    69      14.050   3.061  10.049  1.00 49.56      A    O
ATOM     72  N   ASP    70      12.429   3.424  11.537  1.00 51.12      A    N
ATOM     73  CA  ASP    70      11.407   3.714  10.555  1.00 49.83      A    C
ATOM     74  CB  ASP    70      10.109   4.015  11.279  1.00 54.30      A    C
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ATOM	75	CG	ASP	70	8.902	3.574	10.492	1.00	60.23	A	C
ATOM	76	OD1	ASP	70	8.761	2.334	10.283	1.00	59.99	A	O
ATOM	77	OD2	ASP	70	8.109	4.472	10.087	1.00	61.75	A	O
ATOM	78	C	ASP	70	11.715	4.837	9.567	1.00	45.91	A	C
ATOM	79	O	ASP	70	11.398	4.736	8.386	1.00	43.79	A	O
ATOM	80	N	PRO	71	12.322	5.931	10.040	1.00	43.28	A	N
ATOM	81	CD	PRO	71	12.626	6.241	11.448	1.00	40.22	A	C
ATOM	82	CA	PRO	71	12.627	7.057	9.148	1.00	41.43	A	C
ATOM	83	CB	PRO	71	12.989	8.182	10.108	1.00	39.87	A	C
ATOM	84	CG	PRO	71	13.420	7.490	11.346	1.00	39.36	A	C
ATOM	85	C	PRO	71	13.734	6.767	8.125	1.00	40.85	A	C
ATOM	86	O	PRO	71	13.979	7.562	7.204	1.00	41.69	A	O
ATOM	87	N	TRP	72	14.380	5.614	8.279	1.00	39.17	A	N
ATOM	88	CA	TRP	72	15.426	5.175	7.362	1.00	37.15	A	C
ATOM	89	CB	TRP	72	16.523	4.486	8.133	1.00	33.24	A	C
ATOM	90	CG	TRP	72	17.216	5.367	9.057	1.00	30.60	A	C
ATOM	91	CD2	TRP	72	18.253	4.995	9.959	1.00	28.45	A	C
ATOM	92	CE2	TRP	72	18.678	6.174	10.615	1.00	29.35	A	C
ATOM	93	CE3	TRP	72	18.863	3.785	10.273	1.00	24.28	A	C
ATOM	94	CD1	TRP	72	17.045	6.712	9.194	1.00	32.13	A	C
ATOM	95	NE1	TRP	72	17.923	7.207	10.132	1.00	31.43	A	N
ATOM	96	CZ2	TRP	72	19.684	6.177	11.563	1.00	28.45	A	C
ATOM	97	CZ3	TRP	72	19.860	3.779	11.210	1.00	29.40	A	C
ATOM	98	CH2	TRP	72	20.268	4.971	11.852	1.00	31.73	A	C
ATOM	99	C	TRP	72	14.944	4.217	6.286	1.00	39.33	A	C
ATOM	100	O	TRP	72	15.744	3.741	5.476	1.00	39.93	A	O
ATOM	101	N	ARG	73	13.648	3.916	6.296	1.00	41.01	A	N
ATOM	102	CA	ARG	73	13.065	2.968	5.354	1.00	41.83	A	C
ATOM	103	CB	ARG	73	11.741	2.443	5.903	1.00	43.30	A	C
ATOM	104	CG	ARG	73	11.891	1.798	7.263	1.00	47.92	A	C
ATOM	105	CD	ARG	73	10.582	1.222	7.765	1.00	52.48	A	C
ATOM	106	NE	ARG	73	10.042	0.197	6.879	1.00	56.62	A	N
ATOM	107	CZ	ARG	73	8.761	0.126	6.541	1.00	58.69	A	C
ATOM	108	NH1	ARG	73	7.911	1.025	7.021	1.00	60.74	A	N
ATOM	109	NH2	ARG	73	8.332	-0.841	5.741	1.00	57.82	A	N
ATOM	110	C	ARG	73	12.840	3.653	4.024	1.00	40.96	A	C
ATOM	111	O	ARG	73	12.572	4.841	3.988	1.00	40.86	A	O
ATOM	112	N	LEU	74	12.966	2.913	2.932	1.00	41.26	A	N
ATOM	113	CA	LEU	74	12.781	3.500	1.606	1.00	43.15	A	C
ATOM	114	CB	LEU	74	14.137	3.683	0.914	1.00	42.05	A	C
ATOM	115	CG	LEU	74	15.200	4.372	1.776	1.00	41.09	A	C
ATOM	116	CD1	LEU	74	16.600	4.020	1.259	1.00	40.47	A	C
ATOM	117	CD2	LEU	74	14.945	5.868	1.781	1.00	36.19	A	C
ATOM	118	C	LEU	74	11.885	2.601	0.767	1.00	43.59	A	C
ATOM	119	O	LEU	74	12.341	1.943	-0.180	1.00	45.64	A	O
ATOM	120	N	PRO	75	10.595	2.547	1.122	1.00	42.35	A	N
ATOM	121	CD	PRO	75	10.053	3.206	2.322	1.00	40.81	A	C
ATOM	122	CA	PRO	75	9.605	1.691	0.465	1.00	41.19	A	C
ATOM	123	CB	PRO	75	8.377	1.796	1.370	1.00	40.54	A	C
ATOM	124	CG	PRO	75	8.581	3.023	2.174	1.00	39.32	A	C
ATOM	125	C	PRO	75	9.306	2.079	-0.981	1.00	40.71	A	C
ATOM	126	O	PRO	75	9.277	3.258	-1.329	1.00	39.79	A	O
ATOM	127	N	GLY	76	9.090	1.064	-1.817	1.00	40.71	A	N
ATOM	128	CA	GLY	76	8.969	1.277	-3.250	1.00	38.99	A	C
ATOM	129	C	GLY	76	10.105	0.572	-3.974	1.00	36.52	A	C
ATOM	130	O	GLY	76	10.163	0.540	-5.205	1.00	35.56	A	O
ATOM	131	N	THR	77	11.017	-0.005	-3.206	1.00	34.53	A	N
ATOM	132	CA	THR	77	12.237	-0.530	-3.795	1.00	35.21	A	C
ATOM	133	CB	THR	77	13.366	0.542	-3.812	1.00	34.11	A	C
ATOM	134	OG1	THR	77	12.900	1.742	-4.462	1.00	34.86	A	O
ATOM	135	CG2	THR	77	14.572	0.009	-4.557	1.00	31.17	A	C
ATOM	136	C	THR	77	12.713	-1.726	-2.996	1.00	34.73	A	C
ATOM	137	O	THR	77	12.969	-1.625	-1.790	1.00	34.99	A	O
ATOM	138	N	TYR	78	12.825	-2.869	-3.655	1.00	33.72	A	N
ATOM	139	CA	TYR	78	13.063	-4.087	-2.911	1.00	34.19	A	C
ATOM	140	CB	TYR	78	11.755	-4.904	-2.837	1.00	32.06	A	C
ATOM	141	CG	TYR	78	10.585	-4.090	-2.321	1.00	29.68	A	C
ATOM	142	CD1	TYR	78	9.907	-3.183	-3.152	1.00	29.12	A	C
ATOM	143	CE1	TYR	78	8.916	-2.317	-2.648	1.00	28.24	A	C
ATOM	144	CD2	TYR	78	10.183	-4.134	-0.967	1.00	28.00	A	C
ATOM	145	CE2	TYR	78	9.199	-3.289	-0.463	1.00	24.98	A	C
ATOM	146	CZ	TYR	78	8.577	-2.374	-1.307	1.00	28.24	A	C
ATOM	147	OH	TYR	78	7.659	-1.470	-0.810	1.00	31.65	A	O
ATOM	148	C	TYR	78	14.193	-4.873	-3.557	1.00	35.43	A	C
ATOM	149	O	TYR	78	14.278	-4.993	-4.780	1.00	37.75	A	O
ATOM	150	N	VAL	79	15.094	-5.377	-2.731	1.00	36.65	A	N

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ATOM    151  CA  VAL    79      16.068  -6.347  -3.204  1.00 36.46      A    C
ATOM    152  CB  VAL    79      17.358  -6.304  -2.395  1.00 37.20      A    C
ATOM    153  CG1 VAL    79      18.349  -7.295  -2.972  1.00 37.26      A    C
ATOM    154  CG2 VAL    79      17.924  -4.895  -2.383  1.00 38.46      A    C
ATOM    155  C   VAL    79      15.469  -7.719  -3.001  1.00 38.67      A    C
ATOM    156  O   VAL    79      15.494  -8.246  -1.884  1.00 35.22      A    O
ATOM    157  N   VAL    80      14.915  -8.297  -4.063  1.00 41.04      A    N
ATOM    158  CA  VAL    80      14.581  -9.717  -4.028  1.00 43.11      A    C
ATOM    159  CB  VAL    80      13.775 -10.146  -5.256  1.00 44.56      A    C
ATOM    160  CG1 VAL    80      13.627 -11.649  -5.251  1.00 47.28      A    C
ATOM    161  CG2 VAL    80      12.413  -9.483  -5.257  1.00 47.02      A    C
ATOM    162  C   VAL    80      15.878 -10.520  -4.016  1.00 46.29      A    C
ATOM    163  O   VAL    80      16.741 -10.342  -4.873  1.00 47.03      A    O
ATOM    164  N   VAL    81      16.025 -11.391  -3.034  1.00 49.18      A    N
ATOM    165  CA  VAL    81      17.173 -12.267  -2.967  1.00 52.16      A    C
ATOM    166  CB  VAL    81      17.789 -12.222  -1.565  1.00 51.26      A    C
ATOM    167  CG1 VAL    81      19.027 -13.107  -1.501  1.00 50.81      A    C
ATOM    168  CG2 VAL    81      18.108 -10.791  -1.202  1.00 52.80      A    C
ATOM    169  C   VAL    81      16.687 -13.676  -3.260  1.00 58.30      A    C
ATOM    170  O   VAL    81      15.652 -14.112  -2.732  1.00 58.40      A    O
ATOM    171  N   LEU    82      17.419 -14.393  -4.105  1.00 62.94      A    N
ATOM    172  CA  LEU    82      16.997 -15.734  -4.454  1.00 67.59      A    C
ATOM    173  CB  LEU    82      17.176 -15.949  -5.944  1.00 64.19      A    C
ATOM    174  CG  LEU    82      16.407 -14.837  -6.666  1.00 61.66      A    C
ATOM    175  CD1 LEU    82      16.570 -14.951  -8.164  1.00 60.74      A    C
ATOM    176  CD2 LEU    82      14.938 -14.911  -6.290  1.00 62.60      A    C
ATOM    177  C   LEU    82      17.778 -16.752  -3.648  1.00 73.85      A    C
ATOM    178  O   LEU    82      18.800 -16.419  -3.042  1.00 72.57      A    O
ATOM    179  N   LYS    83      17.268 -17.977  -3.596  1.00 82.70      A    N
ATOM    180  CA  LYS    83      17.995 -19.055  -2.955  1.00 91.24      A    C
ATOM    181  CB  LYS    83      17.184 -20.350  -3.050  1.00 92.11      A    C
ATOM    182  CG  LYS    83      17.946 -21.613  -2.649  1.00 95.32      A    C
ATOM    183  CD  LYS    83      18.130 -21.757  -1.140  1.00 96.21      A    C
ATOM    184  CE  LYS    83      18.758 -23.120  -0.816  1.00 96.12      A    C
ATOM    185  NZ  LYS    83      18.707 -23.483   0.629  1.00 99.11      A    N
ATOM    186  C   LYS    83      19.320 -19.163  -3.694  1.00 95.80      A    C
ATOM    187  O   LYS    83      19.366 -19.052  -4.919  1.00 98.45      A    O
ATOM    188  N   GLU    84      20.399 -19.348  -2.945  1.00100.77      A    N
ATOM    189  CA  GLU    84      21.731 -19.311  -3.532  1.00106.04      A    C
ATOM    190  CB  GLU    84      22.793 -19.394  -2.441  1.00108.26      A    C
ATOM    191  CG  GLU    84      23.164 -20.797  -2.008  1.00112.62      A    C
ATOM    192  CD  GLU    84      24.181 -20.801  -0.937  1.00116.07      A    C
ATOM    193  OE1 GLU    84      25.398 -20.803  -1.237  1.00119.19      A    O
ATOM    194  OE2 GLU    84      23.763 -20.784   0.241  1.00118.28      A    O
ATOM    195  C   GLU    84      21.899 -20.459  -4.512  1.00109.04      A    C
ATOM    196  O   GLU    84      21.010 -21.299  -4.639  1.00108.23      A    O
ATOM    197  N   GLU    85      23.031 -20.490  -5.208  1.00113.81      A    N
ATOM    198  CA  GLU    85      23.242 -21.474  -6.261  1.00118.74      A    C
ATOM    199  CB  GLU    85      23.060 -22.890  -5.704  1.00119.63      A    C
ATOM    200  CG  GLU    85      24.229 -23.390  -4.870  1.00122.20      A    C
ATOM    201  CD  GLU    85      24.788 -24.704  -5.390  1.00123.94      A    C
ATOM    202  OE1 GLU    85      26.009 -24.936  -5.246  1.00124.52      A    O
ATOM    203  OE2 GLU    85      24.011 -25.512  -5.947  1.00125.46      A    O
ATOM    204  C   GLU    85      22.292 -21.252  -7.441  1.00120.43      A    C
ATOM    205  O   GLU    85      22.400 -21.919  -8.470  1.00122.77      A    O
ATOM    206  N   THR    86      21.364 -20.316  -7.290  1.00120.31      A    N
ATOM    207  CA  THR    86      20.463 -19.968  -8.373  1.00119.91      A    C
ATOM    208  CB  THR    86      19.357 -19.023  -7.869  1.00117.92      A    C
ATOM    209  OG1 THR    86      18.543 -19.715  -6.917  1.00115.24      A    O
ATOM    210  CG2 THR    86      18.483 -18.543  -9.021  1.00115.66      A    C
ATOM    211  C   THR    86      21.208 -19.325  -9.541  1.00121.48      A    C
ATOM    212  O   THR    86      22.160 -18.573  -9.353  1.00122.30      A    O
ATOM    213  N   HIS    87      20.764 -19.646 -10.752  1.00123.45      A    N
ATOM    214  CA  HIS    87      21.418 -19.198 -11.980  1.00124.91      A    C
ATOM    215  CB  HIS    87      21.213 -20.227 -13.090  1.00127.94      A    C
ATOM    216  CG  HIS    87      22.040 -21.453 -12.904  1.00131.80      A    C
ATOM    217  CD2 HIS    87      22.930 -21.773 -11.937  1.00133.11      A    C
ATOM    218  ND1 HIS    87      22.017 -22.507 -13.783  1.00133.19      A    N
ATOM    219  CE1 HIS    87      22.867 -23.433 -13.366  1.00134.19      A    C
ATOM    220  NE2 HIS    87      23.431 -23.012 -12.252  1.00134.17      A    N
ATOM    221  C   HIS    87      20.986 -17.836 -12.473  1.00123.89      A    C
ATOM    222  O   HIS    87      19.822 -17.466 -12.361  1.00123.47      A    O
ATOM    223  N   LEU    88      21.939 -17.106 -13.045  1.00123.03      A    N
ATOM    224  CA  LEU    88      21.688 -15.773 -13.574  1.00122.10      A    C
ATOM    225  CB  LEU    88      22.979 -15.195 -14.162  1.00124.68      A    C
ATOM    226  CG  LEU    88      24.215 -15.377 -13.273  1.00126.86      A    C
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ATOM    227  CD1 LEU    88      25.432 -14.711 -13.909  1.00127.21      A    C
ATOM    228  CD2 LEU    88      23.935 -14.794 -11.890  1.00128.06      A    C
ATOM    229  C   LEU    88      20.603 -15.835 -14.641  1.00119.59      A    C
ATOM    230  O   LEU    88      19.887 -14.861 -14.882  1.00118.95      A    O
ATOM    231  N   SER    89      20.495 -16.992 -15.282  1.00117.17      A    N
ATOM    232  CA  SER    89      19.381 -17.268 -16.174  1.00114.74      A    C
ATOM    233  CB  SER    89      19.550 -18.648 -16.814  1.00115.37      A    C
ATOM    234  OG  SER    89      19.719 -19.653 -15.825  1.00116.88      A    O
ATOM    235  C   SER    89      18.111 -17.236 -15.341  1.00112.77      A    C
ATOM    236  O   SER    89      17.099 -16.668 -15.752  1.00111.49      A    O
ATOM    237  N   GLN    90      18.185 -17.840 -14.158  1.00110.68      A    N
ATOM    238  CA  GLN    90      17.033 -17.967 -13.279  1.00108.49      A    C
ATOM    239  CB  GLN    90      17.300 -19.042 -12.223  1.00108.59      A    C
ATOM    240  CG  GLN    90      17.416 -20.449 -12.811  1.00109.07      A    C
ATOM    241  CD  GLN    90      17.691 -21.518 -11.764  1.00109.15      A    C
ATOM    242  OE1 GLN    90      18.790 -21.596 -11.212  1.00109.15      A    O
ATOM    243  NE2 GLN    90      16.693 -22.353 -11.491  1.00108.63      A    N
ATOM    244  C   GLN    90      16.680 -16.642 -12.611  1.00107.20      A    C
ATOM    245  O   GLN    90      15.513 -16.250 -12.589  1.00106.82      A    O
ATOM    246  N   SER    91      17.683 -15.949 -12.078  1.00105.29      A    N
ATOM    247  CA  SER    91      17.472 -14.595 -11.569  1.00103.51      A    C
ATOM    248  CB  SER    91      18.816 -13.871 -11.370  1.00102.29      A    C
ATOM    249  OG  SER    91      19.158 -13.724 -10.000  1.00 98.21      A    O
ATOM    250  C   SER    91      16.630 -13.833 -12.588  1.00103.23      A    C
ATOM    251  O   SER    91      15.618 -13.227 -12.248  1.00102.98      A    O
ATOM    252  N   GLU    92      17.045 -13.889 -13.847  1.00103.82      A    N
ATOM    253  CA  GLU    92      16.451 -13.064 -14.884  1.00104.85      A    C
ATOM    254  CB  GLU    92      17.181 -13.299 -16.211  1.00106.74      A    C
ATOM    255  CG  GLU    92      16.427 -12.816 -17.447  1.00111.04      A    C
ATOM    256  CD  GLU    92      15.480 -13.870 -18.016  1.00112.76      A    C
ATOM    257  OE1 GLU    92      14.269 -13.583 -18.143  1.00114.41      A    O
ATOM    258  OE2 GLU    92      15.948 -14.985 -18.336  1.00114.56      A    O
ATOM    259  C   GLU    92      14.953 -13.302 -15.055  1.00104.14      A    C
ATOM    260  O   GLU    92      14.180 -12.350 -15.106  1.00104.75      A    O
ATOM    261  N   ARG    93      14.540 -14.566 -15.137  1.00103.50      A    N
ATOM    262  CA  ARG    93      13.145 -14.899 -15.445  1.00101.47      A    C
ATOM    263  CB  ARG    93      13.025 -16.375 -15.844  1.00106.48      A    C
ATOM    264  CG  ARG    93      13.405 -16.665 -17.298  1.00113.16      A    C
ATOM    265  CD  ARG    93      13.327 -18.161 -17.623  1.00120.42      A    C
ATOM    266  NE  ARG    93      14.627 -18.826 -17.516  1.00128.11      A    N
ATOM    267  CZ  ARG    93      14.826 -19.991 -16.906  1.00131.80      A    C
ATOM    268  NH1 ARG    93      16.044 -20.518 -16.858  1.00134.30      A    N
ATOM    269  NH2 ARG    93      13.807 -20.629 -16.345  1.00134.64      A    N
ATOM    270  C   ARG    93      12.196 -14.600 -14.286  1.00 96.81      A    C
ATOM    271  O   ARG    93      11.139 -13.990 -14.472  1.00 95.99      A    O
ATOM    272  N   THR    94      12.577 -15.028 -13.089  1.00 91.16      A    N
ATOM    273  CA  THR    94      11.852 -14.643 -11.890  1.00 86.23      A    C
ATOM    274  CB  THR    94      12.723 -14.824 -10.640  1.00 83.21      A    C
ATOM    275  OG1 THR    94      13.153 -16.185 -10.554  1.00 80.97      A    O
ATOM    276  CG2 THR    94      11.949 -14.461  -9.388  1.00 80.14      A    C
ATOM    277  C   THR    94      11.478 -13.171 -12.016  1.00 85.96      A    C
ATOM    278  O   THR    94      10.414 -12.742 -11.577  1.00 85.37      A    O
ATOM    279  N   ALA    95      12.361 -12.404 -12.641  1.00 85.71      A    N
ATOM    280  CA  ALA    95      12.224 -10.960 -12.685  1.00 87.24      A    C
ATOM    281  CB  ALA    95      13.581 -10.329 -12.931  1.00 85.86      A    C
ATOM    282  C   ALA    95      11.239 -10.515 -13.758  1.00 88.89      A    C
ATOM    283  O   ALA    95      10.460  -9.577 -13.554  1.00 88.87      A    O
ATOM    284  N   ARG    96      11.282 -11.178 -14.909  1.00 91.47      A    N
ATOM    285  CA  ARG    96      10.322 -10.896 -15.967  1.00 93.56      A    C
ATOM    286  CB  ARG    96      10.772 -11.535 -17.286  1.00 97.53      A    C
ATOM    287  CG  ARG    96      12.113 -11.029 -17.803  1.00102.81      A    C
ATOM    288  CD  ARG    96      12.042  -9.567 -18.230  1.00108.59      A    C
ATOM    289  NE  ARG    96      13.359  -9.039 -18.581  1.00114.78      A    N
ATOM    290  CZ  ARG    96      13.603  -7.767 -18.876  1.00118.68      A    C
ATOM    291  NH1 ARG    96      14.833  -7.380 -19.182  1.00121.30      A    N
ATOM    292  NH2 ARG    96      12.617  -6.880 -18.864  1.00121.58      A    N
ATOM    293  C   ARG    96       8.985 -11.477 -15.528  1.00 92.22      A    C
ATOM    294  O   ARG    96       7.922 -10.918 -15.818  1.00 91.98      A    O
ATOM    295  N   ARG    97       9.058 -12.595 -14.807  1.00 89.71      A    N
ATOM    296  CA  ARG    97       7.876 -13.237 -14.245  1.00 87.42      A    C
ATOM    297  CB  ARG    97       8.281 -14.483 -13.453  1.00 90.04      A    C
ATOM    298  CG  ARG    97       7.103 -15.307 -12.940  1.00 94.32      A    C
ATOM    299  CD  ARG    97       7.554 -16.678 -12.440  1.00 98.95      A    C
ATOM    300  NE  ARG    97       6.539 -17.704 -12.666  1.00104.16      A    N
ATOM    301  CZ  ARG    97       6.496 -18.489 -13.739  1.00107.61      A    C
ATOM    302  NH1 ARG    97       5.533 -19.394 -13.860  1.00110.25      A    N
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ATOM	303	NH2	ARG	97	7.415	-18.372	-14.690	1.00	109.39	A	N
ATOM	304	C	ARG	97	7.126	-12.271	-13.334	1.00	83.61	A	C
ATOM	305	O	ARG	97	5.972	-11.936	-13.582	1.00	83.78	A	O
ATOM	306	N	LEU	98	7.799	-11.822	-12.279	1.00	78.62	A	N
ATOM	307	CA	LEU	98	7.213	-10.848	-11.376	1.00	73.43	A	C
ATOM	308	CB	LEU	98	8.281	-10.279	-10.446	1.00	72.56	A	C
ATOM	309	CG	LEU	98	8.084	-8.825	-10.006	1.00	71.98	A	C
ATOM	310	CD1	LEU	98	6.792	-8.673	-9.218	1.00	70.37	A	C
ATOM	311	CD2	LEU	98	9.282	-8.400	-9.171	1.00	71.13	A	C
ATOM	312	C	LEU	98	6.592	-9.729	-12.184	1.00	70.64	A	C
ATOM	313	O	LEU	98	5.426	-9.397	-12.011	1.00	69.36	A	O
ATOM	314	N	GLN	99	7.376	-9.152	-13.080	1.00	69.00	A	N
ATOM	315	CA	GLN	99	6.895	-8.026	-13.861	1.00	67.29	A	C
ATOM	316	CB	GLN	99	7.986	-7.537	-14.808	1.00	67.31	A	C
ATOM	317	CG	GLN	99	7.697	-6.182	-15.420	1.00	66.57	A	C
ATOM	318	CD	GLN	99	8.960	-5.387	-15.623	1.00	66.92	A	C
ATOM	319	OE1	GLN	99	8.930	-4.238	-16.068	1.00	67.28	A	O
ATOM	320	NE2	GLN	99	10.092	-6.001	-15.294	1.00	66.83	A	N
ATOM	321	C	GLN	99	5.671	-8.450	-14.660	1.00	65.61	A	C
ATOM	322	O	GLN	99	4.781	-7.641	-14.931	1.00	64.48	A	O
ATOM	323	N	ALA	100	5.645	-9.732	-15.024	1.00	64.28	A	N
ATOM	324	CA	ALA	100	4.546	-10.316	-15.793	1.00	62.89	A	C
ATOM	325	CB	ALA	100	4.904	-11.749	-16.201	1.00	61.93	A	C
ATOM	326	C	ALA	100	3.248	-10.309	-14.980	1.00	62.26	A	C
ATOM	327	O	ALA	100	2.250	-9.671	-15.364	1.00	60.92	A	O
ATOM	328	N	GLN	101	3.271	-11.017	-13.856	1.00	59.83	A	N
ATOM	329	CA	GLN	101	2.189	-10.931	-12.897	1.00	58.75	A	C
ATOM	330	CB	GLN	101	2.619	-11.603	-11.610	1.00	56.30	A	C
ATOM	331	CG	GLN	101	3.498	-12.803	-11.852	1.00	55.38	A	C
ATOM	332	CD	GLN	101	3.346	-13.837	-10.770	1.00	55.44	A	C
ATOM	333	OE1	GLN	101	2.402	-13.785	-9.989	1.00	57.12	A	O
ATOM	334	NE2	GLN	101	4.275	-14.787	-10.712	1.00	55.95	A	N
ATOM	335	C	GLN	101	1.888	-9.457	-12.638	1.00	59.50	A	C
ATOM	336	O	GLN	101	0.918	-8.904	-13.158	1.00	60.05	A	O
ATOM	337	N	ALA	102	2.745	-8.826	-11.843	1.00	59.21	A	N
ATOM	338	CA	ALA	102	2.649	-7.403	-11.557	1.00	59.10	A	C
ATOM	339	CB	ALA	102	4.030	-6.840	-11.318	1.00	56.43	A	C
ATOM	340	C	ALA	102	1.974	-6.634	-12.681	1.00	60.79	A	C
ATOM	341	O	ALA	102	1.054	-5.848	-12.447	1.00	60.89	A	O
ATOM	342	N	ALA	103	2.440	-6.857	-13.904	1.00	63.30	A	N
ATOM	343	CA	ALA	103	1.897	-6.143	-15.052	1.00	64.47	A	C
ATOM	344	CB	ALA	103	2.489	-6.684	-16.347	1.00	65.31	A	C
ATOM	345	C	ALA	103	0.392	-6.315	-15.061	1.00	64.88	A	C
ATOM	346	O	ALA	103	-0.346	-5.346	-15.236	1.00	64.54	A	O
ATOM	347	N	ARG	104	-0.052	-7.554	-14.848	1.00	65.14	A	N
ATOM	348	CA	ARG	104	-1.458	-7.918	-15.003	1.00	65.17	A	C
ATOM	349	CB	ARG	104	-1.598	-9.440	-15.097	1.00	63.37	A	C
ATOM	350	CG	ARG	104	-1.198	-9.991	-16.454	1.00	62.23	A	C
ATOM	351	CD	ARG	104	-1.540	-11.459	-16.593	1.00	61.80	A	C
ATOM	352	NE	ARG	104	-0.642	-12.339	-15.846	1.00	61.29	A	N
ATOM	353	CZ	ARG	104	0.397	-12.968	-16.387	1.00	61.76	A	C
ATOM	354	NH1	ARG	104	1.164	-13.761	-15.644	1.00	61.55	A	N
ATOM	355	NH2	ARG	104	0.675	-12.794	-17.674	1.00	61.90	A	N
ATOM	356	C	ARG	104	-2.373	-7.381	-13.904	1.00	66.10	A	C
ATOM	357	O	ARG	104	-3.469	-6.886	-14.195	1.00	65.62	A	O
ATOM	358	N	ARG	105	-1.935	-7.473	-12.650	1.00	66.65	A	N
ATOM	359	CA	ARG	105	-2.724	-6.935	-11.546	1.00	66.20	A	C
ATOM	360	CB	ARG	105	-2.048	-7.221	-10.211	1.00	64.90	A	C
ATOM	361	CG	ARG	105	-2.350	-8.581	-9.638	1.00	65.02	A	C
ATOM	362	CD	ARG	105	-1.519	-9.681	-10.267	1.00	66.10	A	C
ATOM	363	NE	ARG	105	-1.460	-10.832	-9.365	1.00	70.31	A	N
ATOM	364	CZ	ARG	105	-1.306	-12.102	-9.742	1.00	72.23	A	C
ATOM	365	NH1	ARG	105	-1.267	-13.060	-8.821	1.00	73.45	A	N
ATOM	366	NH2	ARG	105	-1.191	-12.421	-11.028	1.00	73.04	A	N
ATOM	367	C	ARG	105	-2.925	-5.437	-11.703	1.00	66.79	A	C
ATOM	368	O	ARG	105	-3.706	-4.833	-10.979	1.00	67.02	A	O
ATOM	369	N	GLY	106	-2.212	-4.836	-12.650	1.00	69.01	A	N
ATOM	370	CA	GLY	106	-2.379	-3.415	-12.913	1.00	72.19	A	C
ATOM	371	C	GLY	106	-1.244	-2.580	-12.353	1.00	73.50	A	C
ATOM	372	O	GLY	106	-1.375	-1.367	-12.157	1.00	73.14	A	O
ATOM	373	N	TYR	107	-0.126	-3.245	-12.086	1.00	75.51	A	N
ATOM	374	CA	TYR	107	1.040	-2.590	-11.525	1.00	78.05	A	C
ATOM	375	CB	TYR	107	1.454	-3.285	-10.236	1.00	77.61	A	C
ATOM	376	CG	TYR	107	0.475	-3.021	-9.137	1.00	77.39	A	C
ATOM	377	CD1	TYR	107	0.173	-1.725	-8.766	1.00	77.40	A	C
ATOM	378	CE1	TYR	107	-0.771	-1.464	-7.808	1.00	78.07	A	C

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ATOM    379  CD2 TYR   107      -0.195  -4.058  -8.512  1.00 78.22      A    C
ATOM    380  CE2 TYR   107      -1.147  -3.812  -7.550  1.00 77.52      A    C
ATOM    381  CZ  TYR   107      -1.432  -2.512  -7.203  1.00 78.47      A    C
ATOM    382  OH  TYR   107      -2.392  -2.253  -6.255  1.00 78.73      A    O
ATOM    383  C   TYR   107       2.214  -2.548 -12.485  1.00 80.62      A    C
ATOM    384  O   TYR   107       2.787  -3.583 -12.849  1.00 79.63      A    O
ATOM    385  N   LEU   108       2.565  -1.332 -12.889  1.00 83.48      A    N
ATOM    386  CA  LEU   108       3.802  -1.095 -13.610  1.00 84.85      A    C
ATOM    387  CB  LEU   108       3.848   0.358 -14.084  1.00 86.63      A    C
ATOM    388  CG  LEU   108       4.652   0.694 -15.342  1.00 88.85      A    C
ATOM    389  CD1 LEU   108       4.615   2.204 -15.564  1.00 89.67      A    C
ATOM    390  CD2 LEU   108       6.082   0.210 -15.194  1.00 89.84      A    C
ATOM    391  C   LEU   108       4.937  -1.370 -12.624  1.00 84.33      A    C
ATOM    392  O   LEU   108       4.827  -1.060 -11.440  1.00 85.33      A    O
ATOM    393  N   THR   109       6.018  -1.964 -13.103  1.00 82.38      A    N
ATOM    394  CA  THR   109       7.155  -2.227 -12.245  1.00 82.68      A    C
ATOM    395  CB  THR   109       7.104  -3.653 -11.700  1.00 84.37      A    C
ATOM    396  OG1 THR   109       7.043  -4.581 -12.789  1.00 86.52      A    O
ATOM    397  CG2 THR   109       5.897  -3.823 -10.830  1.00 85.93      A    C
ATOM    398  C   THR   109       8.461  -2.033 -13.000  1.00 81.95      A    C
ATOM    399  O   THR   109       8.508  -2.201 -14.222  1.00 79.46      A    O
ATOM    400  N   LYS   110       9.521  -1.676 -12.275  1.00 81.51      A    N
ATOM    401  CA  LYS   110      10.832  -1.569 -12.890  1.00 81.24      A    C
ATOM    402  CB  LYS   110      11.340  -0.129 -12.852  1.00 83.45      A    C
ATOM    403  CG  LYS   110      12.617   0.074 -13.662  1.00 88.37      A    C
ATOM    404  CD  LYS   110      13.088   1.523 -13.651  1.00 92.43      A    C
ATOM    405  CE  LYS   110      12.099   2.444 -14.332  1.00 95.58      A    C
ATOM    406  NZ  LYS   110      12.518   3.867 -14.197  1.00100.67      A    N
ATOM    407  C   LYS   110      11.854  -2.475 -12.241  1.00 79.14      A    C
ATOM    408  O   LYS   110      11.905  -2.610 -11.022  1.00 79.79      A    O
ATOM    409  N   ILE   111      12.666  -3.107 -13.076  1.00 75.80      A    N
ATOM    410  CA  ILE   111      13.853  -3.785 -12.600  1.00 72.25      A    C
ATOM    411  CB  ILE   111      14.150  -5.029 -13.430  1.00 71.02      A    C
ATOM    412  CG2 ILE   111      15.397  -5.711 -12.898  1.00 69.49      A    C
ATOM    413  CG1 ILE   111      12.952  -5.978 -13.391  1.00 69.43      A    C
ATOM    414  CD1 ILE   111      12.768  -6.686 -12.071  1.00 67.40      A    C
ATOM    415  C   ILE   111      15.024  -2.818 -12.738  1.00 71.50      A    C
ATOM    416  O   ILE   111      15.347  -2.371 -13.841  1.00 71.60      A    O
ATOM    417  N   LEU   112      15.655  -2.494 -11.615  1.00 69.75      A    N
ATOM    418  CA  LEU   112      16.715  -1.496 -11.598  1.00 67.25      A    C
ATOM    419  CB  LEU   112      16.689  -0.726 -10.279  1.00 65.97      A    C
ATOM    420  CG  LEU   112      15.329  -0.103  -9.976  1.00 66.90      A    C
ATOM    421  CD1 LEU   112      15.357   0.600  -8.627  1.00 67.88      A    C
ATOM    422  CD2 LEU   112      14.962   0.850 -11.092  1.00 65.29      A    C
ATOM    423  C   LEU   112      18.069  -2.166 -11.778  1.00 66.52      A    C
ATOM    424  O   LEU   112      18.996  -1.589 -12.355  1.00 65.11      A    O
ATOM    425  N   HIS   113      18.184  -3.387 -11.278  1.00 65.05      A    N
ATOM    426  CA  HIS   113      19.458  -4.064 -11.327  1.00 66.25      A    C
ATOM    427  CB  HIS   113      20.428  -3.433 -10.330  1.00 65.11      A    C
ATOM    428  CG  HIS   113      21.807  -4.010 -10.387  1.00 63.95      A    C
ATOM    429  CD2 HIS   113      22.500  -4.757  -9.495  1.00 63.35      A    C
ATOM    430  ND1 HIS   113      22.638  -3.841 -11.473  1.00 63.07      A    N
ATOM    431  CE1 HIS   113      23.783  -4.459 -11.248  1.00 63.01      A    C
ATOM    432  NE2 HIS   113      23.725  -5.023 -10.056  1.00 63.38      A    N
ATOM    433  C   HIS   113      19.321  -5.543 -11.035  1.00 67.36      A    C
ATOM    434  O   HIS   113      18.547  -5.953 -10.176  1.00 67.90      A    O
ATOM    435  N   VAL   114      20.087  -6.341 -11.760  1.00 68.11      A    N
ATOM    436  CA  VAL   114      20.031  -7.774 -11.611  1.00 69.21      A    C
ATOM    437  CB  VAL   114      19.688  -8.443 -12.955  1.00 68.54      A    C
ATOM    438  CG1 VAL   114      19.579  -9.949 -12.775  1.00 67.55      A    C
ATOM    439  CG2 VAL   114      18.388  -7.861 -13.507  1.00 64.17      A    C
ATOM    440  C   VAL   114      21.387  -8.244 -11.129  1.00 71.04      A    C
ATOM    441  O   VAL   114      22.392  -8.048 -11.804  1.00 71.37      A    O
ATOM    442  N   PHE   115      21.415  -8.852  -9.953  1.00 73.34      A    N
ATOM    443  CA  PHE   115      22.674  -9.255  -9.353  1.00 76.69      A    C
ATOM    444  CB  PHE   115      22.500  -9.484  -7.856  1.00 77.67      A    C
ATOM    445  CG  PHE   115      22.268  -8.233  -7.087  1.00 79.20      A    C
ATOM    446  CD1 PHE   115      20.986  -7.852  -6.737  1.00 79.95      A    C
ATOM    447  CD2 PHE   115      23.330  -7.423  -6.734  1.00 79.39      A    C
ATOM    448  CE1 PHE   115      20.766  -6.686  -6.052  1.00 81.16      A    C
ATOM    449  CE2 PHE   115      23.120  -6.250  -6.046  1.00 80.76      A    C
ATOM    450  CZ  PHE   115      21.838  -5.880  -5.704  1.00 82.11      A    C
ATOM    451  C   PHE   115      23.217 -10.518  -9.981  1.00 78.95      A    C
ATOM    452  O   PHE   115      22.708 -11.618  -9.739  1.00 79.39      A    O
ATOM    453  N   HIS   116      24.257 -10.364 -10.791  1.00 81.04      A    N
ATOM    454  CA  HIS   116      25.087 -11.504 -11.131  1.00 82.56      A    C
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ATOM    455  CB  HIS   116      25.073 -11.768 -12.647  1.00 86.34      A    C
ATOM    456  CG  HIS   116      25.224 -10.544 -13.498  1.00 89.89      A    C
ATOM    457  CD2 HIS   116      26.095 -10.260 -14.496  1.00 91.04      A    C
ATOM    458  ND1 HIS   116      24.360  -9.472 -13.428  1.00 91.27      A    N
ATOM    459  CE1 HIS   116      24.689  -8.583 -14.349  1.00 92.06      A    C
ATOM    460  NE2 HIS   116      25.737  -9.038 -15.011  1.00 91.87      A    N
ATOM    461  C   HIS   116      26.510 -11.314 -10.621  1.00 80.27      A    C
ATOM    462  O   HIS   116      27.413 -10.946 -11.370  1.00 80.70      A    O
ATOM    463  N   GLY   117      26.696 -11.572  -9.330  1.00 76.62      A    N
ATOM    464  CA  GLY   117      27.997 -11.369  -8.729  1.00 72.53      A    C
ATOM    465  C   GLY   117      28.215 -12.075  -7.406  1.00 69.71      A    C
ATOM    466  O   GLY   117      28.740 -13.186  -7.363  1.00 68.94      A    O
ATOM    467  N   LEU   118      27.834 -11.437  -6.310  1.00 67.10      A    N
ATOM    468  CA  LEU   118      28.133 -12.025  -5.019  1.00 65.67      A    C
ATOM    469  CB  LEU   118      28.931 -11.040  -4.154  1.00 64.83      A    C
ATOM    470  CG  LEU   118      28.219  -9.904  -3.427  1.00 62.63      A    C
ATOM    471  CD1 LEU   118      29.171  -9.337  -2.418  1.00 61.32      A    C
ATOM    472  CD2 LEU   118      27.746  -8.834  -4.398  1.00 62.31      A    C
ATOM    473  C   LEU   118      26.845 -12.441  -4.333  1.00 64.76      A    C
ATOM    474  O   LEU   118      26.839 -12.867  -3.180  1.00 65.29      A    O
ATOM    475  N   LEU   119      25.749 -12.334  -5.071  1.00 63.71      A    N
ATOM    476  CA  LEU   119      24.451 -12.710  -4.548  1.00 62.88      A    C
ATOM    477  CB  LEU   119      23.998 -11.653  -3.549  1.00 65.56      A    C
ATOM    478  CG  LEU   119      22.666 -11.858  -2.840  1.00 66.68      A    C
ATOM    479  CD1 LEU   119      22.840 -12.943  -1.799  1.00 68.24      A    C
ATOM    480  CD2 LEU   119      22.226 -10.553  -2.188  1.00 67.19      A    C
ATOM    481  C   LEU   119      23.424 -12.846  -5.682  1.00 61.76      A    C
ATOM    482  O   LEU   119      23.265 -11.942  -6.505  1.00 61.17      A    O
ATOM    483  N   PRO   120      22.717 -13.988  -5.739  1.00 60.22      A    N
ATOM    484  CD  PRO   120      22.785 -15.117  -4.795  1.00 59.04      A    C
ATOM    485  CA  PRO   120      21.706 -14.209  -6.774  1.00 58.33      A    C
ATOM    486  CB  PRO   120      21.360 -15.677  -6.621  1.00 57.41      A    C
ATOM    487  CG  PRO   120      21.598 -15.948  -5.181  1.00 58.24      A    C
ATOM    488  C   PRO   120      20.513 -13.336  -6.472  1.00 58.32      A    C
ATOM    489  O   PRO   120      20.147 -13.197  -5.301  1.00 60.08      A    O
ATOM    490  N   GLY   121      19.906 -12.754  -7.507  1.00 55.72      A    N
ATOM    491  CA  GLY   121      18.684 -11.998  -7.301  1.00 51.86      A    C
ATOM    492  C   GLY   121      18.687 -10.663  -8.012  1.00 51.21      A    C
ATOM    493  O   GLY   121      19.580 -10.378  -8.803  1.00 51.96      A    O
ATOM    494  N   PHE   122      17.689  -9.833  -7.737  1.00 49.58      A    N
ATOM    495  CA  PHE   122      17.612  -8.536  -8.383  1.00 47.46      A    C
ATOM    496  CB  PHE   122      16.813  -8.664  -9.676  1.00 45.66      A    C
ATOM    497  CG  PHE   122      15.418  -9.178  -9.473  1.00 44.33      A    C
ATOM    498  CD1 PHE   122      14.341  -8.307  -9.473  1.00 42.51      A    C
ATOM    499  CD2 PHE   122      15.182 -10.536  -9.268  1.00 41.45      A    C
ATOM    500  CE1 PHE   122      13.056  -8.787  -9.273  1.00 42.67      A    C
ATOM    501  CE2 PHE   122      13.906 -11.013  -9.070  1.00 37.88      A    C
ATOM    502  CZ  PHE   122      12.843 -10.143  -9.071  1.00 38.44      A    C
ATOM    503  C   PHE   122      16.990  -7.463  -7.486  1.00 47.82      A    C
ATOM    504  O   PHE   122      16.384  -7.773  -6.455  1.00 46.93      A    O
ATOM    505  N   LEU   123      17.160  -6.201  -7.882  1.00 48.30      A    N
ATOM    506  CA  LEU   123      16.489  -5.073  -7.229  1.00 49.55      A    C
ATOM    507  CB  LEU   123      17.468  -3.902  -6.993  1.00 50.43      A    C
ATOM    508  CG  LEU   123      16.892  -2.637  -6.317  1.00 50.76      A    C
ATOM    509  CD1 LEU   123      16.958  -2.807  -4.803  1.00 50.11      A    C
ATOM    510  CD2 LEU   123      17.666  -1.383  -6.744  1.00 47.52      A    C
ATOM    511  C   LEU   123      15.342  -4.592  -8.109  1.00 49.64      A    C
ATOM    512  O   LEU   123      15.496  -4.450  -9.326  1.00 49.87      A    O
ATOM    513  N   VAL   124      14.199  -4.331  -7.483  1.00 50.02      A    N
ATOM    514  CA  VAL   124      12.982  -3.987  -8.203  1.00 49.18      A    C
ATOM    515  CB  VAL   124      11.981  -5.159  -8.182  1.00 47.70      A    C
ATOM    516  CG1 VAL   124      11.675  -5.567  -6.728  1.00 43.41      A    C
ATOM    517  CG2 VAL   124      10.692  -4.746  -8.896  1.00 46.32      A    C
ATOM    518  C   VAL   124      12.298  -2.774  -7.587  1.00 50.07      A    C
ATOM    519  O   VAL   124      12.255  -2.628  -6.362  1.00 49.64      A    O
ATOM    520  N   LYS   125      11.751  -1.921  -8.447  1.00 50.54      A    N
ATOM    521  CA  LYS   125      10.944  -0.785  -8.021  1.00 51.53      A    C
ATOM    522  CB  LYS   125      11.382   0.467  -8.796  1.00 53.26      A    C
ATOM    523  CG  LYS   125      10.786   1.778  -8.315  1.00 55.05      A    C
ATOM    524  CD  LYS   125      11.670   2.951  -8.741  1.00 58.38      A    C
ATOM    525  CE  LYS   125      11.108   4.289  -8.255  1.00 60.15      A    C
ATOM    526  NZ  LYS   125      11.497   5.417  -9.155  1.00 61.63      A    N
ATOM    527  C   LYS   125       9.462  -1.088  -8.284  1.00 51.78      A    C
ATOM    528  O   LYS   125       8.991  -1.027  -9.426  1.00 52.65      A    O
ATOM    529  N   MET   126       8.733  -1.431  -7.225  1.00 51.29      A    N
ATOM    530  CA  MET   126       7.319  -1.765  -7.342  1.00 49.75      A    C
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ATOM    531  CB  MET   126       7.119  -3.285  -7.354  1.00 49.94      A    C
ATOM    532  CG  MET   126       7.627  -4.011  -6.116  1.00 51.30      A    C
ATOM    533  SD  MET   126       7.626  -5.837  -6.339  1.00 56.10      A    S
ATOM    534  CE  MET   126       7.973  -6.399  -4.675  1.00 54.16      A    C
ATOM    535  C   MET   126       6.565  -1.169  -6.172  1.00 49.16      A    C
ATOM    536  O   MET   126       7.180  -0.762  -5.185  1.00 47.47      A    O
ATOM    537  N   SER   127       5.236  -1.117  -6.287  1.00 48.54      A    N
ATOM    538  CA  SER   127       4.383  -0.842  -5.137  1.00 46.28      A    C
ATOM    539  CB  SER   127       2.917  -0.775  -5.537  1.00 47.48      A    C
ATOM    540  OG  SER   127       2.103  -0.803  -4.375  1.00 47.10      A    O
ATOM    541  C   SER   127       4.540  -1.934  -4.096  1.00 44.81      A    C
ATOM    542  O   SER   127       4.769  -3.094  -4.436  1.00 44.11      A    O
ATOM    543  N   GLY   128       4.416  -1.548  -2.828  1.00 42.54      A    N
ATOM    544  CA  GLY   128       4.503  -2.504  -1.748  1.00 39.84      A    C
ATOM    545  C   GLY   128       3.353  -3.484  -1.829  1.00 40.64      A    C
ATOM    546  O   GLY   128       3.475  -4.625  -1.378  1.00 40.33      A    O
ATOM    547  N   ASP   129       2.241  -3.050  -2.417  1.00 40.43      A    N
ATOM    548  CA  ASP   129       1.109  -3.932  -2.637  1.00 41.66      A    C
ATOM    549  CB  ASP   129       0.170  -3.337  -3.683  1.00 41.97      A    C
ATOM    550  CG  ASP   129      -0.675  -2.192  -3.133  1.00 45.41      A    C
ATOM    551  OD1 ASP   129      -0.446  -1.771  -1.978  1.00 48.87      A    O
ATOM    552  OD2 ASP   129      -1.577  -1.708  -3.853  1.00 45.68      A    O
ATOM    553  C   ASP   129       1.570  -5.311  -3.093  1.00 43.90      A    C
ATOM    554  O   ASP   129       0.985  -6.325  -2.704  1.00 46.09      A    O
ATOM    555  N   LEU   130       2.627  -5.360  -3.898  1.00 43.76      A    N
ATOM    556  CA  LEU   130       3.029  -6.610  -4.533  1.00 45.35      A    C
ATOM    557  CB  LEU   130       3.659  -6.327  -5.905  1.00 43.26      A    C
ATOM    558  CG  LEU   130       2.677  -5.821  -6.978  1.00 43.78      A    C
ATOM    559  CD1 LEU   130       3.369  -4.889  -7.958  1.00 42.09      A    C
ATOM    560  CD2 LEU   130       2.057  -7.012  -7.700  1.00 40.85      A    C
ATOM    561  C   LEU   130       3.985  -7.428  -3.676  1.00 48.24      A    C
ATOM    562  O   LEU   130       4.656  -8.344  -4.167  1.00 47.63      A    O
ATOM    563  N   LEU   131       4.054  -7.116  -2.390  1.00 52.01      A    N
ATOM    564  CA  LEU   131       4.981  -7.844  -1.541  1.00 56.99      A    C
ATOM    565  CB  LEU   131       5.274  -7.056  -0.269  1.00 55.21      A    C
ATOM    566  CG  LEU   131       6.139  -5.830  -0.544  1.00 53.21      A    C
ATOM    567  CD1 LEU   131       5.883  -4.777   0.502  1.00 51.08      A    C
ATOM    568  CD2 LEU   131       7.591  -6.244  -0.589  1.00 51.08      A    C
ATOM    569  C   LEU   131       4.442  -9.217  -1.199  1.00 61.74      A    C
ATOM    570  O   LEU   131       5.199 -10.188  -1.150  1.00 62.05      A    O
ATOM    571  N   GLU   132       3.133  -9.303  -0.969  1.00 67.01      A    N
ATOM    572  CA  GLU   132       2.494 -10.600  -0.757  1.00 70.97      A    C
ATOM    573  CB  GLU   132       0.973 -10.437  -0.698  1.00 76.59      A    C
ATOM    574  CG  GLU   132       0.198 -11.744  -0.504  1.00 85.24      A    C
ATOM    575  CD  GLU   132      -1.324 -11.542  -0.472  1.00 90.08      A    C
ATOM    576  OE1 GLU   132      -2.060 -12.555  -0.564  1.00 93.16      A    O
ATOM    577  OE2 GLU   132      -1.782 -10.376  -0.354  1.00 91.94      A    O
ATOM    578  C   GLU   132       2.870 -11.473  -1.946  1.00 71.12      A    C
ATOM    579  O   GLU   132       3.437 -12.563  -1.797  1.00 70.53      A    O
ATOM    580  N   LEU   133       2.574 -10.948  -3.133  1.00 70.73      A    N
ATOM    581  CA  LEU   133       2.831 -11.636  -4.390  1.00 70.60      A    C
ATOM    582  CB  LEU   133       2.546 -10.687  -5.554  1.00 67.75      A    C
ATOM    583  CG  LEU   133       1.958 -11.328  -6.807  1.00 66.81      A    C
ATOM    584  CD1 LEU   133       1.959 -10.320  -7.933  1.00 65.64      A    C
ATOM    585  CD2 LEU   133       2.765 -12.559  -7.190  1.00 66.81      A    C
ATOM    586  C   LEU   133       4.264 -12.162  -4.494  1.00 72.24      A    C
ATOM    587  O   LEU   133       4.491 -13.370  -4.554  1.00 73.30      A    O
ATOM    588  N   ALA   134       5.230 -11.248  -4.504  1.00 73.74      A    N
ATOM    589  CA  ALA   134       6.602 -11.575  -4.885  1.00 74.48      A    C
ATOM    590  CB  ALA   134       7.446 -10.314  -4.889  1.00 75.03      A    C
ATOM    591  C   ALA   134       7.255 -12.628  -4.001  1.00 75.73      A    C
ATOM    592  O   ALA   134       8.214 -13.278  -4.411  1.00 73.90      A    O
ATOM    593  N   LEU   135       6.743 -12.791  -2.786  1.00 78.60      A    N
ATOM    594  CA  LEU   135       7.309 -13.761  -1.860  1.00 81.85      A    C
ATOM    595  CB  LEU   135       6.842 -13.463  -0.439  1.00 78.00      A    C
ATOM    596  CG  LEU   135       7.491 -12.298   0.306  1.00 74.37      A    C
ATOM    597  CD1 LEU   135       6.520 -11.772   1.325  1.00 71.67      A    C
ATOM    598  CD2 LEU   135       8.775 -12.745   0.971  1.00 72.66      A    C
ATOM    599  C   LEU   135       6.876 -15.164  -2.258  1.00 86.28      A    C
ATOM    600  O   LEU   135       7.490 -16.155  -1.857  1.00 86.58      A    O
ATOM    601  N   LYS   136       5.807 -15.237  -3.047  1.00 92.64      A    N
ATOM    602  CA  LYS   136       5.318 -16.511  -3.560  1.00 99.95      A    C
ATOM    603  CB  LYS   136       3.922 -16.342  -4.172  1.00101.68      A    C
ATOM    604  CG  LYS   136       2.818 -16.042  -3.164  1.00104.97      A    C
ATOM    605  CD  LYS   136       1.451 -16.030  -3.834  1.00108.72      A    C
ATOM    606  CE  LYS   136       0.374 -15.490  -2.906  1.00112.39      A    C
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ATOM    607  NZ  LYS   136       0.413 -16.148  -1.572 1.00114.91      A    N
ATOM    608  C   LYS   136       6.276 -17.056  -4.613 1.00104.36      A    C
ATOM    609  O   LYS   136       6.599 -18.246  -4.616 1.00104.13      A    O
ATOM    610  N   LEU   137       6.735 -16.173  -5.495 1.00109.27      A    N
ATOM    611  CA  LEU   137       7.615 -16.559  -6.589 1.00112.82      A    C
ATOM    612  CB  LEU   137       8.316 -15.325  -7.149 1.00113.33      A    C
ATOM    613  CG  LEU   137       7.332 -14.245  -7.588 1.00114.59      A    C
ATOM    614  CD1 LEU   137       8.040 -13.223  -8.463 1.00113.74      A    C
ATOM    615  CD2 LEU   137       6.183 -14.900  -8.343 1.00113.79      A    C
ATOM    616  C   LEU   137       8.647 -17.590  -6.163 1.00113.26      A    C
ATOM    617  O   LEU   137       9.098 -17.601  -5.017 1.00116.86      A    O
ATOM    618  N   PRO   138       9.021 -18.482  -7.090 1.00111.23      A    N
ATOM    619  CD  PRO   138       8.460 -18.468  -8.452 1.00111.87      A    C
ATOM    620  CA  PRO   138       9.942 -19.607  -6.891 1.00109.31      A    C
ATOM    621  CB  PRO   138       9.671 -20.501  -8.091 1.00110.49      A    C
ATOM    622  CG  PRO   138       9.241 -19.542  -9.157 1.00111.00      A    C
ATOM    623  C   PRO   138      11.411 -19.196  -6.815 1.00106.05      A    C
ATOM    624  O   PRO   138      11.811 -18.178  -7.378 1.00104.55      A    O
ATOM    625  N   HIS   139      12.203 -20.014  -6.124 1.00103.94      A    N
ATOM    626  CA  HIS   139      13.634 -19.771  -5.942 1.00101.12      A    C
ATOM    627  CB  HIS   139      14.288 -19.368  -7.275 1.00106.04      A    C
ATOM    628  CG  HIS   139      14.255 -20.439  -8.326 1.00110.34      A    C
ATOM    629  CD2 HIS   139      13.765 -20.433  -9.589 1.00112.17      A    C
ATOM    630  ND1 HIS   139      14.799 -21.690  -8.133 1.00112.32      A    N
ATOM    631  CE1 HIS   139      14.647 -22.409  -9.232 1.00113.63      A    C
ATOM    632  NE2 HIS   139      14.023 -21.669 -10.130 1.00113.61      A    N
ATOM    633  C   HIS   139      13.894 -18.687  -4.885 1.00 95.84      A    C
ATOM    634  O   HIS   139      14.899 -18.738  -4.166 1.00 95.18      A    O
ATOM    635  N   VAL   140      12.977 -17.722  -4.794 1.00 87.12      A    N
ATOM    636  CA  VAL   140      13.104 -16.590  -3.881 1.00 78.27      A    C
ATOM    637  CB  VAL   140      11.864 -15.710  -3.945 1.00 76.69      A    C
ATOM    638  CG1 VAL   140      11.989 -14.581  -2.963 1.00 76.29      A    C
ATOM    639  CG2 VAL   140      11.684 -15.182  -5.340 1.00 76.45      A    C
ATOM    640  C   VAL   140      13.314 -16.999  -2.425 1.00 73.68      A    C
ATOM    641  O   VAL   140      12.556 -17.800  -1.886 1.00 74.17      A    O
ATOM    642  N   ASP   141      14.334 -16.425  -1.790 1.00 67.84      A    N
ATOM    643  CA  ASP   141      14.716 -16.781  -0.421 1.00 61.64      A    C
ATOM    644  CB  ASP   141      16.243 -16.806  -0.307 1.00 60.92      A    C
ATOM    645  CG  ASP   141      16.726 -17.205   1.077 1.00 60.19      A    C
ATOM    646  OD1 ASP   141      15.931 -17.128   2.038 1.00 62.25      A    O
ATOM    647  OD2 ASP   141      17.908 -17.594   1.204 1.00 57.97      A    O
ATOM    648  C   ASP   141      14.141 -15.800   0.605 1.00 58.24      A    C
ATOM    649  O   ASP   141      13.682 -16.199   1.677 1.00 56.40      A    O
ATOM    650  N   TYR   142      14.192 -14.514   0.269 1.00 54.75      A    N
ATOM    651  CA  TYR   142      13.602 -13.460   1.087 1.00 50.27      A    C
ATOM    652  CB  TYR   142      14.327 -13.346   2.429 1.00 49.53      A    C
ATOM    653  CG  TYR   142      15.789 -12.950   2.337 1.00 50.08      A    C
ATOM    654  CD1 TYR   142      16.173 -11.607   2.243 1.00 49.37      A    C
ATOM    655  CE1 TYR   142      17.505 -11.247   2.179 1.00 45.77      A    C
ATOM    656  CD2 TYR   142      16.787 -13.910   2.364 1.00 48.54      A    C
ATOM    657  CE2 TYR   142      18.119 -13.552   2.300 1.00 48.20      A    C
ATOM    658  CZ  TYR   142      18.470 -12.223   2.209 1.00 47.16      A    C
ATOM    659  OH  TYR   142      19.804 -11.890   2.162 1.00 47.82      A    O
ATOM    660  C   TYR   142      13.717 -12.153   0.335 1.00 48.09      A    C
ATOM    661  O   TYR   142      14.582 -12.011  -0.520 1.00 48.03      A    O
ATOM    662  N   ILE   143      12.847 -11.201   0.650 1.00 46.96      A    N
ATOM    663  CA  ILE   143      12.880  -9.894   0.002 1.00 45.43      A    C
ATOM    664  CB  ILE   143      11.558  -9.602  -0.718 1.00 43.93      A    C
ATOM    665  CG2 ILE   143      11.539  -8.159  -1.213 1.00 42.29      A    C
ATOM    666  CG1 ILE   143      11.377 -10.588  -1.877 1.00 44.21      A    C
ATOM    667  CD1 ILE   143      10.176 -10.287  -2.765 1.00 43.87      A    C
ATOM    668  C   ILE   143      13.122  -8.783   1.010 1.00 45.82      A    C
ATOM    669  O   ILE   143      12.432  -8.701   2.024 1.00 47.06      A    O
ATOM    670  N   GLU   144      14.090  -7.913   0.740 1.00 45.93      A    N
ATOM    671  CA  GLU   144      14.306  -6.784   1.634 1.00 45.04      A    C
ATOM    672  CB  GLU   144      15.774  -6.671   2.033 1.00 44.77      A    C
ATOM    673  CG  GLU   144      15.934  -6.424   3.525 1.00 46.74      A    C
ATOM    674  CD  GLU   144      17.360  -6.169   3.944 1.00 49.18      A    C
ATOM    675  OE1 GLU   144      18.061  -7.157   4.293 1.00 49.24      A    O
ATOM    676  OE2 GLU   144      17.774  -4.981   3.927 1.00 48.88      A    O
ATOM    677  C   GLU   144      13.832  -5.456   1.075 1.00 44.39      A    C
ATOM    678  O   GLU   144      13.848  -5.222  -0.133 1.00 45.65      A    O
ATOM    679  N   GLU   145      13.399  -4.581   1.970 1.00 43.72      A    N
ATOM    680  CA  GLU   145      12.995  -3.247   1.568 1.00 42.02      A    C
ATOM    681  CB  GLU   145      11.828  -2.797   2.427 1.00 43.89      A    C
ATOM    682  CG  GLU   145      11.463  -1.345   2.287 1.00 46.61      A    C
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ATOM    683  CD  GLU   145      10.678  -0.873   3.478  1.00 47.51      A    C
ATOM    684  OE1 GLU   145      11.226  -0.933   4.601  1.00 48.29      A    O
ATOM    685  OE2 GLU   145       9.518  -0.455   3.294  1.00 49.39      A    O
ATOM    686  C   GLU   145      14.168  -2.289   1.730  1.00 40.96      A    C
ATOM    687  O   GLU   145      14.794  -2.218   2.795  1.00 38.91      A    O
ATOM    688  N   ASP   146      14.459  -1.554   0.661  1.00 40.13      A    N
ATOM    689  CA  ASP   146      15.592  -0.625   0.625  1.00 38.49      A    C
ATOM    690  CB  ASP   146      15.517   0.226  -0.651  1.00 39.89      A    C
ATOM    691  CG  ASP   146      16.890   0.626  -1.178  1.00 40.94      A    C
ATOM    692  OD1 ASP   146      17.912   0.084  -0.697  1.00 41.19      A    O
ATOM    693  OD2 ASP   146      16.936   1.486  -2.085  1.00 41.04      A    O
ATOM    694  C   ASP   146      15.607   0.287   1.853  1.00 36.42      A    C
ATOM    695  O   ASP   146      14.557   0.627   2.406  1.00 37.06      A    O
ATOM    696  N   SER   147      16.801   0.690   2.262  1.00 33.43      A    N
ATOM    697  CA  SER   147      16.970   1.466   3.480  1.00 33.76      A    C
ATOM    698  CB  SER   147      17.033   0.516   4.675  1.00 30.49      A    C
ATOM    699  OG  SER   147      17.929   1.016   5.653  1.00 34.97      A    O
ATOM    700  C   SER   147      18.232   2.367   3.436  1.00 34.29      A    C
ATOM    701  O   SER   147      19.248   2.018   2.835  1.00 32.95      A    O
ATOM    702  N   SER   148      18.165   3.525   4.084  1.00 33.24      A    N
ATOM    703  CA  SER   148      19.235   4.508   3.973  1.00 30.80      A    C
ATOM    704  CB  SER   148      18.746   5.872   4.486  1.00 30.12      A    C
ATOM    705  OG  SER   148      17.859   6.474   3.555  1.00 30.98      A    O
ATOM    706  C   SER   148      20.513   4.110   4.704  1.00 29.06      A    C
ATOM    707  O   SER   148      20.469   3.465   5.749  1.00 28.62      A    O
ATOM    708  N   VAL   149      21.653   4.493   4.141  1.00 26.45      A    N
ATOM    709  CA  VAL   149      22.893   4.467   4.882  1.00 26.25      A    C
ATOM    710  CB  VAL   149      23.937   3.507   4.236  1.00 28.16      A    C
ATOM    711  CG1 VAL   149      23.389   2.082   4.231  1.00 27.27      A    C
ATOM    712  CG2 VAL   149      24.289   3.964   2.826  1.00 26.42      A    C
ATOM    713  C   VAL   149      23.479   5.875   4.976  1.00 26.13      A    C
ATOM    714  O   VAL   149      23.208   6.743   4.132  1.00 23.35      A    O
ATOM    715  N   PHE   150      24.277   6.083   6.024  1.00 27.27      A    N
ATOM    716  CA  PHE   150      24.789   7.399   6.408  1.00 26.72      A    C
ATOM    717  CB  PHE   150      24.070   7.889   7.654  1.00 23.59      A    C
ATOM    718  CG  PHE   150      22.611   7.992   7.489  1.00 23.95      A    C
ATOM    719  CD1 PHE   150      21.804   6.893   7.709  1.00 24.72      A    C
ATOM    720  CD2 PHE   150      22.034   9.191   7.112  1.00 25.04      A    C
ATOM    721  CE1 PHE   150      20.433   6.983   7.557  1.00 26.92      A    C
ATOM    722  CE2 PHE   150      20.676   9.295   6.957  1.00 27.24      A    C
ATOM    723  CZ  PHE   150      19.867   8.182   7.181  1.00 27.37      A    C
ATOM    724  C   PHE   150      26.287   7.388   6.714  1.00 27.96      A    C
ATOM    725  O   PHE   150      26.819   6.463   7.379  1.00 26.24      A    O
ATOM    726  N   ALA   151      26.952   8.438   6.256  1.00 27.06      A    N
ATOM    727  CA  ALA   151      28.330   8.673   6.627  1.00 29.75      A    C
ATOM    728  CB  ALA   151      28.763  10.047   6.132  1.00 29.52      A    C
ATOM    729  C   ALA   151      28.460   8.592   8.149  1.00 32.40      A    C
ATOM    730  O   ALA   151      27.541   8.990   8.876  1.00 34.17      A    O
ATOM    731  N   GLN   152      29.593   8.075   8.634  1.00 33.63      A    N
ATOM    732  CA  GLN   152      29.904   8.146  10.060  1.00 34.27      A    C
ATOM    733  CB  GLN   152      30.119   6.748  10.632  1.00 32.18      A    C
ATOM    734  CG  GLN   152      28.947   5.804  10.408  1.00 32.72      A    C
ATOM    735  CD  GLN   152      27.627   6.331  10.972  1.00 28.64      A    C
ATOM    736  OE1 GLN   152      27.453   6.448  12.193  1.00 21.85      A    O
ATOM    737  NE2 GLN   152      26.690   6.651  10.075  1.00 27.12      A    N
ATOM    738  C   GLN   152      31.141   9.000  10.346  1.00 37.30      A    C
ATOM    739  O   GLN   152      31.946   8.571  11.216  1.00 38.53      A    O
ATOM    740  OXT GLN   152      31.277  10.088   9.725  1.00 38.27      A    O
TER     741      GLN   152                                              A
ATOM    742  CB  SER   153      56.497  17.251  14.876  1.00115.65      B    C
ATOM    743  OG  SER   153      57.007  16.956  13.585  1.00117.41      B    O
ATOM    744  C   SER   153      54.149  16.488  14.361  1.00110.66      B    C
ATOM    745  O   SER   153      53.706  16.469  13.214  1.00110.93      B    O
ATOM    746  N   SER   153      54.578  18.106  16.192  1.00114.12      B    N
ATOM    747  CA  SER   153      55.014  17.658  14.839  1.00113.11      B    C
ATOM    748  N   ILE   154      53.900  15.523  15.245  1.00107.72      B    N
ATOM    749  CA  ILE   154      53.158  14.315  14.877  1.00104.32      B    C
ATOM    750  CB  ILE   154      53.840  13.049  15.430  1.00107.91      B    C
ATOM    751  CG2 ILE   154      53.052  11.818  15.029  1.00109.50      B    C
ATOM    752  CG1 ILE   154      55.262  12.947  14.887  1.00111.40      B    C
ATOM    753  CD1 ILE   154      55.319  12.871  13.381  1.00116.02      B    C
ATOM    754  C   ILE   154      51.719  14.330  15.383  1.00100.10      B    C
ATOM    755  O   ILE   154      51.448  14.755  16.506  1.00 97.82      B    O
ATOM    756  N   PRO   155      50.775  13.860  14.553  1.00 96.93      B    N
ATOM    757  CD  PRO   155      50.980  13.534  13.131  1.00 95.03      B    C
ATOM    758  CA  PRO   155      49.363  13.744  14.940  1.00 91.06      B    C
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ATOM    759  CB  PRO   155      48.678  13.254  13.666  1.00 92.51      B    C
ATOM    760  CG  PRO   155      49.602  13.653  12.564  1.00 94.48      B    C
ATOM    761  C   PRO   155      49.187  12.758  16.085  1.00 85.90      B    C
ATOM    762  O   PRO   155      49.609  11.611  15.983  1.00 86.91      B    O
ATOM    763  N   TRP   156      48.553  13.199  17.166  1.00 78.41      B    N
ATOM    764  CA  TRP   156      48.441  12.380  18.366  1.00 71.33      B    C
ATOM    765  CB  TRP   156      47.418  12.984  19.335  1.00 67.02      B    C
ATOM    766  CG  TRP   156      45.987  12.685  18.982  1.00 64.20      B    C
ATOM    767  CD2 TRP   156      45.260  11.474  19.254  1.00 62.56      B    C
ATOM    768  CE2 TRP   156      43.956  11.652  18.760  1.00 61.45      B    C
ATOM    769  CE3 TRP   156      45.588  10.258  19.865  1.00 61.09      B    C
ATOM    770  CD1 TRP   156      45.113  13.516  18.351  1.00 63.97      B    C
ATOM    771  NE1 TRP   156      43.892  12.906  18.214  1.00 62.71      B    N
ATOM    772  CZ2 TRP   156      42.980  10.667  18.861  1.00 59.75      B    C
ATOM    773  CZ3 TRP   156      44.620   9.283  19.962  1.00 59.49      B    C
ATOM    774  CH2 TRP   156      43.331   9.493  19.463  1.00 60.30      B    C
ATOM    775  C   TRP   156      48.043  10.945  18.036  1.00 69.39      B    C
ATOM    776  O   TRP   156      48.377  10.011  18.766  1.00 69.00      B    O
ATOM    777  N   ASN   157      47.329  10.777  16.930  1.00 68.55      B    N
ATOM    778  CA  ASN   157      46.667   9.521  16.621  1.00 67.01      B    C
ATOM    779  CB  ASN   157      45.359   9.807  15.890  1.00 65.67      B    C
ATOM    780  CG  ASN   157      45.544  10.738  14.714  1.00 64.99      B    C
ATOM    781  OD1 ASN   157      45.730  10.295  13.578  1.00 64.19      B    O
ATOM    782  ND2 ASN   157      45.491  12.041  14.978  1.00 63.63      B    N
ATOM    783  C   ASN   157      47.526   8.577  15.786  1.00 66.89      B    C
ATOM    784  O   ASN   157      47.284   7.373  15.748  1.00 66.52      B    O
ATOM    785  N   LEU   158      48.523   9.122  15.104  1.00 67.36      B    N
ATOM    786  CA  LEU   158      49.500   8.291  14.416  1.00 69.81      B    C
ATOM    787  CB  LEU   158      50.192   9.099  13.314  1.00 67.04      B    C
ATOM    788  CG  LEU   158      49.313   9.536  12.140  1.00 64.16      B    C
ATOM    789  CD1 LEU   158      50.117  10.432  11.209  1.00 61.97      B    C
ATOM    790  CD2 LEU   158      48.795   8.316  11.395  1.00 61.15      B    C
ATOM    791  C   LEU   158      50.523   7.796  15.442  1.00 72.86      B    C
ATOM    792  O   LEU   158      51.132   6.734  15.287  1.00 73.28      B    O
ATOM    793  N   GLU   159      50.692   8.580  16.500  1.00 76.18      B    N
ATOM    794  CA  GLU   159      51.581   8.234  17.597  1.00 78.78      B    C
ATOM    795  CB  GLU   159      51.771   9.459  18.499  1.00 80.32      B    C
ATOM    796  CG  GLU   159      52.326   9.149  19.870  1.00 87.64      B    C
ATOM    797  CD  GLU   159      53.837   9.022  19.872  1.00 92.62      B    C
ATOM    798  OE1 GLU   159      54.432   9.032  20.971  1.00 96.21      B    O
ATOM    799  OE2 GLU   159      54.436   8.913  18.778  1.00 97.33      B    O
ATOM    800  C   GLU   159      51.013   7.064  18.399  1.00 78.60      B    C
ATOM    801  O   GLU   159      51.761   6.270  18.967  1.00 78.52      B    O
ATOM    802  N   ARG   160      49.688   6.957  18.428  1.00 78.98      B    N
ATOM    803  CA  ARG   160      49.012   5.959  19.253  1.00 80.10      B    C
ATOM    804  CB  ARG   160      47.604   6.426  19.609  1.00 77.90      B    C
ATOM    805  CG  ARG   160      46.908   5.513  20.602  1.00 74.99      B    C
ATOM    806  CD  ARG   160      47.510   5.691  21.973  1.00 72.60      B    C
ATOM    807  NE  ARG   160      47.759   7.107  22.225  1.00 72.12      B    N
ATOM    808  CZ  ARG   160      46.859   7.951  22.734  1.00 70.97      B    C
ATOM    809  NH1 ARG   160      47.178   9.229  22.921  1.00 69.95      B    N
ATOM    810  NH2 ARG   160      45.644   7.523  23.068  1.00 67.47      B    N
ATOM    811  C   ARG   160      48.921   4.584  18.602  1.00 82.37      B    C
ATOM    812  O   ARG   160      48.769   3.576  19.288  1.00 81.03      B    O
ATOM    813  N   ILE   161      48.996   4.537  17.280  1.00 87.22      B    N
ATOM    814  CA  ILE   161      49.033   3.255  16.598  1.00 93.06      B    C
ATOM    815  CB  ILE   161      48.143   3.260  15.341  1.00 91.27      B    C
ATOM    816  CG2 ILE   161      46.691   3.417  15.742  1.00 91.17      B    C
ATOM    817  CG1 ILE   161      48.548   4.396  14.408  1.00 90.97      B    C
ATOM    818  CD1 ILE   161      47.707   4.457  13.157  1.00 89.92      B    C
ATOM    819  C   ILE   161      50.464   2.886  16.221  1.00 98.36      B    C
ATOM    820  O   ILE   161      50.711   1.922  15.512  1.00 98.21      B    O
ATOM    821  N   THR   162      51.406   3.693  16.714  1.00104.69      B    N
ATOM    822  CA  THR   162      52.823   3.383  16.580  1.00111.43      B    C
ATOM    823  CB  THR   162      53.593   4.576  16.012  1.00112.56      B    C
ATOM    824  OG1 THR   162      53.175   4.817  14.663  1.00112.68      B    O
ATOM    825  CG2 THR   162      55.085   4.300  16.043  1.00113.86      B    C
ATOM    826  C   THR   162      53.457   2.986  17.913  1.00117.00      B    C
ATOM    827  O   THR   162      53.682   3.827  18.787  1.00115.50      B    O
ATOM    828  N   PRO   163      53.769   1.690  18.070  1.00123.18      B    N
ATOM    829  CD  PRO   163      53.698   0.725  16.959  1.00126.42      B    C
ATOM    830  CA  PRO   163      54.269   1.069  19.302  1.00128.43      B    C
ATOM    831  CB  PRO   163      54.318  -0.418  18.957  1.00129.86      B    C
ATOM    832  CG  PRO   163      54.471  -0.449  17.477  1.00129.27      B    C
ATOM    833  C   PRO   163      55.634   1.598  19.745  1.00133.09      B    C
ATOM    834  O   PRO   163      56.310   2.300  18.995  1.00133.34      B    O
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ATOM	835	N	PRO	164	56.049	1.263	20.981	1.00	136.38	B	N
ATOM	836	CD	PRO	164	55.196	0.623	21.997	1.00	137.29	B	C
ATOM	837	CA	PRO	164	57.361	1.628	21.533	1.00	138.48	B	C
ATOM	838	CB	PRO	164	57.315	1.092	22.964	1.00	138.17	B	C
ATOM	839	CG	PRO	164	55.866	1.003	23.285	1.00	137.88	B	C
ATOM	840	C	PRO	164	58.523	1.033	20.747	1.00	140.28	B	C
ATOM	841	O	PRO	164	59.685	1.303	21.046	1.00	142.21	B	O
ATOM	842	N	LEU	179	50.723	-2.852	-3.782	1.00	82.64	B	N
ATOM	843	CA	LEU	179	50.541	-4.298	-3.720	1.00	82.37	B	C
ATOM	844	CB	LEU	179	51.617	-4.927	-2.836	1.00	84.42	B	C
ATOM	845	CG	LEU	179	51.895	-6.429	-2.990	1.00	86.63	B	C
ATOM	846	CD1	LEU	179	52.687	-6.898	-1.772	1.00	87.95	B	C
ATOM	847	CD2	LEU	179	50.605	-7.234	-3.119	1.00	85.71	B	C
ATOM	848	C	LEU	179	49.167	-4.612	-3.135	1.00	80.73	B	C
ATOM	849	O	LEU	179	48.333	-5.267	-3.773	1.00	81.56	B	O
ATOM	850	N	VAL	180	48.940	-4.150	-1.908	1.00	76.81	B	N
ATOM	851	CA	VAL	180	47.650	-4.322	-1.251	1.00	72.33	B	C
ATOM	852	CB	VAL	180	47.816	-4.340	0.282	1.00	74.74	B	C
ATOM	853	CG1	VAL	180	48.717	-3.206	0.709	1.00	77.30	B	C
ATOM	854	CG2	VAL	180	46.467	-4.213	0.956	1.00	77.06	B	C
ATOM	855	C	VAL	180	46.706	-3.194	-1.653	1.00	68.18	B	C
ATOM	856	O	VAL	180	47.070	-2.013	-1.620	1.00	66.03	B	O
ATOM	857	N	GLU	181	45.494	-3.574	-2.042	1.00	64.37	B	N
ATOM	858	CA	GLU	181	44.511	-2.642	-2.592	1.00	59.68	B	C
ATOM	859	CB	GLU	181	43.767	-3.316	-3.750	1.00	62.45	B	C
ATOM	860	CG	GLU	181	43.358	-2.389	-4.874	1.00	68.37	B	C
ATOM	861	CD	GLU	181	44.526	-1.982	-5.766	1.00	72.14	B	C
ATOM	862	OE1	GLU	181	44.296	-1.238	-6.755	1.00	73.58	B	O
ATOM	863	OE2	GLU	181	45.672	-2.404	-5.475	1.00	73.22	B	O
ATOM	864	C	GLU	181	43.522	-2.258	-1.495	1.00	54.39	B	C
ATOM	865	O	GLU	181	43.151	-3.092	-0.668	1.00	52.76	B	O
ATOM	866	N	VAL	182	43.108	-0.997	-1.475	1.00	48.00	B	N
ATOM	867	CA	VAL	182	42.099	-0.558	-0.514	1.00	41.39	B	C
ATOM	868	CB	VAL	182	42.700	0.379	0.559	1.00	39.28	B	C
ATOM	869	CG1	VAL	182	41.694	0.587	1.647	1.00	36.23	B	C
ATOM	870	CG2	VAL	182	44.007	-0.198	1.122	1.00	35.21	B	C
ATOM	871	C	VAL	182	40.956	0.178	-1.212	1.00	39.09	B	C
ATOM	872	O	VAL	182	41.119	1.307	-1.673	1.00	37.28	B	O
ATOM	873	N	TYR	183	39.804	-0.477	-1.305	1.00	37.90	B	N
ATOM	874	CA	TYR	183	38.612	0.145	-1.863	1.00	36.33	B	C
ATOM	875	CB	TYR	183	37.627	-0.918	-2.321	1.00	35.00	B	C
ATOM	876	CG	TYR	183	38.035	-1.581	-3.609	1.00	36.79	B	C
ATOM	877	CD1	TYR	183	39.028	-2.548	-3.636	1.00	36.52	B	C
ATOM	878	CE1	TYR	183	39.395	-3.162	-4.818	1.00	37.26	B	C
ATOM	879	CD2	TYR	183	37.421	-1.245	-4.804	1.00	38.38	B	C
ATOM	880	CE2	TYR	183	37.782	-1.855	-5.994	1.00	39.27	B	C
ATOM	881	CZ	TYR	183	38.767	-2.813	-5.997	1.00	38.74	B	C
ATOM	882	OH	TYR	183	39.106	-3.424	-7.185	1.00	39.99	B	O
ATOM	883	C	TYR	183	37.966	1.015	-0.807	1.00	36.03	B	C
ATOM	884	O	TYR	183	37.905	0.641	0.355	1.00	37.83	B	O
ATOM	885	N	LEU	184	37.493	2.187	-1.201	1.00	35.68	B	N
ATOM	886	CA	LEU	184	36.837	3.073	-0.255	1.00	34.48	B	C
ATOM	887	CB	LEU	184	37.638	4.370	-0.100	1.00	35.48	B	C
ATOM	888	CG	LEU	184	36.889	5.451	0.685	1.00	37.28	B	C
ATOM	889	CD1	LEU	184	36.613	4.911	2.098	1.00	37.59	B	C
ATOM	890	CD2	LEU	184	37.680	6.775	0.708	1.00	35.89	B	C
ATOM	891	C	LEU	184	35.437	3.376	-0.772	1.00	33.43	B	C
ATOM	892	O	LEU	184	35.282	3.925	-1.864	1.00	31.65	B	O
ATOM	893	N	LEU	185	34.419	3.003	0.000	1.00	33.22	B	N
ATOM	894	CA	LEU	185	33.041	3.311	-0.380	1.00	33.87	B	C
ATOM	895	CB	LEU	185	32.148	2.081	-0.216	1.00	31.98	B	C
ATOM	896	CG	LEU	185	32.315	1.032	-1.314	1.00	30.98	B	C
ATOM	897	CD1	LEU	185	33.624	0.314	-1.105	1.00	30.09	B	C
ATOM	898	CD2	LEU	185	31.151	0.062	-1.291	1.00	31.06	B	C
ATOM	899	C	LEU	185	32.480	4.470	0.430	1.00	34.21	B	C
ATOM	900	O	LEU	185	32.153	4.308	1.608	1.00	33.81	B	O
ATOM	901	N	ASP	186	32.358	5.628	-0.219	1.00	34.50	B	N
ATOM	902	CA	ASP	186	32.154	6.889	0.486	1.00	34.92	B	C
ATOM	903	CB	ASP	186	33.502	7.377	1.027	1.00	39.32	B	C
ATOM	904	CG	ASP	186	33.365	8.366	2.196	1.00	45.74	B	C
ATOM	905	OD1	ASP	186	33.661	7.946	3.344	1.00	50.09	B	O
ATOM	906	OD2	ASP	186	32.990	9.553	1.985	1.00	46.44	B	O
ATOM	907	C	ASP	186	31.553	7.968	-0.428	1.00	33.07	B	C
ATOM	908	O	ASP	186	30.912	7.685	-1.451	1.00	31.20	B	O
ATOM	909	N	THR	187	31.780	9.214	-0.033	1.00	29.79	B	N
ATOM	910	CA	THR	187	31.400	10.368	-0.808	1.00	28.62	B	C

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ATOM    911  CB  THR   187      31.427  11.636   0.071  1.00 28.27      B    C
ATOM    912  OG1 THR   187      32.779  11.985   0.392  1.00 31.09      B    O
ATOM    913  CG2 THR   187      30.692  11.386   1.376  1.00 29.63      B    C
ATOM    914  C   THR   187      32.376  10.512  -1.964  1.00 28.24      B    C
ATOM    915  O   THR   187      33.222   9.657  -2.167  1.00 27.28      B    O
ATOM    916  N   SER   188      32.247  11.577  -2.734  1.00 29.91      B    N
ATOM    917  CA  SER   188      33.194  11.839  -3.786  1.00 35.55      B    C
ATOM    918  CB  SER   188      32.696  12.975  -4.687  1.00 36.16      B    C
ATOM    919  OG  SER   188      31.476  13.520  -4.223  1.00 37.67      B    O
ATOM    920  C   SER   188      34.544  12.215  -3.185  1.00 39.38      B    C
ATOM    921  O   SER   188      34.634  12.600  -2.014  1.00 39.88      B    O
ATOM    922  N   ILE   189      35.592  12.096  -3.994  1.00 42.96      B    N
ATOM    923  CA  ILE   189      36.948  12.355  -3.530  1.00 45.75      B    C
ATOM    924  CB  ILE   189      37.794  11.082  -3.594  1.00 46.16      B    C
ATOM    925  CG2 ILE   189      39.070  11.281  -2.810  1.00 45.77      B    C
ATOM    926  CG1 ILE   189      37.018   9.901  -3.020  1.00 44.55      B    C
ATOM    927  CD1 ILE   189      37.195   9.736  -1.550  1.00 46.37      B    C
ATOM    928  C   ILE   189      37.643  13.408  -4.392  1.00 48.83      B    C
ATOM    929  O   ILE   189      37.635  13.327  -5.626  1.00 48.75      B    O
ATOM    930  N   GLN   190      38.254  14.390  -3.735  1.00 52.85      B    N
ATOM    931  CA  GLN   190      39.228  15.268  -4.382  1.00 55.02      B    C
ATOM    932  CB  GLN   190      39.485  16.498  -3.516  1.00 57.93      B    C
ATOM    933  CG  GLN   190      40.325  17.544  -4.212  1.00 64.58      B    C
ATOM    934  CD  GLN   190      40.435  18.828  -3.438  1.00 67.20      B    C
ATOM    935  OE1 GLN   190      41.333  19.625  -3.678  1.00 70.49      B    O
ATOM    936  NE2 GLN   190      39.519  19.041  -2.506  1.00 70.62      B    N
ATOM    937  C   GLN   190      40.529  14.492  -4.551  1.00 54.50      B    C
ATOM    938  O   GLN   190      41.373  14.483  -3.673  1.00 54.01      B    O
ATOM    939  N   SER   191      40.694  13.827  -5.678  1.00 54.52      B    N
ATOM    940  CA  SER   191      41.750  12.835  -5.761  1.00 54.55      B    C
ATOM    941  CB  SER   191      41.455  11.862  -6.895  1.00 51.67      B    C
ATOM    942  OG  SER   191      41.141  12.506  -8.102  1.00 44.78      B    O
ATOM    943  C   SER   191      43.127  13.451  -5.936  1.00 55.93      B    C
ATOM    944  O   SER   191      44.137  12.780  -5.770  1.00 54.97      B    O
ATOM    945  N   ASP   192      43.175  14.731  -6.272  1.00 58.49      B    N
ATOM    946  CA  ASP   192      44.462  15.349  -6.511  1.00 61.41      B    C
ATOM    947  CB  ASP   192      44.475  16.033  -7.861  1.00 56.12      B    C
ATOM    948  CG  ASP   192      43.554  17.212  -7.957  1.00 51.17      B    C
ATOM    949  OD1 ASP   192      42.863  17.489  -6.959  1.00 50.15      B    O
ATOM    950  OD2 ASP   192      43.521  17.862  -9.023  1.00 44.73      B    O
ATOM    951  C   ASP   192      44.964  16.303  -5.424  1.00 66.24      B    C
ATOM    952  O   ASP   192      45.887  17.063  -5.663  1.00 65.77      B    O
ATOM    953  N   HIS   193      44.371  16.244  -4.234  1.00 73.45      B    N
ATOM    954  CA  HIS   193      44.968  16.813  -3.023  1.00 81.62      B    C
ATOM    955  CB  HIS   193      44.239  16.265  -1.793  1.00 82.94      B    C
ATOM    956  CG  HIS   193      44.367  17.120  -0.574  1.00 84.93      B    C
ATOM    957  CD2 HIS   193      43.639  18.178  -0.155  1.00 86.25      B    C
ATOM    958  ND1 HIS   193      45.324  16.906   0.392  1.00 85.89      B    N
ATOM    959  CE1 HIS   193      45.180  17.796   1.356  1.00 87.01      B    C
ATOM    960  NE2 HIS   193      44.164  18.579   1.048  1.00 87.56      B    N
ATOM    961  C   HIS   193      46.434  16.390  -2.985  1.00 86.32      B    C
ATOM    962  O   HIS   193      46.780  15.335  -3.506  1.00 89.16      B    O
ATOM    963  N   ARG   194      47.305  17.187  -2.377  1.00 90.27      B    N
ATOM    964  CA  ARG   194      48.720  16.832  -2.396  1.00 94.87      B    C
ATOM    965  CB  ARG   194      49.606  18.023  -2.003  1.00103.52      B    C
ATOM    966  CG  ARG   194      49.614  18.358  -0.519  1.00114.19      B    C
ATOM    967  CD  ARG   194      51.028  18.385   0.039  1.00122.81      B    C
ATOM    968  NE  ARG   194      51.653  19.700  -0.052  1.00129.82      B    N
ATOM    969  CZ  ARG   194      51.872  20.489   0.992  1.00133.99      B    C
ATOM    970  NH1 ARG   194      52.447  21.666   0.822  1.00136.60      B    N
ATOM    971  NH2 ARG   194      51.515  20.099   2.206  1.00136.97      B    N
ATOM    972  C   ARG   194      48.973  15.676  -1.450  1.00 92.97      B    C
ATOM    973  O   ARG   194      49.986  14.993  -1.547  1.00 93.22      B    O
ATOM    974  N   GLU   195      48.038  15.454  -0.539  1.00 90.26      B    N
ATOM    975  CA  GLU   195      48.202  14.438   0.483  1.00 87.06      B    C
ATOM    976  CB  GLU   195      47.253  14.733   1.644  1.00 87.51      B    C
ATOM    977  CG  GLU   195      47.374  13.788   2.814  1.00 89.28      B    C
ATOM    978  CD  GLU   195      48.483  14.157   3.778  1.00 88.92      B    C
ATOM    979  OE1 GLU   195      49.400  14.915   3.402  1.00 88.50      B    O
ATOM    980  OE2 GLU   195      48.434  13.679   4.927  1.00 90.60      B    O
ATOM    981  C   GLU   195      47.955  13.040  -0.074  1.00 84.15      B    C
ATOM    982  O   GLU   195      48.557  12.069   0.377  1.00 84.55      B    O
ATOM    983  N   ILE   196      47.080  12.937  -1.065  1.00 79.94      B    N
ATOM    984  CA  ILE   196      46.715  11.640  -1.609  1.00 77.04      B    C
ATOM    985  CB  ILE   196      45.258  11.283  -1.261  1.00 76.26      B    C
ATOM    986  CG2 ILE   196      45.079  11.203   0.241  1.00 73.41      B    C
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ATOM    987  CG1 ILE   196      44.319  12.332  -1.844  1.00 74.97      B    C
ATOM    988  CD1 ILE   196      42.868  11.946  -1.783  1.00 75.89      B    C
ATOM    989  C   ILE   196      46.881  11.561  -3.125  1.00 76.54      B    C
ATOM    990  O   ILE   196      46.491  10.571  -3.743  1.00 74.38      B    O
ATOM    991  N   GLU   197      47.456  12.597  -3.729  1.00 77.49      B    N
ATOM    992  CA  GLU   197      47.596  12.625  -5.181  1.00 78.27      B    C
ATOM    993  CB  GLU   197      48.293  13.911  -5.637  1.00 79.61      B    C
ATOM    994  CG  GLU   197      49.760  14.018  -5.283  1.00 85.72      B    C
ATOM    995  CD  GLU   197      50.448  15.218  -5.934  1.00 88.78      B    C
ATOM    996  OE1 GLU   197      51.690  15.174  -6.079  1.00 89.81      B    O
ATOM    997  OE2 GLU   197      49.752  16.200  -6.299  1.00 90.86      B    O
ATOM    998  C   GLU   197      48.356  11.405  -5.701  1.00 76.80      B    C
ATOM    999  O   GLU   197      49.454  11.099  -5.248  1.00 76.81      B    O
ATOM   1000  N   GLY   198      47.749  10.707  -6.653  1.00 75.14      B    N
ATOM   1001  CA  GLY   198      48.355   9.511  -7.202  1.00 72.34      B    C
ATOM   1002  C   GLY   198      47.946   8.230  -6.493  1.00 70.89      B    C
ATOM   1003  O   GLY   198      48.153   7.143  -7.013  1.00 69.29      B    O
ATOM   1004  N   ARG   199      47.352   8.357  -5.312  1.00 71.26      B    N
ATOM   1005  CA  ARG   199      47.126   7.214  -4.436  1.00 70.32      B    C
ATOM   1006  CB  ARG   199      47.618   7.544  -3.031  1.00 74.10      B    C
ATOM   1007  CG  ARG   199      49.106   7.797  -3.006  1.00 83.59      B    C
ATOM   1008  CD  ARG   199      49.602   8.217  -1.650  1.00 91.99      B    C
ATOM   1009  NE  ARG   199      51.056   8.139  -1.592  1.00100.21      B    N
ATOM   1010  CZ  ARG   199      51.760   8.332  -0.487  1.00104.98      B    C
ATOM   1011  NH1 ARG   199      53.079   8.243  -0.516  1.00108.28      B    N
ATOM   1012  NH2 ARG   199      51.138   8.617   0.646  1.00109.18      B    N
ATOM   1013  C   ARG   199      45.680   6.750  -4.378  1.00 67.21      B    C
ATOM   1014  O   ARG   199      45.382   5.730  -3.777  1.00 67.23      B    O
ATOM   1015  N   VAL   200      44.786   7.507  -5.001  1.00 63.27      B    N
ATOM   1016  CA  VAL   200      43.364   7.183  -5.016  1.00 58.38      B    C
ATOM   1017  CB  VAL   200      42.529   8.240  -4.274  1.00 57.40      B    C
ATOM   1018  CG1 VAL   200      41.049   7.967  -4.484  1.00 55.82      B    C
ATOM   1019  CG2 VAL   200      42.861   8.226  -2.799  1.00 55.70      B    C
ATOM   1020  C   VAL   200      42.859   7.126  -6.442  1.00 56.33      B    C
ATOM   1021  O   VAL   200      42.698   8.153  -7.095  1.00 56.53      B    O
ATOM   1022  N   MET   201      42.601   5.921  -6.924  1.00 55.20      B    N
ATOM   1023  CA  MET   201      42.062   5.744  -8.260  1.00 55.32      B    C
ATOM   1024  CB  MET   201      42.500   4.398  -8.830  1.00 55.79      B    C
ATOM   1025  CG  MET   201      42.155   4.221 -10.283  1.00 58.69      B    C
ATOM   1026  SD  MET   201      40.673   3.243 -10.522  1.00 64.01      B    S
ATOM   1027  CE  MET   201      40.321   3.598 -12.296  1.00 63.64      B    C
ATOM   1028  C   MET   201      40.554   5.794  -8.163  1.00 53.91      B    C
ATOM   1029  O   MET   201      39.950   5.013  -7.440  1.00 55.03      B    O
ATOM   1030  N   VAL   202      39.940   6.723  -8.875  1.00 53.21      B    N
ATOM   1031  CA  VAL   202      38.504   6.899  -8.744  1.00 52.50      B    C
ATOM   1032  CB  VAL   202      38.121   8.387  -8.828  1.00 47.24      B    C
ATOM   1033  CG1 VAL   202      36.670   8.529  -9.158  1.00 41.41      B    C
ATOM   1034  CG2 VAL   202      38.421   9.053  -7.509  1.00 41.18      B    C
ATOM   1035  C   VAL   202      37.768   6.120  -9.813  1.00 54.26      B    C
ATOM   1036  O   VAL   202      37.839   6.465 -10.992  1.00 54.98      B    O
ATOM   1037  N   THR   203      37.069   5.064  -9.394  1.00 55.90      B    N
ATOM   1038  CA  THR   203      36.356   4.210 -10.336  1.00 59.12      B    C
ATOM   1039  CB  THR   203      35.896   2.875  -9.690  1.00 57.76      B    C
ATOM   1040  OG1 THR   203      34.699   3.080  -8.924  1.00 55.18      B    O
ATOM   1041  CG2 THR   203      36.987   2.336  -8.768  1.00 55.56      B    C
ATOM   1042  C   THR   203      35.139   4.953 -10.855  1.00 61.90      B    C
ATOM   1043  O   THR   203      34.806   6.041 -10.382  1.00 63.12      B    O
ATOM   1044  N   ASP   204      34.477   4.374 -11.839  1.00 64.10      B    N
ATOM   1045  CA  ASP   204      33.344   5.038 -12.446  1.00 66.16      B    C
ATOM   1046  CB  ASP   204      33.337   4.745 -13.941  1.00 70.85      B    C
ATOM   1047  CG  ASP   204      34.107   3.476 -14.280  1.00 77.17      B    C
ATOM   1048  OD1 ASP   204      33.557   2.369 -14.052  1.00 77.27      B    O
ATOM   1049  OD2 ASP   204      35.265   3.588 -14.764  1.00 80.38      B    O
ATOM   1050  C   ASP   204      32.043   4.578 -11.797  1.00 64.17      B    C
ATOM   1051  O   ASP   204      30.981   4.625 -12.413  1.00 66.46      B    O
ATOM   1052  N   PHE   205      32.124   4.136 -10.547  1.00 60.33      B    N
ATOM   1053  CA  PHE   205      30.941   3.653  -9.848  1.00 57.09      B    C
ATOM   1054  CB  PHE   205      31.305   2.508  -8.904  1.00 58.44      B    C
ATOM   1055  CG  PHE   205      30.137   1.993  -8.137  1.00 58.36      B    C
ATOM   1056  CD1 PHE   205      29.870   2.465  -6.867  1.00 59.33      B    C
ATOM   1057  CD2 PHE   205      29.246   1.106  -8.722  1.00 60.27      B    C
ATOM   1058  CE1 PHE   205      28.727   2.071  -6.192  1.00 60.30      B    C
ATOM   1059  CE2 PHE   205      28.100   0.704  -8.054  1.00 61.35      B    C
ATOM   1060  CZ  PHE   205      27.839   1.190  -6.787  1.00 61.17      B    C
ATOM   1061  C   PHE   205      30.202   4.738  -9.055  1.00 54.53      B    C
ATOM   1062  O   PHE   205      30.717   5.279  -8.077  1.00 52.72      B    O
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ATOM   1063  N   GLU   206      28.981   5.040  -9.478  1.00 52.93      B    N
ATOM   1064  CA  GLU   206      28.152   6.010  -8.774  1.00 51.13      B    C
ATOM   1065  CB  GLU   206      28.037   7.300  -9.595  1.00 53.70      B    C
ATOM   1066  CG  GLU   206      27.201   8.385  -8.934  1.00 57.85      B    C
ATOM   1067  CD  GLU   206      27.691   9.796  -9.245  1.00 59.92      B    C
ATOM   1068  OE1 GLU   206      27.017  10.770  -8.839  1.00 61.92      B    O
ATOM   1069  OE2 GLU   206      28.748   9.936  -9.893  1.00 60.90      B    O
ATOM   1070  C   GLU   206      26.763   5.440  -8.513  1.00 47.24      B    C
ATOM   1071  O   GLU   206      25.986   5.234  -9.429  1.00 47.31      B    O
ATOM   1072  N   ASN   207      26.459   5.175  -7.254  1.00 43.16      B    N
ATOM   1073  CA  ASN   207      25.086   4.929  -6.854  1.00 37.74      B    C
ATOM   1074  CB  ASN   207      24.857   3.447  -6.593  1.00 39.88      B    C
ATOM   1075  CG  ASN   207      23.494   2.998  -7.059  1.00 41.36      B    C
ATOM   1076  OD1 ASN   207      23.376   2.105  -7.914  1.00 40.04      B    O
ATOM   1077  ND2 ASN   207      22.446   3.634  -6.523  1.00 39.17      B    N
ATOM   1078  C   ASN   207      24.736   5.714  -5.606  1.00 34.09      B    C
ATOM   1079  O   ASN   207      25.011   5.278  -4.484  1.00 32.93      B    O
ATOM   1080  N   VAL   208      24.131   6.877  -5.814  1.00 31.06      B    N
ATOM   1081  CA  VAL   208      23.780   7.766  -4.719  1.00 28.94      B    C
ATOM   1082  CB  VAL   208      24.660   9.046  -4.728  1.00 29.67      B    C
ATOM   1083  CG1 VAL   208      26.124   8.677  -4.589  1.00 30.43      B    C
ATOM   1084  CG2 VAL   208      24.436   9.825  -6.006  1.00 24.70      B    C
ATOM   1085  C   VAL   208      22.327   8.181  -4.864  1.00 29.88      B    C
ATOM   1086  O   VAL   208      21.841   8.390  -5.973  1.00 31.23      B    O
ATOM   1087  N   PRO   209      21.599   8.298  -3.747  1.00 31.45      B    N
ATOM   1088  CD  PRO   209      21.915   7.782  -2.406  1.00 31.28      B    C
ATOM   1089  CA  PRO   209      20.289   8.958  -3.785  1.00 28.45      B    C
ATOM   1090  CB  PRO   209      19.697   8.638  -2.426  1.00 28.25      B    C
ATOM   1091  CG  PRO   209      20.892   8.463  -1.549  1.00 30.42      B    C
ATOM   1092  C   PRO   209      20.387  10.467  -4.020  1.00 29.17      B    C
ATOM   1093  O   PRO   209      21.474  11.040  -4.003  1.00 27.51      B    O
ATOM   1094  N   GLU   210      19.235  11.102  -4.220  1.00 32.85      B    N
ATOM   1095  CA  GLU   210      19.148  12.539  -4.455  1.00 33.71      B    C
ATOM   1096  CB  GLU   210      17.762  12.914  -4.962  1.00 39.61      B    C
ATOM   1097  CG  GLU   210      17.373  12.334  -6.302  1.00 51.73      B    C
ATOM   1098  CD  GLU   210      16.081  12.973  -6.857  1.00 61.74      B    C
ATOM   1099  OE1 GLU   210      15.148  13.298  -6.064  1.00 64.35      B    O
ATOM   1100  OE2 GLU   210      16.003  13.157  -8.097  1.00 67.46      B    O
ATOM   1101  C   GLU   210      19.427  13.355  -3.202  1.00 31.17      B    C
ATOM   1102  O   GLU   210      19.150  12.909  -2.089  1.00 28.89      B    O
ATOM   1103  N   GLU   211      19.979  14.553  -3.398  1.00 29.36      B    N
ATOM   1104  CA  GLU   211      20.047  15.546  -2.333  1.00 30.97      B    C
ATOM   1105  CB  GLU   211      20.886  16.759  -2.766  1.00 31.25      B    C
ATOM   1106  CG  GLU   211      22.143  16.448  -3.591  1.00 33.76      B    C
ATOM   1107  CD  GLU   211      23.166  15.622  -2.829  1.00 36.82      B    C
ATOM   1108  OE1 GLU   211      23.159  15.657  -1.576  1.00 37.51      B    O
ATOM   1109  OE2 GLU   211      23.980  14.928  -3.485  1.00 39.74      B    O
ATOM   1110  C   GLU   211      18.619  16.009  -2.014  1.00 32.21      B    C
ATOM   1111  O   GLU   211      17.769  16.094  -2.900  1.00 32.92      B    O
ATOM   1112  N   ASP   212      18.356  16.296  -0.746  1.00 33.11      B    N
ATOM   1113  CA  ASP   212      17.104  16.919  -0.326  1.00 31.22      B    C
ATOM   1114  CB  ASP   212      17.006  16.843   1.207  1.00 33.18      B    C
ATOM   1115  CG  ASP   212      15.875  17.694   1.788  1.00 37.82      B    C
ATOM   1116  OD1 ASP   212      15.010  18.208   1.032  1.00 38.10      B    O
ATOM   1117  OD2 ASP   212      15.861  17.846   3.030  1.00 39.62      B    O
ATOM   1118  C   ASP   212      17.094  18.372  -0.805  1.00 29.26      B    C
ATOM   1119  O   ASP   212      17.763  19.234  -0.228  1.00 27.81      B    O
ATOM   1120  N   GLY   213      16.330  18.626  -1.863  1.00 27.67      B    N
ATOM   1121  CA  GLY   213      16.248  19.952  -2.463  1.00 26.36      B    C
ATOM   1122  C   GLY   213      15.807  21.106  -1.576  1.00 26.31      B    C
ATOM   1123  O   GLY   213      16.199  22.252  -1.805  1.00 23.54      B    O
ATOM   1124  N   THR   214      14.996  20.820  -0.561  1.00 29.38      B    N
ATOM   1125  CA  THR   214      14.552  21.854   0.380  1.00 32.08      B    C
ATOM   1126  CB  THR   214      13.732  21.268   1.535  1.00 32.59      B    C
ATOM   1127  OG1 THR   214      14.618  20.560   2.414  1.00 33.19      B    O
ATOM   1128  CG2 THR   214      12.684  20.311   1.027  1.00 32.64      B    C
ATOM   1129  C   THR   214      15.788  22.468   1.033  1.00 33.20      B    C
ATOM   1130  O   THR   214      15.720  23.556   1.613  1.00 34.45      B    O
ATOM   1131  N   ARG   215      16.908  21.750   0.965  1.00 32.69      B    N
ATOM   1132  CA  ARG   215      17.994  21.986   1.903  1.00 32.87      B    C
ATOM   1133  CB  ARG   215      17.946  20.917   2.987  1.00 31.54      B    C
ATOM   1134  CG  ARG   215      18.150  21.465   4.372  1.00 31.05      B    C
ATOM   1135  CD  ARG   215      17.327  20.692   5.370  1.00 31.52      B    C
ATOM   1136  NE  ARG   215      17.244  19.263   5.060  1.00 30.75      B    N
ATOM   1137  CZ  ARG   215      17.219  18.325   6.004  1.00 31.15      B    C
ATOM   1138  NH1 ARG   215      17.135  17.031   5.704  1.00 24.64      B    N
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ATOM   1139  NH2 ARG   215      17.290  18.708   7.274  1.00 33.67      B    N
ATOM   1140  C   ARG   215      19.395  22.043   1.285  1.00 33.56      B    C
ATOM   1141  O   ARG   215      20.372  22.345   1.974  1.00 32.49      B    O
ATOM   1142  N   PHE   216      19.488  21.768  -0.012  1.00 35.28      B    N
ATOM   1143  CA  PHE   216      20.788  21.626  -0.657  1.00 38.92      B    C
ATOM   1144  CB  PHE   216      20.867  20.293  -1.390  1.00 35.82      B    C
ATOM   1145  CG  PHE   216      22.161  20.080  -2.116  1.00 30.16      B    C
ATOM   1146  CD1 PHE   216      22.198  20.076  -3.499  1.00 25.91      B    C
ATOM   1147  CD2 PHE   216      23.337  19.866  -1.411  1.00 30.02      B    C
ATOM   1148  CE1 PHE   216      23.383  19.862  -4.179  1.00 25.89      B    C
ATOM   1149  CE2 PHE   216      24.539  19.647  -2.083  1.00 30.04      B    C
ATOM   1150  CZ  PHE   216      24.565  19.646  -3.471  1.00 26.63      B    C
ATOM   1151  C   PHE   216      21.128  22.726  -1.640  1.00 42.83      B    C
ATOM   1152  O   PHE   216      20.336  23.043  -2.528  1.00 43.81      B    O
ATOM   1153  N   HIS   217      22.326  23.284  -1.498  1.00 47.52      B    N
ATOM   1154  CA  HIS   217      22.812  24.292  -2.431  1.00 52.66      B    C
ATOM   1155  CB  HIS   217      22.869  25.642  -1.731  1.00 50.71      B    C
ATOM   1156  CG  HIS   217      21.531  26.130  -1.273  1.00 50.46      B    C
ATOM   1157  CD2 HIS   217      21.019  26.293  -0.030  1.00 50.17      B    C
ATOM   1158  ND1 HIS   217      20.531  26.491  -2.152  1.00 49.45      B    N
ATOM   1159  CE1 HIS   217      19.461  26.854  -1.467  1.00 50.58      B    C
ATOM   1160  NE2 HIS   217      19.729  26.743  -0.178  1.00 48.98      B    N
ATOM   1161  C   HIS   217      24.188  23.930  -2.979  1.00 57.43      B    C
ATOM   1162  O   HIS   217      25.180  24.121  -2.308  1.00 59.37      B    O
ATOM   1163  N   ARG   218      24.216  23.419  -4.206  1.00 64.96      B    N
ATOM   1164  CA  ARG   218      25.442  22.892  -4.802  1.00 72.47      B    C
ATOM   1165  CB  ARG   218      25.220  22.646  -6.300  1.00 78.99      B    C
ATOM   1166  CG  ARG   218      26.190  21.661  -6.954  1.00 90.87      B    C
ATOM   1167  CD  ARG   218      25.535  20.276  -7.165  1.00101.66      B    C
ATOM   1168  NE  ARG   218      26.490  19.288  -7.654  1.00114.14      B    N
ATOM   1169  CZ  ARG   218      26.150  18.068  -8.073  1.00121.60      B    C
ATOM   1170  NH1 ARG   218      24.872  17.680  -8.065  1.00127.19      B    N
ATOM   1171  NH2 ARG   218      27.081  17.221  -8.505  1.00127.03      B    N
ATOM   1172  C   ARG   218      26.608  23.862  -4.603  1.00 73.27      B    C
ATOM   1173  O   ARG   218      27.765  23.464  -4.575  1.00 74.75      B    O
ATOM   1174  N   GLN   219      26.274  25.134  -4.441  1.00 73.30      B    N
ATOM   1175  CA  GLN   219      27.223  26.237  -4.488  1.00 70.94      B    C
ATOM   1176  CB  GLN   219      26.467  27.502  -4.879  1.00 75.90      B    C
ATOM   1177  CG  GLN   219      25.114  27.586  -4.175  1.00 83.68      B    C
ATOM   1178  CD  GLN   219      24.098  28.401  -4.942  1.00 89.19      B    C
ATOM   1179  OE1 GLN   219      22.915  28.422  -4.599  1.00 94.35      B    O
ATOM   1180  NE2 GLN   219      24.554  29.083  -5.991  1.00 93.47      B    N
ATOM   1181  C   GLN   219      27.926  26.450  -3.151  1.00 66.77      B    C
ATOM   1182  O   GLN   219      29.042  26.950  -3.105  1.00 65.68      B    O
ATOM   1183  N   ALA   220      27.250  26.096  -2.065  1.00 62.72      B    N
ATOM   1184  CA  ALA   220      27.798  26.247  -0.724  1.00 58.15      B    C
ATOM   1185  CB  ALA   220      26.714  26.721   0.250  1.00 58.33      B    C
ATOM   1186  C   ALA   220      28.304  24.897  -0.295  1.00 55.57      B    C
ATOM   1187  O   ALA   220      29.020  24.768   0.694  1.00 54.79      B    O
ATOM   1188  N   SER   221      27.901  23.882  -1.046  1.00 52.69      B    N
ATOM   1189  CA  SER   221      28.148  22.512  -0.657  1.00 48.98      B    C
ATOM   1190  CB  SER   221      27.234  21.572  -1.442  1.00 48.25      B    C
ATOM   1191  OG  SER   221      27.591  20.221  -1.196  1.00 52.95      B    O
ATOM   1192  C   SER   221      29.612  22.145  -0.876  1.00 45.54      B    C
ATOM   1193  O   SER   221      30.244  22.604  -1.823  1.00 43.51      B    O
ATOM   1194  N   LYS   222      30.140  21.329   0.029  1.00 43.16      B    N
ATOM   1195  CA  LYS   222      31.494  20.805  -0.064  1.00 40.62      B    C
ATOM   1196  CB  LYS   222      32.293  21.215   1.176  1.00 42.07      B    C
ATOM   1197  CG  LYS   222      32.478  22.733   1.292  1.00 45.25      B    C
ATOM   1198  CD  LYS   222      32.723  23.204   2.724  1.00 48.85      B    C
ATOM   1199  CE  LYS   222      34.164  22.947   3.161  1.00 51.16      B    C
ATOM   1200  NZ  LYS   222      34.554  23.815   4.315  1.00 55.09      B    N
ATOM   1201  C   LYS   222      31.409  19.285  -0.183  1.00 38.00      B    C
ATOM   1202  O   LYS   222      31.643  18.554   0.777  1.00 35.80      B    O
ATOM   1203  N   CYS   223      31.073  18.833  -1.383  1.00 36.43      B    N
ATOM   1204  CA  CYS   223      30.694  17.455  -1.618  1.00 37.63      B    C
ATOM   1205  C   CYS   223      31.746  16.399  -1.325  1.00 36.41      B    C
ATOM   1206  O   CYS   223      31.424  15.223  -1.246  1.00 36.97      B    O
ATOM   1207  CB  CYS   223      30.228  17.288  -3.057  1.00 37.34      B    C
ATOM   1208  SG  CYS   223      28.629  18.083  -3.444  1.00 45.36      B    S
ATOM   1209  N   ASP   224      32.998  16.808  -1.173  1.00 35.94      B    N
ATOM   1210  CA  ASP   224      34.088  15.846  -1.114  1.00 34.10      B    C
ATOM   1211  CB  ASP   224      35.177  16.192  -2.129  1.00 33.40      B    C
ATOM   1212  CG  ASP   224      34.719  16.017  -3.560  1.00 33.62      B    C
ATOM   1213  OD1 ASP   224      33.643  15.412  -3.775  1.00 33.19      B    O
ATOM   1214  OD2 ASP   224      35.445  16.478  -4.468  1.00 33.20      B    O
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ATOM	1215	C	ASP	224	34.712	15.808	0.258	1.00	34.14	B	C
ATOM	1216	O	ASP	224	35.558	14.953	0.532	1.00	34.57	B	O
ATOM	1217	N	SER	225	34.311	16.743	1.116	1.00	31.75	B	N
ATOM	1218	CA	SER	225	34.855	16.810	2.468	1.00	31.41	B	C
ATOM	1219	CB	SER	225	33.947	17.663	3.351	1.00	30.07	B	C
ATOM	1220	OG	SER	225	33.806	17.048	4.627	1.00	34.20	B	O
ATOM	1221	C	SER	225	35.119	15.458	3.184	1.00	30.28	B	C
ATOM	1222	O	SER	225	36.241	15.195	3.603	1.00	30.27	B	O
ATOM	1223	N	HIS	226	34.091	14.629	3.338	1.00	29.35	B	N
ATOM	1224	CA	HIS	226	34.120	13.509	4.272	1.00	29.88	B	C
ATOM	1225	CB	HIS	226	32.686	13.036	4.527	1.00	30.92	B	C
ATOM	1226	CG	HIS	226	32.582	11.721	5.236	1.00	32.20	B	C
ATOM	1227	CD2	HIS	226	32.413	11.423	6.548	1.00	32.53	B	C
ATOM	1228	ND1	HIS	226	32.571	10.514	4.568	1.00	32.58	B	N
ATOM	1229	CE1	HIS	226	32.397	9.531	5.436	1.00	31.14	B	C
ATOM	1230	NE2	HIS	226	32.299	10.056	6.645	1.00	31.62	B	N
ATOM	1231	C	HIS	226	34.969	12.355	3.748	1.00	32.22	B	C
ATOM	1232	O	HIS	226	35.771	11.760	4.493	1.00	32.16	B	O
ATOM	1233	N	GLY	227	34.801	12.043	2.465	1.00	32.20	B	N
ATOM	1234	CA	GLY	227	35.568	10.969	1.869	1.00	31.70	B	C
ATOM	1235	C	GLY	227	37.021	11.318	1.607	1.00	31.69	B	C
ATOM	1236	O	GLY	227	37.911	10.485	1.821	1.00	34.79	B	O
ATOM	1237	N	THR	228	37.282	12.536	1.145	1.00	29.42	B	N
ATOM	1238	CA	THR	228	38.658	12.930	0.843	1.00	29.53	B	C
ATOM	1239	CB	THR	228	38.732	14.352	0.270	1.00	27.13	B	C
ATOM	1240	OG1	THR	228	37.899	14.454	-0.891	1.00	28.15	B	O
ATOM	1241	CG2	THR	228	40.151	14.674	-0.146	1.00	25.42	B	C
ATOM	1242	C	THR	228	39.526	12.868	2.105	1.00	29.67	B	C
ATOM	1243	O	THR	228	40.704	12.486	2.059	1.00	30.29	B	O
ATOM	1244	N	HIS	229	38.923	13.232	3.229	1.00	27.72	B	N
ATOM	1245	CA	HIS	229	39.562	13.128	4.525	1.00	27.44	B	C
ATOM	1246	CB	HIS	229	38.653	13.773	5.573	1.00	25.75	B	C
ATOM	1247	CG	HIS	229	39.267	13.858	6.936	1.00	26.14	B	C
ATOM	1248	CD2	HIS	229	39.884	14.883	7.572	1.00	23.57	B	C
ATOM	1249	ND1	HIS	229	39.261	12.801	7.821	1.00	23.19	B	N
ATOM	1250	CE1	HIS	229	39.846	13.174	8.945	1.00	25.43	B	C
ATOM	1251	NE2	HIS	229	40.232	14.432	8.820	1.00	23.55	B	N
ATOM	1252	C	HIS	229	39.841	11.661	4.895	1.00	29.45	B	C
ATOM	1253	O	HIS	229	40.891	11.345	5.477	1.00	29.48	B	O
ATOM	1254	N	LEU	230	38.914	10.765	4.553	1.00	29.61	B	N
ATOM	1255	CA	LEU	230	39.074	9.354	4.877	1.00	30.30	B	C
ATOM	1256	CB	LEU	230	37.769	8.605	4.682	1.00	31.34	B	C
ATOM	1257	CG	LEU	230	36.876	8.864	5.886	1.00	32.18	B	C
ATOM	1258	CD1	LEU	230	35.554	8.130	5.753	1.00	31.32	B	C
ATOM	1259	CD2	LEU	230	37.653	8.438	7.135	1.00	30.90	B	C
ATOM	1260	C	LEU	230	40.166	8.671	4.082	1.00	31.00	B	C
ATOM	1261	O	LEU	230	40.971	7.934	4.664	1.00	30.12	B	O
ATOM	1262	N	ALA	231	40.206	8.913	2.770	1.00	30.56	B	N
ATOM	1263	CA	ALA	231	41.387	8.543	1.975	1.00	31.98	B	C
ATOM	1264	CB	ALA	231	41.272	9.067	0.541	1.00	32.60	B	C
ATOM	1265	C	ALA	231	42.629	9.133	2.633	1.00	31.85	B	C
ATOM	1266	O	ALA	231	43.650	8.462	2.771	1.00	31.71	B	O
ATOM	1267	N	GLY	232	42.520	10.389	3.042	1.00	30.90	B	N
ATOM	1268	CA	GLY	232	43.612	11.045	3.717	1.00	33.63	B	C
ATOM	1269	C	GLY	232	44.108	10.305	4.953	1.00	35.98	B	C
ATOM	1270	O	GLY	232	45.328	10.284	5.231	1.00	35.65	B	O
ATOM	1271	N	VAL	233	43.195	9.705	5.717	1.00	35.53	B	N
ATOM	1272	CA	VAL	233	43.666	8.971	6.881	1.00	37.07	B	C
ATOM	1273	CB	VAL	233	42.533	8.704	7.908	1.00	35.57	B	C
ATOM	1274	CG1	VAL	233	43.009	7.758	8.968	1.00	32.92	B	C
ATOM	1275	CG2	VAL	233	42.113	10.001	8.582	1.00	34.02	B	C
ATOM	1276	C	VAL	233	44.294	7.647	6.440	1.00	37.64	B	C
ATOM	1277	O	VAL	233	45.170	7.120	7.102	1.00	39.28	B	O
ATOM	1278	N	VAL	234	43.875	7.103	5.313	1.00	38.29	B	N
ATOM	1279	CA	VAL	234	44.338	5.766	5.008	1.00	39.89	B	C
ATOM	1280	CB	VAL	234	43.514	5.068	3.086	1.00	36.19	B	C
ATOM	1281	CG1	VAL	234	44.048	3.637	3.661	1.00	33.10	B	C
ATOM	1282	CG2	VAL	234	42.055	5.024	4.258	1.00	34.67	B	C
ATOM	1283	C	VAL	234	45.776	5.855	4.558	1.00	43.08	B	C
ATOM	1284	O	VAL	234	46.623	5.070	4.997	1.00	43.79	B	O
ATOM	1285	N	SER	235	46.053	6.814	3.682	1.00	44.77	B	N
ATOM	1286	CA	SER	235	47.283	6.758	2.910	1.00	46.23	B	C
ATOM	1287	CB	SER	235	47.056	5.964	1.620	1.00	44.76	B	C
ATOM	1288	OG	SER	235	46.021	6.540	0.839	1.00	42.23	B	O
ATOM	1289	C	SER	235	47.784	8.152	2.590	1.00	48.21	B	C
ATOM	1290	O	SER	235	48.595	8.343	1.684	1.00	48.69	B	O

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ATOM   1291  N   GLY   236      47.304   9.130   3.345  1.00 49.94      B    N
ATOM   1292  CA  GLY   236      47.832  10.469   3.192  1.00 52.58      B    C
ATOM   1293  C   GLY   236      49.332  10.484   3.393  1.00 53.64      B    C
ATOM   1294  O   GLY   236      49.845   9.799   4.279  1.00 53.07      B    O
ATOM   1295  N   ARG   237      50.030  11.272   2.577  1.00 55.39      B    N
ATOM   1296  CA  ARG   237      51.495  11.329   2.611  1.00 56.99      B    C
ATOM   1297  CB  ARG   237      52.010  12.277   1.517  1.00 58.98      B    C
ATOM   1298  CG  ARG   237      52.163  11.617   0.149  1.00 64.90      B    C
ATOM   1299  CD  ARG   237      52.103  12.625  -0.988  1.00 70.94      B    C
ATOM   1300  NE  ARG   237      52.670  12.109  -2.229  1.00 78.50      B    N
ATOM   1301  CZ  ARG   237      51.961  11.550  -3.203  1.00 82.50      B    C
ATOM   1302  NH1 ARG   237      52.563  11.111  -4.299  1.00 86.46      B    N
ATOM   1303  NH2 ARG   237      50.653  11.425  -3.079  1.00 84.78      B    N
ATOM   1304  C   ARG   237      52.083  11.730   3.967  1.00 55.17      B    C
ATOM   1305  O   ARG   237      53.036  11.123   4.434  1.00 53.63      B    O
ATOM   1306  N   ASP   238      51.499  12.742   4.599  1.00 55.50      B    N
ATOM   1307  CA  ASP   238      52.045  13.309   5.831  1.00 53.79      B    C
ATOM   1308  CB  ASP   238      52.199  14.832   5.692  1.00 57.66      B    C
ATOM   1309  CG  ASP   238      53.374  15.241   4.779  1.00 61.12      B    C
ATOM   1310  OD1 ASP   238      54.346  14.462   4.622  1.00 60.35      B    O
ATOM   1311  OD2 ASP   238      53.323  16.362   4.219  1.00 63.48      B    O
ATOM   1312  C   ASP   238      51.199  13.008   7.063  1.00 50.51      B    C
ATOM   1313  O   ASP   238      51.606  13.315   8.178  1.00 50.36      B    O
ATOM   1314  N   ALA   239      50.022  12.425   6.857  1.00 47.11      B    N
ATOM   1315  CA  ALA   239      49.095  12.138   7.950  1.00 44.37      B    C
ATOM   1316  CB  ALA   239      48.060  13.257   8.077  1.00 38.70      B    C
ATOM   1317  C   ALA   239      48.394  10.798   7.734  1.00 44.34      B    C
ATOM   1318  O   ALA   239      47.540  10.399   8.524  1.00 44.60      B    O
ATOM   1319  N   GLY   240      48.751  10.109   6.656  1.00 44.64      B    N
ATOM   1320  CA  GLY   240      48.206   8.784   6.427  1.00 46.73      B    C
ATOM   1321  C   GLY   240      48.697   7.731   7.411  1.00 48.30      B    C
ATOM   1322  O   GLY   240      49.718   7.914   8.086  1.00 49.76      B    O
ATOM   1323  N   VAL   241      47.969   6.624   7.501  1.00 48.91      B    N
ATOM   1324  CA  VAL   241      48.395   5.502   8.323  1.00 50.59      B    C
ATOM   1325  CB  VAL   241      47.190   4.674   8.818  1.00 51.60      B    C
ATOM   1326  CG1 VAL   241      47.677   3.372   9.448  1.00 51.09      B    C
ATOM   1327  CG2 VAL   241      46.370   5.482   9.824  1.00 50.77      B    C
ATOM   1328  C   VAL   241      49.293   4.602   7.490  1.00 51.82      B    C
ATOM   1329  O   VAL   241      50.505   4.561   7.690  1.00 52.95      B    O
ATOM   1330  N   ALA   242      48.685   3.885   6.552  1.00 52.19      B    N
ATOM   1331  CA  ALA   242      49.427   3.079   5.596  1.00 53.00      B    C
ATOM   1332  CB  ALA   242      48.617   1.854   5.210  1.00 50.35      B    C
ATOM   1333  C   ALA   242      49.728   3.915   4.360  1.00 54.75      B    C
ATOM   1334  O   ALA   242      49.099   3.743   3.311  1.00 55.41      B    O
ATOM   1335  N   LYS   243      50.693   4.818   4.494  1.00 57.04      B    N
ATOM   1336  CA  LYS   243      51.026   5.750   3.427  1.00 59.49      B    C
ATOM   1337  CB  LYS   243      52.189   6.644   3.865  1.00 62.11      B    C
ATOM   1338  CG  LYS   243      52.031   7.180   5.285  1.00 64.29      B    C
ATOM   1339  CD  LYS   243      53.160   8.135   5.673  1.00 64.68      B    C
ATOM   1340  CE  LYS   243      52.959   8.684   7.088  1.00 65.19      B    C
ATOM   1341  NZ  LYS   243      54.113   9.505   7.542  1.00 65.01      B    N
ATOM   1342  C   LYS   243      51.400   4.993   2.158  1.00 60.19      B    C
ATOM   1343  O   LYS   243      52.058   3.951   2.211  1.00 60.85      B    O
ATOM   1344  N   GLY   244      50.970   5.509   1.013  1.00 59.50      B    N
ATOM   1345  CA  GLY   244      51.328   4.867  -0.238  1.00 58.54      B    C
ATOM   1346  C   GLY   244      50.309   3.847  -0.704  1.00 58.79      B    C
ATOM   1347  O   GLY   244      50.189   3.605  -1.908  1.00 59.80      B    O
ATOM   1348  N   ALA   245      49.571   3.254   0.235  1.00 56.96      B    N
ATOM   1349  CA  ALA   245      48.507   2.321  -0.116  1.00 53.85      B    C
ATOM   1350  CB  ALA   245      47.545   2.160   1.038  1.00 53.74      B    C
ATOM   1351  C   ALA   245      47.777   2.848  -1.340  1.00 51.91      B    C
ATOM   1352  O   ALA   245      47.669   4.055  -1.544  1.00 50.88      B    O
ATOM   1353  N   SER   246      47.292   1.929  -2.158  1.00 49.93      B    N
ATOM   1354  CA  SER   246      46.958   2.244  -3.534  1.00 48.39      B    C
ATOM   1355  CB  SER   246      47.694   1.276  -4.455  1.00 50.45      B    C
ATOM   1356  OG  SER   246      47.393   1.537  -5.808  1.00 55.36      B    O
ATOM   1357  C   SER   246      45.461   2.124  -3.737  1.00 46.93      B    C
ATOM   1358  O   SER   246      44.965   1.082  -4.167  1.00 46.77      B    O
ATOM   1359  N   MET   247      44.745   3.204  -3.444  1.00 46.10      B    N
ATOM   1360  CA  MET   247      43.304   3.138  -3.195  1.00 42.61      B    C
ATOM   1361  CB  MET   247      42.901   4.220  -2.210  1.00 41.03      B    C
ATOM   1362  CG  MET   247      43.382   3.946  -0.814  1.00 41.28      B    C
ATOM   1363  SD  MET   247      42.988   5.289   0.244  1.00 41.25      B    S
ATOM   1364  CE  MET   247      41.356   5.749  -0.392  1.00 44.40      B    C
ATOM   1365  C   MET   247      42.401   3.226  -4.401  1.00 40.94      B    C
ATOM   1366  O   MET   247      42.695   3.891  -5.385  1.00 40.67      B    O
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ATOM   1367  N   ARG   248      41.279   2.540  -4.305  1.00 40.61      B    N
ATOM   1368  CA  ARG   248      40.247   2.656  -5.312  1.00 42.13      B    C
ATOM   1369  CB  ARG   248      40.098   1.304  -6.012  1.00 46.07      B    C
ATOM   1370  CG  ARG   248      41.411   0.802  -6.591  1.00 52.02      B    C
ATOM   1371  CD  ARG   248      41.305  -0.609  -7.134  1.00 60.10      B    C
ATOM   1372  NE  ARG   248      41.744  -0.681  -8.524  1.00 68.77      B    N
ATOM   1373  CZ  ARG   248      40.919  -0.813  -9.560  1.00 74.48      B    C
ATOM   1374  NH1 ARG   248      41.407  -0.864 -10.795  1.00 78.20      B    N
ATOM   1375  NH2 ARG   248      39.606  -0.908  -9.364  1.00 76.74      B    N
ATOM   1376  C   ARG   248      38.950   3.083  -4.601  1.00 39.77      B    C
ATOM   1377  O   ARG   248      38.554   2.480  -3.609  1.00 40.40      B    O
ATOM   1378  N   SER   249      38.297   4.136  -5.073  1.00 35.62      B    N
ATOM   1379  CA  SER   249      37.070   4.561  -4.404  1.00 31.39      B    C
ATOM   1380  CB  SER   249      37.144   6.027  -3.983  1.00 28.56      B    C
ATOM   1381  OG  SER   249      36.827   6.873  -5.066  1.00 27.85      B    O
ATOM   1382  C   SER   249      35.844   4.362  -5.272  1.00 28.48      B    C
ATOM   1383  O   SER   249      35.906   4.501  -6.498  1.00 27.14      B    O
ATOM   1384  N   LEU   250      34.740   4.019  -4.612  1.00 25.37      B    N
ATOM   1385  CA  LEU   250      33.421   3.919  -5.234  1.00 25.79      B    C
ATOM   1386  CB  LEU   250      32.789   2.544  -4.928  1.00 26.82      B    C
ATOM   1387  CG  LEU   250      33.376   1.311  -5.646  1.00 27.29      B    C
ATOM   1388  CD1 LEU   250      34.903   1.214  -5.464  1.00 25.27      B    C
ATOM   1389  CD2 LEU   250      32.710   0.065  -5.115  1.00 28.81      B    C
ATOM   1390  C   LEU   250      32.597   5.013  -4.585  1.00 24.78      B    C
ATOM   1391  O   LEU   250      32.872   5.387  -3.439  1.00 27.07      B    O
ATOM   1392  N   ARG   251      31.599   5.539  -5.285  1.00 22.58      B    N
ATOM   1393  CA  ARG   251      30.750   6.566  -4.664  1.00 20.82      B    C
ATOM   1394  CB  ARG   251      30.669   7.787  -5.554  1.00 18.39      B    C
ATOM   1395  CG  ARG   251      30.297   9.005  -4.774  1.00 20.45      B    C
ATOM   1396  CD  ARG   251      29.335   9.883  -5.561  1.00 21.67      B    C
ATOM   1397  NE  ARG   251      28.997  11.093  -4.839  1.00 19.94      B    N
ATOM   1398  CZ  ARG   251      28.209  12.031  -5.331  1.00 22.92      B    C
ATOM   1399  NH1 ARG   251      27.697  11.877  -6.542  1.00 22.45      B    N
ATOM   1400  NH2 ARG   251      27.919  13.114  -4.610  1.00 28.30      B    N
ATOM   1401  C   ARG   251      29.329   6.124  -4.295  1.00 20.66      B    C
ATOM   1402  O   ARG   251      28.442   6.030  -5.156  1.00 16.50      B    O
ATOM   1403  N   VAL   252      29.127   5.875  -3.000  1.00 21.54      B    N
ATOM   1404  CA  VAL   252      27.852   5.381  -2.482  1.00 21.20      B    C
ATOM   1405  CB  VAL   252      27.993   4.033  -1.688  1.00 19.63      B    C
ATOM   1406  CG1 VAL   252      28.692   3.016  -2.525  1.00 17.44      B    C
ATOM   1407  CG2 VAL   252      28.721   4.248  -0.374  1.00 18.14      B    C
ATOM   1408  C   VAL   252      27.199   6.384  -1.552  1.00 21.35      B    C
ATOM   1409  O   VAL   252      26.128   6.119  -1.029  1.00 24.50      B    O
ATOM   1410  N   LEU   253      27.840   7.522  -1.333  1.00 20.92      B    N
ATOM   1411  CA  LEU   253      27.232   8.594  -0.550  1.00 20.41      B    C
ATOM   1412  CB  LEU   253      28.058   8.861   0.698  1.00 19.82      B    C
ATOM   1413  CG  LEU   253      28.172   7.724   1.711  1.00 22.59      B    C
ATOM   1414  CD1 LEU   253      28.840   8.270   2.980  1.00 21.86      B    C
ATOM   1415  CD2 LEU   253      26.769   7.150   2.020  1.00 21.15      B    C
ATOM   1416  C   LEU   253      27.099   9.896  -1.334  1.00 20.52      B    C
ATOM   1417  O   LEU   253      28.061  10.350  -1.956  1.00 19.48      B    O
ATOM   1418  N   ASN   254      25.924  10.516  -1.303  1.00 21.23      B    N
ATOM   1419  CA  ASN   254      25.798  11.814  -1.964  1.00 23.98      B    C
ATOM   1420  CB  ASN   254      24.346  12.110  -2.349  1.00 26.23      B    C
ATOM   1421  CG  ASN   254      23.416  12.164  -1.157  1.00 25.36      B    C
ATOM   1422  OD1 ASN   254      23.809  12.505  -0.039  1.00 23.00      B    O
ATOM   1423  ND2 ASN   254      22.161  11.823  -1.398  1.00 28.55      B    N
ATOM   1424  C   ASN   254      26.339  12.941  -1.102  1.00 25.22      B    C
ATOM   1425  O   ASN   254      26.926  12.700  -0.028  1.00 24.34      B    O
ATOM   1426  N   CYS   255      26.151  14.171  -1.571  1.00 25.88      B    N
ATOM   1427  CA  CYS   255      26.802  15.302  -0.916  1.00 28.48      B    C
ATOM   1428  C   CYS   255      26.204  15.465   0.466  1.00 29.01      B    C
ATOM   1429  O   CYS   255      26.793  16.101   1.330  1.00 30.18      B    O
ATOM   1430  CB  CYS   255      26.629  16.588  -1.724  1.00 28.76      B    C
ATOM   1431  SG  CYS   255      27.276  16.517  -3.442  1.00 36.42      B    S
ATOM   1432  N   GLN   256      25.033  14.879   0.685  1.00 28.37      B    N
ATOM   1433  CA  GLN   256      24.396  15.027   1.973  1.00 28.98      B    C
ATOM   1434  CB  GLN   256      22.874  15.122   1.820  1.00 28.59      B    C
ATOM   1435  CG  GLN   256      22.359  16.549   2.025  1.00 27.47      B    C
ATOM   1436  CD  GLN   256      21.063  16.820   1.300  1.00 25.99      B    C
ATOM   1437  OE1 GLN   256      20.388  15.895   0.843  1.00 27.71      B    O
ATOM   1438  NE2 GLN   256      20.710  18.088   1.182  1.00 21.77      B    N
ATOM   1439  C   GLN   256      24.767  13.893   2.901  1.00 28.76      B    C
ATOM   1440  O   GLN   256      24.233  13.796   4.005  1.00 28.16      B    O
ATOM   1441  N   GLY   257      25.690  13.045   2.447  1.00 29.54      B    N
ATOM   1442  CA  GLY   257      26.200  11.964   3.280  1.00 28.50      B    C
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ATOM   1443  C   GLY   257      25.263  10.774   3.347  1.00 29.00      B    C
ATOM   1444  O   GLY   257      25.513   9.795   4.060  1.00 27.09      B    O
ATOM   1445  N   LYS   258      24.169  10.872   2.600  1.00 30.82      B    N
ATOM   1446  CA  LYS   258      23.143   9.844   2.595  1.00 31.63      B    C
ATOM   1447  CB  LYS   258      21.757  10.497   2.455  1.00 32.83      B    C
ATOM   1448  CG  LYS   258      20.572   9.539   2.594  1.00 35.03      B    C
ATOM   1449  CD  LYS   258      19.575  10.016   3.632  1.00 33.55      B    C
ATOM   1450  CE  LYS   258      18.826  11.265   3.169  1.00 38.29      B    C
ATOM   1451  NZ  LYS   258      17.669  10.963   2.256  1.00 33.52      B    N
ATOM   1452  C   LYS   258      23.420   8.904   1.433  1.00 30.72      B    C
ATOM   1453  O   LYS   258      23.941   9.323   0.390  1.00 31.26      B    O
ATOM   1454  N   GLY   259      23.089   7.633   1.627  1.00 28.86      B    N
ATOM   1455  CA  GLY   259      23.292   6.635   0.586  1.00 28.66      B    C
ATOM   1456  C   GLY   259      22.214   5.590   0.739  1.00 27.16      B    C
ATOM   1457  O   GLY   259      21.327   5.774   1.556  1.00 29.64      B    O
ATOM   1458  N   THR   260      22.263   4.499  -0.010  1.00 25.94      B    N
ATOM   1459  CA  THR   260      21.243   3.470   0.174  1.00 24.06      B    C
ATOM   1460  CB  THR   260      20.185   3.497  -0.956  1.00 23.07      B    C
ATOM   1461  OG1 THR   260      20.716   2.860  -2.120  1.00 23.27      B    O
ATOM   1462  CG2 THR   260      19.803   4.929  -1.294  1.00 23.82      B    C
ATOM   1463  C   THR   260      21.803   2.059   0.254  1.00 23.68      B    C
ATOM   1464  O   THR   260      22.853   1.774  -0.299  1.00 23.66      B    O
ATOM   1465  N   VAL   261      21.083   1.174   0.938  1.00 24.37      B    N
ATOM   1466  CA  VAL   261      21.509  -0.201   1.075  1.00 24.48      B    C
ATOM   1467  CB  VAL   261      20.511  -1.049   1.868  1.00 24.12      B    C
ATOM   1468  CG1 VAL   261      20.950  -2.486   1.814  1.00 22.98      B    C
ATOM   1469  CG2 VAL   261      20.445  -0.589   3.307  1.00 26.24      B    C
ATOM   1470  C   VAL   261      21.664  -0.856  -0.276  1.00 25.55      B    C
ATOM   1471  O   VAL   261      22.598  -1.631  -0.476  1.00 25.67      B    O
ATOM   1472  N   SER   262      20.754  -0.560  -1.202  1.00 25.26      B    N
ATOM   1473  CA  SER   262      20.823  -1.198  -2.510  1.00 28.13      B    C
ATOM   1474  CB  SER   262      19.559  -0.928  -3.301  1.00 26.07      B    C
ATOM   1475  OG  SER   262      19.262   0.447  -3.272  1.00 30.83      B    O
ATOM   1476  C   SER   262      22.038  -0.704  -3.293  1.00 30.74      B    C
ATOM   1477  O   SER   262      22.734  -1.485  -3.974  1.00 30.74      B    O
ATOM   1478  N   GLY   263      22.296   0.594  -3.172  1.00 32.08      B    N
ATOM   1479  CA  GLY   263      23.413   1.196  -3.876  1.00 34.48      B    C
ATOM   1480  C   GLY   263      24.735   0.711  -3.328  1.00 35.11      B    C
ATOM   1481  O   GLY   263      25.684   0.517  -4.075  1.00 36.07      B    O
ATOM   1482  N   THR   264      24.801   0.514  -2.019  1.00 34.36      B    N
ATOM   1483  CA  THR   264      26.023   0.040  -1.409  1.00 34.13      B    C
ATOM   1484  CB  THR   264      25.886  -0.023   0.126  1.00 32.04      B    C
ATOM   1485  OG1 THR   264      25.904   1.301   0.657  1.00 29.99      B    O
ATOM   1486  CG2 THR   264      27.019  -0.816   0.741  1.00 30.83      B    C
ATOM   1487  C   THR   264      26.216  -1.361  -1.942  1.00 36.25      B    C
ATOM   1488  O   THR   264      27.305  -1.749  -2.384  1.00 36.88      B    O
ATOM   1489  N   LEU   265      25.122  -2.112  -1.900  1.00 36.49      B    N
ATOM   1490  CA  LEU   265      25.119  -3.489  -2.338  1.00 36.94      B    C
ATOM   1491  CB  LEU   265      23.682  -3.980  -2.408  1.00 38.26      B    C
ATOM   1492  CG  LEU   265      23.306  -5.152  -1.522  1.00 36.02      B    C
ATOM   1493  CD1 LEU   265      21.836  -5.460  -1.766  1.00 37.13      B    C
ATOM   1494  CD2 LEU   265      24.178  -6.351  -1.857  1.00 36.13      B    C
ATOM   1495  C   LEU   265      25.773  -3.583  -3.711  1.00 36.28      B    C
ATOM   1496  O   LEU   265      26.781  -4.256  -3.885  1.00 36.69      B    O
ATOM   1497  N   ILE   266      25.196  -2.891  -4.683  1.00 36.29      B    N
ATOM   1498  CA  ILE   266      25.706  -2.929  -6.044  1.00 35.45      B    C
ATOM   1499  CB  ILE   266      24.879  -1.982  -6.950  1.00 31.67      B    C
ATOM   1500  CG2 ILE   266      25.385  -2.047  -8.363  1.00 29.36      B    C
ATOM   1501  CG1 ILE   266      23.390  -2.379  -6.899  1.00 31.50      B    C
ATOM   1502  CD1 ILE   266      22.469  -1.530  -7.756  1.00 25.14      B    C
ATOM   1503  C   ILE   266      27.197  -2.546  -6.067  1.00 37.49      B    C
ATOM   1504  O   ILE   266      27.964  -3.033  -6.904  1.00 37.39      B    O
ATOM   1505  N   GLY   267      27.612  -1.694  -5.134  1.00 37.97      B    N
ATOM   1506  CA  GLY   267      29.013  -1.332  -5.061  1.00 38.87      B    C
ATOM   1507  C   GLY   267      29.894  -2.504  -4.644  1.00 41.23      B    C
ATOM   1508  O   GLY   267      31.043  -2.618  -5.086  1.00 42.31      B    O
ATOM   1509  N   LEU   268      29.367  -3.375  -3.787  1.00 41.65      B    N
ATOM   1510  CA  LEU   268      30.122  -4.524  -3.298  1.00 42.08      B    C
ATOM   1511  CB  LEU   268      29.424  -5.170  -2.095  1.00 42.22      B    C
ATOM   1512  CG  LEU   268      29.354  -4.290  -0.844  1.00 43.99      B    C
ATOM   1513  CD1 LEU   268      28.426  -4.921   0.173  1.00 43.26      B    C
ATOM   1514  CD2 LEU   268      30.756  -4.085  -0.276  1.00 42.42      B    C
ATOM   1515  C   LEU   268      30.204  -5.529  -4.416  1.00 43.00      B    C
ATOM   1516  O   LEU   268      31.120  -6.352  -4.451  1.00 41.06      B    O
ATOM   1517  N   GLU   269      29.238  -5.460  -5.331  1.00 43.87      B    N
ATOM   1518  CA  GLU   269      29.213  -6.378  -6.459  1.00 46.33      B    C
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ATOM   1519  CB  GLU   269      27.818  -6.445  -7.064  1.00 43.85      B    C
ATOM   1520  CG  GLU   269      27.622  -7.621  -7.999  1.00 42.78      B    C
ATOM   1521  CD  GLU   269      26.619  -7.328  -9.103  1.00 43.42      B    C
ATOM   1522  OE1 GLU   269      25.667  -6.554  -8.869  1.00 42.07      B    O
ATOM   1523  OE2 GLU   269      26.788  -7.873 -10.214  1.00 45.39      B    O
ATOM   1524  C   GLU   269      30.220  -5.929  -7.517  1.00 48.39      B    C
ATOM   1525  O   GLU   269      30.919  -6.754  -8.110  1.00 49.72      B    O
ATOM   1526  N   PHE   270      30.297  -4.619  -7.739  1.00 49.23      B    N
ATOM   1527  CA  PHE   270      31.364  -4.030  -8.538  1.00 48.36      B    C
ATOM   1528  CB  PHE   270      31.211  -2.505  -8.522  1.00 47.34      B    C
ATOM   1529  CG  PHE   270      32.293  -1.761  -9.257  1.00 48.28      B    C
ATOM   1530  CD1 PHE   270      32.085  -1.306 -10.552  1.00 48.57      B    C
ATOM   1531  CD2 PHE   270      33.504  -1.475  -8.641  1.00 46.97      B    C
ATOM   1532  CE1 PHE   270      33.059  -0.581 -11.217  1.00 46.26      B    C
ATOM   1533  CE2 PHE   270      34.479  -0.752  -9.302  1.00 47.34      B    C
ATOM   1534  CZ  PHE   270      34.254  -0.306 -10.591  1.00 47.47      B    C
ATOM   1535  C   PHE   270      32.730  -4.454  -7.976  1.00 48.75      B    C
ATOM   1536  O   PHE   270      33.635  -4.802  -8.727  1.00 48.85      B    O
ATOM   1537  N   ILE   271      32.882  -4.442  -6.658  1.00 49.09      B    N
ATOM   1538  CA  ILE   271      34.162  -4.801  -6.071  1.00 51.75      B    C
ATOM   1539  CB  ILE   271      34.193  -4.501  -4.566  1.00 52.01      B    C
ATOM   1540  CG2 ILE   271      35.275  -5.324  -3.886  1.00 49.86      B    C
ATOM   1541  CG1 ILE   271      34.422  -3.005  -4.346  1.00 52.71      B    C
ATOM   1542  CD1 ILE   271      34.765  -2.638  -2.916  1.00 54.18      B    C
ATOM   1543  C   ILE   271      34.484  -6.272  -6.291  1.00 54.91      B    C
ATOM   1544  O   ILE   271      35.629  -6.627  -6.577  1.00 54.18      B    O
ATOM   1545  N   ARG   272      33.467  -7.121  -6.164  1.00 58.06      B    N
ATOM   1546  CA  ARG   272      33.610  -8.557  -6.376  1.00 60.91      B    C
ATOM   1547  CB  ARG   272      32.333  -9.274  -5.934  1.00 62.27      B    C
ATOM   1548  CG  ARG   272      32.265 -10.749  -6.317  1.00 62.63      B    C
ATOM   1549  CD  ARG   272      33.258 -11.598  -5.531  1.00 62.94      B    C
ATOM   1550  NE  ARG   272      33.136 -13.000  -5.914  1.00 66.12      B    N
ATOM   1551  CZ  ARG   272      34.131 -13.878  -5.876  1.00 67.67      B    C
ATOM   1552  NH1 ARG   272      33.923 -15.132  -6.252  1.00 67.95      B    N
ATOM   1553  NH2 ARG   272      35.329 -13.503  -5.456  1.00 69.21      B    N
ATOM   1554  C   ARG   272      33.870  -8.846  -7.847  1.00 62.21      B    C
ATOM   1555  O   ARG   272      34.764  -9.616  -8.196  1.00 63.03      B    O
ATOM   1556  N   LYS   273      33.074  -8.221  -8.701  1.00 62.61      B    N
ATOM   1557  CA  LYS   273      33.251  -8.323 -10.132  1.00 64.08      B    C
ATOM   1558  CB  LYS   273      32.341  -7.313 -10.825  1.00 64.38      B    C
ATOM   1559  CG  LYS   273      32.239  -7.493 -12.320  1.00 67.49      B    C
ATOM   1560  CD  LYS   273      31.427  -8.739 -12.660  1.00 73.19      B    C
ATOM   1561  CE  LYS   273      29.976  -8.594 -12.216  1.00 75.81      B    C
ATOM   1562  NZ  LYS   273      29.238  -7.554 -13.000  1.00 81.17      B    N
ATOM   1563  C   LYS   273      34.708  -8.048 -10.482  1.00 65.41      B    C
ATOM   1564  O   LYS   273      35.328  -8.792 -11.229  1.00 68.02      B    O
ATOM   1565  N   SER   274      35.253  -6.974  -9.925  1.00 68.14      B    N
ATOM   1566  CA  SER   274      36.656  -6.628 -10.118  1.00 69.59      B    C
ATOM   1567  CB  SER   274      37.015  -5.384  -9.310  1.00 67.55      B    C
ATOM   1568  OG  SER   274      36.241  -4.284  -9.727  1.00 67.09      B    O
ATOM   1569  C   SER   274      37.551  -7.764  -9.670  1.00 71.55      B    C
ATOM   1570  O   SER   274      38.246  -8.373 -10.475  1.00 72.19      B    O
ATOM   1571  N   GLN   275      37.525  -8.040  -8.372  1.00 74.22      B    N
ATOM   1572  CA  GLN   275      38.417  -9.023  -7.777  1.00 77.85      B    C
ATOM   1573  CB  GLN   275      37.936  -9.379  -6.369  1.00 76.85      B    C
ATOM   1574  CG  GLN   275      38.908 -10.221  -5.574  1.00 76.18      B    C
ATOM   1575  CD  GLN   275      38.196 -11.209  -4.675  1.00 76.76      B    C
ATOM   1576  OE1 GLN   275      37.049 -11.572  -4.928  1.00 77.71      B    O
ATOM   1577  NE2 GLN   275      38.870 -11.648  -3.619  1.00 76.78      B    N
ATOM   1578  C   GLN   275      38.524 -10.289  -8.629  1.00 80.71      B    C
ATOM   1579  O   GLN   275      39.545 -10.967  -8.606  1.00 80.41      B    O
ATOM   1580  N   LEU   276      37.481 -10.602  -9.388  1.00 84.76      B    N
ATOM   1581  CA  LEU   276      37.564 -11.713 -10.324  1.00 89.24      B    C
ATOM   1582  CB  LEU   276      36.204 -11.978 -10.971  1.00 87.78      B    C
ATOM   1583  CG  LEU   276      35.112 -12.504 -10.039  1.00 87.27      B    C
ATOM   1584  CD1 LEU   276      33.844 -12.767 -10.833  1.00 85.86      B    C
ATOM   1585  CD2 LEU   276      35.591 -13.768  -9.351  1.00 85.57      B    C
ATOM   1586  C   LEU   276      38.598 -11.420 -11.405  1.00 92.06      B    C
ATOM   1587  O   LEU   276      39.735 -11.884 -11.324  1.00 94.30      B    O
ATOM   1588  N   VAL   277      38.206 -10.638 -12.406  1.00 94.47      B    N
ATOM   1589  CA  VAL   277      39.058 -10.389 -13.567  1.00 96.12      B    C
ATOM   1590  CB  VAL   277      38.306  -9.555 -14.652  1.00 95.15      B    C
ATOM   1591  CG1 VAL   277      37.071 -10.292 -15.118  1.00 93.14      B    C
ATOM   1592  CG2 VAL   277      37.928  -8.196 -14.109  1.00 93.32      B    C
ATOM   1593  C   VAL   277      40.378  -9.686 -13.222  1.00 97.93      B    C
ATOM   1594  O   VAL   277      41.159  -9.353 -14.107  1.00 97.39      B    O
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ATOM	1595	N	GLN	278	40.630	-9.467	-11.939	1.00101.32	B	N
ATOM	1596	CA	GLN	278	41.830	-8.754	-11.522	1.00105.97	B	C
ATOM	1597	CB	GLN	278	41.490	-7.782	-10.392	1.00105.35	B	C
ATOM	1598	CG	GLN	278	41.943	-6.352	-10.628	1.00103.69	B	C
ATOM	1599	CD	GLN	278	41.433	-5.404	-9.557	1.00102.60	B	C
ATOM	1600	OE1	GLN	278	41.461	-4.182	-9.726	1.00101.28	B	O
ATOM	1601	NE2	GLN	278	40.959	-5.965	-8.447	1.00100.13	B	N
ATOM	1602	C	GLN	278	42.909	-9.723	-11.057	1.00108.32	B	C
ATOM	1603	O	GLN	278	42.614	-10.809	-10.563	1.00109.28	B	O
ATOM	1604	N	PRO	279	44.181	-9.335	-11.206	1.00109.09	B	N
ATOM	1605	CD	PRO	279	44.637	-8.017	-11.676	1.00107.47	B	C
ATOM	1606	CA	PRO	279	45.300	-10.176	-10.768	1.00110.86	B	C
ATOM	1607	CB	PRO	279	46.530	-9.402	-11.226	1.00107.87	B	C
ATOM	1608	CG	PRO	279	46.078	-7.984	-11.247	1.00106.50	B	C
ATOM	1609	C	PRO	279	45.271	-10.355	-9.257	1.00111.43	B	C
ATOM	1610	O	PRO	279	44.978	-9.413	-8.520	1.00113.76	B	O
ATOM	1611	N	VAL	280	45.588	-11.564	-8.804	1.00110.51	B	N
ATOM	1612	CA	VAL	280	45.386	-11.947	-7.411	1.00109.51	B	C
ATOM	1613	CB	VAL	280	45.741	-13.435	-7.179	1.00111.61	B	C
ATOM	1614	CG1	VAL	280	44.843	-14.319	-8.025	1.00112.81	B	C
ATOM	1615	CG2	VAL	280	47.198	-13.688	-7.524	1.00113.25	B	C
ATOM	1616	C	VAL	280	46.188	-11.099	-6.434	1.00106.06	B	C
ATOM	1617	O	VAL	280	47.391	-10.925	-6.583	1.00106.05	B	O
ATOM	1618	N	GLY	281	45.502	-10.572	-5.429	1.00102.30	B	N
ATOM	1619	CA	GLY	281	46.161	-9.771	-4.420	1.00 96.08	B	C
ATOM	1620	C	GLY	281	45.214	-9.508	-3.272	1.00 90.39	B	C
ATOM	1621	O	GLY	281	44.008	-9.683	-3.415	1.00 91.92	B	O
ATOM	1622	N	PRO	282	45.731	-9.088	-2.114	1.00 85.97	B	N
ATOM	1623	CD	PRO	282	47.160	-8.858	-1.846	1.00 87.84	B	C
ATOM	1624	CA	PRO	282	44.896	-8.825	-0.937	1.00 80.46	B	C
ATOM	1625	CB	PRO	282	45.908	-8.715	0.199	1.00 84.08	B	C
ATOM	1626	CG	PRO	282	47.160	-8.263	-0.471	1.00 87.49	B	C
ATOM	1627	C	PRO	282	44.049	-7.556	-1.006	1.00 74.39	B	C
ATOM	1628	O	PRO	282	44.553	-6.508	-1.516	1.00 72.54	B	O
ATOM	1629	N	LEU	283	42.767	-7.655	-0.730	1.00 68.07	B	N
ATOM	1630	CA	LEU	283	41.861	-6.506	-0.783	1.00 61.93	B	C
ATOM	1631	CB	LEU	283	40.682	-6.782	-1.726	1.00 60.52	B	C
ATOM	1632	CG	LEU	283	41.009	-7.186	-3.163	1.00 60.95	B	C
ATOM	1633	CD1	LEU	283	39.834	-6.834	-4.064	1.00 60.01	B	C
ATOM	1634	CD2	LEU	283	42.266	-6.475	-3.631	1.00 62.63	B	C
ATOM	1635	C	LEU	283	41.315	-6.118	0.585	1.00 57.20	B	C
ATOM	1636	O	LEU	283	40.773	-6.943	1.316	1.00 54.88	B	O
ATOM	1637	N	VAL	284	41.472	-4.844	0.919	1.00 52.54	B	N
ATOM	1638	CA	VAL	284	40.759	-4.245	2.037	1.00 47.36	B	C
ATOM	1639	CB	VAL	284	41.674	-3.304	2.846	1.00 45.36	B	C
ATOM	1640	CG1	VAL	284	40.929	-2.737	4.029	1.00 45.93	B	C
ATOM	1641	CG2	VAL	284	42.887	-4.050	3.322	1.00 42.54	B	C
ATOM	1642	C	VAL	284	39.591	-3.440	1.473	1.00 44.41	B	C
ATOM	1643	O	VAL	284	39.694	-2.814	0.419	1.00 44.27	B	O
ATOM	1644	N	VAL	285	38.465	-3.472	2.166	1.00 41.51	B	N
ATOM	1645	CA	VAL	285	37.341	-2.636	1.769	1.00 38.92	B	C
ATOM	1646	CB	VAL	285	36.171	-3.503	1.299	1.00 37.13	B	C
ATOM	1647	CG1	VAL	285	35.051	-2.645	0.775	1.00 35.52	B	C
ATOM	1648	CG2	VAL	285	36.660	-4.442	0.240	1.00 36.78	B	C
ATOM	1649	C	VAL	285	36.935	-1.794	2.975	1.00 35.14	B	C
ATOM	1650	O	VAL	285	36.583	-2.329	4.033	1.00 36.12	B	O
ATOM	1651	N	LEU	286	37.024	-0.480	2.835	1.00 29.16	B	N
ATOM	1652	CA	LEU	286	36.689	0.388	3.943	1.00 27.92	B	C
ATOM	1653	CB	LEU	286	37.665	1.552	4.026	1.00 24.63	B	C
ATOM	1654	CG	LEU	286	37.299	2.689	4.975	1.00 23.37	B	C
ATOM	1655	CD1	LEU	286	37.193	2.208	6.431	1.00 21.39	B	C
ATOM	1656	CD2	LEU	286	38.360	3.758	4.829	1.00 23.59	B	C
ATOM	1657	C	LEU	286	35.278	0.908	3.742	1.00 28.41	B	C
ATOM	1658	O	LEU	286	34.984	1.517	2.714	1.00 29.12	B	O
ATOM	1659	N	LEU	287	34.414	0.663	4.726	1.00 28.37	B	N
ATOM	1660	CA	LEU	287	32.998	1.003	4.635	1.00 28.51	B	C
ATOM	1661	CB	LEU	287	32.175	-0.275	4.784	1.00 32.94	B	C
ATOM	1662	CG	LEU	287	32.022	-1.197	3.572	1.00 36.53	B	C
ATOM	1663	CD1	LEU	287	31.746	-2.615	4.052	1.00 37.57	B	C
ATOM	1664	CD2	LEU	287	30.878	-0.692	2.667	1.00 37.50	B	C
ATOM	1665	C	LEU	287	32.560	2.022	5.701	1.00 26.32	B	C
ATOM	1666	O	LEU	287	31.825	1.682	6.636	1.00 25.55	B	O
ATOM	1667	N	PRO	288	32.989	3.285	5.564	1.00 25.64	B	N
ATOM	1668	CD	PRO	288	33.477	3.896	4.314	1.00 26.22	B	C
ATOM	1669	CA	PRO	288	32.781	4.299	6.603	1.00 25.01	B	C
ATOM	1670	CB	PRO	288	33.736	5.411	6.196	1.00 25.67	B	C

ATOM	1671	CG	PRO	288	33.688	5.361	4.683	1.00	24.95	B	C
ATOM	1672	C	PRO	288	31.336	4.799	6.659	1.00	24.78	B	C
ATOM	1673	O	PRO	288	31.099	6.009	6.677	1.00	24.06	B	O
ATOM	1674	N	LEU	289	30.391	3.871	6.682	1.00	24.21	B	N
ATOM	1675	CA	LEU	289	29.961	4.213	6.708	1.00	24.93	B	C
ATOM	1676	CB	LEU	289	28.414	4.234	5.277	1.00	21.80	B	C
ATOM	1677	CG	LEU	289	28.726	2.999	4.424	1.00	21.73	B	C
ATOM	1678	CD1	LEU	289	27.882	1.824	4.854	1.00	22.83	B	C
ATOM	1679	CD2	LEU	289	28.461	3.318	2.970	1.00	21.63	B	C
ATOM	1680	C	LEU	289	28.165	3.210	7.557	1.00	25.83	B	C
ATOM	1681	O	LEU	289	28.615	2.087	7.787	1.00	26.10	B	O
ATOM	1682	N	ALA	290	26.987	3.613	8.022	1.00	26.31	B	N
ATOM	1683	CA	ALA	290	26.088	2.683	8.709	1.00	25.80	B	C
ATOM	1684	CB	ALA	290	26.142	2.912	10.220	1.00	22.49	B	C
ATOM	1685	C	ALA	290	24.658	2.860	8.220	1.00	27.47	B	C
ATOM	1686	O	ALA	290	24.266	3.938	7.756	1.00	26.91	B	O
ATOM	1687	N	GLY	291	23.881	1.789	8.327	1.00	29.54	B	N
ATOM	1688	CA	GLY	291	22.435	1.910	8.311	1.00	30.47	B	C
ATOM	1689	C	GLY	291	21.837	1.040	9.397	1.00	31.26	B	C
ATOM	1690	O	GLY	291	22.557	0.577	10.275	1.00	31.62	B	O
ATOM	1691	N	GLY	292	20.530	0.801	9.340	1.00	33.20	B	N
ATOM	1692	CA	GLY	292	19.939	-0.177	10.239	1.00	31.71	B	C
ATOM	1693	C	GLY	292	20.446	-1.550	9.851	1.00	32.14	B	C
ATOM	1694	O	GLY	292	20.856	-1.719	8.707	1.00	31.02	B	O
ATOM	1695	N	TYR	293	20.437	-2.508	10.784	1.00	33.24	B	N
ATOM	1696	CA	TYR	293	20.782	-3.900	10.476	1.00	33.89	B	C
ATOM	1697	CB	TYR	293	20.391	-4.856	11.627	1.00	36.33	B	C
ATOM	1698	CG	TYR	293	20.208	-6.306	11.192	1.00	37.79	B	C
ATOM	1699	CD1	TYR	293	21.258	-7.219	11.258	1.00	38.75	B	C
ATOM	1700	CE1	TYR	293	21.117	-8.501	10.764	1.00	38.29	B	C
ATOM	1701	CD2	TYR	293	19.008	-6.730	10.631	1.00	38.47	B	C
ATOM	1702	CE2	TYR	293	18.853	-7.998	10.139	1.00	37.87	B	C
ATOM	1703	CZ	TYR	293	19.905	-8.884	10.200	1.00	40.01	B	C
ATOM	1704	OH	TYR	293	19.731	-10.145	9.667	1.00	40.12	B	O
ATOM	1705	C	TYR	293	20.064	-4.318	9.208	1.00	34.23	B	C
ATOM	1706	O	TYR	293	18.855	-4.168	9.090	1.00	32.68	B	O
ATOM	1707	N	SER	294	20.831	-4.825	8.252	1.00	36.49	B	N
ATOM	1708	CA	SER	294	20.296	-5.257	6.971	1.00	37.92	B	C
ATOM	1709	CB	SER	294	20.802	-4.365	5.842	1.00	37.02	B	C
ATOM	1710	OG	SER	294	20.222	-4.748	4.601	1.00	36.27	B	O
ATOM	1711	C	SER	294	20.785	-6.666	6.730	1.00	40.68	B	C
ATOM	1712	O	SER	294	21.995	-6.919	6.637	1.00	39.86	B	O
ATOM	1713	N	ARG	295	19.848	-7.593	6.621	1.00	43.03	B	N
ATOM	1714	CA	ARG	295	20.250	-8.969	6.436	1.00	44.95	B	C
ATOM	1715	CB	ARG	295	19.035	-9.900	6.559	1.00	46.57	B	C
ATOM	1716	CG	ARG	295	18.604	-10.555	5.276	1.00	49.25	B	C
ATOM	1717	CD	ARG	295	18.893	-12.030	5.331	1.00	51.43	B	C
ATOM	1718	NE	ARG	295	17.661	-12.797	5.434	1.00	54.06	B	N
ATOM	1719	CZ	ARG	295	17.620	-14.104	5.653	1.00	54.98	B	C
ATOM	1720	NH1	ARG	295	18.751	-14.786	5.790	1.00	56.63	B	N
ATOM	1721	NH2	ARG	295	16.449	-14.722	5.733	1.00	55.08	B	N
ATOM	1722	C	ARG	295	20.925	-9.084	5.069	1.00	43.35	B	C
ATOM	1723	O	ARG	295	21.975	-9.718	4.948	1.00	44.39	B	O
ATOM	1724	N	VAL	296	20.353	-8.446	4.053	1.00	40.22	B	N
ATOM	1725	CA	VAL	296	20.917	-8.555	2.715	1.00	41.05	B	C
ATOM	1726	CB	VAL	296	20.026	-7.833	1.653	1.00	41.81	B	C
ATOM	1727	CG1	VAL	296	19.931	-6.339	1.964	1.00	43.65	B	C
ATOM	1728	CG2	VAL	296	20.590	-8.056	0.251	1.00	41.55	B	C
ATOM	1729	C	VAL	296	22.342	-7.981	2.665	1.00	41.93	B	C
ATOM	1730	O	VAL	296	23.240	-8.588	2.058	1.00	40.42	B	O
ATOM	1731	N	LEU	297	22.555	-6.833	3.317	1.00	40.51	B	N
ATOM	1732	CA	LEU	297	23.873	-6.198	3.334	1.00	39.85	B	C
ATOM	1733	CB	LEU	297	23.808	-4.809	3.992	1.00	36.62	B	C
ATOM	1734	CG	LEU	297	25.093	-3.965	4.013	1.00	33.08	B	C
ATOM	1735	CD1	LEU	297	25.497	-3.593	2.611	1.00	32.37	B	C
ATOM	1736	CD2	LEU	297	24.875	-2.712	4.827	1.00	32.91	B	C
ATOM	1737	C	LEU	297	24.850	-7.073	4.100	1.00	41.61	B	C
ATOM	1738	O	LEU	297	26.014	-7.193	3.724	1.00	42.84	B	O
ATOM	1739	N	ASN	298	24.381	-7.699	5.173	1.00	43.04	B	N
ATOM	1740	CA	ASN	298	25.259	-8.577	5.935	1.00	44.53	B	C
ATOM	1741	CB	ASN	298	24.556	-9.083	7.192	1.00	43.88	B	C
ATOM	1742	CG	ASN	298	24.704	-8.136	8.358	1.00	43.04	B	C
ATOM	1743	OD1	ASN	298	24.706	-6.919	8.186	1.00	43.95	B	O
ATOM	1744	ND2	ASN	298	24.837	-8.689	9.554	1.00	42.99	B	N
ATOM	1745	C	ASN	298	25.710	-9.756	5.083	1.00	45.13	B	C
ATOM	1746	O	ASN	298	26.889	-10.114	5.074	1.00	47.16	B	O

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ATOM   1747  N   ALA   299      24.763 -10.345   4.363  1.00 44.07      B    N
ATOM   1748  CA  ALA   299      25.055 -11.440   3.456  1.00 43.47      B    C
ATOM   1749  CB  ALA   299      23.842 -11.721   2.605  1.00 42.59      B    C
ATOM   1750  C   ALA   299      26.245 -11.098   2.562  1.00 43.77      B    C
ATOM   1751  O   ALA   299      27.290 -11.760   2.616  1.00 43.07      B    O
ATOM   1752  N   ALA   300      26.071 -10.056   1.747  1.00 42.89      B    N
ATOM   1753  CA  ALA   300      27.083  -9.631   0.782  1.00 41.86      B    C
ATOM   1754  CB  ALA   300      26.675  -8.321   0.123  1.00 37.77      B    C
ATOM   1755  C   ALA   300      28.419  -9.459   1.474  1.00 42.97      B    C
ATOM   1756  O   ALA   300      29.466  -9.787   0.917  1.00 44.18      B    O
ATOM   1757  N   CYS   301      28.398  -8.947   2.694  1.00 44.23      B    N
ATOM   1758  CA  CYS   301      29.648  -8.840   3.414  1.00 45.71      B    C
ATOM   1759  CB  CYS   301      29.430  -8.131   4.738  1.00 45.13      B    C
ATOM   1760  SG  CYS   301      29.438  -6.352   4.500  1.00 43.18      B    S
ATOM   1761  C   CYS   301      30.228 -10.224   3.612  1.00 47.63      B    C
ATOM   1762  O   CYS   301      31.383 -10.466   3.287  1.00 47.12      B    O
ATOM   1763  N   GLN   302      29.402 -11.137   4.111  1.00 51.11      B    N
ATOM   1764  CA  GLN   302      29.784 -12.532   4.238  1.00 54.23      B    C
ATOM   1765  CB  GLN   302      28.601 -13.345   4.729  1.00 55.02      B    C
ATOM   1766  CG  GLN   302      28.766 -14.830   4.489  1.00 57.93      B    C
ATOM   1767  CD  GLN   302      27.821 -15.644   5.325  1.00 59.06      B    C
ATOM   1768  OE1 GLN   302      26.660 -15.823   4.964  1.00 60.77      B    O
ATOM   1769  NE2 GLN   302      28.307 -16.140   6.460  1.00 61.10      B    N
ATOM   1770  C   GLN   302      30.290 -13.132   2.929  1.00 56.72      B    C
ATOM   1771  O   GLN   302      31.296 -13.840   2.913  1.00 56.31      B    O
ATOM   1772  N   ARG   303      29.596 -12.862   1.830  1.00 59.07      B    N
ATOM   1773  CA  ARG   303      30.067 -13.342   0.542  1.00 62.00      B    C
ATOM   1774  CB  ARG   303      29.103 -12.924  -0.564  1.00 65.78      B    C
ATOM   1775  CG  ARG   303      29.606 -13.207  -1.966  1.00 73.70      B    C
ATOM   1776  CD  ARG   303      30.386 -14.512  -2.031  1.00 81.75      B    C
ATOM   1777  NE  ARG   303      30.633 -14.969  -3.399  1.00 89.51      B    N
ATOM   1778  CZ  ARG   303      31.436 -15.984  -3.699  1.00 94.37      B    C
ATOM   1779  NH1 ARG   303      31.611 -16.348  -4.963  1.00 98.35      B    N
ATOM   1780  NH2 ARG   303      32.070 -16.631  -2.727  1.00 97.91      B    N
ATOM   1781  C   ARG   303      31.477 -12.811   0.255  1.00 61.71      B    C
ATOM   1782  O   ARG   303      32.407 -13.602   0.077  1.00 62.96      B    O
ATOM   1783  N   LEU   304      31.647 -11.488   0.227  1.00 59.43      B    N
ATOM   1784  CA  LEU   304      32.971 -10.898   0.007  1.00 57.53      B    C
ATOM   1785  CB  LEU   304      32.891  -9.371  -0.032  1.00 54.34      B    C
ATOM   1786  CG  LEU   304      32.255  -8.734  -1.268  1.00 52.05      B    C
ATOM   1787  CD1 LEU   304      32.077  -7.257  -1.037  1.00 52.38      B    C
ATOM   1788  CD2 LEU   304      33.112  -8.967  -2.490  1.00 50.60      B    C
ATOM   1789  C   LEU   304      33.997 -11.309   1.066  1.00 57.95      B    C
ATOM   1790  O   LEU   304      35.203 -11.297   0.809  1.00 57.93      B    O
ATOM   1791  N   ALA   305      33.531 -11.666   2.257  1.00 58.05      B    N
ATOM   1792  CA  ALA   305      34.442 -12.134   3.287  1.00 59.23      B    C
ATOM   1793  CB  ALA   305      33.734 -12.190   4.635  1.00 56.74      B    C
ATOM   1794  C   ALA   305      34.965 -13.515   2.905  1.00 61.55      B    C
ATOM   1795  O   ALA   305      36.163 -13.785   3.025  1.00 60.10      B    O
ATOM   1796  N   ARG   306      34.061 -14.374   2.428  1.00 65.01      B    N
ATOM   1797  CA  ARG   306      34.389 -15.766   2.098  1.00 67.69      B    C
ATOM   1798  CB  ARG   306      33.117 -16.562   1.772  1.00 70.93      B    C
ATOM   1799  CG  ARG   306      32.222 -16.781   2.978  1.00 76.23      B    C
ATOM   1800  CD  ARG   306      31.075 -17.736   2.703  1.00 81.71      B    C
ATOM   1801  NE  ARG   306      30.394 -18.093   3.948  1.00 88.37      B    N
ATOM   1802  CZ  ARG   306      29.232 -18.738   4.016  1.00 91.88      B    C
ATOM   1803  NH1 ARG   306      28.693 -19.016   5.202  1.00 93.39      B    N
ATOM   1804  NH2 ARG   306      28.609 -19.102   2.899  1.00 94.36      B    N
ATOM   1805  C   ARG   306      35.348 -15.843   0.921  1.00 66.50      B    C
ATOM   1806  O   ARG   306      36.274 -16.655   0.911  1.00 67.91      B    O
ATOM   1807  N   ALA   307      35.119 -14.989  -0.068  1.00 63.54      B    N
ATOM   1808  CA  ALA   307      36.020 -14.861  -1.205  1.00 59.80      B    C
ATOM   1809  CB  ALA   307      35.355 -14.024  -2.301  1.00 57.91      B    C
ATOM   1810  C   ALA   307      37.357 -14.232  -0.804  1.00 57.82      B    C
ATOM   1811  O   ALA   307      38.131 -13.823  -1.660  1.00 59.09      B    O
ATOM   1812  N   GLY   308      37.626 -14.134   0.492  1.00 54.92      B    N
ATOM   1813  CA  GLY   308      38.963 -13.757   0.909  1.00 51.44      B    C
ATOM   1814  C   GLY   308      39.210 -12.286   1.178  1.00 49.16      B    C
ATOM   1815  O   GLY   308      40.329 -11.899   1.516  1.00 48.32      B    O
ATOM   1816  N   VAL   309      38.180 -11.456   1.047  1.00 48.13      B    N
ATOM   1817  CA  VAL   309      38.370 -10.021   1.235  1.00 46.11      B    C
ATOM   1818  CB  VAL   309      37.356  -9.209   0.421  1.00 44.93      B    C
ATOM   1819  CG1 VAL   309      37.720  -7.748   0.489  1.00 44.68      B    C
ATOM   1820  CG2 VAL   309      37.329  -9.689  -1.027  1.00 42.19      B    C
ATOM   1821  C   VAL   309      38.264  -9.607   2.704  1.00 46.55      B    C
ATOM   1822  O   VAL   309      37.516 -10.209   3.472  1.00 49.09      B    O
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ATOM	1823	N	VAL	310	39.030	-8.587	3.091	1.00	45.59	B	N
ATOM	1824	CA	VAL	310	38.948	-8.015	4.437	1.00	44.13	B	C
ATOM	1825	CB	VAL	310	40.353	-7.668	4.994	1.00	45.86	B	C
ATOM	1826	CG1	VAL	310	40.219	-6.873	6.295	1.00	46.71	B	C
ATOM	1827	CG2	VAL	310	41.164	-8.939	5.221	1.00	46.65	B	C
ATOM	1828	C	VAL	310	38.125	-6.726	4.433	1.00	42.98	B	C
ATOM	1829	O	VAL	310	38.601	-5.679	3.982	1.00	43.01	B	O
ATOM	1830	N	LEU	311	36.898	-6.799	4.942	1.00	41.73	B	N
ATOM	1831	CA	LEU	311	36.047	-5.612	5.081	1.00	38.57	B	C
ATOM	1832	CB	LEU	311	34.578	-5.978	4.883	1.00	38.91	B	C
ATOM	1833	CG	LEU	311	34.197	-6.356	3.455	1.00	41.16	B	C
ATOM	1834	CD1	LEU	311	35.473	-6.615	2.644	1.00	43.08	B	C
ATOM	1835	CD2	LEU	311	33.280	-7.585	3.481	1.00	40.54	B	C
ATOM	1836	C	LEU	311	36.210	-4.974	6.448	1.00	36.32	B	C
ATOM	1837	O	LEU	311	35.990	-5.630	7.471	1.00	33.22	B	O
ATOM	1838	N	VAL	312	36.598	-3.696	6.454	1.00	36.10	B	N
ATOM	1839	CA	VAL	312	36.634	-2.901	7.682	1.00	36.13	B	C
ATOM	1840	CB	VAL	312	37.971	-2.156	7.835	1.00	33.57	B	C
ATOM	1841	CG1	VAL	312	37.949	-1.274	9.061	1.00	30.27	B	C
ATOM	1842	CG2	VAL	312	39.094	-3.163	7.964	1.00	34.18	B	C
ATOM	1843	C	VAL	312	35.507	-1.893	7.646	1.00	37.18	B	C
ATOM	1844	O	VAL	312	35.316	-1.213	6.643	1.00	39.23	B	O
ATOM	1845	N	THR	313	34.740	-1.803	8.728	1.00	37.86	B	N
ATOM	1846	CA	THR	313	33.611	-0.882	8.744	1.00	37.25	B	C
ATOM	1847	CB	THR	313	32.291	-1.645	8.538	1.00	36.13	B	C
ATOM	1848	OG1	THR	313	31.263	-0.721	8.168	1.00	37.57	B	O
ATOM	1849	CG2	THR	313	31.887	-2.358	9.814	1.00	33.52	B	C
ATOM	1850	C	THR	313	33.531	-0.102	10.053	1.00	36.40	B	C
ATOM	1851	O	THR	313	34.036	-0.561	11.079	1.00	36.86	B	O
ATOM	1852	N	ALA	314	32.894	1.071	10.008	1.00	34.39	B	N
ATOM	1853	CA	ALA	314	32.627	1.873	11.205	1.00	31.94	B	C
ATOM	1854	CB	ALA	314	32.299	3.285	10.802	1.00	32.67	B	C
ATOM	1855	C	ALA	314	31.488	1.308	12.050	1.00	31.52	B	C
ATOM	1856	O	ALA	314	30.443	0.911	11.521	1.00	31.50	B	O
ATOM	1857	N	ALA	315	31.682	1.295	13.364	1.00	31.10	B	N
ATOM	1858	CA	ALA	315	30.728	0.667	14.277	1.00	32.53	B	C
ATOM	1859	CB	ALA	315	31.354	0.496	15.654	1.00	30.26	B	C
ATOM	1860	C	ALA	315	29.437	1.468	14.400	1.00	35.34	B	C
ATOM	1861	O	ALA	315	28.383	0.913	14.708	1.00	36.03	B	O
ATOM	1862	N	GLY	316	29.520	2.774	14.165	1.00	36.52	B	N
ATOM	1863	CA	GLY	316	28.340	3.606	14.262	1.00	37.34	B	C
ATOM	1864	C	GLY	316	28.494	4.613	15.378	1.00	39.62	B	C
ATOM	1865	O	GLY	316	29.157	4.317	16.376	1.00	41.26	B	O
ATOM	1866	N	ASN	317	27.870	5.782	15.216	1.00	38.59	B	N
ATOM	1867	CA	ASN	317	28.134	6.964	16.041	1.00	38.47	B	C
ATOM	1868	CB	ASN	317	28.289	8.205	15.168	1.00	38.39	B	C
ATOM	1869	CG	ASN	317	29.539	8.187	14.381	1.00	39.27	B	C
ATOM	1870	OD1	ASN	317	30.506	7.535	14.764	1.00	40.16	B	O
ATOM	1871	ND2	ASN	317	29.548	8.902	13.268	1.00	40.64	B	N
ATOM	1872	C	ASN	317	27.033	7.264	17.029	1.00	38.47	B	C
ATOM	1873	O	ASN	317	26.906	8.408	17.488	1.00	37.31	B	O
ATOM	1874	N	PHE	318	26.217	6.272	17.343	1.00	38.66	B	N
ATOM	1875	CA	PHE	318	24.974	6.581	18.027	1.00	39.58	B	C
ATOM	1876	CB	PHE	318	23.816	5.899	17.304	1.00	38.09	B	C
ATOM	1877	CG	PHE	318	23.744	6.244	15.839	1.00	35.73	B	C
ATOM	1878	CD1	PHE	318	23.340	7.502	15.430	1.00	33.00	B	C
ATOM	1879	CD2	PHE	318	24.089	5.313	14.869	1.00	34.65	B	C
ATOM	1880	CE1	PHE	318	23.280	7.821	14.083	1.00	30.08	B	C
ATOM	1881	CE2	PHE	318	24.026	5.640	13.516	1.00	32.23	B	C
ATOM	1882	CZ	PHE	318	23.620	6.893	13.132	1.00	28.85	B	C
ATOM	1883	C	PHE	318	24.987	6.218	19.499	1.00	39.58	B	C
ATOM	1884	O	PHE	318	23.973	6.342	20.172	1.00	39.71	B	O
ATOM	1885	N	ARG	319	26.150	5.800	19.996	1.00	41.36	B	N
ATOM	1886	CA	ARG	319	26.316	5.383	21.392	1.00	43.43	B	C
ATOM	1887	CB	ARG	319	26.128	6.579	22.341	1.00	44.77	B	C
ATOM	1888	CG	ARG	319	26.990	6.509	23.603	1.00	48.54	B	C
ATOM	1889	CD	ARG	319	26.728	7.651	24.620	1.00	51.31	B	C
ATOM	1890	NE	ARG	319	27.431	7.397	25.884	1.00	55.25	B	N
ATOM	1891	CZ	ARG	319	28.457	8.111	26.349	1.00	58.43	B	C
ATOM	1892	NH1	ARG	319	28.914	9.152	25.661	1.00	59.31	B	N
ATOM	1893	NH2	ARG	319	29.051	7.763	27.490	1.00	57.79	B	N
ATOM	1894	C	ARG	319	25.308	4.271	21.720	1.00	43.91	B	C
ATOM	1895	O	ARG	319	24.741	4.212	22.820	1.00	44.45	B	O
ATOM	1896	N	ASP	320	25.101	3.396	20.743	1.00	43.13	B	N
ATOM	1897	CA	ASP	320	24.102	2.348	20.809	1.00	43.73	B	C
ATOM	1898	CB	ASP	320	23.065	2.577	19.715	1.00	42.02	B	C

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ATOM   1899  CG  ASP   320      21.795   1.807  19.954  1.00 45.94      B    C
ATOM   1900  OD1 ASP   320      21.657   1.219  21.049  1.00 48.90      B    O
ATOM   1901  OD2 ASP   320      20.926   1.785  19.048  1.00 48.43      B    O
ATOM   1902  C   ASP   320      24.826   1.020  20.580  1.00 45.02      B    C
ATOM   1903  O   ASP   320      26.034   1.010  20.291  1.00 45.17      B    O
ATOM   1904  N   ASP   321      24.111  -0.095  20.714  1.00 45.07      B    N
ATOM   1905  CA  ASP   321      24.692  -1.392  20.387  1.00 46.17      B    C
ATOM   1906  CB  ASP   321      23.789  -2.522  20.884  1.00 47.34      B    C
ATOM   1907  CG  ASP   321      24.383  -3.898  20.619  1.00 51.14      B    C
ATOM   1908  OD1 ASP   321      25.355  -3.996  19.828  1.00 51.98      B    O
ATOM   1909  OD2 ASP   321      23.888  -4.892  21.202  1.00 53.20      B    O
ATOM   1910  C   ASP   321      24.899  -1.522  18.871  1.00 46.01      B    C
ATOM   1911  O   ASP   321      23.972  -1.311  18.081  1.00 46.19      B    O
ATOM   1912  N   ALA   322      26.118  -1.879  18.473  1.00 45.06      B    N
ATOM   1913  CA  ALA   322      26.484  -1.976  17.059  1.00 44.18      B    C
ATOM   1914  CB  ALA   322      27.964  -2.302  16.936  1.00 42.99      B    C
ATOM   1915  C   ALA   322      25.666  -3.011  16.292  1.00 44.26      B    C
ATOM   1916  O   ALA   322      25.483  -2.898  15.082  1.00 42.38      B    O
ATOM   1917  N   CYS   323      26.170  -4.020  16.996  1.00 46.17      B    N
ATOM   1918  CA  CYS   323      24.540  -5.154  16.335  1.00 47.75      B    C
ATOM   1919  C   CYS   323      23.225  -4.782  15.655  1.00 47.19      B    C
ATOM   1920  O   CYS   323      22.621  -5.596  14.947  1.00 49.21      B    O
ATOM   1921  CB  CYS   323      24.302  -6.277  17.337  1.00 47.92      B    C
ATOM   1922  SG  CYS   323      25.755  -6.815  18.317  1.00 57.92      B    S
ATOM   1923  N   LEU   324      22.786  -3.547  15.859  1.00 44.42      B    N
ATOM   1924  CA  LEU   324      21.537  -3.106  15.275  1.00 41.85      B    C
ATOM   1925  CB  LEU   324      20.781  -2.256  16.288  1.00 42.78      B    C
ATOM   1926  CG  LEU   324      19.899  -3.109  17.215  1.00 42.58      B    C
ATOM   1927  CD1 LEU   324      20.409  -4.537  17.241  1.00 41.91      B    C
ATOM   1928  CD2 LEU   324      19.893  -2.514  18.621  1.00 41.27      B    C
ATOM   1929  C   LEU   324      21.746  -2.348  13.969  1.00 41.25      B    C
ATOM   1930  O   LEU   324      20.801  -1.845  13.367  1.00 38.50      B    O
ATOM   1931  N   TYR   325      22.992  -2.299  13.519  1.00 40.98      B    N
ATOM   1932  CA  TYR   325      23.347  -1.560  12.315  1.00 39.73      B    C
ATOM   1933  CB  TYR   325      24.155  -0.328  12.696  1.00 37.77      B    C
ATOM   1934  CG  TYR   325      23.499   0.517  13.742  1.00 33.29      B    C
ATOM   1935  CD1 TYR   325      22.650   1.557  13.377  1.00 34.00      B    C
ATOM   1936  CE1 TYR   325      22.022   2.334  14.321  1.00 33.26      B    C
ATOM   1937  CD2 TYR   325      23.709   0.271  15.086  1.00 30.61      B    C
ATOM   1938  CE2 TYR   325      23.087   1.038  16.039  1.00 33.72      B    C
ATOM   1939  CZ  TYR   325      22.239   2.074  15.649  1.00 33.01      B    C
ATOM   1940  OH  TYR   325      21.597   2.844  16.588  1.00 32.19      B    O
ATOM   1941  C   TYR   325      24.188  -2.407  11.377  1.00 40.49      B    C
ATOM   1942  O   TYR   325      25.078  -3.134  11.823  1.00 41.28      B    O
ATOM   1943  N   SER   326      23.936  -2.294  10.080  1.00 40.51      B    N
ATOM   1944  CA  SER   326      24.848  -2.889   9.106  1.00 41.58      B    C
ATOM   1945  CB  SER   326      24.062  -3.730   8.093  1.00 42.97      B    C
ATOM   1946  OG  SER   326      23.315  -4.736   8.753  1.00 42.86      B    O
ATOM   1947  C   SER   326      25.653  -1.791   8.393  1.00 41.09      B    C
ATOM   1948  O   SER   326      25.202  -0.655   8.270  1.00 42.79      B    O
ATOM   1949  N   PRO   327      26.860  -2.123   7.924  1.00 40.03      B    N
ATOM   1950  CD  PRO   327      27.684  -1.256   7.062  1.00 39.65      B    C
ATOM   1951  CA  PRO   327      27.418  -3.473   8.060  1.00 38.97      B    C
ATOM   1952  CB  PRO   327      28.303  -3.620   6.821  1.00 38.89      B    C
ATOM   1953  CG  PRO   327      28.693  -2.200   6.454  1.00 38.79      B    C
ATOM   1954  C   PRO   327      28.168  -3.781   9.358  1.00 37.64      B    C
ATOM   1955  O   PRO   327      28.985  -4.697   9.392  1.00 38.69      B    O
ATOM   1956  N   ALA   328      27.890  -3.037  10.424  1.00 35.56      B    N
ATOM   1957  CA  ALA   328      28.624  -3.219  11.672  1.00 34.25      B    C
ATOM   1958  CB  ALA   328      28.270  -2.119  12.651  1.00 31.78      B    C
ATOM   1959  C   ALA   328      28.316  -4.571  12.287  1.00 34.71      B    C
ATOM   1960  O   ALA   328      29.181  -5.236  12.848  1.00 32.22      B    O
ATOM   1961  N   SER   329      27.064  -4.972  12.185  1.00 37.03      B    N
ATOM   1962  CA  SER   329      26.602  -6.104  12.955  1.00 39.06      B    C
ATOM   1963  CB  SER   329      25.068  -6.120  13.024  1.00 39.81      B    C
ATOM   1964  OG  SER   329      24.482  -6.166  11.734  1.00 39.62      B    O
ATOM   1965  C   SER   329      27.123  -7.343  12.282  1.00 39.36      B    C
ATOM   1966  O   SER   329      27.029  -8.439  12.827  1.00 41.11      B    O
ATOM   1967  N   ALA   330      27.691  -7.163  11.096  1.00 38.72      B    N
ATOM   1968  CA  ALA   330      28.173  -8.304  10.324  1.00 39.45      B    C
ATOM   1969  CB  ALA   330      28.489  -7.871   8.901  1.00 36.46      B    C
ATOM   1970  C   ALA   330      29.391  -8.988  10.962  1.00 40.24      B    C
ATOM   1971  O   ALA   330      30.410  -8.355  11.276  1.00 39.63      B    O
ATOM   1972  N   PRO   331      29.287 -10.308  11.159  1.00 41.42      B    N
ATOM   1973  CD  PRO   331      28.083 -11.075  10.788  1.00 42.18      B    C
ATOM   1974  CA  PRO   331      30.275 -11.147  11.842  1.00 42.22      B    C
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ATOM	1975	CB	PRO	331	29.609	-12.530	11.681	1.00	42.39	B	C
ATOM	1976	CG	PRO	331	28.163	-12.270	11.698	1.00	42.38	B	C
ATOM	1977	C	PRO	331	31.613	-11.193	11.112	1.00	42.21	B	C
ATOM	1978	O	PRO	331	32.677	-11.112	11.727	1.00	41.29	B	O
ATOM	1979	N	GLU	332	31.552	-11.340	9.797	1.00	42.77	B	N
ATOM	1980	CA	GLU	332	32.755	-11.545	9.014	1.00	45.40	B	C
ATOM	1981	CB	GLU	332	32.432	-12.446	7.825	1.00	46.55	B	C
ATOM	1982	CG	GLU	332	31.067	-12.180	7.218	1.00	50.46	B	C
ATOM	1983	CD	GLU	332	30.035	-13.245	7.573	1.00	52.20	B	C
ATOM	1984	OE1	GLU	332	30.196	-14.405	7.128	1.00	50.77	B	O
ATOM	1985	OE2	GLU	332	29.058	-12.916	8.292	1.00	52.83	B	O
ATOM	1986	C	GLU	332	33.344	-10.209	8.551	1.00	45.69	B	C
ATOM	1987	O	GLU	332	34.237	-10.159	7.698	1.00	47.38	B	O
ATOM	1988	N	VAL	333	32.832	-9.127	9.129	1.00	45.52	B	N
ATOM	1989	CA	VAL	333	33.361	-7.777	8.926	1.00	44.61	B	C
ATOM	1990	CB	VAL	333	32.211	-6.768	8.689	1.00	43.61	B	C
ATOM	1991	CG1	VAL	333	32.749	-5.345	8.738	1.00	43.34	B	C
ATOM	1992	CG2	VAL	333	31.545	-7.051	7.350	1.00	42.09	B	C
ATOM	1993	C	VAL	333	34.169	-7.311	10.148	1.00	44.86	B	C
ATOM	1994	O	VAL	333	33.864	-7.694	11.283	1.00	44.91	B	O
ATOM	1995	N	ILE	334	35.193	-6.488	9.915	1.00	43.56	B	N
ATOM	1996	CA	ILE	334	35.937	-5.873	11.009	1.00	42.14	B	C
ATOM	1997	CB	ILE	334	37.403	-5.597	10.618	1.00	41.32	B	C
ATOM	1998	CG2	ILE	334	38.148	-4.925	11.771	1.00	39.45	B	C
ATOM	1999	CG1	ILE	334	38.108	-6.917	10.323	1.00	42.05	B	C
ATOM	2000	CD1	ILE	334	38.789	-7.525	11.528	1.00	41.74	B	C
ATOM	2001	C	ILE	334	35.276	-4.564	11.391	1.00	42.30	B	C
ATOM	2002	O	ILE	334	35.264	-3.604	10.617	1.00	44.99	B	O
ATOM	2003	N	THR	335	34.724	-4.528	12.593	1.00	40.57	B	N
ATOM	2004	CA	THR	335	33.899	-3.411	13.012	1.00	39.86	B	C
ATOM	2005	CB	THR	335	32.565	-3.934	13.560	1.00	38.34	B	C
ATOM	2006	OG1	THR	335	31.900	-4.682	12.527	1.00	35.93	B	O
ATOM	2007	CG2	THR	335	31.676	-2.783	14.016	1.00	37.95	B	C
ATOM	2008	C	THR	335	34.653	-2.615	14.064	1.00	39.84	B	C
ATOM	2009	O	THR	335	35.212	-3.185	14.999	1.00	40.88	B	O
ATOM	2010	N	VAL	336	34.695	-1.297	13.903	1.00	39.68	B	N
ATOM	2011	CA	VAL	336	35.676	-0.501	14.628	1.00	38.86	B	C
ATOM	2012	CB	VAL	336	36.670	0.190	13.674	1.00	36.14	B	C
ATOM	2013	CG1	VAL	336	37.660	1.015	14.475	1.00	34.81	B	C
ATOM	2014	CG2	VAL	336	37.398	-0.843	12.830	1.00	34.08	B	C
ATOM	2015	C	VAL	336	35.029	0.576	15.455	1.00	39.97	B	C
ATOM	2016	O	VAL	336	34.361	1.463	14.923	1.00	41.64	B	O
ATOM	2017	N	GLY	337	35.239	0.518	16.763	1.00	40.20	B	N
ATOM	2018	CA	GLY	337	34.802	1.610	17.606	1.00	39.73	B	C
ATOM	2019	C	GLY	337	35.746	2.787	17.486	1.00	39.43	B	C
ATOM	2020	O	GLY	337	36.780	2.690	16.820	1.00	39.46	B	O
ATOM	2021	N	ALA	338	35.391	3.901	18.126	1.00	39.12	B	N
ATOM	2022	CA	ALA	338	36.217	5.106	18.096	1.00	38.05	B	C
ATOM	2023	CB	ALA	338	35.435	6.256	17.479	1.00	34.55	B	C
ATOM	2024	C	ALA	338	36.667	5.466	19.509	1.00	38.46	B	C
ATOM	2025	O	ALA	338	35.876	5.430	20.451	1.00	39.47	B	O
ATOM	2026	N	THR	339	37.944	5.795	19.651	1.00	39.40	B	N
ATOM	2027	CA	THR	339	38.479	6.249	20.922	1.00	42.52	B	C
ATOM	2028	CB	THR	339	39.537	5.277	21.517	1.00	44.20	B	C
ATOM	2029	OG1	THR	339	40.599	5.071	20.576	1.00	46.75	B	O
ATOM	2030	CG2	THR	339	38.907	3.935	21.861	1.00	45.49	B	C
ATOM	2031	C	THR	339	39.154	7.569	20.684	1.00	44.11	B	C
ATOM	2032	O	THR	339	39.425	7.931	19.542	1.00	43.96	B	O
ATOM	2033	N	ASN	340	39.422	8.284	21.767	1.00	46.79	B	N
ATOM	2034	CA	ASN	340	40.081	9.577	21.685	1.00	48.44	B	C
ATOM	2035	CB	ASN	340	39.285	10.626	22.465	1.00	46.64	B	C
ATOM	2036	CG	ASN	340	39.236	10.334	23.943	1.00	45.72	B	C
ATOM	2037	OD1	ASN	340	39.719	9.294	24.401	1.00	43.67	B	O
ATOM	2038	ND2	ASN	340	38.651	11.254	24.706	1.00	45.68	B	N
ATOM	2039	C	ASN	340	41.515	9.510	22.210	1.00	47.97	B	C
ATOM	2040	O	ASN	340	42.037	8.434	22.482	1.00	46.71	B	O
ATOM	2041	N	ALA	341	42.134	10.676	22.337	1.00	49.51	B	N
ATOM	2042	CA	ALA	341	43.538	10.786	22.698	1.00	51.79	B	C
ATOM	2043	CB	ALA	341	43.997	12.243	22.534	1.00	49.71	B	C
ATOM	2044	C	ALA	341	43.801	10.296	24.110	1.00	54.25	B	C
ATOM	2045	O	ALA	341	44.954	10.227	24.547	1.00	54.89	B	O
ATOM	2046	N	GLN	342	42.730	9.964	24.829	1.00	56.61	B	N
ATOM	2047	CA	GLN	342	42.838	9.357	26.158	1.00	58.83	B	C
ATOM	2048	CB	GLN	342	41.779	9.946	27.084	1.00	60.38	B	C
ATOM	2049	CG	GLN	342	41.374	11.352	26.715	1.00	63.15	B	C
ATOM	2050	CD	GLN	342	42.561	12.272	26.637	1.00	65.06	B	C

ATOM	2051	OE1	GLN	342	42.539	13.276	25.918	1.00	66.99	B	O
ATOM	2052	NE2	GLN	342	43.616	11.939	27.381	1.00	66.05	B	N
ATOM	2053	C	GLN	342	42.660	7.832	26.105	1.00	59.71	B	C
ATOM	2054	O	GLN	342	42.625	7.165	27.143	1.00	59.74	B	O
ATOM	2055	N	ASP	343	42.547	7.289	24.895	1.00	59.69	B	N
ATOM	2056	CA	ASP	343	42.193	5.888	24.706	1.00	59.97	B	C
ATOM	2057	CB	ASP	343	43.276	4.988	25.315	1.00	60.43	B	C
ATOM	2058	CG	ASP	343	44.410	4.685	24.328	1.00	62.18	B	C
ATOM	2059	OD1	ASP	343	44.125	4.483	23.130	1.00	61.55	B	O
ATOM	2060	OD2	ASP	343	45.591	4.644	24.739	1.00	64.06	B	O
ATOM	2061	C	ASP	343	40.804	5.552	25.281	1.00	59.58	B	C
ATOM	2062	O	ASP	343	40.562	4.440	25.756	1.00	58.20	B	O
ATOM	2063	N	GLN	344	39.896	6.527	25.218	1.00	58.40	B	N
ATOM	2064	CA	GLN	344	38.531	6.362	25.712	1.00	56.68	B	C
ATOM	2065	CB	GLN	344	38.226	7.428	26.768	1.00	57.01	B	C
ATOM	2066	CG	GLN	344	38.911	7.171	28.100	1.00	59.32	B	C
ATOM	2067	CD	GLN	344	38.507	5.837	28.714	1.00	61.30	B	C
ATOM	2068	OE1	GLN	344	37.395	5.687	29.225	1.00	61.08	B	O
ATOM	2069	NE2	GLN	344	39.410	4.858	28.660	1.00	61.88	B	N
ATOM	2070	C	GLN	344	37.473	6.420	24.607	1.00	54.73	B	C
ATOM	2071	O	GLN	344	37.713	6.939	23.520	1.00	55.32	B	O
ATOM	2072	N	PRO	345	36.276	5.887	24.884	1.00	52.34	B	N
ATOM	2073	CD	PRO	345	35.864	5.218	26.130	1.00	52.86	B	C
ATOM	2074	CA	PRO	345	35.208	5.872	23.881	1.00	51.32	B	C
ATOM	2075	CB	PRO	345	34.080	5.115	24.572	1.00	52.46	B	C
ATOM	2076	CG	PRO	345	34.752	4.348	25.676	1.00	53.13	B	C
ATOM	2077	C	PRO	345	34.782	7.278	23.486	1.00	48.74	B	C
ATOM	2078	O	PRO	345	34.501	8.114	24.345	1.00	47.73	B	O
ATOM	2079	N	VAL	346	34.732	7.535	22.185	1.00	46.48	B	N
ATOM	2080	CA	VAL	346	34.432	8.875	21.700	1.00	45.16	B	C
ATOM	2081	CB	VAL	346	34.719	8.994	20.195	1.00	44.09	B	C
ATOM	2082	CG1	VAL	346	33.991	10.203	19.640	1.00	45.77	B	C
ATOM	2083	CG2	VAL	346	36.215	9.134	19.961	1.00	43.38	B	C
ATOM	2084	C	VAL	346	32.989	9.327	21.953	1.00	43.26	B	C
ATOM	2085	O	VAL	346	32.032	8.604	21.695	1.00	43.80	B	O
ATOM	2086	N	THR	347	32.833	10.535	22.466	1.00	40.93	B	N
ATOM	2087	CA	THR	347	31.529	11.168	22.437	1.00	38.99	B	C
ATOM	2088	CB	THR	347	31.267	11.969	23.727	1.00	38.98	B	C
ATOM	2089	OG1	THR	347	32.410	12.783	24.025	1.00	39.16	B	O
ATOM	2090	CG2	THR	347	31.000	11.024	24.877	1.00	37.77	B	C
ATOM	2091	C	THR	347	31.437	12.095	21.225	1.00	36.77	B	C
ATOM	2092	O	THR	347	32.324	12.917	20.981	1.00	35.24	B	O
ATOM	2093	N	LEU	348	30.359	11.933	20.463	1.00	34.41	B	N
ATOM	2094	CA	LEU	348	30.085	12.764	19.300	1.00	32.42	B	C
ATOM	2095	CB	LEU	348	29.981	11.869	18.057	1.00	30.11	B	C
ATOM	2096	CG	LEU	348	31.248	11.023	17.803	1.00	29.43	B	C
ATOM	2097	CD1	LEU	348	30.916	9.801	16.981	1.00	26.50	B	C
ATOM	2098	CD2	LEU	348	32.315	11.869	17.112	1.00	25.20	B	C
ATOM	2099	C	LEU	348	28.760	13.441	19.609	1.00	31.86	B	C
ATOM	2100	O	LEU	348	27.766	12.762	19.801	1.00	33.15	B	O
ATOM	2101	N	GLY	349	28.735	14.764	19.702	1.00	31.16	B	N
ATOM	2102	CA	GLY	349	27.519	15.402	20.194	1.00	32.64	B	C
ATOM	2103	C	GLY	349	27.086	14.817	21.534	1.00	32.03	B	C
ATOM	2104	O	GLY	349	27.831	14.868	22.484	1.00	34.71	B	O
ATOM	2105	N	THR	350	25.895	14.251	21.642	1.00	33.25	B	N
ATOM	2106	CA	THR	350	25.549	13.563	22.890	1.00	32.37	B	C
ATOM	2107	CB	THR	350	24.172	14.014	23.433	1.00	31.72	B	C
ATOM	2108	OG1	THR	350	23.142	13.664	22.502	1.00	33.43	B	O
ATOM	2109	CG2	THR	350	24.156	15.502	23.644	1.00	32.71	B	C
ATOM	2110	C	THR	350	25.553	12.039	22.721	1.00	31.43	B	C
ATOM	2111	O	THR	350	25.330	11.292	23.679	1.00	30.78	B	O
ATOM	2112	N	LEU	351	25.808	11.586	21.497	1.00	29.11	B	N
ATOM	2113	CA	LEU	351	26.001	10.177	21.238	1.00	27.69	B	C
ATOM	2114	CB	LEU	351	25.421	9.831	19.878	1.00	27.14	B	C
ATOM	2115	CG	LEU	351	23.939	10.206	19.774	1.00	28.37	B	C
ATOM	2116	CD1	LEU	351	23.314	9.489	18.590	1.00	29.77	B	C
ATOM	2117	CD2	LEU	351	23.218	9.806	21.054	1.00	27.08	B	C
ATOM	2118	C	LEU	351	27.482	9.824	21.298	1.00	29.91	B	C
ATOM	2119	O	LEU	351	28.173	10.139	22.277	1.00	31.43	B	O
ATOM	2120	N	GLY	352	27.981	9.178	20.248	1.00	30.92	B	N
ATOM	2121	CA	GLY	352	29.376	8.774	20.232	1.00	30.77	B	C
ATOM	2122	C	GLY	352	29.534	7.340	19.770	1.00	30.80	B	C
ATOM	2123	O	GLY	352	28.656	6.788	19.104	1.00	31.81	B	O
ATOM	2124	N	THR	353	30.647	6.715	20.121	1.00	30.12	B	N
ATOM	2125	CA	THR	353	30.980	5.456	19.488	1.00	30.91	B	C
ATOM	2126	CB	THR	353	32.410	5.035	19.835	1.00	29.91	B	C

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ATOM   2127  OG1 THR   353      32.743   3.818  19.152  1.00 26.96      B    O
ATOM   2128  CG2 THR   353      32.539   4.829  21.332  1.00 31.40      B    C
ATOM   2129  C   THR   353      30.020   4.381  19.955  1.00 32.86      B    C
ATOM   2130  O   THR   353      29.869   4.171  21.162  1.00 32.88      B    O
ATOM   2131  N   ASN   354      29.371   3.710  19.002  1.00 32.48      B    N
ATOM   2132  CA  ASN   354      28.714   2.430  19.277  1.00 34.53      B    C
ATOM   2133  CB  ASN   354      28.125   1.825  17.994  1.00 32.64      B    C
ATOM   2134  CG  ASN   354      26.811   2.451  17.611  1.00 32.64      B    C
ATOM   2135  OD1 ASN   354      26.421   3.484  18.158  1.00 34.86      B    O
ATOM   2136  ND2 ASN   354      26.116   1.838  16.669  1.00 30.53      B    N
ATOM   2137  C   ASN   354      29.656   1.399  19.910  1.00 34.76      B    C
ATOM   2138  O   ASN   354      30.882   1.558  19.919  1.00 32.98      B    O
ATOM   2139  N   PHE   355      29.059   0.326  20.413  1.00 36.79      B    N
ATOM   2140  CA  PHE   355      29.770  -0.671  21.207  1.00 39.22      B    C
ATOM   2141  CB  PHE   355      29.805  -0.217  22.661  1.00 39.08      B    C
ATOM   2142  CG  PHE   355      28.478   0.292  23.150  1.00 39.28      B    C
ATOM   2143  CD1 PHE   355      28.297   1.635  23.433  1.00 37.87      B    C
ATOM   2144  CD2 PHE   355      27.399  -0.570  23.280  1.00 39.36      B    C
ATOM   2145  CE1 PHE   355      27.073   2.107  23.831  1.00 38.35      B    C
ATOM   2146  CE2 PHE   355      26.166  -0.100  23.681  1.00 39.17      B    C
ATOM   2147  CZ  PHE   355      26.004   1.241  23.955  1.00 38.97      B    C
ATOM   2148  C   PHE   355      29.058  -2.022  21.128  1.00 41.42      B    C
ATOM   2149  O   PHE   355      28.135  -2.222  20.327  1.00 42.06      B    O
ATOM   2150  N   GLY   356      29.472  -2.945  21.985  1.00 43.05      B    N
ATOM   2151  CA  GLY   356      28.791  -4.217  22.030  1.00 46.32      B    C
ATOM   2152  C   GLY   356      29.615  -5.316  21.418  1.00 49.86      B    C
ATOM   2153  O   GLY   356      30.734  -5.080  20.960  1.00 49.34      B    O
ATOM   2154  N   ARG   357      29.067  -6.525  21.415  1.00 52.59      B    N
ATOM   2155  CA  ARG   357      29.853  -7.687  21.050  1.00 55.32      B    C
ATOM   2156  CB  ARG   357      29.054  -8.957  21.316  1.00 56.96      B    C
ATOM   2157  CG  ARG   357      27.733  -9.016  20.601  1.00 57.12      B    C
ATOM   2158  CD  ARG   357      26.958 -10.270  20.970  1.00 60.05      B    C
ATOM   2159  NE  ARG   357      25.646 -10.272  20.324  1.00 66.70      B    N
ATOM   2160  CZ  ARG   357      25.436 -10.567  19.041  1.00 69.70      B    C
ATOM   2161  NH1 ARG   357      24.202 -10.535  18.550  1.00 71.70      B    N
ATOM   2162  NH2 ARG   357      26.455 -10.900  18.249  1.00 70.96      B    N
ATOM   2163  C   ARG   357      30.282  -7.628  19.584  1.00 55.87      B    C
ATOM   2164  O   ARG   357      31.312  -8.195  19.208  1.00 58.14      B    O
ATOM   2165  N   CYS   358      29.493  -6.925  18.770  1.00 53.84      B    N
ATOM   2166  CA  CYS   358      29.711  -6.841  17.328  1.00 50.17      B    C
ATOM   2167  C   CYS   358      30.828  -5.854  16.927  1.00 48.24      B    C
ATOM   2168  O   CYS   358      31.055  -5.601  15.744  1.00 46.71      B    O
ATOM   2169  CB  CYS   358      28.381  -6.484  16.648  1.00 50.59      B    C
ATOM   2170  SG  CYS   358      27.087  -7.713  17.021  1.00 53.47      B    S
ATOM   2171  N   VAL   359      31.531  -5.306  17.912  1.00 46.32      B    N
ATOM   2172  CA  VAL   359      32.716  -4.503  17.630  1.00 45.82      B    C
ATOM   2173  CB  VAL   359      32.806  -3.282  18.567  1.00 44.83      B    C
ATOM   2174  CG1 VAL   359      34.050  -2.469  18.240  1.00 42.10      B    C
ATOM   2175  CG2 VAL   359      31.541  -2.426  18.436  1.00 44.41      B    C
ATOM   2176  C   VAL   359      33.968  -5.344  17.829  1.00 46.74      B    C
ATOM   2177  O   VAL   359      34.087  -6.061  18.821  1.00 48.03      B    O
ATOM   2178  N   ASP   360      34.908  -5.262  16.893  1.00 46.35      B    N
ATOM   2179  CA  ASP   360      36.122  -6.067  16.991  1.00 44.78      B    C
ATOM   2180  CB  ASP   360      36.624  -6.448  15.597  1.00 44.72      B    C
ATOM   2181  CG  ASP   360      35.649  -7.354  14.859  1.00 46.97      B    C
ATOM   2182  OD1 ASP   360      35.617  -8.576  15.133  1.00 48.10      B    O
ATOM   2183  OD2 ASP   360      34.902  -6.842  14.005  1.00 49.17      B    O
ATOM   2184  C   ASP   360      37.197  -5.326  17.756  1.00 43.17      B    C
ATOM   2185  O   ASP   360      38.004  -5.931  18.438  1.00 44.20      B    O
ATOM   2186  N   LEU   361      37.192  -4.008  17.653  1.00 42.15      B    N
ATOM   2187  CA  LEU   361      38.119  -3.190  18.417  1.00 42.07      B    C
ATOM   2188  CB  LEU   361      39.577  -3.571  18.098  1.00 42.02      B    C
ATOM   2189  CG  LEU   361      40.202  -3.182  16.739  1.00 43.57      B    C
ATOM   2190  CD1 LEU   361      41.713  -3.002  16.874  1.00 42.62      B    C
ATOM   2191  CD2 LEU   361      39.894  -4.237  15.704  1.00 41.99      B    C
ATOM   2192  C   LEU   361      37.879  -1.715  18.092  1.00 42.18      B    C
ATOM   2193  O   LEU   361      36.995  -1.375  17.297  1.00 40.87      B    O
ATOM   2194  N   PHE   362      38.669  -0.850  18.717  1.00 42.23      B    N
ATOM   2195  CA  PHE   362      38.589   0.587  18.498  1.00 43.61      B    C
ATOM   2196  CB  PHE   362      38.373   1.307  19.834  1.00 42.71      B    C
ATOM   2197  CG  PHE   362      37.078   0.961  20.497  1.00 43.77      B    C
ATOM   2198  CD1 PHE   362      36.206   1.953  20.907  1.00 44.12      B    C
ATOM   2199  CD2 PHE   362      36.716  -0.361  20.688  1.00 43.83      B    C
ATOM   2200  CE1 PHE   362      34.987   1.627  21.494  1.00 44.59      B    C
ATOM   2201  CE2 PHE   362      35.505  -0.693  21.272  1.00 44.45      B    C
ATOM   2202  CZ  PHE   362      34.640   0.300  21.677  1.00 43.51      B    C
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ATOM	2203	C	PHE	362	39.873	1.091	17.844	1.00	43.97	B	C
ATOM	2204	O	PHE	362	40.813	0.327	17.648	1.00	44.43	B	O
ATOM	2205	N	ALA	363	39.896	2.374	17.498	1.00	44.29	B	N
ATOM	2206	CA	ALA	363	41.108	3.047	17.043	1.00	42.94	B	C
ATOM	2207	CB	ALA	363	41.400	2.678	15.606	1.00	41.34	B	C
ATOM	2208	C	ALA	363	40.849	4.544	17.170	1.00	42.78	B	C
ATOM	2209	O	ALA	363	39.703	4.971	17.259	1.00	40.55	B	O
ATOM	2210	N	PRO	364	41.909	5.362	17.184	1.00	44.63	B	N
ATOM	2211	CD	PRO	364	43.317	5.026	16.940	1.00	45.92	B	C
ATOM	2212	CA	PRO	364	41.721	6.807	17.314	1.00	45.95	B	C
ATOM	2213	CB	PRO	364	43.094	7.378	16.984	1.00	45.73	B	C
ATOM	2214	CG	PRO	364	44.027	6.299	17.301	1.00	45.76	B	C
ATOM	2215	C	PRO	364	40.669	7.297	16.329	1.00	47.71	B	C
ATOM	2216	O	PRO	364	40.728	6.973	15.138	1.00	46.96	B	O
ATOM	2217	N	GLY	365	39.716	8.076	16.839	1.00	48.81	B	N
ATOM	2218	CA	GLY	365	38.632	8.575	16.016	1.00	50.95	B	C
ATOM	2219	C	GLY	365	38.112	9.945	16.425	1.00	52.82	B	C
ATOM	2220	O	GLY	365	37.095	10.417	15.896	1.00	54.20	B	O
ATOM	2221	N	GLU	366	38.798	10.585	17.367	1.00	53.12	B	N
ATOM	2222	CA	GLU	366	38.454	11.940	17.772	1.00	53.18	B	C
ATOM	2223	CB	GLU	366	38.184	11.985	19.284	1.00	55.16	B	C
ATOM	2224	CG	GLU	366	37.350	13.188	19.757	1.00	57.81	B	C
ATOM	2225	CD	GLU	366	37.236	13.294	21.288	1.00	59.11	B	C
ATOM	2226	OE1	GLU	366	36.271	12.740	21.874	1.00	59.52	B	O
ATOM	2227	OE2	GLU	366	38.120	13.938	21.902	1.00	58.89	B	O
ATOM	2228	C	GLU	366	39.627	12.840	17.416	1.00	51.60	B	C
ATOM	2229	O	GLU	366	40.750	12.567	17.799	1.00	53.06	B	O
ATOM	2230	N	ASP	367	39.375	13.900	16.663	1.00	50.68	B	N
ATOM	2231	CA	ASP	367	40.416	14.884	16.377	1.00	48.72	B	C
ATOM	2232	CB	ASP	367	41.114	15.288	17.680	1.00	51.98	B	C
ATOM	2233	CG	ASP	367	42.050	16.475	17.506	1.00	55.60	B	C
ATOM	2234	OD1	ASP	367	41.637	17.490	16.888	1.00	57.07	B	O
ATOM	2235	OD2	ASP	367	43.200	16.388	17.998	1.00	57.74	B	O
ATOM	2236	C	ASP	367	41.442	14.320	15.403	1.00	45.06	B	C
ATOM	2237	O	ASP	367	42.648	14.432	15.614	1.00	44.79	B	O
ATOM	2238	N	ILE	368	40.977	13.694	14.337	1.00	40.15	B	N
ATOM	2239	CA	ILE	368	41.924	13.092	13.417	1.00	35.41	B	C
ATOM	2240	CB	ILE	368	41.340	11.830	12.776	1.00	32.03	B	C
ATOM	2241	CG2	ILE	368	42.346	11.236	11.818	1.00	30.65	B	C
ATOM	2242	CG1	ILE	368	40.900	10.843	13.860	1.00	28.33	B	C
ATOM	2243	CD1	ILE	368	41.884	10.653	14.993	1.00	22.93	B	C
ATOM	2244	C	ILE	368	42.275	14.099	12.318	1.00	35.65	B	C
ATOM	2245	O	ILE	368	41.392	14.566	11.584	1.00	34.20	B	O
ATOM	2246	N	ILE	369	43.558	14.444	12.209	1.00	33.12	B	N
ATOM	2247	CA	ILE	369	43.975	15.381	11.180	1.00	31.13	B	C
ATOM	2248	CB	ILE	369	45.308	16.061	11.529	1.00	31.54	B	C
ATOM	2249	CG2	ILE	369	46.469	15.233	11.000	1.00	29.64	B	C
ATOM	2250	CG1	ILE	369	45.366	17.448	10.875	1.00	31.98	B	C
ATOM	2251	CD1	ILE	369	46.541	18.283	11.339	1.00	32.59	B	C
ATOM	2252	C	ILE	369	44.140	14.625	9.867	1.00	30.83	B	C
ATOM	2253	O	ILE	369	44.605	13.483	9.850	1.00	31.37	B	O
ATOM	2254	N	GLY	370	43.747	15.264	8.773	1.00	30.51	B	N
ATOM	2255	CA	GLY	370	43.750	14.606	7.482	1.00	32.26	B	C
ATOM	2256	C	GLY	370	43.444	15.582	6.365	1.00	32.89	B	C
ATOM	2257	O	GLY	370	43.140	16.755	6.619	1.00	33.94	B	O
ATOM	2258	N	ALA	371	43.524	15.102	5.126	1.00	32.44	B	N
ATOM	2259	CA	ALA	371	43.276	15.930	3.937	1.00	31.69	B	C
ATOM	2260	CB	ALA	371	43.233	15.044	2.699	1.00	28.10	B	C
ATOM	2261	C	ALA	371	41.974	16.729	4.029	1.00	32.16	B	C
ATOM	2262	O	ALA	371	40.945	16.180	4.402	1.00	34.89	B	O
ATOM	2263	N	SER	372	42.009	18.015	3.684	1.00	31.83	B	N
ATOM	2264	CA	SER	372	40.774	18.781	3.552	1.00	31.63	B	C
ATOM	2265	CB	SER	372	40.803	20.030	4.414	1.00	29.79	B	C
ATOM	2266	OG	SER	372	39.646	20.803	4.140	1.00	27.55	B	O
ATOM	2267	C	SER	372	40.489	19.198	2.121	1.00	31.94	B	C
ATOM	2268	O	SER	372	41.357	19.687	1.428	1.00	33.59	B	O
ATOM	2269	N	SER	373	39.256	19.019	1.681	1.00	33.37	B	N
ATOM	2270	CA	SER	373	38.922	19.262	0.285	1.00	32.32	B	C
ATOM	2271	CB	SER	373	37.592	18.575	-0.069	1.00	29.34	B	C
ATOM	2272	OG	SER	373	36.498	19.166	0.624	1.00	29.96	B	O
ATOM	2273	C	SER	373	38.846	20.767	0.034	1.00	32.38	B	C
ATOM	2274	O	SER	373	38.728	21.215	-1.111	1.00	28.01	B	O
ATOM	2275	N	ASP	374	38.922	21.537	1.120	1.00	35.17	B	N
ATOM	2276	CA	ASP	374	38.882	23.005	1.040	1.00	38.35	B	C
ATOM	2277	CB	ASP	374	39.132	23.641	2.416	1.00	39.54	B	C
ATOM	2278	CG	ASP	374	37.900	23.614	3.316	1.00	42.83	B	C

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ATOM   2279  OD1 ASP   374      36.992  22.785   3.091  1.00 44.31      B    O
ATOM   2280  OD2 ASP   374      37.837  24.429   4.261  1.00 46.41      B    O
ATOM   2281  C   ASP   374      39.910  23.533   0.047  1.00 39.30      B    C
ATOM   2282  O   ASP   374      39.578  24.303  -0.846  1.00 40.00      B    O
ATOM   2283  N   CYS   375      41.159  23.116   0.189  1.00 39.54      B    N
ATOM   2284  CA  CYS   375      42.150  23.476  -0.807  1.00 41.25      B    C
ATOM   2285  C   CYS   375      43.162  22.362  -0.892  1.00 42.24      B    C
ATOM   2286  O   CYS   375      43.130  21.444  -0.071  1.00 44.57      B    O
ATOM   2287  CB  CYS   375      42.836  24.787  -0.423  1.00 42.60      B    C
ATOM   2288  SG  CYS   375      44.265  24.654   0.698  1.00 43.30      B    S
ATOM   2289  N   SER   376      44.059  22.445  -1.872  1.00 40.33      B    N
ATOM   2290  CA  SER   376      44.898  21.307  -2.254  1.00 37.25      B    C
ATOM   2291  CB  SER   376      45.615  21.602  -3.573  1.00 39.26      B    C
ATOM   2292  OG  SER   376      46.458  22.749  -3.441  1.00 40.65      B    O
ATOM   2293  C   SER   376      45.947  20.922  -1.222  1.00 35.77      B    C
ATOM   2294  O   SER   376      46.542  19.855  -1.319  1.00 34.73      B    O
ATOM   2295  N   THR   377      46.216  21.798  -0.259  1.00 34.14      B    N
ATOM   2296  CA  THR   377      47.278  21.526   0.710  1.00 32.96      B    C
ATOM   2297  CB  THR   377      48.426  22.559   0.603  1.00 29.40      B    C
ATOM   2298  OG1 THR   377      47.982  23.838   1.082  1.00 29.82      B    O
ATOM   2299  CG2 THR   377      48.869  22.695  -0.842  1.00 24.15      B    C
ATOM   2300  C   THR   377      46.685  21.582   2.103  1.00 34.62      B    C
ATOM   2301  O   THR   377      47.349  21.282   3.099  1.00 34.86      B    O
ATOM   2302  N   CYS   378      45.416  21.961   2.138  1.00 35.87      B    N
ATOM   2303  CA  CYS   378      44.663  22.146   3.364  1.00 39.20      B    C
ATOM   2304  C   CYS   378      44.469  20.854   4.157  1.00 38.82      B    C
ATOM   2305  O   CYS   378      44.251  19.800   3.583  1.00 38.85      B    O
ATOM   2306  CB  CYS   378      43.310  22.784   3.006  1.00 41.18      B    C
ATOM   2307  SG  CYS   378      43.483  24.562   2.585  1.00 47.55      B    S
ATOM   2308  N   PHE   379      44.573  20.955   5.481  1.00 40.10      B    N
ATOM   2309  CA  PHE   379      44.255  19.858   6.404  1.00 40.73      B    C
ATOM   2310  CB  PHE   379      45.438  19.563   7.336  1.00 42.07      B    C
ATOM   2311  CG  PHE   379      46.518  18.732   6.711  1.00 44.32      B    C
ATOM   2312  CD1 PHE   379      47.646  19.330   6.174  1.00 45.53      B    C
ATOM   2313  CD2 PHE   379      46.411  17.349   6.667  1.00 44.45      B    C
ATOM   2314  CE1 PHE   379      48.652  18.557   5.601  1.00 48.10      B    C
ATOM   2315  CE2 PHE   379      47.405  16.579   6.100  1.00 45.09      B    C
ATOM   2316  CZ  PHE   379      48.529  17.180   5.567  1.00 46.79      B    C
ATOM   2317  C   PHE   379      43.062  20.238   7.268  1.00 40.73      B    C
ATOM   2318  O   PHE   379      42.756  21.421   7.438  1.00 39.27      B    O
ATOM   2319  N   VAL   380      42.391  19.236   7.827  1.00 41.60      B    N
ATOM   2320  CA  VAL   380      41.339  19.503   8.814  1.00 41.83      B    C
ATOM   2321  CB  VAL   380      39.963  19.731   8.147  1.00 39.80      B    C
ATOM   2322  CG1 VAL   380      39.411  18.415   7.625  1.00 37.70      B    C
ATOM   2323  CG2 VAL   380      39.017  20.360   9.128  1.00 38.24      B    C
ATOM   2324  C   VAL   380      41.200  18.347   9.787  1.00 42.06      B    C
ATOM   2325  O   VAL   380      41.609  17.223   9.496  1.00 40.84      B    O
ATOM   2326  N   SER   381      40.620  18.634  10.944  1.00 43.79      B    N
ATOM   2327  CA  SER   381      40.349  17.600  11.936  1.00 45.78      B    C
ATOM   2328  CB  SER   381      40.716  18.123  13.330  1.00 46.59      B    C
ATOM   2329  OG  SER   381      40.806  17.069  14.279  1.00 51.19      B    O
ATOM   2330  C   SER   381      38.869  17.145  11.900  1.00 45.40      B    C
ATOM   2331  O   SER   381      37.958  17.954  11.746  1.00 43.60      B    O
ATOM   2332  N   GLN   382      38.645  15.841  12.035  1.00 46.71      B    N
ATOM   2333  CA  GLN   382      37.298  15.291  12.031  1.00 47.56      B    C
ATOM   2334  CB  GLN   382      36.971  14.768  10.648  1.00 47.78      B    C
ATOM   2335  CG  GLN   382      37.028  15.855   9.630  1.00 53.25      B    C
ATOM   2336  CD  GLN   382      36.169  15.571   8.438  1.00 56.07      B    C
ATOM   2337  OE1 GLN   382      35.414  16.443   7.989  1.00 58.19      B    O
ATOM   2338  NE2 GLN   382      36.269  14.345   7.905  1.00 57.86      B    N
ATOM   2339  C   GLN   382      37.155  14.172  13.041  1.00 47.35      B    C
ATOM   2340  O   GLN   382      38.134  13.492  13.362  1.00 49.75      B    O
ATOM   2341  N   SER   383      35.936  13.978  13.538  1.00 44.94      B    N
ATOM   2342  CA  SER   383      35.651  12.855  14.426  1.00 42.17      B    C
ATOM   2343  CB  SER   383      35.265  13.366  15.810  1.00 40.00      B    C
ATOM   2344  OG  SER   383      36.295  14.160  16.353  1.00 35.61      B    O
ATOM   2345  C   SER   383      34.550  11.933  13.903  1.00 41.99      B    C
ATOM   2346  O   SER   383      33.657  12.354  13.174  1.00 40.39      B    O
ATOM   2347  N   GLY   384      34.626  10.667  14.295  1.00 42.52      B    N
ATOM   2348  CA  GLY   384      33.608   9.706  13.907  1.00 42.23      B    C
ATOM   2349  C   GLY   384      34.158   8.293  13.905  1.00 41.28      B    C
ATOM   2350  O   GLY   384      35.375   8.104  13.941  1.00 39.89      B    O
ATOM   2351  N   THR   385      33.285   7.294  13.862  1.00 40.09      B    N
ATOM   2352  CA  THR   385      33.784   5.945  13.746  1.00 41.46      B    C
ATOM   2353  CB  THR   385      32.691   4.873  13.979  1.00 40.99      B    C
ATOM   2354  OG1 THR   385      31.490   5.217  13.268  1.00 41.65      B    O
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ATOM	2355	CG2	THR	385	32.414	4.731	15.454	1.00	41.03	B	C
ATOM	2356	C	THR	385	34.405	5.741	12.375	1.00	43.11	B	C
ATOM	2357	O	THR	385	35.218	4.836	12.187	1.00	45.39	B	O
ATOM	2358	N	SER	386	34.043	6.579	11.414	1.00	42.03	B	N
ATOM	2359	CA	SER	386	34.672	6.478	10.112	1.00	40.89	B	C
ATOM	2360	CB	SER	386	34.002	7.424	9.132	1.00	42.12	B	C
ATOM	2361	OG	SER	386	32.735	6.907	8.763	1.00	45.70	B	O
ATOM	2362	C	SER	386	36.167	6.760	10.182	1.00	39.87	B	C
ATOM	2363	O	SER	386	36.960	6.079	9.529	1.00	38.99	B	O
ATOM	2364	N	GLN	387	36.563	7.747	10.979	1.00	38.57	B	N
ATOM	2365	CA	GLN	387	37.986	8.019	11.164	1.00	38.45	B	C
ATOM	2366	CB	GLN	387	38.179	9.266	12.025	1.00	38.28	B	C
ATOM	2367	CG	GLN	387	37.233	10.404	11.670	1.00	37.87	B	C
ATOM	2368	CD	GLN	387	37.271	10.745	10.198	1.00	37.13	B	C
ATOM	2369	OE1	GLN	387	38.334	10.910	9.629	1.00	41.19	B	O
ATOM	2370	NE2	GLN	387	36.107	10.849	9.575	1.00	38.64	B	N
ATOM	2371	C	GLN	387	38.641	6.810	11.846	1.00	38.88	B	C
ATOM	2372	O	GLN	387	39.767	6.427	11.518	1.00	39.69	B	O
ATOM	2373	N	ALA	388	37.922	6.199	12.786	1.00	37.00	B	N
ATOM	2374	CA	ALA	388	38.434	5.042	13.507	1.00	34.63	B	C
ATOM	2375	CB	ALA	388	37.467	4.654	14.623	1.00	31.74	B	C
ATOM	2376	C	ALA	388	38.647	3.865	12.556	1.00	34.56	B	C
ATOM	2377	O	ALA	388	39.647	3.155	12.640	1.00	35.06	B	O
ATOM	2378	N	ALA	389	37.702	3.657	11.648	1.00	34.57	B	N
ATOM	2379	CA	ALA	389	37.788	2.550	10.703	1.00	33.13	B	C
ATOM	2380	CB	ALA	389	36.454	2.356	9.991	1.00	32.48	B	C
ATOM	2381	C	ALA	389	38.883	2.795	9.680	1.00	33.71	B	C
ATOM	2382	O	ALA	389	39.514	1.855	9.214	1.00	34.66	B	O
ATOM	2383	N	ALA	390	39.109	4.056	9.319	1.00	34.06	B	N
ATOM	2384	CA	ALA	390	40.198	4.367	8.407	1.00	35.02	B	C
ATOM	2385	CB	ALA	390	40.296	5.872	8.210	1.00	32.15	B	C
ATOM	2386	C	ALA	390	41.523	3.795	8.973	1.00	35.11	B	C
ATOM	2387	O	ALA	390	42.236	3.071	8.279	1.00	33.97	B	O
ATOM	2388	N	HIS	391	41.820	4.092	10.238	1.00	34.71	B	N
ATOM	2389	CA	HIS	391	43.073	3.561	10.846	1.00	37.06	B	C
ATOM	2390	CB	HIS	391	43.133	4.076	12.321	1.00	37.89	B	C
ATOM	2391	CG	HIS	391	43.435	5.528	12.536	1.00	41.44	B	C
ATOM	2392	CD2	HIS	391	44.268	6.146	13.407	1.00	42.12	B	C
ATOM	2393	ND1	HIS	391	42.808	6.534	11.833	1.00	41.80	B	N
ATOM	2394	CE1	HIS	391	43.237	7.707	12.263	1.00	42.62	B	C
ATOM	2395	NE2	HIS	391	44.124	7.501	13.219	1.00	42.69	B	N
ATOM	2396	C	HIS	391	43.230	2.152	10.749	1.00	37.12	B	C
ATOM	2397	O	HIS	391	44.272	1.646	10.336	1.00	38.23	B	O
ATOM	2398	N	VAL	392	42.191	1.429	11.142	1.00	36.22	B	N
ATOM	2399	CA	VAL	392	42.251	-0.016	11.115	1.00	33.77	B	C
ATOM	2400	CB	VAL	392	40.988	-0.626	11.701	1.00	31.11	B	C
ATOM	2401	CG1	VAL	392	40.991	-2.135	11.491	1.00	27.16	B	C
ATOM	2402	CG2	VAL	392	40.905	-0.282	13.170	1.00	28.41	B	C
ATOM	2403	C	VAL	392	42.401	-0.488	9.681	1.00	34.43	B	C
ATOM	2404	O	VAL	392	43.086	-1.479	9.415	1.00	35.64	B	O
ATOM	2405	N	ALA	393	41.758	0.227	8.759	1.00	33.11	B	N
ATOM	2406	CA	ALA	393	41.821	-0.130	7.354	1.00	33.65	B	C
ATOM	2407	CB	ALA	393	40.934	0.809	6.525	1.00	31.34	B	C
ATOM	2408	C	ALA	393	43.278	-0.025	6.911	1.00	34.97	B	C
ATOM	2409	O	ALA	393	43.755	-0.830	6.105	1.00	33.98	B	O
ATOM	2410	N	GLY	394	43.977	0.969	7.454	1.00	35.70	B	N
ATOM	2411	CA	GLY	394	45.345	1.232	7.045	1.00	36.60	B	C
ATOM	2412	C	GLY	394	46.253	0.163	7.604	1.00	38.29	B	C
ATOM	2413	O	GLY	394	47.080	-0.405	6.887	1.00	37.89	B	O
ATOM	2414	N	ILE	395	46.080	-0.123	8.890	1.00	39.54	B	N
ATOM	2415	CA	ILE	395	46.869	-1.141	9.553	1.00	39.77	B	C
ATOM	2416	CB	ILE	395	46.439	-1.312	11.010	1.00	39.22	B	C
ATOM	2417	CG2	ILE	395	47.192	-2.480	11.636	1.00	36.68	B	C
ATOM	2418	CG1	ILE	395	46.666	-0.005	11.780	1.00	37.42	B	C
ATOM	2419	CD1	ILE	395	46.141	-0.037	13.210	1.00	34.61	B	C
ATOM	2420	C	ILE	395	46.634	-2.441	8.824	1.00	41.71	B	C
ATOM	2421	O	ILE	395	47.568	-3.143	8.476	1.00	43.58	B	O
ATOM	2422	N	ALA	396	45.376	-2.756	8.578	1.00	43.04	B	N
ATOM	2423	CA	ALA	396	45.054	-3.961	7.833	1.00	45.09	B	C
ATOM	2424	CB	ALA	396	43.574	-3.968	7.464	1.00	43.55	B	C
ATOM	2425	C	ALA	396	45.903	-4.019	6.573	1.00	46.33	B	C
ATOM	2426	O	ALA	396	46.203	-5.099	6.062	1.00	47.54	B	O
ATOM	2427	N	ALA	397	46.276	-2.844	6.073	1.00	48.10	B	N
ATOM	2428	CA	ALA	397	46.967	-2.743	4.797	1.00	49.04	B	C
ATOM	2429	CB	ALA	397	46.744	-1.374	4.195	1.00	47.71	B	C
ATOM	2430	C	ALA	397	48.454	-3.004	4.968	1.00	50.53	B	C

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ATOM   2431  O   ALA   397      49.049  -3.730   4.174  1.00 51.84      B    O
ATOM   2432  N   MET   398      49.045  -2.403   6.000  1.00 51.67      B    N
ATOM   2433  CA  MET   398      50.423  -2.682   6.388  1.00 53.66      B    C
ATOM   2434  CB  MET   398      50.750  -1.959   7.687  1.00 54.03      B    C
ATOM   2435  CG  MET   398      50.318  -0.514   7.716  1.00 56.66      B    C
ATOM   2436  SD  MET   398      51.552   0.591   7.036  1.00 60.87      B    S
ATOM   2437  CE  MET   398      51.735  -0.099   5.342  1.00 61.35      B    C
ATOM   2438  C   MET   398      50.607  -4.182   6.601  1.00 55.46      B    C
ATOM   2439  O   MET   398      51.606  -4.775   6.183  1.00 56.38      B    O
ATOM   2440  N   MET   399      49.624  -4.795   7.243  1.00 55.72      B    N
ATOM   2441  CA  MET   399      49.743  -6.179   7.642  1.00 56.05      B    C
ATOM   2442  CB  MET   399      48.808  -6.448   8.822  1.00 52.69      B    C
ATOM   2443  CG  MET   399      49.220  -5.660  10.083  1.00 49.75      B    C
ATOM   2444  SD  MET   399      48.186  -5.968  11.517  1.00 43.73      B    S
ATOM   2445  CE  MET   399      47.122  -7.288  10.807  1.00 41.37      B    C
ATOM   2446  C   MET   399      49.471  -7.141   6.496  1.00 58.09      B    C
ATOM   2447  O   MET   399      50.184  -8.123   6.328  1.00 60.78      B    O
ATOM   2448  N   LEU   400      48.444  -6.866   5.705  1.00 60.18      B    N
ATOM   2449  CA  LEU   400      48.212  -7.651   4.503  1.00 61.96      B    C
ATOM   2450  CB  LEU   400      46.845  -7.325   3.905  1.00 60.96      B    C
ATOM   2451  CG  LEU   400      45.716  -8.301   4.218  1.00 59.67      B    C
ATOM   2452  CD1 LEU   400      44.429  -7.814   3.591  1.00 59.61      B    C
ATOM   2453  CD2 LEU   400      46.061  -9.670   3.677  1.00 59.08      B    C
ATOM   2454  C   LEU   400      49.304  -7.349   3.483  1.00 65.48      B    C
ATOM   2455  O   LEU   400      49.422  -8.034   2.473  1.00 65.74      B    O
ATOM   2456  N   SER   401      50.103  -6.320   3.743  1.00 69.84      B    N
ATOM   2457  CA  SER   401      51.162  -5.956   2.811  1.00 75.12      B    C
ATOM   2458  CB  SER   401      51.500  -4.468   2.937  1.00 76.14      B    C
ATOM   2459  OG  SER   401      52.416  -4.065   1.931  1.00 82.39      B    O
ATOM   2460  C   SER   401      52.398  -6.791   3.112  1.00 77.81      B    C
ATOM   2461  O   SER   401      53.097  -7.239   2.197  1.00 77.60      B    O
ATOM   2462  N   ALA   402      52.659  -6.987   4.405  1.00 80.22      B    N
ATOM   2463  CA  ALA   402      53.724  -7.872   4.858  1.00 81.56      B    C
ATOM   2464  CB  ALA   402      53.906  -7.733   6.365  1.00 81.70      B    C
ATOM   2465  C   ALA   402      53.383  -9.318   4.492  1.00 83.56      B    C
ATOM   2466  O   ALA   402      54.068  -9.938   3.687  1.00 83.97      B    O
ATOM   2467  N   GLU   403      52.312  -9.846   5.071  1.00 86.39      B    N
ATOM   2468  CA  GLU   403      51.893 -11.207   4.781  1.00 90.62      B    C
ATOM   2469  CB  GLU   403      51.516 -11.935   6.072  1.00 94.08      B    C
ATOM   2470  CG  GLU   403      52.532 -11.811   7.195  1.00101.30      B    C
ATOM   2471  CD  GLU   403      52.184 -10.705   8.179  1.00105.07      B    C
ATOM   2472  OE1 GLU   403      51.312 -10.929   9.047  1.00107.17      B    O
ATOM   2473  OE2 GLU   403      52.782  -9.610   8.087  1.00107.51      B    O
ATOM   2474  C   GLU   403      50.694 -11.212   3.849  1.00 89.62      B    C
ATOM   2475  O   GLU   403      49.560 -11.324   4.298  1.00 89.74      B    O
ATOM   2476  N   PRO   404      50.930 -11.117   2.535  1.00 87.70      B    N
ATOM   2477  CD  PRO   404      52.238 -11.106   1.863  1.00 82.55      B    C
ATOM   2478  CA  PRO   404      49.823 -11.086   1.574  1.00 86.92      B    C
ATOM   2479  CB  PRO   404      50.514 -11.236   0.221  1.00 82.11      B    C
ATOM   2480  CG  PRO   404      51.883 -10.718   0.453  1.00 81.28      B    C
ATOM   2481  C   PRO   404      48.826 -12.201   1.817  1.00 86.62      B    C
ATOM   2482  O   PRO   404      47.623 -12.004   1.699  1.00 89.69      B    O
ATOM   2483  N   GLU   405      49.332 -13.376   2.170  1.00 84.71      B    N
ATOM   2484  CA  GLU   405      48.497 -14.571   2.216  1.00 82.30      B    C
ATOM   2485  CB  GLU   405      49.339 -15.811   1.885  1.00 83.56      B    C
ATOM   2486  CG  GLU   405      49.728 -15.903   0.415  1.00 84.66      B    C
ATOM   2487  CD  GLU   405      50.509 -17.162   0.076  1.00 86.54      B    C
ATOM   2488  OE1 GLU   405      51.337 -17.113  -0.862  1.00 86.91      B    O
ATOM   2489  OE2 GLU   405      50.294 -18.200   0.742  1.00 88.12      B    O
ATOM   2490  C   GLU   405      47.772 -14.756   3.549  1.00 80.19      B    C
ATOM   2491  O   GLU   405      47.171 -15.803   3.800  1.00 79.93      B    O
ATOM   2492  N   LEU   406      47.818 -13.740   4.402  1.00 78.24      B    N
ATOM   2493  CA  LEU   406      47.071 -13.795   5.648  1.00 76.72      B    C
ATOM   2494  CB  LEU   406      47.211 -12.484   6.410  1.00 74.80      B    C
ATOM   2495  CG  LEU   406      48.424 -12.421   7.327  1.00 74.00      B    C
ATOM   2496  CD1 LEU   406      48.719 -10.994   7.740  1.00 73.34      B    C
ATOM   2497  CD2 LEU   406      48.144 -13.286   8.529  1.00 74.12      B    C
ATOM   2498  C   LEU   406      45.605 -14.060   5.365  1.00 76.33      B    C
ATOM   2499  O   LEU   406      45.077 -13.652   4.330  1.00 77.41      B    O
ATOM   2500  N   THR   407      44.948 -14.748   6.291  1.00 75.06      B    N
ATOM   2501  CA  THR   407      43.540 -15.085   6.126  1.00 73.55      B    C
ATOM   2502  CB  THR   407      43.229 -16.503   6.668  1.00 72.50      B    C
ATOM   2503  OG1 THR   407      43.044 -16.453   8.088  1.00 70.12      B    O
ATOM   2504  CG2 THR   407      44.382 -17.435   6.372  1.00 70.92      B    C
ATOM   2505  C   THR   407      42.660 -14.080   6.866  1.00 73.16      B    C
ATOM   2506  O   THR   407      43.125 -13.364   7.762  1.00 71.83      B    O
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ATOM   2507  N   LEU   408      41.390 -14.039   6.472  1.00 72.79      B    N
ATOM   2508  CA  LEU   408      40.370 -13.276   7.176  1.00 71.72      B    C
ATOM   2509  CB  LEU   408      38.987 -13.813   6.797  1.00 70.47      B    C
ATOM   2510  CG  LEU   408      37.747 -12.922   6.920  1.00 69.40      B    C
ATOM   2511  CD1 LEU   408      37.916 -11.902   8.053  1.00 67.99      B    C
ATOM   2512  CD2 LEU   408      37.528 -12.232   5.584  1.00 69.52      B    C
ATOM   2513  C   LEU   408      40.589 -13.449   8.677  1.00 71.42      B    C
ATOM   2514  O   LEU   408      40.722 -12.474   9.419  1.00 71.69      B    O
ATOM   2515  N   ALA   409      40.640 -14.707   9.107  1.00 70.70      B    N
ATOM   2516  CA  ALA   409      40.632 -15.048  10.525  1.00 69.24      B    C
ATOM   2517  CB  ALA   409      40.137 -16.485  10.701  1.00 68.18      B    C
ATOM   2518  C   ALA   409      41.997 -14.867  11.192  1.00 68.50      B    C
ATOM   2519  O   ALA   409      42.080 -14.711  12.409  1.00 67.64      B    O
ATOM   2520  N   GLU   410      43.060 -14.882  10.394  1.00 68.72      B    N
ATOM   2521  CA  GLU   410      44.407 -14.683  10.918  1.00 69.75      B    C
ATOM   2522  CB  GLU   410      45.442 -15.290   9.970  1.00 69.56      B    C
ATOM   2523  CG  GLU   410      45.349 -16.800   9.836  1.00 70.17      B    C
ATOM   2524  CD  GLU   410      46.369 -17.356   8.868  1.00 70.08      B    C
ATOM   2525  OE1 GLU   410      46.232 -18.532   8.455  1.00 71.00      B    O
ATOM   2526  OE2 GLU   410      47.309 -16.610   8.521  1.00 70.53      B    O
ATOM   2527  C   GLU   410      44.734 -13.209  11.132  1.00 70.53      B    C
ATOM   2528  O   GLU   410      45.324 -12.838  12.151  1.00 70.22      B    O
ATOM   2529  N   LEU   411      44.362 -12.371  10.168  1.00 70.72      B    N
ATOM   2530  CA  LEU   411      44.614 -10.941  10.292  1.00 70.59      B    C
ATOM   2531  CB  LEU   411      44.163 -10.203   9.028  1.00 69.74      B    C
ATOM   2532  CG  LEU   411      44.300  -8.676   8.969  1.00 67.81      B    C
ATOM   2533  CD1 LEU   411      44.390  -8.235   7.528  1.00 66.98      B    C
ATOM   2534  CD2 LEU   411      43.113  -8.014   9.634  1.00 67.36      B    C
ATOM   2535  C   LEU   411      43.821 -10.460  11.484  1.00 71.74      B    C
ATOM   2536  O   LEU   411      44.301  -9.662  12.289  1.00 71.67      B    O
ATOM   2537  N   ARG   412      42.603 -10.970  11.597  1.00 73.60      B    N
ATOM   2538  CA  ARG   412      41.745 -10.648  12.719  1.00 76.27      B    C
ATOM   2539  CB  ARG   412      40.459 -11.485  12.617  1.00 73.69      B    C
ATOM   2540  CG  ARG   412      39.320 -10.901  13.422  1.00 70.65      B    C
ATOM   2541  CD  ARG   412      38.098 -11.772  13.268  1.00 67.93      B    C
ATOM   2542  NE  ARG   412      36.875 -10.988  13.159  1.00 66.79      B    N
ATOM   2543  CZ  ARG   412      36.239 -10.764  12.017  1.00 66.69      B    C
ATOM   2544  NH1 ARG   412      35.130 -10.042  12.007  1.00 65.72      B    N
ATOM   2545  NH2 ARG   412      36.717 -11.264  10.885  1.00 68.01      B    N
ATOM   2546  C   ARG   412      42.488 -11.005  13.997  1.00 80.83      B    C
ATOM   2547  O   ARG   412      42.543 -10.216  14.930  1.00 80.65      B    O
ATOM   2548  N   GLN   413      43.077 -12.196  14.018  1.00 86.79      B    N
ATOM   2549  CA  GLN   413      43.835 -12.665  15.170  1.00 92.24      B    C
ATOM   2550  CB  GLN   413      44.276 -14.114  14.951  1.00 94.97      B    C
ATOM   2551  CG  GLN   413      44.915 -14.766  16.170  1.00 99.47      B    C
ATOM   2552  CD  GLN   413      44.028 -14.675  17.399  1.00102.34      B    C
ATOM   2553  OE1 GLN   413      44.204 -13.792  18.245  1.00103.95      B    O
ATOM   2554  NE2 GLN   413      43.062 -15.588  17.501  1.00104.17      B    N
ATOM   2555  C   GLN   413      45.062 -11.792  15.410  1.00 94.56      B    C
ATOM   2556  O   GLN   413      45.477 -11.580  16.550  1.00 95.17      B    O
ATOM   2557  N   ARG   414      45.638 -11.288  14.324  1.00 97.12      B    N
ATOM   2558  CA  ARG   414      46.857 -10.498  14.405  1.00 98.76      B    C
ATOM   2559  CB  ARG   414      47.512 -10.401  13.029  1.00101.11      B    C
ATOM   2560  CG  ARG   414      48.259 -11.653  12.624  1.00104.98      B    C
ATOM   2561  CD  ARG   414      49.105 -11.403  11.395  1.00109.69      B    C
ATOM   2562  NE  ARG   414      50.249 -12.307  11.305  1.00114.56      B    N
ATOM   2563  CZ  ARG   414      50.158 -13.631  11.227  1.00118.01      B    C
ATOM   2564  NH1 ARG   414      48.968 -14.220  11.233  1.00120.46      B    N
ATOM   2565  NH2 ARG   414      51.258 -14.367  11.131  1.00120.17      B    N
ATOM   2566  C   ARG   414      46.611  -9.102  14.950  1.00 98.39      B    C
ATOM   2567  O   ARG   414      47.365  -8.619  15.794  1.00 98.42      B    O
ATOM   2568  N   LEU   415      45.556  -8.457  14.465  1.00 98.39      B    N
ATOM   2569  CA  LEU   415      45.194  -7.126  14.934  1.00 99.38      B    C
ATOM   2570  CB  LEU   415      43.926  -6.639  14.239  1.00 96.98      B    C
ATOM   2571  CG  LEU   415      44.046  -6.188  12.784  1.00 94.64      B    C
ATOM   2572  CD1 LEU   415      42.667  -5.991  12.184  1.00 93.57      B    C
ATOM   2573  CD2 LEU   415      44.838  -4.905  12.725  1.00 93.89      B    C
ATOM   2574  C   LEU   415      44.964  -7.142  16.430  1.00102.00      B    C
ATOM   2575  O   LEU   415      45.363  -6.224  17.130  1.00100.71      B    O
ATOM   2576  N   ILE   416      44.325  -8.200  16.916  1.00106.93      B    N
ATOM   2577  CA  ILE   416      43.995  -8.306  18.332  1.00111.24      B    C
ATOM   2578  CB  ILE   416      43.220  -9.593  18.642  1.00109.49      B    C
ATOM   2579  CG2 ILE   416      42.848  -9.620  20.110  1.00107.39      B    C
ATOM   2580  CG1 ILE   416      41.958  -9.668  17.789  1.00109.48      B    C
ATOM   2581  CD1 ILE   416      41.160 -10.926  18.002  1.00110.78      B    C
ATOM   2582  C   ILE   416      45.253  -8.319  19.179  1.00114.47      B    C
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ATOM   2583  O   ILE   416      45.367  -7.568  20.146  1.00117.39      B    O
ATOM   2584  N   HIS   417      46.195  -9.181  18.812  1.00116.77      B    N
ATOM   2585  CA  HIS   417      47.420  -9.315  19.578  1.00117.72      B    C
ATOM   2586  CB  HIS   417      48.350 -10.343  18.929  1.00124.49      B    C
ATOM   2587  CG  HIS   417      49.677 -10.471  19.611  1.00130.81      B    C
ATOM   2588  CD2 HIS   417      50.035 -11.091  20.760  1.00133.74      B    C
ATOM   2589  ND1 HIS   417      50.829  -9.905  19.107  1.00133.21      B    N
ATOM   2590  CE1 HIS   417      51.839 -10.170  19.916  1.00135.24      B    C
ATOM   2591  NE2 HIS   417      51.384 -10.888  20.928  1.00135.71      B    N
ATOM   2592  C   HIS   417      48.125  -7.976  19.669  1.00115.09      B    C
ATOM   2593  O   HIS   417      48.408  -7.491  20.761  1.00113.36      B    O
ATOM   2594  N   PHE   418      48.394  -7.371  18.517  1.00112.88      B    N
ATOM   2595  CA  PHE   418      49.206  -6.162  18.469  1.00110.87      B    C
ATOM   2596  CB  PHE   418      49.750  -5.943  17.059  1.00115.25      B    C
ATOM   2597  CG  PHE   418      51.070  -6.611  16.814  1.00121.12      B    C
ATOM   2598  CD1 PHE   418      51.145  -7.792  16.098  1.00123.17      B    C
ATOM   2599  CD2 PHE   418      52.238  -6.059  17.312  1.00123.35      B    C
ATOM   2600  CE1 PHE   418      52.359  -8.408  15.884  1.00125.37      B    C
ATOM   2601  CE2 PHE   418      53.456  -6.672  17.101  1.00125.90      B    C
ATOM   2602  CZ  PHE   418      53.514  -7.852  16.383  1.00126.36      B    C
ATOM   2603  C   PHE   418      48.472  -4.913  18.927  1.00106.04      B    C
ATOM   2604  O   PHE   418      49.033  -3.820  18.896  1.00106.28      B    O
ATOM   2605  N   SER   419      47.224  -5.084  19.353  1.00 99.46      B    N
ATOM   2606  CA  SER   419      46.413  -3.987  19.875  1.00 92.29      B    C
ATOM   2607  CB  SER   419      44.937  -4.359  19.839  1.00 90.97      B    C
ATOM   2608  OG  SER   419      44.506  -4.607  18.522  1.00 89.32      B    O
ATOM   2609  C   SER   419      46.781  -3.652  21.311  1.00 88.83      B    C
ATOM   2610  O   SER   419      46.976  -4.547  22.124  1.00 89.15      B    O
ATOM   2611  N   ALA   420      46.858  -2.363  21.627  1.00 85.37      B    N
ATOM   2612  CA  ALA   420      47.016  -1.927  23.010  1.00 81.97      B    C
ATOM   2613  CB  ALA   420      47.044  -0.406  23.082  1.00 81.91      B    C
ATOM   2614  C   ALA   420      45.842  -2.469  23.816  1.00 79.89      B    C
ATOM   2615  O   ALA   420      44.713  -2.475  23.330  1.00 79.29      B    O
ATOM   2616  N   LYS   421      46.100  -2.926  25.039  1.00 77.60      B    N
ATOM   2617  CA  LYS   421      45.101  -3.706  25.762  1.00 76.65      B    C
ATOM   2618  CB  LYS   421      45.629  -5.120  26.005  1.00 75.21      B    C
ATOM   2619  CG  LYS   421      46.034  -5.842  24.730  1.00 74.25      B    C
ATOM   2620  CD  LYS   421      45.404  -7.221  24.634  1.00 72.30      B    C
ATOM   2621  CE  LYS   421      45.912  -7.973  23.414  1.00 71.65      B    C
ATOM   2622  NZ  LYS   421      47.403  -7.974  23.350  1.00 69.08      B    N
ATOM   2623  C   LYS   421      44.606  -3.108  27.080  1.00 76.30      B    C
ATOM   2624  O   LYS   421      45.388  -2.590  27.874  1.00 75.81      B    O
ATOM   2625  N   ASP   422      43.292  -3.193  27.291  1.00 76.07      B    N
ATOM   2626  CA  ASP   422      42.654  -2.840  28.557  1.00 75.90      B    C
ATOM   2627  CB  ASP   422      43.025  -3.859  29.635  1.00 76.51      B    C
ATOM   2628  CG  ASP   422      42.547  -5.259  29.293  1.00 77.29      B    C
ATOM   2629  OD1 ASP   422      41.496  -5.684  29.823  1.00 77.24      B    O
ATOM   2630  OD2 ASP   422      43.218  -5.936  28.485  1.00 78.52      B    O
ATOM   2631  C   ASP   422      42.978  -1.434  29.032  1.00 75.72      B    C
ATOM   2632  O   ASP   422      43.111  -1.185  30.231  1.00 75.77      B    O
ATOM   2633  N   VAL   423      43.094  -0.516  28.078  1.00 75.77      B    N
ATOM   2634  CA  VAL   423      43.330   0.893  28.373  1.00 75.46      B    C
ATOM   2635  CB  VAL   423      44.279   1.529  27.323  1.00 74.23      B    C
ATOM   2636  CG1 VAL   423      45.572   0.739  27.256  1.00 73.29      B    C
ATOM   2637  CG2 VAL   423      43.616   1.555  25.953  1.00 72.59      B    C
ATOM   2638  C   VAL   423      42.008   1.666  28.385  1.00 75.67      B    C
ATOM   2639  O   VAL   423      41.911   2.734  28.997  1.00 75.74      B    O
ATOM   2640  N   ILE   424      40.999   1.118  27.706  1.00 74.56      B    N
ATOM   2641  CA  ILE   424      39.674   1.733  27.641  1.00 74.42      B    C
ATOM   2642  CB  ILE   424      38.810   1.097  26.512  1.00 73.36      B    C
ATOM   2643  CG2 ILE   424      37.506   1.865  26.345  1.00 74.09      B    C
ATOM   2644  CG1 ILE   424      39.558   1.125  25.183  1.00 71.55      B    C
ATOM   2645  CD1 ILE   424      38.737   0.586  24.031  1.00 68.53      B    C
ATOM   2646  C   ILE   424      38.934   1.526  28.969  1.00 74.42      B    C
ATOM   2647  O   ILE   424      38.736   0.384  29.397  1.00 74.56      B    O
ATOM   2648  N   ASN   425      38.516   2.612  29.618  1.00 73.22      B    N
ATOM   2649  CA  ASN   425      37.707   2.466  30.822  1.00 73.77      B    C
ATOM   2650  CB  ASN   425      37.545   3.803  31.550  1.00 72.83      B    C
ATOM   2651  CG  ASN   425      36.794   3.659  32.863  1.00 71.81      B    C
ATOM   2652  OD1 ASN   425      35.814   2.925  32.949  1.00 72.71      B    O
ATOM   2653  ND2 ASN   425      37.258   4.350  33.891  1.00 71.45      B    N
ATOM   2654  C   ASN   425      36.338   1.944  30.426  1.00 74.65      B    C
ATOM   2655  O   ASN   425      35.618   2.603  29.686  1.00 74.07      B    O
ATOM   2656  N   GLU   426      35.976   0.766  30.924  1.00 76.81      B    N
ATOM   2657  CA  GLU   426      34.741   0.111  30.501  1.00 79.15      B    C
ATOM   2658  CB  GLU   426      34.712  -1.337  30.986  1.00 80.99      B    C
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ATOM   2659  CG  GLU   426      35.579  -2.282  30.184  1.00 83.84      B    C
ATOM   2660  CD  GLU   426      35.147  -3.729  30.345  1.00 86.17      B    C
ATOM   2661  OE1 GLU   426      35.400  -4.529  29.418  1.00 87.57      B    O
ATOM   2662  OE2 GLU   426      34.551  -4.063  31.395  1.00 85.46      B    O
ATOM   2663  C   GLU   426      33.480   0.824  30.984  1.00 79.50      B    C
ATOM   2664  O   GLU   426      32.386   0.569  30.474  1.00 80.47      B    O
ATOM   2665  N   ALA   427      33.637   1.722  31.956  1.00 78.27      B    N
ATOM   2666  CA  ALA   427      32.501   2.376  32.599  1.00 76.26      B    C
ATOM   2667  CB  ALA   427      32.936   2.969  33.924  1.00 75.03      B    C
ATOM   2668  C   ALA   427      31.835   3.452  31.740  1.00 75.60      B    C
ATOM   2669  O   ALA   427      30.987   4.205  32.224  1.00 75.77      B    O
ATOM   2670  N   TRP   428      32.226   3.530  30.473  1.00 74.75      B    N
ATOM   2671  CA  TRP   428      31.555   4.397  29.512  1.00 74.44      B    C
ATOM   2672  CB  TRP   428      32.542   4.811  28.423  1.00 74.02      B    C
ATOM   2673  CG  TRP   428      32.114   5.942  27.534  1.00 73.99      B    C
ATOM   2674  CD2 TRP   428      31.476   5.843  26.247  1.00 74.18      B    C
ATOM   2675  CE2 TRP   428      31.407   7.143  25.715  1.00 73.61      B    C
ATOM   2676  CE3 TRP   428      30.968   4.781  25.492  1.00 74.77      B    C
ATOM   2677  CD1 TRP   428      32.379   7.258  27.723  1.00 73.84      B    C
ATOM   2678  NE1 TRP   428      31.963   7.988  26.637  1.00 74.39      B    N
ATOM   2679  CZ2 TRP   428      30.853   7.414  24.462  1.00 72.79      B    C
ATOM   2680  CZ3 TRP   428      30.417   5.055  24.245  1.00 74.68      B    C
ATOM   2681  CH2 TRP   428      30.367   6.362  23.746  1.00 73.12      B    C
ATOM   2682  C   TRP   428      30.424   3.579  28.905  1.00 74.60      B    C
ATOM   2683  O   TRP   428      29.370   4.106  28.555  1.00 73.81      B    O
ATOM   2684  N   PHE   429      30.656   2.276  28.799  1.00 75.15      B    N
ATOM   2685  CA  PHE   429      29.715   1.370  28.146  1.00 76.08      B    C
ATOM   2686  CB  PHE   429      30.447   0.117  27.654  1.00 75.20      B    C
ATOM   2687  CG  PHE   429      31.632   0.409  26.773  1.00 73.39      B    C
ATOM   2688  CD1 PHE   429      32.897  -0.041  27.119  1.00 72.50      B    C
ATOM   2689  CD2 PHE   429      31.478   1.127  25.594  1.00 72.64      B    C
ATOM   2690  CE1 PHE   429      33.989   0.214  26.304  1.00 71.63      B    C
ATOM   2691  CE2 PHE   429      32.562   1.386  24.775  1.00 72.20      B    C
ATOM   2692  CZ  PHE   429      33.822   0.928  25.131  1.00 71.45      B    C
ATOM   2693  C   PHE   429      28.588   0.958  29.083  1.00 76.15      B    C
ATOM   2694  O   PHE   429      28.820   0.627  30.242  1.00 76.42      B    O
ATOM   2695  N   PRO   430      27.346   0.961  28.587  1.00 76.64      B    N
ATOM   2696  CD  PRO   430      26.911   1.204  27.203  1.00 76.64      B    C
ATOM   2697  CA  PRO   430      26.237   0.571  29.458  1.00 78.80      B    C
ATOM   2698  CB  PRO   430      25.006   0.663  28.549  1.00 77.13      B    C
ATOM   2699  CG  PRO   430      25.536   0.602  27.168  1.00 76.00      B    C
ATOM   2700  C   PRO   430      26.445  -0.825  30.039  1.00 80.99      B    C
ATOM   2701  O   PRO   430      26.865  -1.747  29.341  1.00 81.23      B    O
ATOM   2702  N   GLU   431      26.140  -0.961  31.325  1.00 83.65      B    N
ATOM   2703  CA  GLU   431      26.550  -2.107  32.130  1.00 85.46      B    C
ATOM   2704  CB  GLU   431      25.639  -2.199  33.363  1.00 89.78      B    C
ATOM   2705  CG  GLU   431      26.071  -3.227  34.405  1.00 95.94      B    C
ATOM   2706  CD  GLU   431      25.317  -3.091  35.721  1.00 98.87      B    C
ATOM   2707  OE1 GLU   431      25.879  -3.512  36.758  1.00100.92      B    O
ATOM   2708  OE2 GLU   431      24.174  -2.570  35.717  1.00100.66      B    O
ATOM   2709  C   GLU   431      26.569  -3.446  31.388  1.00 83.87      B    C
ATOM   2710  O   GLU   431      27.593  -4.133  31.337  1.00 83.49      B    O
ATOM   2711  N   ASP   432      25.429  -3.799  30.808  1.00 81.63      B    N
ATOM   2712  CA  ASP   432      25.220  -5.121  30.232  1.00 79.14      B    C
ATOM   2713  CB  ASP   432      23.722  -5.335  29.977  1.00 79.86      B    C
ATOM   2714  CG  ASP   432      22.998  -4.047  29.596  1.00 79.96      B    C
ATOM   2715  OD1 ASP   432      23.092  -3.046  30.346  1.00 79.78      B    O
ATOM   2716  OD2 ASP   432      22.329  -4.039  28.542  1.00 79.18      B    O
ATOM   2717  C   ASP   432      26.009  -5.339  28.944  1.00 77.30      B    C
ATOM   2718  O   ASP   432      26.070  -6.451  28.413  1.00 76.75      B    O
ATOM   2719  N   GLN   433      26.612  -4.268  28.445  1.00 75.05      B    N
ATOM   2720  CA  GLN   433      27.378  -4.327  27.213  1.00 72.42      B    C
ATOM   2721  CB  GLN   433      27.122  -3.074  26.375  1.00 71.49      B    C
ATOM   2722  CG  GLN   433      25.835  -3.110  25.591  1.00 70.56      B    C
ATOM   2723  CD  GLN   433      25.896  -4.114  24.468  1.00 71.25      B    C
ATOM   2724  OE1 GLN   433      24.888  -4.442  23.845  1.00 71.67      B    O
ATOM   2725  NE2 GLN   433      27.091  -4.615  24.204  1.00 72.06      B    N
ATOM   2726  C   GLN   433      28.867  -4.461  27.484  1.00 71.41      B    C
ATOM   2727  O   GLN   433      29.608  -4.934  26.627  1.00 71.95      B    O
ATOM   2728  N   ARG   434      29.296  -4.047  28.675  1.00 69.73      B    N
ATOM   2729  CA  ARG   434      30.717  -3.900  28.985  1.00 68.37      B    C
ATOM   2730  CB  ARG   434      30.900  -3.423  30.424  1.00 68.00      B    C
ATOM   2731  CG  ARG   434      30.438  -2.003  30.678  1.00 68.67      B    C
ATOM   2732  CD  ARG   434      31.044  -1.449  31.956  1.00 69.05      B    C
ATOM   2733  NE  ARG   434      30.069  -0.678  32.713  1.00 70.49      B    N
ATOM   2734  CZ  ARG   434      29.112  -1.228  33.449  1.00 72.47      B    C
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ATOM   2735  NH1 ARG   434      28.255  -0.462  34.113  1.00 73.59      B    N
ATOM   2736  NH2 ARG   434      29.018  -2.551  33.521  1.00 73.57      B    N
ATOM   2737  C   ARG   434      31.510  -5.184  28.795  1.00 68.33      B    C
ATOM   2738  O   ARG   434      32.589  -5.180  28.190  1.00 68.56      B    O
ATOM   2739  N   VAL   435      30.981  -6.284  29.319  1.00 67.14      B    N
ATOM   2740  CA  VAL   435      31.681  -7.555  29.233  1.00 66.36      B    C
ATOM   2741  CB  VAL   435      30.884  -8.689  29.945  1.00 66.62      B    C
ATOM   2742  CG1 VAL   435      30.754  -8.367  31.431  1.00 64.88      B    C
ATOM   2743  CG2 VAL   435      29.498  -8.857  29.307  1.00 64.17      B    C
ATOM   2744  C   VAL   435      31.934  -7.936  27.776  1.00 65.17      B    C
ATOM   2745  O   VAL   435      32.990  -8.475  27.442  1.00 66.15      B    O
ATOM   2746  N   LEU   436      30.971  -7.635  26.910  1.00 63.04      B    N
ATOM   2747  CA  LEU   436      31.051  -8.010  25.497  1.00 61.71      B    C
ATOM   2748  CB  LEU   436      29.648  -7.993  24.871  1.00 62.24      B    C
ATOM   2749  CG  LEU   436      28.556  -8.820  25.561  1.00 61.83      B    C
ATOM   2750  CD1 LEU   436      27.225  -8.631  24.823  1.00 60.35      B    C
ATOM   2751  CD2 LEU   436      28.970 -10.294  25.593  1.00 60.07      B    C
ATOM   2752  C   LEU   436      31.976  -7.099  24.675  1.00 59.12      B    C
ATOM   2753  O   LEU   436      32.655  -7.554  23.759  1.00 57.04      B    O
ATOM   2754  N   THR   437      31.988  -5.812  25.000  1.00 57.08      B    N
ATOM   2755  CA  THR   437      32.728  -4.844  24.207  1.00 56.21      B    C
ATOM   2756  CB  THR   437      32.403  -3.415  24.648  1.00 53.88      B    C
ATOM   2757  OG1 THR   437      31.009  -3.163  24.442  1.00 55.23      B    O
ATOM   2758  CG2 THR   437      33.216  -2.426  23.858  1.00 50.77      B    C
ATOM   2759  C   THR   437      34.229  -5.064  24.338  1.00 55.95      B    C
ATOM   2760  O   THR   437      34.822  -4.747  25.374  1.00 56.67      B    O
ATOM   2761  N   PRO   438      34.862  -5.596  23.274  1.00 54.46      B    N
ATOM   2762  CD  PRO   438      34.210  -5.902  21.990  1.00 54.63      B    C
ATOM   2763  CA  PRO   438      36.283  -5.949  23.244  1.00 53.85      B    C
ATOM   2764  CB  PRO   438      36.515  -6.417  21.807  1.00 52.36      B    C
ATOM   2765  CG  PRO   438      35.187  -6.851  21.329  1.00 52.88      B    C
ATOM   2766  C   PRO   438      37.156  -4.764  23.581  1.00 54.24      B    C
ATOM   2767  O   PRO   438      37.476  -3.952  22.721  1.00 55.50      B    O
ATOM   2768  N   ASN   439      37.553  -4.658  24.835  1.00 54.28      B    N
ATOM   2769  CA  ASN   439      38.460  -3.600  25.203  1.00 54.38      B    C
ATOM   2770  CB  ASN   439      38.716  -3.662  26.700  1.00 53.17      B    C
ATOM   2771  CG  ASN   439      39.413  -2.438  27.206  1.00 52.95      B    C
ATOM   2772  OD1 ASN   439      40.452  -2.054  26.674  1.00 54.56      B    O
ATOM   2773  ND2 ASN   439      38.846  -1.804  28.233  1.00 52.08      B    N
ATOM   2774  C   ASN   439      39.769  -3.758  24.417  1.00 55.78      B    C
ATOM   2775  O   ASN   439      40.655  -4.519  24.814  1.00 57.73      B    O
ATOM   2776  N   LEU   440      39.877  -3.043  23.295  1.00 56.06      B    N
ATOM   2777  CA  LEU   440      41.041  -3.142  22.397  1.00 55.86      B    C
ATOM   2778  CB  LEU   440      40.833  -4.263  21.379  1.00 55.14      B    C
ATOM   2779  CG  LEU   440      41.127  -5.704  21.783  1.00 54.08      B    C
ATOM   2780  CD1 LEU   440      40.647  -6.666  20.684  1.00 51.94      B    C
ATOM   2781  CD2 LEU   440      42.614  -5.845  22.028  1.00 53.63      B    C
ATOM   2782  C   LEU   440      41.263  -1.837  21.627  1.00 56.41      B    C
ATOM   2783  O   LEU   440      40.359  -1.354  20.947  1.00 56.93      B    O
ATOM   2784  N   VAL   441      42.461  -1.271  21.717  1.00 56.47      B    N
ATOM   2785  CA  VAL   441      42.772  -0.074  20.944  1.00 56.91      B    C
ATOM   2786  CB  VAL   441      43.250   1.057  21.851  1.00 55.91      B    C
ATOM   2787  CG1 VAL   441      43.613   2.270  21.009  1.00 55.40      B    C
ATOM   2788  CG2 VAL   441      42.174   1.396  22.864  1.00 54.73      B    C
ATOM   2789  C   VAL   441      43.857  -0.357  19.916  1.00 58.09      B    C
ATOM   2790  O   VAL   441      45.021  -0.486  20.272  1.00 58.32      B    O
ATOM   2791  N   ALA   442      43.472  -0.433  18.643  1.00 60.35      B    N
ATOM   2792  CA  ALA   442      44.369  -0.877  17.576  1.00 62.99      B    C
ATOM   2793  CB  ALA   442      43.740  -0.611  16.219  1.00 62.78      B    C
ATOM   2794  C   ALA   442      45.743  -0.232  17.630  1.00 65.68      B    C
ATOM   2795  O   ALA   442      45.908   0.885  18.121  1.00 65.64      B    O
ATOM   2796  N   ALA   443      46.730  -0.952  17.113  1.00 69.73      B    N
ATOM   2797  CA  ALA   443      48.092  -0.449  17.028  1.00 74.87      B    C
ATOM   2798  CB  ALA   443      48.802  -0.639  18.356  1.00 70.53      B    C
ATOM   2799  C   ALA   443      48.838  -1.182  15.923  1.00 79.99      B    C
ATOM   2800  O   ALA   443      48.650  -2.383  15.726  1.00 81.17      B    O
ATOM   2801  N   LEU   444      49.677  -0.454  15.196  1.00 86.92      B    N
ATOM   2802  CA  LEU   444      50.523  -1.065  14.185  1.00 94.56      B    C
ATOM   2803  CB  LEU   444      51.269   0.016  13.404  1.00 91.22      B    C
ATOM   2804  CG  LEU   444      50.583   0.555  12.150  1.00 88.84      B    C
ATOM   2805  CD1 LEU   444      51.176   1.898  11.774  1.00 86.05      B    C
ATOM   2806  CD2 LEU   444      50.739  -0.447  11.017  1.00 85.93      B    C
ATOM   2807  C   LEU   444      51.523  -2.009  14.840  1.00100.27      B    C
ATOM   2808  O   LEU   444      51.990  -1.762  15.955  1.00101.97      B    O
ATOM   2809  N   PRO   445      51.854  -3.115  14.155  1.00102.71      B    N
ATOM   2810  CD  PRO   445      51.177  -3.601  12.942  1.00102.47      B    C
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ATOM   2811  CA  PRO   445      52.880  -4.056  14.611  1.00108.96      B     C
ATOM   2812  CB  PRO   445      52.808  -5.199  13.598  1.00103.83      B     C
ATOM   2813  CG  PRO   445      51.472  -5.072  12.969  1.00101.93      B     C
ATOM   2814  C   PRO   445      54.257  -3.416  14.618  1.00113.37      B     C
ATOM   2815  O   PRO   445      54.698  -2.856  13.611  1.00112.80      B     O
ATOM   2816  N   PRO   446      54.952  -3.485  15.760  1.00116.22      B     N
ATOM   2817  CD  PRO   446      54.384  -3.732  17.095  1.00119.98      B     C
ATOM   2818  CA  PRO   446      56.374  -3.135  15.797  1.00122.39      B     C
ATOM   2819  CB  PRO   446      56.702  -3.083  17.291  1.00122.67      B     C
ATOM   2820  CG  PRO   446      55.580  -3.813  17.975  1.00120.33      B     C
ATOM   2821  C   PRO   446      57.221  -4.149  15.017  1.00125.56      B     C
ATOM   2822  O   PRO   446      58.166  -3.766  14.326  1.00129.91      B     O
ATOM   2823  N   SER   447      56.864  -5.433  15.111  1.00131.30      B     N
ATOM   2824  CA  SER   447      57.519  -6.498  14.335  1.00136.24      B     C
ATOM   2825  CB  SER   447      58.228  -7.485  15.270  1.00141.23      B     C
ATOM   2826  OG  SER   447      57.298  -8.310  15.952  1.00144.06      B     O
ATOM   2827  C   SER   447      56.556  -7.277  13.423  1.00141.04      B     C
ATOM   2828  O   SER   447      55.822  -8.152  13.934  1.00131.68      B     O
ATOM   2829  OXT SER   447      56.535  -7.006  12.201  1.00113.76      B     O
TER    2830      SER   447                                              B
ATOM   2831  CB  GLU     1       3.711  32.350  16.970  1.00138.52      L31H  C
ATOM   2832  CG  GLU     1       3.439  33.848  16.919  1.00141.85      L31H  C
ATOM   2833  CD  GLU     1       4.640  34.640  16.438  1.00143.45      L31H  C
ATOM   2834  OE1 GLU     1       4.445  35.731  15.860  1.00143.83      L31H  O
ATOM   2835  OE2 GLU     1       5.780  34.170  16.635  1.00143.91      L31H  O
ATOM   2836  C   GLU     1       4.715  30.381  18.137  1.00130.94      L31H  C
ATOM   2837  O   GLU     1       4.120  29.647  17.346  1.00131.43      L31H  O
ATOM   2838  N   GLU     1       5.502  32.678  18.664  1.00135.71      L31H  N
ATOM   2839  CA  GLU     1       4.317  31.849  18.288  1.00134.83      L31H  C
ATOM   2840  N   SER     2       5.722  29.967  18.902  1.00125.13      L31H  N
ATOM   2841  CA  SER     2       6.323  28.637  18.781  1.00119.15      L31H  C
ATOM   2842  CB  SER     2       7.056  28.280  20.080  1.00120.98      L31H  C
ATOM   2843  OG  SER     2       6.217  28.431  21.212  1.00124.69      L31H  O
ATOM   2844  C   SER     2       5.320  27.539  18.440  1.00113.88      L31H  C
ATOM   2845  O   SER     2       4.159  27.597  18.849  1.00112.42      L31H  O
ATOM   2846  N   VAL     3       5.771  26.538  17.688  1.00107.83      L31H  N
ATOM   2847  CA  VAL     3       4.917  25.405  17.342  1.00101.42      L31H  C
ATOM   2848  CB  VAL     3       5.242  24.870  15.924  1.00101.94      L31H  C
ATOM   2849  CG1 VAL     3       6.704  24.464  15.839  1.00102.39      L31H  C
ATOM   2850  CG2 VAL     3       4.319  23.704  15.579  1.00102.00      L31H  C
ATOM   2851  C   VAL     3       5.069  24.295  18.375  1.00 95.09      L31H  C
ATOM   2852  O   VAL     3       4.316  23.326  18.386  1.00 95.30      L31H  O
ATOM   2853  N   LEU     4       6.051  24.453  19.252  1.00 87.11      L31H  N
ATOM   2854  CA  LEU     4       6.090  23.680  20.483  1.00 80.33      L31H  C
ATOM   2855  CB  LEU     4       7.521  23.244  20.799  1.00 78.74      L31H  C
ATOM   2856  CG  LEU     4       8.079  22.079  19.989  1.00 75.03      L31H  C
ATOM   2857  CD1 LEU     4       9.349  21.583  20.651  1.00 73.88      L31H  C
ATOM   2858  CD2 LEU     4       7.052  20.973  19.908  1.00 73.52      L31H  C
ATOM   2859  C   LEU     4       5.547  24.506  21.646  1.00 76.26      L31H  C
ATOM   2860  O   LEU     4       6.084  25.570  21.962  1.00 74.35      L31H  O
ATOM   2861  N   THR     5       4.490  24.010  22.288  1.00 72.42      L31H  N
ATOM   2862  CA  THR     5       3.856  24.736  23.388  1.00 67.66      L31H  C
ATOM   2863  CB  THR     5       2.350  24.890  23.143  1.00 65.52      L31H  C
ATOM   2864  OG1 THR     5       1.842  23.672  22.591  1.00 64.87      L31H  O
ATOM   2865  CG2 THR     5       2.075  26.056  22.184  1.00 64.39      L31H  C
ATOM   2866  C   THR     5       4.069  24.115  24.761  1.00 65.49      L31H  C
ATOM   2867  O   THR     5       3.962  22.903  24.936  1.00 65.15      L31H  O
ATOM   2868  N   GLN     6       4.378  24.969  25.730  1.00 64.29      L31H  N
ATOM   2869  CA  GLN     6       4.589  24.555  27.114  1.00 63.73      L31H  C
ATOM   2870  CB  GLN     6       6.057  24.170  27.326  1.00 61.83      L31H  C
ATOM   2871  CG  GLN     6       7.018  25.346  27.379  1.00 59.76      L31H  C
ATOM   2872  CD  GLN     6       8.361  25.017  26.745  1.00 60.41      L31H  C
ATOM   2873  OE1 GLN     6       8.433  24.730  25.554  1.00 61.76      L31H  O
ATOM   2874  NE2 GLN     6       9.430  25.056  27.540  1.00 58.87      L31H  N
ATOM   2875  C   GLN     6       4.202  25.692  28.074  1.00 64.47      L31H  C
ATOM   2876  O   GLN     6       4.228  26.867  27.702  1.00 64.01      L31H  O
ATOM   2877  N   PRO     7       3.861  25.353  29.330  1.00 66.46      L31H  N
ATOM   2878  CD  PRO     7       4.220  24.113  30.038  1.00 67.84      L31H  C
ATOM   2879  CA  PRO     7       3.326  26.353  30.254  1.00 65.35      L31H  C
ATOM   2880  CB  PRO     7       3.116  25.574  31.547  1.00 65.91      L31H  C
ATOM   2881  CG  PRO     7       4.129  24.508  31.494  1.00 67.09      L31H  C
ATOM   2882  C   PRO     7       4.305  27.496  30.442  1.00 64.17      L31H  C
ATOM   2883  O   PRO     7       5.506  27.349  30.222  1.00 65.08      L31H  O
ATOM   2884  N   PRO     8       3.791  28.663  30.834  1.00 63.81      L31H  N
ATOM   2885  CD  PRO     8       2.356  28.982  30.785  1.00 61.18      L31H  C
ATOM   2886  CA  PRO     8       4.604  29.855  31.085  1.00 63.42      L31H  C
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ATOM   2887  CB  PRO     8       3.577  30.996  31.075  1.00 61.24      L31H C
ATOM   2888  CG  PRO     8       2.367  30.419  30.395  1.00 60.71      L31H C
ATOM   2889  C   PRO     8       5.369  29.786  32.411  1.00 63.36      L31H C
ATOM   2890  O   PRO     8       6.443  30.382  32.568  1.00 63.66      L31H O
ATOM   2891  N   SER     9       4.798  29.062  33.365  1.00 63.25      L31H N
ATOM   2892  CA  SER     9       5.182  29.218  34.757  1.00 63.19      L31H C
ATOM   2893  CB  SER     9       4.466  30.446  35.350  1.00 60.99      L31H C
ATOM   2894  OG  SER     9       4.887  30.729  36.678  1.00 57.73      L31H O
ATOM   2895  C   SER     9       4.866  27.961  35.578  1.00 64.48      L31H C
ATOM   2896  O   SER     9       3.784  27.360  35.478  1.00 65.45      L31H O
ATOM   2897  N   VAL    10       5.837  27.556  36.382  1.00 63.89      L31H N
ATOM   2898  CA  VAL    10       5.607  26.517  37.362  1.00 63.42      L31H C
ATOM   2899  CB  VAL    10       6.233  25.170  36.910  1.00 63.39      L31H C
ATOM   2900  CG1 VAL    10       5.360  24.514  35.836  1.00 62.57      L31H C
ATOM   2901  CG2 VAL    10       7.633  25.415  36.349  1.00 63.77      L31H C
ATOM   2902  C   VAL    10       6.224  26.980  38.670  1.00 63.10      L31H C
ATOM   2903  O   VAL    10       6.939  27.989  38.708  1.00 62.36      L31H O
ATOM   2904  N   SER    11       5.917  26.258  39.741  1.00 63.31      L31H N
ATOM   2905  CA  SER    11       6.542  26.485  41.035  1.00 64.07      L31H C
ATOM   2906  CB  SER    11       6.227  27.894  41.545  1.00 65.21      L31H C
ATOM   2907  OG  SER    11       4.873  28.237  41.310  1.00 68.61      L31H O
ATOM   2908  C   SER    11       6.107  25.443  42.059  1.00 63.28      L31H C
ATOM   2909  O   SER    11       4.965  24.979  42.068  1.00 63.10      L31H O
ATOM   2910  N   GLY    12       7.055  25.064  42.903  1.00 62.91      L31H N
ATOM   2911  CA  GLY    12       6.799  24.139  43.992  1.00 62.96      L31H C
ATOM   2912  C   GLY    12       7.831  24.493  45.040  1.00 62.06      L31H C
ATOM   2913  O   GLY    12       8.833  25.129  44.706  1.00 62.73      L31H O
ATOM   2914  N   ALA    13       7.608  24.119  46.295  1.00 59.54      L31H N
ATOM   2915  CA  ALA    13       8.553  24.482  47.341  1.00 57.29      L31H C
ATOM   2916  CB  ALA    13       7.914  24.294  48.692  1.00 54.68      L31H C
ATOM   2917  C   ALA    13       9.824  23.627  47.217  1.00 57.41      L31H C
ATOM   2918  O   ALA    13       9.863  22.648  46.462  1.00 57.04      L31H O
ATOM   2919  N   PRO    14      10.891  24.000  47.938  1.00 56.71      L31H N
ATOM   2920  CD  PRO    14      11.122  25.236  48.702  1.00 56.70      L31H C
ATOM   2921  CA  PRO    14      12.071  23.137  47.937  1.00 57.23      L31H C
ATOM   2922  CB  PRO    14      13.043  23.867  48.860  1.00 57.17      L31H C
ATOM   2923  CG  PRO    14      12.608  25.283  48.806  1.00 55.90      L31H C
ATOM   2924  C   PRO    14      11.699  21.758  48.464  1.00 58.32      L31H C
ATOM   2925  O   PRO    14      10.950  21.646  49.432  1.00 58.21      L31H O
ATOM   2926  N   GLY    15      12.211  20.714  47.817  1.00 60.26      L31H N
ATOM   2927  CA  GLY    15      11.854  19.358  48.195  1.00 62.96      L31H C
ATOM   2928  C   GLY    15      10.772  18.728  47.332  1.00 65.28      L31H C
ATOM   2929  O   GLY    15      10.696  17.500  47.221  1.00 64.05      L31H O
ATOM   2930  N   GLN    16       9.936  19.560  46.710  1.00 68.62      L31H N
ATOM   2931  CA  GLN    16       8.802  19.065  45.926  1.00 71.60      L31H C
ATOM   2932  CB  GLN    16       7.661  20.072  45.957  1.00 75.87      L31H C
ATOM   2933  CG  GLN    16       6.908  20.112  47.252  1.00 82.09      L31H C
ATOM   2934  CD  GLN    16       5.727  21.055  47.177  1.00 87.55      L31H C
ATOM   2935  OE1 GLN    16       4.583  20.625  46.977  1.00 90.03      L31H O
ATOM   2936  NE2 GLN    16       5.994  22.355  47.326  1.00 89.14      L31H N
ATOM   2937  C   GLN    16       9.123  18.735  44.471  1.00 70.08      L31H C
ATOM   2938  O   GLN    16      10.101  19.220  43.902  1.00 69.52      L31H O
ATOM   2939  N   ARG    17       8.278  17.904  43.876  1.00 68.13      L31H N
ATOM   2940  CA  ARG    17       8.411  17.566  42.471  1.00 66.35      L31H C
ATOM   2941  CB  ARG    17       8.093  16.084  42.248  1.00 65.46      L31H C
ATOM   2942  CG  ARG    17       8.223  15.626  40.804  1.00 65.51      L31H C
ATOM   2943  CD  ARG    17       7.961  14.138  40.688  1.00 66.05      L31H C
ATOM   2944  NE  ARG    17       7.644  13.722  39.326  1.00 68.32      L31H N
ATOM   2945  CZ  ARG    17       6.444  13.852  38.764  1.00 71.52      L31H C
ATOM   2946  NH1 ARG    17       5.440  14.397  39.446  1.00 73.00      L31H N
ATOM   2947  NH2 ARG    17       6.239  13.418  37.526  1.00 72.19      L31H N
ATOM   2948  C   ARG    17       7.481  18.434  41.620  1.00 65.81      L31H C
ATOM   2949  O   ARG    17       6.403  18.850  42.068  1.00 64.65      L31H O
ATOM   2950  N   VAL    18       7.930  18.699  40.392  1.00 64.40      L31H N
ATOM   2951  CA  VAL    18       7.210  19.499  39.411  1.00 63.29      L31H C
ATOM   2952  CB  VAL    18       7.816  20.908  39.293  1.00 65.09      L31H C
ATOM   2953  CG1 VAL    18       7.832  21.588  40.648  1.00 65.44      L31H C
ATOM   2954  CG2 VAL    18       9.222  20.811  38.727  1.00 66.78      L31H C
ATOM   2955  C   VAL    18       7.320  18.827  38.044  1.00 62.02      L31H C
ATOM   2956  O   VAL    18       8.181  17.980  37.838  1.00 60.97      L31H O
ATOM   2957  N   THR    19       6.451  19.204  37.112  1.00 61.73      L31H N
ATOM   2958  CA  THR    19       6.657  18.845  35.712  1.00 61.82      L31H C
ATOM   2959  CB  THR    19       5.694  17.739  35.261  1.00 61.35      L31H C
ATOM   2960  OG1 THR    19       4.376  18.279  35.116  1.00 60.99      L31H O
ATOM   2961  CG2 THR    19       5.687  16.610  36.264  1.00 61.60      L31H C
ATOM   2962  C   THR    19       6.465  20.049  34.781  1.00 63.20      L31H C
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ATOM   2963  O   THR  19       5.754  21.002  35.124  1.00 64.11      L31H O
ATOM   2964  N   ILE  20       7.113  20.003  33.613  1.00 62.60      L31H N
ATOM   2965  CA  ILE  20       6.843  20.935  32.512  1.00 62.83      L31H C
ATOM   2966  CB  ILE  20       8.089  21.772  32.198  1.00 63.61      L31H C
ATOM   2967  CG2 ILE  20       7.815  22.690  31.032  1.00 65.17      L31H C
ATOM   2968  CG1 ILE  20       8.487  22.588  33.429  1.00 64.22      L31H C
ATOM   2969  CD1 ILE  20       9.698  23.492  33.213  1.00 64.62      L31H C
ATOM   2970  C   ILE  20       6.432  20.123  31.275  1.00 63.78      L31H C
ATOM   2971  O   ILE  20       6.787  18.945  31.162  1.00 62.92      L31H O
ATOM   2972  N   SER  21       5.682  20.729  30.354  1.00 64.09      L31H N
ATOM   2973  CA  SER  21       4.990  19.935  29.336  1.00 64.86      L31H C
ATOM   2974  CB  SER  21       3.536  19.776  29.754  1.00 63.74      L31H C
ATOM   2975  OG  SER  21       3.435  19.888  31.163  1.00 64.14      L31H O
ATOM   2976  C   SER  21       5.058  20.444  27.887  1.00 65.67      L31H C
ATOM   2977  O   SER  21       4.306  21.337  27.489  1.00 68.38      L31H O
ATOM   2978  N   CYS  22       5.945  19.842  27.099  1.00 64.14      L31H N
ATOM   2979  CA  CYS  22       6.069  20.125  25.671  1.00 63.02      L31H C
ATOM   2980  C   CYS  22       5.038  19.313  24.866  1.00 63.37      L31H C
ATOM   2981  O   CYS  22       5.077  18.080  24.848  1.00 62.28      L31H O
ATOM   2982  CB  CYS  22       7.483  19.755  25.210  1.00 61.52      L31H C
ATOM   2983  SG  CYS  22       8.208  20.677  23.808  1.00 61.05      L31H S
ATOM   2984  N   THR  23       4.117  20.005  24.202  1.00 64.37      L31H N
ATOM   2985  CA  THR  23       3.184  19.348  23.290  1.00 64.83      L31H C
ATOM   2986  CB  THR  23       1.713  19.709  23.634  1.00 65.15      L31H C
ATOM   2987  OG1 THR  23       1.553  21.133  23.655  1.00 65.49      L31H O
ATOM   2988  CG2 THR  23       1.337  19.152  25.006  1.00 65.92      L31H C
ATOM   2989  C   THR  23       3.492  19.757  21.852  1.00 64.35      L31H C
ATOM   2990  O   THR  23       3.929  20.882  21.596  1.00 64.00      L31H O
ATOM   2991  N   GLY  24       3.269  18.841  20.917  1.00 62.67      L31H N
ATOM   2992  CA  GLY  24       3.772  19.038  19.572  1.00 61.57      L31H C
ATOM   2993  C   GLY  24       2.763  18.800  18.472  1.00 61.13      L31H C
ATOM   2994  O   GLY  24       1.659  19.313  18.532  1.00 62.34      L31H O
ATOM   2995  N   SER  25       3.157  18.029  17.461  1.00 61.63      L31H N
ATOM   2996  CA  SER  25       2.299  17.697  16.319  1.00 62.41      L31H C
ATOM   2997  CB  SER  25       1.914  18.961  15.557  1.00 62.65      L31H C
ATOM   2998  OG  SER  25       2.996  19.425  14.770  1.00 64.84      L31H O
ATOM   2999  C   SER  25       3.025  16.737  15.362  1.00 62.86      L31H C
ATOM   3000  O   SER  25       4.250  16.601  15.416  1.00 63.34      L31H O
ATOM   3001  N   SER  26       2.274  16.092  14.474  1.00 62.26      L31H N
ATOM   3002  CA  SER  26       2.812  14.979  13.695  1.00 62.09      L31H C
ATOM   3003  CB  SER  26       1.695  14.304  12.899  1.00 62.24      L31H C
ATOM   3004  OG  SER  26       1.213  15.157  11.880  1.00 60.48      L31H O
ATOM   3005  C   SER  26       3.939  15.353  12.735  1.00 62.28      L31H C
ATOM   3006  O   SER  26       4.438  14.497  12.006  1.00 61.32      L31H O
ATOM   3007  N   SER  27       4.333  16.620  12.717  1.00 62.13      L31H N
ATOM   3008  CA  SER  27       5.432  17.031  11.854  1.00 61.96      L31H C
ATOM   3009  CB  SER  27       5.100  18.346  11.141  1.00 61.55      L31H C
ATOM   3010  OG  SER  27       5.675  19.458  11.804  1.00 57.20      L31H O
ATOM   3011  C   SER  27       6.732  17.180  12.642  1.00 61.99      L31H C
ATOM   3012  O   SER  27       7.776  17.510  12.072  1.00 62.44      L31H O
ATOM   3013  N   ASN  28       6.664  16.940  13.950  1.00 60.60      L31H N
ATOM   3014  CA  ASN  28       7.861  16.930  14.785  1.00 59.48      L31H C
ATOM   3015  CB  ASN  28       8.044  18.275  15.486  1.00 59.78      L31H C
ATOM   3016  CG  ASN  28       6.746  18.856  15.983  1.00 58.38      L31H C
ATOM   3017  OD1 ASN  28       6.023  19.505  15.235  1.00 58.36      L31H O
ATOM   3018  ND2 ASN  28       6.444  18.631  17.250  1.00 58.43      L31H N
ATOM   3019  C   ASN  28       7.858  15.830  15.822  1.00 59.65      L31H C
ATOM   3020  O   ASN  28       8.286  14.712  15.548  1.00 57.88      L31H O
ATOM   3021  N   ILE  29       7.377  16.151  17.019  1.00 61.65      L31H N
ATOM   3022  CA  ILE  29       7.398  15.196  18.123  1.00 64.05      L31H C
ATOM   3023  CB  ILE  29       6.712  15.764  19.384  1.00 63.61      L31H C
ATOM   3024  CG2 ILE  29       6.839  14.760  20.530  1.00 65.10      L31H C
ATOM   3025  CG1 ILE  29       7.353  17.093  19.781  1.00 63.17      L31H C
ATOM   3026  CD1 ILE  29       6.811  17.686  21.061  1.00 61.54      L31H C
ATOM   3027  C   ILE  29       6.686  13.904  17.730  1.00 65.35      L31H C
ATOM   3028  O   ILE  29       7.096  12.813  18.120  1.00 65.26      L31H O
ATOM   3029  N   GLY  30       5.617  14.043  16.953  1.00 66.80      L31H N
ATOM   3030  CA  GLY  30       4.897  12.886  16.472  1.00 69.29      L31H C
ATOM   3031  C   GLY  30       5.367  12.481  15.093  1.00 70.79      L31H C
ATOM   3032  O   GLY  30       4.609  11.920  14.308  1.00 73.79      L31H O
ATOM   3033  N   ALA  31       6.623  12.772  14.783  1.00 70.57      L31H N
ATOM   3034  CA  ALA  31       7.179  12.388  13.496  1.00 68.69      L31H C
ATOM   3035  CB  ALA  31       7.541  13.623  12.705  1.00 69.46      L31H C
ATOM   3036  C   ALA  31       8.405  11.502  13.689  1.00 67.46      L31H C
ATOM   3037  O   ALA  31       9.124  11.205  12.737  1.00 66.80      L31H O
ATOM   3038  N   GLY  32       8.634  11.081  14.930  1.00 65.06      L31H N
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ATOM   3039  CA  GLY    32       9.744  10.192  15.212  1.00 61.49      L31H C
ATOM   3040  C   GLY    32      10.998  10.939  15.632  1.00 60.15      L31H C
ATOM   3041  O   GLY    32      12.018  10.327  15.972  1.00 58.78      L31H O
ATOM   3042  N   TYR    33      10.923  12.268  15.608  1.00 57.00      L31H N
ATOM   3043  CA  TYR    33      12.055  13.103  15.974  1.00 52.69      L31H C
ATOM   3044  CB  TYR    33      11.947  14.450  15.255  1.00 50.83      L31H C
ATOM   3045  CG  TYR    33      11.990  14.315  13.741  1.00 49.75      L31H C
ATOM   3046  CD1 TYR    33      13.156  13.907  13.099  1.00 49.31      L31H C
ATOM   3047  CE1 TYR    33      13.196  13.721  11.729  1.00 48.18      L31H C
ATOM   3048  CD2 TYR    33      10.857  14.544  12.959  1.00 47.71      L31H C
ATOM   3049  CE2 TYR    33      10.886  14.359  11.583  1.00 47.03      L31H C
ATOM   3050  CZ  TYR    33      12.063  13.942  10.979  1.00 48.69      L31H C
ATOM   3051  OH  TYR    33      12.120  13.707   9.629  1.00 50.47      L31H O
ATOM   3052  C   TYR    33      12.064  13.272  17.487  1.00 50.76      L31H C
ATOM   3053  O   TYR    33      11.014  13.254  18.111  1.00 52.52      L31H O
ATOM   3054  N   ASP    34      13.251  13.407  18.074  1.00 48.14      L31H N
ATOM   3055  CA  ASP    34      13.417  13.437  19.531  1.00 45.26      L31H C
ATOM   3056  CB  ASP    34      14.791  12.898  19.920  1.00 45.18      L31H C
ATOM   3057  CG  ASP    34      14.910  11.409  19.734  1.00 44.74      L31H C
ATOM   3058  OD1 ASP    34      15.934  10.849  20.197  1.00 43.60      L31H O
ATOM   3059  OD2 ASP    34      13.989  10.803  19.133  1.00 45.00      L31H O
ATOM   3060  C   ASP    34      13.269  14.817  20.165  1.00 45.11      L31H C
ATOM   3061  O   ASP    34      13.605  15.849  19.564  1.00 44.68      L31H O
ATOM   3062  N   VAL    35      12.798  14.829  21.407  1.00 44.99      L31H N
ATOM   3063  CA  VAL    35      12.738  16.064  22.178  1.00 43.90      L31H C
ATOM   3064  CB  VAL    35      11.457  16.141  22.997  1.00 44.15      L31H C
ATOM   3065  CG1 VAL    35      11.532  17.331  23.965  1.00 43.72      L31H C
ATOM   3066  CG2 VAL    35      10.256  16.254  22.050  1.00 44.74      L31H C
ATOM   3067  C   VAL    35      13.908  16.221  23.128  1.00 43.44      L31H C
ATOM   3068  O   VAL    35      14.232  15.306  23.888  1.00 42.76      L31H O
ATOM   3069  N   HIS    36      14.534  17.393  23.075  1.00 43.03      L31H N
ATOM   3070  CA  HIS    36      15.642  17.736  23.956  1.00 42.46      L31H C
ATOM   3071  CB  HIS    36      16.868  18.136  23.124  1.00 42.43      L31H C
ATOM   3072  CG  HIS    36      17.111  17.275  21.912  1.00 43.91      L31H C
ATOM   3073  CD2 HIS    36      17.089  17.573  20.587  1.00 43.23      L31H C
ATOM   3074  ND1 HIS    36      17.489  15.948  21.995  1.00 43.31      L31H N
ATOM   3075  CE1 HIS    36      17.689  15.469  20.778  1.00 41.56      L31H C
ATOM   3076  NE2 HIS    36      17.454  16.435  19.906  1.00 41.42      L31H N
ATOM   3077  C   HIS    36      15.167  18.926  24.796  1.00 42.73      L31H C
ATOM   3078  O   HIS    36      14.271  19.649  24.363  1.00 44.22      L31H O
ATOM   3079  N   TRP    37      15.753  19.130  25.981  1.00 43.29      L31H N
ATOM   3080  CA  TRP    37      15.326  20.197  26.916  1.00 43.33      L31H C
ATOM   3081  CB  TRP    37      14.601  19.599  28.124  1.00 45.93      L31H C
ATOM   3082  CG  TRP    37      13.257  18.980  27.843  1.00 49.34      L31H C
ATOM   3083  CD2 TRP    37      11.982  19.600  28.015  1.00 50.13      L31H C
ATOM   3084  CE2 TRP    37      11.002  18.624  27.743  1.00 51.37      L31H C
ATOM   3085  CE3 TRP    37      11.571  20.888  28.378  1.00 52.24      L31H C
ATOM   3086  CD1 TRP    37      13.006  17.686  27.471  1.00 51.27      L31H C
ATOM   3087  NE1 TRP    37      11.651  17.465  27.412  1.00 51.06      L31H N
ATOM   3088  CZ2 TRP    37       9.640  18.895  27.826  1.00 52.35      L31H C
ATOM   3089  CZ3 TRP    37      10.220  21.154  28.457  1.00 52.72      L31H C
ATOM   3090  CH2 TRP    37       9.269  20.162  28.182  1.00 52.64      L31H C
ATOM   3091  C   TRP    37      16.485  21.044  27.457  1.00 43.28      L31H C
ATOM   3092  O   TRP    37      17.541  20.514  27.809  1.00 42.97      L31H O
ATOM   3093  N   TYR    38      16.285  22.355  27.560  1.00 43.01      L31H N
ATOM   3094  CA  TYR    38      17.362  23.232  28.014  1.00 43.05      L31H C
ATOM   3095  CB  TYR    38      17.771  24.181  26.901  1.00 38.88      L31H C
ATOM   3096  CG  TYR    38      18.088  23.478  25.624  1.00 33.77      L31H C
ATOM   3097  CD1 TYR    38      17.073  23.103  24.743  1.00 28.65      L31H C
ATOM   3098  CE1 TYR    38      17.364  22.421  23.583  1.00 25.10      L31H C
ATOM   3099  CD2 TYR    38      19.409  23.154  25.305  1.00 32.66      L31H C
ATOM   3100  CE2 TYR    38      19.713  22.476  24.144  1.00 29.28      L31H C
ATOM   3101  CZ  TYR    38      18.688  22.114  23.289  1.00 27.75      L31H C
ATOM   3102  OH  TYR    38      19.009  21.469  22.127  1.00 26.39      L31H O
ATOM   3103  C   TYR    38      17.007  24.056  29.234  1.00 46.95      L31H C
ATOM   3104  O   TYR    38      15.897  24.577  29.357  1.00 48.67      L31H O
ATOM   3105  N   GLN    39      17.974  24.192  30.127  1.00 49.84      L31H N
ATOM   3106  CA  GLN    39      17.837  25.060  31.281  1.00 52.09      L31H C
ATOM   3107  CB  GLN    39      18.437  24.377  32.506  1.00 52.21      L31H C
ATOM   3108  CG  GLN    39      18.538  25.261  33.725  1.00 54.03      L31H C
ATOM   3109  CD  GLN    39      19.329  24.604  34.840  1.00 55.27      L31H C
ATOM   3110  OE1 GLN    39      20.550  24.734  34.907  1.00 56.25      L31H O
ATOM   3111  NE2 GLN    39      18.635  23.892  35.722  1.00 55.16      L31H N
ATOM   3112  C   GLN    39      18.567  26.371  30.999  1.00 54.46      L31H C
ATOM   3113  O   GLN    39      19.650  26.382  30.401  1.00 52.58      L31H O
ATOM   3114  N   GLN    40      17.965  27.478  31.415  1.00 57.66      L31H N
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ATOM	3115	CA	GLN	40	18.592	28.779	31.254	1.00	61.56	L31H C
ATOM	3116	CB	GLN	40	17.964	29.560	30.092	1.00	59.03	L31H C
ATOM	3117	CG	GLN	40	18.774	30.790	29.713	1.00	56.95	L31H C
ATOM	3118	CD	GLN	40	18.186	31.591	28.563	1.00	56.21	L31H C
ATOM	3119	OE1	GLN	40	16.964	31.695	28.405	1.00	55.30	L31H O
ATOM	3120	NE2	GLN	40	19.066	32.175	27.756	1.00	53.83	L31H N
ATOM	3121	C	GLN	40	18.451	29.584	32.525	1.00	66.50	L31H C
ATOM	3122	O	GLN	40	17.384	30.123	32.813	1.00	66.89	L31H O
ATOM	3123	N	LEU	41	19.533	29.660	33.285	1.00	72.91	L31H N
ATOM	3124	CA	LEU	41	19.607	30.596	34.389	1.00	79.05	L31H C
ATOM	3125	CB	LEU	41	20.938	30.436	35.108	1.00	79.25	L31H C
ATOM	3126	CG	LEU	41	21.403	28.984	35.224	1.00	81.00	L31H C
ATOM	3127	CD1	LEU	41	22.063	28.530	33.916	1.00	80.82	L31H C
ATOM	3128	CD2	LEU	41	22.382	28.871	36.379	1.00	80.82	L31H C
ATOM	3129	C	LEU	41	19.507	31.980	33.773	1.00	82.23	L31H C
ATOM	3130	O	LEU	41	20.261	32.311	32.859	1.00	83.72	L31H O
ATOM	3131	N	PRO	42	18.562	32.803	34.251	1.00	83.71	L31H N
ATOM	3132	CD	PRO	42	17.570	32.557	35.313	1.00	84.10	L31H C
ATOM	3133	CA	PRO	42	18.400	34.127	33.646	1.00	84.51	L31H C
ATOM	3134	CB	PRO	42	17.306	34.780	34.497	1.00	83.82	L31H C
ATOM	3135	CG	PRO	42	16.529	33.621	35.057	1.00	83.24	L31H C
ATOM	3136	C	PRO	42	19.718	34.891	33.692	1.00	84.11	L31H C
ATOM	3137	O	PRO	42	20.413	34.884	34.709	1.00	84.12	L31H O
ATOM	3138	N	GLY	43	20.066	35.527	32.576	1.00	83.31	L31H N
ATOM	3139	CA	GLY	43	21.333	36.227	32.487	1.00	82.03	L31H C
ATOM	3140	C	GLY	43	22.437	35.422	31.823	1.00	81.46	L31H C
ATOM	3141	O	GLY	43	23.581	35.873	31.732	1.00	81.22	L31H O
ATOM	3142	N	THR	44	22.106	34.222	31.357	1.00	79.80	L31H N
ATOM	3143	CA	THR	44	23.093	33.409	30.668	1.00	75.75	L31H C
ATOM	3144	CB	THR	44	23.760	32.413	31.624	1.00	76.28	L31H C
ATOM	3145	OG1	THR	44	24.190	33.104	32.799	1.00	76.33	L31H O
ATOM	3146	CG2	THR	44	24.981	31.776	30.959	1.00	75.30	L31H C
ATOM	3147	C	THR	44	22.492	32.633	29.512	1.00	72.48	L31H C
ATOM	3148	O	THR	44	21.270	32.567	29.352	1.00	73.96	L31H O
ATOM	3149	N	ALA	45	23.375	32.048	28.709	1.00	66.86	L31H N
ATOM	3150	CA	ALA	45	22.990	31.144	27.635	1.00	60.76	L31H C
ATOM	3151	CB	ALA	45	24.237	30.700	26.885	1.00	57.60	L31H C
ATOM	3152	C	ALA	45	22.222	29.918	28.145	1.00	56.71	L31H C
ATOM	3153	O	ALA	45	22.384	29.485	29.294	1.00	54.89	L31H O
ATOM	3154	N	PRO	46	21.366	29.344	27.290	1.00	54.07	L31H N
ATOM	3155	CD	PRO	46	20.704	29.972	26.134	1.00	53.12	L31H C
ATOM	3156	CA	PRO	46	20.794	28.029	27.590	1.00	51.10	L31H C
ATOM	3157	CB	PRO	46	19.846	27.776	26.420	1.00	49.83	L31H C
ATOM	3158	CG	PRO	46	19.452	29.137	25.977	1.00	51.00	L31H C
ATOM	3159	C	PRO	46	21.872	26.961	27.708	1.00	48.33	L31H C
ATOM	3160	O	PRO	46	22.848	26.967	26.969	1.00	45.76	L31H O
ATOM	3161	N	LYS	47	21.696	26.059	28.665	1.00	46.26	L31H N
ATOM	3162	CA	LYS	47	22.559	24.902	28.792	1.00	43.92	L31H C
ATOM	3163	CB	LYS	47	23.201	24.875	30.189	1.00	45.52	L31H C
ATOM	3164	CG	LYS	47	23.152	23.519	30.910	1.00	48.84	L31H C
ATOM	3165	CD	LYS	47	23.908	23.554	32.248	1.00	49.89	L31H C
ATOM	3166	CE	LYS	47	23.940	22.169	32.922	1.00	54.13	L31H C
ATOM	3167	NZ	LYS	47	22.786	21.909	33.858	1.00	56.30	L31H N
ATOM	3168	C	LYS	47	21.695	23.671	28.560	1.00	42.72	L31H C
ATOM	3169	O	LYS	47	20.528	23.630	28.967	1.00	42.20	L31H O
ATOM	3170	N	LEU	48	22.256	22.680	27.875	1.00	41.33	L31H N
ATOM	3171	CA	LEU	48	21.549	21.423	27.651	1.00	39.44	L31H C
ATOM	3172	CB	LEU	48	22.369	20.515	26.729	1.00	36.39	L31H C
ATOM	3173	CG	LEU	48	21.898	19.070	26.538	1.00	33.98	L31H C
ATOM	3174	CD1	LEU	48	20.513	19.024	25.858	1.00	30.93	L31H C
ATOM	3175	CD2	LEU	48	22.951	18.338	25.716	1.00	31.01	L31H C
ATOM	3176	C	LEU	48	21.291	20.713	28.978	1.00	39.99	L31H C
ATOM	3177	O	LEU	48	22.205	20.573	29.813	1.00	37.73	L31H O
ATOM	3178	N	LEU	49	20.046	20.266	29.155	1.00	40.04	L31H N
ATOM	3179	CA	LEU	49	19.604	19.633	30.395	1.00	40.75	L31H C
ATOM	3180	CB	LEU	49	18.407	20.377	30.968	1.00	41.14	L31H C
ATOM	3181	CG	LEU	49	17.894	19.879	32.320	1.00	43.03	L31H C
ATOM	3182	CD1	LEU	49	18.888	20.293	33.442	1.00	40.83	L31H C
ATOM	3183	CD2	LEU	49	16.481	20.451	32.556	1.00	41.04	L31H C
ATOM	3184	C	LEU	49	19.197	18.181	30.196	1.00	42.11	L31H C
ATOM	3185	O	LEU	49	19.422	17.350	31.071	1.00	43.22	L31H O
ATOM	3186	N	ILE	50	18.567	17.889	29.060	1.00	42.58	L31H N
ATOM	3187	CA	ILE	50	18.181	16.526	28.714	1.00	43.24	L31H C
ATOM	3188	CB	ILE	50	16.872	16.118	29.406	1.00	43.38	L31H C
ATOM	3189	CG2	ILE	50	16.229	14.953	28.663	1.00	41.25	L31H C
ATOM	3190	CG1	ILE	50	17.156	15.752	30.855	1.00	42.23	L31H C

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ATOM   3191  CD1 ILE    50      16.212  14.721  31.385  1.00 43.01      L31H C
ATOM   3192  C   ILE    50      18.005  16.304  27.213  1.00 45.56      L31H C
ATOM   3193  O   ILE    50      17.188  16.973  26.569  1.00 43.85      L31H O
ATOM   3194  N   SER    51      18.759  15.340  26.673  1.00 48.30      L31H N
ATOM   3195  CA  SER    51      18.711  15.008  25.246  1.00 49.71      L31H C
ATOM   3196  CB  SER    51      20.130  14.952  24.675  1.00 49.26      L31H C
ATOM   3197  OG  SER    51      20.939  14.047  25.403  1.00 52.11      L31H O
ATOM   3198  C   SER    51      17.991  13.686  24.961  1.00 50.08      L31H C
ATOM   3199  O   SER    51      17.737  12.896  25.865  1.00 50.16      L31H O
ATOM   3200  N   GLY    52      17.648  13.468  23.695  1.00 51.31      L31H N
ATOM   3201  CA  GLY    52      17.056  12.209  23.284  1.00 52.07      L31H C
ATOM   3202  C   GLY    52      15.903  11.773  24.164  1.00 54.09      L31H C
ATOM   3203  O   GLY    52      15.778  10.590  24.499  1.00 54.32      L31H O
ATOM   3204  N   ASN    53      15.069  12.737  24.551  1.00 55.48      L31H N
ATOM   3205  CA  ASN    53      13.841  12.470  25.310  1.00 55.60      L31H C
ATOM   3206  CB  ASN    53      13.102  11.243  24.751  1.00 52.09      L31H C
ATOM   3207  CG  ASN    53      12.894  11.306  23.248  1.00 49.39      L31H C
ATOM   3208  OD1 ASN    53      12.568  12.353  22.692  1.00 50.31      L31H O
ATOM   3209  ND2 ASN    53      13.081  10.176  22.584  1.00 46.57      L31H N
ATOM   3210  C   ASN    53      14.102  12.245  26.799  1.00 56.53      L31H C
ATOM   3211  O   ASN    53      13.345  12.709  27.647  1.00 57.59      L31H O
ATOM   3212  N   SER    54      15.177  11.537  27.115  1.00 58.06      L31H N
ATOM   3213  CA  SER    54      15.289  10.921  28.421  1.00 59.90      L31H C
ATOM   3214  CB  SER    54      14.677   9.518  28.357  1.00 60.64      L31H C
ATOM   3215  OG  SER    54      15.248   8.732  27.315  1.00 59.19      L31H O
ATOM   3216  C   SER    54      16.709  10.844  29.000  1.00 62.28      L31H C
ATOM   3217  O   SER    54      16.905  10.242  30.055  1.00 62.32      L31H O
ATOM   3218  N   ASN    55      17.692  11.444  28.330  1.00 64.32      L31H N
ATOM   3219  CA  ASN    55      19.087  11.293  28.744  1.00 67.64      L31H C
ATOM   3220  CB  ASN    55      19.938  10.886  27.545  1.00 67.43      L31H C
ATOM   3221  CG  ASN    55      19.450   9.603  26.892  1.00 67.97      L31H C
ATOM   3222  OD1 ASN    55      19.565   8.509  27.458  1.00 66.43      L31H O
ATOM   3223  ND2 ASN    55      18.898   9.731  25.695  1.00 68.44      L31H N
ATOM   3224  C   ASN    55      19.700  12.524  29.401  1.00 69.95      L31H C
ATOM   3225  O   ASN    55      19.502  13.638  28.937  1.00 71.67      L31H O
ATOM   3226  N   ARG    56      20.457  12.307  30.477  1.00 72.79      L31H N
ATOM   3227  CA  ARG    56      21.089  13.387  31.238  1.00 75.45      L31H C
ATOM   3228  CB  ARG    56      20.883  13.179  32.739  1.00 77.48      L31H C
ATOM   3229  CG  ARG    56      19.439  13.153  33.188  1.00 83.20      L31H C
ATOM   3230  CD  ARG    56      19.324  12.706  34.637  1.00 88.80      L31H C
ATOM   3231  NE  ARG    56      19.736  11.315  34.816  1.00 95.67      L31H N
ATOM   3232  CZ  ARG    56      20.967  10.935  35.153  1.00 99.28      L31H C
ATOM   3233  NH1 ARG    56      21.260   9.643  35.293  1.00100.99      L31H N
ATOM   3234  NH2 ARG    56      21.908  11.851  35.352  1.00101.75      L31H N
ATOM   3235  C   ARG    56      22.587  13.481  30.978  1.00 75.94      L31H C
ATOM   3236  O   ARG    56      23.318  12.505  31.139  1.00 76.36      L31H O
ATOM   3237  N   PRO    57      23.070  14.668  30.586  1.00 76.18      L31H N
ATOM   3238  CD  PRO    57      22.322  15.811  30.039  1.00 75.75      L31H C
ATOM   3239  CA  PRO    57      24.519  14.895  30.545  1.00 77.03      L31H C
ATOM   3240  CB  PRO    57      24.656  16.255  29.859  1.00 75.92      L31H C
ATOM   3241  CG  PRO    57      23.346  16.482  29.176  1.00 75.20      L31H C
ATOM   3242  C   PRO    57      25.064  14.931  31.965  1.00 77.22      L31H C
ATOM   3243  O   PRO    57      24.490  15.587  32.832  1.00 78.83      L31H O
ATOM   3244  N   SER    58      26.166  14.232  32.208  1.00 76.68      L31H N
ATOM   3245  CA  SER    58      26.803  14.297  33.521  1.00 75.29      L31H C
ATOM   3246  CB  SER    58      28.198  13.664  33.488  1.00 75.46      L31H C
ATOM   3247  OG  SER    58      29.072  14.426  32.675  1.00 72.13      L31H O
ATOM   3248  C   SER    58      26.917  15.768  33.910  1.00 73.26      L31H C
ATOM   3249  O   SER    58      27.172  16.625  33.067  1.00 73.30      L31H O
ATOM   3250  N   GLY    59      26.732  16.046  35.192  1.00 71.02      L31H N
ATOM   3251  CA  GLY    59      26.434  17.396  35.623  1.00 66.74      L31H C
ATOM   3252  C   GLY    59      24.984  17.447  36.062  1.00 63.73      L31H C
ATOM   3253  O   GLY    59      24.676  17.851  37.181  1.00 63.24      L31H O
ATOM   3254  N   VAL    60      24.089  17.013  35.182  1.00 60.90      L31H N
ATOM   3255  CA  VAL    60      22.662  17.015  35.480  1.00 58.68      L31H C
ATOM   3256  CB  VAL    60      21.850  17.011  34.178  1.00 57.78      L31H C
ATOM   3257  CG1 VAL    60      20.391  17.176  34.475  1.00 56.47      L31H C
ATOM   3258  CG2 VAL    60      22.336  18.108  33.268  1.00 58.00      L31H C
ATOM   3259  C   VAL    60      22.277  15.789  36.325  1.00 58.24      L31H C
ATOM   3260  O   VAL    60      22.498  14.646  35.923  1.00 55.12      L31H O
ATOM   3261  N   PRO    61      21.706  16.023  37.519  1.00 58.08      L31H N
ATOM   3262  CD  PRO    61      21.552  17.349  38.137  1.00 57.75      L31H C
ATOM   3263  CA  PRO    61      21.353  14.951  38.459  1.00 58.26      L31H C
ATOM   3264  CB  PRO    61      21.030  15.696  39.760  1.00 58.11      L31H C
ATOM   3265  CG  PRO    61      21.652  17.034  39.600  1.00 58.28      L31H C
ATOM   3266  C   PRO    61      20.172  14.116  37.976  1.00 58.36      L31H C
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ATOM   3267  O   PRO   61      19.438  14.526  37.074  1.00 56.83      L31H O
ATOM   3268  N   ASP   62      19.991  12.947  38.579  1.00 58.64      L31H N
ATOM   3269  CA  ASP   62      18.872  12.098  38.205  1.00 59.95      L31H C
ATOM   3270  CB  ASP   62      19.073  10.657  38.707  1.00 62.80      L31H C
ATOM   3271  CG  ASP   62      19.202  10.556  40.234  1.00 67.16      L31H C
ATOM   3272  OD1 ASP   62      19.770   9.540  40.700  1.00 67.81      L31H O
ATOM   3273  OD2 ASP   62      18.742  11.471  40.969  1.00 69.22      L31H O
ATOM   3274  C   ASP   62      17.587  12.686  38.758  1.00 58.55      L31H C
ATOM   3275  O   ASP   62      16.500  12.160  38.530  1.00 57.80      L31H O
ATOM   3276  N   ARG   63      17.721  13.787  39.487  1.00 57.77      L31H N
ATOM   3277  CA  ARG   63      16.561  14.546  39.933  1.00 57.68      L31H C
ATOM   3278  CB  ARG   63      17.006  15.733  40.781  1.00 59.02      L31H C
ATOM   3279  CG  ARG   63      17.583  15.367  42.130  1.00 60.91      L31H C
ATOM   3280  CD  ARG   63      17.748  16.618  42.979  1.00 64.55      L31H C
ATOM   3281  NE  ARG   63      18.504  17.660  42.280  1.00 67.17      L31H N
ATOM   3282  CZ  ARG   63      18.006  18.840  41.915  1.00 67.43      L31H C
ATOM   3283  NH1 ARG   63      16.741  19.145  42.178  1.00 67.62      L31H N
ATOM   3284  NH2 ARG   63      18.779  19.719  41.297  1.00 67.50      L31H N
ATOM   3285  C   ARG   63      15.761  15.054  38.734  1.00 56.49      L31H C
ATOM   3286  O   ARG   63      14.541  15.224  38.817  1.00 56.13      L31H O
ATOM   3287  N   PHE   64      16.471  15.297  37.632  1.00 55.21      L31H N
ATOM   3288  CA  PHE   64      15.873  15.705  36.360  1.00 54.46      L31H C
ATOM   3289  CB  PHE   64      16.782  16.699  35.623  1.00 53.69      L31H C
ATOM   3290  CG  PHE   64      17.058  17.949  36.385  1.00 51.65      L31H C
ATOM   3291  CD1 PHE   64      16.251  19.067  36.220  1.00 50.28      L31H C
ATOM   3292  CD2 PHE   64      18.131  18.010  37.269  1.00 50.64      L31H C
ATOM   3293  CE1 PHE   64      16.507  20.226  36.920  1.00 48.84      L31H C
ATOM   3294  CE2 PHE   64      18.394  19.165  37.974  1.00 48.98      L31H C
ATOM   3295  CZ  PHE   64      17.577  20.280  37.798  1.00 48.52      L31H C
ATOM   3296  C   PHE   64      15.690  14.490  35.459  1.00 53.96      L31H C
ATOM   3297  O   PHE   64      16.627  13.713  35.262  1.00 53.07      L31H O
ATOM   3298  N   SER   65      14.495  14.344  34.896  1.00 54.00      L31H N
ATOM   3299  CA  SER   65      14.223  13.248  33.970  1.00 54.18      L31H C
ATOM   3300  CB  SER   65      13.755  12.013  34.748  1.00 53.56      L31H C
ATOM   3301  OG  SER   65      12.887  12.390  35.802  1.00 52.90      L31H O
ATOM   3302  C   SER   65      13.182  13.653  32.927  1.00 54.21      L31H C
ATOM   3303  O   SER   65      12.299  14.463  33.202  1.00 54.41      L31H O
ATOM   3304  N   GLY   66      13.303  13.100  31.725  1.00 54.11      L31H N
ATOM   3305  CA  GLY   66      12.403  13.472  30.649  1.00 53.71      L31H C
ATOM   3306  C   GLY   66      11.674  12.267  30.103  1.00 54.08      L31H C
ATOM   3307  O   GLY   66      11.973  11.136  30.478  1.00 54.45      L31H O
ATOM   3308  N   SER   67      10.708  12.500  29.224  1.00 54.58      L31H N
ATOM   3309  CA  SER   67       9.993  11.398  28.591  1.00 55.72      L31H C
ATOM   3310  CB  SER   67       9.188  10.609  29.632  1.00 55.62      L31H C
ATOM   3311  OG  SER   67       8.017  11.305  30.019  1.00 57.25      L31H O
ATOM   3312  C   SER   67       9.067  11.916  27.496  1.00 56.30      L31H C
ATOM   3313  O   SER   67       8.618  13.060  27.535  1.00 55.91      L31H O
ATOM   3314  N   LYS   68       8.797  11.064  26.515  1.00 56.98      L31H N
ATOM   3315  CA  LYS   68       8.024  11.453  25.347  1.00 57.91      L31H C
ATOM   3316  CB  LYS   68       8.933  11.564  24.118  1.00 57.46      L31H C
ATOM   3317  CG  LYS   68       8.182  11.490  22.801  1.00 56.11      L31H C
ATOM   3318  CD  LYS   68       9.093  11.156  21.637  1.00 56.19      L31H C
ATOM   3319  CE  LYS   68       8.287  10.873  20.368  1.00 55.24      L31H C
ATOM   3320  NZ  LYS   68       9.135  10.836  19.137  1.00 51.72      L31H N
ATOM   3321  C   LYS   68       6.949  10.416  25.084  1.00 59.65      L31H C
ATOM   3322  O   LYS   68       7.141   9.224  25.340  1.00 60.17      L31H O
ATOM   3323  N   SER   69       5.817  10.874  24.564  1.00 61.33      L31H N
ATOM   3324  CA  SER   69       4.696   9.995  24.280  1.00 61.60      L31H C
ATOM   3325  CB  SER   69       3.870   9.787  25.550  1.00 60.95      L31H C
ATOM   3326  OG  SER   69       2.891   8.791  25.352  1.00 59.59      L31H O
ATOM   3327  C   SER   69       3.822  10.619  23.203  1.00 61.91      L31H C
ATOM   3328  O   SER   69       3.270  11.708  23.397  1.00 64.14      L31H O
ATOM   3329  N   GLY   70       3.694   9.934  22.071  1.00 60.05      L31H N
ATOM   3330  CA  GLY   70       2.820  10.421  21.018  1.00 56.96      L31H C
ATOM   3331  C   GLY   70       3.285  11.749  20.447  1.00 56.37      L31H C
ATOM   3332  O   GLY   70       4.269  11.810  19.709  1.00 56.19      L31H O
ATOM   3333  N   THR   71       2.587  12.825  20.786  1.00 55.57      L31H N
ATOM   3334  CA  THR   71       2.895  14.124  20.212  1.00 55.61      L31H C
ATOM   3335  CB  THR   71       1.685  14.708  19.503  1.00 54.53      L31H C
ATOM   3336  OG1 THR   71       0.597  14.758  20.428  1.00 56.64      L31H O
ATOM   3337  CG2 THR   71       1.289  13.850  18.316  1.00 54.65      L31H C
ATOM   3338  C   THR   71       3.328  15.098  21.289  1.00 56.84      L31H C
ATOM   3339  O   THR   71       3.385  16.308  21.061  1.00 57.59      L31H O
ATOM   3340  N   SER   72       3.628  14.572  22.470  1.00 57.58      L31H N
ATOM   3341  CA  SER   72       4.041  15.421  23.579  1.00 57.61      L31H C
ATOM   3342  CB  SER   72       2.875  15.631  24.542  1.00 57.91      L31H C
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ATOM   3343  OG  SER    72       2.211  14.410  24.795  1.00 58.50      L31H O
ATOM   3344  C   SER    72       5.221  14.830  24.323  1.00 57.22      L31H C
ATOM   3345  O   SER    72       5.757  13.796  23.937  1.00 57.93      L31H O
ATOM   3346  N   ALA    73       3.625  15.502  25.392  1.00 56.55      L31H N
ATOM   3347  CA  ALA    73       6.766  15.073  26.187  1.00 56.17      L31H C
ATOM   3348  CB  ALA    73       8.063  15.426  25.466  1.00 55.12      L31H C
ATOM   3349  C   ALA    73       6.681  15.802  27.514  1.00 56.46      L31H C
ATOM   3350  O   ALA    73       5.891  16.719  27.659  1.00 56.55      L31H O
ATOM   3351  N   SER    74       7.484  15.401  28.487  1.00 58.75      L31H N
ATOM   3352  CA  SER    74       7.588  16.178  29.716  1.00 60.34      L31H C
ATOM   3353  CB  SER    74       6.575  15.712  30.738  1.00 60.09      L31H C
ATOM   3354  OG  SER    74       6.962  16.205  32.010  1.00 58.83      L31H O
ATOM   3355  C   SER    74       8.953  16.163  30.389  1.00 60.92      L31H C
ATOM   3356  O   SER    74       9.735  15.224  30.228  1.00 63.33      L31H O
ATOM   3357  N   LEU    75       9.216  17.211  31.162  1.00 59.31      L31H N
ATOM   3358  CA  LEU    75      10.383  17.259  32.032  1.00 58.44      L31H C
ATOM   3359  CB  LEU    75      11.150  18.574  31.803  1.00 59.47      L31H C
ATOM   3360  CG  LEU    75      12.456  18.834  32.565  1.00 57.74      L31H C
ATOM   3361  CD1 LEU    75      13.397  17.653  32.379  1.00 58.12      L31H C
ATOM   3362  CD2 LEU    75      13.102  20.118  32.067  1.00 56.35      L31H C
ATOM   3363  C   LEU    75       9.924  17.163  33.492  1.00 58.84      L31H C
ATOM   3364  O   LEU    75       9.086  17.945  33.938  1.00 58.64      L31H O
ATOM   3365  N   ALA    76      10.470  16.201  34.226  1.00 59.13      L31H N
ATOM   3366  CA  ALA    76      10.194  16.059  35.650  1.00 60.32      L31H C
ATOM   3367  CB  ALA    76       9.776  14.632  35.951  1.00 59.16      L31H C
ATOM   3368  C   ALA    76      11.427  16.423  36.480  1.00 61.43      L31H C
ATOM   3369  O   ALA    76      12.535  15.935  36.220  1.00 62.96      L31H O
ATOM   3370  N   ILE    77      11.237  17.276  37.482  1.00 61.31      L31H N
ATOM   3371  CA  ILE    77      12.317  17.597  38.413  1.00 61.17      L31H C
ATOM   3372  CB  ILE    77      12.624  19.120  38.434  1.00 61.96      L31H C
ATOM   3373  CG2 ILE    77      13.872  19.387  39.253  1.00 61.34      L31H C
ATOM   3374  CG1 ILE    77      12.863  19.633  37.015  1.00 61.57      L31H C
ATOM   3375  CD1 ILE    77      13.330  21.061  36.956  1.00 60.74      L31H C
ATOM   3376  C   ILE    77      11.932  17.163  39.821  1.00 60.56      L31H C
ATOM   3377  O   ILE    77      10.914  17.592  40.341  1.00 60.68      L31H O
ATOM   3378  N   THR    78      12.733  16.315  40.447  1.00 60.81      L31H N
ATOM   3379  CA  THR    78      12.422  15.930  41.815  1.00 63.27      L31H C
ATOM   3380  CB  THR    78      12.493  14.402  41.993  1.00 62.92      L31H C
ATOM   3381  OG1 THR    78      13.677  13.902  41.366  1.00 65.95      L31H O
ATOM   3382  CG2 THR    78      11.287  13.739  41.375  1.00 62.85      L31H C
ATOM   3383  C   THR    78      13.315  16.603  42.861  1.00 64.30      L31H C
ATOM   3384  O   THR    78      14.312  17.251  42.523  1.00 63.28      L31H O
ATOM   3385  N   GLY    79      12.929  16.460  44.128  1.00 64.43      L31H N
ATOM   3386  CA  GLY    79      13.737  16.961  45.222  1.00 64.73      L31H C
ATOM   3387  C   GLY    79      14.187  18.371  44.944  1.00 65.69      L31H C
ATOM   3388  O   GLY    79      15.382  18.678  44.980  1.00 66.58      L31H O
ATOM   3389  N   LEU    80      13.219  19.233  44.662  1.00 66.28      L31H N
ATOM   3390  CA  LEU    80      13.501  20.552  44.116  1.00 67.14      L31H C
ATOM   3391  CB  LEU    80      12.182  21.315  43.940  1.00 66.33      L31H C
ATOM   3392  CG  LEU    80      12.006  22.208  42.707  1.00 65.90      L31H C
ATOM   3393  CD1 LEU    80      12.826  21.693  41.522  1.00 64.85      L31H C
ATOM   3394  CD2 LEU    80      10.531  22.256  42.364  1.00 63.32      L31H C
ATOM   3395  C   LEU    80      14.473  21.333  45.008  1.00 67.62      L31H C
ATOM   3396  O   LEU    80      14.247  21.501  46.203  1.00 66.07      L31H O
ATOM   3397  N   GLN    81      15.571  21.782  44.412  1.00 69.02      L31H N
ATOM   3398  CA  GLN    81      16.511  22.664  45.090  1.00 71.32      L31H C
ATOM   3399  CB  GLN    81      17.956  22.251  44.784  1.00 71.81      L31H C
ATOM   3400  CG  GLN    81      18.387  20.908  45.359  1.00 72.58      L31H C
ATOM   3401  CD  GLN    81      19.656  20.370  44.703  1.00 73.31      L31H C
ATOM   3402  OE1 GLN    81      20.123  20.900  43.695  1.00 73.31      L31H O
ATOM   3403  NE2 GLN    81      20.212  19.309  45.274  1.00 75.44      L31H N
ATOM   3404  C   GLN    81      16.262  24.082  44.586  1.00 71.73      L31H C
ATOM   3405  O   GLN    81      15.703  24.269  43.508  1.00 71.27      L31H O
ATOM   3406  N   ALA    82      16.675  25.076  45.365  1.00 73.50      L31H N
ATOM   3407  CA  ALA    82      16.338  26.464  45.064  1.00 75.37      L31H C
ATOM   3408  CB  ALA    82      16.599  27.340  46.277  1.00 76.64      L31H C
ATOM   3409  C   ALA    82      17.146  26.961  43.882  1.00 76.22      L31H C
ATOM   3410  O   ALA    82      16.703  27.823  43.123  1.00 76.26      L31H O
ATOM   3411  N   GLU    83      18.336  26.399  43.732  1.00 76.93      L31H N
ATOM   3412  CA  GLU    83      19.222  26.755  42.645  1.00 77.92      L31H C
ATOM   3413  CB  GLU    83      20.657  26.390  43.025  1.00 82.68      L31H C
ATOM   3414  CG  GLU    83      21.235  27.232  44.165  1.00 90.24      L31H C
ATOM   3415  CD  GLU    83      20.590  26.959  45.522  1.00 94.97      L31H C
ATOM   3416  OE1 GLU    83      20.480  25.777  45.917  1.00 99.12      L31H O
ATOM   3417  OE2 GLU    83      20.199  27.934  46.199  1.00 98.16      L31H O
ATOM   3418  C   GLU    83      18.802  26.050  41.356  1.00 75.65      L31H C
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ATOM   3419  O   GLU  83      19.563  25.980  40.395  1.00 74.80      L31H O
ATOM   3420  N   ASP  84      17.579  25.527  41.351  1.00 73.41      L31H N
ATOM   3421  CA  ASP  84      16.981  24.964  40.145  1.00 71.11      L31H C
ATOM   3422  CB  ASP  84      16.245  23.661  40.471  1.00 70.69      L31H C
ATOM   3423  CG  ASP  84      17.181  22.466  40.499  1.00 70.12      L31H C
ATOM   3424  OD1 ASP  84      18.176  22.487  39.741  1.00 70.45      L31H O
ATOM   3425  OD2 ASP  84      16.935  21.514  41.271  1.00 67.92      L31H O
ATOM   3426  C   ASP  84      16.032  25.949  39.495  1.00 69.37      L31H C
ATOM   3427  O   ASP  84      15.559  25.735  38.385  1.00 69.35      L31H O
ATOM   3428  N   GLU  85      15.765  27.040  40.197  1.00 67.68      L31H N
ATOM   3429  CA  GLU  85      14.955  28.115  39.653  1.00 65.89      L31H C
ATOM   3430  CB  GLU  85      14.859  29.253  40.680  1.00 67.36      L31H C
ATOM   3431  CG  GLU  85      14.009  30.448  40.250  1.00 69.50      L31H C
ATOM   3432  CD  GLU  85      13.407  31.190  41.433  1.00 71.65      L31H C
ATOM   3433  OE1 GLU  85      13.686  32.402  41.596  1.00 73.05      L31H O
ATOM   3434  OE2 GLU  85      12.650  30.554  42.200  1.00 73.05      L31H O
ATOM   3435  C   GLU  85      15.585  28.613  38.346  1.00 63.80      L31H C
ATOM   3436  O   GLU  85      16.765  28.993  38.316  1.00 62.73      L31H O
ATOM   3437  N   ALA  86      14.795  28.601  37.270  1.00 61.53      L31H N
ATOM   3438  CA  ALA  86      15.285  28.998  35.949  1.00 58.22      L31H C
ATOM   3439  CB  ALA  86      16.519  28.179  35.595  1.00 59.72      L31H C
ATOM   3440  C   ALA  86      14.251  28.888  34.819  1.00 56.10      L31H C
ATOM   3441  O   ALA  86      13.093  28.519  35.037  1.00 55.00      L31H O
ATOM   3442  N   ASP  87      14.691  29.225  33.608  1.00 53.32      L31H N
ATOM   3443  CA  ASP  87      13.868  29.104  32.415  1.00 51.34      L31H C
ATOM   3444  CB  ASP  87      14.125  30.272  31.475  1.00 51.11      L31H C
ATOM   3445  CG  ASP  87      13.594  31.583  32.020  1.00 53.30      L31H C
ATOM   3446  OD1 ASP  87      12.788  31.549  32.972  1.00 53.43      L31H O
ATOM   3447  OD2 ASP  87      13.983  32.649  31.491  1.00 55.18      L31H O
ATOM   3448  C   ASP  87      14.154  27.801  31.675  1.00 51.29      L31H C
ATOM   3449  O   ASP  87      15.285  27.556  31.221  1.00 50.39      L31H O
ATOM   3450  N   TYR  88      13.113  26.977  31.544  1.00 49.52      L31H N
ATOM   3451  CA  TYR  88      13.207  25.706  30.848  1.00 49.11      L31H C
ATOM   3452  CB  TYR  88      12.560  24.624  31.706  1.00 48.57      L31H C
ATOM   3453  CG  TYR  88      13.368  24.390  32.957  1.00 47.75      L31H C
ATOM   3454  CD1 TYR  88      12.958  24.884  34.188  1.00 47.35      L31H C
ATOM   3455  CE1 TYR  88      13.767  24.761  35.309  1.00 46.21      L31H C
ATOM   3456  CD2 TYR  88      14.599  23.760  32.884  1.00 47.54      L31H C
ATOM   3457  CE2 TYR  88      15.411  23.633  33.986  1.00 46.69      L31H C
ATOM   3458  CZ  TYR  88      15.000  24.136  35.197  1.00 46.67      L31H C
ATOM   3459  OH  TYR  88      15.849  24.031  36.274  1.00 43.55      L31H O
ATOM   3460  C   TYR  88      12.585  25.768  29.457  1.00 49.99      L31H C
ATOM   3461  O   TYR  88      11.539  26.388  29.257  1.00 50.31      L31H O
ATOM   3462  N   TYR  89      13.262  25.157  28.490  1.00 49.96      L31H N
ATOM   3463  CA  TYR  89      12.795  25.166  27.115  1.00 51.75      L31H C
ATOM   3464  CB  TYR  89      13.607  26.157  26.259  1.00 53.70      L31H C
ATOM   3465  CG  TYR  89      13.533  27.636  26.644  1.00 56.56      L31H C
ATOM   3466  CD1 TYR  89      14.474  28.196  27.509  1.00 57.34      L31H C
ATOM   3467  CE1 TYR  89      14.466  29.550  27.814  1.00 58.06      L31H C
ATOM   3468  CD2 TYR  89      12.567  28.482  26.095  1.00 56.23      L31H C
ATOM   3469  CE2 TYR  89      12.548  29.843  26.396  1.00 57.93      L31H C
ATOM   3470  CZ  TYR  89      13.507  30.371  27.257  1.00 58.84      L31H C
ATOM   3471  OH  TYR  89      13.533  31.723  27.547  1.00 59.54      L31H O
ATOM   3472  C   TYR  89      12.955  23.760  26.541  1.00 52.19      L31H C
ATOM   3473  O   TYR  89      13.932  23.063  26.830  1.00 51.07      L31H O
ATOM   3474  N   CYS  90      11.988  23.356  25.722  1.00 51.93      L31H N
ATOM   3475  CA  CYS  90      12.106  22.135  24.939  1.00 51.12      L31H C
ATOM   3476  C   CYS  90      12.345  22.414  23.460  1.00 50.42      L31H C
ATOM   3477  O   CYS  90      12.111  23.526  22.983  1.00 50.17      L31H O
ATOM   3478  CB  CYS  90      10.848  21.292  25.097  1.00 53.40      L31H C
ATOM   3479  SG  CYS  90       9.280  22.130  24.728  1.00 51.64      L31H S
ATOM   3480  N   GLN  91      12.802  21.395  22.733  1.00 49.04      L31H N
ATOM   3481  CA  GLN  91      13.132  21.550  21.320  1.00 46.80      L31H C
ATOM   3482  CB  GLN  91      14.568  22.037  21.204  1.00 46.39      L31H C
ATOM   3483  CG  GLN  91      15.174  21.980  19.822  1.00 43.37      L31H C
ATOM   3484  CD  GLN  91      16.675  21.962  19.925  1.00 43.12      L31H C
ATOM   3485  OE1 GLN  91      17.231  21.108  20.604  1.00 42.92      L31H O
ATOM   3486  NE2 GLN  91      17.343  22.909  19.273  1.00 41.91      L31H N
ATOM   3487  C   GLN  91      12.964  20.257  20.531  1.00 46.33      L31H C
ATOM   3488  O   GLN  91      13.317  19.183  21.015  1.00 45.84      L31H O
ATOM   3489  N   SER  92      12.433  20.371  19.313  1.00 45.77      L31H N
ATOM   3490  CA  SER  92      12.363  19.246  18.381  1.00 45.24      L31H C
ATOM   3491  CB  SER  92      10.977  18.603  18.438  1.00 45.03      L31H C
ATOM   3492  OG  SER  92      10.870  17.510  17.537  1.00 47.80      L31H O
ATOM   3493  C   SER  92      12.679  19.673  16.937  1.00 45.02      L31H C
ATOM   3494  O   SER  92      12.694  20.860  16.622  1.00 44.46      L31H O
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ATOM   3495  N   TYR    93      12.948  18.695  16.073  1.00 44.70      L31H N
ATOM   3496  CA  TYR    93      13.093  18.933  14.640  1.00 43.04      L31H C
ATOM   3497  CB  TYR    93      13.960  17.835  13.993  1.00 40.79      L31H C
ATOM   3498  CG  TYR    93      14.200  18.002  12.493  1.00 38.37      L31H C
ATOM   3499  CD1 TYR    93      15.219  18.832  12.014  1.00 36.91      L31H C
ATOM   3500  CE1 TYR    93      15.435  19.013  10.642  1.00 32.83      L31H C
ATOM   3501  CD2 TYR    93      13.400  17.349  11.558  1.00 35.95      L31H C
ATOM   3502  CE2 TYR    93      13.609  17.521  10.180  1.00 35.00      L31H C
ATOM   3503  CZ  TYR    93      14.631  18.360   9.732  1.00 34.67      L31H C
ATOM   3504  OH  TYR    93      14.833  18.558   8.381  1.00 32.18      L31H O
ATOM   3505  C   TYR    93      11.692  18.892  14.055  1.00 43.84      L31H C
ATOM   3506  O   TYR    93      10.820  18.220  14.592  1.00 42.51      L31H O
ATOM   3507  N   ASP    94      11.482  19.615  12.964  1.00 45.57      L31H N
ATOM   3508  CA  ASP    94      10.186  19.654  12.308  1.00 47.61      L31H C
ATOM   3509  CB  ASP    94       9.550  21.027  12.531  1.00 49.33      L31H C
ATOM   3510  CG  ASP    94       8.067  21.041  12.222  1.00 51.78      L31H C
ATOM   3511  OD1 ASP    94       7.277  21.314  13.156  1.00 52.86      L31H O
ATOM   3512  OD2 ASP    94       7.693  20.783  11.056  1.00 51.71      L31H O
ATOM   3513  C   ASP    94      10.399  19.402  10.818  1.00 48.27      L31H C
ATOM   3514  O   ASP    94      11.058  20.187  10.141  1.00 48.27      L31H O
ATOM   3515  N   SER    95       9.848  18.312  10.299  1.00 49.99      L31H N
ATOM   3516  CA  SER    95      10.228  17.877   8.964  1.00 52.79      L31H C
ATOM   3517  CB  SER    95       9.762  16.446   8.702  1.00 52.89      L31H C
ATOM   3518  OG  SER    95       8.526  16.436   8.005  1.00 54.06      L31H O
ATOM   3519  C   SER    95       9.663  18.797   7.890  1.00 55.32      L31H C
ATOM   3520  O   SER    95      10.082  18.735   6.734  1.00 56.36      L31H O
ATOM   3521  N   SER    96       8.712  19.649   8.261  1.00 57.01      L31H N
ATOM   3522  CA  SER    96       8.219  20.657   7.330  1.00 59.06      L31H C
ATOM   3523  CB  SER    96       6.765  20.981   7.624  1.00 57.61      L31H C
ATOM   3524  OG  SER    96       6.662  21.651   8.865  1.00 56.06      L31H O
ATOM   3525  C   SER    96       9.048  21.943   7.412  1.00 61.43      L31H C
ATOM   3526  O   SER    96       9.380  22.542   6.391  1.00 63.10      L31H O
ATOM   3527  N   LEU    97       9.380  22.369   8.625  1.00 62.38      L31H N
ATOM   3528  CA  LEU    97      10.125  23.608   8.802  1.00 63.01      L31H C
ATOM   3529  CB  LEU    97       9.970  24.118  10.238  1.00 66.93      L31H C
ATOM   3530  CG  LEU    97       8.521  24.274  10.714  1.00 69.67      L31H C
ATOM   3531  CD1 LEU    97       8.477  25.078  12.010  1.00 69.75      L31H C
ATOM   3532  CD2 LEU    97       7.709  24.963   9.636  1.00 69.77      L31H C
ATOM   3533  C   LEU    97      11.606  23.422   8.477  1.00 61.49      L31H C
ATOM   3534  O   LEU    97      12.307  24.384   8.146  1.00 59.90      L31H O
ATOM   3535  N   SER    98      12.073  22.181   8.577  1.00 58.39      L31H N
ATOM   3536  CA  SER    98      13.452  21.858   8.262  1.00 55.51      L31H C
ATOM   3537  CB  SER    98      13.793  22.338   6.856  1.00 56.09      L31H C
ATOM   3538  OG  SER    98      13.189  21.493   5.898  1.00 59.25      L31H O
ATOM   3539  C   SER    98      14.422  22.457   9.264  1.00 53.09      L31H C
ATOM   3540  O   SER    98      15.576  22.727   8.945  1.00 51.29      L31H O
ATOM   3541  N   GLY    99      13.950  22.663  10.484  1.00 51.30      L31H N
ATOM   3542  CA  GLY    99      14.827  23.149  11.532  1.00 50.67      L31H C
ATOM   3543  C   GLY    99      14.406  22.584  12.868  1.00 50.25      L31H C
ATOM   3544  O   GLY    99      13.335  21.998  12.991  1.00 49.46      L31H O
ATOM   3545  N   SER   100      15.245  22.754  13.877  1.00 50.20      L31H N
ATOM   3546  CA  SER   100      14.920  22.233  15.193  1.00 50.66      L31H C
ATOM   3547  CB  SER   100      16.177  21.667  15.858  1.00 51.28      L31H C
ATOM   3548  OG  SER   100      16.337  20.294  15.521  1.00 52.02      L31H O
ATOM   3549  C   SER   100      14.294  23.321  16.055  1.00 50.29      L31H C
ATOM   3550  O   SER   100      14.995  24.184  16.582  1.00 50.01      L31H O
ATOM   3551  N   VAL   101      12.970  23.266  16.190  1.00 48.60      L31H N
ATOM   3552  CA  VAL   101      12.189  24.384  16.701  1.00 47.52      L31H C
ATOM   3553  CB  VAL   101      10.881  24.535  15.918  1.00 46.73      L31H C
ATOM   3554  CG1 VAL   101      11.192  24.568  14.442  1.00 47.59      L31H C
ATOM   3555  CG2 VAL   101       9.920  23.403  16.256  1.00 45.46      L31H C
ATOM   3556  C   VAL   101      11.853  24.264  18.184  1.00 47.82      L31H C
ATOM   3557  O   VAL   101      11.604  23.167  18.691  1.00 48.23      L31H O
ATOM   3558  N   PHE   102      11.841  25.399  18.876  1.00 46.37      L31H N
ATOM   3559  CA  PHE   102      11.718  25.388  20.318  1.00 47.35      L31H C
ATOM   3560  CB  PHE   102      12.631  26.421  20.932  1.00 44.20      L31H C
ATOM   3561  CG  PHE   102      14.061  26.242  20.590  1.00 41.82      L31H C
ATOM   3562  CD1 PHE   102      14.551  26.688  19.380  1.00 39.65      L31H C
ATOM   3563  CD2 PHE   102      14.944  25.730  21.522  1.00 40.97      L31H C
ATOM   3564  CE1 PHE   102      15.893  26.641  19.113  1.00 38.50      L31H C
ATOM   3565  CE2 PHE   102      16.292  25.680  21.260  1.00 38.80      L31H C
ATOM   3566  CZ  PHE   102      16.769  26.139  20.057  1.00 39.41      L31H C
ATOM   3567  C   PHE   102      10.315  25.646  20.828  1.00 50.48      L31H C
ATOM   3568  O   PHE   102       9.402  25.958  20.067  1.00 52.07      L31H O
ATOM   3569  N   GLY   103      10.166  25.515  22.140  1.00 53.14      L31H N
ATOM   3570  CA  GLY   103       8.894  25.767  22.779  1.00 55.97      L31H C
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ATOM   3571  C   GLY   103       8.882  27.133  23.425  1.00 57.19      L31H C
ATOM   3572  O   GLY   103       9.941  27.744  23.611  1.00 56.34      L31H O
ATOM   3573  N   GLY   104       7.679  27.606  23.761  1.00 58.19      L31H N
ATOM   3574  CA  GLY   104       7.525  28.924  24.349  1.00 58.63      L31H C
ATOM   3575  C   GLY   104       8.458  29.139  25.523  1.00 59.14      L31H C
ATOM   3576  O   GLY   104       9.114  30.177  25.626  1.00 59.20      L31H O
ATOM   3577  N   GLY   105       8.532  28.151  26.406  1.00 60.02      L31H N
ATOM   3578  CA  GLY   105       9.420  28.257  27.547  1.00 61.89      L31H C
ATOM   3579  C   GLY   105       8.647  28.213  28.850  1.00 63.99      L31H C
ATOM   3580  O   GLY   105       7.412  28.344  28.866  1.00 65.05      L31H O
ATOM   3581  N   THR   106       9.369  28.016  29.951  1.00 63.25      L31H N
ATOM   3582  CA  THR   106       8.743  28.026  31.264  1.00 61.53      L31H C
ATOM   3583  CB  THR   106       8.311  26.620  31.690  1.00 60.33      L31H C
ATOM   3584  OG1 THR   106       7.374  26.095  30.740  1.00 60.09      L31H O
ATOM   3585  CG2 THR   106       7.661  26.667  33.058  1.00 58.71      L31H C
ATOM   3586  C   THR   106       9.652  28.608  32.342  1.00 62.44      L31H C
ATOM   3587  O   THR   106      10.829  28.256  32.458  1.00 61.28      L31H O
ATOM   3588  N   LYS   107       9.087  29.522  33.122  1.00 63.21      L31H N
ATOM   3589  CA  LYS   107       9.755  30.039  34.299  1.00 62.64      L31H C
ATOM   3590  CB  LYS   107       9.161  31.391  34.689  1.00 64.98      L31H C
ATOM   3591  CG  LYS   107       9.643  31.922  36.030  1.00 67.60      L31H C
ATOM   3592  CD  LYS   107      10.962  32.670  35.901  1.00 72.42      L31H C
ATOM   3593  CE  LYS   107      11.213  33.560  37.126  1.00 76.31      L31H C
ATOM   3594  NZ  LYS   107      12.373  34.501  36.969  1.00 79.44      L31H N
ATOM   3595  C   LYS   107       9.543  29.042  35.426  1.00 61.19      L31H C
ATOM   3596  O   LYS   107       8.401  28.794  35.846  1.00 62.04      L31H O
ATOM   3597  N   LEU   108      10.638  28.457  35.904  1.00 57.46      L31H N
ATOM   3598  CA  LEU   108      10.570  27.633  37.102  1.00 53.84      L31H C
ATOM   3599  CB  LEU   108      11.556  26.466  37.040  1.00 54.15      L31H C
ATOM   3600  CG  LEU   108      11.541  25.642  38.328  1.00 50.02      L31H C
ATOM   3601  CD1 LEU   108      10.186  25.017  38.503  1.00 50.06      L31H C
ATOM   3602  CD2 LEU   108      12.597  24.589  38.290  1.00 51.57      L31H C
ATOM   3603  C   LEU   108      10.902  28.478  38.309  1.00 53.02      L31H C
ATOM   3604  O   LEU   108      12.016  28.981  38.440  1.00 51.31      L31H O
ATOM   3605  N   THR   109       9.925  28.629  39.192  1.00 52.73      L31H N
ATOM   3606  CA  THR   109      10.118  29.355  40.440  1.00 52.49      L31H C
ATOM   3607  CB  THR   109       9.000  30.416  40.628  1.00 52.18      L31H C
ATOM   3608  OG1 THR   109       9.260  31.542  39.777  1.00 50.86      L31H O
ATOM   3609  CG2 THR   109       8.918  30.866  42.076  1.00 53.04      L31H C
ATOM   3610  C   THR   109      10.113  28.375  41.617  1.00 53.48      L31H C
ATOM   3611  O   THR   109       9.120  27.680  41.851  1.00 51.71      L31H O
ATOM   3612  N   VAL   110      11.228  28.304  42.342  1.00 54.50      L31H N
ATOM   3613  CA  VAL   110      11.286  27.475  43.546  1.00 56.59      L31H C
ATOM   3614  CB  VAL   110      12.727  26.972  43.863  1.00 55.19      L31H C
ATOM   3615  CG1 VAL   110      12.700  26.076  45.091  1.00 51.04      L31H C
ATOM   3616  CG2 VAL   110      13.302  26.227  42.673  1.00 54.74      L31H C
ATOM   3617  C   VAL   110      10.824  28.353  44.690  1.00 59.73      L31H C
ATOM   3618  O   VAL   110      11.501  29.312  45.047  1.00 59.28      L31H O
ATOM   3619  N   LEU   111       9.670  28.027  45.260  1.00 63.84      L31H N
ATOM   3620  CA  LEU   111       8.976  28.955  46.141  1.00 69.11      L31H C
ATOM   3621  CB  LEU   111       7.519  28.520  46.288  1.00 64.19      L31H C
ATOM   3622  CG  LEU   111       6.819  28.414  44.932  1.00 60.56      L31H C
ATOM   3623  CD1 LEU   111       5.553  27.609  45.066  1.00 58.84      L31H C
ATOM   3624  CD2 LEU   111       6.546  29.800  44.386  1.00 55.98      L31H C
ATOM   3625  C   LEU   111       9.645  29.066  47.503  1.00 75.53      L31H C
ATOM   3626  O   LEU   111       9.702  28.101  48.259  1.00 78.23      L31H O
ATOM   3627  N   GLY   112      10.159  30.254  47.801  1.00 81.81      L31H N
ATOM   3628  CA  GLY   112      10.745  30.510  49.103  1.00 90.35      L31H C
ATOM   3629  C   GLY   112      10.012  31.619  49.837  1.00 95.92      L31H C
ATOM   3630  O   GLY   112      10.472  32.096  50.874  1.00 95.75      L31H O
ATOM   3631  N   GLN   113       8.870  32.029  49.287  1.00102.79      L31H N
ATOM   3632  CA  GLN   113       8.019  33.058  49.885  1.00110.31      L31H C
ATOM   3633  CB  GLN   113       8.334  34.429  49.280  1.00113.77      L31H C
ATOM   3634  CG  GLN   113       9.794  34.826  49.315  1.00118.20      L31H C
ATOM   3635  CD  GLN   113      10.067  36.072  48.494  1.00120.53      L31H C
ATOM   3636  OE1 GLN   113       9.723  37.186  48.896  1.00121.11      L31H O
ATOM   3637  NE2 GLN   113      10.687  35.890  47.332  1.00121.57      L31H N
ATOM   3638  C   GLN   113       6.545  32.727  49.625  1.00113.82      L31H C
ATOM   3639  O   GLN   113       6.214  32.017  48.679  1.00113.64      L31H O
ATOM   3640  N   PRO   114       5.640  33.248  50.462  1.00117.28      L31H N
ATOM   3641  CD  PRO   114       5.936  34.027  51.673  1.00117.03      L31H C
ATOM   3642  CA  PRO   114       4.210  32.935  50.357  1.00119.60      L31H C
ATOM   3643  CB  PRO   114       3.609  33.570  51.609  1.00119.09      L31H C
ATOM   3644  CG  PRO   114       4.761  33.732  52.542  1.00118.20      L31H C
ATOM   3645  C   PRO   114       3.566  33.484  49.094  1.00121.80      L31H C
ATOM   3646  O   PRO   114       4.237  34.052  48.239  1.00122.55      L31H O
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ATOM   3647  N   LYS  115       2.254  33.304  48.988  1.00123.51      L31H N
ATOM   3648  CA  LYS  115       1.465  33.997  47.980  1.00124.74      L31H C
ATOM   3649  CB  LYS  115      -0.025  33.715  48.192  1.00125.92      L31H C
ATOM   3650  CG  LYS  115      -0.961  34.837  47.740  1.00125.60      L31H C
ATOM   3651  CD  LYS  115      -1.039  34.958  46.224  1.00124.69      L31H C
ATOM   3652  CE  LYS  115      -1.896  33.860  45.617  1.00124.00      L31H C
ATOM   3653  NZ  LYS  115      -3.289  33.894  46.139  1.00123.01      L31H N
ATOM   3654  C   LYS  115       1.722  35.492  48.094  1.00125.34      L31H C
ATOM   3655  O   LYS  115       2.067  35.988  49.166  1.00126.21      L31H O
ATOM   3656  N   ALA  116       1.559  36.206  46.986  1.00125.78      L31H N
ATOM   3657  CA  ALA  116       1.657  37.657  46.994  1.00125.51      L31H C
ATOM   3658  CB  ALA  116       3.117  38.080  46.981  1.00124.48      L31H C
ATOM   3659  C   ALA  116       0.933  38.250  45.794  1.00125.03      L31H C
ATOM   3660  O   ALA  116       1.418  38.161  44.667  1.00125.35      L31H O
ATOM   3661  N   ALA  117      -0.229  38.848  46.043  1.00123.76      L31H N
ATOM   3662  CA  ALA  117      -0.939  39.596  45.011  1.00120.64      L31H C
ATOM   3663  CB  ALA  117      -2.425  39.662  45.336  1.00124.29      L31H C
ATOM   3664  C   ALA  117      -0.352  41.001  44.940  1.00117.39      L31H C
ATOM   3665  O   ALA  117      -0.156  41.654  45.964  1.00117.55      L31H O
ATOM   3666  N   PRO  118      -0.051  41.480  43.724  1.00113.58      L31H N
ATOM   3667  CD  PRO  118      -0.303  40.835  42.425  1.00113.39      L31H C
ATOM   3668  CA  PRO  118       0.628  42.768  43.574  1.00109.03      L31H C
ATOM   3669  CB  PRO  118       1.009  42.804  42.096  1.00110.84      L31H C
ATOM   3670  CG  PRO  118       0.030  41.917  41.436  1.00112.56      L31H C
ATOM   3671  C   PRO  118      -0.236  43.959  43.974  1.00104.68      L31H C
ATOM   3672  O   PRO  118      -1.468  43.915  43.878  1.00103.62      L31H O
ATOM   3673  N   SER  119       0.429  45.015  44.434  1.00 98.81      L31H N
ATOM   3674  CA  SER  119      -0.215  46.302  44.650  1.00 92.70      L31H C
ATOM   3675  CB  SER  119       0.456  47.045  45.805  1.00 93.69      L31H C
ATOM   3676  OG  SER  119       1.866  47.017  45.677  1.00 95.23      L31H O
ATOM   3677  C   SER  119      -0.112  47.119  43.373  1.00 87.43      L31H C
ATOM   3678  O   SER  119       0.908  47.093  42.684  1.00 85.84      L31H O
ATOM   3679  N   VAL  120      -1.180  47.835  43.053  1.00 82.08      L31H N
ATOM   3680  CA  VAL  120      -1.258  48.520  41.780  1.00 78.07      L31H C
ATOM   3681  CB  VAL  120      -2.216  47.795  40.833  1.00 78.62      L31H C
ATOM   3682  CG1 VAL  120      -2.413  48.604  39.569  1.00 79.64      L31H C
ATOM   3683  CG2 VAL  120      -1.661  46.424  40.502  1.00 79.29      L31H C
ATOM   3684  C   VAL  120      -1.725  49.949  41.944  1.00 75.01      L31H C
ATOM   3685  O   VAL  120      -2.768  50.201  42.530  1.00 73.87      L31H O
ATOM   3686  N   THR  121      -0.943  50.888  41.432  1.00 72.79      L31H N
ATOM   3687  CA  THR  121      -1.383  52.276  41.371  1.00 71.71      L31H C
ATOM   3688  CB  THR  121      -0.371  53.227  42.045  1.00 69.80      L31H C
ATOM   3689  OG1 THR  121      -0.160  52.821  43.403  1.00 67.98      L31H O
ATOM   3690  CG2 THR  121      -0.900  54.647  42.039  1.00 69.16      L31H C
ATOM   3691  C   THR  121      -1.543  52.676  39.909  1.00 71.78      L31H C
ATOM   3692  O   THR  121      -0.764  52.242  39.054  1.00 72.43      L31H O
ATOM   3693  N   LEU  122      -2.558  53.487  39.621  1.00 70.64      L31H N
ATOM   3694  CA  LEU  122      -2.806  53.917  38.252  1.00 70.47      L31H C
ATOM   3695  CB  LEU  122      -4.121  53.331  37.737  1.00 69.96      L31H C
ATOM   3696  CG  LEU  122      -4.416  53.622  36.263  1.00 69.35      L31H C
ATOM   3697  CD1 LEU  122      -3.214  53.213  35.412  1.00 68.20      L31H C
ATOM   3698  CD2 LEU  122      -5.663  52.865  35.823  1.00 69.21      L31H C
ATOM   3699  C   LEU  122      -2.848  55.432  38.124  1.00 70.57      L31H C
ATOM   3700  O   LEU  122      -3.630  56.098  38.805  1.00 71.23      L31H O
ATOM   3701  N   PHE  123      -2.008  55.967  37.239  1.00 69.59      L31H N
ATOM   3702  CA  PHE  123      -1.916  57.410  37.026  1.00 67.96      L31H C
ATOM   3703  CB  PHE  123      -0.460  57.865  37.114  1.00 68.86      L31H C
ATOM   3704  CG  PHE  123       0.025  58.076  38.516  1.00 70.50      L31H C
ATOM   3705  CD1 PHE  123       0.808  57.124  39.144  1.00 71.72      L31H C
ATOM   3706  CD2 PHE  123      -0.301  59.230  39.206  1.00 71.39      L31H C
ATOM   3707  CE1 PHE  123       1.256  57.318  40.431  1.00 72.59      L31H C
ATOM   3708  CE2 PHE  123       0.145  59.432  40.495  1.00 71.92      L31H C
ATOM   3709  CZ  PHE  123       0.924  58.475  41.107  1.00 73.15      L31H C
ATOM   3710  C   PHE  123      -2.499  57.859  35.690  1.00 66.49      L31H C
ATOM   3711  O   PHE  123      -2.185  57.293  34.636  1.00 66.68      L31H O
ATOM   3712  N   PRO  124      -3.360  58.890  35.728  1.00 64.71      L31H N
ATOM   3713  CD  PRO  124      -3.896  59.403  36.999  1.00 64.64      L31H C
ATOM   3714  CA  PRO  124      -3.870  59.650  34.579  1.00 62.32      L31H C
ATOM   3715  CB  PRO  124      -4.992  60.483  35.175  1.00 64.01      L31H C
ATOM   3716  CG  PRO  124      -4.579  60.680  36.592  1.00 64.63      L31H C
ATOM   3717  C   PRO  124      -2.792  60.544  33.954  1.00 62.33      L31H C
ATOM   3718  O   PRO  124      -1.774  60.847  34.587  1.00 61.20      L31H O
ATOM   3719  N   PRO  125      -3.002  60.978  32.700  1.00 62.33      L31H N
ATOM   3720  CD  PRO  125      -3.860  60.367  31.677  1.00 62.42      L31H C
ATOM   3721  CA  PRO  125      -2.120  61.991  32.117  1.00 60.75      L31H C
ATOM   3722  CB  PRO  125      -2.604  62.094  30.672  1.00 61.44      L31H C
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ATOM   3723  CG  PRO  125      -3.179  60.762  30.395  1.00 62.48      L31H C
ATOM   3724  C   PRO  125      -2.189  63.329  32.854  1.00 60.75      L31H C
ATOM   3725  O   PRO  125      -3.210  63.680  33.447  1.00 59.69      L31H O
ATOM   3726  N   SER  126      -1.085  64.066  32.819  1.00 60.32      L31H N
ATOM   3727  CA  SER  126      -1.027  65.390  33.416  1.00 58.57      L31H C
ATOM   3728  CB  SER  126       0.402  65.714  33.827  1.00 58.32      L31H C
ATOM   3729  OG  SER  126       1.278  65.586  32.719  1.00 57.04      L31H O
ATOM   3730  C   SER  126      -1.509  66.446  32.436  1.00 57.83      L31H C
ATOM   3731  O   SER  126      -1.473  66.261  31.223  1.00 56.21      L31H O
ATOM   3732  N   SER  127      -1.959  67.563  32.981  1.00 58.56      L31H N
ATOM   3733  CA  SER  127      -2.261  68.716  32.165  1.00 59.40      L31H C
ATOM   3734  CB  SER  127      -2.431  69.952  33.049  1.00 60.04      L31H C
ATOM   3735  OG  SER  127      -2.162  71.136  32.312  1.00 61.03      L31H O
ATOM   3736  C   SER  127      -1.102  68.934  31.207  1.00 59.25      L31H C
ATOM   3737  O   SER  127      -1.295  69.145  30.009  1.00 59.39      L31H O
ATOM   3738  N   GLU  128       0.106  68.878  31.748  1.00 58.37      L31H N
ATOM   3739  CA  GLU  128       1.282  69.233  30.976  1.00 57.49      L31H C
ATOM   3740  CB  GLU  128       2.533  69.082  31.830  1.00 58.17      L31H C
ATOM   3741  CG  GLU  128       2.688  70.142  32.877  1.00 58.94      L31H C
ATOM   3742  CD  GLU  128       4.104  70.643  32.938  1.00 60.68      L31H C
ATOM   3743  OE1 GLU  128       4.807  70.541  31.902  1.00 61.85      L31H O
ATOM   3744  OE2 GLU  128       4.510  71.133  34.016  1.00 60.55      L31H O
ATOM   3745  C   GLU  128       1.417  68.382  29.728  1.00 55.84      L31H C
ATOM   3746  O   GLU  128       1.695  68.900  28.652  1.00 53.53      L31H O
ATOM   3747  N   GLU  129       1.220  67.074  29.883  1.00 55.94      L31H N
ATOM   3748  CA  GLU  129       1.441  66.128  28.796  1.00 55.91      L31H C
ATOM   3749  CB  GLU  129       1.426  64.694  29.316  1.00 57.14      L31H C
ATOM   3750  CG  GLU  129       2.033  63.697  28.336  1.00 59.19      L31H C
ATOM   3751  CD  GLU  129       1.519  62.287  28.538  1.00 60.92      L31H C
ATOM   3752  OE1 GLU  129       2.071  61.358  27.906  1.00 60.62      L31H O
ATOM   3753  OE2 GLU  129       0.560  62.112  29.328  1.00 61.87      L31H O
ATOM   3754  C   GLU  129       0.363  66.285  27.747  1.00 55.89      L31H C
ATOM   3755  O   GLU  129       0.588  66.057  26.556  1.00 54.96      L31H O
ATOM   3756  N   LEU  130      -0.819  66.671  28.203  1.00 56.93      L31H N
ATOM   3757  CA  LEU  130      -1.917  66.963  27.304  1.00 57.51      L31H C
ATOM   3758  CB  LEU  130      -3.198  67.191  28.112  1.00 57.09      L31H C
ATOM   3759  CG  LEU  130      -4.161  65.989  28.097  1.00 58.06      L31H C
ATOM   3760  CD1 LEU  130      -3.374  64.692  27.940  1.00 56.91      L31H C
ATOM   3761  CD2 LEU  130      -4.999  65.970  29.366  1.00 57.58      L31H C
ATOM   3762  C   LEU  130      -1.578  68.185  26.464  1.00 57.33      L31H C
ATOM   3763  O   LEU  130      -1.741  68.172  25.251  1.00 58.08      L31H O
ATOM   3764  N   GLN  131      -1.082  69.231  27.111  1.00 57.04      L31H N
ATOM   3765  CA  GLN  131      -0.650  70.424  26.402  1.00 56.50      L31H C
ATOM   3766  CB  GLN  131      -0.101  71.447  27.389  1.00 56.13      L31H C
ATOM   3767  CG  GLN  131      -1.142  72.340  27.972  1.00 59.45      L31H C
ATOM   3768  CD  GLN  131      -1.880  73.113  26.905  1.00 61.61      L31H C
ATOM   3769  OE1 GLN  131      -3.018  73.544  27.114  1.00 61.71      L31H O
ATOM   3770  NE2 GLN  131      -1.237  73.294  25.746  1.00 60.83      L31H N
ATOM   3771  C   GLN  131       0.412  70.139  25.340  1.00 57.13      L31H C
ATOM   3772  O   GLN  131       0.656  70.959  24.457  1.00 58.38      L31H O
ATOM   3773  N   ALA  132       1.063  68.989  25.431  1.00 56.27      L31H N
ATOM   3774  CA  ALA  132       2.122  68.660  24.493  1.00 54.83      L31H C
ATOM   3775  CB  ALA  132       3.210  67.886  25.199  1.00 53.35      L31H C
ATOM   3776  C   ALA  132       1.517  67.830  23.380  1.00 54.99      L31H C
ATOM   3777  O   ALA  132       2.218  67.361  22.484  1.00 53.23      L31H O
ATOM   3778  N   ASN  133       0.199  67.664  23.465  1.00 56.66      L31H N
ATOM   3779  CA  ASN  133      -0.592  66.906  22.499  1.00 58.35      L31H C
ATOM   3780  CB  ASN  133      -0.380  67.464  21.083  1.00 61.86      L31H C
ATOM   3781  CG  ASN  133      -1.567  67.198  20.149  1.00 63.84      L31H C
ATOM   3782  OD1 ASN  133      -2.663  67.756  20.322  1.00 64.92      L31H O
ATOM   3783  ND2 ASN  133      -1.345  66.348  19.148  1.00 63.87      L31H N
ATOM   3784  C   ASN  133      -0.235  65.419  22.562  1.00 58.50      L31H C
ATOM   3785  O   ASN  133      -0.227  64.719  21.542  1.00 58.17      L31H O
ATOM   3786  N   LYS  134       0.070  64.950  23.772  1.00 57.34      L31H N
ATOM   3787  CA  LYS  134       0.210  63.523  24.045  1.00 55.52      L31H C
ATOM   3788  CB  LYS  134       1.676  63.178  24.266  1.00 54.57      L31H C
ATOM   3789  CG  LYS  134       2.589  63.429  23.065  1.00 53.32      L31H C
ATOM   3790  CD  LYS  134       2.539  62.241  22.131  1.00 53.44      L31H C
ATOM   3791  CE  LYS  134       3.597  62.364  21.049  1.00 52.60      L31H C
ATOM   3792  NZ  LYS  134       4.983  62.412  21.589  1.00 51.62      L31H N
ATOM   3793  C   LYS  134      -0.595  63.141  25.284  1.00 55.69      L31H C
ATOM   3794  O   LYS  134      -1.126  64.002  25.987  1.00 56.06      L31H O
ATOM   3795  N   ALA  135      -0.696  61.845  25.545  1.00 55.25      L31H N
ATOM   3796  CA  ALA  135      -1.287  61.365  26.793  1.00 55.37      L31H C
ATOM   3797  CB  ALA  135      -2.810  61.336  26.681  1.00 53.00      L31H C
ATOM   3798  C   ALA  135      -0.749  59.973  27.120  1.00 55.35      L31H C
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ATOM   3799  O   ALA   135      -0.805  59.060  26.292  1.00 56.00      L31H O
ATOM   3800  N   THR   136      -0.202  59.818  28.319  1.00 54.87      L31H N
ATOM   3801  CA  THR   136       0.250  58.511  28.754  1.00 54.77      L31H C
ATOM   3802  CB  THR   136       1.772  58.447  28.819  1.00 53.90      L31H C
ATOM   3803  OG1 THR   136       2.306  58.645  27.506  1.00 55.20      L31H O
ATOM   3804  CG2 THR   136       2.223  57.086  29.318  1.00 53.01      L31H C
ATOM   3805  C   THR   136      -0.323  58.080  30.095  1.00 55.19      L31H C
ATOM   3806  O   THR   136      -0.081  58.710  31.121  1.00 55.30      L31H O
ATOM   3807  N   LEU   137      -1.091  56.998  30.063  1.00 56.06      L31H N
ATOM   3808  CA  LEU   137      -1.559  56.333  31.267  1.00 57.51      L31H C
ATOM   3809  CB  LEU   137      -2.723  55.396  30.931  1.00 58.17      L31H C
ATOM   3810  CG  LEU   137      -3.954  56.062  30.301  1.00 60.61      L31H C
ATOM   3811  CD1 LEU   137      -5.070  55.038  30.171  1.00 60.26      L31H C
ATOM   3812  CD2 LEU   137      -4.420  57.243  31.159  1.00 60.91      L31H C
ATOM   3813  C   LEU   137      -0.418  55.534  31.885  1.00 58.44      L31H C
ATOM   3814  O   LEU   137       0.310  54.813  31.193  1.00 58.78      L31H O
ATOM   3815  N   VAL   138      -0.258  55.663  33.195  1.00 59.31      L31H N
ATOM   3816  CA  VAL   138       0.802  54.944  33.890  1.00 60.22      L31H C
ATOM   3817  CB  VAL   138       1.685  55.916  34.681  1.00 58.67      L31H C
ATOM   3818  CG1 VAL   138       2.717  55.158  35.481  1.00 57.67      L31H C
ATOM   3819  CG2 VAL   138       2.366  56.864  33.724  1.00 59.34      L31H C
ATOM   3820  C   VAL   138       0.203  53.921  34.841  1.00 62.22      L31H C
ATOM   3821  O   VAL   138      -0.596  54.272  35.705  1.00 62.49      L31H O
ATOM   3822  N   CYS   139       0.577  52.656  34.674  1.00 64.39      L31H N
ATOM   3823  CA  CYS   139       0.107  51.607  35.574  1.00 66.45      L31H C
ATOM   3824  C   CYS   139       1.281  50.907  36.234  1.00 63.76      L31H C
ATOM   3825  O   CYS   139       2.050  50.208  35.569  1.00 63.60      L31H O
ATOM   3826  CB  CYS   139      -0.727  50.580  34.808  1.00 72.30      L31H C
ATOM   3827  SG  CYS   139      -1.495  49.302  35.852  1.00 80.69      L31H S
ATOM   3828  N   LEU   140       1.411  51.086  37.543  1.00 61.42      L31H N
ATOM   3829  CA  LEU   140       2.547  50.527  38.267  1.00 61.83      L31H C
ATOM   3830  CB  LEU   140       3.230  51.624  39.082  1.00 59.77      L31H C
ATOM   3831  CG  LEU   140       3.587  52.822  38.205  1.00 57.02      L31H C
ATOM   3832  CD1 LEU   140       3.474  54.095  38.989  1.00 56.73      L31H C
ATOM   3833  CD2 LEU   140       4.975  52.644  37.659  1.00 58.13      L31H C
ATOM   3834  C   LEU   140       2.139  49.368  39.174  1.00 62.96      L31H C
ATOM   3835  O   LEU   140       1.163  49.458  39.926  1.00 61.87      L31H O
ATOM   3836  N   ILE   141       2.895  48.278  39.080  1.00 64.43      L31H N
ATOM   3837  CA  ILE   141       2.601  47.052  39.812  1.00 65.71      L31H C
ATOM   3838  CB  ILE   141       2.208  45.926  38.839  1.00 65.35      L31H C
ATOM   3839  CG2 ILE   141       2.084  44.602  39.581  1.00 65.06      L31H C
ATOM   3840  CG1 ILE   141       0.905  46.292  38.129  1.00 65.30      L31H C
ATOM   3841  CD1 ILE   141       1.111  46.923  36.776  1.00 65.52      L31H C
ATOM   3842  C   ILE   141       3.813  46.593  40.629  1.00 67.11      L31H C
ATOM   3843  O   ILE   141       4.911  46.420  40.094  1.00 66.16      L31H O
ATOM   3844  N   SER   142       3.614  46.394  41.926  1.00 68.80      L31H N
ATOM   3845  CA  SER   142       4.703  45.941  42.774  1.00 69.99      L31H C
ATOM   3846  CB  SER   142       5.371  47.130  43.465  1.00 69.56      L31H C
ATOM   3847  OG  SER   142       4.482  47.762  44.364  1.00 69.04      L31H O
ATOM   3848  C   SER   142       4.237  44.946  43.824  1.00 72.02      L31H C
ATOM   3849  O   SER   142       3.041  44.825  44.099  1.00 70.57      L31H O
ATOM   3850  N   ASP   143       5.205  44.232  44.395  1.00 75.76      L31H N
ATOM   3851  CA  ASP   143       4.990  43.387  45.561  1.00 79.15      L31H C
ATOM   3852  CB  ASP   143       4.247  44.172  46.652  1.00 79.95      L31H C
ATOM   3853  CG  ASP   143       5.084  45.303  47.236  1.00 81.27      L31H C
ATOM   3854  OD1 ASP   143       6.124  45.652  46.640  1.00 83.43      L31H O
ATOM   3855  OD2 ASP   143       4.703  45.845  48.296  1.00 82.20      L31H O
ATOM   3856  C   ASP   143       4.222  42.122  45.217  1.00 81.34      L31H C
ATOM   3857  O   ASP   143       3.249  41.777  45.886  1.00 81.70      L31H O
ATOM   3858  N   PHE   144       4.654  41.430  44.169  1.00 84.09      L31H N
ATOM   3859  CA  PHE   144       4.002  40.183  43.793  1.00 88.16      L31H C
ATOM   3860  CB  PHE   144       3.227  40.352  42.485  1.00 86.52      L31H C
ATOM   3861  CG  PHE   144       4.030  40.953  41.371  1.00 84.52      L31H C
ATOM   3862  CD1 PHE   144       4.561  40.153  40.376  1.00 83.55      L31H C
ATOM   3863  CD2 PHE   144       4.236  42.319  41.305  1.00 83.02      L31H C
ATOM   3864  CE1 PHE   144       5.282  40.705  39.338  1.00 81.29      L31H C
ATOM   3865  CE2 PHE   144       4.956  42.875  40.269  1.00 81.03      L31H C
ATOM   3866  CZ  PHE   144       5.479  42.069  39.285  1.00 80.66      L31H C
ATOM   3867  C   PHE   144       4.974  39.018  43.670  1.00 91.37      L31H C
ATOM   3868  O   PHE   144       6.103  39.178  43.200  1.00 92.48      L31H O
ATOM   3869  N   TYR   145       4.523  37.846  44.109  1.00 93.82      L31H N
ATOM   3870  CA  TYR   145       5.322  36.629  44.038  1.00 94.58      L31H C
ATOM   3871  CB  TYR   145       6.216  36.500  45.271  1.00 95.09      L31H C
ATOM   3872  CG  TYR   145       7.203  35.361  45.180  1.00 95.11      L31H C
ATOM   3873  CD1 TYR   145       7.035  34.210  45.932  1.00 95.46      L31H C
ATOM   3874  CE1 TYR   145       7.939  33.165  45.850  1.00 95.67      L31H C
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ATOM   3875  CD2 TYR   145       8.306  35.437  44.336  1.00 95.57      L31H C
ATOM   3876  CE2 TYR   145       9.215  34.397  44.245  1.00 95.35      L31H C
ATOM   3877  CZ  TYR   145       9.027  33.263  45.007  1.00 95.54      L31H C
ATOM   3878  OH  TYR   145       9.933  32.230  44.935  1.00 93.89      L31H O
ATOM   3879  C   TYR   145       4.411  35.416  43.947  1.00 96.10      L31H C
ATOM   3880  O   TYR   145       3.405  35.334  44.649  1.00 95.23      L31H O
ATOM   3881  N   PRO   146       4.754  34.459  43.072  1.00 98.97      L31H N
ATOM   3882  CD  PRO   146       3.974  33.236  42.816  1.00102.65      L31H C
ATOM   3883  CA  PRO   146       5.945  34.535  42.217  1.00100.63      L31H C
ATOM   3884  CB  PRO   146       5.996  33.161  41.552  1.00103.16      L31H C
ATOM   3885  CG  PRO   146       4.586  32.692  41.560  1.00103.82      L31H C
ATOM   3886  C   PRO   146       5.880  35.673  41.196  1.00101.63      L31H C
ATOM   3887  O   PRO   146       4.796  36.118  40.814  1.00101.06      L31H O
ATOM   3888  N   GLY   147       7.048  36.141  40.764  1.00103.99      L31H N
ATOM   3889  CA  GLY   147       7.104  37.257  39.837  1.00106.69      L31H C
ATOM   3890  C   GLY   147       6.669  36.921  38.428  1.00107.75      L31H C
ATOM   3891  O   GLY   147       7.498  36.741  37.535  1.00109.16      L31H O
ATOM   3892  N   ALA   148       5.358  36.841  38.231  1.00106.88      L31H N
ATOM   3893  CA  ALA   148       4.796  36.566  36.918  1.00106.36      L31H C
ATOM   3894  CB  ALA   148       4.810  35.069  36.644  1.00106.24      L31H C
ATOM   3895  C   ALA   148       3.375  37.101  36.844  1.00106.36      L31H C
ATOM   3896  O   ALA   148       2.477  36.607  37.521  1.00105.85      L31H O
ATOM   3897  N   VAL   149       3.183  38.118  36.014  1.00107.52      L31H N
ATOM   3898  CA  VAL   149       1.890  38.766  35.871  1.00108.34      L31H C
ATOM   3899  CB  VAL   149       1.857  40.102  36.625  1.00108.35      L31H C
ATOM   3900  CG1 VAL   149       1.858  39.856  38.121  1.00108.86      L31H C
ATOM   3901  CG2 VAL   149       3.069  40.938  36.237  1.00108.49      L31H C
ATOM   3902  C   VAL   149       1.609  39.033  34.405  1.00108.11      L31H C
ATOM   3903  O   VAL   149       2.528  39.094  33.591  1.00108.80      L31H O
ATOM   3904  N   THR   150       0.332  39.186  34.075  1.00107.28      L31H N
ATOM   3905  CA  THR   150      -0.069  39.576  32.733  1.00106.06      L31H C
ATOM   3906  CB  THR   150      -0.798  38.439  32.011  1.00105.51      L31H C
ATOM   3907  OG1 THR   150       0.127  37.386  31.725  1.00104.72      L31H O
ATOM   3908  CG2 THR   150      -1.410  38.938  30.715  1.00104.90      L31H C
ATOM   3909  C   THR   150      -1.005  40.760  32.824  1.00105.63      L31H C
ATOM   3910  O   THR   150      -1.935  40.756  33.628  1.00106.26      L31H O
ATOM   3911  N   VAL   151      -0.758  41.770  31.995  1.00104.86      L31H N
ATOM   3912  CA  VAL   151      -1.558  42.992  32.022  1.00103.61      L31H C
ATOM   3913  CB  VAL   151      -0.663  44.254  31.995  1.00103.81      L31H C
ATOM   3914  CG1 VAL   151      -1.530  45.510  32.047  1.00103.60      L31H C
ATOM   3915  CG2 VAL   151       0.305  44.226  33.166  1.00102.99      L31H C
ATOM   3916  C   VAL   151      -2.544  43.072  30.861  1.00101.69      L31H C
ATOM   3917  O   VAL   151      -2.207  42.743  29.721  1.00101.44      L31H O
ATOM   3918  N   ALA   152      -3.763  43.506  31.166  1.00 99.44      L31H N
ATOM   3919  CA  ALA   152      -4.775  43.731  30.147  1.00 97.98      L31H C
ATOM   3920  CB  ALA   152      -5.788  42.600  30.161  1.00 96.47      L31H C
ATOM   3921  C   ALA   152      -5.466  45.056  30.423  1.00 97.32      L31H C
ATOM   3922  O   ALA   152      -6.002  45.269  31.511  1.00 98.16      L31H O
ATOM   3923  N   TRP   153      -5.452  45.947  29.436  1.00 95.82      L31H N
ATOM   3924  CA  TRP   153      -6.007  47.285  29.607  1.00 94.83      L31H C
ATOM   3925  CB  TRP   153      -5.149  48.311  28.857  1.00 89.61      L31H C
ATOM   3926  CG  TRP   153      -3.833  48.634  29.516  1.00 83.09      L31H C
ATOM   3927  CD2 TRP   153      -3.550  49.759  30.356  1.00 80.15      L31H C
ATOM   3928  CE2 TRP   153      -2.188  49.674  30.724  1.00 78.88      L31H C
ATOM   3929  CE3 TRP   153      -4.315  50.829  30.832  1.00 78.10      L31H C
ATOM   3930  CD1 TRP   153      -2.665  47.932  29.413  1.00 81.18      L31H C
ATOM   3931  NE1 TRP   153      -1.672  48.551  30.136  1.00 78.50      L31H N
ATOM   3932  CZ2 TRP   153      -1.576  50.618  31.544  1.00 78.36      L31H C
ATOM   3933  CZ3 TRP   153      -3.709  51.765  31.645  1.00 78.62      L31H C
ATOM   3934  CH2 TRP   153      -2.349  51.654  31.994  1.00 79.04      L31H C
ATOM   3935  C   TRP   153      -7.444  47.361  29.102  1.00 97.21      L31H C
ATOM   3936  O   TRP   153      -7.772  46.810  28.054  1.00 97.31      L31H O
ATOM   3937  N   LYS   154      -8.301  48.046  29.846  1.00100.61      L31H N
ATOM   3938  CA  LYS   154      -9.645  48.314  29.366  1.00104.68      L31H C
ATOM   3939  CB  LYS   154     -10.688  47.950  30.422  1.00104.13      L31H C
ATOM   3940  CG  LYS   154     -10.824  46.462  30.667  1.00104.81      L31H C
ATOM   3941  CD  LYS   154     -10.929  45.698  29.358  1.00106.43      L31H C
ATOM   3942  CE  LYS   154     -10.925  44.196  29.604  1.00108.33      L31H C
ATOM   3943  NZ  LYS   154      -9.774  43.768  30.447  1.00112.27      L31H N
ATOM   3944  C   LYS   154      -9.791  49.775  29.011  1.00108.13      L31H C
ATOM   3945  O   LYS   154      -9.726  50.643  29.883  1.00108.54      L31H O
ATOM   3946  N   ALA   155      -9.987  50.038  27.723  1.00112.59      L31H N
ATOM   3947  CA  ALA   155     -10.381  51.360  27.258  1.00115.22      L31H C
ATOM   3948  CB  ALA   155     -10.121  51.488  25.761  1.00118.02      L31H C
ATOM   3949  C   ALA   155     -11.865  51.528  27.555  1.00116.52      L31H C
ATOM   3950  O   ALA   155     -12.712  50.944  26.879  1.00115.14      L31H O
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ATOM   3951  N   ASP   156     -12.170  52.324  28.575  1.00118.91      L31H N
ATOM   3952  CA  ASP   156     -13.519  52.392  29.120  1.00122.96      L31H C
ATOM   3953  CB  ASP   156     -14.509  52.810  28.028  1.00123.77      L31H C
ATOM   3954  CG  ASP   156     -15.840  53.254  28.589  1.00125.04      L31H C
ATOM   3955  OD1 ASP   156     -16.106  54.475  28.587  1.00126.49      L31H O
ATOM   3956  OD2 ASP   156     -16.619  52.382  29.031  1.00126.62      L31H O
ATOM   3957  C   ASP   156     -13.885  51.016  29.676  1.00125.66      L31H C
ATOM   3958  O   ASP   156     -13.474  50.653  30.779  1.00124.90      L31H O
ATOM   3959  N   SER   157     -14.649  50.251  28.905  1.00129.31      L31H N
ATOM   3960  CA  SER   157     -14.951  48.869  29.257  1.00133.75      L31H C
ATOM   3961  CB  SER   157     -16.420  48.739  29.660  1.00133.49      L31H C
ATOM   3962  OG  SER   157     -16.729  49.610  30.736  1.00132.03      L31H O
ATOM   3963  C   SER   157     -14.651  47.984  28.055  1.00136.62      L31H C
ATOM   3964  O   SER   157     -14.649  46.755  28.146  1.00137.92      L31H O
ATOM   3965  N   SER   158     -14.401  48.632  26.923  1.00138.76      L31H N
ATOM   3966  CA  SER   158     -13.950  47.944  25.725  1.00139.39      L31H C
ATOM   3967  CB  SER   158     -14.216  48.808  24.492  1.00138.66      L31H C
ATOM   3968  OG  SER   158     -15.557  49.263  24.468  1.00135.17      L31H O
ATOM   3969  C   SER   158     -12.456  47.666  25.845  1.00140.86      L31H C
ATOM   3970  O   SER   158     -11.651  48.583  26.018  1.00142.39      L31H O
ATOM   3971  N   PRO   159     -12.071  46.386  25.760  1.00141.72      L31H N
ATOM   3972  CD  PRO   159     -13.033  45.270  25.740  1.00142.52      L31H C
ATOM   3973  CA  PRO   159     -10.681  45.914  25.858  1.00142.97      L31H C
ATOM   3974  CB  PRO   159     -10.818  44.391  25.810  1.00142.77      L31H C
ATOM   3975  CG  PRO   159     -12.225  44.122  26.262  1.00142.30      L31H C
ATOM   3976  C   PRO   159      -9.746  46.433  24.755  1.00144.51      L31H C
ATOM   3977  O   PRO   159      -9.762  45.930  23.629  1.00144.32      L31H O
ATOM   3978  N   VAL   160      -8.920  47.423  25.088  1.00145.75      L31H N
ATOM   3979  CA  VAL   160      -7.951  47.968  24.140  1.00146.54      L31H C
ATOM   3980  CB  VAL   160      -7.373  49.294  24.640  1.00145.44      L31H C
ATOM   3981  CG1 VAL   160      -6.897  49.139  26.068  1.00144.76      L31H C
ATOM   3982  CG2 VAL   160      -6.214  49.715  23.754  1.00144.65      L31H C
ATOM   3983  C   VAL   160      -6.786  47.017  23.892  1.00147.40      L31H C
ATOM   3984  O   VAL   160      -6.278  46.393  24.820  1.00149.15      L31H O
ATOM   3985  N   LYS   161      -6.360  46.921  22.637  1.00147.17      L31H N
ATOM   3986  CA  LYS   161      -5.296  45.995  22.257  1.00146.00      L31H C
ATOM   3987  CB  LYS   161      -5.794  45.039  21.166  1.00149.06      L31H C
ATOM   3988  CG  LYS   161      -7.090  44.313  21.506  1.00152.58      L31H C
ATOM   3989  CD  LYS   161      -6.931  43.408  22.719  1.00155.68      L31H C
ATOM   3990  CE  LYS   161      -8.256  42.765  23.106  1.00157.97      L31H C
ATOM   3991  NZ  LYS   161      -8.862  42.009  21.976  1.00159.35      L31H N
ATOM   3992  C   LYS   161      -4.035  46.712  21.766  1.00143.65      L31H C
ATOM   3993  O   LYS   161      -2.954  46.121  21.722  1.00142.97      L31H O
ATOM   3994  N   ALA   162      -4.175  47.981  21.391  1.00141.12      L31H N
ATOM   3995  CA  ALA   162      -3.054  48.748  20.855  1.00137.51      L31H C
ATOM   3996  CB  ALA   162      -3.354  49.183  19.427  1.00139.76      L31H C
ATOM   3997  C   ALA   162      -2.743  49.966  21.716  1.00134.24      L31H C
ATOM   3998  O   ALA   162      -3.619  50.502  22.395  1.00132.85      L31H O
ATOM   3999  N   GLY   163      -1.489  50.399  21.680  1.00130.13      L31H N
ATOM   4000  CA  GLY   163      -1.082  51.547  22.469  1.00125.56      L31H C
ATOM   4001  C   GLY   163      -0.547  51.148  23.829  1.00122.13      L31H C
ATOM   4002  O   GLY   163      -0.385  51.983  24.719  1.00122.41      L31H O
ATOM   4003  N   VAL   164      -0.274  49.859  23.989  1.00117.65      L31H N
ATOM   4004  CA  VAL   164       0.178  49.329  25.264  1.00112.18      L31H C
ATOM   4005  CB  VAL   164      -0.770  48.246  25.771  1.00111.56      L31H C
ATOM   4006  CG1 VAL   164      -0.274  47.711  27.098  1.00110.84      L31H C
ATOM   4007  CG2 VAL   164      -2.169  48.805  25.897  1.00110.72      L31H C
ATOM   4008  C   VAL   164       1.563  48.721  25.145  1.00108.74      L31H C
ATOM   4009  O   VAL   164       1.923  48.178  24.104  1.00109.80      L31H O
ATOM   4010  N   GLU   165       2.340  48.810  26.214  1.00103.31      L31H N
ATOM   4011  CA  GLU   165       3.605  48.099  26.271  1.00 97.93      L31H C
ATOM   4012  CB  GLU   165       4.683  48.891  25.534  1.00 99.42      L31H C
ATOM   4013  CG  GLU   165       5.841  48.038  25.045  1.00102.44      L31H C
ATOM   4014  CD  GLU   165       6.165  48.284  23.583  1.00104.36      L31H C
ATOM   4015  OE1 GLU   165       6.303  47.296  22.829  1.00105.42      L31H O
ATOM   4016  OE2 GLU   165       6.281  49.467  23.191  1.00105.49      L31H O
ATOM   4017  C   GLU   165       4.004  47.874  27.725  1.00 93.34      L31H C
ATOM   4018  O   GLU   165       4.181  48.826  28.484  1.00 92.89      L31H O
ATOM   4019  N   THR   166       4.132  46.610  28.113  1.00 87.51      L31H N
ATOM   4020  CA  THR   166       4.417  46.263  29.498  1.00 82.22      L31H C
ATOM   4021  CB  THR   166       3.504  45.138  29.970  1.00 79.71      L31H C
ATOM   4022  OG1 THR   166       2.136  45.535  29.821  1.00 77.59      L31H O
ATOM   4023  CG2 THR   166       3.784  44.814  31.415  1.00 78.67      L31H C
ATOM   4024  C   THR   166       5.862  45.805  29.647  1.00 81.16      L31H C
ATOM   4025  O   THR   166       6.469  45.332  28.687  1.00 81.15      L31H O
ATOM   4026  N   THR   167       6.425  45.953  30.842  1.00 79.71      L31H N
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ATOM	4027	CA	THR	167	7.755	45.411	31.101	1.00	78.78	L31H C
ATOM	4028	CB	THR	167	8.555	46.266	32.125	1.00	77.41	L31H C
ATOM	4029	OG1	THR	167	7.949	46.166	33.418	1.00	75.88	L31H O
ATOM	4030	CG2	THR	167	8.588	47.723	31.701	1.00	75.26	L31H C
ATOM	4031	C	THR	167	7.640	43.990	31.650	1.00	79.22	L31H C
ATOM	4032	O	THR	167	6.599	43.588	32.174	1.00	79.15	L31H O
ATOM	4033	N	THR	168	8.714	43.226	31.515	1.00	79.89	L31H N
ATOM	4034	CA	THR	168	8.823	41.965	32.220	1.00	80.51	L31H C
ATOM	4035	CB	THR	168	9.904	41.072	31.594	1.00	80.02	L31H C
ATOM	4036	OG1	THR	168	11.170	41.744	31.649	1.00	80.02	L31H O
ATOM	4037	CG2	THR	168	9.553	40.758	30.139	1.00	78.70	L31H C
ATOM	4038	C	THR	168	9.194	42.275	33.664	1.00	81.33	L31H C
ATOM	4039	O	THR	168	10.021	43.149	33.937	1.00	80.31	L31H O
ATOM	4040	N	PRO	169	8.570	41.565	34.609	1.00	82.60	L31H N
ATOM	4041	CD	PRO	169	7.606	40.479	34.373	1.00	80.47	L31H C
ATOM	4042	CA	PRO	169	8.725	41.854	36.035	1.00	85.29	L31H C
ATOM	4043	CB	PRO	169	7.885	40.775	36.706	1.00	82.01	L31H C
ATOM	4044	CG	PRO	169	6.876	40.399	35.673	1.00	79.24	L31H C
ATOM	4045	C	PRO	169	10.179	41.824	36.483	1.00	89.48	L31H C
ATOM	4046	O	PRO	169	11.051	41.286	35.797	1.00	91.81	L31H O
ATOM	4047	N	SER	170	10.437	42.412	37.642	1.00	92.27	L31H N
ATOM	4048	CA	SER	170	11.800	42.582	38.105	1.00	94.31	L31H C
ATOM	4049	CB	SER	170	12.258	44.010	37.807	1.00	94.07	L31H C
ATOM	4050	OG	SER	170	13.452	44.326	38.498	1.00	89.69	L31H O
ATOM	4051	C	SER	170	11.886	42.300	39.596	1.00	96.61	L31H C
ATOM	4052	O	SER	170	10.964	42.609	40.347	1.00	97.99	L31H O
ATOM	4053	N	LYS	171	12.996	41.707	40.020	1.00	98.61	L31H N
ATOM	4054	CA	LYS	171	13.235	41.485	41.439	1.00	99.91	L31H C
ATOM	4055	CB	LYS	171	14.609	40.856	41.664	1.00	100.64	L31H C
ATOM	4056	CG	LYS	171	14.768	39.457	41.092	1.00	103.49	L31H C
ATOM	4057	CD	LYS	171	14.183	38.400	42.018	1.00	108.13	L31H C
ATOM	4058	CE	LYS	171	14.654	37.002	41.624	1.00	111.94	L31H C
ATOM	4059	NZ	LYS	171	14.016	35.924	42.438	1.00	117.51	L31H N
ATOM	4060	C	LYS	171	13.173	42.814	42.164	1.00	99.93	L31H C
ATOM	4061	O	LYS	171	13.773	43.791	41.727	1.00	99.84	L31H O
ATOM	4062	N	GLN	172	12.443	42.848	43.271	1.00	100.47	L31H N
ATOM	4063	CA	GLN	172	12.382	44.039	44.101	1.00	101.48	L31H C
ATOM	4064	CB	GLN	172	11.002	44.178	44.740	1.00	98.99	L31H C
ATOM	4065	CG	GLN	172	9.949	44.798	43.840	1.00	94.44	L31H C
ATOM	4066	CD	GLN	172	8.802	45.394	44.630	1.00	91.26	L31H C
ATOM	4067	OE1	GLN	172	7.771	44.757	44.825	1.00	89.71	L31H O
ATOM	4068	NE2	GLN	172	8.980	46.626	45.093	1.00	89.08	L31H N
ATOM	4069	C	GLN	172	13.436	43.996	45.196	1.00	103.90	L31H C
ATOM	4070	O	GLN	172	13.942	42.926	45.546	1.00	104.53	L31H O
ATOM	4071	N	SER	173	13.763	45.167	45.735	1.00	106.09	L31H N
ATOM	4072	CA	SER	173	14.639	45.244	46.892	1.00	107.30	L31H C
ATOM	4073	CB	SER	173	14.693	46.685	47.431	1.00	108.22	L31H C
ATOM	4074	OG	SER	173	13.400	47.231	47.635	1.00	109.67	L31H O
ATOM	4075	C	SER	173	14.121	44.284	47.960	1.00	107.59	L31H C
ATOM	4076	O	SER	173	14.899	43.592	48.609	1.00	107.41	L31H O
ATOM	4077	N	ASN	174	12.802	44.221	48.118	1.00	108.22	L31H N
ATOM	4078	CA	ASN	174	12.195	43.317	49.091	1.00	109.03	L31H C
ATOM	4079	CB	ASN	174	10.791	43.792	49.460	1.00	111.08	L31H C
ATOM	4080	CG	ASN	174	9.730	43.232	48.535	1.00	113.10	L31H C
ATOM	4081	OD1	ASN	174	9.974	43.021	47.348	1.00	114.08	L31H O
ATOM	4082	ND2	ASN	174	8.542	42.986	49.077	1.00	114.85	L31H N
ATOM	4083	C	ASN	174	12.115	41.888	48.559	1.00	108.23	L31H C
ATOM	4084	O	ASN	174	11.580	41.004	49.227	1.00	108.50	L31H O
ATOM	4085	N	ASN	175	12.624	41.672	47.350	1.00	106.34	L31H N
ATOM	4086	CA	ASN	175	12.779	40.328	46.804	1.00	103.04	L31H C
ATOM	4087	CB	ASN	175	13.367	39.407	47.868	1.00	105.63	L31H C
ATOM	4088	CG	ASN	175	13.858	38.101	47.294	1.00	108.23	L31H C
ATOM	4089	OD1	ASN	175	13.064	37.240	46.912	1.00	110.12	L31H O
ATOM	4090	ND2	ASN	175	15.175	37.942	47.226	1.00	108.79	L31H N
ATOM	4091	C	ASN	175	11.488	39.719	46.243	1.00	98.67	L31H C
ATOM	4092	O	ASN	175	11.485	38.590	45.754	1.00	98.94	L31H O
ATOM	4093	N	LYS	176	10.391	40.467	46.316	1.00	92.52	L31H N
ATOM	4094	CA	LYS	176	9.232	40.193	45.467	1.00	84.85	L31H C
ATOM	4095	CB	LYS	176	7.957	40.777	46.076	1.00	82.79	L31H C
ATOM	4096	CG	LYS	176	7.607	40.260	47.446	1.00	78.63	L31H C
ATOM	4097	CD	LYS	176	6.213	40.708	47.808	1.00	76.31	L31H C
ATOM	4098	CE	LYS	176	5.944	40.605	49.289	1.00	74.93	L31H C
ATOM	4099	NZ	LYS	176	4.626	41.208	49.610	1.00	73.02	L31H N
ATOM	4100	C	LYS	176	9.475	40.837	44.101	1.00	81.53	L31H C
ATOM	4101	O	LYS	176	10.603	41.215	43.777	1.00	79.55	L31H O
ATOM	4102	N	TYR	177	8.422	40.980	43.303	1.00	78.56	L31H N

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ATOM   4103  CA  TYR   177       8.594  41.559  41.979  1.00 75.70      L31H C
ATOM   4104  CB  TYR   177       8.269  40.518  40.916  1.00 75.16      L31H C
ATOM   4105  CG  TYR   177       9.271  39.394  40.898  1.00 76.26      L31H C
ATOM   4106  CD1 TYR   177      10.276  39.349  39.948  1.00 75.58      L31H C
ATOM   4107  CE1 TYR   177      11.193  38.328  39.940  1.00 76.47      L31H C
ATOM   4108  CD2 TYR   177       9.216  38.380  41.843  1.00 76.97      L31H C
ATOM   4109  CE2 TYR   177      10.126  37.357  41.839  1.00 76.87      L31H C
ATOM   4110  CZ  TYR   177      11.110  37.333  40.888  1.00 76.76      L31H C
ATOM   4111  OH  TYR   177      12.005  36.293  40.877  1.00 79.13      L31H O
ATOM   4112  C   TYR   177       7.812  42.834  41.716  1.00 73.16      L31H C
ATOM   4113  O   TYR   177       6.801  43.111  42.357  1.00 73.48      L31H O
ATOM   4114  N   ALA   178       8.318  43.607  40.762  1.00 69.40      L31H N
ATOM   4115  CA  ALA   178       7.723  44.872  40.352  1.00 65.29      L31H C
ATOM   4116  CB  ALA   178       8.625  46.036  40.787  1.00 64.49      L31H C
ATOM   4117  C   ALA   178       7.592  44.852  38.833  1.00 62.74      L31H C
ATOM   4118  O   ALA   178       8.361  44.165  38.156  1.00 62.95      L31H O
ATOM   4119  N   ALA   179       6.626  45.596  38.303  1.00 59.52      L31H N
ATOM   4120  CA  ALA   179       6.476  45.723  36.858  1.00 57.65      L31H C
ATOM   4121  CB  ALA   179       5.974  44.401  36.267  1.00 57.21      L31H C
ATOM   4122  C   ALA   179       5.532  46.863  36.473  1.00 56.23      L31H C
ATOM   4123  O   ALA   179       4.528  47.116  37.144  1.00 56.72      L31H O
ATOM   4124  N   SER   180       5.854  47.546  35.384  1.00 53.90      L31H N
ATOM   4125  CA  SER   180       5.019  48.640  34.912  1.00 53.09      L31H C
ATOM   4126  CB  SER   180       5.809  49.954  34.967  1.00 51.19      L31H C
ATOM   4127  OG  SER   180       7.114  49.791  34.444  1.00 48.03      L31H O
ATOM   4128  C   SER   180       4.470  48.407  33.501  1.00 53.56      L31H C
ATOM   4129  O   SER   180       5.167  47.884  32.629  1.00 52.95      L31H O
ATOM   4130  N   SER   181       3.216  48.796  33.288  1.00 58.75      L31H N
ATOM   4131  CA  SER   181       2.627  48.783  31.956  1.00 56.72      L31H C
ATOM   4132  CB  SER   181       1.489  47.762  31.910  1.00 58.34      L31H C
ATOM   4133  OG  SER   181       1.028  47.549  30.587  1.00 61.46      L31H O
ATOM   4134  C   SER   181       2.091  50.181  31.607  1.00 58.40      L31H C
ATOM   4135  O   SER   181       1.392  50.804  32.417  1.00 58.88      L31H O
ATOM   4136  N   TYR   182       2.415  50.674  30.410  1.00 59.24      L31H N
ATOM   4137  CA  TYR   182       1.992  52.020  29.994  1.00 60.58      L31H C
ATOM   4138  CB  TYR   182       3.190  52.839  29.526  1.00 59.97      L31H C
ATOM   4139  CG  TYR   182       4.257  53.031  30.567  1.00 61.07      L31H C
ATOM   4140  CD1 TYR   182       5.161  52.024  30.852  1.00 61.25      L31H C
ATOM   4141  CE1 TYR   182       6.159  52.208  31.783  1.00 63.96      L31H C
ATOM   4142  CD2 TYR   182       4.377  54.231  31.243  1.00 62.41      L31H C
ATOM   4143  CE2 TYR   182       5.369  54.430  32.177  1.00 64.17      L31H C
ATOM   4144  CZ  TYR   182       6.264  53.417  32.449  1.00 65.02      L31H C
ATOM   4145  OH  TYR   182       7.264  53.619  33.386  1.00 64.46      L31H O
ATOM   4146  C   TYR   182       0.971  51.977  28.863  1.00 61.39      L31H C
ATOM   4147  O   TYR   182       1.008  51.070  28.025  1.00 61.61      L31H O
ATOM   4148  N   LEU   183       0.077  52.965  28.831  1.00 61.88      L31H N
ATOM   4149  CA  LEU   183      -0.935  53.053  27.777  1.00 63.70      L31H C
ATOM   4150  CB  LEU   183      -2.328  52.847  28.354  1.00 62.49      L31H C
ATOM   4151  CG  LEU   183      -3.392  52.953  27.265  1.00 61.82      L31H C
ATOM   4152  CD1 LEU   183      -2.992  52.064  26.097  1.00 60.30      L31H C
ATOM   4153  CD2 LEU   183      -4.749  52.545  27.815  1.00 63.19      L31H C
ATOM   4154  C   LEU   183      -0.930  54.384  27.044  1.00 65.27      L31H C
ATOM   4155  O   LEU   183      -1.397  55.384  27.587  1.00 66.49      L31H O
ATOM   4156  N   SER   184      -0.439  54.390  25.805  1.00 66.26      L31H N
ATOM   4157  CA  SER   184      -0.326  55.629  25.029  1.00 65.98      L31H C
ATOM   4158  CB  SER   184       0.806  55.524  23.999  1.00 64.96      L31H C
ATOM   4159  OG  SER   184       2.085  55.477  24.613  1.00 60.60      L31H O
ATOM   4160  C   SER   184      -1.611  55.993  24.299  1.00 66.76      L31H C
ATOM   4161  O   SER   184      -2.030  55.295  23.379  1.00 67.45      L31H O
ATOM   4162  N   LEU   185      -2.217  57.104  24.705  1.00 67.44      L31H N
ATOM   4163  CA  LEU   185      -3.396  57.644  24.035  1.00 66.92      L31H C
ATOM   4164  CB  LEU   185      -4.493  57.932  25.054  1.00 65.96      L31H C
ATOM   4165  CG  LEU   185      -5.146  56.727  25.720  1.00 65.79      L31H C
ATOM   4166  CD1 LEU   185      -6.109  57.201  26.794  1.00 65.42      L31H C
ATOM   4167  CD2 LEU   185      -5.871  55.903  24.673  1.00 66.19      L31H C
ATOM   4168  C   LEU   185      -3.103  58.930  23.262  1.00 67.86      L31H C
ATOM   4169  O   LEU   185      -1.968  59.419  23.227  1.00 66.43      L31H O
ATOM   4170  N   THR   186      -4.149  59.461  22.637  1.00 69.39      L31H N
ATOM   4171  CA  THR   186      -4.165  60.833  22.143  1.00 70.50      L31H C
ATOM   4172  CB  THR   186      -4.752  60.912  20.739  1.00 71.20      L31H C
ATOM   4173  OG1 THR   186      -6.127  60.504  20.780  1.00 73.03      L31H O
ATOM   4174  CG2 THR   186      -3.981  60.006  19.792  1.00 70.92      L31H C
ATOM   4175  C   THR   186      -5.093  61.591  23.071  1.00 72.25      L31H C
ATOM   4176  O   THR   186      -6.005  61.002  23.648  1.00 71.28      L31H O
ATOM   4177  N   PRO   187      -4.882  62.907  23.227  1.00 74.63      L31H N
ATOM   4178  CD  PRO   187      -3.868  63.776  22.607  1.00 75.83      L31H C
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ATOM   4179  CA  PRO   187      -5.702  63.639  24.193  1.00 77.83      L31H C
ATOM   4180  CB  PRO   187      -5.257  65.091  24.016  1.00 76.66      L31H C
ATOM   4181  CG  PRO   187      -3.865  64.986  23.509  1.00 75.65      L31H C
ATOM   4182  C   PRO   187      -7.183  63.447  23.912  1.00 81.70      L31H C
ATOM   4183  O   PRO   187      -7.983  63.246  24.828  1.00 82.51      L31H O
ATOM   4184  N   GLU   188      -7.546  63.487  22.638  1.00 86.40      L31H N
ATOM   4185  CA  GLU   188      -8.949  63.405  22.266  1.00 90.69      L31H C
ATOM   4186  CB  GLU   188      -9.095  63.677  20.764  1.00 93.16      L31H C
ATOM   4187  CG  GLU   188      -8.586  65.065  20.335  1.00 96.35      L31H C
ATOM   4188  CD  GLU   188      -7.112  65.069  19.929  1.00 98.21      L31H C
ATOM   4189  OE1 GLU   188      -6.420  66.079  20.197  1.00 98.38      L31H O
ATOM   4190  OE2 GLU   188      -6.648  64.066  19.337  1.00 99.43      L31H O
ATOM   4191  C   GLU   188      -9.540  62.037  22.646  1.00 91.37      L31H C
ATOM   4192  O   GLU   188     -10.648  61.957  23.185  1.00 91.61      L31H O
ATOM   4193  N   GLN   189      -8.790  60.969  22.385  1.00 91.22      L31H N
ATOM   4194  CA  GLN   189      -9.160  59.642  22.872  1.00 90.89      L31H C
ATOM   4195  CB  GLN   189      -8.070  58.625  22.520  1.00 90.99      L31H C
ATOM   4196  CG  GLN   189      -8.070  58.177  21.077  1.00 92.41      L31H C
ATOM   4197  CD  GLN   189      -7.153  56.987  20.842  1.00 94.60      L31H C
ATOM   4198  OE1 GLN   189      -5.963  57.035  21.157  1.00 96.03      L31H O
ATOM   4199  NE2 GLN   189      -7.706  55.910  20.285  1.00 95.79      L31H N
ATOM   4200  C   GLN   189      -9.364  59.655  24.390  1.00 91.43      L31H C
ATOM   4201  O   GLN   189     -10.352  59.129  24.900  1.00 90.47      L31H O
ATOM   4202  N   TRP   190      -8.419  60.263  25.101  1.00 92.68      L31H N
ATOM   4203  CA  TRP   190      -8.422  60.283  26.559  1.00 93.50      L31H C
ATOM   4204  CB  TRP   190      -7.126  60.914  27.068  1.00 89.68      L31H C
ATOM   4205  CG  TRP   190      -7.192  61.346  28.504  1.00 84.95      L31H C
ATOM   4206  CD2 TRP   190      -7.416  60.504  29.642  1.00 81.90      L31H C
ATOM   4207  CE2 TRP   190      -7.386  61.329  30.781  1.00 80.42      L31H C
ATOM   4208  CE3 TRP   190      -7.637  59.132  29.806  1.00 79.74      L31H C
ATOM   4209  CD1 TRP   190      -7.042  62.613  28.988  1.00 82.64      L31H C
ATOM   4210  NE1 TRP   190      -7.157  62.611  30.355  1.00 81.46      L31H N
ATOM   4211  CZ2 TRP   190      -7.570  60.826  32.068  1.00 78.89      L31H C
ATOM   4212  CZ3 TRP   190      -7.818  58.637  31.081  1.00 77.91      L31H C
ATOM   4213  CH2 TRP   190      -7.783  59.482  32.195  1.00 78.15      L31H C
ATOM   4214  C   TRP   190      -9.610  61.045  27.135  1.00 96.42      L31H C
ATOM   4215  O   TRP   190     -10.117  60.709  28.210  1.00 97.82      L31H O
ATOM   4216  N   LYS   191     -10.047  62.077  26.422  1.00 98.25      L31H N
ATOM   4217  CA  LYS   191     -11.203  62.852  26.846  1.00100.49      L31H C
ATOM   4218  CB  LYS   191     -11.041  64.311  26.415  1.00101.18      L31H C
ATOM   4219  CG  LYS   191      -9.788  64.978  26.969  1.00102.90      L31H C
ATOM   4220  CD  LYS   191      -9.867  66.503  26.892  1.00106.93      L31H C
ATOM   4221  CE  LYS   191      -9.818  67.013  25.448  1.00110.65      L31H C
ATOM   4222  NZ  LYS   191      -8.442  67.022  24.860  1.00114.77      L31H N
ATOM   4223  C   LYS   191     -12.477  62.266  26.249  1.00102.09      L31H C
ATOM   4224  O   LYS   191     -13.582  62.683  26.591  1.00101.51      L31H O
ATOM   4225  N   SER   192     -12.306  61.287  25.363  1.00104.72      L31H N
ATOM   4226  CA  SER   192     -13.417  60.682  24.629  1.00106.72      L31H C
ATOM   4227  CB  SER   192     -12.921  60.124  23.295  1.00106.20      L31H C
ATOM   4228  OG  SER   192     -13.778  59.094  22.834  1.00104.28      L31H O
ATOM   4229  C   SER   192     -14.151  59.571  25.381  1.00108.14      L31H C
ATOM   4230  O   SER   192     -15.380  59.490  25.333  1.00109.03      L31H O
ATOM   4231  N   HIS   193     -13.398  58.708  26.057  1.00109.30      L31H N
ATOM   4232  CA  HIS   193     -13.983  57.583  26.780  1.00111.11      L31H C
ATOM   4233  CB  HIS   193     -12.986  56.425  26.830  1.00111.11      L31H C
ATOM   4234  CG  HIS   193     -12.560  55.939  25.476  1.00111.14      L31H C
ATOM   4235  CD2 HIS   193     -12.706  54.734  24.873  1.00111.10      L31H C
ATOM   4236  ND1 HIS   193     -11.882  56.736  24.578  1.00111.45      L31H N
ATOM   4237  CE1 HIS   193     -11.629  56.044  23.480  1.00111.53      L31H C
ATOM   4238  NE2 HIS   193     -12.118  54.826  23.634  1.00111.26      L31H N
ATOM   4239  C   HIS   193     -14.397  57.990  28.195  1.00113.91      L31H C
ATOM   4240  O   HIS   193     -14.072  59.088  28.655  1.00111.98      L31H O
ATOM   4241  N   ARG   194     -15.127  57.112  28.878  1.00118.43      L31H N
ATOM   4242  CA  ARG   194     -15.711  57.466  30.168  1.00123.44      L31H C
ATOM   4243  CB  ARG   194     -16.983  56.645  30.416  1.00125.26      L31H C
ATOM   4244  CG  ARG   194     -18.237  57.237  29.774  1.00128.96      L31H C
ATOM   4245  CD  ARG   194     -18.480  58.659  30.274  1.00132.25      L31H C
ATOM   4246  NE  ARG   194     -19.711  59.247  29.752  1.00136.13      L31H N
ATOM   4247  CZ  ARG   194     -20.074  60.510  29.952  1.00138.35      L31H C
ATOM   4248  NH1 ARG   194     -21.209  60.967  29.442  1.00140.09      L31H N
ATOM   4249  NH2 ARG   194     -19.300  61.319  30.662  1.00140.23      L31H N
ATOM   4250  C   ARG   194     -14.728  57.273  31.315  1.00124.67      L31H C
ATOM   4251  O   ARG   194     -14.760  58.008  32.304  1.00126.55      L31H O
ATOM   4252  N   SER   195     -13.850  56.287  31.169  1.00124.75      L31H N
ATOM   4253  CA  SER   195     -12.792  56.039  32.144  1.00123.68      L31H C
ATOM   4254  CB  SER   195     -13.382  55.629  33.485  1.00124.49      L31H C
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ATOM	4255	OG	SER	195	-13.668	54.242	33.477	1.00123.08	L31H O
ATOM	4256	C	SER	195	-11.930	54.900	31.638	1.00122.61	L31H C
ATOM	4257	O	SER	195	-12.385	54.081	30.847	1.00121.79	L31H O
ATOM	4258	N	TYR	196	-10.680	54.840	32.105	1.00122.31	L31H N
ATOM	4259	CA	TYR	196	-9.808	53.740	31.740	1.00122.26	L31H C
ATOM	4260	CB	TYR	196	-8.525	54.277	31.096	1.00126.23	L31H C
ATOM	4261	CG	TYR	196	-8.693	54.755	29.663	1.00130.75	L31H C
ATOM	4262	CD1	TYR	196	-8.219	53.997	28.596	1.00132.15	L31H C
ATOM	4263	CE1	TYR	196	-8.356	54.434	27.290	1.00134.17	L31H C
ATOM	4264	CD2	TYR	196	-9.316	55.968	29.378	1.00132.83	L31H C
ATOM	4265	CE2	TYR	196	-9.457	56.411	28.073	1.00134.67	L31H C
ATOM	4266	CZ	TYR	196	-8.974	55.641	27.036	1.00134.93	L31H C
ATOM	4267	OH	TYR	196	-9.100	56.086	25.742	1.00136.58	L31H O
ATOM	4268	C	TYR	196	-9.463	52.899	32.957	1.00119.64	L31H C
ATOM	4269	O	TYR	196	-9.385	53.410	34.073	1.00119.26	L31H O
ATOM	4270	N	SER	197	-9.265	51.604	32.735	1.00116.33	L31H N
ATOM	4271	CA	SER	197	-8.875	50.705	33.812	1.00112.39	L31H C
ATOM	4272	CB	SER	197	-10.025	49.765	34.160	1.00112.14	L31H C
ATOM	4273	OG	SER	197	-11.131	50.498	34.647	1.00111.27	L31H O
ATOM	4274	C	SER	197	-7.650	49.882	33.461	1.00109.10	L31H C
ATOM	4275	O	SER	197	-7.510	49.400	32.337	1.00110.35	L31H O
ATOM	4276	N	CYS	198	-6.763	49.728	34.437	1.00103.72	L31H N
ATOM	4277	CA	CYS	198	-5.640	48.814	34.309	1.00 98.46	L31H C
ATOM	4278	C	CYS	198	-5.983	47.497	35.017	1.00 98.24	L31H C
ATOM	4279	O	CYS	198	-6.588	47.506	36.093	1.00 96.63	L31H O
ATOM	4280	CB	CYS	198	-4.387	49.441	34.923	1.00 94.19	L31H C
ATOM	4281	SG	CYS	198	-2.880	48.471	34.620	1.00 88.73	L31H S
ATOM	4282	N	GLN	199	-5.613	46.368	34.412	1.00 98.30	L31H N
ATOM	4283	CA	GLN	199	-5.964	45.059	34.964	1.00 98.37	L31H C
ATOM	4284	CB	GLN	199	-7.214	44.521	34.279	1.00 97.13	L31H C
ATOM	4285	CG	GLN	199	-8.459	45.312	34.584	1.00 95.47	L31H C
ATOM	4286	CD	GLN	199	-9.671	44.731	33.914	1.00 93.77	L31H C
ATOM	4287	OE1	GLN	199	-9.569	43.761	33.165	1.00 91.94	L31H O
ATOM	4288	NE2	GLN	199	-10.831	45.320	34.176	1.00 93.03	L31H N
ATOM	4289	C	GLN	199	-4.852	44.027	34.845	1.00 99.27	L31H C
ATOM	4290	O	GLN	199	-4.604	43.481	33.768	1.00 99.69	L31H O
ATOM	4291	N	VAL	200	-4.197	43.748	35.967	1.00 99.34	L31H N
ATOM	4292	CA	VAL	200	-3.076	42.820	35.983	1.00 99.87	L31H C
ATOM	4293	CB	VAL	200	-1.931	43.346	36.881	1.00101.07	L31H C
ATOM	4294	CG1	VAL	200	-0.764	42.369	36.866	1.00100.65	L31H C
ATOM	4295	CG2	VAL	200	-1.484	44.716	36.398	1.00101.41	L31H C
ATOM	4296	C	VAL	200	-3.496	41.434	36.467	1.00 99.94	L31H C
ATOM	4297	O	VAL	200	-4.264	41.295	37.418	1.00 98.21	L31H O
ATOM	4298	N	THR	201	-2.984	40.410	35.798	1.00101.27	L31H N
ATOM	4299	CA	THR	201	-3.321	39.039	36.125	1.00103.26	L31H C
ATOM	4300	CB	THR	201	-3.668	38.261	34.848	1.00103.65	L31H C
ATOM	4301	OG1	THR	201	-4.591	39.028	34.060	1.00105.27	L31H O
ATOM	4302	CG2	THR	201	-4.292	36.918	35.195	1.00104.43	L31H C
ATOM	4303	C	THR	201	-2.121	38.383	36.808	1.00105.17	L31H C
ATOM	4304	O	THR	201	-1.079	38.181	36.185	1.00104.71	L31H O
ATOM	4305	N	HIS	202	-2.265	38.070	38.094	1.00107.14	L31H N
ATOM	4306	CA	HIS	202	-1.243	37.316	38.819	1.00109.43	L31H C
ATOM	4307	CB	HIS	202	-0.565	38.188	39.877	1.00107.76	L31H C
ATOM	4308	CG	HIS	202	0.532	37.488	40.622	1.00105.81	L31H C
ATOM	4309	CD2	HIS	202	1.730	37.012	40.206	1.00104.55	L31H C
ATOM	4310	ND1	HIS	202	0.460	37.210	41.970	1.00104.95	L31H N
ATOM	4311	CE1	HIS	202	1.566	36.595	42.352	1.00104.13	L31H C
ATOM	4312	NE2	HIS	202	2.353	36.464	41.300	1.00103.58	L31H N
ATOM	4313	C	HIS	202	-1.856	36.104	39.500	1.00112.26	L31H C
ATOM	4314	O	HIS	202	-2.775	36.239	40.307	1.00112.96	L31H O
ATOM	4315	N	GLU	203	-1.337	34.924	39.172	1.00115.35	L31H N
ATOM	4316	CA	GLU	203	-1.795	33.681	39.780	1.00117.88	L31H C
ATOM	4317	CB	GLU	203	-1.393	33.627	41.258	1.00119.01	L31H C
ATOM	4318	CG	GLU	203	0.065	33.274	41.501	1.00120.93	L31H C
ATOM	4319	CD	GLU	203	0.423	31.886	40.997	1.00122.12	L31H C
ATOM	4320	OE1	GLU	203	0.839	31.036	41.814	1.00122.96	L31H O
ATOM	4321	OE2	GLU	203	0.288	31.641	39.780	1.00122.34	L31H O
ATOM	4322	C	GLU	203	-3.302	33.509	39.658	1.00118.66	L31H C
ATOM	4323	O	GLU	203	-3.943	32.972	40.561	1.00119.67	L31H O
ATOM	4324	N	GLY	203	-3.866	33.967	38.543	1.00118.23	L31H N
ATOM	4325	CA	GLY	203	-5.292	33.802	38.312	1.00116.73	L31H C
ATOM	4326	C	GLY	203	-6.158	34.822	39.031	1.00116.04	L31H C
ATOM	4327	O	GLY	203	-7.383	34.772	38.938	1.00115.24	L31H O
ATOM	4328	N	SER	205	-5.522	35.741	39.755	1.00116.14	L31H N
ATOM	4329	CA	SER	205	-6.219	36.869	40.372	1.00116.01	L31H C
ATOM	4330	CB	SER	205	-5.801	37.024	41.837	1.00115.22	L31H C

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ATOM   4331  OG  SER   205      -6.212  35.913  42.612  1.00112.64      L31H O
ATOM   4332  C   SER   205      -5.890  38.153  39.623  1.00115.80      L31H C
ATOM   4333  O   SER   205      -4.725  38.435  39.347  1.00116.83      L31H O
ATOM   4334  N   THR   206      -6.920  38.930  39.299  1.00114.25      L31H N
ATOM   4335  CA  THR   206      -6.726  40.193  38.598  1.00111.42      L31H C
ATOM   4336  CB  THR   206      -7.560  40.258  37.306  1.00109.53      L31H C
ATOM   4337  OG1 THR   206      -7.060  39.302  36.364  1.00105.70      L31H O
ATOM   4338  CG2 THR   206      -7.481  41.646  36.694  1.00107.19      L31H C
ATOM   4339  C   THR   206      -7.062  41.408  39.447  1.00110.26      L31H C
ATOM   4340  O   THR   206      -8.218  41.640  39.801  1.00110.98      L31H O
ATOM   4341  N   VAL   207      -6.031  42.182  39.765  1.00108.60      L31H N
ATOM   4342  CA  VAL   207      -6.202  43.479  40.402  1.00107.27      L31H C
ATOM   4343  CB  VAL   207      -4.883  43.963  41.054  1.00106.79      L31H C
ATOM   4344  CG1 VAL   207      -5.067  45.355  41.615  1.00106.40      L31H C
ATOM   4345  CG2 VAL   207      -4.455  43.010  42.157  1.00105.72      L31H C
ATOM   4346  C   VAL   207      -6.606  44.480  39.331  1.00106.10      L31H C
ATOM   4347  O   VAL   207      -6.032  44.488  38.242  1.00105.64      L31H O
ATOM   4348  N   GLU   208      -7.595  45.315  39.638  1.00105.37      L31H N
ATOM   4349  CA  GLU   208      -8.014  46.370  38.721  1.00104.39      L31H C
ATOM   4350  CB  GLU   208      -9.452  46.134  38.250  1.00104.64      L31H C
ATOM   4351  CG  GLU   208      -9.922  47.115  37.182  1.00106.41      L31H C
ATOM   4352  CD  GLU   208     -11.408  47.442  37.281  1.00107.65      L31H C
ATOM   4353  OE1 GLU   208     -12.221  46.736  36.646  1.00108.81      L31H O
ATOM   4354  OE2 GLU   208     -11.758  48.411  37.990  1.00107.97      L31H O
ATOM   4355  C   GLU   208      -7.915  47.733  39.397  1.00103.40      L31H C
ATOM   4356  O   GLU   208      -8.078  47.843  40.611  1.00103.44      L31H O
ATOM   4357  N   LYS   209      -7.632  48.760  38.599  1.00102.70      L31H N
ATOM   4358  CA  LYS   209      -7.599  50.146  39.064  1.00102.80      L31H C
ATOM   4359  CB  LYS   209      -6.167  50.550  39.443  1.00101.95      L31H C
ATOM   4360  CG  LYS   209      -5.551  49.745  40.587  1.00 99.94      L31H C
ATOM   4361  CD  LYS   209      -5.467  50.558  41.876  1.00 98.45      L31H C
ATOM   4362  CE  LYS   209      -5.392  49.647  43.096  1.00 98.07      L31H C
ATOM   4363  NZ  LYS   209      -4.863  50.344  44.300  1.00 97.51      L31H N
ATOM   4364  C   LYS   209      -8.099  51.040  37.932  1.00103.44      L31H C
ATOM   4365  O   LYS   209      -7.729  50.839  36.776  1.00103.27      L31H O
ATOM   4366  N   THR   210      -8.935  52.022  38.258  1.00104.82      L31H N
ATOM   4367  CA  THR   210      -9.552  52.865  37.234  1.00106.53      L31H C
ATOM   4368  CB  THR   210     -11.077  52.683  37.218  1.00104.13      L31H C
ATOM   4369  OG1 THR   210     -11.391  51.309  36.967  1.00101.83      L31H O
ATOM   4370  CG2 THR   210     -11.706  53.554  36.143  1.00102.17      L31H C
ATOM   4371  C   THR   210      -9.263  54.355  37.411  1.00108.67      L31H C
ATOM   4372  O   THR   210      -9.142  54.841  38.535  1.00110.86      L31H O
ATOM   4373  N   VAL   211      -9.161  55.076  36.295  1.00109.56      L31H N
ATOM   4374  CA  VAL   211      -9.075  56.539  36.314  1.00109.07      L31H C
ATOM   4375  CB  VAL   211      -7.650  57.027  35.925  1.00108.56      L31H C
ATOM   4376  CG1 VAL   211      -6.654  56.676  37.022  1.00106.89      L31H C
ATOM   4377  CG2 VAL   211      -7.220  56.385  34.614  1.00107.30      L31H C
ATOM   4378  C   VAL   211     -10.099  57.171  35.360  1.00109.17      L31H C
ATOM   4379  O   VAL   211     -10.532  56.537  34.393  1.00108.84      L31H O
ATOM   4380  N   ALA   212     -10.489  58.415  35.639  1.00110.64      L31H N
ATOM   4381  CA  ALA   212     -11.443  59.126  34.786  1.00113.75      L31H C
ATOM   4382  CB  ALA   212     -12.842  59.003  35.359  1.00114.93      L31H C
ATOM   4383  C   ALA   212     -11.085  60.598  34.611  1.00115.04      L31H C
ATOM   4384  O   ALA   212     -10.995  61.344  35.583  1.00114.62      L31H O
ATOM   4385  N   PRO   213     -10.905  61.036  33.353  1.00115.69      L31H N
ATOM   4386  CD  PRO   213     -11.212  60.200  32.179  1.00111.97      L31H C
ATOM   4387  CA  PRO   213     -10.408  62.364  32.964  1.00117.21      L31H C
ATOM   4388  CB  PRO   213     -10.372  62.312  31.434  1.00113.00      L31H C
ATOM   4389  CG  PRO   213     -11.269  61.194  31.055  1.00111.14      L31H C
ATOM   4390  C   PRO   213     -11.198  63.570  33.459  1.00119.54      L31H C
ATOM   4391  O   PRO   213     -10.971  64.686  32.998  1.00121.28      L31H O
ATOM   4392  N   THR   214     -12.116  63.356  34.395  1.00120.41      L31H N
ATOM   4393  CA  THR   214     -12.907  64.459  34.933  1.00120.39      L31H C
ATOM   4394  CB  THR   214     -14.039  63.943  35.857  1.00121.60      L31H C
ATOM   4395  OG1 THR   214     -14.770  62.911  35.184  1.00119.92      L31H O
ATOM   4396  CG2 THR   214     -15.005  65.073  36.207  1.00119.36      L31H C
ATOM   4397  C   THR   214     -12.003  65.407  35.725  1.00119.41      L31H C
ATOM   4398  O   THR   214     -11.989  66.621  35.410  1.00121.23      L31H O
ATOM   4399  OXT THR   214     -11.314  64.922  36.649  1.00120.59      L31H O
TER    4400       THR   214                                             L31H
ATOM   4401  CB  GLU     1      36.818  24.019  27.876  1.00 91.51      H31H C
ATOM   4402  CG  GLU     1      36.442  25.509  27.850  1.00 97.41      H31H C
ATOM   4403  CD  GLU     1      35.215  25.841  28.707  1.00100.73      H31H C
ATOM   4404  OE1 GLU     1      35.228  26.889  29.392  1.00102.86      H31H O
ATOM   4405  OE2 GLU     1      34.237  25.061  28.693  1.00102.54      H31H O
ATOM   4406  C   GLU     1      36.959  23.770  25.393  1.00 84.82      H31H C
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ATOM   4407  O   GLU     1      37.515  24.326  24.446  1.00 85.09      H31H O
ATOM   4408  N   GLU     1      38.978  24.294  26.678  1.00 88.13      H31H N
ATOM   4409  CA  GLU     1      37.693  23.546  26.704  1.00 87.70      H31H C
ATOM   4410  N   VAL     2      35.705  23.337  25.343  1.00 80.58      H31H N
ATOM   4411  CA  VAL     2      34.873  23.538  24.168  1.00 75.75      H31H C
ATOM   4412  CB  VAL     2      33.865  22.405  24.026  1.00 74.92      H31H C
ATOM   4413  CG1 VAL     2      33.140  22.534  22.704  1.00 73.70      H31H C
ATOM   4414  CG2 VAL     2      34.575  21.067  24.147  1.00 74.46      H31H C
ATOM   4415  C   VAL     2      34.112  24.859  24.255  1.00 72.89      H31H C
ATOM   4416  O   VAL     2      33.272  25.045  25.136  1.00 71.91      H31H O
ATOM   4417  N   GLN     3      34.401  25.767  23.327  1.00 68.98      H31H N
ATOM   4418  CA  GLN     3      33.936  27.139  23.439  1.00 63.56      H31H C
ATOM   4419  CB  GLN     3      35.091  28.014  23.915  1.00 64.18      H31H C
ATOM   4420  CG  GLN     3      34.683  29.323  24.562  1.00 64.34      H31H C
ATOM   4421  CD  GLN     3      35.868  30.013  25.193  1.00 64.70      H31H C
ATOM   4422  OE1 GLN     3      35.795  31.180  25.602  1.00 64.94      H31H O
ATOM   4423  NE2 GLN     3      36.980  29.292  25.276  1.00 64.00      H31H N
ATOM   4424  C   GLN     3      33.375  27.689  22.131  1.00 59.78      H31H C
ATOM   4425  O   GLN     3      33.905  27.416  21.051  1.00 59.58      H31H O
ATOM   4426  N   LEU     4      32.297  28.465  22.244  1.00 55.16      H31H N
ATOM   4427  CA  LEU     4      31.736  29.217  21.122  1.00 51.64      H31H C
ATOM   4428  CB  LEU     4      30.289  28.797  20.884  1.00 48.66      H31H C
ATOM   4429  CG  LEU     4      30.063  27.411  20.293  1.00 46.75      H31H C
ATOM   4430  CD1 LEU     4      28.572  27.169  20.180  1.00 47.44      H31H C
ATOM   4431  CD2 LEU     4      30.716  27.303  18.927  1.00 45.40      H31H C
ATOM   4432  C   LEU     4      31.768  30.694  21.485  1.00 51.24      H31H C
ATOM   4433  O   LEU     4      31.480  31.040  22.627  1.00 51.53      H31H O
ATOM   4434  N   VAL     5      32.112  31.571  20.543  1.00 51.07      H31H N
ATOM   4435  CA  VAL     5      32.099  33.009  20.843  1.00 52.18      H31H C
ATOM   4436  CB  VAL     5      33.505  33.560  21.180  1.00 50.09      H31H C
ATOM   4437  CG1 VAL     5      33.387  35.028  21.543  1.00 49.53      H31H C
ATOM   4438  CG2 VAL     5      34.141  32.794  22.322  1.00 49.00      H31H C
ATOM   4439  C   VAL     5      31.563  33.882  19.713  1.00 53.87      H31H C
ATOM   4440  O   VAL     5      32.284  34.195  18.766  1.00 55.08      H31H O
ATOM   4441  N   GLU     6      30.311  34.306  19.819  1.00 54.66      H31H N
ATOM   4442  CA  GLU     6      29.732  35.132  18.770  1.00 56.37      H31H C
ATOM   4443  CB  GLU     6      28.226  34.881  18.665  1.00 54.99      H31H C
ATOM   4444  CG  GLU     6      27.495  34.976  19.987  1.00 54.97      H31H C
ATOM   4445  CD  GLU     6      27.518  33.665  20.770  1.00 55.38      H31H C
ATOM   4446  OE1 GLU     6      28.549  33.383  21.431  1.00 53.88      H31H O
ATOM   4447  OE2 GLU     6      26.500  32.925  20.723  1.00 53.41      H31H O
ATOM   4448  C   GLU     6      29.998  36.613  19.010  1.00 57.61      H31H C
ATOM   4449  O   GLU     6      30.002  37.071  20.156  1.00 59.58      H31H O
ATOM   4450  N   SER     7      30.225  37.355  17.926  1.00 57.59      H31H N
ATOM   4451  CA  SER     7      30.437  38.799  18.006  1.00 56.93      H31H C
ATOM   4452  CB  SER     7      31.906  39.125  17.764  1.00 55.14      H31H C
ATOM   4453  OG  SER     7      32.269  38.770  16.441  1.00 54.54      H31H O
ATOM   4454  C   SER     7      29.589  39.533  16.969  1.00 57.75      H31H C
ATOM   4455  O   SER     7      28.690  38.947  16.347  1.00 58.75      H31H O
ATOM   4456  N   GLY     8      29.876  40.820  16.792  1.00 57.04      H31H N
ATOM   4457  CA  GLY     8      29.352  41.535  15.643  1.00 57.05      H31H C
ATOM   4458  C   GLY     8      28.048  42.264  15.900  1.00 57.40      H31H C
ATOM   4459  O   GLY     8      27.650  43.126  15.110  1.00 57.11      H31H O
ATOM   4460  N   GLY     9      27.384  41.923  17.002  1.00 57.05      H31H N
ATOM   4461  CA  GLY     9      26.120  42.558  17.332  1.00 57.07      H31H C
ATOM   4462  C   GLY     9      26.265  44.057  17.523  1.00 57.42      H31H C
ATOM   4463  O   GLY     9      27.372  44.593  17.456  1.00 57.35      H31H O
ATOM   4464  N   GLY    10      25.145  44.737  17.757  1.00 57.20      H31H N
ATOM   4465  CA  GLY    10      25.175  46.177  17.926  1.00 57.17      H31H C
ATOM   4466  C   GLY    10      23.875  46.876  17.566  1.00 58.48      H31H C
ATOM   4467  O   GLY    10      22.830  46.243  17.363  1.00 57.46      H31H O
ATOM   4468  N   LEU    11      23.948  48.204  17.494  1.00 59.75      H31H N
ATOM   4469  CA  LEU    11      22.814  49.044  17.119  1.00 59.30      H31H C
ATOM   4470  CB  LEU    11      22.908  50.378  17.860  1.00 61.04      H31H C
ATOM   4471  CG  LEU    11      21.886  51.491  17.591  1.00 63.37      H31H C
ATOM   4472  CD1 LEU    11      20.477  51.090  18.065  1.00 60.92      H31H C
ATOM   4473  CD2 LEU    11      22.372  52.757  18.308  1.00 63.68      H31H C
ATOM   4474  C   LEU    11      22.830  49.274  15.606  1.00 59.16      H31H C
ATOM   4475  O   LEU    11      23.904  49.326  14.995  1.00 58.55      H31H O
ATOM   4476  N   VAL    12      21.640  49.399  15.013  1.00 57.89      H31H N
ATOM   4477  CA  VAL    12      21.489  49.473  13.560  1.00 57.68      H31H C
ATOM   4478  CB  VAL    12      21.935  48.151  12.891  1.00 58.73      H31H C
ATOM   4479  CG1 VAL    12      21.083  47.863  11.667  1.00 57.50      H31H C
ATOM   4480  CG2 VAL    12      23.416  48.249  12.488  1.00 58.66      H31H C
ATOM   4481  C   VAL    12      20.056  49.780  13.134  1.00 57.57      H31H C
ATOM   4482  O   VAL    12      19.106  49.170  13.624  1.00 57.48      H31H O
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ATOM   4483  N   LYS   13      19.905  50.725  12.212  1.00 56.16      H31B N
ATOM   4484  CA  LYS   13      18.583  51.217  11.835  1.00 54.93      H31H C
ATOM   4485  CB  LYS   13      18.701  52.571  11.111  1.00 57.48      H31H C
ATOM   4486  CG  LYS   13      19.294  53.705  11.951  1.00 58.72      H31H C
ATOM   4487  CD  LYS   13      19.230  55.040  11.220  0.50 60.60      H31H C
ATOM   4488  CE  LYS   13      17.828  55.638  11.249  1.00 61.98      H31H C
ATOM   4489  NZ  LYS   13      16.837  54.802  10.504  1.00 65.29      H31H N
ATOM   4490  C   LYS   13      17.846  50.237  10.938  1.00 52.27      H31H C
ATOM   4491  O   LYS   13      18.456  49.409  10.276  1.00 51.72      H31H O
ATOM   4492  N   PRO   14      16.513  50.335  10.888  1.00 51.07      H31H N
ATOM   4493  CD  PRO   14      15.629  51.184  11.701  1.00 50.36      H31H C
ATOM   4494  CA  PRO   14      15.756  49.463   9.981  1.00 51.56      H31H C
ATOM   4495  CB  PRO   14      14.315  49.950  10.136  1.00 49.93      H31H C
ATOM   4496  CG  PRO   14      14.283  50.547  11.491  1.00 50.40      H31H C
ATOM   4497  C   PRO   14      16.245  49.535   8.526  1.00 52.20      H31H C
ATOM   4498  O   PRO   14      16.571  50.616   8.016  1.00 51.66      H31H O
ATOM   4499  N   GLY   15      16.293  48.373   7.871  1.00 53.10      H31H N
ATOM   4500  CA  GLY   15      16.747  48.295   6.493  1.00 53.00      H31H C
ATOM   4501  C   GLY   15      18.237  48.032   6.398  1.00 54.02      H31H C
ATOM   4502  O   GLY   15      18.736  47.583   5.368  1.00 55.51      H31H O
ATOM   4503  N   GLY   16      18.959  48.310   7.477  1.00 53.44      H31H N
ATOM   4504  CA  GLY   16      20.401  48.146   7.455  1.00 52.71      H31H C
ATOM   4505  C   GLY   16      20.827  46.695   7.437  1.00 53.74      H31H C
ATOM   4506  O   GLY   16      20.030  45.795   7.131  1.00 53.76      H31H O
ATOM   4507  N   SER   17      22.088  46.456   7.781  1.00 53.93      H31H N
ATOM   4508  CA  SER   17      22.700  45.171   7.494  1.00 52.93      H31H C
ATOM   4509  CB  SER   17      23.081  45.114   6.008  1.00 51.53      H31H C
ATOM   4510  OG  SER   17      21.949  45.342   5.180  1.00 49.99      H31H O
ATOM   4511  C   SER   17      23.919  44.885   8.363  1.00 53.75      H31H C
ATOM   4512  O   SER   17      25.036  45.301   8.051  1.00 54.05      H31H O
ATOM   4513  N   LEU   18      23.694  44.154   9.451  1.00 54.07      H31H N
ATOM   4514  CA  LEU   18      24.773  43.741  10.346  1.00 52.61      H31H C
ATOM   4515  CB  LEU   18      24.274  43.776  11.799  1.00 53.51      H31H C
ATOM   4516  CG  LEU   18      25.328  43.451  12.864  1.00 55.27      H31H C
ATOM   4517  CD1 LEU   18      26.373  44.567  12.903  1.00 54.03      H31H C
ATOM   4518  CD2 LEU   18      24.658  43.258  14.229  1.00 53.34      H31H C
ATOM   4519  C   LEU   18      25.255  42.329   9.977  1.00 50.58      H31H C
ATOM   4520  O   LEU   18      24.562  41.596   9.269  1.00 50.59      H31H O
ATOM   4521  N   ARG   19      26.440  41.948  10.447  1.00 47.55      H31H N
ATOM   4522  CA  ARG   19      26.975  40.621  10.144  1.00 45.24      H31H C
ATOM   4523  CB  ARG   19      27.973  40.697   8.998  1.00 44.78      H31H C
ATOM   4524  CG  ARG   19      28.777  39.417   8.827  1.00 46.47      H31H C
ATOM   4525  CD  ARG   19      29.711  39.545   7.655  1.00 46.31      H31H C
ATOM   4526  NE  ARG   19      30.114  38.261   7.109  1.00 47.93      H31H N
ATOM   4527  CZ  ARG   19      31.321  37.743   7.278  1.00 50.43      H31H C
ATOM   4528  NH1 ARG   19      31.630  36.560   6.743  1.00 48.32      H31H N
ATOM   4529  NH2 ARG   19      32.217  38.420   7.992  1.00 50.58      H31H N
ATOM   4530  C   ARG   19      27.646  39.908  11.312  1.00 43.40      H31H C
ATOM   4531  O   ARG   19      28.622  40.399  11.881  1.00 43.30      H31H O
ATOM   4532  N   LEU   20      27.141  38.722  11.630  1.00 41.57      H31H N
ATOM   4533  CA  LEU   20      27.496  38.040  12.866  1.00 40.63      H31H C
ATOM   4534  CB  LEU   20      26.259  37.394  13.482  1.00 38.45      H31H C
ATOM   4535  CG  LEU   20      25.242  38.429  13.925  1.00 37.41      H31H C
ATOM   4536  CD1 LEU   20      24.000  37.758  14.531  1.00 35.80      H31H C
ATOM   4537  CD2 LEU   20      25.945  39.365  14.911  1.00 34.57      H31H C
ATOM   4538  C   LEU   20      28.553  36.988  12.664  1.00 40.56      H31H C
ATOM   4539  O   LEU   20      28.522  36.244  11.696  1.00 41.52      H31H O
ATOM   4540  N   SER   21      29.489  36.932  13.597  1.00 42.24      H31H N
ATOM   4541  CA  SER   21      30.476  35.874  13.614  1.00 44.84      H31H C
ATOM   4542  CB  SER   21      31.877  36.454  13.484  1.00 45.20      H31H C
ATOM   4543  OG  SER   21      32.052  37.046  12.210  1.00 45.61      H31H O
ATOM   4544  C   SER   21      30.361  35.102  14.913  1.00 46.37      H31H C
ATOM   4545  O   SER   21      29.816  35.616  15.897  1.00 48.51      H31H O
ATOM   4546  N   CYS   22      30.872  33.874  14.902  1.00 45.31      H31H N
ATOM   4547  CA  CYS   22      30.833  32.997  16.053  1.00 45.48      H31H C
ATOM   4548  C   CYS   22      32.020  32.044  15.972  1.00 45.47      H31H C
ATOM   4549  O   CYS   22      32.082  31.197  15.081  1.00 45.24      H31H O
ATOM   4550  CB  CYS   22      29.529  32.211  16.053  1.00 46.72      H31H C
ATOM   4551  SG  CYS   22      29.593  30.638  16.967  1.00 52.03      H31H S
ATOM   4552  N   ALA   23      32.962  32.195  16.903  1.00 45.04      H31H N
ATOM   4553  CA  ALA   23      34.255  31.520  16.821  1.00 45.19      H31H C
ATOM   4554  CB  ALA   23      35.374  32.514  17.065  1.00 43.20      H31H C
ATOM   4555  C   ALA   23      34.367  30.360  17.806  1.00 45.82      H31H C
ATOM   4556  O   ALA   23      34.373  30.561  19.023  1.00 47.17      H31H O
ATOM   4557  N   ALA   24      34.473  29.147  17.272  1.00 44.91      H31H N
ATOM   4558  CA  ALA   24      34.486  27.953  18.098  1.00 44.44      H31H C
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ATOM   4559  CB  ALA   24      33.590  26.910  17.502  1.00 45.51      H31H C
ATOM   4560  C   ALA   24      35.881  27.398  18.230  1.00 44.69      H31H C
ATOM   4561  O   ALA   24      36.701  27.533  17.323  1.00 44.96      H31H O
ATOM   4562  N   SER   25      36.137  26.754  19.362  1.00 44.80      H31H N
ATOM   4563  CA  SER   25      37.454  26.204  19.654  1.00 44.29      H31H C
ATOM   4564  CB  SER   25      38.327  27.287  20.285  1.00 44.22      H31H C
ATOM   4565  OG  SER   25      37.541  28.248  20.983  1.00 44.39      H31H O
ATOM   4566  C   SER   25      37.366  24.994  20.580  1.00 44.72      H31H C
ATOM   4567  O   SER   25      36.397  24.823  21.322  1.00 44.63      H31H O
ATOM   4568  N   GLY   26      38.384  24.145  20.526  1.00 44.89      H31H N
ATOM   4569  CA  GLY   26      38.428  23.012  21.430  1.00 44.97      H31H C
ATOM   4570  C   GLY   26      37.508  21.877  21.024  1.00 44.73      H31H C
ATOM   4571  O   GLY   26      37.004  21.145  21.872  1.00 45.24      H31H O
ATOM   4572  N   PHE   27      37.280  21.732  19.725  1.00 44.04      H31H N
ATOM   4573  CA  PHE   27      36.499  20.615  19.220  1.00 42.33      H31H C
ATOM   4574  CB  PHE   27      35.074  20.703  19.739  1.00 39.84      H31H C
ATOM   4575  CG  PHE   27      34.256  21.745  19.078  1.00 38.30      H31H C
ATOM   4576  CD1 PHE   27      34.255  23.037  19.549  1.00 38.44      H31H C
ATOM   4577  CD2 PHE   27      33.405  21.411  18.038  1.00 39.02      H31H C
ATOM   4578  CE1 PHE   27      33.402  23.985  18.999  1.00 38.76      H31H C
ATOM   4579  CE2 PHE   27      32.553  22.352  17.484  1.00 39.23      H31H C
ATOM   4580  CZ  PHE   27      32.549  23.638  17.968  1.00 38.31      H31H C
ATOM   4581  C   PHE   27      36.495  20.575  17.697  1.00 42.00      H31H C
ATOM   4582  O   PHE   27      36.708  21.596  17.042  1.00 44.48      H31H O
ATOM   4583  N   THR   28      36.266  19.398  17.131  1.00 39.42      H31H N
ATOM   4584  CA  THR   28      36.311  19.263  15.692  1.00 40.18      H31H C
ATOM   4585  CB  THR   28      36.369  17.775  15.307  1.00 40.22      H31H C
ATOM   4586  OG1 THR   28      35.447  17.047  16.120  1.00 43.67      H31H O
ATOM   4587  CG2 THR   28      37.764  17.211  15.536  1.00 39.15      H31H C
ATOM   4588  C   THR   28      35.125  19.972  15.010  1.00 40.83      H31H C
ATOM   4589  O   THR   28      34.120  19.360  14.642  1.00 41.68      H31H O
ATOM   4590  N   PHE   29      35.271  21.282  14.844  1.00 39.15      H31H N
ATOM   4591  CA  PHE   29      34.258  22.129  14.237  1.00 36.37      H31H C
ATOM   4592  CB  PHE   29      34.820  23.537  14.064  1.00 34.08      H31H C
ATOM   4593  CG  PHE   29      33.869  24.504  13.420  1.00 32.57      H31H C
ATOM   4594  CD1 PHE   29      32.871  25.108  14.161  1.00 32.26      H31H C
ATOM   4595  CD2 PHE   29      34.033  24.877  12.098  1.00 31.00      H31H C
ATOM   4596  CE1 PHE   29      32.055  26.081  13.595  1.00 33.72      H31H C
ATOM   4597  CE2 PHE   29      33.229  25.842  11.523  1.00 30.04      H31H C
ATOM   4598  CZ  PHE   29      32.238  26.451  12.269  1.00 32.68      H31H C
ATOM   4599  C   PHE   29      33.802  21.606  12.891  1.00 35.94      H31H C
ATOM   4600  O   PHE   29      32.664  21.837  12.489  1.00 37.05      H31H O
ATOM   4601  N   SER   30      34.689  20.919  12.185  1.00 34.97      H31H N
ATOM   4602  CA  SER   30      34.351  20.421  10.866  1.00 37.04      H31H C
ATOM   4603  CB  SER   30      35.615  20.115  10.079  1.00 38.55      H31H C
ATOM   4604  OG  SER   30      36.576  21.137  10.284  1.00 45.91      H31H O
ATOM   4605  C   SER   30      33.495  19.163  10.995  1.00 38.11      H31H C
ATOM   4606  O   SER   30      33.013  18.611  10.000  1.00 40.04      H31H O
ATOM   4607  N   SER   31      33.285  18.718  12.226  1.00 37.14      H31H N
ATOM   4608  CA  SER   31      32.404  17.589  12.449  1.00 37.25      H31H C
ATOM   4609  CB  SER   31      33.079  16.573  13.379  1.00 39.60      H31H C
ATOM   4610  OG  SER   31      34.428  16.335  12.995  1.00 41.49      H31H O
ATOM   4611  C   SER   31      31.043  17.997  13.021  1.00 36.03      H31H C
ATOM   4612  O   SER   31      30.155  17.156  13.164  1.00 36.13      H31H O
ATOM   4613  N   TYR   32      30.863  19.271  13.350  1.00 34.15      H31H N
ATOM   4614  CA  TYR   32      29.593  19.683  13.924  1.00 33.10      H31H C
ATOM   4615  CB  TYR   32      29.824  20.398  15.257  1.00 32.49      H31H C
ATOM   4616  CG  TYR   32      30.191  19.432  16.369  1.00 33.31      H31H C
ATOM   4617  CD1 TYR   32      31.384  18.741  16.336  1.00 32.16      H31H C
ATOM   4618  CE1 TYR   32      31.691  17.817  17.293  1.00 34.34      H31H C
ATOM   4619  CD2 TYR   32      29.313  19.168  17.410  1.00 32.93      H31H C
ATOM   4620  CE2 TYR   32      29.614  18.239  18.379  1.00 33.46      H31H C
ATOM   4621  CZ  TYR   32      30.809  17.564  18.315  1.00 33.85      H31H C
ATOM   4622  OH  TYR   32      31.143  16.635  19.278  1.00 35.15      H31H O
ATOM   4623  C   TYR   32      28.758  20.541  12.999  1.00 32.60      H31H C
ATOM   4624  O   TYR   32      29.277  21.281  12.172  1.00 35.97      H31H O
ATOM   4625  N   SER   33      27.449  20.417  13.124  1.00 30.53      H31H N
ATOM   4626  CA  SER   33      26.532  21.322  12.463  1.00 28.81      H31H C
ATOM   4627  CB  SER   33      25.173  20.652  12.302  1.00 25.79      H31H C
ATOM   4628  OG  SER   33      25.209  19.655  11.302  1.00 22.79      H31H O
ATOM   4629  C   SER   33      26.390  22.561  13.335  1.00 29.79      H31H C
ATOM   4630  O   SER   33      26.681  22.519  14.530  1.00 29.74      H31H O
ATOM   4631  N   MET   34      25.939  23.661  12.745  1.00 30.43      H31H N
ATOM   4632  CA  MET   34      25.839  24.916  13.483  1.00 31.05      H31H C
ATOM   4633  CB  MET   34      26.990  25.837  13.106  1.00 31.89      H31H C
ATOM   4634  CG  MET   34      28.361  25.200  13.325  1.00 32.27      H31H C
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ATOM   4635  SD  MET   34      28.799  25.192  15.077  1.00 32.96      H31H S
ATOM   4636  CE  MET   34      28.136  26.768  15.651  1.00 34.63      H31H C
ATOM   4637  C   MET   34      24.519  25.612  13.235  1.00 31.84      H31H C
ATOM   4638  O   MET   34      23.991  25.620  12.120  1.00 30.49      H31H O
ATOM   4639  N   ASN   35      23.981  26.188  14.300  1.00 34.23      H31H N
ATOM   4640  CA  ASN   35      22.678  26.835  14.254  1.00 35.04      H31H C
ATOM   4641  CB  ASN   35      21.686  26.033  15.069  1.00 33.61      H31H C
ATOM   4642  CG  ASN   35      21.946  24.564  14.982  1.00 30.94      H31H C
ATOM   4643  OD1 ASN   35      21.254  23.847  14.272  1.00 30.44      H31H O
ATOM   4644  ND2 ASN   35      22.961  24.098  15.704  1.00 30.36      H31H N
ATOM   4645  C   ASN   35      22.822  28.218  14.851  1.00 36.26      H31H C
ATOM   4646  O   ASN   35      23.693  28.445  15.703  1.00 38.02      H31H O
ATOM   4647  N   TRP   36      21.990  29.143  14.380  1.00 35.97      H31H N
ATOM   4648  CA  TRP   36      21.827  30.419  15.043  1.00 35.60      H31H C
ATOM   4649  CB  TRP   36      21.871  31.563  14.046  1.00 33.77      H31H C
ATOM   4650  CG  TRP   36      23.227  31.892  13.566  1.00 34.67      H31H C
ATOM   4651  CD2 TRP   36      24.252  32.593  14.265  1.00 34.96      H31H C
ATOM   4652  CE2 TRP   36      25.363  32.692  13.398  1.00 35.76      H31H C
ATOM   4653  CE3 TRP   36      24.341  33.149  15.543  1.00 36.02      H31H C
ATOM   4654  CD1 TRP   36      23.743  31.599  12.329  1.00 35.15      H31H C
ATOM   4655  NE1 TRP   36      25.020  32.071  12.225  1.00 34.37      H31H N
ATOM   4656  CZ2 TRP   36      26.556  33.331  13.768  1.00 34.70      H31H C
ATOM   4657  CZ3 TRP   36      25.526  33.783  15.914  1.00 37.06      H31H C
ATOM   4658  CH2 TRP   36      26.618  33.869  15.024  1.00 36.44      H31H C
ATOM   4659  C   TRP   36      20.472  30.381  15.698  1.00 38.32      H31H C
ATOM   4660  O   TRP   36      19.465  30.072  15.043  1.00 38.04      H31H O
ATOM   4661  N   VAL   37      20.455  30.680  16.993  1.00 40.27      H31H N
ATOM   4662  CA  VAL   37      19.212  30.858  17.730  1.00 43.42      H31H C
ATOM   4663  CB  VAL   37      19.034  29.749  18.807  1.00 42.89      H31H C
ATOM   4664  CG1 VAL   37      17.703  29.938  19.523  1.00 39.41      H31H C
ATOM   4665  CG2 VAL   37      19.123  28.358  18.170  1.00 41.48      H31H C
ATOM   4666  C   VAL   37      19.187  32.233  18.416  1.00 46.02      H31H C
ATOM   4667  O   VAL   37      20.165  32.655  19.056  1.00 44.43      H31H O
ATOM   4668  N   ARG   38      18.057  32.921  18.291  1.00 48.68      H31H N
ATOM   4669  CA  ARG   38      17.945  34.276  18.807  1.00 52.25      H31H C
ATOM   4670  CB  ARG   38      17.639  35.253  17.668  1.00 51.00      H31H C
ATOM   4671  CG  ARG   38      16.240  35.109  17.069  1.00 47.89      H31H C
ATOM   4672  CD  ARG   38      16.041  36.123  15.962  1.00 46.37      H31H C
ATOM   4673  NE  ARG   38      14.706  36.073  15.372  1.00 45.46      H31H N
ATOM   4674  CZ  ARG   38      14.329  36.822  14.337  1.00 43.87      H31H C
ATOM   4675  NH1 ARG   38      15.190  37.672  13.791  1.00 43.17      H31H N
ATOM   4676  NH2 ARG   38      13.101  36.720  13.840  1.00 40.65      H31H N
ATOM   4677  C   ARG   38      16.868  34.399  19.872  1.00 56.24      H31H C
ATOM   4678  O   ARG   38      15.848  33.707  19.829  1.00 54.41      H31H O
ATOM   4679  N   GLN   39      17.106  35.311  20.811  1.00 62.26      H31H N
ATOM   4680  CA  GLN   39      16.211  35.540  21.936  1.00 68.98      H31H C
ATOM   4681  CB  GLN   39      16.793  34.868  23.175  1.00 65.66      H31H C
ATOM   4682  CG  GLN   39      15.815  34.613  24.288  1.00 61.47      H31H C
ATOM   4683  CD  GLN   39      16.497  33.981  25.467  1.00 59.61      H31H C
ATOM   4684  OE1 GLN   39      17.640  34.311  25.768  1.00 58.86      H31H O
ATOM   4685  NE2 GLN   39      15.813  33.060  26.139  1.00 59.10      H31H N
ATOM   4686  C   GLN   39      16.018  37.037  22.206  1.00 75.28      H31H C
ATOM   4687  O   GLN   39      16.893  37.691  22.776  1.00 76.14      H31H O
ATOM   4688  N   ALA   40      14.870  37.569  21.798  1.00 82.49      H31H N
ATOM   4689  CA  ALA   40      14.456  38.915  22.195  1.00 88.68      H31H C
ATOM   4690  CB  ALA   40      13.087  39.221  21.614  1.00 86.83      H31H C
ATOM   4691  C   ALA   40      14.403  38.999  23.722  1.00 93.41      H31H C
ATOM   4692  O   ALA   40      14.006  38.041  24.383  1.00 96.08      H31H O
ATOM   4693  N   PRO   41      14.790  40.151  24.304  1.00 96.76      H31H N
ATOM   4694  CD  PRO   41      15.211  41.406  23.659  1.00 99.21      H31H C
ATOM   4695  CA  PRO   41      14.848  40.250  25.769  1.00 98.49      H31H C
ATOM   4696  CB  PRO   41      15.227  41.709  26.015  1.00 99.06      H31H C
ATOM   4697  CG  PRO   41      15.917  42.125  24.767  1.00 99.43      H31H C
ATOM   4698  C   PRO   41      13.519  39.878  26.427  1.00 99.33      H31H C
ATOM   4699  O   PRO   41      12.459  40.368  26.032  1.00 98.92      H31H O
ATOM   4700  N   GLY   42      13.580  39.001  27.423  1.00100.58      H31H N
ATOM   4701  CA  GLY   42      12.365  38.532  28.062  1.00101.63      H31H C
ATOM   4702  C   GLY   42      11.418  37.838  27.102  1.00101.74      H31H C
ATOM   4703  O   GLY   42      10.220  37.766  27.354  1.00103.13      H31H O
ATOM   4704  N   LYS   43      11.950  37.333  25.994  1.00100.03      H31H N
ATOM   4705  CA  LYS   43      11.163  36.535  25.059  1.00 97.53      H31H C
ATOM   4706  CB  LYS   43      11.258  37.115  23.644  1.00100.83      H31H C
ATOM   4707  CG  LYS   43      10.223  38.182  23.322  1.00105.01      H31H C
ATOM   4708  CD  LYS   43       8.805  37.627  23.437  1.00110.42      H31H C
ATOM   4709  CE  LYS   43       7.809  38.418  22.588  1.00115.16      H31H C
ATOM   4710  NZ  LYS   43       7.763  39.871  22.929  1.00120.90      H31H N
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ATOM	4711	C	LYS	43	11.645	35.088	25.049	1.00	93.32	H31H C
ATOM	4712	O	LYS	43	12.586	34.734	25.759	1.00	92.70	H31H O
ATOM	4713	N	GLY	44	10.994	34.257	24.240	1.00	88.87	H31H N
ATOM	4714	CA	GLY	44	11.399	32.867	24.125	1.00	82.75	H31H C
ATOM	4715	C	GLY	44	12.655	32.674	23.294	1.00	77.82	H31H C
ATOM	4716	O	GLY	44	13.386	33.626	23.025	1.00	77.56	H31H O
ATOM	4717	N	LEU	45	12.909	31.428	22.900	1.00	73.49	H31H N
ATOM	4718	CA	LEU	45	13.967	31.093	21.945	1.00	67.33	H31H C
ATOM	4719	CB	LEU	45	14.739	29.858	22.413	1.00	65.36	H31H C
ATOM	4720	CG	LEU	45	15.730	30.019	23.567	1.00	62.88	H31H C
ATOM	4721	CD1	LEU	45	16.271	28.667	23.952	1.00	63.05	H31H C
ATOM	4722	CD2	LEU	45	16.872	30.921	23.158	1.00	62.71	H31H C
ATOM	4723	C	LEU	45	13.368	30.810	20.566	1.00	65.07	H31H C
ATOM	4724	O	LEU	45	12.401	30.055	20.440	1.00	63.83	H31H O
ATOM	4725	N	GLU	46	13.936	31.430	19.536	1.00	62.28	H31H N
ATOM	4726	CA	GLU	46	13.544	31.126	18.170	1.00	59.26	H31H C
ATOM	4727	CB	GLU	46	12.874	32.332	17.522	1.00	59.59	H31H C
ATOM	4728	CG	GLU	46	12.279	31.999	16.166	1.00	64.07	H31H C
ATOM	4729	CD	GLU	46	11.623	33.185	15.462	1.00	66.06	H31H C
ATOM	4730	OE1	GLU	46	10.599	32.954	14.777	1.00	67.56	H31H O
ATOM	4731	OE2	GLU	46	12.126	34.331	15.581	1.00	65.32	H31H O
ATOM	4732	C	GLU	46	14.739	30.681	17.325	1.00	56.58	H31H C
ATOM	4733	O	GLU	46	15.717	31.417	17.160	1.00	56.38	H31H O
ATOM	4734	N	TRP	47	14.653	29.459	16.801	1.00	52.74	H31H N
ATOM	4735	CA	TRP	47	15.666	28.928	15.893	1.00	47.43	H31H C
ATOM	4736	CB	TRP	47	15.390	27.453	15.559	1.00	45.19	H31H C
ATOM	4737	CG	TRP	47	16.115	27.022	14.328	1.00	40.33	H31H C
ATOM	4738	CD2	TRP	47	15.560	26.829	13.019	1.00	38.45	H31H C
ATOM	4739	CE2	TRP	47	16.639	26.587	12.141	1.00	37.27	H31H C
ATOM	4740	CE3	TRP	47	14.260	26.841	12.503	1.00	36.40	H31H C
ATOM	4741	CD1	TRP	47	17.458	26.875	14.197	1.00	39.72	H31H C
ATOM	4742	NE1	TRP	47	17.785	26.620	12.888	1.00	39.52	H31H N
ATOM	4743	CZ2	TRP	47	16.462	26.363	10.775	1.00	34.81	H31H C
ATOM	4744	CZ3	TRP	47	14.084	26.618	11.148	1.00	35.90	H31H C
ATOM	4745	CH2	TRP	47	15.183	26.383	10.296	1.00	34.98	H31H C
ATOM	4746	C	TRP	47	15.659	29.726	14.601	1.00	45.07	H31H C
ATOM	4747	O	TRP	47	14.613	29.880	13.970	1.00	43.66	H31H O
ATOM	4748	N	VAL	48	16.830	30.212	14.200	1.00	43.15	H31H N
ATOM	4749	CA	VAL	48	16.926	31.072	13.025	1.00	41.49	H31H C
ATOM	4750	CB	VAL	48	17.852	32.261	13.274	1.00	42.05	H31H C
ATOM	4751	CG1	VAL	48	18.160	32.943	11.963	1.00	40.58	H31H C
ATOM	4752	CG2	VAL	48	17.191	33.232	14.232	1.00	42.96	H31H C
ATOM	4753	C	VAL	48	17.407	30.374	11.770	1.00	39.83	H31H C
ATOM	4754	O	VAL	48	16.730	30.404	10.736	1.00	37.72	H31H O
ATOM	4755	N	SER	49	18.580	29.752	11.864	1.00	37.12	H31H N
ATOM	4756	CA	SER	49	19.169	29.059	10.722	1.00	35.46	H31H C
ATOM	4757	CB	SER	49	19.985	30.042	9.893	1.00	33.71	H31H C
ATOM	4758	OG	SER	49	20.437	29.445	8.701	1.00	37.92	H31H O
ATOM	4759	C	SER	49	20.060	27.898	11.173	1.00	34.93	H31H C
ATOM	4760	O	SER	49	20.574	27.900	12.291	1.00	34.06	H31H O
ATOM	4761	N	SER	50	20.229	26.897	10.313	1.00	33.84	H31H N
ATOM	4762	CA	SER	50	21.244	25.865	10.552	1.00	31.90	H31H C
ATOM	4763	CB	SER	50	20.598	24.527	10.919	1.00	32.87	H31H C
ATOM	4764	OG	SER	50	20.026	24.569	12.211	1.00	36.99	H31H O
ATOM	4765	C	SER	50	22.101	25.676	9.318	1.00	29.80	H31H C
ATOM	4766	O	SER	50	21.675	25.992	8.208	1.00	30.95	H31H O
ATOM	4767	N	ILE	51	23.306	25.156	9.516	1.00	27.14	H31H N
ATOM	4768	CA	ILE	51	24.194	24.797	8.410	1.00	24.85	H31H C
ATOM	4769	CB	ILE	51	25.101	25.998	8.040	1.00	24.72	H31H C
ATOM	4770	CG2	ILE	51	25.868	26.476	9.254	1.00	24.95	H31H C
ATOM	4771	CG1	ILE	51	26.066	25.604	6.929	1.00	27.20	H31H C
ATOM	4772	CD1	ILE	51	26.529	26.779	6.101	1.00	28.08	H31H C
ATOM	4773	C	ILE	51	25.045	23.587	8.817	1.00	25.34	H31H C
ATOM	4774	O	ILE	51	25.720	23.620	9.851	1.00	24.66	H31H O
ATOM	4775	N	SER	52	25.010	22.509	8.033	1.00	25.68	H31H N
ATOM	4776	CA	SER	52	25.716	21.275	8.435	1.00	24.38	H31H C
ATOM	4777	CB	SER	52	25.185	20.051	7.683	1.00	22.57	H31H C
ATOM	4778	OG	SER	52	25.317	20.181	6.272	1.00	22.51	H31H O
ATOM	4779	C	SER	52	27.203	21.399	8.193	1.00	25.36	H31H C
ATOM	4780	O	SER	52	27.659	22.349	7.547	1.00	27.15	H31H O
ATOM	4781	N	SER	53	27.962	20.429	8.691	1.00	26.78	H31H N
ATOM	4782	CA	SER	53	29.414	20.558	8.715	1.00	27.05	H31H C
ATOM	4783	CB	SER	53	30.061	19.303	9.317	1.00	30.95	H31H C
ATOM	4784	OG	SER	53	29.791	18.147	8.553	1.00	33.86	H31H O
ATOM	4785	C	SER	53	29.998	20.840	7.339	1.00	24.84	H31H C
ATOM	4786	O	SER	53	30.897	21.674	7.200	1.00	23.50	H31H O

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ATOM   4787  N   SER   54      29.475  20.165   6.323  1.00 24.23      H31H N
ATOM   4788  CA  SER   54      29.955  20.384   4.967  1.00 24.68      H31H C
ATOM   4789  CB  SER   54      30.116  19.060   4.223  1.00 23.52      H31H C
ATOM   4790  OG  SER   54      28.858  18.490   3.938  1.00 27.63      H31H O
ATOM   4791  C   SER   54      29.070  21.307   4.146  1.00 26.14      H31H C
ATOM   4792  O   SER   54      29.316  21.469   2.945  1.00 25.03      H31H O
ATOM   4793  N   SER   55      28.054  21.911   4.780  1.00 26.34      H31H N
ATOM   4794  CA  SER   55      27.186  22.904   4.126  1.00 27.36      H31H C
ATOM   4795  CB  SER   55      28.004  24.042   3.500  1.00 24.26      H31H C
ATOM   4796  OG  SER   55      28.836  24.687   4.449  1.00 27.22      H31H O
ATOM   4797  C   SER   55      26.310  23.319   3.025  1.00 30.48      H31H C
ATOM   4798  O   SER   55      26.060  22.985   2.026  1.00 31.05      H31H O
ATOM   4799  N   SER   56      25.843  21.085   3.196  1.00 33.26      H31H N
ATOM   4800  CA  SER   56      25.012  20.433   2.180  1.00 34.73      H31H C
ATOM   4801  CB  SER   56      25.446  18.986   2.021  1.00 35.72      H31H C
ATOM   4802  OG  SER   56      26.824  18.900   2.313  1.00 39.26      H31H O
ATOM   4803  C   SER   56      23.594  20.481   2.688  1.00 34.95      H31H C
ATOM   4804  O   SER   56      22.655  20.002   2.043  1.00 35.68      H31H O
ATOM   4805  N   TYR   57      23.460  21.039   3.882  1.00 34.46      H31H N
ATOM   4806  CA  TYR   57      22.165  21.241   4.499  1.00 34.44      H31H C
ATOM   4807  CB  TYR   57      22.027  20.369   5.740  1.00 33.79      H31H C
ATOM   4808  CG  TYR   57      21.899  18.888   5.474  1.00 35.21      H31H C
ATOM   4809  CD1 TYR   57      22.920  18.020   5.831  1.00 35.98      H31H C
ATOM   4810  CE1 TYR   57      22.790  16.663   5.671  1.00 35.95      H31H C
ATOM   4811  CD2 TYR   57      20.733  18.347   4.931  1.00 33.79      H31H C
ATOM   4812  CE2 TYR   57      20.596  16.980   4.765  1.00 33.48      H31H C
ATOM   4813  CZ  TYR   57      21.632  16.148   5.146  1.00 34.19      H31H C
ATOM   4814  OH  TYR   57      21.526  14.787   5.059  1.00 34.50      H31H O
ATOM   4815  C   TYR   57      22.116  22.683   4.927  1.00 34.00      H31H C
ATOM   4816  O   TYR   57      22.869  23.078   5.808  1.00 37.57      H31H O
ATOM   4817  N   ILE   58      21.247  23.477   4.319  1.00 32.02      H31H N
ATOM   4818  CA  ILE   58      21.036  24.828   4.811  1.00 30.95      H31H C
ATOM   4819  CB  ILE   58      21.641  25.871   3.851  1.00 28.79      H31H C
ATOM   4820  CG2 ILE   58      21.097  27.230   4.152  1.00 29.09      H31H C
ATOM   4821  CG1 ILE   58      23.157  25.900   3.993  1.00 28.87      H31H C
ATOM   4822  CD1 ILE   58      23.842  26.837   3.009  1.00 26.40      H31H C
ATOM   4823  C   ILE   58      19.554  25.102   5.011  1.00 31.64      H31H C
ATOM   4824  O   ILE   58      18.759  24.967   4.087  1.00 32.16      H31H O
ATOM   4825  N   SER   59      19.186  25.469   6.233  1.00 31.86      H31H N
ATOM   4826  CA  SER   59      17.811  25.837   6.528  1.00 32.29      H31H C
ATOM   4827  CB  SER   59      17.191  24.920   7.576  1.00 32.65      H31H C
ATOM   4828  OG  SER   59      17.150  23.589   7.139  1.00 36.33      H31H O
ATOM   4829  C   SER   59      17.739  27.240   7.063  1.00 31.69      H31H C
ATOM   4830  O   SER   59      18.696  27.747   7.661  1.00 30.20      H31H O
ATOM   4831  N   TYR   60      16.568  27.839   6.874  1.00 30.90      H31H N
ATOM   4832  CA  TYR   60      16.235  29.110   7.495  1.00 32.01      H31H C
ATOM   4833  CB  TYR   60      16.383  30.252   6.473  1.00 28.90      H31H C
ATOM   4834  CG  TYR   60      17.790  30.432   5.922  1.00 25.82      H31H C
ATOM   4835  CD1 TYR   60      18.719  31.233   6.586  1.00 24.60      H31H C
ATOM   4836  CE1 TYR   60      19.980  31.462   6.059  1.00 25.44      H31H C
ATOM   4837  CD2 TYR   60      18.165  29.850   4.713  1.00 21.26      H31H C
ATOM   4838  CE2 TYR   60      19.415  30.063   4.176  1.00 23.84      H31H C
ATOM   4839  CZ  TYR   60      20.331  30.878   4.846  1.00 27.34      H31H C
ATOM   4840  OH  TYR   60      21.583  31.133   4.295  1.00 24.44      H31H O
ATOM   4841  C   TYR   60      14.798  29.065   8.034  1.00 34.54      H31H C
ATOM   4842  O   TYR   60      13.929  28.376   7.482  1.00 34.40      H31H O
ATOM   4843  N   ALA   61      14.551  29.796   9.118  1.00 37.78      H31H N
ATOM   4844  CA  ALA   61      13.187  30.030   9.576  1.00 40.86      H31H C
ATOM   4845  CB  ALA   61      13.214  30.679  10.946  1.00 42.20      H31H C
ATOM   4846  C   ALA   61      12.468  30.937   8.573  1.00 42.66      H31H C
ATOM   4847  O   ALA   61      13.079  31.851   8.009  1.00 40.39      H31H O
ATOM   4848  N   ASP   62      11.175  30.681   8.357  1.00 46.05      H31H N
ATOM   4849  CA  ASP   62      10.362  31.465   7.409  1.00 49.44      H31H C
ATOM   4850  CB  ASP   62       8.874  31.073   7.494  1.00 52.61      H31H C
ATOM   4851  CG  ASP   62       8.480  29.984   6.482  1.00 56.56      H31H C
ATOM   4852  OD1 ASP   62       7.576  29.165   6.791  1.00 58.22      H31H O
ATOM   4853  OD2 ASP   62       9.062  29.946   5.376  1.00 58.42      H31H O
ATOM   4854  C   ASP   62      10.488  32.976   7.609  1.00 49.31      H31H C
ATOM   4855  O   ASP   62      10.516  33.727   6.641  1.00 48.62      H31H O
ATOM   4856  N   SER   63      10.573  33.418   8.859  1.00 49.39      H31H N
ATOM   4857  CA  SER   63      10.617  34.847   9.152  1.00 49.60      H31H C
ATOM   4858  CB  SER   63      10.284  35.089  10.625  1.00 51.26      H31H C
ATOM   4859  OG  SER   63      11.163  34.376  11.473  1.00 52.27      H31H O
ATOM   4860  C   SER   63      11.946  35.526   8.810  1.00 49.28      H31H C
ATOM   4861  O   SER   63      12.054  36.754   8.864  1.00 49.16      H31H O
ATOM   4862  N   VAL   64      12.964  34.739   8.477  1.00 49.23      H31H N
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ATOM	4863	CA	VAL	64	14.203	35.316	7.978	1.00	48.95	H31H C
ATOM	4864	CB	VAL	64	15.410	34.952	8.851	1.00	47.82	H31H C
ATOM	4865	CG1	VAL	64	15.263	35.543	10.224	1.00	46.76	H31H C
ATOM	4866	CG2	VAL	64	15.542	33.458	8.924	1.00	49.37	H31H C
ATOM	4867	C	VAL	64	14.482	34.840	6.569	1.00	49.25	H31H C
ATOM	4868	O	VAL	64	15.322	35.401	5.881	1.00	49.52	H31H O
ATOM	4869	N	LYS	65	13.784	33.803	6.136	1.00	49.86	H31H N
ATOM	4870	CA	LYS	65	14.060	33.254	4.826	1.00	52.89	H31H C
ATOM	4871	CB	LYS	65	13.015	32.197	4.467	1.00	55.35	H31H C
ATOM	4872	CG	LYS	65	13.477	31.183	3.436	1.00	60.33	H31H C
ATOM	4873	CD	LYS	65	12.342	30.252	3.010	1.00	66.57	H31H C
ATOM	4874	CE	LYS	65	11.667	29.602	4.212	1.00	71.34	H31H C
ATOM	4875	NZ	LYS	65	12.526	28.612	4.925	1.00	79.97	H31H N
ATOM	4876	C	LYS	65	14.034	34.402	3.822	1.00	52.76	H31H C
ATOM	4877	O	LYS	65	13.028	35.079	3.675	1.00	54.46	H31H O
ATOM	4878	N	GLY	66	15.156	34.637	3.153	1.00	52.42	H31H N
ATOM	4879	CA	GLY	66	15.207	35.685	2.158	1.00	49.76	H31H C
ATOM	4880	C	GLY	66	16.102	36.835	2.565	1.00	50.14	H31H C
ATOM	4881	O	GLY	66	16.658	37.526	1.717	1.00	50.75	H31H O
ATOM	4882	N	ARG	67	16.255	37.045	3.866	1.00	49.59	H31H N
ATOM	4883	CA	ARG	67	16.979	38.208	4.368	1.00	47.70	H31H C
ATOM	4884	CB	ARG	67	16.146	38.939	5.429	1.00	48.66	H31H C
ATOM	4885	CG	ARG	67	15.156	39.953	4.890	1.00	47.79	H31H C
ATOM	4886	CD	ARG	67	13.700	39.689	5.337	1.00	49.90	H31H C
ATOM	4887	NE	ARG	67	13.499	39.282	6.738	1.00	49.54	H31H N
ATOM	4888	CZ	ARG	67	13.916	39.954	7.808	1.00	48.90	H31H C
ATOM	4889	NH1	ARG	67	14.590	41.090	7.674	1.00	46.55	H31H N
ATOM	4890	NH2	ARG	67	13.622	39.500	9.019	1.00	47.78	H31H N
ATOM	4891	C	ARG	67	18.326	37.841	4.969	1.00	46.20	H31H C
ATOM	4892	O	ARG	67	19.294	38.596	4.863	1.00	46.99	H31H O
ATOM	4893	N	PHE	68	18.391	36.697	5.630	1.00	44.42	H31H N
ATOM	4894	CA	PHE	68	19.646	36.302	6.262	1.00	43.90	H31H C
ATOM	4895	CB	PHE	68	19.390	35.737	7.658	1.00	46.31	H31H C
ATOM	4896	CG	PHE	68	18.923	36.755	8.655	1.00	48.27	H31H C
ATOM	4897	CD1	PHE	68	18.783	36.410	9.991	1.00	48.40	H31H C
ATOM	4898	CD2	PHE	68	18.633	38.058	8.265	1.00	49.75	H31H C
ATOM	4899	CE1	PHE	68	18.361	37.344	10.925	1.00	50.45	H31H C
ATOM	4900	CE2	PHE	68	18.207	38.998	9.193	1.00	49.99	H31H C
ATOM	4901	CZ	PHE	68	18.073	38.642	10.525	1.00	50.51	H31H C
ATOM	4902	C	PHE	68	20.393	35.265	5.434	1.00	41.81	H31H C
ATOM	4903	O	PHE	68	19.790	34.417	4.782	1.00	42.54	H31H O
ATOM	4904	N	THR	69	21.712	35.318	5.453	1.00	39.78	H31H N
ATOM	4905	CA	THR	69	22.451	34.211	4.884	1.00	37.73	H31H C
ATOM	4906	CB	THR	69	23.226	34.631	3.637	1.00	37.83	H31H C
ATOM	4907	OG1	THR	69	22.324	35.241	2.707	1.00	40.88	H31H O
ATOM	4908	CG2	THR	69	23.843	33.418	2.980	1.00	36.18	H31H C
ATOM	4909	C	THR	69	23.404	33.638	5.902	1.00	36.00	H31H C
ATOM	4910	O	THR	69	24.206	34.365	6.497	1.00	34.40	H31H O
ATOM	4911	N	ILE	70	23.287	32.331	6.107	1.00	33.23	H31H N
ATOM	4912	CA	ILE	70	24.162	31.610	7.015	1.00	32.07	H31H C
ATOM	4913	CB	ILE	70	23.375	30.489	7.735	1.00	31.36	H31H C
ATOM	4914	CG2	ILE	70	22.789	29.499	6.722	1.00	28.51	H31H C
ATOM	4915	CG1	ILE	70	24.263	29.778	8.739	1.00	28.22	H31H C
ATOM	4916	CD1	ILE	70	23.492	28.722	9.500	1.00	26.73	H31H C
ATOM	4917	C	ILE	70	25.327	31.020	6.221	1.00	33.56	H31H C
ATOM	4918	O	ILE	70	25.172	30.647	5.052	1.00	32.66	H31H O
ATOM	4919	N	SER	71	26.499	30.962	6.845	1.00	34.11	H31H N
ATOM	4920	CA	SER	71	27.667	30.376	6.196	1.00	35.09	H31H C
ATOM	4921	CB	SER	71	28.220	31.321	5.130	1.00	34.90	H31H C
ATOM	4922	OG	SER	71	28.839	32.451	5.718	1.00	38.50	H31H O
ATOM	4923	C	SER	71	28.734	30.114	7.225	1.00	35.70	H31H C
ATOM	4924	O	SER	71	28.878	30.872	8.167	1.00	37.92	H31H O
ATOM	4925	N	ARG	72	29.490	29.043	7.053	1.00	36.94	H31H N
ATOM	4926	CA	ARG	72	30.624	28.806	7.931	1.00	37.72	H31H C
ATOM	4927	CB	ARG	72	30.459	27.472	8.667	1.00	36.35	H31H C
ATOM	4928	CG	ARG	72	29.515	26.487	7.986	1.00	33.89	H31H C
ATOM	4929	CD	ARG	72	30.152	25.131	7.804	1.00	31.35	H31H C
ATOM	4930	NE	ARG	72	30.836	24.658	9.002	1.00	30.33	H31H N
ATOM	4931	CZ	ARG	72	30.230	24.045	10.015	1.00	31.11	H31H C
ATOM	4932	NH1	ARG	72	28.920	23.839	9.976	1.00	32.29	H31H N
ATOM	4933	NH2	ARG	72	30.934	23.608	11.055	1.00	30.74	H31H N
ATOM	4934	C	ARG	72	31.939	28.813	7.165	1.00	38.69	H31H C
ATOM	4935	O	ARG	72	31.957	28.750	5.944	1.00	35.95	H31H O
ATOM	4936	N	ASP	73	33.037	28.892	7.906	1.00	42.23	H31H N
ATOM	4937	CA	ASP	73	34.380	28.780	7.344	1.00	44.50	H31H C
ATOM	4938	CB	ASP	73	35.056	30.154	7.307	1.00	46.65	H31H C

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ATOM   4939  CG  ASP   73      36.385  30.126   6.577  1.00 50.59      H31H C
ATOM   4940  OD1 ASP   73      36.846  29.015   6.248  1.00 53.52      H31H O
ATOM   4941  OD2 ASP   73      36.969  31.207   6.323  1.00 53.00      H31H O
ATOM   4942  C   ASP   73      35.206  27.839   8.219  1.00 45.16      H31H C
ATOM   4943  O   ASP   73      35.841  28.279   9.180  1.00 44.90      H31H O
ATOM   4944  N   ASN   74      35.206  26.550   7.887  1.00 46.03      H31H N
ATOM   4945  CA  ASN   74      35.841  25.545   8.742  1.00 45.42      H31H C
ATOM   4946  CB  ASN   74      35.761  24.167   8.108  1.00 43.35      H31H C
ATOM   4947  CG  ASN   74      34.398  23.569   8.208  1.00 40.70      H31H C
ATOM   4948  OD1 ASN   74      33.479  24.181   8.738  1.00 38.41      H31H O
ATOM   4949  ND2 ASN   74      34.254  22.353   7.696  1.00 42.55      H31H N
ATOM   4950  C   ASN   74      37.297  25.859   8.995  1.00 45.96      H31H C
ATOM   4951  O   ASN   74      37.777  25.768  10.126  1.00 47.31      H31H O
ATOM   4952  N   ALA   75      37.998  26.209   7.925  1.00 44.61      H31H N
ATOM   4953  CA  ALA   75      39.411  26.523   8.003  1.00 43.81      H31H C
ATOM   4954  CB  ALA   75      39.864  27.138   6.693  1.00 41.11      H31H C
ATOM   4955  C   ALA   75      39.617  27.503   9.138  1.00 45.05      H31H C
ATOM   4956  O   ALA   75      40.545  27.359   9.941  1.00 45.68      H31H O
ATOM   4957  N   LYS   76      38.725  28.490   9.194  1.00 46.12      H31H N
ATOM   4958  CA  LYS   76      38.763  29.549  10.189  1.00 46.69      H31H C
ATOM   4959  CB  LYS   76      38.127  30.815   9.626  1.00 48.88      H31H C
ATOM   4960  CG  LYS   76      38.887  31.472   8.494  1.00 49.92      H31H C
ATOM   4961  CD  LYS   76      38.191  32.768   8.107  1.00 51.68      H31H C
ATOM   4962  CE  LYS   76      39.084  33.660   7.272  1.00 52.61      H31H C
ATOM   4963  NZ  LYS   76      38.491  35.025   7.225  1.00 56.36      H31H N
ATOM   4964  C   LYS   76      38.034  29.169  11.470  1.00 46.37      H31H C
ATOM   4965  O   LYS   76      38.020  29.934  12.432  1.00 48.52      H31H O
ATOM   4966  N   ASN   77      37.410  27.999  11.488  1.00 44.21      H31H N
ATOM   4967  CA  ASN   77      36.691  27.587  12.685  1.00 41.76      H31H C
ATOM   4968  CB  ASN   77      37.656  27.462  13.870  1.00 41.56      H31H C
ATOM   4969  CG  ASN   77      38.109  26.038  14.116  1.00 41.99      H31H C
ATOM   4970  OD1 ASN   77      37.684  25.402  15.078  1.00 45.05      H31H O
ATOM   4971  ND2 ASN   77      38.980  25.532  13.255  1.00 41.31      H31H N
ATOM   4972  C   ASN   77      35.637  28.636  13.034  1.00 40.31      H31H C
ATOM   4973  O   ASN   77      35.521  29.024  14.187  1.00 39.76      H31H O
ATOM   4974  N   SER   78      34.873  29.106  12.056  1.00 38.73      H31H N
ATOM   4975  CA  SER   78      33.932  30.171  12.348  1.00 38.78      H31H C
ATOM   4976  CB  SER   78      34.564  31.524  12.031  1.00 37.52      H31H C
ATOM   4977  OG  SER   78      35.815  31.629  12.700  1.00 40.76      H31H O
ATOM   4978  C   SER   78      32.589  30.047  11.655  1.00 39.34      H31H C
ATOM   4979  O   SER   78      32.450  29.364  10.632  1.00 38.37      H31H O
ATOM   4980  N   LEU   79      31.601  30.709  12.249  1.00 38.57      H31H N
ATOM   4981  CA  LEU   79      30.232  30.704  11.759  1.00 38.65      H31H C
ATOM   4982  CB  LEU   79      29.313  30.097  12.811  1.00 37.25      H31H C
ATOM   4983  CG  LEU   79      27.843  30.034  12.420  1.00 36.78      H31H C
ATOM   4984  CD1 LEU   79      27.708  29.259  11.119  1.00 34.19      H31H C
ATOM   4985  CD2 LEU   79      27.043  29.381  13.547  1.00 37.12      H31H C
ATOM   4986  C   LEU   79      29.840  32.147  11.517  1.00 39.27      H31H C
ATOM   4987  O   LEU   79      30.407  33.045  12.136  1.00 40.45      H31H O
ATOM   4988  N   TYR   80      28.884  32.377  10.620  1.00 38.92      H31H N
ATOM   4989  CA  TYR   80      28.496  33.740  10.241  1.00 37.79      H31H C
ATOM   4990  CB  TYR   80      29.241  34.233   8.993  1.00 36.09      H31H C
ATOM   4991  CG  TYR   80      30.730  34.123   9.073  1.00 36.73      H31H C
ATOM   4992  CD1 TYR   80      31.421  33.202   8.289  1.00 37.10      H31H C
ATOM   4993  CE1 TYR   80      32.800  33.070   8.391  1.00 37.42      H31H C
ATOM   4994  CD2 TYR   80      31.455  34.913   9.957  1.00 36.52      H31H C
ATOM   4995  CE2 TYR   80      32.837  34.791  10.065  1.00 37.01      H31H C
ATOM   4996  CZ  TYR   80      33.501  33.866   9.285  1.00 36.52      H31H C
ATOM   4997  OH  TYR   80      34.860  33.706   9.429  1.00 36.87      H31H O
ATOM   4998  C   TYR   80      27.033  33.766   9.919  1.00 37.36      H31H C
ATOM   4999  O   TYR   80      26.479  32.778   9.415  1.00 38.45      H31H O
ATOM   5000  N   LEU   81      26.424  34.913  10.191  1.00 36.86      H31H N
ATOM   5001  CA  LEU   81      25.080  35.214   9.731  1.00 38.28      H31H C
ATOM   5002  CB  LEU   81      24.131  35.156  10.923  1.00 38.22      H31H C
ATOM   5003  CG  LEU   81      22.641  35.249  10.624  1.00 42.17      H31H C
ATOM   5004  CD1 LEU   81      22.202  34.037   9.801  1.00 42.79      H31H C
ATOM   5005  CD2 LEU   81      21.868  35.317  11.946  1.00 42.84      H31H C
ATOM   5006  C   LEU   81      25.048  36.608   9.066  1.00 39.75      H31H C
ATOM   5007  O   LEU   81      25.212  37.633   9.734  1.00 40.10      H31H O
ATOM   5008  N   GLN   82      24.850  36.651   7.751  1.00 40.37      H31H N
ATOM   5009  CA  GLN   82      24.705  37.930   7.061  1.00 41.87      H31H C
ATOM   5010  CB  GLN   82      24.978  37.783   5.569  1.00 42.62      H31H C
ATOM   5011  CG  GLN   82      26.194  38.551   5.084  1.00 42.97      H31H C
ATOM   5012  CD  GLN   82      26.244  39.958   5.634  1.00 43.64      H31H C
ATOM   5013  OE1 GLN   82      27.320  40.479   5.913  1.00 38.06      H31H O
ATOM   5014  NE2 GLN   82      25.068  40.586   5.794  1.00 44.28      H31H N
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ATOM   5015  C   GLN   82      23.291  38.408   7.219  1.00 42.93      H31H C
ATOM   5016  O   GLN   82      22.388  37.803   6.666  1.00 45.25      H31H O
ATOM   5017  N   MET   83      23.083  39.489   7.958  1.00 44.37      H31H N
ATOM   5018  CA  MET   83      21.722  39.984   8.172  1.00 46.84      H31H C
ATOM   5019  CB  MET   83      21.500  40.300   9.651  1.00 48.94      H31H C
ATOM   5020  CG  MET   83      21.310  39.076  10.530  1.00 51.24      H31H C
ATOM   5021  SD  MET   83      21.149  39.501  12.271  1.00 54.99      H31H S
ATOM   5022  CE  MET   83      22.766  40.207  12.592  1.00 49.92      H31H C
ATOM   5023  C   MET   83      21.410  41.222   7.330  1.00 46.55      H31H C
ATOM   5024  O   MET   83      22.081  42.247   7.434  1.00 47.66      H31H O
ATOM   5025  N   ASN   84      20.389  41.119   6.494  1.00 45.09      H31H N
ATOM   5026  CA  ASN   84      20.077  42.179   5.558  1.00 44.96      H31H C
ATOM   5027  CB  ASN   84      20.227  41.676   4.119  1.00 46.67      H31H C
ATOM   5028  CG  ASN   84      21.666  41.478   3.722  1.00 49.33      H31H C
ATOM   5029  OD1 ASN   84      22.441  42.440   3.639  1.00 50.57      H31H O
ATOM   5030  ND2 ASN   84      22.045  40.226   3.475  1.00 50.44      H31H N
ATOM   5031  C   ASN   84      18.653  42.614   5.790  1.00 44.59      H31H C
ATOM   5032  O   ASN   84      17.833  41.829   6.257  1.00 42.65      H31H O
ATOM   5033  N   SER   85      18.356  43.861   5.443  1.00 44.77      H31H N
ATOM   5034  CA  SER   85      16.991  44.357   5.525  1.00 45.45      H31H C
ATOM   5035  CB  SER   85      16.067  43.579   4.595  1.00 42.38      H31H C
ATOM   5036  OG  SER   85      16.474  43.734   3.259  1.00 40.36      H31H O
ATOM   5037  C   SER   85      16.488  44.194   6.935  1.00 47.12      H31H C
ATOM   5038  O   SER   85      15.347  43.761   7.151  1.00 47.21      H31H O
ATOM   5039  N   LEU   86      17.336  44.529   7.896  1.00 47.75      H31H N
ATOM   5040  CA  LEU   86      16.960  44.316   9.266  1.00 51.11      H31H C
ATOM   5041  CB  LEU   86      18.081  44.791  10.184  1.00 47.93      H31H C
ATOM   5042  CG  LEU   86      19.154  43.706  10.351  1.00 45.15      H31H C
ATOM   5043  CD1 LEU   86      20.334  44.222  11.166  1.00 43.75      H31H C
ATOM   5044  CD2 LEU   86      18.526  42.496  11.017  1.00 41.27      H31H C
ATOM   5045  C   LEU   86      15.629  45.008   9.578  1.00 55.36      H31H C
ATOM   5046  O   LEU   86      15.174  45.903   8.854  1.00 55.80      H31H O
ATOM   5047  N   ARG   87      14.987  44.543  10.639  1.00 59.60      H31H N
ATOM   5048  CA  ARG   87      13.719  45.090  11.079  1.00 63.34      H31H C
ATOM   5049  CB  ARG   87      12.580  44.161  10.673  1.00 63.61      H31H C
ATOM   5050  CG  ARG   87      12.534  43.801   9.202  1.00 66.77      H31H C
ATOM   5051  CD  ARG   87      11.671  42.563   9.001  1.00 71.73      H31H C
ATOM   5052  NE  ARG   87      11.220  42.384   7.620  1.00 76.44      H31H N
ATOM   5053  CZ  ARG   87      10.722  41.246   7.134  1.00 78.44      H31H C
ATOM   5054  NH1 ARG   87      10.333  41.178   5.866  1.00 79.88      H31H N
ATOM   5055  NH2 ARG   87      10.621  40.172   7.910  1.00 77.64      H31H N
ATOM   5056  C   ARG   87      13.795  45.179  12.603  1.00 66.43      H31H C
ATOM   5057  O   ARG   87      14.652  44.544  13.222  1.00 66.59      H31H O
ATOM   5058  N   ALA   88      12.911  45.970  13.210  1.00 69.22      H31H N
ATOM   5059  CA  ALA   88      12.838  46.026  14.669  1.00 70.35      H31H C
ATOM   5060  CB  ALA   88      11.691  46.937  15.101  1.00 71.65      H31H C
ATOM   5061  C   ALA   88      12.626  44.603  15.208  1.00 71.16      H31H C
ATOM   5062  O   ALA   88      13.061  44.263  16.317  1.00 70.52      H31H O
ATOM   5063  N   GLU   89      11.979  43.778  14.386  1.00 70.86      H31H N
ATOM   5064  CA  GLU   89      11.535  42.442  14.767  1.00 70.01      H31H C
ATOM   5065  CB  GLU   89      10.529  41.912  13.743  1.00 72.23      H31H C
ATOM   5066  CG  GLU   89       9.345  42.832  13.487  1.00 79.51      H31H C
ATOM   5067  CD  GLU   89       9.752  44.288  13.250  1.00 83.99      H31H C
ATOM   5068  OE1 GLU   89       9.627  45.104  14.191  1.00 86.96      H31H O
ATOM   5069  OE2 GLU   89      10.198  44.620  12.127  1.00 86.67      H31H O
ATOM   5070  C   GLU   89      12.706  41.477  14.859  1.00 66.80      H31H C
ATOM   5071  O   GLU   89      12.559  40.355  15.325  1.00 68.37      H31H O
ATOM   5072  N   ASP   90      13.871  41.906  14.404  1.00 61.86      H31H N
ATOM   5073  CA  ASP   90      15.043  41.058  14.496  1.00 58.04      H31H C
ATOM   5074  CB  ASP   90      15.880  41.127  13.207  1.00 56.31      H31H C
ATOM   5075  CG  ASP   90      15.065  40.877  11.949  1.00 54.61      H31H C
ATOM   5076  OD1 ASP   90      14.182  39.988  11.952  1.00 53.40      H31H O
ATOM   5077  OD2 ASP   90      15.326  41.580  10.948  1.00 52.88      H31H O
ATOM   5078  C   ASP   90      15.904  41.481  15.675  1.00 56.02      H31H C
ATOM   5079  O   ASP   90      17.039  41.038  15.808  1.00 56.72      H31H O
ATOM   5080  N   THR   91      15.397  42.346  16.540  1.00 54.77      H31H N
ATOM   5081  CA  THR   91      16.217  42.699  17.693  1.00 53.43      H31H C
ATOM   5082  CB  THR   91      15.783  44.029  18.353  1.00 53.62      H31H C
ATOM   5083  OG1 THR   91      15.867  45.102  17.400  1.00 52.95      H31H O
ATOM   5084  CG2 THR   91      16.703  44.349  19.531  1.00 52.72      H31H C
ATOM   5085  C   THR   91      16.113  41.580  18.712  1.00 51.54      H31H C
ATOM   5086  O   THR   91      15.016  41.114  19.023  1.00 52.16      H31H O
ATOM   5087  N   ALA   92      17.256  41.142  19.218  1.00 48.98      H31H N
ATOM   5088  CA  ALA   92      17.306  39.982  20.096  1.00 48.01      H31H C
ATOM   5089  CB  ALA   92      16.595  38.804  19.447  1.00 48.07      H31H C
ATOM   5090  C   ALA   92      18.752  39.615  20.385  1.00 47.42      H31H C
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ATOM	5091	O	ALA	92	19.667	40.032	19.660	1.00	46.91	H31H O
ATOM	5092	N	VAL	93	18.973	38.843	21.443	1.00	47.37	H31H N
ATOM	5093	CA	VAL	93	20.286	38.247	21.599	1.00	48.79	H31H C
ATOM	5094	CB	VAL	93	20.576	37.784	23.057	1.00	48.74	H31H C
ATOM	5095	CG1	VAL	93	21.909	37.018	23.110	1.00	46.43	H31H C
ATOM	5096	CG2	VAL	93	20.676	39.004	23.973	1.00	48.32	H31H C
ATOM	5097	C	VAL	93	20.340	37.073	20.634	1.00	48.44	H31H C
ATOM	5098	O	VAL	93	19.307	36.484	20.299	1.00	47.17	H31H O
ATOM	5099	N	TYR	94	21.550	36.779	20.163	1.00	48.07	H31H N
ATOM	5100	CA	TYR	94	21.765	35.885	19.039	1.00	46.52	H31H C
ATOM	5101	CB	TYR	94	22.207	36.690	17.811	1.00	45.40	H31H C
ATOM	5102	CG	TYR	94	21.060	37.203	16.970	1.00	44.60	H31H C
ATOM	5103	CD1	TYR	94	20.530	36.420	15.960	1.00	45.87	H31H C
ATOM	5104	CE1	TYR	94	19.435	36.836	15.214	1.00	45.29	H31H C
ATOM	5105	CD2	TYR	94	20.466	38.440	17.217	1.00	44.37	H31H C
ATOM	5106	CE2	TYR	94	19.357	38.868	16.469	1.00	43.30	H31H C
ATOM	5107	CZ	TYR	94	18.852	38.046	15.470	1.00	43.79	H31H C
ATOM	5108	OH	TYR	94	17.748	38.377	14.726	1.00	42.59	H31H O
ATOM	5109	C	TYR	94	22.849	34.899	19.428	1.00	47.42	H31H C
ATOM	5110	O	TYR	94	24.013	35.290	19.592	1.00	47.83	H31H O
ATOM	5111	N	PHE	95	22.464	33.628	19.582	1.00	46.90	H31H N
ATOM	5112	CA	PHE	95	23.404	32.567	19.947	1.00	45.06	H31H C
ATOM	5113	CB	PHE	95	22.786	31.652	20.997	1.00	44.52	H31H C
ATOM	5114	CG	PHE	95	22.223	32.371	22.185	1.00	44.23	H31H C
ATOM	5115	CD1	PHE	95	23.018	32.644	23.290	1.00	42.47	H31H C
ATOM	5116	CD2	PHE	95	20.885	32.724	22.225	1.00	43.04	H31H C
ATOM	5117	CE1	PHE	95	22.482	33.251	24.414	1.00	41.09	H31H C
ATOM	5118	CE2	PHE	95	20.347	33.333	23.351	1.00	41.79	H31H C
ATOM	5119	CZ	PHE	95	21.146	33.595	24.443	1.00	40.34	H31H C
ATOM	5120	C	PHE	95	23.772	31.710	18.738	1.00	45.62	H31H C
ATOM	5121	O	PHE	95	22.952	31.490	17.847	1.00	44.98	H31H O
ATOM	5122	N	CYS	96	24.999	31.201	18.713	1.00	46.61	H31H N
ATOM	5123	CA	CYS	96	25.268	29.992	17.943	1.00	47.13	H31H C
ATOM	5124	C	CYS	96	25.289	28.826	18.903	1.00	46.71	H31H C
ATOM	5125	O	CYS	96	25.674	28.979	20.065	1.00	48.53	H31H O
ATOM	5126	CB	CYS	96	26.613	30.064	17.213	1.00	48.93	H31H C
ATOM	5127	SG	CYS	96	27.983	30.710	18.215	1.00	51.77	H31H S
ATOM	5128	N	ALA	97	24.858	27.667	18.419	1.00	45.19	H31H N
ATOM	5129	CA	ALA	97	25.093	26.411	19.114	1.00	43.27	H31H C
ATOM	5130	CB	ALA	97	23.829	25.984	19.849	1.00	43.29	H31H C
ATOM	5131	C	ALA	97	25.502	25.350	18.089	1.00	42.74	H31H C
ATOM	5132	O	ALA	97	25.105	25.410	16.921	1.00	43.45	H31H O
ATOM	5133	N	ARG	98	26.291	24.375	18.522	1.00	40.05	H31H N
ATOM	5134	CA	ARG	98	26.621	23.242	17.668	1.00	35.65	H31H C
ATOM	5135	CB	ARG	98	28.018	22.754	17.967	1.00	32.17	H31H C
ATOM	5136	CG	ARG	98	28.119	22.079	19.301	1.00	31.31	H31H C
ATOM	5137	CD	ARG	98	29.532	21.582	19.568	1.00	33.70	H31H C
ATOM	5138	NE	ARG	98	29.505	20.508	20.553	1.00	35.51	H31H N
ATOM	5139	CZ	ARG	98	30.575	19.838	20.966	1.00	35.38	H31H C
ATOM	5140	NH1	ARG	98	31.776	20.127	20.484	1.00	35.88	H31H N
ATOM	5141	NH2	ARG	98	30.443	18.868	21.857	1.00	35.44	H31H N
ATOM	5142	C	ARG	98	25.660	22.086	17.877	1.00	35.56	H31H C
ATOM	5143	O	ARG	98	24.942	22.026	18.867	1.00	34.36	H31H O
ATOM	5144	N	ASP	99	25.656	21.173	16.918	1.00	36.76	H31H N
ATOM	5145	CA	ASP	99	25.133	19.830	17.108	1.00	37.67	H31H C
ATOM	5146	CB	ASP	99	23.600	19.794	16.903	1.00	40.46	H31H C
ATOM	5147	CG	ASP	99	23.156	20.217	15.498	1.00	43.61	H31H C
ATOM	5148	OD1	ASP	99	23.413	21.369	15.081	1.00	46.81	H31H O
ATOM	5149	OD2	ASP	99	22.519	19.392	14.808	1.00	47.79	H31H O
ATOM	5150	C	ASP	99	25.843	18.915	16.116	1.00	38.10	H31H C
ATOM	5151	O	ASP	99	25.848	19.191	14.922	1.00	38.27	H31H O
ATOM	5152	N	TYR	100	26.459	17.844	16.615	1.00	38.50	H31H N
ATOM	5153	CA	TYR	100	27.167	16.884	15.774	1.00	38.52	H31H C
ATOM	5154	CB	TYR	100	27.589	15.658	16.596	1.00	39.27	H31H C
ATOM	5155	CG	TYR	100	28.459	14.673	15.824	1.00	42.22	H31H C
ATOM	5156	CD1	TYR	100	29.841	14.817	15.770	1.00	42.22	H31H C
ATOM	5157	CE1	TYR	100	30.622	13.949	15.012	1.00	44.08	H31H C
ATOM	5158	CD2	TYR	100	27.887	13.629	15.105	1.00	42.57	H31H C
ATOM	5159	CE2	TYR	100	28.656	12.765	14.355	1.00	43.34	H31H C
ATOM	5160	CZ	TYR	100	30.014	12.929	14.308	1.00	44.31	H31H C
ATOM	5161	OH	TYR	100	30.743	12.066	13.525	1.00	47.41	H31H O
ATOM	5162	C	TYR	100	26.339	16.435	14.569	1.00	38.28	H31H C
ATOM	5163	O	TYR	100	25.145	16.127	14.681	1.00	39.03	H31H O
ATOM	5164	N	ASP	101	27.010	16.382	13.423	1.00	36.46	H31H N
ATOM	5165	CA	ASP	101	26.376	16.244	12.120	1.00	34.34	H31H C
ATOM	5166	CB	ASP	101	27.209	17.043	11.104	1.00	35.01	H31H C

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ATOM   5167  CG  ASP  101      26.672  16.953   9.702  1.00 35.89      H31H C
ATOM   5168  OD1 ASP  101      25.668  16.233   9.508  1.00 37.29      H31H O
ATOM   5169  OD2 ASP  101      27.255  17.603   8.794  1.00 36.30      H31H O
ATOM   5170  C   ASP  101      26.261  14.764  11.717  1.00 33.54      H31H C
ATOM   5171  O   ASP  101      27.227  14.139  11.288  1.00 34.19      H31H O
ATOM   5172  N   PHE  102      25.066  14.209  11.867  1.00 32.47      H31H N
ATOM   5173  CA  PHE  102      24.817  12.807  11.553  1.00 32.51      H31H C
ATOM   5174  CB  PHE  102      23.784  12.216  12.528  1.00 30.44      H31H C
ATOM   5175  CG  PHE  102      24.275  12.156  13.954  1.00 30.38      H31H C
ATOM   5176  CD1 PHE  102      25.064  11.100  14.387  1.00 29.51      H31H C
ATOM   5177  CD2 PHE  102      23.982  13.178  14.849  1.00 29.52      H31H C
ATOM   5178  CE1 PHE  102      25.551  11.073  15.686  1.00 29.97      H31H C
ATOM   5179  CE2 PHE  102      24.459  13.157  16.144  1.00 27.61      H31H C
ATOM   5180  CZ  PHE  102      25.244  12.112  16.567  1.00 28.39      H31H C
ATOM   5181  C   PHE  102      24.332  12.687  10.125  1.00 32.53      H31H C
ATOM   5182  O   PHE  102      23.776  11.597   9.776  1.00 33.61      H31H O
ATOM   5183  N   TRP  103      24.530  13.662   9.304  1.00 31.13      H31H N
ATOM   5184  CA  TRP  103      24.184  13.562   7.895  1.00 30.67      H31H C
ATOM   5185  CB  TRP  103      25.121  12.554   7.252  1.00 31.31      H31H C
ATOM   5186  CG  TRP  103      26.556  12.931   7.490  1.00 31.50      H31H C
ATOM   5187  CD2 TRP  103      27.346  13.801   6.686  1.00 29.89      H31H C
ATOM   5188  CE2 TRP  103      28.607  13.916   7.310  1.00 28.91      H31H C
ATOM   5189  CE3 TRP  103      27.112  14.496   5.491  1.00 28.60      H31H C
ATOM   5190  CD1 TRP  103      27.348  12.556   8.544  1.00 31.47      H31H C
ATOM   5191  NE1 TRP  103      28.578  13.147   8.443  1.00 28.09      H31H N
ATOM   5192  CZ2 TRP  103      29.634  14.700   6.779  1.00 27.83      H31H C
ATOM   5193  CZ3 TRP  103      28.132  15.274   4.960  1.00 27.26      H31H C
ATOM   5194  CH2 TRP  103      29.382  15.369   5.606  1.00 26.36      H31H C
ATOM   5195  C   TRP  103      22.714  13.168   7.704  1.00 29.90      H31H C
ATOM   5196  O   TRP  103      22.367  12.385   6.815  1.00 30.62      H31H O
ATOM   5197  N   SER  104      21.878  13.725   8.579  1.00 28.52      H31H N
ATOM   5198  CA  SER  104      20.420  13.590   8.576  1.00 28.62      H31H C
ATOM   5199  CB  SER  104      19.846  13.762   7.144  1.00 27.84      H31H C
ATOM   5200  OG  SER  104      19.749  12.558   6.387  1.00 27.27      H31H O
ATOM   5201  C   SER  104      19.918  12.298   9.222  1.00 29.81      H31H C
ATOM   5202  O   SER  104      18.702  12.069   9.311  1.00 31.22      H31H O
ATOM   5203  N   ALA  105      20.844  11.459   9.684  1.00 28.06      H31H N
ATOM   5204  CA  ALA  105      20.452  10.163  10.233  1.00 27.77      H31H C
ATOM   5205  CB  ALA  105      21.681   9.306  10.506  1.00 21.95      H31H C
ATOM   5206  C   ALA  105      19.667  10.370  11.525  1.00 28.00      H31H C
ATOM   5207  O   ALA  105      18.822   9.557  11.908  1.00 28.93      H31H O
ATOM   5208  N   TYR  106      19.957  11.480  12.187  1.00 29.04      H31H N
ATOM   5209  CA  TYR  106      19.459  11.743  13.522  1.00 29.36      H31H C
ATOM   5210  CB  TYR  106      20.144  10.823  14.532  1.00 27.64      H31H C
ATOM   5211  CG  TYR  106      19.677  11.081  15.939  1.00 26.39      H31H C
ATOM   5212  CD1 TYR  106      20.543  11.580  16.907  1.00 24.73      H31H C
ATOM   5213  CE1 TYR  106      20.087  11.839  18.199  1.00 26.54      H31H C
ATOM   5214  CD2 TYR  106      18.348  10.847  16.297  1.00 24.05      H31H C
ATOM   5215  CE2 TYR  106      17.885  11.098  17.574  1.00 24.69      H31H C
ATOM   5216  CZ  TYR  106      18.748  11.594  18.526  1.00 26.04      H31H C
ATOM   5217  OH  TYR  106      18.269  11.851  19.795  1.00 25.58      H31H O
ATOM   5218  C   TYR  106      19.773  13.189  13.858  1.00 30.98      H31H C
ATOM   5219  O   TYR  106      20.887  13.666  13.605  1.00 33.21      H31H O
ATOM   5220  N   TYR  107      18.793  13.892  14.415  1.00 32.40      H31H N
ATOM   5221  CA  TYR  107      19.002  15.270  14.847  1.00 31.17      H31H C
ATOM   5222  CB  TYR  107      17.799  16.120  14.440  1.00 30.89      H31H C
ATOM   5223  CG  TYR  107      17.611  16.171  12.936  1.00 30.51      H31H C
ATOM   5224  CD1 TYR  107      16.553  15.511  12.326  1.00 31.26      H31H C
ATOM   5225  CE1 TYR  107      16.430  15.479  10.946  1.00 31.80      H31H C
ATOM   5226  CD2 TYR  107      18.545  16.816  12.119  1.00 28.58      H31H C
ATOM   5227  CE2 TYR  107      18.432  16.792  10.743  1.00 28.43      H31H C
ATOM   5228  CZ  TYR  107      17.374  16.118  10.161  1.00 30.95      H31H C
ATOM   5229  OH  TYR  107      17.275  16.046   8.794  1.00 31.71      H31H O
ATOM   5230  C   TYR  107      19.257  15.337  16.351  1.00 31.17      H31H C
ATOM   5231  O   TYR  107      18.357  15.246  17.163  1.00 30.52      H31H O
ATOM   5232  N   ASP  108      20.521  15.470  16.714  1.00 34.73      H31H N
ATOM   5233  CA  ASP  108      20.916  15.519  18.112  1.00 35.64      H31H C
ATOM   5234  CB  ASP  108      22.427  15.347  18.208  1.00 36.20      H31H C
ATOM   5235  CG  ASP  108      22.883  15.015  19.607  1.00 36.50      H31H C
ATOM   5236  OD1 ASP  108      22.011  14.673  20.434  1.00 32.41      H31H O
ATOM   5237  OD2 ASP  108      24.108  15.097  19.870  1.00 39.02      H31H O
ATOM   5238  C   ASP  108      20.500  16.867  18.708  1.00 36.58      H31H C
ATOM   5239  O   ASP  108      20.003  17.745  17.988  1.00 39.14      H31H O
ATOM   5240  N   ALA  109      20.695  17.033  20.012  1.00 35.61      H31H N
ATOM   5241  CA  ALA  109      20.442  18.320  20.654  1.00 35.70      H31H C
ATOM   5242  CB  ALA  109      20.263  18.135  22.155  1.00 35.91      H31H C
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ATOM   5243  C   ALA   109      21.616  19.247  20.382  1.00 35.45      H31H C
ATOM   5244  O   ALA   109      22.711  18.781  20.079  1.00 37.43      H31H O
ATOM   5245  N   PHE   110      21.379  20.549  20.479  1.00 34.40      H31H N
ATOM   5246  CA  PHE   110      22.445  21.546  20.407  1.00 35.00      H31H C
ATOM   5247  CB  PHE   110      21.872  22.963  20.195  1.00 34.51      H31H C
ATOM   5248  CG  PHE   110      20.961  23.102  18.997  1.00 33.42      H31H C
ATOM   5249  CD1 PHE   110      20.489  24.342  18.618  1.00 33.80      H31H C
ATOM   5250  CD2 PHE   110      20.560  22.003  18.266  1.00 34.66      H31H C
ATOM   5251  CE1 PHE   110      19.638  24.482  17.544  1.00 33.62      H31H C
ATOM   5252  CE2 PHE   110      19.703  22.146  17.186  1.00 34.07      H31H C
ATOM   5253  CZ  PHE   110      19.246  23.382  16.831  1.00 33.99      H31H C
ATOM   5254  C   PHE   110      23.181  21.523  21.742  1.00 35.07      H31H C
ATOM   5255  O   PHE   110      22.827  22.255  22.665  1.00 36.68      H31H O
ATOM   5256  N   ASP   111      24.204  20.698  21.863  1.00 34.77      H31H N
ATOM   5257  CA  ASP   111      24.739  20.446  23.185  1.00 37.29      H31H C
ATOM   5258  CB  ASP   111      25.618  19.183  23.190  1.00 36.40      H31H C
ATOM   5259  CG  ASP   111      26.789  19.262  22.221  1.00 35.66      H31H C
ATOM   5260  OD1 ASP   111      26.798  20.147  21.339  1.00 34.96      H31H O
ATOM   5261  OD2 ASP   111      27.708  18.419  22.349  1.00 37.27      H31H O
ATOM   5262  C   ASP   111      25.501  21.613  23.790  1.00 38.76      H31H C
ATOM   5263  O   ASP   111      25.620  21.695  25.006  1.00 40.16      H31H O
ATOM   5264  N   VAL   112      26.017  22.516  22.963  1.00 40.27      H31H N
ATOM   5265  CA  VAL   112      26.888  23.577  23.473  1.00 41.93      H31H C
ATOM   5266  CB  VAL   112      28.369  23.296  23.147  1.00 43.13      H31H C
ATOM   5267  CG1 VAL   112      29.266  24.239  23.943  1.00 40.95      H31H C
ATOM   5268  CG2 VAL   112      28.709  21.828  23.435  1.00 40.65      H31H C
ATOM   5269  C   VAL   112      26.526  24.909  22.843  1.00 43.13      H31H C
ATOM   5270  O   VAL   112      26.478  25.009  21.616  1.00 44.68      H31H O
ATOM   5271  N   TRP   113      26.275  25.928  23.668  1.00 43.43      H31H N
ATOM   5272  CA  TRP   113      25.983  27.269  23.149  1.00 43.70      H31H C
ATOM   5273  CB  TRP   113      24.672  27.793  23.712  1.00 40.08      H31H C
ATOM   5274  CG  TRP   113      23.485  26.921  23.464  1.00 37.13      H31H C
ATOM   5275  CD2 TRP   113      22.154  27.366  23.183  1.00 34.61      H31H C
ATOM   5276  CE2 TRP   113      21.323  26.235  23.184  1.00 34.01      H31H C
ATOM   5277  CE3 TRP   113      21.588  28.613  22.936  1.00 34.32      H31H C
ATOM   5278  CD1 TRP   113      23.414  25.568  23.604  1.00 35.50      H31H C
ATOM   5279  NE1 TRP   113      22.112  25.147  23.444  1.00 35.84      H31H N
ATOM   5280  CZ2 TRP   113      19.955  26.313  22.946  1.00 36.16      H31H C
ATOM   5281  CZ3 TRP   113      20.224  28.690  22.699  1.00 36.33      H31H C
ATOM   5282  CH2 TRP   113      19.424  27.552  22.706  1.00 35.59      H31H C
ATOM   5283  C   TRP   113      27.071  28.316  23.429  1.00 45.51      H31H C
ATOM   5284  O   TRP   113      27.778  28.251  24.446  1.00 44.71      H31H O
ATOM   5285  N   GLY   114      27.193  29.282  22.519  1.00 46.53      H31H N
ATOM   5286  CA  GLY   114      27.996  30.458  22.795  1.00 49.42      H31H C
ATOM   5287  C   GLY   114      27.275  31.384  23.754  1.00 52.13      H31H C
ATOM   5288  O   GLY   114      26.102  31.175  24.058  1.00 51.42      H31H O
ATOM   5289  N   GLN   115      27.973  32.416  24.221  1.00 56.29      H31H N
ATOM   5290  CA  GLN   115      27.459  33.318  25.260  1.00 59.63      H31H C
ATOM   5291  CB  GLN   115      28.611  34.141  25.840  1.00 63.79      H31H C
ATOM   5292  CG  GLN   115      29.457  33.380  26.847  1.00 71.42      H31H C
ATOM   5293  CD  GLN   115      28.620  32.832  27.993  1.00 76.37      H31H C
ATOM   5294  OE1 GLN   115      28.448  33.496  29.022  1.00 79.37      H31H O
ATOM   5295  NE2 GLN   115      28.086  31.617  27.818  1.00 80.03      H31H N
ATOM   5296  C   GLN   115      26.337  34.260  24.820  1.00 58.74      H31H C
ATOM   5297  O   GLN   115      25.461  34.613  25.610  1.00 57.37      H31H O
ATOM   5298  N   GLY   116      26.374  34.666  23.559  1.00 58.43      H31H N
ATOM   5299  CA  GLY   116      25.326  35.507  23.019  1.00 58.13      H31H C
ATOM   5300  C   GLY   116      25.876  36.841  22.544  1.00 59.56      H31H C
ATOM   5301  O   GLY   116      26.974  37.260  22.938  1.00 59.14      H31H O
ATOM   5302  N   THR   117      25.110  37.514  21.690  1.00 60.54      H31H N
ATOM   5303  CA  THR   117      25.468  38.853  21.226  1.00 59.73      H31H C
ATOM   5304  CB  THR   117      26.301  38.779  19.926  1.00 60.22      H31H C
ATOM   5305  OG1 THR   117      27.299  39.801  19.943  1.00 60.73      H31H O
ATOM   5306  CG2 THR   117      25.410  38.972  18.710  1.00 59.64      H31H C
ATOM   5307  C   THR   117      24.165  39.616  20.975  1.00 58.58      H31H C
ATOM   5308  O   THR   117      23.145  39.014  20.634  1.00 55.70      H31H O
ATOM   5309  N   MET   118      24.188  40.931  21.175  1.00 58.61      H31H N
ATOM   5310  CA  MET   118      22.944  41.700  21.156  1.00 58.29      H31H C
ATOM   5311  CB  MET   118      22.865  42.626  22.377  1.00 55.41      H31H C
ATOM   5312  CG  MET   118      21.515  43.324  22.541  1.00 54.00      H31H C
ATOM   5313  SD  MET   118      20.114  42.172  22.578  1.00 54.63      H31H S
ATOM   5314  CE  MET   118      18.732  43.182  21.995  1.00 51.15      H31H C
ATOM   5315  C   MET   118      22.777  42.504  19.867  1.00 59.12      H31H C
ATOM   5316  O   MET   118      23.583  43.371  19.538  1.00 58.18      H31H O
ATOM   5317  N   VAL   119      21.724  42.188  19.129  1.00 61.17      H31H N
ATOM   5318  CA  VAL   119      21.431  42.899  17.907  1.00 63.75      H31H C
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ATOM   5319  CB  VAL   119      21.186  41.930  16.730  1.00 62.58      H31H C
ATOM   5320  CG1 VAL   119      20.577  42.678  15.548  1.00 61.18      H31H C
ATOM   5321  CG2 VAL   119      22.502  41.298  16.309  1.00 61.02      H31H C
ATOM   5322  C   VAL   119      20.191  43.728  18.145  1.00 66.03      H31H C
ATOM   5323  O   VAL   119      19.113  43.188  18.410  1.00 67.24      H31H O
ATOM   5324  N   THR   120      20.360  45.046  18.061  1.00 67.50      H31H N
ATOM   5325  CA  THR   120      19.261  45.983  18.247  1.00 66.26      H31H C
ATOM   5326  CB  THR   120      19.589  47.049  19.299  1.00 65.93      H31H C
ATOM   5327  OG1 THR   120      20.422  46.489  20.329  1.00 64.19      H31H O
ATOM   5328  CG2 THR   120      18.298  47.570  19.908  1.00 62.66      H31H C
ATOM   5329  C   THR   120      19.009  46.707  16.942  1.00 65.33      H31H C
ATOM   5330  O   THR   120      19.924  47.309  16.373  1.00 64.94      H31H O
ATOM   5331  N   VAL   121      17.767  46.651  16.478  1.00 64.42      H31H N
ATOM   5332  CA  VAL   121      17.370  47.370  15.282  1.00 64.31      H31H C
ATOM   5333  CB  VAL   121      16.748  46.407  14.242  1.00 64.18      H31H C
ATOM   5334  CG1 VAL   121      16.259  47.176  13.018  1.00 63.06      H31H C
ATOM   5335  CG2 VAL   121      17.771  45.370  13.839  1.00 62.69      H31H C
ATOM   5336  C   VAL   121      16.368  48.462  15.641  1.00 65.00      H31H C
ATOM   5337  O   VAL   121      15.201  48.181  15.953  1.00 64.07      H31H O
ATOM   5338  N   SER   122      16.837  49.709  15.610  1.00 65.59      H31H N
ATOM   5339  CA  SER   122      15.948  50.861  15.712  1.00 66.99      H31H C
ATOM   5340  CB  SER   122      15.411  51.015  17.134  1.00 67.36      H31H C
ATOM   5341  OG  SER   122      14.609  52.182  17.233  1.00 67.09      H31H O
ATOM   5342  C   SER   122      16.563  52.190  15.284  1.00 67.53      H31H C
ATOM   5343  O   SER   122      17.785  52.371  15.235  1.00 66.70      H31H O
ATOM   5344  N   SER   123      15.672  53.125  14.992  1.00 67.46      H31H N
ATOM   5345  CA  SER   123      16.044  54.457  14.568  1.00 65.91      H31H C
ATOM   5346  CB  SER   123      14.839  55.103  13.898  1.00 65.83      H31H C
ATOM   5347  OG  SER   123      13.660  54.388  14.237  1.00 63.82      H31H O
ATOM   5348  C   SER   123      16.526  55.306  15.741  1.00 64.19      H31H C
ATOM   5349  O   SER   123      17.133  56.350  15.543  1.00 64.69      H31H O
ATOM   5350  N   ALA   124      16.262  54.848  16.959  1.00 62.50      H31H N
ATOM   5351  CA  ALA   124      16.615  55.594  18.163  1.00 61.24      H31H C
ATOM   5352  CB  ALA   124      16.471  54.709  19.382  1.00 61.18      H31H C
ATOM   5353  C   ALA   124      18.016  56.184  18.124  1.00 61.63      H31H C
ATOM   5354  O   ALA   124      18.869  55.742  17.355  1.00 62.06      H31H O
ATOM   5355  N   SER   125      18.243  57.185  18.970  1.00 61.89      H31H N
ATOM   5356  CA  SER   125      19.499  57.927  18.985  1.00 61.67      H31H C
ATOM   5357  CB  SER   125      19.631  58.731  17.687  1.00 63.06      H31H C
ATOM   5358  OG  SER   125      20.889  59.378  17.586  1.00 65.82      H31H O
ATOM   5359  C   SER   125      19.496  58.854  20.208  1.00 60.83      H31H C
ATOM   5360  O   SER   125      18.436  59.271  20.668  1.00 60.45      H31H O
ATOM   5361  N   THR   126      20.683  59.165  20.725  1.00 59.55      H31H N
ATOM   5362  CA  THR   126      20.843  59.763  22.055  1.00 58.53      H31H C
ATOM   5363  CB  THR   126      22.361  60.096  22.324  1.00 59.12      H31H C
ATOM   5364  OG1 THR   126      22.496  60.854  23.539  1.00 57.87      H31H O
ATOM   5365  CG2 THR   126      22.968  60.874  21.156  1.00 57.69      H31H C
ATOM   5366  C   THR   126      19.973  61.005  22.314  1.00 57.27      H31H C
ATOM   5367  O   THR   126      20.164  62.051  21.693  1.00 58.39      H31H O
ATOM   5368  N   LYS   127      19.026  60.878  23.247  1.00 54.41      H31H N
ATOM   5369  CA  LYS   127      17.996  61.901  23.478  1.00 50.77      H31H C
ATOM   5370  CB  LYS   127      16.834  61.696  22.505  1.00 49.90      H31H C
ATOM   5371  CG  LYS   127      15.469  61.689  23.161  1.00 48.86      H31H C
ATOM   5372  CD  LYS   127      14.393  61.926  22.117  1.00 49.04      H31H C
ATOM   5373  CE  LYS   127      13.002  62.077  22.731  1.00 49.43      H31H C
ATOM   5374  NZ  LYS   127      12.866  63.275  23.602  1.00 48.62      H31H N
ATOM   5375  C   LYS   127      17.441  61.934  24.908  1.00 48.34      H31H C
ATOM   5376  O   LYS   127      17.107  60.895  25.480  1.00 48.08      H31H O
ATOM   5377  N   GLY   128      17.321  63.138  25.464  1.00 45.40      H31H N
ATOM   5378  CA  GLY   128      16.933  63.289  26.858  1.00 41.87      H31H C
ATOM   5379  C   GLY   128      15.431  63.204  27.066  1.00 40.01      H31H C
ATOM   5380  O   GLY   128      14.664  63.391  26.124  1.00 38.07      H31H O
ATOM   5381  N   PRO   129      14.978  62.914  28.296  1.00 40.21      H31H N
ATOM   5382  CD  PRO   129      15.828  62.783  29.492  1.00 40.34      H31H C
ATOM   5383  CA  PRO   129      13.566  62.665  28.617  1.00 39.64      H31H C
ATOM   5384  CB  PRO   129      13.636  61.979  29.967  1.00 38.84      H31H C
ATOM   5385  CG  PRO   129      14.824  62.610  30.603  1.00 39.50      H31H C
ATOM   5386  C   PRO   129      12.685  63.929  28.677  1.00 39.72      H31H C
ATOM   5387  O   PRO   129      13.155  65.020  28.963  1.00 38.60      H31H O
ATOM   5388  N   SER   130      11.401  63.753  28.397  1.00 40.04      H31H N
ATOM   5389  CA  SER   130      10.380  64.702  28.806  1.00 39.31      H31H C
ATOM   5390  CB  SER   130       9.172  64.635  27.857  1.00 39.01      H31H C
ATOM   5391  OG  SER   130       9.521  64.950  26.520  1.00 40.33      H31H O
ATOM   5392  C   SER   130       9.944  64.253  30.190  1.00 38.84      H31H C
ATOM   5393  O   SER   130       9.720  63.073  30.395  1.00 39.02      H31H O
ATOM   5394  N   VAL   131       9.812  65.179  31.133  1.00 38.94      H31H N
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ATOM   5395  CA  VAL   131       9.342  64.822  32.462  1.00 37.98      H31H C
ATOM   5396  CB  VAL   131      10.370  65.220  33.533  1.00 38.94      H31H C
ATOM   5397  CG1 VAL   131      10.189  64.377  34.794  1.00 37.15      H31H C
ATOM   5398  CG2 VAL   131      11.776  65.066  32.982  1.00 37.53      H31H C
ATOM   5399  C   VAL   131       8.010  65.493  32.798  1.00 39.20      H31H C
ATOM   5400  O   VAL   131       7.889  66.717  32.855  1.00 39.59      H31H O
ATOM   5401  N   PHE   132       6.997  64.686  33.036  1.00 40.76      H31H N
ATOM   5402  CA  PHE   132       5.727  65.231  33.454  1.00 41.89      H31H C
ATOM   5403  CB  PHE   132       4.625  64.733  32.531  1.00 45.07      H31H C
ATOM   5404  CG  PHE   132       4.873  65.039  31.086  1.00 47.51      H31H C
ATOM   5405  CD1 PHE   132       4.524  66.274  30.556  1.00 45.96      H31H C
ATOM   5406  CD2 PHE   132       5.457  64.091  30.255  1.00 47.99      H31H C
ATOM   5407  CE1 PHE   132       4.752  66.554  29.224  1.00 47.37      H31H C
ATOM   5408  CE2 PHE   132       5.687  64.370  28.921  1.00 49.35      H31H C
ATOM   5409  CZ  PHE   132       5.332  65.605  28.405  1.00 48.08      H31H C
ATOM   5410  C   PHE   132       5.468  64.782  34.867  1.00 42.41      H31H C
ATOM   5411  O   PHE   132       5.802  63.667  35.239  1.00 40.02      H31H O
ATOM   5412  N   PRO   133       4.874  65.651  35.681  1.00 44.73      H31H N
ATOM   5413  CD  PRO   133       4.401  67.008  35.358  1.00 42.60      H31H C
ATOM   5414  CA  PRO   133       4.565  65.279  37.059  1.00 45.76      H31H C
ATOM   5415  CB  PRO   133       4.235  66.612  37.709  1.00 44.76      H31H C
ATOM   5416  CG  PRO   133       3.636  67.410  36.595  1.00 41.73      H31H C
ATOM   5417  C   PRO   133       3.374  64.345  37.062  1.00 47.64      H31H C
ATOM   5418  O   PRO   133       2.421  64.559  36.325  1.00 49.24      H31H O
ATOM   5419  N   LEU   134       3.430  63.306  37.880  1.00 48.88      H31H N
ATOM   5420  CA  LEU   134       2.235  62.537  38.190  1.00 50.70      H31H C
ATOM   5421  CB  LEU   134       2.554  61.046  38.207  1.00 50.15      H31H C
ATOM   5422  CG  LEU   134       3.341  60.531  36.999  1.00 49.91      H31H C
ATOM   5423  CD1 LEU   134       3.959  59.175  37.321  1.00 48.75      H31H C
ATOM   5424  CD2 LEU   134       2.424  60.455  35.788  1.00 49.17      H31H C
ATOM   5425  C   LEU   134       1.798  62.984  39.575  1.00 52.49      H31H C
ATOM   5426  O   LEU   134       2.431  62.629  40.567  1.00 53.95      H31H O
ATOM   5427  N   ALA   135       0.726  63.768  39.639  1.00 54.85      H31H N
ATOM   5428  CA  ALA   135       0.292  64.375  40.892  1.00 57.86      H31H C
ATOM   5429  CB  ALA   135      -0.610  65.564  40.606  1.00 54.42      H31H C
ATOM   5430  C   ALA   135      -0.436  63.387  41.784  1.00 60.60      H31H C
ATOM   5431  O   ALA   135      -0.947  62.377  41.318  1.00 60.28      H31H O
ATOM   5432  N   PRO   136      -0.488  63.674  43.088  1.00 64.55      H31H N
ATOM   5433  CD  PRO   136       0.300  64.741  43.725  1.00 67.29      H31H C
ATOM   5434  CA  PRO   136      -1.275  62.919  44.067  1.00 69.15      H31H C
ATOM   5435  CB  PRO   136      -0.755  63.421  45.409  1.00 69.65      H31H C
ATOM   5436  CG  PRO   136      -0.256  64.790  45.123  1.00 68.88      H31H C
ATOM   5437  C   PRO   136      -2.775  63.155  43.917  1.00 72.89      H31H C
ATOM   5438  O   PRO   136      -3.215  64.290  43.691  1.00 73.33      H31H O
ATOM   5439  N   SER   137      -3.552  62.080  44.058  1.00 77.84      H31H N
ATOM   5440  CA  SER   137      -5.003  62.131  43.878  1.00 82.41      H31H C
ATOM   5441  CB  SER   137      -5.583  60.712  43.864  1.00 82.42      H31H C
ATOM   5442  OG  SER   137      -5.241  60.017  45.050  1.00 83.20      H31H O
ATOM   5443  C   SER   137      -5.682  62.948  44.973  1.00 84.96      H31H C
ATOM   5444  O   SER   137      -5.100  63.201  46.028  1.00 87.33      H31H O
ATOM   5445  N   GLY   144      -3.876  56.071  54.474  1.00 84.33      H31H N
ATOM   5446  CA  GLY   144      -3.690  57.342  55.159  1.00 83.90      H31H C
ATOM   5447  C   GLY   144      -2.769  58.322  54.437  1.00 82.39      H31H C
ATOM   5448  O   GLY   144      -2.568  59.450  54.882  1.00 82.07      H31H O
ATOM   5449  N   THR   145      -2.202  57.898  53.316  1.00 80.81      H31H N
ATOM   5450  CA  THR   145      -1.246  58.729  52.603  1.00 79.74      H31H C
ATOM   5451  CB  THR   145       0.194  58.274  52.897  1.00 81.45      H31H C
ATOM   5452  OG1 THR   145       0.388  56.939  52.413  1.00 82.63      H31H O
ATOM   5453  CG2 THR   145       0.460  58.315  54.389  1.00 81.35      H31H C
ATOM   5454  C   THR   145      -1.490  58.699  51.093  1.00 78.23      H31H C
ATOM   5455  O   THR   145      -2.372  57.985  50.607  1.00 76.79      H31H O
ATOM   5456  N   ALA   146      -0.708  59.486  50.357  1.00 76.75      H31H N
ATOM   5457  CA  ALA   146      -0.910  59.648  48.922  1.00 74.80      H31H C
ATOM   5458  CB  ALA   146      -1.355  61.071  48.617  1.00 74.54      H31H C
ATOM   5459  C   ALA   146       0.359  59.320  48.151  1.00 72.36      H31H C
ATOM   5460  O   ALA   146       1.468  59.474  48.662  1.00 72.82      H31H O
ATOM   5461  N   ALA   147       0.187  58.857  46.917  1.00 68.95      H31H N
ATOM   5462  CA  ALA   147       1.320  58.517  46.062  1.00 64.43      H31H C
ATOM   5463  CB  ALA   147       1.154  57.116  45.501  1.00 65.16      H31H C
ATOM   5464  C   ALA   147       1.447  59.513  44.921  1.00 61.83      H31H C
ATOM   5465  O   ALA   147       0.531  59.677  44.116  1.00 61.04      H31H O
ATOM   5466  N   LEU   148       2.587  60.180  44.851  1.00 58.23      H31H N
ATOM   5467  CA  LEU   148       2.890  60.984  43.689  1.00 55.96      H31H C
ATOM   5468  CB  LEU   148       3.086  62.448  44.072  1.00 55.54      H31H C
ATOM   5469  CG  LEU   148       4.068  62.755  45.194  1.00 54.79      H31H C
ATOM   5470  CD1 LEU   148       5.140  63.728  44.718  1.00 57.06      H31H C
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ATOM   5471  CD2 LEU   148       3.289  63.337  46.346  1.00 54.72      H31H C
ATOM   5472  C   LEU   148       4.149  60.448  43.063  1.00 54.70      H31H C
ATOM   5473  O   LEU   148       4.822  59.596  43.633  1.00 54.10      H31H O
ATOM   5474  N   GLY   149       4.462  60.946  41.879  1.00 53.02      H31H N
ATOM   5475  CA  GLY   149       5.675  60.521  41.221  1.00 51.68      H31H C
ATOM   5476  C   GLY   149       5.905  61.418  40.034  1.00 49.79      H31H C
ATOM   5477  O   GLY   149       5.166  62.368  39.832  1.00 50.10      H31H O
ATOM   5478  N   CYS   150       6.927  61.131  39.246  1.00 48.52      H31H N
ATOM   5479  CA  CYS   150       7.064  61.821  37.986  1.00 47.23      H31H C
ATOM   5480  C   CYS   150       7.447  60.877  36.839  1.00 43.82      H31H C
ATOM   5481  O   CYS   150       8.290  60.002  36.996  1.00 43.71      H31H O
ATOM   5482  CB  CYS   150       8.048  62.988  38.150  1.00 49.36      H31H C
ATOM   5483  SG  CYS   150       9.812  62.571  38.243  1.00 56.29      H31H S
ATOM   5484  N   LEU   151       6.772  61.050  35.701  1.00 41.91      H31H N
ATOM   5485  CA  LEU   151       6.853  60.144  34.545  1.00 40.54      H31H C
ATOM   5486  CB  LEU   151       5.505  60.099  33.823  1.00 40.30      H31H C
ATOM   5487  CG  LEU   151       5.616  59.625  32.371  1.00 40.58      H31H C
ATOM   5488  CD1 LEU   151       6.148  58.202  32.343  1.00 42.03      H31H C
ATOM   5489  CD2 LEU   151       4.256  59.696  31.685  1.00 40.84      H31H C
ATOM   5490  C   LEU   151       7.924  60.558  33.533  1.00 40.01      H31H C
ATOM   5491  O   LEU   151       7.765  61.548  32.833  1.00 39.20      H31H O
ATOM   5492  N   VAL   152       9.000  59.789  33.444  1.00 40.24      H31H N
ATOM   5493  CA  VAL   152      10.169  60.190  32.664  1.00 41.29      H31H C
ATOM   5494  CB  VAL   152      11.463  59.823  33.443  1.00 40.87      H31H C
ATOM   5495  CG1 VAL   152      12.663  60.541  32.856  1.00 41.31      H31H C
ATOM   5496  CG2 VAL   152      11.292  60.147  34.921  1.00 38.03      H31H C
ATOM   5497  C   VAL   152      10.164  59.463  31.312  1.00 42.74      H31H C
ATOM   5498  O   VAL   152      10.623  58.321  31.225  1.00 45.26      H31H O
ATOM   5499  N   LYS   153       9.654  60.098  30.257  1.00 43.18      H31H N
ATOM   5500  CA  LYS   153       9.476  59.375  28.991  1.00 45.09      H31H C
ATOM   5501  CB  LYS   153       8.035  59.501  28.491  1.00 44.98      H31H C
ATOM   5502  CG  LYS   153       7.594  60.909  28.168  1.00 46.95      H31H C
ATOM   5503  CD  LYS   153       6.077  60.974  27.919  1.00 49.28      H31H C
ATOM   5504  CE  LYS   153       5.711  60.748  26.441  1.00 49.98      H31H C
ATOM   5505  NZ  LYS   153       4.315  61.196  26.093  1.00 48.65      H31H N
ATOM   5506  C   LYS   153      10.425  59.736  27.848  1.00 45.72      H31H C
ATOM   5507  O   LYS   153      11.028  60.813  27.822  1.00 46.61      H31H O
ATOM   5508  N   ASP   154      10.568  58.798  26.920  1.00 45.12      H31H N
ATOM   5509  CA  ASP   154      11.169  59.055  25.618  1.00 45.38      H31H C
ATOM   5510  CB  ASP   154      10.349  60.101  24.882  1.00 44.90      H31H C
ATOM   5511  CG  ASP   154       8.960  59.602  24.521  1.00 48.08      H31H C
ATOM   5512  OD1 ASP   154       8.805  58.376  24.299  1.00 49.40      H31H O
ATOM   5513  OD2 ASP   154       8.019  60.434  24.455  1.00 51.13      H31H O
ATOM   5514  C   ASP   154      12.639  59.453  25.601  1.00 45.46      H31H C
ATOM   5515  O   ASP   154      13.013  60.401  24.932  1.00 44.22      H31H O
ATOM   5516  N   TYR   155      13.476  58.712  26.318  1.00 47.16      H31H N
ATOM   5517  CA  TYR   155      14.921  58.918  26.259  1.00 49.79      H31H C
ATOM   5518  CB  TYR   155      15.467  59.204  27.659  1.00 51.22      H31H C
ATOM   5519  CG  TYR   155      15.315  58.042  28.623  1.00 54.62      H31H C
ATOM   5520  CD1 TYR   155      16.430  57.412  29.156  1.00 55.10      H31H C
ATOM   5521  CE1 TYR   155      16.294  56.327  30.002  1.00 57.77      H31H C
ATOM   5522  CD2 TYR   155      14.055  57.556  28.970  1.00 54.59      H31H C
ATOM   5523  CE2 TYR   155      13.909  56.479  29.811  1.00 55.80      H31H C
ATOM   5524  CZ  TYR   155      15.031  55.860  30.325  1.00 57.60      H31H C
ATOM   5525  OH  TYR   155      14.905  54.749  31.139  1.00 59.92      H31H O
ATOM   5526  C   TYR   155      15.669  57.716  25.654  1.00 50.37      H31H C
ATOM   5527  O   TYR   155      15.057  56.698  25.296  1.00 48.82      H31H O
ATOM   5528  N   PHE   156      16.991  57.866  25.549  1.00 49.83      H31H N
ATOM   5529  CA  PHE   156      17.889  56.872  24.984  1.00 50.40      H31H C
ATOM   5530  CB  PHE   156      17.494  56.525  23.545  1.00 51.14      H31H C
ATOM   5531  CG  PHE   156      18.311  55.411  22.949  1.00 52.08      H31H C
ATOM   5532  CD1 PHE   156      19.596  55.651  22.470  1.00 53.14      H31H C
ATOM   5533  CD2 PHE   156      17.818  54.112  22.922  1.00 52.74      H31H C
ATOM   5534  CE1 PHE   156      20.379  54.618  21.982  1.00 54.59      H31H C
ATOM   5535  CE2 PHE   156      18.592  53.068  22.437  1.00 54.30      H31H C
ATOM   5536  CZ  PHE   156      19.875  53.317  21.966  1.00 54.91      H31H C
ATOM   5537  C   PHE   156      19.306  57.436  24.975  1.00 52.12      H31H C
ATOM   5538  O   PHE   156      19.518  58.620  24.715  1.00 51.42      H31H O
ATOM   5539  N   PRO   157      20.303  56.586  25.246  1.00 52.79      H31H N
ATOM   5540  CD  PRO   157      21.723  56.973  25.178  1.00 55.23      H31H C
ATOM   5541  CA  PRO   157      20.151  55.191  25.668  1.00 56.36      H31H C
ATOM   5542  CB  PRO   157      21.487  54.576  25.285  1.00 54.78      H31H C
ATOM   5543  CG  PRO   157      22.456  55.688  25.531  1.00 54.58      H31H C
ATOM   5544  C   PRO   157      19.921  55.158  27.179  1.00 58.64      H31H C
ATOM   5545  O   PRO   157      19.828  56.220  27.807  1.00 60.21      H31H O
ATOM   5546  N   GLU   158      19.850  53.960  27.761  1.00 59.18      H31H N
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ATOM   5547  CA  GLU   158      19.939  53.803  29.217  1.00 60.12      H31H C
ATOM   5548  CB  GLU   158      19.800  52.330  29.593  1.00 60.44      H31H C
ATOM   5549  CG  GLU   158      18.442  51.733  29.306  1.00 61.07      H31H C
ATOM   5550  CD  GLU   158      17.480  51.975  30.436  1.00 61.97      H31H C
ATOM   5551  OE1 GLU   158      17.604  53.037  31.095  1.00 62.07      H31H O
ATOM   5552  OE2 GLU   158      16.611  51.105  30.668  1.00 61.87      H31H O
ATOM   5553  C   GLU   158      21.266  54.330  29.781  1.00 59.69      H31H C
ATOM   5554  O   GLU   158      22.245  54.482  29.057  1.00 60.37      H31H O
ATOM   5555  N   PRO   159      21.315  54.619  31.091  1.00 58.97      H31H N
ATOM   5556  CD  PRO   159      22.618  54.767  31.768  1.00 59.60      H31H C
ATOM   5557  CA  PRO   159      20.205  54.718  32.051  1.00 58.30      H31H C
ATOM   5558  CB  PRO   159      20.813  54.145  33.319  1.00 60.20      H31H C
ATOM   5559  CG  PRO   159      22.263  54.657  33.248  1.00 60.02      H31H C
ATOM   5560  C   PRO   159      19.769  56.184  32.257  1.00 56.56      H31H C
ATOM   5561  O   PRO   159      20.471  57.112  31.838  1.00 55.30      H31H O
ATOM   5562  N   VAL   160      18.630  56.394  32.913  1.00 55.04      H31H N
ATOM   5563  CA  VAL   160      18.411  57.644  33.649  1.00 54.39      H31H C
ATOM   5564  CB  VAL   160      16.999  58.228  33.447  1.00 55.26      H31H C
ATOM   5565  CG1 VAL   160      16.883  58.885  32.090  1.00 58.59      H31H C
ATOM   5566  CG2 VAL   160      15.970  57.133  33.612  1.00 55.92      H31H C
ATOM   5567  C   VAL   160      18.552  57.384  35.131  1.00 53.67      H31H C
ATOM   5568  O   VAL   160      18.305  56.277  35.607  1.00 53.08      H31H O
ATOM   5569  N   THR   161      18.933  58.402  35.878  1.00 50.80      H31H N
ATOM   5570  CA  THR   161      18.797  58.269  37.305  1.00 49.44      H31H C
ATOM   5571  CB  THR   161      20.164  58.345  37.983  1.00 49.51      H31H C
ATOM   5572  OG1 THR   161      20.751  59.632  37.771  1.00 50.02      H31H O
ATOM   5573  CG2 THR   161      21.073  57.297  37.382  1.00 50.69      H31H C
ATOM   5574  C   THR   161      17.836  59.318  37.839  1.00 49.75      H31H C
ATOM   5575  O   THR   161      17.912  60.497  37.492  1.00 49.46      H31H O
ATOM   5576  N   VAL   162      16.900  58.867  38.662  1.00 49.63      H31H N
ATOM   5577  CA  VAL   162      15.885  59.750  39.218  1.00 48.46      H31H C
ATOM   5578  CB  VAL   162      14.470  59.172  39.007  1.00 50.10      H31H C
ATOM   5579  CG1 VAL   162      13.426  60.152  39.528  1.00 51.21      H31H C
ATOM   5580  CG2 VAL   162      14.253  58.865  37.527  1.00 49.13      H31H C
ATOM   5581  C   VAL   162      16.123  59.923  40.707  1.00 46.96      H31H C
ATOM   5582  O   VAL   162      16.495  58.988  41.399  1.00 44.46      H31H O
ATOM   5583  N   SER   163      15.909  61.131  41.196  1.00 47.63      H31H N
ATOM   5584  CA  SER   163      16.055  61.397  42.619  1.00 47.37      H31H C
ATOM   5585  CB  SER   163      17.421  62.010  42.892  1.00 45.40      H31H C
ATOM   5586  OG  SER   163      17.530  62.317  44.262  1.00 47.31      H31H O
ATOM   5587  C   SER   163      14.966  62.336  43.125  1.00 46.40      H31H C
ATOM   5588  O   SER   163      14.323  63.032  42.339  1.00 47.13      H31H O
ATOM   5589  N   TRP   164      14.754  62.361  44.435  1.00 44.44      H31H N
ATOM   5590  CA  TRP   164      13.855  63.359  45.007  1.00 42.49      H31H C
ATOM   5591  CB  TRP   164      12.660  62.675  45.676  1.00 36.70      H31H C
ATOM   5592  CG  TRP   164      11.840  61.877  44.694  1.00 34.26      H31H C
ATOM   5593  CD2 TRP   164      10.677  62.318  43.979  1.00 34.05      H31H C
ATOM   5594  CE2 TRP   164      10.229  61.228  43.192  1.00 34.15      H31H C
ATOM   5595  CE3 TRP   164       9.969  63.528  43.925  1.00 32.87      H31H C
ATOM   5596  CD1 TRP   164      12.045  60.576  44.315  1.00 33.54      H31H C
ATOM   5597  NE1 TRP   164      11.082  60.179  43.416  1.00 30.36      H31H N
ATOM   5598  CZ2 TRP   164       9.103  61.317  42.362  1.00 33.86      H31H C
ATOM   5599  CZ3 TRP   164       8.855  63.613  43.102  1.00 32.86      H31H C
ATOM   5600  CH2 TRP   164       8.432  62.512  42.331  1.00 34.03      H31H C
ATOM   5601  C   TRP   164      14.559  64.302  45.981  1.00 43.04      H31H C
ATOM   5602  O   TRP   164      15.408  63.894  46.783  1.00 41.69      H31H O
ATOM   5603  N   ASN   165      14.211  65.580  45.882  1.00 44.35      H31H N
ATOM   5604  CA  ASN   165      14.805  66.593  46.735  1.00 43.74      H31H C
ATOM   5605  CB  ASN   165      14.254  66.453  48.145  1.00 45.04      H31H C
ATOM   5606  CG  ASN   165      12.780  66.757  48.205  1.00 46.52      H31H C
ATOM   5607  OD1 ASN   165      12.330  67.728  47.603  1.00 49.82      H31H O
ATOM   5608  ND2 ASN   165      12.016  65.933  48.920  1.00 45.08      H31H N
ATOM   5609  C   ASN   165      16.320  66.473  46.728  1.00 43.80      H31H C
ATOM   5610  O   ASN   165      16.996  66.852  47.687  1.00 43.40      H31H O
ATOM   5611  N   SER   166      16.848  65.939  45.632  1.00 43.57      H31H N
ATOM   5612  CA  SER   166      18.277  66.007  45.387  1.00 44.05      H31H C
ATOM   5613  CB  SER   166      18.802  67.385  45.778  1.00 44.33      H31H C
ATOM   5614  OG  SER   166      18.185  68.364  44.978  1.00 48.35      H31H O
ATOM   5615  C   SER   166      19.030  64.947  46.164  1.00 42.94      H31H C
ATOM   5616  O   SER   166      20.245  65.039  46.326  1.00 41.74      H31H O
ATOM   5617  N   GLY   167      18.301  63.952  46.655  1.00 42.43      H31H N
ATOM   5618  CA  GLY   167      18.929  62.897  47.418  1.00 41.74      H31H C
ATOM   5619  C   GLY   167      18.292  62.682  48.772  1.00 42.63      H31H C
ATOM   5620  O   GLY   167      18.144  61.533  49.209  1.00 43.39      H31H O
ATOM   5621  N   ALA   168      17.896  63.769  49.436  1.00 42.33      H31H N
ATOM   5622  CA  ALA   168      17.324  63.656  50.778  1.00 40.60      H31H C
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ATOM   5623  CB  ALA  168      17.054  65.021  51.346  1.00 39.45      H31H C
ATOM   5624  C   ALA  168      16.061  62.799  50.846  1.00 39.62      H31H C
ATOM   5625  O   ALA  168      15.870  62.080  51.809  1.00 39.68      H31H O
ATOM   5626  N   LEU  169      15.204  62.846  49.830  1.00 39.52      H31H N
ATOM   5627  CA  LEU  169      13.955  62.089  49.897  1.00 40.32      H31H C
ATOM   5628  CB  LEU  169      12.848  62.777  49.076  1.00 38.82      H31H C
ATOM   5629  CG  LEU  169      11.506  62.016  49.118  1.00 38.57      H31H C
ATOM   5630  CD1 LEU  169      11.201  61.626  50.544  1.00 36.39      H31H C
ATOM   5631  CD2 LEU  169      10.372  62.854  48.547  1.00 38.11      H31H C
ATOM   5632  C   LEU  169      14.109  60.636  49.437  1.00 42.25      H31H C
ATOM   5633  O   LEU  169      13.874  60.301  48.268  1.00 41.44      H31H O
ATOM   5634  N   THR  170      14.479  59.759  50.363  1.00 44.70      H31H N
ATOM   5635  CA  THR  170      14.772  58.391  49.961  1.00 46.53      H31H C
ATOM   5636  CB  THR  170      16.187  57.944  50.439  1.00 45.23      H31H C
ATOM   5637  OG1 THR  170      16.368  58.249  51.831  1.00 46.22      H31H O
ATOM   5638  CG2 THR  170      17.262  58.664  49.625  1.00 45.00      H31H C
ATOM   5639  C   THR  170      13.731  57.367  50.373  1.00 47.49      H31H C
ATOM   5640  O   THR  170      13.345  56.528  49.568  1.00 49.33      H31H O
ATOM   5641  N   SER  171      13.250  57.431  51.604  1.00 48.12      H31H N
ATOM   5642  CA  SER  171      12.273  56.440  52.035  1.00 49.02      H31H C
ATOM   5643  CB  SER  171      11.928  56.639  53.502  1.00 50.34      H31H C
ATOM   5644  OG  SER  171      11.290  57.891  53.670  1.00 49.96      H31H O
ATOM   5645  C   SER  171      11.005  56.580  51.206  1.00 48.93      H31H C
ATOM   5646  O   SER  171      10.606  57.690  50.859  1.00 50.01      H31H O
ATOM   5647  N   GLY  172      10.370  55.458  50.887  1.00 48.36      H31H N
ATOM   5648  CA  GLY  172       9.101  55.513  50.184  1.00 46.99      H31H C
ATOM   5649  C   GLY  172       9.206  55.745  48.687  1.00 45.95      H31H C
ATOM   5650  O   GLY  172       8.187  55.845  48.015  1.00 46.41      H31H O
ATOM   5651  N   VAL  173      10.430  55.822  48.168  1.00 45.18      H31H N
ATOM   5652  CA  VAL  173      10.683  56.062  46.748  1.00 44.79      H31H C
ATOM   5653  CB  VAL  173      11.999  56.839  46.546  1.00 43.99      H31H C
ATOM   5654  CG1 VAL  173      12.280  56.991  45.068  1.00 43.36      H31H C
ATOM   5655  CG2 VAL  173      11.933  58.192  47.224  1.00 45.03      H31H C
ATOM   5656  C   VAL  173      10.837  54.765  45.970  1.00 46.04      H31H C
ATOM   5657  O   VAL  173      11.828  54.070  46.157  1.00 46.61      H31H O
ATOM   5658  N   HIS  174       9.899  54.424  45.091  1.00 48.02      H31H N
ATOM   5659  CA  HIS  174      10.165  53.321  44.167  1.00 50.67      H31H C
ATOM   5660  CB  HIS  174       9.160  52.188  44.299  1.00 56.04      H31H C
ATOM   5661  CG  HIS  174       9.618  50.887  43.707  1.00 64.33      H31H C
ATOM   5662  CD2 HIS  174      10.847  50.313  43.652  1.00 67.01      H31H C
ATOM   5663  ND1 HIS  174       8.755  49.998  43.097  1.00 67.38      H31H N
ATOM   5664  CE1 HIS  174       9.432  48.933  42.694  1.00 67.43      H31H C
ATOM   5665  NE2 HIS  174      10.702  49.099  43.019  1.00 68.12      H31H N
ATOM   5666  C   HIS  174      10.243  53.765  42.715  1.00 48.63      H31H C
ATOM   5667  O   HIS  174       9.222  54.087  42.110  1.00 48.68      H31H O
ATOM   5668  N   THR  175      11.450  53.771  42.167  1.00 46.86      H31H N
ATOM   5669  CA  THR  175      11.689  54.015  40.747  1.00 45.90      H31H C
ATOM   5670  CB  THR  175      13.096  54.614  40.559  1.00 45.07      H31H C
ATOM   5671  OG1 THR  175      13.205  55.827  41.317  1.00 42.71      H31H O
ATOM   5672  CG2 THR  175      13.383  54.873  39.077  1.00 44.22      H31H C
ATOM   5673  C   THR  175      11.608  52.702  39.946  1.00 45.53      H31H C
ATOM   5674  O   THR  175      12.380  51.769  40.193  1.00 44.10      H31H O
ATOM   5675  N   PHE  176      10.686  52.630  38.991  1.00 45.00      H31H N
ATOM   5676  CA  PHE  176      10.489  51.395  38.236  1.00 47.18      H31H C
ATOM   5677  CB  PHE  176       9.036  51.253  37.827  1.00 46.51      H31H C
ATOM   5678  CG  PHE  176       8.102  51.087  38.986  1.00 48.17      H31H C
ATOM   5679  CD1 PHE  176       7.654  52.191  39.694  1.00 45.97      H31H C
ATOM   5680  CD2 PHE  176       7.647  49.827  39.353  1.00 47.11      H31H C
ATOM   5681  CE1 PHE  176       6.771  52.039  40.738  1.00 45.93      H31H C
ATOM   5682  CE2 PHE  176       6.762  49.675  40.400  1.00 46.13      H31H C
ATOM   5683  CZ  PHE  176       6.325  50.781  41.090  1.00 46.30      H31H C
ATOM   5684  C   PHE  176      11.364  51.265  36.999  1.00 48.64      H31H C
ATOM   5685  O   PHE  176      11.760  52.261  36.394  1.00 48.01      H31H O
ATOM   5686  N   PRO  177      11.688  50.013  36.613  1.00 48.62      H31H N
ATOM   5687  CD  PRO  177      11.330  48.782  37.338  1.00 50.33      H31H C
ATOM   5688  CA  PRO  177      12.573  49.722  35.474  1.00 49.81      H31H C
ATOM   5689  CB  PRO  177      12.832  48.216  35.590  1.00 49.25      H31H C
ATOM   5690  CG  PRO  177      12.471  47.872  37.005  1.00 49.07      H31H C
ATOM   5691  C   PRO  177      11.898  50.093  34.157  1.00 48.15      H31H C
ATOM   5692  O   PRO  177      10.692  49.864  33.977  1.00 44.95      H31H O
ATOM   5693  N   ALA  178      12.681  50.671  33.250  1.00 47.67      H31H N
ATOM   5694  CA  ALA  178      12.133  51.303  32.055  1.00 49.62      H31H C
ATOM   5695  CB  ALA  178      13.218  52.103  31.354  1.00 50.13      H31H C
ATOM   5696  C   ALA  178      11.530  50.298  31.087  1.00 50.13      H31H C
ATOM   5697  O   ALA  178      11.955  49.153  31.023  1.00 50.26      H31H O
ATOM   5698  N   VAL  179      10.538  50.728  30.325  1.00 50.60      H31H N
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ATOM   5699  CA  VAL   179      10.123  49.941  29.181  1.00 52.63      H31H C
ATOM   5700  CB  VAL   179       8.585  49.921  29.017  1.00 49.49      H31H C
ATOM   5701  CG1 VAL   179       8.075  51.264  28.523  1.00 47.79      H31H C
ATOM   5702  CG2 VAL   179       8.204  48.828  28.057  1.00 48.76      H31H C
ATOM   5703  C   VAL   179      10.760  50.485  27.890  1.00 57.03      H31H C
ATOM   5704  O   VAL   179      11.205  51.639  27.820  1.00 55.93      H31H O
ATOM   5705  N   LEU   180      10.824  49.625  26.880  1.00 61.40      H31H N
ATOM   5706  CA  LEU   180      11.157  50.033  25.528  1.00 63.99      H31H C
ATOM   5707  CB  LEU   180      12.042  48.976  24.878  1.00 61.84      H31H C
ATOM   5708  CG  LEU   180      13.014  49.469  23.813  1.00 60.90      H31H C
ATOM   5709  CD1 LEU   180      13.527  50.842  24.190  1.00 58.46      H31H C
ATOM   5710  CD2 LEU   180      14.172  48.482  23.684  1.00 60.70      H31H C
ATOM   5711  C   LEU   180       9.829  50.137  24.788  1.00 67.05      H31H C
ATOM   5712  O   LEU   180       9.032  49.200  24.813  1.00 68.83      H31H O
ATOM   5713  N   GLN   181       9.576  51.281  24.158  1.00 69.86      H31H N
ATOM   5714  CA  GLN   181       8.376  51.441  23.341  1.00 71.70      H31H C
ATOM   5715  CB  GLN   181       7.882  52.898  23.384  1.00 74.68      H31H C
ATOM   5716  CG  GLN   181       7.427  53.378  24.772  1.00 79.74      H31H C
ATOM   5717  CD  GLN   181       7.670  54.877  25.014  1.00 83.20      H31H C
ATOM   5718  OE1 GLN   181       6.731  55.643  25.273  1.00 85.51      H31H O
ATOM   5719  NE2 GLN   181       8.933  55.293  24.938  1.00 84.18      H31H N
ATOM   5720  C   GLN   181       8.741  51.039  21.912  1.00 72.09      H31H C
ATOM   5721  O   GLN   181       9.930  50.913  21.574  1.00 70.09      H31H O
ATOM   5722  N   SER   182       7.723  50.830  21.080  1.00 72.37      H31H N
ATOM   5723  CA  SER   182       7.937  50.423  19.694  1.00 72.49      H31H C
ATOM   5724  CB  SER   182       6.599  50.335  18.967  1.00 71.69      H31H C
ATOM   5725  OG  SER   182       5.733  49.438  19.639  1.00 72.82      H31H O
ATOM   5726  C   SER   182       8.853  51.407  18.976  1.00 72.06      H31H C
ATOM   5727  O   SER   182       9.297  51.164  17.855  1.00 73.81      H31H O
ATOM   5728  N   SER   183       9.131  52.519  19.640  1.00 69.30      H31H N
ATOM   5729  CA  SER   183       9.965  53.554  19.076  1.00 67.21      H31H C
ATOM   5730  CB  SER   183       9.624  54.890  19.735  1.00 69.40      H31H C
ATOM   5731  OG  SER   183      10.483  55.921  19.280  1.00 71.75      H31H O
ATOM   5732  C   SER   183      11.445  53.239  19.269  1.00 65.60      H31H C
ATOM   5733  O   SER   183      12.305  53.765  18.552  1.00 64.41      H31H O
ATOM   5734  N   GLY   184      11.747  52.386  20.245  1.00 64.00      H31H N
ATOM   5735  CA  GLY   184      13.137  52.179  20.626  1.00 62.60      H31H C
ATOM   5736  C   GLY   184      13.634  53.272  21.562  1.00 61.18      H31H C
ATOM   5737  O   GLY   184      14.828  53.376  21.850  1.00 60.60      H31H O
ATOM   5738  N   LEU   185      12.703  54.097  22.030  1.00 59.98      H31H N
ATOM   5739  CA  LEU   185      12.987  55.063  23.079  1.00 59.22      H31H C
ATOM   5740  CB  LEU   185      12.357  56.422  22.744  1.00 59.91      H31H C
ATOM   5741  CG  LEU   185      12.913  57.128  21.502  1.00 59.10      H31H C
ATOM   5742  CD1 LEU   185      12.278  58.502  21.353  1.00 57.09      H31H C
ATOM   5743  CD2 LEU   185      14.431  57.239  21.621  1.00 57.14      H31H C
ATOM   5744  C   LEU   185      12.398  54.522  24.372  1.00 58.54      H31H C
ATOM   5745  O   LEU   185      11.390  53.805  24.343  1.00 58.03      H31H O
ATOM   5746  N   TYR   186      13.030  54.861  25.495  1.00 56.77      H31H N
ATOM   5747  CA  TYR   186      12.629  54.334  26.792  1.00 54.56      H31H C
ATOM   5748  CB  TYR   186      13.853  54.115  27.678  1.00 53.50      H31H C
ATOM   5749  CG  TYR   186      14.782  53.017  27.218  1.00 53.54      H31H C
ATOM   5750  CD1 TYR   186      14.423  51.672  27.341  1.00 53.96      H31H C
ATOM   5751  CE1 TYR   186      15.281  50.655  26.928  1.00 53.54      H31H C
ATOM   5752  CD2 TYR   186      16.023  53.319  26.673  1.00 52.95      H31H C
ATOM   5753  CE2 TYR   186      16.885  52.317  26.259  1.00 55.74      H31H C
ATOM   5754  CZ  TYR   186      16.509  50.987  26.386  1.00 54.83      H31H C
ATOM   5755  OH  TYR   186      17.366  50.006  25.944  1.00 54.90      H31H O
ATOM   5756  C   TYR   186      11.663  55.249  27.526  1.00 53.17      H31H C
ATOM   5757  O   TYR   186      11.697  56.463  27.374  1.00 55.58      H31H O
ATOM   5758  N   SER   187      10.798  54.656  28.331  1.00 50.88      H31H N
ATOM   5759  CA  SER   187      10.108  55.397  29.376  1.00 49.39      H31H C
ATOM   5760  CB  SER   187       8.619  55.528  29.057  1.00 48.12      H31H C
ATOM   5761  OG  SER   187       8.416  56.389  27.956  1.00 49.32      H31H O
ATOM   5762  C   SER   187      10.278  54.665  30.699  1.00 48.74      H31H C
ATOM   5763  O   SER   187      10.304  53.432  30.741  1.00 49.37      H31H O
ATOM   5764  N   LEU   188      10.408  55.421  31.779  1.00 46.61      H31H N
ATOM   5765  CA  LEU   188      10.223  54.852  33.099  1.00 45.05      H31H C
ATOM   5766  CB  LEU   188      11.572  54.451  33.708  1.00 41.87      H31H C
ATOM   5767  CG  LEU   188      12.446  55.532  34.348  1.00 41.43      H31H C
ATOM   5768  CD1 LEU   188      11.988  55.808  35.773  1.00 41.51      H31H C
ATOM   5769  CD2 LEU   188      13.876  55.058  34.363  1.00 40.21      H31H C
ATOM   5770  C   LEU   188       9.549  55.917  33.943  1.00 44.78      H31H C
ATOM   5771  O   LEU   188       9.594  57.093  33.599  1.00 44.93      H31H O
ATOM   5772  N   SER   189       8.920  55.509  35.038  1.00 44.18      H31H N
ATOM   5773  CA  SER   189       8.382  56.461  35.993  1.00 43.74      H31H C
ATOM   5774  CB  SER   189       6.857  56.474  35.905  1.00 42.81      H31H C
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ATOM   5775  OG  SER   189       6.384  55.250  35.391  1.00 40.88      H31H O
ATOM   5776  C   SER   189       8.835  56.138  37.416  1.00 44.62      H31H C
ATOM   5777  O   SER   189       9.054  54.978  37.758  1.00 44.06      H31H O
ATOM   5778  N   SER   190       8.986  57.175  38.235  1.00 45.13      H31H N
ATOM   5779  CA  SER   190       9.349  57.004  39.637  1.00 47.28      H31H C
ATOM   5780  CB  SER   190      10.651  57.759  39.910  1.00 47.70      H31H C
ATOM   5781  OG  SER   190      10.959  57.794  41.291  1.00 51.62      H31H O
ATOM   5782  C   SER   190       8.228  57.514  40.559  1.00 48.63      H31H C
ATOM   5783  O   SER   190       7.793  58.654  40.449  1.00 49.40      H31H O
ATOM   5784  N   VAL   191       7.749  56.669  41.460  1.00 48.61      H31H N
ATOM   5785  CA  VAL   191       6.716  57.091  42.404  1.00 48.08      H31H C
ATOM   5786  CB  VAL   191       5.628  56.022  42.579  1.00 48.63      H31H C
ATOM   5787  CG1 VAL   191       4.361  56.655  43.120  1.00 49.16      H31H C
ATOM   5788  CG2 VAL   191       5.378  55.322  41.276  1.00 50.66      H31H C
ATOM   5789  C   VAL   191       7.344  57.302  43.770  1.00 48.14      H31H C
ATOM   5790  O   VAL   191       8.455  56.846  44.032  1.00 48.84      H31H O
ATOM   5791  N   VAL   192       6.625  57.980  44.649  1.00 47.74      H31H N
ATOM   5792  CA  VAL   192       7.071  58.124  46.020  1.00 49.09      H31H C
ATOM   5793  CB  VAL   192       7.993  59.347  46.168  1.00 47.37      H31H C
ATOM   5794  CG1 VAL   192       7.188  60.616  46.033  1.00 49.41      H31H C
ATOM   5795  CG2 VAL   192       8.716  59.299  47.494  1.00 48.43      H31H C
ATOM   5796  C   VAL   192       5.833  58.285  46.893  1.00 51.29      H31H C
ATOM   5797  O   VAL   192       4.869  58.925  46.491  1.00 52.06      H31H O
ATOM   5798  N   THR   193       5.845  57.683  48.076  1.00 53.99      H31H N
ATOM   5799  CA  THR   193       4.707  57.791  48.978  1.00 55.72      H31H C
ATOM   5800  CB  THR   193       4.442  56.475  49.712  1.00 54.82      H31H C
ATOM   5801  OG1 THR   193       4.299  55.412  48.762  1.00 52.78      H31H O
ATOM   5802  CG2 THR   193       3.174  56.593  50.529  1.00 53.19      H31H C
ATOM   5803  C   THR   193       4.949  58.865  50.024  1.00 57.60      H31H C
ATOM   5804  O   THR   193       6.014  58.915  50.638  1.00 60.19      H31H O
ATOM   5805  N   VAL   194       3.951  59.718  50.223  1.00 59.12      H31H N
ATOM   5806  CA  VAL   194       4.007  60.758  51.244  1.00 59.31      H31H C
ATOM   5807  CB  VAL   194       4.182  62.149  50.614  1.00 57.34      H31H C
ATOM   5808  CG1 VAL   194       5.482  62.222  49.839  1.00 56.49      H31H C
ATOM   5809  CG2 VAL   194       3.001  62.443  49.714  1.00 53.69      H31H C
ATOM   5810  C   VAL   194       2.705  60.776  52.041  1.00 61.25      H31H C
ATOM   5811  O   VAL   194       1.692  60.235  51.604  1.00 61.92      H31H O
ATOM   5812  N   PRO   195       2.721  61.416  53.222  1.00 62.39      H31H N
ATOM   5813  CD  PRO   195       3.960  61.784  53.933  1.00 60.34      H31H C
ATOM   5814  CA  PRO   195       1.521  61.810  53.980  1.00 63.17      H31H C
ATOM   5815  CB  PRO   195       2.088  62.549  55.197  1.00 59.42      H31H C
ATOM   5816  CG  PRO   195       3.473  62.076  55.326  1.00 58.40      H31H C
ATOM   5817  C   PRO   195       0.575  62.718  53.182  1.00 64.78      H31H C
ATOM   5818  O   PRO   195       1.016  63.634  52.486  1.00 65.00      H31H O
ATOM   5819  N   SER   196      -0.725  62.476  53.297  1.00 65.72      H31H N
ATOM   5820  CA  SER   196      -1.705  63.386  52.714  1.00 66.61      H31H C
ATOM   5821  CB  SER   196      -3.117  62.826  52.895  1.00 65.85      H31H C
ATOM   5822  OG  SER   196      -3.210  61.527  52.356  1.00 61.78      H31H O
ATOM   5823  C   SER   196      -1.607  64.754  53.393  1.00 67.65      H31H C
ATOM   5824  O   SER   196      -1.704  65.795  52.746  1.00 68.06      H31H O
ATOM   5825  N   SER   197      -1.403  64.734  54.704  1.00 69.10      H31H N
ATOM   5826  CA  SER   197      -1.411  65.944  55.516  1.00 69.65      H31H C
ATOM   5827  CB  SER   197      -1.255  65.561  56.990  1.00 69.31      H31H C
ATOM   5828  OG  SER   197      -0.156  64.684  57.169  1.00 68.21      H31H O
ATOM   5829  C   SER   197      -0.329  66.959  55.130  1.00 70.07      H31H C
ATOM   5830  O   SER   197      -0.543  68.174  55.225  1.00 70.52      H31H O
ATOM   5831  N   SER   198       0.835  66.465  54.712  1.00 69.45      H31H N
ATOM   5832  CA  SER   198       1.959  67.342  54.405  1.00 68.99      H31H C
ATOM   5833  CB  SER   198       3.295  66.596  54.588  1.00 70.62      H31H C
ATOM   5834  OG  SER   198       3.418  65.476  53.730  1.00 73.17      H31H O
ATOM   5835  C   SER   198       1.855  67.925  52.996  1.00 68.67      H31H C
ATOM   5836  O   SER   198       2.640  68.798  52.610  1.00 66.31      H31H O
ATOM   5837  N   LEU   199       0.875  67.450  52.233  1.00 69.05      H31H N
ATOM   5838  CA  LEU   199       0.587  68.038  50.938  1.00 70.87      H31H C
ATOM   5839  CB  LEU   199      -0.469  67.219  50.205  1.00 68.03      H31H C
ATOM   5840  CG  LEU   199      -0.068  65.802  49.805  1.00 66.69      H31H C
ATOM   5841  CD1 LEU   199      -1.221  65.107  49.119  1.00 65.13      H31H C
ATOM   5842  CD2 LEU   199       1.127  65.864  48.884  1.00 66.64      H31H C
ATOM   5843  C   LEU   199       0.077  69.448  51.171  1.00 74.01      H31H C
ATOM   5844  O   LEU   199      -0.534  69.736  52.202  1.00 75.02      H31H O
ATOM   5845  N   GLY   200       0.331  70.335  50.224  1.00 76.20      H31H N
ATOM   5846  CA  GLY   200      -0.025  71.720  50.442  1.00 79.03      H31H C
ATOM   5847  C   GLY   200       1.110  72.503  51.069  1.00 81.18      H31H C
ATOM   5848  O   GLY   200       1.260  73.693  50.805  1.00 82.30      H31H O
ATOM   5849  N   THR   201       1.922  71.847  51.889  1.00 82.71      H31H N
ATOM   5850  CA  THR   201       2.984  72.552  52.591  1.00 83.47      H31H C
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ATOM   5851  CB  THR   201       2.869  72.334  54.093  1.00 82.12      H31H C
ATOM   5852  OG1 THR   201       2.774  70.932  54.358  1.00 82.06      H31H O
ATOM   5853  CG2 THR   201       1.643  73.031  54.632  1.00 80.62      H31H C
ATOM   5854  C   THR   201       4.377  72.143  52.142  1.00 84.13      H31H C
ATOM   5855  O   THR   201       5.191  72.988  51.766  1.00 86.17      H31H O
ATOM   5856  N   GLN   202       4.653  70.846  52.189  1.00 83.67      H31H N
ATOM   5857  CA  GLN   202       5.949  70.339  51.760  1.00 81.24      H31H C
ATOM   5858  CB  GLN   202       6.141  68.902  52.250  1.00 84.61      H31H C
ATOM   5859  CG  GLN   202       6.355  68.812  53.740  1.00 89.81      H31H C
ATOM   5860  CD  GLN   202       7.301  69.881  54.232  1.00 93.21      H31H C
ATOM   5861  OE1 GLN   202       6.995  70.603  55.178  1.00 95.82      H31H O
ATOM   5862  NE2 GLN   202       8.458  69.997  53.586  1.00 96.62      H31H N
ATOM   5863  C   GLN   202       6.013  70.393  50.246  1.00 76.65      H31H C
ATOM   5864  O   GLN   202       4.997  70.253  49.575  1.00 75.95      H31H O
ATOM   5865  N   THR   203       7.199  70.607  49.699  1.00 72.04      H31H N
ATOM   5866  CA  THR   203       7.326  70.634  48.253  1.00 67.97      H31H C
ATOM   5867  CB  THR   203       7.958  71.961  47.791  1.00 68.16      H31H C
ATOM   5868  OG1 THR   203       9.304  72.037  48.263  1.00 69.55      H31H O
ATOM   5869  CG2 THR   203       7.170  73.149  48.352  1.00 66.96      H31H C
ATOM   5870  C   THR   203       8.145  69.439  47.737  1.00 64.26      H31H C
ATOM   5871  O   THR   203       9.212  69.119  48.262  1.00 63.14      H31H O
ATOM   5872  N   TYR   204       7.623  68.769  46.713  1.00 60.03      H31H N
ATOM   5873  CA  TYR   204       8.261  67.567  46.183  1.00 56.87      H31H C
ATOM   5874  CB  TYR   204       7.262  66.406  46.201  1.00 56.87      H31H C
ATOM   5875  CG  TYR   204       6.884  66.022  47.605  1.00 58.37      H31H C
ATOM   5876  CD1 TYR   204       5.636  66.324  48.121  1.00 59.73      H31H C
ATOM   5877  CE1 TYR   204       5.318  66.035  49.436  1.00 60.84      H31H C
ATOM   5878  CD2 TYR   204       7.805  65.414  48.440  1.00 60.23      H31H C
ATOM   5879  CE2 TYR   204       7.500  65.119  49.749  1.00 60.88      H31H C
ATOM   5880  CZ  TYR   204       6.257  65.432  50.245  1.00 61.19      H31H C
ATOM   5881  OH  TYR   204       5.964  65.141  51.558  1.00 60.98      H31H O
ATOM   5882  C   TYR   204       8.844  67.751  44.774  1.00 54.04      H31H C
ATOM   5883  O   TYR   204       8.117  67.841  43.784  1.00 52.97      H31H O
ATOM   5884  N   ILE   205      10.167  67.806  44.693  1.00 50.55      H31H N
ATOM   5885  CA  ILE   205      10.831  67.982  43.415  1.00 48.02      H31H C
ATOM   5886  CB  ILE   205      11.880  69.126  43.485  1.00 43.92      H31H C
ATOM   5887  CG2 ILE   205      12.670  69.191  42.178  1.00 39.57      H31H C
ATOM   5888  CG1 ILE   205      11.171  70.449  43.801  1.00 40.24      H31H C
ATOM   5889  CD1 ILE   205      12.090  71.615  44.002  1.00 36.63      H31H C
ATOM   5890  C   ILE   205      11.511  66.685  43.017  1.00 48.10      H31H C
ATOM   5891  O   ILE   205      12.252  66.096  43.801  1.00 49.28      H31H O
ATOM   5892  N   CYS   206      11.256  66.234  41.801  1.00 47.41      H31H N
ATOM   5893  CA  CYS   206      11.926  65.042  41.313  1.00 47.33      H31H C
ATOM   5894  C   CYS   206      13.021  65.467  40.346  1.00 46.16      H31H C
ATOM   5895  O   CYS   206      12.751  66.190  39.395  1.00 46.16      H31H O
ATOM   5896  CB  CYS   206      10.905  64.116  40.637  1.00 48.70      H31H C
ATOM   5897  SG  CYS   206      10.705  64.310  38.841  1.00 49.42      H31H S
ATOM   5898  N   ASN   207      14.259  65.046  40.608  1.00 46.09      H31H N
ATOM   5899  CA  ASN   207      15.400  65.405  39.758  1.00 45.79      H31H C
ATOM   5900  CB  ASN   207      16.598  65.771  40.618  1.00 47.44      H31H C
ATOM   5901  CG  ASN   207      16.194  66.274  41.978  1.00 49.48      H31H C
ATOM   5902  OD1 ASN   207      15.873  65.481  42.884  1.00 49.54      H31H O
ATOM   5903  ND2 ASN   207      16.199  67.597  42.140  1.00 48.35      H31H N
ATOM   5904  C   ASN   207      15.794  64.256  38.841  1.00 45.19      H31H C
ATOM   5905  O   ASN   207      15.981  63.126  39.292  1.00 45.98      H31H O
ATOM   5906  N   VAL   208      15.920  64.545  37.552  1.00 44.59      H31H N
ATOM   5907  CA  VAL   208      16.179  63.499  36.579  1.00 44.44      H31H C
ATOM   5908  CB  VAL   208      15.061  63.395  35.565  1.00 42.93      H31H C
ATOM   5909  CG1 VAL   208      15.378  62.291  34.596  1.00 42.05      H31H C
ATOM   5910  CG2 VAL   208      13.733  63.145  36.275  1.00 42.37      H31H C
ATOM   5911  C   VAL   208      17.477  63.703  35.824  1.00 46.24      H31H C
ATOM   5912  O   VAL   208      17.649  64.669  35.076  1.00 46.31      H31H O
ATOM   5913  N   ASN   209      18.398  62.774  36.027  1.00 49.74      H31H N
ATOM   5914  CA  ASN   209      19.695  62.842  35.389  1.00 51.57      H31H C
ATOM   5915  CB  ASN   209      20.775  62.534  36.408  1.00 53.49      H31H C
ATOM   5916  CG  ASN   209      21.866  63.555  36.397  1.00 58.10      H31H C
ATOM   5917  OD1 ASN   209      22.623  63.663  35.425  1.00 61.93      H31H O
ATOM   5918  ND2 ASN   209      21.961  64.331  37.476  1.00 60.31      H31H N
ATOM   5919  C   ASN   209      19.758  61.854  34.233  1.00 53.30      H31H C
ATOM   5920  O   ASN   209      19.119  60.794  34.266  1.00 52.74      H31H O
ATOM   5921  N   HIS   210      20.521  62.213  33.208  1.00 55.96      H31H N
ATOM   5922  CA  HIS   210      20.682  61.371  32.027  1.00 60.39      H31H C
ATOM   5923  CB  HIS   210      19.506  61.586  31.075  1.00 57.95      H31H C
ATOM   5924  CG  HIS   210      19.554  60.737  29.841  1.00 57.46      H31H C
ATOM   5925  CD2 HIS   210      18.962  59.554  29.550  1.00 56.95      H31H C
ATOM   5926  ND1 HIS   210      20.240  61.111  28.705  1.00 56.62      H31H N
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ATOM   5927  CE1 HIS   210      20.064  60.196  27.768  1.00 56.04      H31H C
ATOM   5928  NE2 HIS   210      19.293  59.241  28.255  1.00 56.13      H31H N
ATOM   5929  C   HIS   210      21.989  61.732  31.332  1.00 64.05      H31H C
ATOM   5930  O   HIS   210      21.999  62.563  30.423  1.00 66.26      H31H O
ATOM   5931  N   LYS   211      23.088  61.105  31.759  1.00 67.42      H31H N
ATOM   5932  CA  LYS   211      24.422  61.500  31.308  1.00 71.12      H31H C
ATOM   5933  CB  LYS   211      25.493  60.800  32.148  1.00 73.47      H31H C
ATOM   5934  CG  LYS   211      25.650  61.391  33.537  1.00 78.05      H31H C
ATOM   5935  CD  LYS   211      26.676  60.635  34.371  1.00 83.09      H31H C
ATOM   5936  CE  LYS   211      26.518  60.963  35.851  1.00 87.25      H31H C
ATOM   5937  NZ  LYS   211      25.078  60.948  36.299  1.00 92.55      H31H N
ATOM   5938  C   LYS   211      24.717  61.288  29.821  1.00 71.44      H31H C
ATOM   5939  O   LYS   211      25.602  61.954  29.265  1.00 70.88      H31H O
ATOM   5940  N   PRO   212      23.988  60.363  29.161  1.00 70.66      H31H N
ATOM   5941  CD  PRO   212      23.075  59.370  29.752  1.00 68.67      H31H C
ATOM   5942  CA  PRO   212      24.151  60.136  27.723  1.00 70.83      H31H C
ATOM   5943  CB  PRO   212      23.177  59.003  27.425  1.00 68.87      H31H C
ATOM   5944  CG  PRO   212      23.014  58.308  28.708  1.00 67.41      H31H C
ATOM   5945  C   PRO   212      23.856  61.369  26.881  1.00 71.39      H31H C
ATOM   5946  O   PRO   212      24.549  61.634  25.897  1.00 73.26      H31H O
ATOM   5947  N   SER   213      22.831  62.124  27.268  1.00 69.78      H31H N
ATOM   5948  CA  SER   213      22.506  63.363  26.573  1.00 67.25      H31H C
ATOM   5949  CB  SER   213      20.998  63.448  26.316  1.00 68.21      H31H C
ATOM   5950  OG  SER   213      20.260  63.475  27.523  1.00 69.83      H31H O
ATOM   5951  C   SER   213      22.976  64.603  27.333  1.00 64.95      H31H C
ATOM   5952  O   SER   213      22.739  65.733  26.895  1.00 64.30      H31H O
ATOM   5953  N   ASN   214      23.651  64.380  28.458  1.00 63.80      H31H N
ATOM   5954  CA  ASN   214      23.966  65.445  29.405  1.00 62.78      H31H C
ATOM   5955  CB  ASN   214      25.135  66.282  28.898  1.00 64.09      H31H C
ATOM   5956  CG  ASN   214      26.432  65.517  28.892  1.00 66.41      H31H C
ATOM   5957  OD1 ASN   214      26.678  64.705  28.003  1.00 68.49      H31H O
ATOM   5958  ND2 ASN   214      27.276  65.771  29.886  1.00 68.13      H31H N
ATOM   5959  C   ASN   214      22.752  66.338  29.632  1.00 60.83      H31H C
ATOM   5960  O   ASN   214      22.733  67.513  29.258  1.00 59.26      H31H O
ATOM   5961  N   THR   215      21.732  65.767  30.257  1.00 58.26      H31H N
ATOM   5962  CA  THR   215      20.473  66.471  30.437  1.00 55.62      H31H C
ATOM   5963  CB  THR   215      19.412  65.954  29.444  1.00 52.20      H31H C
ATOM   5964  OG1 THR   215      19.909  66.088  28.112  1.00 47.16      H31H O
ATOM   5965  CG2 THR   215      18.137  66.741  29.585  1.00 49.57      H31H C
ATOM   5966  C   THR   215      19.937  66.288  31.855  1.00 54.60      H31H C
ATOM   5967  O   THR   215      19.390  65.228  32.174  1.00 57.16      H31H O
ATOM   5968  N   LYS   216      20.069  67.298  32.710  1.00 51.53      H31H N
ATOM   5969  CA  LYS   216      19.366  67.224  33.976  1.00 49.41      H31H C
ATOM   5970  CB  LYS   216      20.251  67.745  35.109  1.00 47.56      H31H C
ATOM   5971  CG  LYS   216      19.748  67.346  36.505  1.00 51.25      H31H C
ATOM   5972  CD  LYS   216      20.613  67.953  37.610  1.00 56.11      H31H C
ATOM   5973  CE  LYS   216      19.792  68.452  38.809  1.00 59.03      H31H C
ATOM   5974  NZ  LYS   216      19.922  69.939  39.020  1.00 66.78      H31H N
ATOM   5975  C   LYS   216      18.032  67.994  33.913  1.00 47.52      H31H C
ATOM   5976  O   LYS   216      17.947  69.086  33.343  1.00 47.37      H31H O
ATOM   5977  N   VAL   217      16.986  67.399  34.475  1.00 45.17      H31H N
ATOM   5978  CA  VAL   217      15.675  68.029  34.495  1.00 45.48      H31H C
ATOM   5979  CB  VAL   217      14.705  67.359  33.504  1.00 42.90      H31H C
ATOM   5980  CG1 VAL   217      13.382  68.094  33.502  1.00 43.01      H31H C
ATOM   5981  CG2 VAL   217      15.291  67.349  32.126  1.00 42.04      H31H C
ATOM   5982  C   VAL   217      15.033  67.935  35.874  1.00 45.53      H31H C
ATOM   5983  O   VAL   217      14.689  66.837  36.325  1.00 44.46      H31H O
ATOM   5984  N   ASP   218      14.844  69.085  36.522  1.00 45.13      H31H N
ATOM   5985  CA  ASP   218      14.068  69.157  37.765  1.00 44.22      H31H C
ATOM   5986  CB  ASP   218      14.646  70.233  38.697  1.00 45.01      H31H C
ATOM   5987  CG  ASP   218      16.039  69.896  39.177  1.00 46.69      H31H C
ATOM   5988  OD1 ASP   218      16.230  68.770  39.690  1.00 48.07      H31H O
ATOM   5989  OD2 ASP   218      16.942  70.752  39.040  1.00 47.24      H31H O
ATOM   5990  C   ASP   218      12.609  69.485  37.468  1.00 41.82      H31H C
ATOM   5991  O   ASP   218      12.325  70.438  36.759  1.00 41.41      H31H O
ATOM   5992  N   LYS   219      11.688  68.690  38.000  1.00 41.58      H31H N
ATOM   5993  CA  LYS   219      10.261  69.006  37.898  1.00 43.18      H31H C
ATOM   5994  CB  LYS   219       9.546  68.005  36.983  1.00 42.71      H31H C
ATOM   5995  CG  LYS   219       8.209  68.494  36.454  1.00 42.46      H31H C
ATOM   5996  CD  LYS   219       8.401  69.794  35.667  1.00 46.00      H31H C
ATOM   5997  CE  LYS   219       7.112  70.264  34.971  1.00 46.61      H31H C
ATOM   5998  NZ  LYS   219       7.108  69.969  33.497  1.00 46.37      H31H N
ATOM   5999  C   LYS   219       9.612  68.982  39.279  1.00 44.12      H31H C
ATOM   6000  O   LYS   219       9.785  68.027  40.035  1.00 43.32      H31H O
ATOM   6001  N   LYS   220       8.879  70.037  39.616  1.00 46.31      H31H N
ATOM   6002  CA  LYS   220       8.173  70.073  40.887  1.00 51.36      H31H C
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ATOM   6003  CB  LYS   220       8.004  71.512  41.356  1.00 51.83      H31H C
ATOM   6004  CG  LYS   220       7.136  71.684  42.587  1.00 53.30      H31H C
ATOM   6005  CD  LYS   220       7.297  73.099  43.134  1.00 55.65      H31H C
ATOM   6006  CE  LYS   220       6.414  73.349  44.343  1.00 58.38      H31H C
ATOM   6007  NZ  LYS   220       6.792  74.612  45.042  1.00 61.74      H31H N
ATOM   6008  C   LYS   220       6.810  69.413  40.731  1.00 55.52      H31H C
ATOM   6009  O   LYS   220       6.127  69.592  39.727  1.00 56.17      H31H O
ATOM   6010  N   VAL   221       6.425  68.622  41.717  1.00 59.39      H31H N
ATOM   6011  CA  VAL   221       5.191  67.885  41.620  1.00 63.66      H31H C
ATOM   6012  CB  VAL   221       5.439  66.388  41.823  1.00 64.08      H31H C
ATOM   6013  CG1 VAL   221       4.117  65.618  41.778  1.00 62.71      H31H C
ATOM   6014  CG2 VAL   221       6.391  65.892  40.743  1.00 61.51      H31H C
ATOM   6015  C   VAL   221       4.234  68.410  42.661  1.00 67.49      H31H C
ATOM   6016  O   VAL   221       4.508  68.365  43.852  1.00 67.95      H31H O
ATOM   6017  N   GLU   222       3.118  68.942  42.182  1.00 72.52      H31H N
ATOM   6018  CA  GLU   222       2.115  69.555  43.032  1.00 77.04      H31H C
ATOM   6019  CB  GLU   222       1.951  71.041  42.689  1.00 79.41      H31H C
ATOM   6020  CG  GLU   222       3.071  71.952  43.174  1.00 87.22      H31H C
ATOM   6021  CD  GLU   222       2.790  73.425  42.900  1.00 91.43      H31H C
ATOM   6022  OE1 GLU   222       3.133  74.277  43.750  1.00 95.60      H31H O
ATOM   6023  OE2 GLU   222       2.224  73.730  41.829  1.00 95.74      H31H O
ATOM   6024  C   GLU   222       0.793  68.843  42.802  1.00 79.22      H31H C
ATOM   6025  O   GLU   222       0.551  68.271  41.737  1.00 78.25      H31H O
ATOM   6026  N   PRO   223      -0.084  68.879  43.803  1.00 81.42      H31H N
ATOM   6027  CD  PRO   223       0.214  69.436  45.129  1.00 83.23      H31H C
ATOM   6028  CA  PRO   223      -1.433  68.318  43.734  1.00 83.45      H31H C
ATOM   6029  CB  PRO   223      -2.002  68.595  45.121  1.00 84.78      H31H C
ATOM   6030  CG  PRO   223      -0.805  68.783  45.992  1.00 85.56      H31H C
ATOM   6031  C   PRO   223      -2.261  68.981  42.635  1.00 84.25      H31H C
ATOM   6032  O   PRO   223      -1.940  70.080  42.185  1.00 82.72      H31H O
ATOM   6033  N   LYS   224      -3.332  68.310  42.221  1.00 85.68      H31H N
ATOM   6034  CA  LYS   224      -4.128  68.730  41.069  1.00 87.49      H31H C
ATOM   6035  CB  LYS   224      -4.799  67.507  40.432  1.00 87.94      H31H C
ATOM   6036  CG  LYS   224      -4.095  66.198  40.749  1.00 88.70      H31H C
ATOM   6037  CD  LYS   224      -4.345  65.129  39.692  1.00 90.62      H31H C
ATOM   6038  CE  LYS   224      -3.572  63.850  40.031  1.00 92.06      H31H C
ATOM   6039  NZ  LYS   224      -3.487  62.866  38.912  1.00 92.45      H31H N
ATOM   6040  C   LYS   224      -5.191  69.764  41.441  1.00 88.37      H31H C
ATOM   6041  O   LYS   224      -6.384  69.388  41.475  1.00 89.80      H31H O
ATOM   6042  OXT LYS   224      -4.825  70.932  41.701  1.00 89.87      H31H O
TER    6043      LYS   224                                              H31H
ATOM   6044  CB  SER     2      63.863  32.817   2.917  1.00 52.61      L21B C
ATOM   6045  OG  SER     2      65.036  32.750   2.112  1.00 56.46      L21B O
ATOM   6046  C   SER     2      63.296  30.542   1.965  1.00 51.89      L21B C
ATOM   6047  O   SER     2      62.911  29.519   2.538  1.00 49.06      L21B O
ATOM   6048  N   SER     2      61.635  31.805   3.365  1.00 50.71      L21B N
ATOM   6049  CA  SER     2      62.738  31.922   2.361  1.00 50.89      L21B C
ATOM   6050  N   ALA     3      64.199  30.533   0.985  1.00 56.94      L21B N
ATOM   6051  CA  ALA     3      64.675  29.305   0.362  1.00 57.67      L21B C
ATOM   6052  CB  ALA     3      65.526  29.649  -0.843  1.00 55.59      L21B C
ATOM   6053  C   ALA     3      65.454  28.390   1.309  1.00 57.11      L21B C
ATOM   6054  O   ALA     3      65.912  28.813   2.368  1.00 57.76      L21B O
ATOM   6055  N   LEU     4      65.585  27.125   0.922  1.00 51.91      L21B N
ATOM   6056  CA  LEU     4      66.411  26.170   1.654  1.00 50.60      L21B C
ATOM   6057  CB  LEU     4      65.999  24.735   1.303  1.00 49.68      L21B C
ATOM   6058  CG  LEU     4      64.497  24.436   1.262  1.00 46.57      L21B C
ATOM   6059  CD1 LEU     4      64.270  22.951   1.014  1.00 42.80      L21B C
ATOM   6060  CD2 LEU     4      63.852  24.877   2.561  1.00 45.01      L21B C
ATOM   6061  C   LEU     4      67.880  26.383   1.282  1.00 51.12      L21B C
ATOM   6062  O   LEU     4      68.178  26.834   0.185  1.00 51.19      L21B O
ATOM   6063  N   THR     5      68.791  26.045   2.191  1.00 53.04      L21B N
ATOM   6064  CA  THR     5      70.207  26.318   1.999  1.00 54.33      L21B C
ATOM   6065  CB  THR     5      70.858  26.820   3.300  1.00 55.17      L21B C
ATOM   6066  OG1 THR     5      70.095  27.914   3.834  1.00 56.53      L21B O
ATOM   6067  CG2 THR     5      72.293  27.267   3.037  1.00 54.92      L21B C
ATOM   6068  C   THR     5      70.958  25.072   1.561  1.00 55.20      L21B C
ATOM   6069  O   THR     5      71.062  24.107   2.315  1.00 53.92      L21B O
ATOM   6070  N   GLN     6      71.480  25.103   0.339  1.00 57.74      L21B N
ATOM   6071  CA  GLN     6      72.399  24.075  -0.142  1.00 60.09      L21B C
ATOM   6072  CB  GLN     6      71.997  23.595  -1.536  1.00 58.92      L21B C
ATOM   6073  CG  GLN     6      70.627  22.949  -1.646  1.00 57.25      L21B C
ATOM   6074  CD  GLN     6      70.367  22.370  -3.041  1.00 56.07      L21B C
ATOM   6075  OE1 GLN     6      69.373  22.697  -3.688  1.00 55.90      L21B O
ATOM   6076  NE2 GLN     6      71.266  21.508  -3.503  1.00 51.99      L21B N
ATOM   6077  C   GLN     6      73.798  24.676  -0.220  1.00 62.99      L21B C
ATOM   6078  O   GLN     6      73.962  25.893  -0.141  1.00 63.57      L21B O
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ATOM   6079  N   PRO     7      74.827  23.831  -0.364  1.00 65.06      L21B N
ATOM   6080  CD  PRO     7      74.858  22.407   0.006  1.00 66.69      L21B C
ATOM   6081  CA  PRO     7      76.150  24.319  -0.761  1.00 66.63      L21B C
ATOM   6082  CB  PRO     7      77.052  23.094  -0.600  1.00 67.18      L21B C
ATOM   6083  CG  PRO     7      76.305  22.179   0.307  1.00 68.34      L21B C
ATOM   6084  C   PRO     7      76.081  24.772  -2.212  1.00 66.86      L21B C
ATOM   6085  O   PRO     7      75.360  24.179  -3.001  1.00 67.36      L21B O
ATOM   6086  N   ALA     8      76.824  25.810  -2.575  1.00 67.52      L21B N
ATOM   6087  CA  ALA     8      76.860  26.225  -3.971  1.00 68.06      L21B C
ATOM   6088  CB  ALA     8      77.642  27.526  -4.121  1.00 70.77      L21B C
ATOM   6089  C   ALA     8      77.482  25.126  -4.831  1.00 68.63      L21B C
ATOM   6090  O   ALA     8      77.122  24.967  -5.993  1.00 67.08      L21B O
ATOM   6091  N   SER     9      78.413  24.364  -4.265  1.00 69.56      L21B N
ATOM   6092  CA  SER     9      78.988  23.232  -4.990  1.00 71.04      L21B C
ATOM   6093  CB  SER     9      80.051  23.718  -5.979  1.00 71.64      L21B C
ATOM   6094  OG  SER     9      81.060  24.455  -5.315  1.00 75.30      L21B O
ATOM   6095  C   SER     9      79.587  22.153  -4.083  1.00 69.92      L21B C
ATOM   6096  O   SER     9      79.840  22.389  -2.902  1.00 69.61      L21B O
ATOM   6097  N   VAL    10      79.771  20.963  -4.651  1.00 68.05      L21B N
ATOM   6098  CA  VAL    10      80.487  19.867  -4.009  1.00 67.25      L21B C
ATOM   6099  CB  VAL    10      79.553  18.805  -3.398  1.00 68.41      L21B C
ATOM   6100  CG1 VAL    10      78.759  19.393  -2.259  1.00 69.36      L21B C
ATOM   6101  CG2 VAL    10      78.639  18.242  -4.478  1.00 68.43      L21B C
ATOM   6102  C   VAL    10      81.273  19.173  -5.093  1.00 66.54      L21B C
ATOM   6103  O   VAL    10      80.874  19.177  -6.262  1.00 67.05      L21B O
ATOM   6104  N   SER    11      82.383  18.560  -4.701  1.00 65.07      L21B N
ATOM   6105  CA  SER    11      83.317  18.029  -5.670  1.00 63.23      L21B C
ATOM   6106  CB  SER    11      84.572  18.897  -5.687  1.00 61.31      L21B C
ATOM   6107  OG  SER    11      85.258  18.728  -6.908  1.00 63.07      L21B O
ATOM   6108  C   SER    11      83.683  16.584  -5.352  1.00 62.63      L21B C
ATOM   6109  O   SER    11      83.759  16.194  -4.184  1.00 62.08      L21B O
ATOM   6110  N   GLY    12      83.911  15.794  -6.397  1.00 61.55      L21B N
ATOM   6111  CA  GLY    12      84.362  14.427  -6.206  1.00 59.55      L21B C
ATOM   6112  C   GLY    12      84.904  13.767  -7.463  1.00 59.44      L21B C
ATOM   6113  O   GLY    12      84.533  14.127  -8.585  1.00 57.13      L21B O
ATOM   6114  N   SER    13      85.784  12.787  -7.256  1.00 60.89      L21B N
ATOM   6115  CA  SER    13      86.412  12.017  -8.334  1.00 61.13      L21B C
ATOM   6116  CB  SER    13      87.909  11.855  -8.050  1.00 61.04      L21B C
ATOM   6117  OG  SER    13      88.515  11.011  -9.010  1.00 60.21      L21B O
ATOM   6118  C   SER    13      85.759  10.646  -8.439  1.00 59.77      L21B C
ATOM   6119  O   SER    13      85.223  10.145  -7.463  1.00 59.25      L21B O
ATOM   6120  N   PRO    14      85.809  10.014  -9.621  1.00 59.99      L21B N
ATOM   6121  CD  PRO    14      86.598  10.379 -10.812  1.00 59.07      L21B C
ATOM   6122  CA  PRO    14      85.040   8.776  -9.823  1.00 62.35      L21B C
ATOM   6123  CB  PRO    14      85.579   8.235 -11.148  1.00 60.39      L21B C
ATOM   6124  CG  PRO    14      86.045   9.474 -11.883  1.00 59.18      L21B C
ATOM   6125  C   PRO    14      85.163   7.759  -8.676  1.00 64.90      L21B C
ATOM   6126  O   PRO    14      86.192   7.676  -8.010  1.00 65.34      L21B O
ATOM   6127  N   GLY    15      84.098   7.001  -8.434  1.00 67.76      L21B N
ATOM   6128  CA  GLY    15      84.110   6.056  -7.336  1.00 71.41      L21B C
ATOM   6129  C   GLY    15      84.108   6.707  -5.963  1.00 75.03      L21B C
ATOM   6130  O   GLY    15      84.199   6.020  -4.950  1.00 74.91      L21B O
ATOM   6131  N   GLN    16      84.006   8.029  -5.908  1.00 79.31      L21B N
ATOM   6132  CA  GLN    16      83.878   8.712  -4.623  1.00 83.11      L21B C
ATOM   6133  CB  GLN    16      84.415  10.143  -4.714  1.00 85.50      L21B C
ATOM   6134  CG  GLN    16      85.929  10.243  -4.740  1.00 90.35      L21B C
ATOM   6135  CD  GLN    16      86.419  11.683  -4.710  1.00 92.66      L21B C
ATOM   6136  OE1 GLN    16      87.482  12.005  -5.256  1.00 92.77      L21B O
ATOM   6137  NE2 GLN    16      85.644  12.560  -4.069  1.00 93.76      L21B N
ATOM   6138  C   GLN    16      82.428   8.746  -4.145  1.00 83.84      L21B C
ATOM   6139  O   GLN    16      81.498   8.490  -4.919  1.00 84.75      L21B O
ATOM   6140  N   SER    17      82.246   9.056  -2.864  1.00 83.83      L21B N
ATOM   6141  CA  SER    17      80.913   9.239  -2.294  1.00 84.75      L21B C
ATOM   6142  CB  SER    17      80.603   8.125  -1.279  1.00 86.48      L21B C
ATOM   6143  OG  SER    17      80.314   6.891  -1.922  1.00 89.21      L21B O
ATOM   6144  C   SER    17      80.758  10.607  -1.620  1.00 83.88      L21B C
ATOM   6145  O   SER    17      81.204  10.812  -0.490  1.00 82.32      L21B O
ATOM   6146  N   ILE    18      80.116  11.539  -2.318  1.00 83.58      L21B N
ATOM   6147  CA  ILE    18      79.808  12.850  -1.751  1.00 83.01      L21B C
ATOM   6148  CB  ILE    18      79.972  13.974  -2.794  1.00 82.86      L21B C
ATOM   6149  CG2 ILE    18      81.394  14.495  -2.776  1.00 80.11      L21B C
ATOM   6150  CG1 ILE    18      79.581  13.462  -4.182  1.00 85.84      L21B C
ATOM   6151  CD1 ILE    18      78.174  12.921  -4.275  1.00 90.99      L21B C
ATOM   6152  C   ILE    18      78.394  12.925  -1.211  1.00 81.22      L21B C
ATOM   6153  O   ILE    18      77.562  12.068  -1.492  1.00 83.09      L21B O
ATOM   6154  N   THR    19      78.117  13.960  -0.433  1.00 77.37      L21B N
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ATOM   6155  CA  THR  19      76.739  14.264  -0.115  1.00 73.65      L21B C
ATOM   6156  CB  THR  19      76.344  13.699   1.250  1.00 72.21      L21B C
ATOM   6157  OG1 THR  19      76.707  14.624   2.277  1.00 70.39      L21B O
ATOM   6158  CG2 THR  19      77.049  12.387   1.488  1.00 70.27      L21B C
ATOM   6159  C   THR  19      76.458  15.765  -0.134  1.00 71.42      L21B C
ATOM   6160  O   THR  19      77.370  16.590  -0.039  1.00 71.11      L21B O
ATOM   6161  N   ILE  20      75.178  16.100  -0.261  1.00 69.05      L21B N
ATOM   6162  CA  ILE  20      74.737  17.473  -0.418  1.00 65.68      L21B C
ATOM   6163  CB  ILE  20      74.059  17.658  -1.789  1.00 61.69      L21B C
ATOM   6164  CG2 ILE  20      73.641  19.087  -1.966  1.00 61.81      L21B C
ATOM   6165  CG1 ILE  20      75.028  17.270  -2.906  1.00 58.15      L21B C
ATOM   6166  CD1 ILE  20      74.481  17.454  -4.286  1.00 54.31      L21B C
ATOM   6167  C   ILE  20      73.773  17.862   0.700  1.00 65.38      L21B C
ATOM   6168  O   ILE  20      72.859  17.120   1.041  1.00 65.12      L21B O
ATOM   6169  N   SER  21      73.988  19.036   1.270  1.00 66.68      L21B N
ATOM   6170  CA  SER  21      73.178  19.513   2.381  1.00 67.79      L21B C
ATOM   6171  CB  SER  21      74.038  20.377   3.303  1.00 68.52      L21B C
ATOM   6172  OG  SER  21      73.230  21.136   4.187  1.00 70.90      L21B O
ATOM   6173  C   SER  21      71.953  20.314   1.930  1.00 67.36      L21B C
ATOM   6174  O   SER  21      71.994  21.035   0.932  1.00 66.91      L21B O
ATOM   6175  N   CYS  22      70.868  20.194   2.689  1.00 67.57      L21B N
ATOM   6176  CA  CYS  22      69.663  20.984   2.452  1.00 68.05      L21B C
ATOM   6177  C   CYS  22      69.109  21.427   3.798  1.00 69.14      L21B C
ATOM   6178  O   CYS  22      68.334  20.711   4.402  1.00 69.10      L21B O
ATOM   6179  CB  CYS  22      68.629  20.118   1.732  1.00 67.60      L21B C
ATOM   6180  SG  CYS  22      67.077  20.933   1.216  1.00 65.97      L21B S
ATOM   6181  N   THR  23      69.495  22.593   4.290  1.00 72.15      L21B N
ATOM   6182  CA  THR  23      69.051  22.958   5.628  1.00 76.33      L21B C
ATOM   6183  CB  THR  23      70.205  23.520   6.473  1.00 75.55      L21B C
ATOM   6184  OG1 THR  23      70.577  24.806   5.972  1.00 78.73      L21B O
ATOM   6185  CG2 THR  23      71.409  22.595   6.407  1.00 75.50      L21B C
ATOM   6186  C   THR  23      67.903  23.967   5.615  1.00 78.37      L21B C
ATOM   6187  O   THR  23      68.068  25.123   5.217  1.00 79.56      L21B O
ATOM   6188  N   GLY  24      66.737  23.509   6.059  1.00 79.38      L21B N
ATOM   6189  CA  GLY  24      65.554  24.344   6.057  1.00 80.39      L21B C
ATOM   6190  C   GLY  24      65.173  24.696   7.472  1.00 81.41      L21B C
ATOM   6191  O   GLY  24      66.042  24.918   8.306  1.00 81.58      L21B O
ATOM   6192  N   THR  25      63.878  24.747   7.757  1.00 83.11      L21B N
ATOM   6193  CA  THR  25      63.429  25.146   9.085  1.00 85.45      L21B C
ATOM   6194  CB  THR  25      62.866  26.591   9.071  1.00 86.14      L21B C
ATOM   6195  OG1 THR  25      61.621  26.629   8.360  1.00 87.19      L21B O
ATOM   6196  CG2 THR  25      63.846  27.528   8.388  1.00 86.73      L21B C
ATOM   6197  C   THR  25      62.372  24.216   9.692  1.00 86.38      L21B C
ATOM   6198  O   THR  25      62.153  23.098   9.217  1.00 85.39      L21B O
ATOM   6199  N   SER  26      61.735  24.693  10.757  1.00 87.61      L21B N
ATOM   6200  CA  SER  26      60.632  23.990  11.394  1.00 89.14      L21B C
ATOM   6201  CB  SER  26      60.383  24.581  12.781  1.00 87.79      L21B C
ATOM   6202  OG  SER  26      60.268  25.992  12.708  1.00 85.49      L21B O
ATOM   6203  C   SER  26      59.375  24.120  10.538  1.00 91.27      L21B C
ATOM   6204  O   SER  26      58.300  23.680  10.923  1.00 92.38      L21B O
ATOM   6205  N   SER  27      59.520  24.737   9.374  1.00 93.31      L21B N
ATOM   6206  CA  SER  27      58.409  24.899   8.452  1.00 94.36      L21B C
ATOM   6207  CB  SER  27      58.176  26.382   8.174  1.00 94.15      L21B C
ATOM   6208  OG  SER  27      58.262  27.140   9.369  1.00 91.23      L21B O
ATOM   6209  C   SER  27      58.730  24.186   7.148  1.00 95.59      L21B C
ATOM   6210  O   SER  27      57.837  23.851   6.376  1.00 97.39      L21B O
ATOM   6211  N   ASP  28      60.019  23.964   6.913  1.00 96.07      L21B N
ATOM   6212  CA  ASP  28      60.507  23.454   5.637  1.00 96.19      L21B C
ATOM   6213  CB  ASP  28      61.825  24.146   5.270  1.00 98.10      L21B C
ATOM   6214  CG  ASP  28      61.718  25.670   5.281  1.00 99.61      L21B C
ATOM   6215  OD1 ASP  28      62.749  26.351   5.067  1.00 97.89      L21B O
ATOM   6216  OD2 ASP  28      60.603  26.188   5.502  1.00103.08      L21B O
ATOM   6217  C   ASP  28      60.724  21.945   5.704  1.00 96.20      L21B C
ATOM   6218  O   ASP  28      59.811  21.163   5.451  1.00 94.14      L21B O
ATOM   6219  N   VAL  29      61.938  21.535   6.046  1.00 97.12      L21B N
ATOM   6220  CA  VAL  29      62.221  20.122   6.218  1.00 99.22      L21B C
ATOM   6221  CB  VAL  29      63.636  19.773   5.739  1.00 98.78      L21B C
ATOM   6222  CG1 VAL  29      63.907  20.450   4.416  1.00 98.60      L21B C
ATOM   6223  CG2 VAL  29      64.651  20.188   6.771  1.00 98.58      L21B C
ATOM   6224  C   VAL  29      62.108  19.788   7.695  1.00100.89      L21B C
ATOM   6225  O   VAL  29      62.414  18.677   8.110  1.00102.16      L21B O
ATOM   6226  N   GLY  30      61.671  20.767   8.481  1.00102.57      L21B N
ATOM   6227  CA  GLY  30      61.450  20.546   9.899  1.00105.19      L21B C
ATOM   6228  C   GLY  30      60.199  19.727  10.159  1.00106.33      L21B C
ATOM   6229  O   GLY  30      60.273  18.632  10.716  1.00106.20      L21B O
ATOM   6230  N   GLY  31      59.047  20.260   9.764  1.00106.57      L21B N
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ATOM   6231  CA  GLY  31      57.841  19.452   9.704  1.00108.37      L21B C
ATOM   6232  C   GLY  31      57.715  18.872   8.308  1.00110.31      L21B C
ATOM   6233  O   GLY  31      58.503  19.220   7.432  1.00110.17      L21B O
ATOM   6234  N   TYR  32      56.739  17.997   8.088  1.00113.44      L21B N
ATOM   6235  CA  TYR  32      56.571  17.350   6.789  1.00116.17      L21B C
ATOM   6236  CB  TYR  32      56.262  18.386   5.701  1.00116.41      L21B C
ATOM   6237  CG  TYR  32      55.157  19.341   6.065  1.00116.91      L21B C
ATOM   6238  CD1 TYR  32      55.442  20.632   6.478  1.00116.46      L21B C
ATOM   6239  CE1 TYR  32      54.436  21.495   6.863  1.00116.02      L21B C
ATOM   6240  CD2 TYR  32      53.832  18.938   6.040  1.00117.30      L21B C
ATOM   6241  CE2 TYR  32      52.819  19.792   6.423  1.00116.92      L21B C
ATOM   6242  CZ  TYR  32      53.127  21.068   6.836  1.00116.30      L21B C
ATOM   6243  OH  TYR  32      52.120  21.910   7.244  1.00115.72      L21B O
ATOM   6244  C   TYR  32      57.824  16.580   6.393  1.00116.81      L21B C
ATOM   6245  O   TYR  32      58.878  17.165   6.156  1.00119.69      L21B O
ATOM   6246  N   ASN  33      57.707  15.262   6.314  1.00115.62      L21B N
ATOM   6247  CA  ASN  33      58.799  14.445   5.812  1.00113.19      L21B C
ATOM   6248  CB  ASN  33      58.848  13.114   6.566  1.00117.95      L21B C
ATOM   6249  CG  ASN  33      59.306  13.282   8.009  1.00119.92      L21B C
ATOM   6250  OD1 ASN  33      60.493  13.158   8.310  1.00121.58      L21B O
ATOM   6251  ND2 ASN  33      58.367  13.572   8.904  1.00121.03      L21B N
ATOM   6252  C   ASN  33      58.608  14.220   4.323  1.00109.46      L21B C
ATOM   6253  O   ASN  33      58.512  13.088   3.852  1.00106.99      L21B O
ATOM   6254  N   SER  34      58.548  15.325   3.591  1.00106.79      L21B N
ATOM   6255  CA  SER  34      58.362  15.283   2.152  1.00104.32      L21B C
ATOM   6256  CB  SER  34      56.946  15.740   1.803  1.00106.50      L21B C
ATOM   6257  OG  SER  34      55.978  14.885   2.390  1.00111.24      L21B O
ATOM   6258  C   SER  34      59.392  16.158   1.439  1.00100.10      L21B C
ATOM   6259  O   SER  34      59.077  17.247   0.958  1.00 99.91      L21B O
ATOM   6260  N   VAL  35      60.626  15.664   1.385  1.00 94.55      L21B N
ATOM   6261  CA  VAL  35      61.709  16.306   0.646  1.00 89.79      L21B C
ATOM   6262  CB  VAL  35      62.984  16.444   1.524  1.00 92.16      L21B C
ATOM   6263  CG1 VAL  35      64.129  16.991   0.699  1.00 93.68      L21B C
ATOM   6264  CG2 VAL  35      62.715  17.366   2.701  1.00 92.64      L21B C
ATOM   6265  C   VAL  35      62.066  15.512  -0.616  1.00 84.92      L21B C
ATOM   6266  O   VAL  35      62.069  14.280  -0.616  1.00 83.72      L21B O
ATOM   6267  N   SER  36      62.367  16.227  -1.694  1.00 79.11      L21B N
ATOM   6268  CA  SER  36      62.736  15.591  -2.953  1.00 72.73      L21B C
ATOM   6269  CB  SER  36      61.610  15.776  -3.973  1.00 73.26      L21B C
ATOM   6270  OG  SER  36      60.416  15.151  -3.537  1.00 71.59      L21B O
ATOM   6271  C   SER  36      64.048  16.150  -3.514  1.00 67.49      L21B C
ATOM   6272  O   SER  36      64.421  17.287  -3.221  1.00 67.42      L21B O
ATOM   6273  N   TRP  37      64.747  15.345  -4.311  1.00 59.72      L21B N
ATOM   6274  CA  TRP  37      65.956  15.799  -4.991  1.00 53.82      L21B C
ATOM   6275  CB  TRP  37      67.188  14.986  -4.533  1.00 52.55      L21B C
ATOM   6276  CG  TRP  37      67.512  15.188  -3.075  1.00 51.75      L21B C
ATOM   6277  CD2 TRP  37      68.289  16.250  -2.492  1.00 51.86      L21B C
ATOM   6278  CE2 TRP  37      68.218  16.097  -1.089  1.00 51.73      L21B C
ATOM   6279  CE3 TRP  37      69.031  17.314  -3.017  1.00 51.67      L21B C
ATOM   6280  CD1 TRP  37      67.032  14.457  -2.033  1.00 50.94      L21B C
ATOM   6281  NE1 TRP  37      67.446  14.996  -0.838  1.00 50.27      L21B N
ATOM   6282  CZ2 TRP  37      68.860  16.969  -0.201  1.00 52.65      L21B C
ATOM   6283  CZ3 TRP  37      69.671  18.180  -2.131  1.00 52.41      L21B C
ATOM   6284  CH2 TRP  37      69.578  18.000  -0.739  1.00 52.64      L21B C
ATOM   6285  C   TRP  37      65.789  15.691  -6.500  1.00 50.21      L21B C
ATOM   6286  O   TRP  37      65.254  14.707  -7.016  1.00 48.35      L21B O
ATOM   6287  N   TYR  38      66.232  16.715  -7.216  1.00 46.26      L21B N
ATOM   6288  CA  TYR  38      66.171  16.653  -8.663  1.00 44.46      L21B C
ATOM   6289  CB  TYR  38      65.166  17.680  -9.204  1.00 43.15      L21B C
ATOM   6290  CG  TYR  38      63.817  17.598  -8.542  1.00 43.72      L21B C
ATOM   6291  CD1 TYR  38      63.592  18.195  -7.300  1.00 43.01      L21B C
ATOM   6292  CE1 TYR  38      62.358  18.109  -6.671  1.00 43.19      L21B C
ATOM   6293  CD2 TYR  38      62.769  16.913  -9.139  1.00 43.98      L21B C
ATOM   6294  CE2 TYR  38      61.528  16.826  -8.514  1.00 45.21      L21B C
ATOM   6295  CZ  TYR  38      61.335  17.427  -7.284  1.00 44.43      L21B C
ATOM   6296  OH  TYR  38      60.110  17.343  -6.672  1.00 46.77      L21B O
ATOM   6297  C   TYR  38      67.538  16.887  -9.283  1.00 43.39      L21B C
ATOM   6298  O   TYR  38      68.369  17.634  -8.765  1.00 41.69      L21B O
ATOM   6299  N   GLN  39      67.756  16.250 -10.415  1.00 42.45      L21B N
ATOM   6300  CA  GLN  39      69.051  16.271 -11.045  1.00 44.62      L21B C
ATOM   6301  CB  GLN  39      69.563  14.830 -11.147  1.00 43.52      L21B C
ATOM   6302  CG  GLN  39      70.923  14.695 -11.736  1.00 43.60      L21B C
ATOM   6303  CD  GLN  39      71.010  13.530 -12.675  1.00 43.02      L21B C
ATOM   6304  OE1 GLN  39      70.373  13.515 -13.729  1.00 44.47      L21B O
ATOM   6305  NE2 GLN  39      71.803  12.541 -12.305  1.00 44.49      L21B N
ATOM   6306  C   GLN  39      68.918  16.917 -12.420  1.00 45.66      L21B C
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ATOM   6307  O   GLN    39      68.207  16.427 -13.288  1.00 46.08      L21B O
ATOM   6308  N   GLN    40      69.595  18.032 -12.624  1.00 47.83      L21B N
ATOM   6309  CA  GLN    40      69.399  18.751 -13.863  1.00 51.69      L21B C
ATOM   6310  CB  GLN    40      68.899  20.160 -13.548  1.00 47.93      L21B C
ATOM   6311  CG  GLN    40      68.128  20.864 -14.645  1.00 40.63      L21B C
ATOM   6312  CD  GLN    40      67.849  22.305 -14.288  1.00 40.95      L21B C
ATOM   6313  OE1 GLN    40      68.289  22.797 -13.245  1.00 38.49      L21B O
ATOM   6314  NE2 GLN    40      67.110  22.997 -15.147  1.00 40.56      L21B N
ATOM   6315  C   GLN    40      70.700  18.786 -14.648  1.00 56.56      L21B C
ATOM   6316  O   GLN    40      71.663  19.444 -14.257  1.00 56.79      L21B O
ATOM   6317  N   HIS    41      70.723  18.049 -15.751  1.00 63.38      L21B N
ATOM   6318  CA  HIS    41      71.838  18.105 -16.682  1.00 70.76      L21B C
ATOM   6319  CB  HIS    41      71.747  16.988 -17.723  1.00 76.62      L21B C
ATOM   6320  CG  HIS    41      72.508  15.752 -17.354  1.00 84.15      L21B C
ATOM   6321  CD2 HIS    41      73.110  14.817 -18.126  1.00 87.55      L21B C
ATOM   6322  ND1 HIS    41      72.715  15.363 -16.047  1.00 85.99      L21B N
ATOM   6323  CE1 HIS    41      73.412  14.241 -16.031  1.00 89.12      L21B C
ATOM   6324  NE2 HIS    41      73.664  13.888 -17.279  1.00 90.00      L21B N
ATOM   6325  C   HIS    41      71.762  19.431 -17.386  1.00 72.18      L21B C
ATOM   6326  O   HIS    41      70.686  19.880 -17.756  1.00 72.48      L21B O
ATOM   6327  N   PRO    42      72.910  20.070 -17.602  1.00 74.10      L21B N
ATOM   6328  CD  PRO    42      74.271  19.528 -17.437  1.00 75.72      L21B C
ATOM   6329  CA  PRO    42      72.911  21.425 -18.160  1.00 74.30      L21B C
ATOM   6330  CB  PRO    42      74.385  21.677 -18.467  1.00 75.42      L21B C
ATOM   6331  CG  PRO    42      75.127  20.753 -17.534  1.00 76.94      L21B C
ATOM   6332  C   PRO    42      72.039  21.510 -19.413  1.00 73.03      L21B C
ATOM   6333  O   PRO    42      72.244  20.767 -20.374  1.00 73.33      L21B O
ATOM   6334  N   GLY    43      71.056  22.406 -19.391  1.00 70.77      L21B N
ATOM   6335  CA  GLY    43      70.264  22.663 -20.580  1.00 69.10      L21B C
ATOM   6336  C   GLY    43      69.196  21.615 -20.847  1.00 68.64      L21B C
ATOM   6337  O   GLY    43      68.511  21.650 -21.877  1.00 68.71      L21B O
ATOM   6338  N   LYS    44      69.050  20.672 -19.924  1.00 66.15      L21B N
ATOM   6339  CA  LYS    44      67.954  19.718 -19.990  1.00 62.25      L21B C
ATOM   6340  CB  LYS    44      68.496  18.293 -19.887  1.00 63.41      L21B C
ATOM   6341  CG  LYS    44      68.098  17.408 -21.052  1.00 63.60      L21B C
ATOM   6342  CD  LYS    44      69.236  17.236 -22.039  1.00 64.54      L21B C
ATOM   6343  CE  LYS    44      70.137  16.088 -21.614  1.00 66.95      L21B C
ATOM   6344  NZ  LYS    44      69.357  14.844 -21.329  1.00 67.67      L21B N
ATOM   6345  C   LYS    44      66.993  20.014 -18.841  1.00 59.21      L21B C
ATOM   6346  O   LYS    44      67.323  20.792 -17.943  1.00 60.37      L21B O
ATOM   6347  N   ALA    45      65.807  19.410 -18.873  1.00 53.31      L21B N
ATOM   6348  CA  ALA    45      64.799  19.666 -17.849  1.00 47.57      L21B C
ATOM   6349  CB  ALA    45      63.420  19.452 -18.421  1.00 46.62      L21B C
ATOM   6350  C   ALA    45      65.016  18.757 -16.653  1.00 44.16      L21B C
ATOM   6351  O   ALA    45      65.471  17.630 -16.813  1.00 45.35      L21B O
ATOM   6352  N   PRO    46      64.699  19.234 -15.436  1.00 40.96      L21B N
ATOM   6353  CD  PRO    46      64.061  20.525 -15.160  1.00 39.13      L21B C
ATOM   6354  CA  PRO    46      64.986  18.509 -14.194  1.00 39.73      L21B C
ATOM   6355  CB  PRO    46      64.389  19.394 -13.109  1.00 37.66      L21B C
ATOM   6356  CG  PRO    46      64.330  20.716 -13.706  1.00 37.73      L21B C
ATOM   6357  C   PRO    46      64.345  17.137 -14.197  1.00 39.44      L21B C
ATOM   6358  O   PRO    46      63.377  16.903 -14.928  1.00 39.36      L21B O
ATOM   6359  N   LYS    47      64.882  16.245 -13.364  1.00 38.72      L21B N
ATOM   6360  CA  LYS    47      64.447  14.848 -13.300  1.00 38.21      L21B C
ATOM   6361  CB  LYS    47      65.377  13.983 -14.145  1.00 38.85      L21B C
ATOM   6362  CG  LYS    47      64.962  12.549 -14.275  1.00 40.70      L21B C
ATOM   6363  CD  LYS    47      66.095  11.628 -13.885  1.00 41.88      L21B C
ATOM   6364  CE  LYS    47      65.859  10.233 -14.427  1.00 41.50      L21B C
ATOM   6365  NZ  LYS    47      66.936   9.330 -13.966  1.00 43.50      L21B N
ATOM   6366  C   LYS    47      64.482  14.372 -11.854  1.00 38.19      L21B C
ATOM   6367  O   LYS    47      65.471  14.592 -11.163  1.00 39.09      L21B O
ATOM   6368  N   LEU    48      63.415  13.708 -11.402  1.00 38.85      L21B N
ATOM   6369  CA  LEU    48      63.282  13.342  -9.998  1.00 38.42      L21B C
ATOM   6370  CB  LEU    48      61.821  13.069  -9.662  1.00 36.60      L21B C
ATOM   6371  CG  LEU    48      61.539  12.550  -8.241  1.00 34.74      L21B C
ATOM   6372  CD1 LEU    48      61.520  13.712  -7.240  1.00 32.22      L21B C
ATOM   6373  CD2 LEU    48      60.213  11.808  -8.233  1.00 30.83      L21B C
ATOM   6374  C   LEU    48      64.121  12.124  -9.609  1.00 42.07      L21B C
ATOM   6375  O   LEU    48      64.150  11.112 -10.323  1.00 39.62      L21B O
ATOM   6376  N   MET    49      64.778  12.242  -8.448  1.00 46.56      L21B N
ATOM   6377  CA  MET    49      65.732  11.263  -7.926  1.00 49.08      L21B C
ATOM   6378  CB  MET    49      67.106  11.906  -7.794  1.00 47.33      L21B C
ATOM   6379  CG  MET    49      68.147  11.319  -8.707  1.00 46.66      L21B C
ATOM   6380  SD  MET    49      67.781  11.564 -10.453  1.00 49.31      L21B S
ATOM   6381  CE  MET    49      69.100  10.529 -11.241  1.00 43.99      L21B C
ATOM   6382  C   MET    49      65.324  10.710  -6.567  1.00 52.99      L21B C
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ATOM   6383  O   MET  49      65.582   9.552  -6.269  1.00 56.08      L21B O
ATOM   6384  N   ILE  50      64.698  11.542  -5.740  1.00 57.38      L21B N
ATOM   6385  CA  ILE  50      64.297  11.149  -4.386  1.00 62.77      L21B C
ATOM   6386  CB  ILE  50      65.429  11.454  -3.365  1.00 61.84      L21B C
ATOM   6387  CG2 ILE  50      64.975  11.128  -1.946  1.00 59.34      L21B C
ATOM   6388  CG1 ILE  50      66.674  10.633  -3.701  1.00 62.37      L21B C
ATOM   6389  CD1 ILE  50      66.568   9.172  -3.330  1.00 62.87      L21B C
ATOM   6390  C   ILE  50      63.024  11.882  -3.937  1.00 67.31      L21B C
ATOM   6391  O   ILE  50      62.874  13.075  -4.172  1.00 67.56      L21B O
ATOM   6392  N   TYR  51      62.108  11.163  -3.297  1.00 72.88      L21B N
ATOM   6393  CA  TYR  51      60.957  11.792  -2.655  1.00 77.81      L21B C
ATOM   6394  CB  TYR  51      59.684  11.567  -3.470  1.00 80.69      L21B C
ATOM   6395  CG  TYR  51      59.343  10.120  -3.728  1.00 85.48      L21B C
ATOM   6396  CD1 TYR  51      58.614   9.378  -2.805  1.00 86.81      L21B C
ATOM   6397  CE1 TYR  51      58.273   8.064  -3.059  1.00 91.20      L21B C
ATOM   6398  CD2 TYR  51      59.723   9.504  -4.913  1.00 87.17      L21B C
ATOM   6399  CE2 TYR  51      59.388   8.196  -5.175  1.00 91.86      L21B C
ATOM   6400  CZ  TYR  51      58.664   7.481  -4.249  1.00 93.04      L21B C
ATOM   6401  OH  TYR  51      58.334   6.177  -4.525  1.00 98.22      L21B O
ATOM   6402  C   TYR  51      60.751  11.251  -1.254  1.00 78.65      L21B C
ATOM   6403  O   TYR  51      61.310  10.221  -0.891  1.00 80.21      L21B O
ATOM   6404  N   GLU  52      59.942  11.949  -0.470  1.00 79.50      L21B N
ATOM   6405  CA  GLU  52      59.763  11.598   0.928  1.00 80.88      L21B C
ATOM   6406  CB  GLU  52      58.756  10.452   1.052  1.00 84.81      L21B C
ATOM   6407  CG  GLU  52      57.474  10.696   0.252  1.00 91.28      L21B C
ATOM   6408  CD  GLU  52      56.273   9.900   0.749  1.00 94.35      L21B C
ATOM   6409  OE1 GLU  52      55.551   9.333  -0.097  1.00 96.91      L21B O
ATOM   6410  OE2 GLU  52      56.043   9.846   1.978  1.00 97.80      L21B O
ATOM   6411  C   GLU  52      61.119  11.191   1.496  1.00 79.08      L21B C
ATOM   6412  O   GLU  52      61.477  10.020   1.495  1.00 78.80      L21B O
ATOM   6413  N   VAL  53      61.856  12.192   1.967  1.00 77.54      L21B N
ATOM   6414  CA  VAL  53      63.236  12.003   2.446  1.00 75.58      L21B C
ATOM   6415  CB  VAL  53      63.293  11.922   3.981  1.00 76.57      L21B C
ATOM   6416  CG1 VAL  53      62.286  12.866   4.630  1.00 76.93      L21B C
ATOM   6417  CG2 VAL  53      63.040  10.470   4.380  1.00 78.24      L21B C
ATOM   6418  C   VAL  53      64.137  11.017   1.839  1.00 72.99      L21B C
ATOM   6419  O   VAL  53      65.296  11.298   1.555  1.00 72.93      L21B O
ATOM   6420  N   SER  54      63.623   9.807   1.638  1.00 70.73      L21B N
ATOM   6421  CA  SER  54      64.491   8.664   1.383  1.00 69.14      L21B C
ATOM   6422  CB  SER  54      64.597   7.798   2.641  1.00 67.96      L21B C
ATOM   6423  OG  SER  54      63.409   7.057   2.862  1.00 66.01      L21B O
ATOM   6424  C   SER  54      64.110   7.774   0.205  1.00 68.64      L21B C
ATOM   6425  O   SER  54      64.851   6.858  -0.132  1.00 68.13      L21B O
ATOM   6426  N   ASN  55      62.975   8.030  -0.430  1.00 68.81      L21B N
ATOM   6427  CA  ASN  55      62.484   7.095  -1.435  1.00 70.32      L21B C
ATOM   6428  CB  ASN  55      60.962   7.070  -1.418  1.00 68.63      L21B C
ATOM   6429  CG  ASN  55      60.417   6.256  -0.273  1.00 66.99      L21B C
ATOM   6430  OD1 ASN  55      60.253   6.756   0.840  1.00 65.31      L21B O
ATOM   6431  ND2 ASN  55      60.143   4.981  -0.536  1.00 67.92      L21B N
ATOM   6432  C   ASN  55      62.974   7.353  -2.850  1.00 72.09      L21B C
ATOM   6433  O   ASN  55      63.184   8.494  -3.242  1.00 72.67      L21B O
ATOM   6434  N   ARG  56      63.143   6.279  -3.614  1.00 74.41      L21B N
ATOM   6435  CA  ARG  56      63.667   6.376  -4.969  1.00 78.24      L21B C
ATOM   6436  CB  ARG  56      64.995   5.619  -5.070  1.00 78.52      L21B C
ATOM   6437  CG  ARG  56      65.465   5.352  -6.488  1.00 79.19      L21B C
ATOM   6438  CD  ARG  56      66.854   4.744  -6.501  1.00 81.29      L21B C
ATOM   6439  NE  ARG  56      66.905   3.448  -5.831  1.00 85.09      L21B N
ATOM   6440  CZ  ARG  56      67.379   3.257  -4.602  1.00 87.03      L21B C
ATOM   6441  NH1 ARG  56      67.382   2.036  -4.078  1.00 89.77      L21B N
ATOM   6442  NH2 ARG  56      67.846   4.281  -3.897  1.00 87.54      L21B N
ATOM   6443  C   ARG  56      62.700   5.845  -6.025  1.00 78.99      L21B C
ATOM   6444  O   ARG  56      62.260   4.700  -5.958  1.00 81.35      L21B O
ATOM   6445  N   PRO  57      62.361   6.679  -7.022  1.00 78.11      L21B N
ATOM   6446  CD  PRO  57      62.335   8.145  -6.931  1.00 74.41      L21B C
ATOM   6447  CA  PRO  57      61.659   6.211  -8.222  1.00 80.05      L21B C
ATOM   6448  CB  PRO  57      61.704   7.410  -9.177  1.00 75.32      L21B C
ATOM   6449  CG  PRO  57      62.227   8.580  -8.372  1.00 73.32      L21B C
ATOM   6450  C   PRO  57      62.341   4.997  -8.834  1.00 82.95      L21B C
ATOM   6451  O   PRO  57      63.301   4.476  -8.279  1.00 85.80      L21B O
ATOM   6452  N   SER  58      61.840   4.549  -9.980  1.00 84.28      L21B N
ATOM   6453  CA  SER  58      62.325   3.322 -10.595  1.00 85.65      L21B C
ATOM   6454  CB  SER  58      61.282   2.768 -11.564  1.00 86.16      L21B C
ATOM   6455  OG  SER  58      60.341   1.955 -10.881  1.00 86.06      L21B O
ATOM   6456  C   SER  58      63.660   3.471 -11.323  1.00 86.71      L21B C
ATOM   6457  O   SER  58      64.612   2.751 -11.028  1.00 87.93      L21B O
ATOM   6458  N   GLY  59      63.733   4.397 -12.272  1.00 87.65      L21B N
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ATOM   6459  CA  GLY    59      64.947   4.547 -13.056  1.00 87.89      L21B C
ATOM   6460  C   GLY    59      66.181   4.868 -12.229  1.00 88.35      L21B C
ATOM   6461  O   GLY    59      67.286   4.431 -12.552  1.00 88.22      L21B O
ATOM   6462  N   VAL    60      66.002   5.628 -11.156  1.00 89.24      L21B N
ATOM   6463  CA  VAL    60      67.131   6.077 -10.355  1.00 90.52      L21B C
ATOM   6464  CB  VAL    60      66.649   6.943  -9.176  1.00 90.94      L21B C
ATOM   6465  CG1 VAL    60      67.830   7.355  -8.309  1.00 90.92      L21B C
ATOM   6466  CG2 VAL    60      65.926   8.165  -9.698  1.00 89.49      L21B C
ATOM   6467  C   VAL    60      67.965   4.917  -9.820  1.00 90.98      L21B C
ATOM   6468  O   VAL    60      67.431   3.962  -9.263  1.00 90.77      L21B O
ATOM   6469  N   SER    61      69.279   5.020 -10.000  1.00 92.18      L21B N
ATOM   6470  CA  SER    61      70.220   3.994  -9.562  1.00 93.58      L21B C
ATOM   6471  CB  SER    61      71.616   4.292 -10.126  1.00 93.45      L21B C
ATOM   6472  OG  SER    61      72.558   3.294  -9.762  1.00 93.35      L21B O
ATOM   6473  C   SER    61      70.284   3.911  -8.038  1.00 94.34      L21B C
ATOM   6474  O   SER    61      69.902   4.844  -7.332  1.00 93.56      L21B O
ATOM   6475  N   ASN    62      70.762   2.779  -7.535  1.00 95.84      L21B N
ATOM   6476  CA  ASN    62      70.855   2.564  -6.098  1.00 96.77      L21B C
ATOM   6477  CB  ASN    62      71.117   1.074  -5.795  1.00 97.06      L21B C
ATOM   6478  CG  ASN    62      71.867   0.350  -6.920  1.00 97.69      L21B C
ATOM   6479  OD1 ASN    62      72.751  -0.470  -6.662  1.00 94.14      L21B O
ATOM   6480  ND2 ASN    62      71.504   0.638  -8.167  1.00 99.77      L21B N
ATOM   6481  C   ASN    62      71.936   3.445  -5.465  1.00 96.37      L21B C
ATOM   6482  O   ASN    62      72.013   3.566  -4.238  1.00 97.21      L21B O
ATOM   6483  N   ARG    63      72.756   4.067  -6.313  1.00 94.95      L21B N
ATOM   6484  CA  ARG    63      73.833   4.953  -5.862  1.00 92.26      L21B C
ATOM   6485  CB  ARG    63      74.654   5.450  -7.056  1.00 96.28      L21B C
ATOM   6486  CG  ARG    63      75.721   4.486  -7.538  1.00102.27      L21B C
ATOM   6487  CD  ARG    63      76.425   5.012  -8.780  1.00109.64      L21B C
ATOM   6488  NE  ARG    63      75.705   4.669 -10.006  1.00117.20      L21B N
ATOM   6489  CZ  ARG    63      76.177   4.858 -11.237  1.00120.97      L21B C
ATOM   6490  NH1 ARG    63      77.379   5.397 -11.416  1.00123.18      L21B N
ATOM   6491  NH2 ARG    63      75.449   4.501 -12.290  1.00122.26      L21B N
ATOM   6492  C   ARG    63      73.279   6.154  -5.107  1.00 88.51      L21B C
ATOM   6493  O   ARG    63      73.897   6.643  -4.162  1.00 87.03      L21B O
ATOM   6494  N   PHE    64      72.112   6.627  -5.533  1.00 84.21      L21B N
ATOM   6495  CA  PHE    64      71.492   7.792  -4.926  1.00 79.46      L21B C
ATOM   6496  CB  PHE    64      70.617   8.514  -5.947  1.00 77.20      L21B C
ATOM   6497  CG  PHE    64      71.371   9.032  -7.139  1.00 73.89      L21B C
ATOM   6498  CD1 PHE    64      72.315  10.034  -6.997  1.00 71.50      L21B C
ATOM   6499  CD2 PHE    64      71.116   8.533  -8.409  1.00 73.34      L21B C
ATOM   6500  CE1 PHE    64      72.995  10.531  -8.099  1.00 69.83      L21B C
ATOM   6501  CE2 PHE    64      71.791   9.026  -9.520  1.00 71.53      L21B C
ATOM   6502  CZ  PHE    64      72.731  10.028  -9.363  1.00 70.20      L21B C
ATOM   6503  C   PHE    64      70.643   7.384  -3.734  1.00 77.58      L21B C
ATOM   6504  O   PHE    64      69.857   6.447  -3.817  1.00 77.96      L21B O
ATOM   6505  N   SER    65      70.805   8.101  -2.629  1.00 75.04      L21B N
ATOM   6506  CA  SER    65      70.039   7.851  -1.413  1.00 74.86      L21B C
ATOM   6507  CB  SER    65      70.851   6.980  -0.459  1.00 74.40      L21B C
ATOM   6508  OG  SER    65      72.221   7.354  -0.478  1.00 74.90      L21B O
ATOM   6509  C   SER    65      69.646   9.142  -0.702  1.00 75.62      L21B C
ATOM   6510  O   SER    65      70.332  10.157  -0.808  1.00 75.41      L21B O
ATOM   6511  N   GLY    66      68.541   9.106   0.030  1.00 76.47      L21B N
ATOM   6512  CA  GLY    66      68.127  10.295   0.745  1.00 77.49      L21B C
ATOM   6513  C   GLY    66      68.129  10.100   2.243  1.00 79.01      L21B C
ATOM   6514  O   GLY    66      68.074   8.975   2.718  1.00 79.00      L21B O
ATOM   6515  N   SER    67      68.187  11.197   2.985  1.00 81.38      L21B N
ATOM   6516  CA  SER    67      68.061  11.146   4.432  1.00 84.01      L21B C
ATOM   6517  CB  SER    67      69.374  10.708   5.067  1.00 83.67      L21B C
ATOM   6518  OG  SER    67      69.386  11.030   6.454  1.00 79.98      L21B O
ATOM   6519  C   SER    67      67.682  12.510   4.981  1.00 86.52      L21B C
ATOM   6520  O   SER    67      67.820  13.520   4.296  1.00 88.88      L21B O
ATOM   6521  N   LYS    68      67.202  12.539   6.219  1.00 87.89      L21B N
ATOM   6522  CA  LYS    68      66.874  13.798   6.879  1.00 89.48      L21B C
ATOM   6523  CB  LYS    68      65.350  13.985   6.906  1.00 90.65      L21B C
ATOM   6524  CG  LYS    68      64.889  15.386   7.302  1.00 91.64      L21B C
ATOM   6525  CD  LYS    68      63.737  15.347   8.302  1.00 93.17      L21B C
ATOM   6526  CE  LYS    68      62.378  15.370   7.615  1.00 94.61      L21B C
ATOM   6527  NZ  LYS    68      62.140  16.629   6.858  1.00 96.05      L21B N
ATOM   6528  C   LYS    68      67.435  13.789   8.309  1.00 90.68      L21B C
ATOM   6529  O   LYS    68      68.008  12.794   8.748  1.00 90.52      L21B O
ATOM   6530  N   SER    69      67.283  14.905   9.017  1.00 92.64      L21B N
ATOM   6531  CA  SER    69      67.631  15.009  10.434  1.00 94.30      L21B C
ATOM   6532  CB  SER    69      69.064  14.526  10.685  1.00 94.54      L21B C
ATOM   6533  OG  SER    69      69.420  14.675  12.050  1.00 94.63      L21B O
ATOM   6534  C   SER    69      67.504  16.465  10.853  1.00 94.68      L21B C
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ATOM   6535  O   SER   69      67.867  17.365  10.099  1.00 95.48      L21B O
ATOM   6536  N   GLY   70      66.988  16.699  12.053  1.00 93.84      L21B N
ATOM   6537  CA  GLY   70      66.749  18.063  12.471  1.00 93.22      L21B C
ATOM   6538  C   GLY   70      66.095  18.822  11.338  1.00 93.55      L21B C
ATOM   6539  O   GLY   70      65.264  18.271  10.616  1.00 92.49      L21B O
ATOM   6540  N   ASN   71      66.476  20.082  11.166  1.00 94.45      L21B N
ATOM   6541  CA  ASN   71      65.925  20.890  10.089  1.00 94.23      L21B C
ATOM   6542  CB  ASN   71      65.702  22.323  10.571  1.00 95.39      L21B C
ATOM   6543  CG  ASN   71      64.724  22.400  11.730  1.00 98.12      L21B C
ATOM   6544  OD1 ASN   71      64.501  21.414  12.438  1.00 98.62      L21B O
ATOM   6545  ND2 ASN   71      64.131  23.574  11.930  1.00 98.55      L21B N
ATOM   6546  C   ASN   71      66.828  20.887   8.863  1.00 92.16      L21B C
ATOM   6547  O   ASN   71      66.873  21.862   8.119  1.00 94.10      L21B O
ATOM   6548  N   THR   72      67.541  19.786   8.648  1.00 87.06      L21B N
ATOM   6549  CA  THR   72      68.445  19.690   7.510  1.00 82.06      L21B C
ATOM   6550  CB  THR   72      69.884  20.104   7.900  1.00 82.77      L21B C
ATOM   6551  OG1 THR   72      70.792  19.717   6.860  1.00 80.39      L21B O
ATOM   6552  CG2 THR   72      70.291  19.458   9.213  1.00 83.47      L21B C
ATOM   6553  C   THR   72      68.495  18.307   6.865  1.00 78.60      L21B C
ATOM   6554  O   THR   72      69.052  17.363   7.418  1.00 77.66      L21B O
ATOM   6555  N   ALA   73      67.919  18.203   5.673  1.00 75.24      L21B N
ATOM   6556  CA  ALA   73      67.911  16.960   4.915  1.00 72.01      L21B C
ATOM   6557  CB  ALA   73      66.668  16.895   4.045  1.00 72.86      L21B C
ATOM   6558  C   ALA   73      69.151  16.890   4.048  1.00 69.53      L21B C
ATOM   6559  O   ALA   73      69.801  17.904   3.812  1.00 70.14      L21B O
ATOM   6560  N   SER   74      69.482  15.699   3.566  1.00 65.99      L21B N
ATOM   6561  CA  SER   74      70.686  15.548   2.761  1.00 62.02      L21B C
ATOM   6562  CB  SER   74      71.920  15.435   3.670  1.00 62.25      L21B C
ATOM   6563  OG  SER   74      71.556  15.188   5.022  1.00 60.36      L21B O
ATOM   6564  C   SER   74      70.648  14.371   1.802  1.00 59.13      L21B C
ATOM   6565  O   SER   74      70.224  13.276   2.153  1.00 57.73      L21B O
ATOM   6566  N   LEU   75      71.100  14.614   0.580  1.00 56.92      L21B N
ATOM   6567  CA  LEU   75      71.247  13.556  -0.405  1.00 56.96      L21B C
ATOM   6568  CB  LEU   75      70.961  14.109  -1.813  1.00 53.44      L21B C
ATOM   6569  CG  LEU   75      71.478  13.368  -3.055  1.00 49.90      L21B C
ATOM   6570  CD1 LEU   75      70.723  12.083  -3.242  1.00 47.92      L21B C
ATOM   6571  CD2 LEU   75      71.333  14.248  -4.283  1.00 47.50      L21B C
ATOM   6572  C   LEU   75      72.683  13.072  -0.319  1.00 59.01      L21B C
ATOM   6573  O   LEU   75      73.597  13.882  -0.273  1.00 58.67      L21B O
ATOM   6574  N   THR   76      72.887  11.760  -0.286  1.00 61.56      L21B N
ATOM   6575  CA  THR   76      74.217  11.203  -0.522  1.00 63.56      L21B C
ATOM   6576  CB  THR   76      74.689  10.351   0.679  1.00 63.91      L21B C
ATOM   6577  OG1 THR   76      75.672   9.404   0.239  1.00 62.00      L21B O
ATOM   6578  CG2 THR   76      73.520   9.630   1.317  1.00 65.76      L21B C
ATOM   6579  C   THR   76      74.301  10.363  -1.806  1.00 63.85      L21B C
ATOM   6580  O   THR   76      73.356   9.682  -2.181  1.00 64.85      L21B O
ATOM   6581  N   ILE   77      75.441  10.433  -2.481  1.00 64.17      L21B N
ATOM   6582  CA  ILE   77      75.659   9.668  -3.700  1.00 65.28      L21B C
ATOM   6583  CB  ILE   77      75.768  10.588  -4.929  1.00 66.15      L21B C
ATOM   6584  CG2 ILE   77      75.926   9.741  -6.193  1.00 65.87      L21B C
ATOM   6585  CG1 ILE   77      74.542  11.505  -5.005  1.00 66.23      L21B C
ATOM   6586  CD1 ILE   77      74.658  12.647  -6.022  1.00 63.85      L21B C
ATOM   6587  C   ILE   77      76.966   8.885  -3.595  1.00 66.96      L21B C
ATOM   6588  O   ILE   77      78.022   9.465  -3.335  1.00 66.30      L21B O
ATOM   6589  N   SER   78      76.900   7.574  -3.804  1.00 68.95      L21B N
ATOM   6590  CA  SER   78      78.101   6.746  -3.865  1.00 71.17      L21B C
ATOM   6591  CB  SER   78      77.860   5.437  -3.129  1.00 72.64      L21B C
ATOM   6592  OG  SER   78      76.778   4.739  -3.723  1.00 77.40      L21B O
ATOM   6593  C   SER   78      78.479   6.442  -5.313  1.00 71.61      L21B C
ATOM   6594  O   SER   78      77.635   6.483  -6.209  1.00 72.46      L21B O
ATOM   6595  N   GLY   79      79.750   6.143  -5.544  1.00 71.26      L21B N
ATOM   6596  CA  GLY   79      80.152   5.652  -6.849  1.00 72.39      L21B C
ATOM   6597  C   GLY   79      80.059   6.655  -7.983  1.00 73.11      L21B C
ATOM   6598  O   GLY   79      79.582   6.323  -9.076  1.00 73.09      L21B O
ATOM   6599  N   LEU   80      80.529   7.872  -7.729  1.00 73.02      L21B N
ATOM   6600  CA  LEU   80      80.457   8.954  -8.701  1.00 75.02      L21B C
ATOM   6601  CB  LEU   80      81.414  10.068  -8.289  1.00 73.44      L21B C
ATOM   6602  CG  LEU   80      80.769  11.379  -7.849  1.00 73.15      L21B C
ATOM   6603  CD1 LEU   80      81.385  11.863  -6.549  1.00 72.79      L21B C
ATOM   6604  CD2 LEU   80      80.951  12.403  -8.949  1.00 73.15      L21B C
ATOM   6605  C   LEU   80      80.799   8.496 -10.109  1.00 76.64      L21B C
ATOM   6606  O   LEU   80      81.746   7.744 -10.294  1.00 77.62      L21B O
ATOM   6607  N   GLN   81      80.033   8.949 -11.101  1.00 78.31      L21B N
ATOM   6608  CA  GLN   81      80.361   8.673 -12.499  1.00 80.13      L21B C
ATOM   6609  CB  GLN   81      79.558   7.482 -13.027  1.00 81.09      L21B C
ATOM   6610  CG  GLN   81      80.408   6.231 -13.313  1.00 86.00      L21B C
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ATOM   6611  CD  GLN    81      81.532   6.458 -14.348  1.00 87.47      L21B C
ATOM   6612  OE1 GLN    81      81.628   5.730 -15.344  1.00 88.68      L21B O
ATOM   6613  NE2 GLN    81      82.384   7.458 -14.105  1.00 87.99      L21B N
ATOM   6614  C   GLN    81      80.158   9.862 -13.425  1.00 80.81      L21B C
ATOM   6615  O   GLN    81      79.674  10.910 -13.011  1.00 80.44      L21B O
ATOM   6616  N   ALA    82      80.532   9.681 -14.688  1.00 83.61      L21B N
ATOM   6617  CA  ALA    82      80.618  10.779 -15.648  1.00 85.92      L21B C
ATOM   6618  CB  ALA    82      81.479  10.362 -16.830  1.00 87.14      L21B C
ATOM   6619  C   ALA    82      79.252  11.250 -16.143  1.00 87.29      L21B C
ATOM   6620  O   ALA    82      79.142  12.285 -16.800  1.00 87.53      L21B O
ATOM   6621  N   GLU    83      78.215  10.480 -15.836  1.00 88.20      L21B N
ATOM   6622  CA  GLU    83      76.853  10.872 -16.168  1.00 86.75      L21B C
ATOM   6623  CB  GLU    83      76.027   9.645 -16.581  1.00 93.51      L21B C
ATOM   6624  CG  GLU    83      75.485   8.968 -17.876  1.00101.49      L21B C
ATOM   6625  CD  GLU    83      76.352   9.864 -19.106  1.00105.65      L21B C
ATOM   6626  OE1 GLU    83      75.306  10.538 -19.249  1.00108.12      L21B O
ATOM   6627  OE2 GLU    83      77.297   9.889 -19.929  1.00108.37      L21B O
ATOM   6628  C   GLU    83      76.203  11.546 -14.967  1.00 87.10      L21B C
ATOM   6629  O   GLU    83      75.026  11.891 -15.004  1.00 85.86      L21B O
ATOM   6630  N   ASP    84      76.970  11.725 -13.899  1.00 85.49      L21B N
ATOM   6631  CA  ASP    84      76.427  12.298 -12.677  1.00 85.32      L21B C
ATOM   6632  CB  ASP    84      76.985  11.571 -11.457  1.00 85.04      L21B C
ATOM   6633  CG  ASP    84      76.359  10.212 -11.263  1.00 85.56      L21B C
ATOM   6634  OD1 ASP    84      75.368   9.921 -11.971  1.00 84.97      L21B O
ATOM   6635  OD2 ASP    84      76.854   9.441 -10.409  1.00 84.74      L21B O
ATOM   6636  C   ASP    84      76.709  13.782 -12.553  1.00 84.15      L21B C
ATOM   6637  O   ASP    84      76.047  14.488 -11.789  1.00 86.30      L21B O
ATOM   6638  N   GLU    85      77.696  14.258 -13.299  1.00 81.29      L21B N
ATOM   6639  CA  GLU    85      78.000  15.675 -13.284  1.00 77.09      L21B C
ATOM   6640  CB  GLU    85      79.103  15.990 -14.298  1.00 80.08      L21B C
ATOM   6641  CG  GLU    85      79.936  17.229 -13.971  1.00 82.41      L21B C
ATOM   6642  CD  GLU    85      81.225  17.298 -14.784  1.00 85.05      L21B C
ATOM   6643  OE1 GLU    85      81.886  18.365 -14.775  1.00 87.67      L21B O
ATOM   6644  OE2 GLU    85      81.579  16.281 -15.433  1.00 86.35      L21B O
ATOM   6645  C   GLU    85      76.707  16.378 -13.663  1.00 72.73      L21B C
ATOM   6646  O   GLU    85      76.233  16.248 -14.795  1.00 72.00      L21B O
ATOM   6647  N   ALA    86      76.126  17.093 -12.700  1.00 67.45      L21B N
ATOM   6648  CA  ALA    86      74.894  17.839 -12.930  1.00 62.87      L21B C
ATOM   6649  CB  ALA    86      73.738  16.877 -13.092  1.00 61.14      L21B C
ATOM   6650  C   ALA    86      74.588  18.828 -11.803  1.00 60.31      L21B C
ATOM   6651  O   ALA    86      75.329  18.920 -10.823  1.00 59.84      L21B O
ATOM   6652  N   ASP    87      73.488  19.569 -11.953  1.00 56.64      L21B N
ATOM   6653  CA  ASP    87      72.981  20.439 -10.892  1.00 51.63      L21B C
ATOM   6654  CB  ASP    87      72.359  21.694 -11.495  1.00 51.56      L21B C
ATOM   6655  CG  ASP    87      73.392  22.714 -11.931  1.00 49.78      L21B C
ATOM   6656  OD1 ASP    87      74.611  22.481 -11.745  1.00 46.62      L21B O
ATOM   6657  OD2 ASP    87      72.965  23.759 -12.464  1.00 49.17      L21B O
ATOM   6658  C   ASP    87      71.925  19.711 -10.067  1.00 48.54      L21B C
ATOM   6659  O   ASP    87      71.115  18.961 -10.616  1.00 47.22      L21B O
ATOM   6660  N   TYR    88      71.928  19.942  -8.756  1.00 44.85      L21B N
ATOM   6661  CA  TYR    88      70.998  19.255  -7.863  1.00 43.21      L21B C
ATOM   6662  CB  TYR    88      71.738  18.231  -7.004  1.00 41.53      L21B C
ATOM   6663  CG  TYR    88      72.244  17.037  -7.777  1.00 41.60      L21B C
ATOM   6664  CD1 TYR    88      73.344  17.148  -8.624  1.00 42.48      L21B C
ATOM   6665  CE1 TYR    88      73.805  16.065  -9.349  1.00 41.25      L21B C
ATOM   6666  CD2 TYR    88      71.618  15.804  -7.674  1.00 39.77      L21B C
ATOM   6667  CE2 TYR    88      72.075  14.718  -8.396  1.00 41.08      L21B C
ATOM   6668  CZ  TYR    88      73.169  14.859  -9.228  1.00 40.42      L21B C
ATOM   6669  OH  TYR    88      73.645  13.785  -9.924  1.00 41.65      L21B O
ATOM   6670  C   TYR    88      70.236  20.199  -6.945  1.00 42.70      L21B C
ATOM   6671  O   TYR    88      70.823  21.011  -6.237  1.00 43.49      L21B O
ATOM   6672  N   TYR    89      68.916  20.082  -6.943  1.00 43.07      L21B N
ATOM   6673  CA  TYR    89      68.089  20.937  -6.095  1.00 41.89      L21B C
ATOM   6674  CB  TYR    89      67.103  21.754  -6.955  1.00 37.85      L21B C
ATOM   6675  CG  TYR    89      67.755  22.596  -8.038  1.00 31.95      L21B C
ATOM   6676  CD1 TYR    89      68.093  22.031  -9.269  1.00 29.53      L21B C
ATOM   6677  CE1 TYR    89      68.727  22.782 -10.259  1.00 28.25      L21B C
ATOM   6678  CD2 TYR    89      68.058  23.946  -7.820  1.00 30.26      L21B C
ATOM   6679  CE2 TYR    89      68.694  24.719  -8.808  1.00 27.69      L21B C
ATOM   6680  CZ  TYR    89      69.031  24.128 -10.026  1.00 27.67      L21B C
ATOM   6681  OH  TYR    89      69.699  24.847 -11.002  1.00 21.69      L21B O
ATOM   6682  C   TYR    89      67.315  20.074  -5.112  1.00 42.66      L21B C
ATOM   6683  O   TYR    89      66.771  19.034  -5.481  1.00 44.45      L21B O
ATOM   6684  N   CYS    90      67.265  20.495  -3.861  1.00 44.04      L21B N
ATOM   6685  CA  CYS    90      66.297  19.909  -2.964  1.00 49.10      L21B C
ATOM   6686  C   CYS    90      65.021  20.751  -2.907  1.00 50.09      L21B C
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ATOM   6687  O   CYS    90      65.039  21.980  -3.019  1.00 47.52      L21B O
ATOM   6688  CB  CYS    90      66.890  19.753  -1.568  1.00 53.04      L21B C
ATOM   6689  SG  CYS    90      67.391  21.301  -0.757  1.00 61.42      L21B S
ATOM   6690  N   ASN    91      63.905  20.068  -2.734  1.00 51.79      L21B N
ATOM   6691  CA  ASN    91      62.625  20.733  -2.620  1.00 53.65      L21B C
ATOM   6692  CB  ASN    91      61.774  20.348  -3.834  1.00 55.81      L21B C
ATOM   6693  CG  ASN    91      60.330  20.139  -3.495  1.00 59.21      L21B C
ATOM   6694  OD1 ASN    91      59.971  19.164  -2.820  1.00 62.36      L21B O
ATOM   6695  ND2 ASN    91      59.476  21.045  -3.967  1.00 60.75      L21B N
ATOM   6696  C   ASN    91      61.984  20.292  -1.307  1.00 52.94      L21B C
ATOM   6697  O   ASN    91      62.323  19.237  -0.777  1.00 52.18      L21B O
ATOM   6698  N   SER    92      61.085  21.108  -0.765  1.00 52.71      L21B N
ATOM   6699  CA  SER    92      60.248  20.669   0.347  1.00 51.08      L21B C
ATOM   6700  CB  SER    92      60.893  20.993   1.680  1.00 50.00      L21B C
ATOM   6701  OG  SER    92      59.963  20.766   2.715  1.00 46.75      L21B O
ATOM   6702  C   SER    92      58.869  21.284   0.347  1.00 51.25      L21B C
ATOM   6703  O   SER    92      58.706  22.470   0.053  1.00 50.09      L21B O
ATOM   6704  N   TYR    93      57.876  20.470   0.699  1.00 52.79      L21B N
ATOM   6705  CA  TYR    93      56.552  20.995   1.003  1.00 54.65      L21B C
ATOM   6706  CB  TYR    93      55.628  19.920   1.579  1.00 50.92      L21B C
ATOM   6707  CG  TYR    93      55.188  18.849   0.616  1.00 48.58      L21B C
ATOM   6708  CD1 TYR    93      55.740  18.762  -0.647  1.00 47.35      L21B C
ATOM   6709  CE1 TYR    93      55.373  17.746  -1.513  1.00 48.50      L21B C
ATOM   6710  CD2 TYR    93      54.243  17.891   0.995  1.00 46.57      L21B C
ATOM   6711  CE2 TYR    93      53.871  16.869   0.143  1.00 46.50      L21B C
ATOM   6712  CZ  TYR    93      54.443  16.798  -1.115  1.00 48.92      L21B C
ATOM   6713  OH  TYR    93      54.110  15.776  -1.982  1.00 50.22      L21B O
ATOM   6714  C   TYR    93      56.733  22.048   2.068  1.00 57.60      L21B C
ATOM   6715  O   TYR    93      57.557  21.896   2.959  1.00 57.59      L21B O
ATOM   6716  N   THR    94      55.951  23.111   1.991  1.00 62.04      L21B N
ATOM   6717  CA  THR    94      55.672  23.890   3.183  1.00 64.72      L21B C
ATOM   6718  CB  THR    94      56.079  25.350   2.997  1.00 62.42      L21B C
ATOM   6719  OG1 THR    94      55.926  26.035   4.243  1.00 64.29      L21B O
ATOM   6720  CG2 THR    94      55.212  26.019   1.955  1.00 59.60      L21B C
ATOM   6721  C   THR    94      54.178  23.803   3.512  1.00 65.94      L21B C
ATOM   6722  O   THR    94      53.428  23.081   2.854  1.00 69.06      L21B O
ATOM   6723  N   SER    95      53.754  24.524   4.538  1.00 65.95      L21B N
ATOM   6724  CA  SER    95      52.365  24.502   4.967  1.00 64.79      L21B C
ATOM   6725  CB  SER    95      52.192  25.481   6.120  1.00 66.94      L21B C
ATOM   6726  OG  SER    95      53.426  25.622   6.818  1.00 69.87      L21B O
ATOM   6727  C   SER    95      51.402  24.855   3.830  1.00 63.59      L21B C
ATOM   6728  O   SER    95      50.291  24.333   3.763  1.00 61.36      L21B O
ATOM   6729  N   THR    96      51.847  25.729   2.932  1.00 63.25      L21B N
ATOM   6730  CA  THR    96      51.009  26.245   1.860  1.00 63.14      L21B C
ATOM   6731  CB  THR    96      50.750  27.731   2.070  1.00 61.83      L21B C
ATOM   6732  OG1 THR    96      51.992  28.381   2.345  1.00 62.39      L21B O
ATOM   6733  CG2 THR    96      49.805  27.950   3.225  1.00 61.90      L21B C
ATOM   6734  C   THR    96      51.567  26.046   0.442  1.00 64.18      L21B C
ATOM   6735  O   THR    96      50.810  26.045  -0.525  1.00 65.09      L21B O
ATOM   6736  N   SER    97      52.878  25.879   0.301  1.00 64.48      L21B N
ATOM   6737  CA  SER    97      53.459  25.738  -1.034  1.00 64.51      L21B C
ATOM   6738  CB  SER    97      53.716  27.118  -1.627  1.00 63.56      L21B C
ATOM   6739  OG  SER    97      54.028  28.042  -0.603  1.00 61.46      L21B O
ATOM   6740  C   SER    97      54.739  24.910  -1.078  1.00 64.82      L21B C
ATOM   6741  O   SER    97      54.927  23.999  -0.270  1.00 66.42      L21B O
ATOM   6742  N   MET    98      55.605  25.217  -2.041  1.00 64.07      L21B N
ATOM   6743  CA  MET    98      56.908  24.557  -2.149  1.00 62.18      L21B C
ATOM   6744  CB  MET    98      57.126  24.032  -3.574  1.00 63.88      L21B C
ATOM   6745  CG  MET    98      55.993  23.171  -4.104  1.00 67.65      L21B C
ATOM   6746  SD  MET    98      55.659  21.710  -3.078  1.00 72.81      L21B S
ATOM   6747  CE  MET    98      56.438  20.416  -4.065  1.00 73.89      L21B C
ATOM   6748  C   MET    98      58.031  25.530  -1.802  1.00 59.11      L21B C
ATOM   6749  O   MET    98      57.833  26.736  -1.831  1.00 60.23      L21B O
ATOM   6750  N   VAL    99      59.201  25.001  -1.459  1.00 55.44      L21B N
ATOM   6751  CA  VAL    99      60.453  25.748  -1.572  1.00 51.98      L21B C
ATOM   6752  CB  VAL    99      60.916  26.368  -0.239  1.00 52.88      L21B C
ATOM   6753  CG1 VAL    99      60.298  27.734  -0.041  1.00 54.75      L21B C
ATOM   6754  CG2 VAL    99      60.547  25.444   0.903  1.00 55.69      L21B C
ATOM   6755  C   VAL    99      61.569  24.839  -2.041  1.00 49.13      L21B C
ATOM   6756  O   VAL    99      61.746  23.728  -1.539  1.00 47.82      L21B O
ATOM   6757  N   PHE   100      62.325  25.325  -3.013  1.00 46.97      L21B N
ATOM   6758  CA  PHE   100      63.500  24.628  -3.481  1.00 44.40      L21B C
ATOM   6759  CB  PHE   100      63.664  24.834  -4.981  1.00 42.82      L21B C
ATOM   6760  CG  PHE   100      62.723  24.021  -5.813  1.00 41.24      L21B C
ATOM   6761  CD1 PHE   100      61.481  24.518  -6.164  1.00 40.72      L21B C
ATOM   6762  CD2 PHE   100      63.107  22.781  -6.298  1.00 40.13      L21B C
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ATOM   6763  CE1 PHE   100      60.632  23.802  -6.996  1.00 38.77      L21B C
ATOM   6764  CE2 PHE   100      62.269  22.054  -7.129  1.00 39.31      L21B C
ATOM   6765  CZ  PHE   100      61.027  22.569  -7.482  1.00 40.01      L21B C
ATOM   6766  C   PHE   100      64.705  25.183  -2.749  1.00 43.67      L21B C
ATOM   6767  O   PHE   100      64.636  26.240  -2.138  1.00 42.08      L21B O
ATOM   6768  N   GLY   101      65.808  24.452  -2.794  1.00 44.87      L21B N
ATOM   6769  CA  GLY   101      67.050  24.990  -2.278  1.00 44.48      L21B C
ATOM   6770  C   GLY   101      67.713  25.786  -3.385  1.00 44.11      L21B C
ATOM   6771  O   GLY   101      67.280  25.735  -4.550  1.00 42.11      L21B O
ATOM   6772  N   GLY   102      68.764  26.517  -3.021  1.00 43.78      L21B N
ATOM   6773  CA  GLY   102      69.452  27.362  -3.973  1.00 42.09      L21B C
ATOM   6774  C   GLY   102      70.110  26.548  -5.064  1.00 41.93      L21B C
ATOM   6775  O   GLY   102      70.423  27.067  -6.133  1.00 39.38      L21B O
ATOM   6776  N   GLY   103      70.305  25.261  -4.792  1.00 43.18      L21B N
ATOM   6777  CA  GLY   103      70.929  24.376  -5.760  1.00 44.49      L21B C
ATOM   6778  C   GLY   103      72.381  24.131  -5.405  1.00 46.09      L21B C
ATOM   6779  O   GLY   103      73.019  24.962  -4.735  1.00 45.13      L21B O
ATOM   6780  N   THR   104      72.898  22.985  -5.839  1.00 47.33      L21B N
ATOM   6781  CA  THR   104      74.308  22.669  -5.691  1.00 49.35      L21B C
ATOM   6782  CB  THR   104      74.541  21.577  -4.622  1.00 46.64      L21B C
ATOM   6783  OG1 THR   104      74.064  22.036  -3.353  1.00 43.30      L21B O
ATOM   6784  CG2 THR   104      76.018  21.274  -4.491  1.00 43.32      L21B C
ATOM   6785  C   THR   104      74.850  22.180  -7.026  1.00 52.94      L21B C
ATOM   6786  O   THR   104      74.168  21.456  -7.752  1.00 53.31      L21B O
ATOM   6787  N   LYS   105      76.074  22.591  -7.342  1.00 55.91      L21B N
ATOM   6788  CA  LYS   105      76.745  22.179  -8.566  1.00 59.05      L21B C
ATOM   6789  CB  LYS   105      77.629  23.326  -9.068  1.00 61.76      L21B C
ATOM   6790  CG  LYS   105      77.714  23.459 -10.584  1.00 65.69      L21B C
ATOM   6791  CD  LYS   105      78.692  22.469 -11.214  1.00 68.17      L21B C
ATOM   6792  CE  LYS   105      79.046  22.903 -12.635  1.00 70.95      L21B C
ATOM   6793  NZ  LYS   105      80.160  22.133 -13.244  1.00 73.15      L21B N
ATOM   6794  C   LYS   105      77.602  20.953  -8.252  1.00 59.65      L21B C
ATOM   6795  O   LYS   105      78.315  20.946  -7.252  1.00 60.37      L21B O
ATOM   6796  N   LEU   106      77.534  19.922  -9.095  1.00 60.39      L21B N
ATOM   6797  CA  LEU   106      78.335  18.710  -8.898  1.00 61.91      L21B C
ATOM   6798  CB  LEU   106      77.446  17.462  -8.881  1.00 62.09      L21B C
ATOM   6799  CG  LEU   106      78.153  16.100  -8.764  1.00 60.56      L21B C
ATOM   6800  CD1 LEU   106      78.509  15.828  -7.314  1.00 60.55      L21B C
ATOM   6801  CD2 LEU   106      77.252  15.000  -9.277  1.00 60.14      L21B C
ATOM   6802  C   LEU   106      79.392  18.521  -9.977  1.00 64.04      L21B C
ATOM   6803  O   LEU   106      79.095  18.572 -11.170  1.00 63.86      L21B O
ATOM   6804  N   THR   107      80.628  18.276  -9.559  1.00 65.89      L21B N
ATOM   6805  CA  THR   107      81.684  18.024 -10.521  1.00 67.61      L21B C
ATOM   6806  CB  THR   107      82.803  19.050 -10.378  1.00 67.75      L21B C
ATOM   6807  OG1 THR   107      82.220  20.345 -10.228  1.00 68.92      L21B O
ATOM   6808  CG2 THR   107      83.692  19.050 -11.613  1.00 67.06      L21B C
ATOM   6809  C   THR   107      82.272  16.644 -10.316  1.00 68.99      L21B C
ATOM   6810  O   THR   107      82.644  16.280  -9.200  1.00 68.71      L21B O
ATOM   6811  N   VAL   108      82.349  15.879 -11.400  1.00 70.46      L21B N
ATOM   6812  CA  VAL   108      83.153  14.661 -11.425  1.00 72.21      L21B C
ATOM   6813  CB  VAL   108      82.594  13.619 -12.423  1.00 73.56      L21B C
ATOM   6814  CG1 VAL   108      83.486  12.397 -12.449  1.00 74.74      L21B C
ATOM   6815  CG2 VAL   108      81.188  13.226 -12.040  1.00 74.55      L21B C
ATOM   6816  C   VAL   108      84.553  15.047 -11.880  1.00 72.19      L21B C
ATOM   6817  O   VAL   108      84.814  15.159 -13.083  1.00 71.74      L21B O
ATOM   6818  N   LEU   109      85.443  15.247 -10.909  1.00 72.49      L21B N
ATOM   6819  CA  LEU   109      86.805  15.715 -11.160  1.00 72.84      L21B C
ATOM   6820  CB  LEU   109      87.622  15.621  -9.871  1.00 70.84      L21B C
ATOM   6821  CG  LEU   109      87.127  16.543  -8.755  1.00 70.37      L21B C
ATOM   6822  CD1 LEU   109      87.677  16.075  -7.441  1.00 69.58      L21B C
ATOM   6823  CD2 LEU   109      87.544  17.991  -9.031  1.00 69.90      L21B C
ATOM   6824  C   LEU   109      87.518  14.962 -12.280  1.00 73.49      L21B C
ATOM   6825  O   LEU   109      87.811  13.778 -12.154  1.00 74.15      L21B O
ATOM   6826  N   GLY   110      87.792  15.666 -13.374  1.00 73.86      L21B N
ATOM   6827  CA  GLY   110      88.496  15.074 -14.497  1.00 74.58      L21B C
ATOM   6828  C   GLY   110      89.870  15.695 -14.688  1.00 74.91      L21B C
ATOM   6829  O   GLY   110      90.465  15.602 -15.762  1.00 73.20      L21B O
ATOM   6830  N   GLN   111      90.362  16.335 -13.631  1.00 77.03      L21B N
ATOM   6831  CA  GLN   111      91.662  17.003 -13.625  1.00 80.05      L21B C
ATOM   6832  CB  GLN   111      91.706  18.108 -14.680  1.00 81.49      L21B C
ATOM   6833  CG  GLN   111      90.660  19.177 -14.485  1.00 82.71      L21B C
ATOM   6834  CD  GLN   111      91.022  20.463 -15.171  1.00 81.90      L21B C
ATOM   6835  OE1 GLN   111      91.918  21.183 -14.726  1.00 81.02      L21B O
ATOM   6836  NE2 GLN   111      90.332  20.764 -16.264  1.00 80.18      L21B N
ATOM   6837  C   GLN   111      91.887  17.599 -12.238  1.00 80.10      L21B C
ATOM   6838  O   GLN   111      91.014  17.524 -11.377  1.00 81.14      L21B O
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ATOM   6839  N   PRO  112      93.073  18.171 -11.990  1.00 78.66      L21B N
ATOM   6840  CD  PRO  112      94.285  18.245 -12.819  1.00 74.89      L21B C
ATOM   6841  CA  PRO  112      93.327  18.636 -10.625  1.00 76.87      L21B C
ATOM   6842  CB  PRO  112      94.845  18.786 -10.569  1.00 74.73      L21B C
ATOM   6843  CG  PRO  112      95.361  18.095 -11.803  1.00 73.97      L21B C
ATOM   6844  C   PRO  112      92.615  19.952 -10.358  1.00 75.57      L21B C
ATOM   6845  O   PRO  112      92.349  20.717 -11.285  1.00 76.36      L21B O
ATOM   6846  N   LYS  113      92.312  20.210  -9.091  1.00 72.84      L21B N
ATOM   6847  CA  LYS  113      91.651  21.448  -8.695  1.00 70.02      L21B C
ATOM   6848  CB  LYS  113      91.273  21.394  -7.210  1.00 71.19      L21B C
ATOM   6849  CG  LYS  113      90.170  20.412  -6.863  1.00 71.89      L21B C
ATOM   6850  CD  LYS  113      89.612  20.699  -5.473  1.00 72.26      L21B C
ATOM   6851  CE  LYS  113      88.519  19.711  -5.076  1.00 73.00      L21B C
ATOM   6852  NZ  LYS  113      88.060  19.907  -3.667  1.00 71.49      L21B N
ATOM   6853  C   LYS  113      92.570  22.649  -8.943  1.00 68.17      L21B C
ATOM   6854  O   LYS  113      93.767  22.585  -8.675  1.00 68.73      L21B O
ATOM   6855  N   ALA  114      92.018  23.747  -9.449  1.00 65.81      L21B N
ATOM   6856  CA  ALA  114      92.836  24.897  -9.802  1.00 62.76      L21B C
ATOM   6857  CB  ALA  114      92.869  25.050 -11.313  1.00 63.57      L21B C
ATOM   6858  C   ALA  114      92.280  26.158  -9.159  1.00 60.79      L21B C
ATOM   6859  O   ALA  114      91.119  26.489  -9.362  1.00 59.21      L21B O
ATOM   6860  N   ALA  115      93.098  26.861  -8.381  1.00 57.86      L21B N
ATOM   6861  CA  ALA  115      92.658  28.117  -7.771  1.00 55.74      L21B C
ATOM   6862  CB  ALA  115      93.698  28.630  -6.786  1.00 54.06      L21B C
ATOM   6863  C   ALA  115      92.432  29.140  -8.879  1.00 54.53      L21B C
ATOM   6864  O   ALA  115      93.166  29.165  -9.861  1.00 56.60      L21B O
ATOM   6865  N   PRO  116      91.409  29.997  -8.739  1.00 52.44      L21B N
ATOM   6866  CD  PRO  116      90.642  30.294  -7.513  1.00 53.94      L21B C
ATOM   6867  CA  PRO  116      91.089  30.936  -9.816  1.00 50.36      L21B C
ATOM   6868  CB  PRO  116      89.689  31.412  -9.447  1.00 52.55      L21B C
ATOM   6869  CG  PRO  116      89.720  31.425  -7.930  1.00 53.47      L21B C
ATOM   6870  C   PRO  116      92.103  32.093  -9.851  1.00 48.65      L21B C
ATOM   6871  O   PRO  116      92.627  32.498  -8.803  1.00 47.48      L21B O
ATOM   6872  N   SER  117      92.375  32.627 -11.043  1.00 45.92      L21B N
ATOM   6873  CA  SER  117      93.083  33.906 -11.154  1.00 43.14      L21B C
ATOM   6874  CB  SER  117      93.978  33.918 -12.384  1.00 42.23      L21B C
ATOM   6875  OG  SER  117      94.987  32.945 -12.258  1.00 47.57      L21B O
ATOM   6876  C   SER  117      92.109  35.073 -11.241  1.00 41.00      L21B C
ATOM   6877  O   SER  117      91.326  35.177 -12.181  1.00 40.97      L21B O
ATOM   6878  N   VAL  118      92.163  35.955 -10.259  1.00 37.34      L21B N
ATOM   6879  CA  VAL  118      91.337  37.143 -10.274  1.00 36.92      L21B C
ATOM   6880  CB  VAL  118      90.715  37.387  -8.878  1.00 36.85      L21B C
ATOM   6881  CG1 VAL  118      89.895  38.663  -8.867  1.00 34.92      L21B C
ATOM   6882  CG2 VAL  118      89.862  36.213  -8.495  1.00 38.22      L21B C
ATOM   6883  C   VAL  118      92.215  38.332 -10.635  1.00 37.10      L21B C
ATOM   6884  O   VAL  118      93.224  38.580  -9.971  1.00 37.46      L21B O
ATOM   6885  N   THR  119      91.846  39.075 -11.673  1.00 36.80      L21B N
ATOM   6886  CA  THR  119      92.433  40.405 -11.836  1.00 38.29      L21B C
ATOM   6887  CB  THR  119      93.380  40.467 -13.072  1.00 40.02      L21B C
ATOM   6888  OG1 THR  119      92.629  40.766 -14.249  1.00 42.31      L21B O
ATOM   6889  CG2 THR  119      94.098  39.115 -13.267  1.00 41.99      L21B C
ATOM   6890  C   THR  119      91.335  41.463 -11.949  1.00 36.59      L21B C
ATOM   6891  O   THR  119      90.328  41.254 -12.631  1.00 35.96      L21B O
ATOM   6892  N   LEU  120      91.511  42.588 -11.262  1.00 34.86      L21B N
ATOM   6893  CA  LEU  120      90.407  43.542 -11.101  1.00 35.65      L21B C
ATOM   6894  CB  LEU  120      89.973  43.569  -9.627  1.00 32.38      L21B C
ATOM   6895  CG  LEU  120      89.176  44.819  -9.256  1.00 32.01      L21B C
ATOM   6896  CD1 LEU  120      87.826  44.808  -9.988  1.00 29.34      L21B C
ATOM   6897  CD2 LEU  120      89.019  44.892  -7.740  1.00 30.16      L21B C
ATOM   6898  C   LEU  120      90.716  44.978 -11.595  1.00 34.81      L21B C
ATOM   6899  O   LEU  120      91.618  45.628 -11.081  1.00 35.69      L21B O
ATOM   6900  N   PHE  121      89.959  45.464 -12.580  1.00 35.05      L21B N
ATOM   6901  CA  PHE  121      90.240  46.752 -13.227  1.00 37.49      L21B C
ATOM   6902  CB  PHE  121      90.054  46.649 -14.733  1.00 37.01      L21B C
ATOM   6903  CG  PHE  121      91.133  45.888 -15.418  1.00 39.94      L21B C
ATOM   6904  CD1 PHE  121      90.885  44.631 -15.950  1.00 41.17      L21B C
ATOM   6905  CD2 PHE  121      92.403  46.431 -15.546  1.00 40.30      L21B C
ATOM   6906  CE1 PHE  121      91.882  43.926 -16.601  1.00 41.70      L21B C
ATOM   6907  CE2 PHE  121      93.407  45.736 -16.193  1.00 41.18      L21B C
ATOM   6908  CZ  PHE  121      93.147  44.477 -16.725  1.00 42.12      L21B C
ATOM   6909  C   PHE  121      89.327  47.840 -12.727  1.00 37.59      L21B C
ATOM   6910  O   PHE  121      88.118  47.679 -12.717  1.00 38.34      L21B O
ATOM   6911  N   PRO  122      89.895  48.980 -12.320  1.00 33.96      L21B N
ATOM   6912  CD  PRO  122      91.343  49.211 -12.217  1.00 33.93      L21B C
ATOM   6913  CA  PRO  122      89.118  50.140 -11.849  1.00 38.29      L21B C
ATOM   6914  CB  PRO  122      90.165  51.012 -11.187  1.00 35.32      L21B C
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ATOM   6915  CG  PRO   122      91.417  50.673 -11.910  1.00 33.59      L21B C
ATOM   6916  C   PRO   122      88.425  50.864 -12.992  1.00 39.54      L21B C
ATOM   6917  O   PRO   122      88.852  50.762 -14.136  1.00 39.03      L21B O
ATOM   6918  N   PRO   123      87.360  51.623 -12.697  1.00 38.00      L21B N
ATOM   6919  CD  PRO   123      86.853  52.011 -11.371  1.00 42.45      L21B C
ATOM   6920  CA  PRO   123      86.729  52.399 -13.778  1.00 40.80      L21B C
ATOM   6921  CB  PRO   123      85.750  53.326 -13.057  1.00 41.73      L21B C
ATOM   6922  CG  PRO   123      85.552  52.698 -11.698  1.00 44.67      L21B C
ATOM   6923  C   PRO   123      87.813  53.192 -14.490  1.00 42.29      L21B C
ATOM   6924  O   PRO   123      88.806  53.588 -13.871  1.00 40.52      L21B O
ATOM   6925  N   SER   124      87.644  53.410 -15.788  1.00 44.28      L21B N
ATOM   6926  CA  SER   124      88.611  54.216 -16.506  1.00 47.54      L21B C
ATOM   6927  CB  SER   124      88.663  53.823 -17.988  1.00 47.47      L21B C
ATOM   6928  OG  SER   124      87.506  54.258 -18.684  1.00 53.94      L21B O
ATOM   6929  C   SER   124      88.267  55.697 -16.348  1.00 49.32      L21B C
ATOM   6930  O   SER   124      87.158  56.071 -15.948  1.00 49.97      L21B O
ATOM   6931  N   SER   125      89.255  56.527 -16.636  1.00 50.71      L21B N
ATOM   6932  CA  SER   125      89.127  57.964 -16.549  1.00 51.79      L21B C
ATOM   6933  CB  SER   125      90.474  58.586 -16.895  1.00 54.13      L21B C
ATOM   6934  OG  SER   125      91.447  57.553 -17.108  1.00 58.67      L21B O
ATOM   6935  C   SER   125      88.062  58.413 -17.530  1.00 52.33      L21B C
ATOM   6936  O   SER   125      87.218  59.240 -17.197  1.00 53.85      L21B O
ATOM   6937  N   GLU   126      88.093  57.852 -18.735  1.00 52.90      L21B N
ATOM   6938  CA  GLU   126      87.157  58.247 -19.780  1.00 53.19      L21B C
ATOM   6939  CB  GLU   126      87.458  57.509 -21.082  1.00 57.40      L21B C
ATOM   6940  CG  GLU   126      88.843  57.708 -21.675  1.00 62.62      L21B C
ATOM   6941  CD  GLU   126      88.903  57.183 -23.100  1.00 66.23      L21B C
ATOM   6942  OE1 GLU   126      88.015  57.561 -23.904  1.00 68.20      L21B O
ATOM   6943  OE2 GLU   126      89.822  56.392 -23.414  1.00 68.73      L21B O
ATOM   6944  C   GLU   126      85.734  57.919 -19.373  1.00 51.91      L21B C
ATOM   6945  O   GLU   126      84.833  58.750 -19.467  1.00 50.90      L21B O
ATOM   6946  N   GLU   127      85.528  56.687 -18.928  1.00 50.99      L21B N
ATOM   6947  CA  GLU   127      84.197  56.264 -18.523  1.00 49.49      L21B C
ATOM   6948  CB  GLU   127      84.213  54.811 -18.046  1.00 48.63      L21B C
ATOM   6949  CG  GLU   127      82.828  54.274 -17.767  1.00 45.52      L21B C
ATOM   6950  CD  GLU   127      82.845  53.032 -16.907  1.00 44.98      L21B C
ATOM   6951  OE1 GLU   127      81.780  52.691 -16.353  1.00 42.76      L21B O
ATOM   6952  OE2 GLU   127      83.920  52.399 -16.794  1.00 43.35      L21B O
ATOM   6953  C   GLU   127      83.656  57.171 -17.422  1.00 48.57      L21B C
ATOM   6954  O   GLU   127      82.534  57.660 -17.516  1.00 49.29      L21B O
ATOM   6955  N   LEU   128      84.447  57.408 -16.382  1.00 47.26      L21B N
ATOM   6956  CA  LEU   128      84.003  58.319 -15.337  1.00 46.49      L21B C
ATOM   6957  CB  LEU   128      85.145  58.634 -14.377  1.00 44.93      L21B C
ATOM   6958  CG  LEU   128      85.440  57.550 -13.341  1.00 44.39      L21B C
ATOM   6959  CD1 LEU   128      86.492  58.057 -12.377  1.00 42.24      L21B C
ATOM   6960  CD2 LEU   128      84.160  57.177 -12.588  1.00 43.91      L21B C
ATOM   6961  C   LEU   128      83.461  59.609 -15.948  1.00 45.72      L21B C
ATOM   6962  O   LEU   128      82.445  60.143 -15.502  1.00 44.90      L21B O
ATOM   6963  N   GLN   129      84.124  60.094 -16.991  1.00 44.74      L21B N
ATOM   6964  CA  GLN   129      83.690  61.326 -17.643  1.00 43.60      L21B C
ATOM   6965  CB  GLN   129      84.730  61.764 -18.673  1.00 44.32      L21B C
ATOM   6966  CG  GLN   129      84.757  63.251 -18.915  1.00 44.45      L21B C
ATOM   6967  CD  GLN   129      84.894  64.018 -17.632  1.00 44.59      L21B C
ATOM   6968  OE1 GLN   129      83.960  64.705 -17.184  1.00 46.48      L21B O
ATOM   6969  NE2 GLN   129      86.065  63.903 -17.014  1.00 43.44      L21B N
ATOM   6970  C   GLN   129      82.335  61.142 -18.318  1.00 43.46      L21B C
ATOM   6971  O   GLN   129      81.613  62.111 -18.567  1.00 42.25      L21B O
ATOM   6972  N   ALA   130      81.999  59.890 -18.612  1.00 43.83      L21B N
ATOM   6973  CA  ALA   130      80.723  59.562 -19.236  1.00 44.55      L21B C
ATOM   6974  CB  ALA   130      80.842  58.252 -19.998  1.00 40.95      L21B C
ATOM   6975  C   ALA   130      79.613  59.472 -18.190  1.00 45.51      L21B C
ATOM   6976  O   ALA   130      78.444  59.242 -18.516  1.00 45.69      L21B O
ATOM   6977  N   ASN   131      79.998  59.658 -16.932  1.00 46.19      L21B N
ATOM   6978  CA  ASN   131      79.057  59.676 -15.814  1.00 46.40      L21B C
ATOM   6979  CB  ASN   131      77.856  60.560 -16.116  1.00 46.67      L21B C
ATOM   6980  CG  ASN   131      77.079  60.903 -14.866  1.00 48.91      L21B C
ATOM   6981  OD1 ASN   131      77.634  60.893 -13.764  1.00 47.77      L21B O
ATOM   6982  ND2 ASN   131      75.788  61.203 -15.021  1.00 51.00      L21B N
ATOM   6983  C   ASN   131      78.572  58.279 -15.467  1.00 46.14      L21B C
ATOM   6984  O   ASN   131      77.431  58.080 -15.028  1.00 45.59      L21B O
ATOM   6985  N   LYS   132      79.456  57.313 -15.687  1.00 44.66      L21B N
ATOM   6986  CA  LYS   132      79.241  55.945 -15.265  1.00 42.08      L21B C
ATOM   6987  CB  LYS   132      78.787  55.099 -16.450  1.00 40.90      L21B C
ATOM   6988  CG  LYS   132      77.627  55.703 -17.217  1.00 43.12      L21B C
ATOM   6989  CD  LYS   132      76.291  55.233 -16.679  1.00 47.00      L21B C
ATOM   6990  CE  LYS   132      75.556  54.388 -17.718  1.00 51.53      L21B C
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ATOM   6991  NZ  LYS   132      75.127  55.214 -18.892  1.00 53.24      L21B N
ATOM   6992  C   LYS   132      80.567  55.436 -14.727  1.00 40.86      L21B C
ATOM   6993  O   LYS   132      81.625  55.974 -15.036  1.00 39.43      L21B O
ATOM   6994  N   ALA   133      80.515  54.410 -13.898  1.00 40.80      L21B N
ATOM   6995  CA  ALA   133      81.742  53.771 -13.469  1.00 40.55      L21B C
ATOM   6996  CB  ALA   133      82.155  54.284 -12.094  1.00 41.61      L21B C
ATOM   6997  C   ALA   133      81.503  52.277 -13.436  1.00 39.52      L21B C
ATOM   6998  O   ALA   133      80.407  51.827 -13.094  1.00 40.03      L21B O
ATOM   6999  N   THR   134      82.517  51.511 -13.816  1.00 37.08      L21B N
ATOM   7000  CA  THR   134      82.387  50.070 -13.826  1.00 35.24      L21B C
ATOM   7001  CB  THR   134      82.402  49.568 -15.261  1.00 34.18      L21B C
ATOM   7002  OG1 THR   134      81.076  50.373 -15.860  1.00 31.16      L21B O
ATOM   7003  CG2 THR   134      81.630  48.113 -15.229  1.00 34.12      L21B C
ATOM   7004  C   THR   134      83.660  49.413 -13.288  1.00 35.11      L21B C
ATOM   7005  O   THR   134      84.756  49.639 -13.809  1.00 34.86      L21B O
ATOM   7006  N   LEU   135      83.527  48.604 -12.240  1.00 35.01      L21B N
ATOM   7007  CA  LEU   135      84.659  47.785 -11.811  1.00 35.41      L21B C
ATOM   7008  CB  LEU   135      84.648  47.559 -10.299  1.00 35.87      L21B C
ATOM   7009  CG  LEU   135      84.344  48.751  -9.393  1.00 35.82      L21B C
ATOM   7010  CD1 LEU   135      84.432  48.303  -7.946  1.00 37.53      L21B C
ATOM   7011  CD2 LEU   135      85.307  49.859  -9.652  1.00 35.35      L21B C
ATOM   7012  C   LEU   135      84.568  46.451 -12.520  1.00 33.80      L21B C
ATOM   7013  O   LEU   135      83.529  45.820 -12.520  1.00 35.43      L21B O
ATOM   7014  N   VAL   136      85.659  46.033 -13.136  1.00 33.39      L21B N
ATOM   7015  CA  VAL   136      85.658  44.817 -13.927  1.00 33.35      L21B C
ATOM   7016  CB  VAL   136      86.196  45.094 -15.343  1.00 31.20      L21B C
ATOM   7017  CG1 VAL   136      86.377  43.785 -16.117  1.00 30.64      L21B C
ATOM   7018  CG2 VAL   136      85.222  46.009 -16.080  1.00 28.09      L21B C
ATOM   7019  C   VAL   136      86.492  43.727 -13.262  1.00 33.93      L21B C
ATOM   7020  O   VAL   136      87.718  43.823 -13.186  1.00 33.89      L21B O
ATOM   7021  N   CYS   137      85.815  42.692 -12.771  1.00 34.36      L21B N
ATOM   7022  CA  CYS   137      86.505  41.597 -12.113  1.00 36.21      L21B C
ATOM   7023  C   CYS   137      86.499  40.375 -13.001  1.00 35.27      L21B C
ATOM   7024  O   CYS   137      85.460  39.765 -13.199  1.00 34.76      L21B O
ATOM   7025  CB  CYS   137      85.848  41.264 -10.772  1.00 38.22      L21B C
ATOM   7026  SG  CYS   137      86.846  40.139  -9.739  1.00 40.11      L21B S
ATOM   7027  N   LEU   138      87.670  40.029 -13.531  1.00 35.98      L21B N
ATOM   7028  CA  LEU   138      87.815  38.912 -14.454  1.00 36.04      L21B C
ATOM   7029  CB  LEU   138      88.747  39.322 -15.594  1.00 37.69      L21B C
ATOM   7030  CG  LEU   138      88.403  40.638 -16.312  1.00 39.81      L21B C
ATOM   7031  CD1 LEU   138      89.536  41.044 -17.241  1.00 40.38      L21B C
ATOM   7032  CD2 LEU   138      87.129  40.481 -17.103  1.00 39.74      L21B C
ATOM   7033  C   LEU   138      88.367  37.670 -13.725  1.00 39.64      L21B C
ATOM   7034  O   LEU   138      89.270  37.784 -12.889  1.00 41.42      L21B O
ATOM   7035  N   ILE   139      87.824  36.490 -14.030  1.00 38.47      L21B N
ATOM   7036  CA  ILE   139      88.251  35.268 -13.351  1.00 38.43      L21B C
ATOM   7037  CB  ILE   139      87.150  34.756 -12.429  1.00 37.11      L21B C
ATOM   7038  CG2 ILE   139      87.707  33.697 -11.485  1.00 38.23      L21B C
ATOM   7039  CG1 ILE   139      86.581  35.914 -11.627  1.00 36.92      L21B C
ATOM   7040  CD1 ILE   139      85.608  35.472 -10.583  1.00 38.37      L21B C
ATOM   7041  C   ILE   139      88.618  34.150 -14.335  1.00 38.50      L21B C
ATOM   7042  O   ILE   139      87.834  33.819 -15.225  1.00 36.72      L21B O
ATOM   7043  N   SER   140      89.803  33.564 -14.172  1.00 39.81      L21B N
ATOM   7044  CA  SER   140      90.312  32.604 -15.156  1.00 42.70      L21B C
ATOM   7045  CB  SER   140      91.266  33.292 -16.145  1.00 42.39      L21B C
ATOM   7046  OG  SER   140      92.065  34.270 -15.498  1.00 43.52      L21B O
ATOM   7047  C   SER   140      91.020  31.398 -14.564  1.00 43.55      L21B C
ATOM   7048  O   SER   140      91.514  31.441 -13.432  1.00 43.83      L21B O
ATOM   7049  N   ASP   141      91.055  30.320 -15.344  1.00 44.57      L21B N
ATOM   7050  CA  ASP   141      91.910  29.173 -15.060  1.00 46.96      L21B C
ATOM   7051  CB  ASP   141      93.388  29.589 -15.151  1.00 49.43      L21B C
ATOM   7052  CG  ASP   141      93.753  30.153 -16.519  1.00 52.67      L21B C
ATOM   7053  OD1 ASP   141      93.943  31.388 -16.623  1.00 55.09      L21B O
ATOM   7054  OD2 ASP   141      93.843  29.362 -17.492  1.00 54.16      L21B O
ATOM   7055  C   ASP   141      91.626  28.547 -13.697  1.00 47.35      L21B C
ATOM   7056  O   ASP   141      92.510  28.431 -12.853  1.00 47.00      L21B O
ATOM   7057  N   PHE   142      90.389  28.134 -13.481  1.00 49.13      L21B N
ATOM   7058  CA  PHE   142      90.066  27.424 -12.260  1.00 51.26      L21B C
ATOM   7059  CB  PHE   142      89.306  28.349 -11.315  1.00 50.93      L21B C
ATOM   7060  CG  PHE   142      88.015  28.882 -11.885  1.00 52.14      L21B C
ATOM   7061  CD1 PHE   142      86.839  28.146 -11.790  1.00 51.98      L21B C
ATOM   7062  CD2 PHE   142      87.960  30.142 -12.457  1.00 51.72      L21B C
ATOM   7063  CE1 PHE   142      85.641  28.661 -12.245  1.00 49.40      L21B C
ATOM   7064  CE2 PHE   142      86.767  30.658 -12.913  1.00 49.64      L21B C
ATOM   7065  CZ  PHE   142      85.608  29.917 -12.804  1.00 50.17      L21B C
ATOM   7066  C   PHE   142      89.255  26.166 -12.561  1.00 51.68      L21B C
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ATOM   7067  O   PHE  142      88.654  26.058 -13.626  1.00 51.50      L21B O
ATOM   7068  N   TYR  143      89.267  25.218 -11.625  1.00 52.56      L21B N
ATOM   7069  CA  TYR  143      88.530  23.962 -11.760  1.00 54.58      L21B C
ATOM   7070  CB  TYR  143      89.282  23.010 -12.699  1.00 56.96      L21B C
ATOM   7071  CG  TYR  143      88.523  21.763 -13.117  1.00 59.81      L21B C
ATOM   7072  CD1 TYR  143      88.524  20.618 -12.324  1.00 62.38      L21B C
ATOM   7073  CE1 TYR  143      87.865  19.450 -12.731  1.00 63.84      L21B C
ATOM   7074  CD2 TYR  143      87.844  21.717 -14.327  1.00 60.41      L21B C
ATOM   7075  CE2 TYR  143      87.185  20.563 -14.744  1.00 63.55      L21B C
ATOM   7076  CZ  TYR  143      87.198  19.431 -13.943  1.00 64.86      L21B C
ATOM   7077  OH  TYR  143      86.534  18.291 -14.356  1.00 65.40      L21B O
ATOM   7078  C   TYR  143      88.382  23.328 -10.378  1.00 53.01      L21B C
ATOM   7079  O   TYR  143      89.303  23.365  -9.569  1.00 53.42      L21B O
ATOM   7080  N   PRO  144      87.206  22.759 -10.079  1.00 50.74      L21B N
ATOM   7081  CD  PRO  144      86.972  22.124  -8.767  1.00 49.09      L21B C
ATOM   7082  CA  PRO  144      86.003  22.734 -10.923  1.00 49.32      L21B C
ATOM   7083  CB  PRO  144      85.062  21.804 -10.164  1.00 47.39      L21B C
ATOM   7084  CG  PRO  144      85.478  21.930  -8.737  1.00 46.20      L21B C
ATOM   7085  C   PRO  144      85.382  24.130 -11.153  1.00 47.94      L21B C
ATOM   7086  O   PRO  144      85.702  25.088 -10.440  1.00 46.61      L21B O
ATOM   7087  N   GLY  145      84.501  24.228 -12.152  1.00 46.23      L21B N
ATOM   7088  CA  GLY  145      83.958  25.512 -12.578  1.00 42.95      L21B C
ATOM   7089  C   GLY  145      82.812  26.070 -11.742  1.00 41.27      L21B C
ATOM   7090  O   GLY  145      81.719  26.310 -12.250  1.00 40.44      L21B O
ATOM   7091  N   ALA  146      83.062  26.275 -10.453  1.00 39.59      L21B N
ATOM   7092  CA  ALA  146      82.094  26.915  -9.571  1.00 38.70      L21B C
ATOM   7093  CB  ALA  146      81.366  25.871  -8.763  1.00 31.97      L21B C
ATOM   7094  C   ALA  146      82.779  27.922  -8.637  1.00 39.60      L21B C
ATOM   7095  O   ALA  146      83.631  27.551  -7.833  1.00 41.88      L21B O
ATOM   7096  N   VAL  147      82.417  29.197  -8.755  1.00 38.94      L21B N
ATOM   7097  CA  VAL  147      82.985  30.227  -7.894  1.00 39.15      L21B C
ATOM   7098  CB  VAL  147      83.970  31.166  -8.630  1.00 40.72      L21B C
ATOM   7099  CG1 VAL  147      85.144  30.392  -9.186  1.00 40.44      L21B C
ATOM   7100  CG2 VAL  147      83.229  31.921  -9.733  1.00 40.31      L21B C
ATOM   7101  C   VAL  147      81.844  31.100  -7.439  1.00 38.68      L21B C
ATOM   7102  O   VAL  147      80.803  31.151  -8.080  1.00 41.11      L21B O
ATOM   7103  N   THR  148      82.033  31.797  -6.335  1.00 36.37      L21B N
ATOM   7104  CA  THR  148      81.051  32.778  -5.955  1.00 36.67      L21B C
ATOM   7105  CB  THR  148      80.323  32.336  -4.659  1.00 35.50      L21B C
ATOM   7106  OG1 THR  148      80.850  33.034  -3.521  1.00 34.72      L21B O
ATOM   7107  CG2 THR  148      80.505  30.836  -4.467  1.00 37.45      L21B C
ATOM   7108  C   THR  148      81.807  34.102  -5.783  1.00 37.12      L21B C
ATOM   7109  O   THR  148      82.966  34.112  -5.342  1.00 36.84      L21B O
ATOM   7110  N   VAL  149      81.178  35.207  -6.169  1.00 33.24      L21B N
ATOM   7111  CA  VAL  149      81.858  36.472  -6.104  1.00 33.40      L21B C
ATOM   7112  CB  VAL  149      81.907  37.135  -7.485  1.00 31.91      L21B C
ATOM   7113  CG1 VAL  149      82.473  38.539  -7.369  1.00 33.40      L21B C
ATOM   7114  CG2 VAL  149      82.767  36.321  -8.404  1.00 30.80      L21B C
ATOM   7115  C   VAL  149      81.145  37.370  -5.121  1.00 34.21      L21B C
ATOM   7116  O   VAL  149      79.944  37.244  -4.921  1.00 34.91      L21B O
ATOM   7117  N   ALA  150      81.886  38.264  -4.487  1.00 33.86      L21B N
ATOM   7118  CA  ALA  150      81.280  39.225  -3.585  1.00 35.73      L21B C
ATOM   7119  CB  ALA  150      81.210  38.656  -2.188  1.00 33.79      L21B C
ATOM   7120  C   ALA  150      82.112  40.501  -3.597  1.00 39.22      L21B C
ATOM   7121  O   ALA  150      83.349  40.462  -3.711  1.00 40.31      L21B O
ATOM   7122  N   TRP  151      81.430  41.633  -3.487  1.00 40.11      L21B N
ATOM   7123  CA  TRP  151      82.080  42.922  -3.632  1.00 41.54      L21B C
ATOM   7124  CB  TRP  151      81.393  43.745  -4.729  1.00 38.77      L21B C
ATOM   7125  CG  TRP  151      81.582  43.201  -6.112  1.00 36.67      L21B C
ATOM   7126  CD2 TRP  151      82.611  43.564  -7.048  1.00 36.37      L21B C
ATOM   7127  CE2 TRP  151      82.391  42.819  -8.219  1.00 34.19      L21B C
ATOM   7128  CE3 TRP  151      83.694  44.448  -7.005  1.00 35.66      L21B C
ATOM   7129  CD1 TRP  151      80.609  42.277  -6.737  1.00 35.59      L21B C
ATOM   7130  NE1 TRP  151      81.286  42.039  -8.006  1.00 35.04      L21B N
ATOM   7131  CZ2 TRP  151      83.205  42.928  -9.330  1.00 34.35      L21B C
ATOM   7132  CZ3 TRP  151      84.503  44.553  -8.107  1.00 35.09      L21B C
ATOM   7133  CH2 TRP  151      84.257  43.798  -9.256  1.00 35.84      L21B C
ATOM   7134  C   TRP  151      81.997  43.664  -2.316  1.00 44.09      L21B C
ATOM   7135  O   TRP  151      80.964  43.656  -1.644  1.00 45.40      L21B O
ATOM   7136  N   LYS  152      83.092  44.308  -1.946  1.00 46.45      L21B N
ATOM   7137  CA  LYS  152      83.072  45.197  -0.811  1.00 48.60      L21B C
ATOM   7138  CB  LYS  152      84.092  44.751   0.225  1.00 48.03      L21B C
ATOM   7139  CG  LYS  152      83.871  43.340   0.723  1.00 50.00      L21B C
ATOM   7140  CD  LYS  152      83.032  43.305   2.003  1.00 51.32      L21B C
ATOM   7141  CE  LYS  152      82.767  41.869   2.424  1.00 52.22      L21B C
ATOM   7142  NZ  LYS  152      82.022  41.119   1.373  1.00 55.40      L21B N
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ATOM	7143	C	LYS	152	83.390	46.601	-1.281	1.00	50.99	L21B C
ATOM	7144	O	LYS	152	84.217	46.799	-2.181	1.00	51.75	L21B O
ATOM	7145	N	ALA	153	82.701	47.565	-0.680	1.00	52.20	L21B N
ATOM	7146	CA	ALA	153	83.076	48.961	-0.762	1.00	53.01	L21B C
ATOM	7147	CB	ALA	153	81.854	49.802	-1.069	1.00	53.76	L21B C
ATOM	7148	C	ALA	153	83.664	49.342	0.596	1.00	53.38	L21B C
ATOM	7149	O	ALA	153	82.970	49.338	1.605	1.00	53.17	L21B O
ATOM	7150	N	ASP	154	84.950	49.671	0.610	1.00	55.41	L21B N
ATOM	7151	CA	ASP	154	85.732	49.644	1.840	1.00	56.87	L21B C
ATOM	7152	CB	ASP	154	85.183	50.652	2.856	1.00	60.15	L21B C
ATOM	7153	CG	ASP	154	85.664	52.060	2.579	1.00	63.17	L21B C
ATOM	7154	OD1	ASP	154	86.880	52.204	2.319	1.00	66.19	L21B O
ATOM	7155	OD2	ASP	154	84.842	53.010	2.609	1.00	63.31	L21B O
ATOM	7156	C	ASP	154	85.659	48.238	2.398	1.00	56.82	L21B C
ATOM	7157	O	ASP	154	86.161	47.297	1.790	1.00	54.87	L21B O
ATOM	7158	N	SER	155	85.010	48.090	3.545	1.00	57.77	L21B N
ATOM	7159	CA	SER	155	84.814	46.771	4.112	1.00	58.15	L21B C
ATOM	7160	CB	SER	155	85.331	46.744	5.542	1.00	57.73	L21B C
ATOM	7161	OG	SER	155	86.728	46.519	5.538	1.00	60.83	L21B O
ATOM	7162	C	SER	155	83.367	46.327	4.068	1.00	58.08	L21B C
ATOM	7163	O	SER	155	83.069	45.151	4.217	1.00	59.53	L21B O
ATOM	7164	N	SER	156	82.461	47.267	3.858	1.00	58.33	L21B N
ATOM	7165	CA	SER	156	81.057	46.918	3.777	1.00	59.90	L21B C
ATOM	7166	CB	SER	156	80.201	48.162	3.965	1.00	60.53	L21B C
ATOM	7167	OG	SER	156	80.336	48.660	5.283	1.00	59.84	L21B O
ATOM	7168	C	SER	156	80.733	46.260	2.445	1.00	60.68	L21B C
ATOM	7169	O	SER	156	81.328	46.583	1.422	1.00	61.89	L21B O
ATOM	7170	N	PRO	157	79.792	45.304	2.453	1.00	60.88	L21B N
ATOM	7171	CD	PRO	157	79.262	44.713	3.690	1.00	58.20	L21B C
ATOM	7172	CA	PRO	157	79.348	44.547	1.278	1.00	60.26	L21B C
ATOM	7173	CB	PRO	157	78.687	43.304	1.886	1.00	58.77	L21B C
ATOM	7174	CG	PRO	157	79.155	43.277	3.324	1.00	56.44	L21B C
ATOM	7175	C	PRO	157	78.402	45.303	0.334	1.00	59.58	L21B C
ATOM	7176	O	PRO	157	77.472	45.984	0.763	1.00	57.31	L21B O
ATOM	7177	N	VAL	158	78.657	45.150	-0.959	1.00	59.32	L21B N
ATOM	7178	CA	VAL	158	77.915	45.844	-2.002	1.00	59.47	L21B C
ATOM	7179	CB	VAL	158	78.700	45.791	-3.331	1.00	57.95	L21B C
ATOM	7180	CG1	VAL	158	77.839	46.290	-4.469	1.00	54.31	L21B C
ATOM	7181	CG2	VAL	158	79.975	46.617	-3.213	1.00	53.69	L21B C
ATOM	7182	C	VAL	158	76.536	45.218	-2.204	1.00	60.73	L21B C
ATOM	7183	O	VAL	158	76.414	44.001	-2.333	1.00	61.06	L21B O
ATOM	7184	N	LYS	159	75.505	46.061	-2.244	1.00	62.19	L21B N
ATOM	7185	CA	LYS	159	74.113	45.602	-2.315	1.00	61.60	L21B C
ATOM	7186	CB	LYS	159	73.236	46.399	-1.341	1.00	63.72	L21B C
ATOM	7187	CG	LYS	159	73.455	46.090	0.126	1.00	69.06	L21B C
ATOM	7188	CD	LYS	159	72.644	47.037	0.998	1.00	75.91	L21B C
ATOM	7189	CE	LYS	159	73.000	46.892	2.478	1.00	82.57	L21B C
ATOM	7190	NZ	LYS	159	72.140	47.750	3.366	1.00	87.68	L21B N
ATOM	7191	C	LYS	159	73.462	45.674	-3.699	1.00	58.86	L21B C
ATOM	7192	O	LYS	159	72.303	45.300	-3.847	1.00	57.81	L21B O
ATOM	7193	N	ALA	160	74.186	46.165	-4.701	1.00	55.78	L21B N
ATOM	7194	CA	ALA	160	73.587	46.415	-6.008	1.00	52.62	L21B C
ATOM	7195	CB	ALA	160	72.592	47.559	-5.900	1.00	51.57	L21B C
ATOM	7196	C	ALA	160	74.622	46.724	-7.090	1.00	51.55	L21B C
ATOM	7197	O	ALA	160	75.718	47.210	-6.803	1.00	52.32	L21B O
ATOM	7198	N	GLY	161	74.269	46.441	-8.339	1.00	49.63	L21B N
ATOM	7199	CA	GLY	161	75.115	46.834	-9.448	1.00	47.88	L21B C
ATOM	7200	C	GLY	161	75.947	45.670	-9.917	1.00	46.51	L21B C
ATOM	7201	O	GLY	161	76.533	45.684	-11.000	1.00	47.18	L21B O
ATOM	7202	N	VAL	162	75.991	44.642	-9.091	1.00	45.68	L21B N
ATOM	7203	CA	VAL	162	76.766	43.467	-9.420	1.00	46.39	L21B C
ATOM	7204	CB	VAL	162	76.994	42.592	-8.177	1.00	45.36	L21B C
ATOM	7205	CG1	VAL	162	77.924	41.457	-8.522	1.00	45.18	L21B C
ATOM	7206	CG2	VAL	162	77.563	43.423	-7.048	1.00	43.58	L21B C
ATOM	7207	C	VAL	162	76.066	42.636	-10.489	1.00	46.49	L21B C
ATOM	7208	O	VAL	162	74.848	42.523	-10.510	1.00	46.00	L21B O
ATOM	7209	N	GLU	163	76.852	42.058	-11.382	1.00	47.79	L21B N
ATOM	7210	CA	GLU	163	76.306	41.230	-12.440	1.00	49.24	L21B C
ATOM	7211	CB	GLU	163	75.884	42.106	-13.625	1.00	52.12	L21B C
ATOM	7212	CG	GLU	163	74.748	41.536	-14.443	1.00	60.05	L21B C
ATOM	7213	CD	GLU	163	73.390	41.886	-13.864	1.00	65.24	L21B C
ATOM	7214	OE1	GLU	163	72.774	41.012	-13.209	1.00	67.90	L21B O
ATOM	7215	OE2	GLU	163	72.940	43.038	-14.068	1.00	70.08	L21B O
ATOM	7216	C	GLU	163	77.401	40.260	-12.865	1.00	46.11	L21B C
ATOM	7217	O	GLU	163	78.410	40.682	-13.419	1.00	47.48	L21B O
ATOM	7218	N	THR	164	77.206	38.970	-12.599	1.00	41.78	L21B N

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ATOM   7219  CA  THR   164      78.282  37.988 -12.730  1.00 38.50      L21B C
ATOM   7220  CB  THR   164      78.531  37.248 -11.400  1.00 34.50      L21B C
ATOM   7221  OG1 THR   164      78.699  38.199 -10.346  1.00 30.64      L21B O
ATOM   7222  CG2 THR   164      79.756  36.372 -11.495  1.00 28.00      L21B C
ATOM   7223  C   THR   164      77.871  36.955 -13.756  1.00 38.26      L21B C
ATOM   7224  O   THR   164      76.696  36.638 -13.868  1.00 39.75      L21B O
ATOM   7225  N   THR   165      78.838  36.416 -14.487  1.00 37.96      L21B N
ATOM   7226  CA  THR   165      78.562  35.385 -15.483  1.00 37.83      L21B C
ATOM   7227  CB  THR   165      79.599  35.420 -16.618  1.00 34.80      L21B C
ATOM   7228  OG1 THR   165      80.880  35.040 -16.097  1.00 36.93      L21B O
ATOM   7229  CG2 THR   165      79.709  36.825 -17.215  1.00 31.26      L21B C
ATOM   7230  C   THR   165      78.593  33.985 -14.866  1.00 37.21      L21B C
ATOM   7231  O   THR   165      79.114  33.770 -13.777  1.00 39.28      L21B O
ATOM   7232  N   THR   166      78.036  33.019 -15.565  1.00 36.76      L21B N
ATOM   7233  CA  THR   166      78.243  31.645 -15.168  1.00 38.24      L21B C
ATOM   7234  CB  THR   166      77.106  30.778 -15.660  1.00 34.50      L21B C
ATOM   7235  OG1 THR   166      77.096  30.796 -17.089  1.00 34.16      L21B O
ATOM   7236  CG2 THR   166      75.787  31.322 -15.164  1.00 33.33      L21B C
ATOM   7237  C   THR   166      79.537  31.194 -15.827  1.00 37.53      L21B C
ATOM   7238  O   THR   166      79.864  31.623 -16.932  1.00 36.25      L21B O
ATOM   7239  N   PRO   167      80.299  30.325 -15.153  1.00 37.69      L21B N
ATOM   7240  CD  PRO   167      80.022  29.728 -13.837  1.00 37.82      L21B C
ATOM   7241  CA  PRO   167      81.614  29.933 -15.668  1.00 41.45      L21B C
ATOM   7242  CB  PRO   167      82.133  28.968 -14.613  1.00 38.56      L21B C
ATOM   7243  CG  PRO   167      81.376  29.323 -13.368  1.00 37.09      L21B C
ATOM   7244  C   PRO   167      81.473  29.277 -17.029  1.00 42.27      L21B C
ATOM   7245  O   PRO   167      80.425  28.735 -17.341  1.00 44.62      L21B O
ATOM   7246  N   SER   168      82.514  29.334 -17.845  1.00 43.91      L21B N
ATOM   7247  CA  SER   168      82.479  28.642 -19.123  1.00 49.02      L21B C
ATOM   7248  CB  SER   168      81.934  29.574 -20.200  1.00 48.31      L21B C
ATOM   7249  OG  SER   168      82.776  30.696 -20.353  1.00 51.04      L21B O
ATOM   7250  C   SER   168      83.860  28.112 -19.517  1.00 52.43      L21B C
ATOM   7251  O   SER   168      84.858  28.810 -19.384  1.00 51.73      L21B O
ATOM   7252  N   LYS   169      83.907  26.866 -19.981  1.00 56.92      L21B N
ATOM   7253  CA  LYS   169      85.169  26.213 -20.283  1.00 61.14      L21B C
ATOM   7254  CB  LYS   169      84.918  24.845 -20.924  1.00 62.84      L21B C
ATOM   7255  CG  LYS   169      84.830  23.687 -19.932  1.00 66.49      L21B C
ATOM   7256  CD  LYS   169      85.821  22.565 -20.294  1.00 70.79      L21B C
ATOM   7257  CE  LYS   169      86.088  21.615 -19.117  1.00 73.34      L21B C
ATOM   7258  NZ  LYS   169      84.856  20.937 -18.570  1.00 78.76      L21B N
ATOM   7259  C   LYS   169      85.955  27.092 -21.235  1.00 62.66      L21B C
ATOM   7260  O   LYS   169      85.383  27.720 -22.115  1.00 63.23      L21B O
ATOM   7261  N   GLN   170      87.269  27.141 -21.053  1.00 64.67      L21B N
ATOM   7262  CA  GLN   170      88.105  28.052 -21.828  1.00 65.86      L21B C
ATOM   7263  CB  GLN   170      89.262  28.556 -20.967  1.00 66.96      L21B C
ATOM   7264  CG  GLN   170      88.852  29.309 -19.725  1.00 68.32      L21B C
ATOM   7265  CD  GLN   170      90.050  29.889 -18.995  1.00 69.91      L21B C
ATOM   7266  OE1 GLN   170      90.010  30.118 -17.783  1.00 71.16      L21B O
ATOM   7267  NE2 GLN   170      91.127  30.130 -19.734  1.00 71.33      L21B N
ATOM   7268  C   GLN   170      88.677  27.413 -23.097  1.00 66.50      L21B C
ATOM   7269  O   GLN   170      87.997  27.295 -24.125  1.00 66.61      L21B O
ATOM   7270  N   SER   171      89.946  27.027 -23.009  1.00 66.58      L21B N
ATOM   7271  CA  SER   171      90.641  26.325 -24.081  1.00 67.32      L21B C
ATOM   7272  CB  SER   171      91.243  27.318 -25.076  1.00 68.07      L21B C
ATOM   7273  OG  SER   171      90.258  27.775 -25.985  1.00 70.61      L21B O
ATOM   7274  C   SER   171      91.744  25.474 -23.466  1.00 66.68      L21B C
ATOM   7275  O   SER   171      92.026  24.376 -23.938  1.00 65.89      L21B O
ATOM   7276  N   ASN   172      92.361  25.994 -22.408  1.00 65.46      L21B N
ATOM   7277  CA  ASN   172      93.171  25.173 -21.517  1.00 63.92      L21B C
ATOM   7278  CB  ASN   172      93.956  26.056 -20.544  1.00 62.95      L21B C
ATOM   7279  CG  ASN   172      93.058  26.891 -19.652  1.00 62.55      L21B C
ATOM   7280  OD1 ASN   172      92.050  27.431 -20.100  1.00 62.38      L21B O
ATOM   7281  ND2 ASN   172      93.426  27.003 -18.378  1.00 61.56      L21B N
ATOM   7282  C   ASN   172      92.230  24.245 -20.754  1.00 63.74      L21B C
ATOM   7283  O   ASN   172      92.646  23.450 -19.916  1.00 63.97      L21B O
ATOM   7284  N   ASN   173      90.946  24.363 -21.062  1.00 63.55      L21B N
ATOM   7285  CA  ASN   173      89.937  23.472 -20.518  1.00 62.96      L21B C
ATOM   7286  CB  ASN   173      90.350  22.014 -20.725  1.00 63.60      L21B C
ATOM   7287  CG  ASN   173      89.203  21.058 -20.483  1.00 65.50      L21B C
ATOM   7288  OD1 ASN   173      89.023  20.565 -19.366  1.00 67.22      L21B O
ATOM   7289  ND2 ASN   173      88.407  20.799 -21.525  1.00 65.19      L21B N
ATOM   7290  C   ASN   173      89.646  23.724 -19.037  1.00 61.57      L21B C
ATOM   7291  O   ASN   173      88.868  22.990 -18.415  1.00 60.70      L21B O
ATOM   7292  N   LYS   174      90.268  24.758 -18.474  1.00 58.80      L21B N
ATOM   7293  CA  LYS   174      89.852  25.262 -17.173  1.00 57.03      L21B C
ATOM   7294  CB  LYS   174      91.047  25.873 -16.436  1.00 56.43      L21B C
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ATOM   7295  CG  LYS   174      92.179  24.893 -16.202  1.00 54.88      L21B C
ATOM   7296  CD  LYS   174      93.293  25.524 -15.389  1.00 55.65      L21B C
ATOM   7297  CE  LYS   174      94.474  24.578 -15.209  1.00 54.27      L21B C
ATOM   7298  NZ  LYS   174      95.762  25.322 -15.121  1.00 55.04      L21B N
ATOM   7299  C   LYS   174      88.739  26.302 -17.352  1.00 56.01      L21B C
ATOM   7300  O   LYS   174      88.342  26.607 -18.482  1.00 54.56      L21B O
ATOM   7301  N   TYR   175      88.227  26.835 -16.245  1.00 54.93      L21B N
ATOM   7302  CA  TYR   175      87.035  27.680 -16.309  1.00 55.31      L21B C
ATOM   7303  CB  TYR   175      85.965  27.195 -15.319  1.00 57.78      L21B C
ATOM   7304  CG  TYR   175      85.217  25.953 -15.760  1.00 60.75      L21B C
ATOM   7305  CD1 TYR   175      84.034  26.051 -16.487  1.00 61.53      L21B C
ATOM   7306  CE1 TYR   175      83.348  24.923 -16.896  1.00 63.42      L21B C
ATOM   7307  CD2 TYR   175      85.695  24.685 -15.452  1.00 60.53      L21B C
ATOM   7308  CE2 TYR   175      85.017  23.552 -15.854  1.00 63.85      L21B C
ATOM   7309  CZ  TYR   175      83.845  23.672 -16.578  1.00 63.99      L21B C
ATOM   7310  OH  TYR   175      83.186  22.537 -16.995  1.00 64.21      L21B O
ATOM   7311  C   TYR   175      87.316  29.158 -16.058  1.00 53.03      L21B C
ATOM   7312  O   TYR   175      88.237  29.523 -15.323  1.00 53.97      L21B O
ATOM   7313  N   ALA   176      86.514  30.008 -16.686  1.00 48.30      L21B N
ATOM   7314  CA  ALA   176      86.626  31.440 -16.486  1.00 43.86      L21B C
ATOM   7315  CB  ALA   176      87.113  32.113 -17.765  1.00 41.69      L21B C
ATOM   7316  C   ALA   176      85.264  31.985 -16.087  1.00 41.19      L21B C
ATOM   7317  O   ALA   176      84.241  31.345 -16.310  1.00 40.23      L21B O
ATOM   7318  N   ALA   177      85.255  33.161 -15.480  1.00 37.62      L21B N
ATOM   7319  CA  ALA   177      84.012  33.839 -15.198  1.00 36.67      L21B C
ATOM   7320  CB  ALA   177      83.549  33.496 -13.807  1.00 36.19      L21B C
ATOM   7321  C   ALA   177      84.303  35.322 -15.312  1.00 36.41      L21B C
ATOM   7322  O   ALA   177      85.440  35.694 -15.603  1.00 38.70      L21B O
ATOM   7323  N   SER   178      83.296  36.169 -15.096  1.00 33.97      L21B N
ATOM   7324  CA  SER   178      83.560  37.567 -14.777  1.00 30.19      L21B C
ATOM   7325  CB  SER   178      83.893  38.359 -16.039  1.00 29.61      L21B C
ATOM   7326  OG  SER   178      82.756  38.462 -16.878  1.00 31.45      L21B O
ATOM   7327  C   SER   178      82.376  38.206 -14.076  1.00 29.26      L21B C
ATOM   7328  O   SER   178      81.254  37.704 -14.138  1.00 28.35      L21B O
ATOM   7329  N   SER   179      82.635  39.330 -13.420  1.00 28.18      L21B N
ATOM   7330  CA  SER   179      81.663  39.969 -12.551  1.00 28.05      L21B C
ATOM   7331  CB  SER   179      81.939  39.533 -11.099  1.00 25.74      L21B C
ATOM   7332  OG  SER   179      81.115  40.215 -10.164  1.00 25.59      L21B O
ATOM   7333  C   SER   179      81.785  41.497 -12.707  1.00 28.42      L21B C
ATOM   7334  O   SER   179      82.882  42.050 -12.632  1.00 30.29      L21B O
ATOM   7335  N   TYR   180      80.670  42.184 -12.919  1.00 28.81      L21B N
ATOM   7336  CA  TYR   180      80.718  43.628 -13.137  1.00 29.20      L21B C
ATOM   7337  CB  TYR   180      80.079  43.980 -14.493  1.00 25.61      L21B C
ATOM   7338  CG  TYR   180      80.858  43.465 -15.684  1.00 22.04      L21B C
ATOM   7339  CD1 TYR   180      80.891  42.115 -15.973  1.00 21.54      L21B C
ATOM   7340  CE1 TYR   180      81.619  41.623 -17.041  1.00 24.06      L21B C
ATOM   7341  CD2 TYR   180      81.585  44.325 -16.503  1.00 22.08      L21B C
ATOM   7342  CE2 TYR   180      82.332  43.841 -17.586  1.00 23.42      L21B C
ATOM   7343  CZ  TYR   180      82.341  42.484 -17.849  1.00 25.10      L21B C
ATOM   7344  OH  TYR   180      83.053  41.964 -18.914  1.00 26.40      L21B O
ATOM   7345  C   TYR   180      80.017  44.396 -12.017  1.00 29.98      L21B C
ATOM   7346  O   TYR   180      78.879  44.115 -11.665  1.00 30.59      L21B O
ATOM   7347  N   LEU   181      80.701  45.369 -11.445  1.00 31.66      L21B N
ATOM   7348  CA  LEU   181      80.075  46.194 -10.434  1.00 34.40      L21B C
ATOM   7349  CB  LEU   181      80.950  46.241  -9.178  1.00 35.91      L21B C
ATOM   7350  CG  LEU   181      80.622  47.301  -8.118  1.00 37.29      L21B C
ATOM   7351  CD1 LEU   181      79.141  47.276  -7.792  1.00 38.10      L21B C
ATOM   7352  CD2 LEU   181      81.458  47.046  -6.874  1.00 35.95      L21B C
ATOM   7353  C   LEU   181      79.860  47.593 -11.007  1.00 35.22      L21B C
ATOM   7354  O   LEU   181      80.823  48.271 -11.396  1.00 35.53      L21B O
ATOM   7355  N   SER   182      78.591  48.001 -11.072  1.00 34.30      L21B N
ATOM   7356  CA  SER   182      78.195  49.265 -11.687  1.00 34.46      L21B C
ATOM   7357  CB  SER   182      76.884  49.110 -12.448  1.00 33.20      L21B C
ATOM   7358  OG  SER   182      77.030  48.265 -13.568  1.00 36.65      L21B O
ATOM   7359  C   SER   182      78.001  50.314 -10.620  1.00 35.64      L21B C
ATOM   7360  O   SER   182      77.212  50.135  -9.687  1.00 37.03      L21B O
ATOM   7361  N   LEU   183      78.717  51.416 -10.770  1.00 35.80      L21B N
ATOM   7362  CA  LEU   183      78.703  52.494  -9.796  1.00 36.71      L21B C
ATOM   7363  CB  LEU   183      80.101  52.637  -9.193  1.00 32.29      L21B C
ATOM   7364  CG  LEU   183      80.297  51.602  -8.101  1.00 32.15      L21B C
ATOM   7365  CD1 LEU   183      81.726  51.602  -7.616  1.00 31.53      L21B C
ATOM   7366  CD2 LEU   183      79.335  51.923  -6.969  1.00 30.51      L21B C
ATOM   7367  C   LEU   183      78.291  53.790 -10.486  1.00 37.88      L21B C
ATOM   7368  O   LEU   183      78.384  53.896 -11.719  1.00 41.35      L21B O
ATOM   7369  N   THR   184      77.836  54.769  -9.708  1.00 37.42      L21B N
ATOM   7370  CA  THR   184      77.936  56.165 -10.137  1.00 36.46      L21B C
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ATOM   7371  CB  THR   184      77.002  57.095  -9.338  1.00 34.03      L21B C
ATOM   7372  OG1 THR   184      77.609  57.385  -8.073  1.00 32.25      L21B O
ATOM   7373  CG2 THR   184      75.644  56.460  -9.121  1.00 27.17      L21B C
ATOM   7374  C   THR   184      79.369  56.584  -9.807  1.00 37.89      L21B C
ATOM   7375  O   THR   184      80.027  55.952  -8.984  1.00 38.09      L21B O
ATOM   7376  N   PRO   185      79.867  57.664 -10.430  1.00 40.32      L21B N
ATOM   7377  CD  PRO   185      79.307  58.350 -11.603  1.00 38.82      L21B C
ATOM   7378  CA  PRO   185      81.209  58.171 -10.093  1.00 41.63      L21B C
ATOM   7379  CB  PRO   185      81.441  59.272 -11.113  1.00 37.99      L21B C
ATOM   7380  CG  PRO   185      80.532  58.935 -12.234  1.00 38.54      L21B C
ATOM   7381  C   PRO   185      81.288  58.701  -8.670  1.00 43.60      L21B C
ATOM   7382  O   PRO   185      82.314  58.619  -8.008  1.00 43.46      L21B O
ATOM   7383  N   GLU   186      80.178  59.239  -8.204  1.00 48.02      L21B N
ATOM   7384  CA  GLU   186      80.127  59.827  -6.887  1.00 53.68      L21B C
ATOM   7385  CB  GLU   186      78.767  60.491  -6.677  1.00 58.48      L21B C
ATOM   7386  CG  GLU   186      78.455  61.609  -7.690  1.00 66.64      L21B C
ATOM   7387  CD  GLU   186      78.399  61.126  -9.148  1.00 70.44      L21B C
ATOM   7388  OE1 GLU   186      78.031  59.955  -9.384  1.00 72.48      L21B O
ATOM   7389  OE2 GLU   186      78.727  61.930 -10.057  1.00 73.54      L21B O
ATOM   7390  C   GLU   186      80.360  58.748  -5.851  1.00 54.01      L21B C
ATOM   7391  O   GLU   186      81.019  58.978  -4.841  1.00 55.79      L21B O
ATOM   7392  N   GLN   187      79.830  57.558  -6.103  1.00 53.63      L21B N
ATOM   7393  CA  GLN   187      79.934  56.503  -5.109  1.00 51.43      L21B C
ATOM   7394  CB  GLN   187      78.681  55.628  -5.131  1.00 50.42      L21B C
ATOM   7395  CG  GLN   187      78.234  55.226  -6.505  1.00 50.59      L21B C
ATOM   7396  CD  GLN   187      76.767  54.823  -6.554  1.00 50.84      L21B C
ATOM   7397  OE1 GLN   187      76.405  53.872  -7.251  1.00 50.48      L21B O
ATOM   7398  NE2 GLN   187      75.917  55.547  -5.821  1.00 49.29      L21B N
ATOM   7399  C   GLN   187      81.189  55.667  -5.298  1.00 50.12      L21B C
ATOM   7400  O   GLN   187      81.622  54.956  -4.387  1.00 50.30      L21B O
ATOM   7401  N   TRP   188      81.782  55.767  -6.479  1.00 47.84      L21B N
ATOM   7402  CA  TRP   188      83.108  55.222  -6.694  1.00 47.52      L21B C
ATOM   7403  CB  TRP   188      83.420  55.191  -8.187  1.00 42.90      L21B C
ATOM   7404  CG  TRP   188      84.880  55.086  -8.485  1.00 39.27      L21B C
ATOM   7405  CD2 TRP   188      85.750  53.999  -8.157  1.00 36.97      L21B C
ATOM   7406  CE2 TRP   188      87.038  54.341  -8.621  1.00 35.12      L21B C
ATOM   7407  CE3 TRP   188      85.569  52.774  -7.515  1.00 36.62      L21B C
ATOM   7408  CD1 TRP   188      85.653  56.015  -9.116  1.00 37.57      L21B C
ATOM   7409  NE1 TRP   188      86.950  55.576  -9.203  1.00 35.22      L21B N
ATOM   7410  CZ2 TRP   188      88.141  53.496  -8.467  1.00 35.08      L21B C
ATOM   7411  CZ3 TRP   188      86.676  51.931  -7.360  1.00 37.18      L21B C
ATOM   7412  CH2 TRP   188      87.940  52.298  -7.834  1.00 35.01      L21B C
ATOM   7413  C   TRP   188      84.135  56.094  -5.962  1.00 50.19      L21B C
ATOM   7414  O   TRP   188      85.096  55.589  -5.366  1.00 51.20      L21B O
ATOM   7415  N   LYS   189      83.926  57.405  -6.004  1.00 50.84      L21B N
ATOM   7416  CA  LYS   189      84.898  58.326  -5.442  1.00 52.62      L21B C
ATOM   7417  CB  LYS   189      84.673  59.745  -5.974  1.00 53.78      L21B C
ATOM   7418  CG  LYS   189      85.073  59.957  -7.425  1.00 58.09      L21B C
ATOM   7419  CD  LYS   189      86.572  59.787  -7.626  1.00 61.78      L21B C
ATOM   7420  CE  LYS   189      86.927  59.741  -9.099  1.00 65.61      L21B C
ATOM   7421  NZ  LYS   189      88.402  59.624  -9.287  1.00 73.62      L21B N
ATOM   7422  C   LYS   189      84.818  58.339  -3.927  1.00 52.30      L21B C
ATOM   7423  O   LYS   189      85.780  58.698  -3.246  1.00 54.71      L21B O
ATOM   7424  N   SER   190      83.671  57.949  -3.393  1.00 49.80      L21B N
ATOM   7425  CA  SER   190      83.425  58.174  -1.984  1.00 47.44      L21B C
ATOM   7426  CB  SER   190      82.007  58.680  -1.772  1.00 45.86      L21B C
ATOM   7427  OG  SER   190      81.100  57.847  -2.455  1.00 46.05      L21B O
ATOM   7428  C   SER   190      83.658  56.935  -1.150  1.00 46.24      L21B C
ATOM   7429  O   SER   190      83.083  56.786  -0.081  1.00 46.23      L21B O
ATOM   7430  N   HIS   191      84.513  56.048  -1.633  1.00 45.30      L21B N
ATOM   7431  CA  HIS   191      85.104  55.051  -0.762  1.00 45.91      L21B C
ATOM   7432  CB  HIS   191      84.467  53.687  -1.037  1.00 44.94      L21B C
ATOM   7433  CG  HIS   191      83.037  53.602  -0.602  1.00 44.01      L21B C
ATOM   7434  CD2 HIS   191      81.888  53.929  -1.241  1.00 42.43      L21B C
ATOM   7435  ND1 HIS   191      82.669  53.191   0.661  1.00 44.59      L21B N
ATOM   7436  CE1 HIS   191      81.356  53.270   0.782  1.00 42.41      L21B C
ATOM   7437  NE2 HIS   191      80.859  53.716  -0.357  1.00 40.13      L21B N
ATOM   7438  C   HIS   191      86.623  54.994  -0.926  1.00 46.85      L21B C
ATOM   7439  O   HIS   191      87.143  55.172  -2.019  1.00 47.33      L21B O
ATOM   7440  N   ARG   192      87.334  54.762   0.170  1.00 47.92      L21B N
ATOM   7441  CA  ARG   192      88.779  54.561   0.102  1.00 49.29      L21B C
ATOM   7442  CB  ARG   192      89.364  54.317   1.496  1.00 53.95      L21B C
ATOM   7443  CG  ARG   192      89.706  55.585   2.258  1.00 62.03      L21B C
ATOM   7444  CD  ARG   192      91.080  56.101   1.869  1.00 68.57      L21B C
ATOM   7445  NE  ARG   192      91.236  57.520   2.178  1.00 76.88      L21B N
ATOM   7446  CZ  ARG   192      92.376  58.194   2.059  1.00 80.97      L21B C
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ATOM   7447  NH1 ARG   192      92.421  59.489   2.363  1.00 83.55      L21B N
ATOM   7448  NH2 ARG   192      93.474  57.574   1.640  1.00 83.12      L21B N
ATOM   7449  C   ARG   192      89.156  53.397  -0.800  1.00 46.81      L21B C
ATOM   7450  O   ARG   192      90.292  53.325  -1.266  1.00 47.72      L21B O
ATOM   7451  N   SER   193      88.222  52.481  -1.037  1.00 42.98      L21B N
ATOM   7452  CA  SER   193      88.496  51.342  -1.907  1.00 41.55      L21B C
ATOM   7453  CB  SER   193      89.566  50.434  -1.287  1.00 39.74      L21B C
ATOM   7454  OG  SER   193      89.199  50.021   0.018  1.00 40.91      L21B O
ATOM   7455  C   SER   193      87.243  50.526  -2.198  1.00 40.51      L21B C
ATOM   7456  O   SER   193      86.157  50.890  -1.774  1.00 40.73      L21B O
ATOM   7457  N   TYR   194      87.427  49.433  -2.938  1.00 38.90      L21B N
ATOM   7458  CA  TYR   194      86.377  48.500  -3.338  1.00 35.74      L21B C
ATOM   7459  CB  TYR   194      85.754  48.931  -4.667  1.00 32.23      L21B C
ATOM   7460  CG  TYR   194      84.533  49.816  -4.524  1.00 31.63      L21B C
ATOM   7461  CD1 TYR   194      83.254  49.267  -4.528  1.00 28.78      L21B C
ATOM   7462  CE1 TYR   194      82.127  50.060  -4.415  1.00 29.78      L21B C
ATOM   7463  CD2 TYR   194      84.656  51.202  -4.397  1.00 31.17      L21B C
ATOM   7464  CE2 TYR   194      83.526  52.017  -4.283  1.00 32.51      L21B C
ATOM   7465  CZ  TYR   194      82.259  51.433  -4.297  1.00 33.20      L21B C
ATOM   7466  OH  TYR   194      81.124  52.220  -4.238  1.00 34.06      L21B O
ATOM   7467  C   TYR   194      87.121  47.190  -3.531  1.00 36.59      L21B C
ATOM   7468  O   TYR   194      88.316  47.207  -3.855  1.00 35.94      L21B O
ATOM   7469  N   SER   195      86.457  46.052  -3.339  1.00 36.59      L21B N
ATOM   7470  CA  SER   195      87.182  44.785  -3.492  1.00 37.13      L21B C
ATOM   7471  CB  SER   195      87.685  44.281  -2.135  1.00 36.06      L21B C
ATOM   7472  OG  SER   195      88.132  45.339  -1.307  1.00 34.68      L21B O
ATOM   7473  C   SER   195      86.384  43.674  -4.162  1.00 37.41      L21B C
ATOM   7474  O   SER   195      85.171  43.554  -3.986  1.00 37.59      L21B O
ATOM   7475  N   CYS   196      87.077  42.853  -4.933  1.00 37.65      L21B N
ATOM   7476  CA  CYS   196      86.443  41.701  -5.528  1.00 38.71      L21B C
ATOM   7477  C   CYS   196      86.891  40.511  -4.690  1.00 39.42      L21B C
ATOM   7478  O   CYS   196      88.093  40.277  -4.532  1.00 40.53      L21B O
ATOM   7479  CB  CYS   196      86.881  41.552  -6.997  1.00 38.27      L21B C
ATOM   7480  SG  CYS   196      85.991  40.237  -7.892  1.00 40.80      L21B S
ATOM   7481  N   GLN   197      85.932  39.779  -4.122  1.00 39.35      L21B N
ATOM   7482  CA  GLN   197      86.259  38.614  -3.303  1.00 38.91      L21B C
ATOM   7483  CB  GLN   197      85.733  38.794  -1.876  1.00 38.51      L21B C
ATOM   7484  CG  GLN   197      86.541  39.823  -1.078  1.00 42.70      L21B C
ATOM   7485  CD  GLN   197      85.947  40.191   0.280  1.00 45.03      L21B C
ATOM   7486  OE1 GLN   197      84.739  40.429   0.416  1.00 47.31      L21B O
ATOM   7487  NE2 GLN   197      86.806  40.251   1.296  1.00 44.75      L21B N
ATOM   7488  C   GLN   197      85.688  37.361  -3.924  1.00 38.57      L21B C
ATOM   7489  O   GLN   197      84.475  37.185  -3.966  1.00 41.15      L21B O
ATOM   7490  N   VAL   198      86.565  36.493  -4.411  1.00 37.99      L21B N
ATOM   7491  CA  VAL   198      86.148  35.335  -5.186  1.00 39.54      L21B C
ATOM   7492  CB  VAL   198      86.892  35.280  -6.527  1.00 36.74      L21B C
ATOM   7493  CG1 VAL   198      86.339  34.164  -7.404  1.00 36.03      L21B C
ATOM   7494  CG2 VAL   198      86.769  36.607  -7.219  1.00 37.28      L21B C
ATOM   7495  C   VAL   198      86.432  34.048  -4.434  1.00 41.58      L21B C
ATOM   7496  O   VAL   198      87.563  33.563  -4.429  1.00 43.11      L21B O
ATOM   7497  N   THR   199      85.411  33.488  -3.798  1.00 43.26      L21B N
ATOM   7498  CA  THR   199      85.579  32.201  -3.133  1.00 45.16      L21B C
ATOM   7499  CB  THR   199      84.428  31.889  -2.144  1.00 43.09      L21B C
ATOM   7500  OG1 THR   199      84.438  32.838  -1.072  1.00 42.60      L21B O
ATOM   7501  CG2 THR   199      84.586  30.490  -1.569  1.00 41.46      L21B C
ATOM   7502  C   THR   199      85.602  31.108  -4.175  1.00 45.66      L21B C
ATOM   7503  O   THR   199      84.915  31.190  -5.184  1.00 47.58      L21B O
ATOM   7504  N   HIS   200      86.395  30.081  -3.921  1.00 47.28      L21B N
ATOM   7505  CA  HIS   200      86.419  28.904  -4.770  1.00 49.91      L21B C
ATOM   7506  CB  HIS   200      87.472  29.077  -5.864  1.00 47.40      L21B C
ATOM   7507  CG  HIS   200      87.659  27.864  -6.713  1.00 44.66      L21B C
ATOM   7508  CD2 HIS   200      86.973  27.417  -7.791  1.00 43.66      L21B C
ATOM   7509  ND1 HIS   200      88.656  26.941  -6.479  1.00 43.35      L21B N
ATOM   7510  CE1 HIS   200      88.573  25.978  -7.379  1.00 44.39      L21B C
ATOM   7511  NE2 HIS   200      87.560  26.242  -8.186  1.00 43.71      L21B N
ATOM   7512  C   HIS   200      86.768  27.716  -3.892  1.00 52.57      L21B C
ATOM   7513  O   HIS   200      87.756  27.768  -3.170  1.00 52.42      L21B O
ATOM   7514  N   GLU   201      85.961  26.658  -3.942  1.00 57.26      L21B N
ATOM   7515  CA  GLU   201      86.200  25.483  -3.098  1.00 63.32      L21B C
ATOM   7516  CB  GLU   201      87.180  24.515  -3.776  1.00 66.42      L21B C
ATOM   7517  CG  GLU   201      86.891  24.244  -5.239  1.00 75.18      L21B C
ATOM   7518  CD  GLU   201      85.769  23.242  -5.448  1.00 79.87      L21B C
ATOM   7519  OE1 GLU   201      84.840  23.536  -6.242  1.00 82.95      L21B O
ATOM   7520  OE2 GLU   201      85.820  22.163  -4.818  1.00 82.36      L21B O
ATOM   7521  C   GLU   201      86.793  25.904  -1.755  1.00 64.56      L21B C
ATOM   7522  O   GLU   201      87.925  25.538  -1.431  1.00 64.47      L21B O
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ATOM   7523  N   GLY  201      86.046  26.689  -0.988  1.00 66.07      L21B N
ATOM   7524  CA  GLY  201      86.506  27.076   0.333  1.00 67.95      L21B C
ATOM   7525  C   GLY  201      87.527  28.201   0.360  1.00 70.16      L21B C
ATOM   7526  O   GLY  201      87.648  28.895   1.373  1.00 71.89      L21B O
ATOM   7527  N   SER  203      88.249  28.394  -0.745  1.00 70.44      L21B N
ATOM   7528  CA  SER  203      89.361  29.358  -0.812  1.00 69.03      L21B C
ATOM   7529  CB  SER  203      90.540  28.731  -1.571  1.00 69.91      L21B C
ATOM   7530  OG  SER  203      90.239  28.519  -2.948  1.00 70.63      L21B O
ATOM   7531  C   SER  203      89.020  30.723  -1.450  1.00 66.90      L21B C
ATOM   7532  O   SER  203      88.819  30.816  -2.661  1.00 66.99      L21B O
ATOM   7533  N   THR  204      88.999  31.779  -0.636  1.00 64.21      L21B N
ATOM   7534  CA  THR  204      88.670  33.136  -1.108  1.00 61.96      L21B C
ATOM   7535  CB  THR  204      87.967  33.949   0.016  1.00 62.44      L21B C
ATOM   7536  OG1 THR  204      86.801  33.248   0.457  1.00 61.41      L21B O
ATOM   7537  CG2 THR  204      87.560  35.324  -0.484  1.00 63.92      L21B C
ATOM   7538  C   THR  204      89.890  33.951  -1.604  1.00 59.78      L21B C
ATOM   7539  O   THR  204      90.821  34.226  -0.847  1.00 57.60      L21B O
ATOM   7540  N   VAL  205      89.874  34.348  -2.874  1.00 57.57      L21B N
ATOM   7541  CA  VAL  205      90.932  35.201  -3.410  1.00 56.13      L21B C
ATOM   7542  CB  VAL  205      91.390  34.727  -4.797  1.00 55.74      L21B C
ATOM   7543  CG1 VAL  205      92.575  35.542  -5.245  1.00 55.66      L21B C
ATOM   7544  CG2 VAL  205      91.724  33.264  -4.768  1.00 56.05      L21B C
ATOM   7545  C   VAL  205      90.420  36.631  -3.555  1.00 55.70      L21B C
ATOM   7546  O   VAL  205      89.401  36.867  -4.202  1.00 57.15      L21B O
ATOM   7547  N   GLU  206      91.139  37.584  -2.973  1.00 53.18      L21B N
ATOM   7548  CA  GLU  206      90.673  38.965  -2.912  1.00 50.02      L21B C
ATOM   7549  CB  GLU  206      90.657  39.419  -1.459  1.00 50.29      L21B C
ATOM   7550  CG  GLU  206      90.759  40.913  -1.272  1.00 51.53      L21B C
ATOM   7551  CD  GLU  206      90.102  41.358   0.009  1.00 52.70      L21B C
ATOM   7552  OE1 GLU  206      89.339  40.550   0.577  1.00 54.78      L21B O
ATOM   7553  OE2 GLU  206      90.339  42.503   0.447  1.00 54.12      L21B O
ATOM   7554  C   GLU  206      91.495  39.950  -3.743  1.00 48.36      L21B C
ATOM   7555  O   GLU  206      92.713  39.824  -3.853  1.00 49.43      L21B O
ATOM   7556  N   LYS  207      90.826  40.938  -4.325  1.00 46.01      L21B N
ATOM   7557  CA  LYS  207      91.519  42.007  -5.033  1.00 43.18      L21B C
ATOM   7558  CB  LYS  207      91.554  41.729  -6.529  1.00 40.80      L21B C
ATOM   7559  CG  LYS  207      92.938  41.528  -7.081  1.00 37.52      L21B C
ATOM   7560  CD  LYS  207      93.639  40.392  -6.382  1.00 39.09      L21B C
ATOM   7561  CE  LYS  207      95.108  40.317  -6.775  1.00 39.83      L21B C
ATOM   7562  NZ  LYS  207      95.307  40.319  -8.260  1.00 39.68      L21B N
ATOM   7563  C   LYS  207      90.827  43.329  -4.782  1.00 43.89      L21B C
ATOM   7564  O   LYS  207      89.633  43.372  -4.487  1.00 45.18      L21B O
ATOM   7565  N   THR  208      91.584  44.411  -4.890  1.00 43.72      L21B N
ATOM   7566  CA  THR  208      91.081  45.721  -4.527  1.00 43.45      L21B C
ATOM   7567  CB  THR  208      91.437  46.051  -3.071  1.00 43.01      L21B C
ATOM   7568  OG1 THR  208      90.722  45.168  -2.199  1.00 43.45      L21B O
ATOM   7569  CG2 THR  208      91.091  47.494  -2.737  1.00 43.14      L21B C
ATOM   7570  C   THR  208      91.655  46.795  -5.437  1.00 45.05      L21B C
ATOM   7571  O   THR  208      92.786  46.690  -5.903  1.00 43.07      L21B O
ATOM   7572  N   VAL  209      90.850  47.820  -5.696  1.00 47.06      L21B N
ATOM   7573  CA  VAL  209      91.288  48.975  -6.448  1.00 49.42      L21B C
ATOM   7574  CB  VAL  209      90.703  48.950  -7.853  1.00 50.14      L21B C
ATOM   7575  CG1 VAL  209      91.166  47.695  -8.573  1.00 47.79      L21B C
ATOM   7576  CG2 VAL  209      89.182  49.003  -7.779  1.00 51.93      L21B C
ATOM   7577  C   VAL  209      90.794  50.208  -5.719  1.00 52.56      L21B C
ATOM   7578  O   VAL  209      89.876  50.120  -4.906  1.00 51.78      L21B O
ATOM   7579  N   ALA  210      91.411  51.354  -5.997  1.00 57.14      L21B N
ATOM   7580  CA  ALA  210      91.030  52.611  -5.343  1.00 60.45      L21B C
ATOM   7581  CB  ALA  210      92.036  52.962  -4.256  1.00 57.24      L21B C
ATOM   7582  C   ALA  210      90.918  53.763  -6.331  1.00 64.66      L21B C
ATOM   7583  O   ALA  210      91.609  53.801  -7.352  1.00 61.92      L21B O
ATOM   7584  N   PRO  211      90.039  54.726  -6.033  1.00 70.80      L21B N
ATOM   7585  CD  PRO  211      89.146  54.779  -4.868  1.00 73.04      L21B C
ATOM   7586  CA  PRO  211      89.878  55.890  -6.898  1.00 76.06      L21B C
ATOM   7587  CB  PRO  211      88.663  56.608  -6.316  1.00 75.09      L21B C
ATOM   7588  CG  PRO  211      88.606  56.179  -4.918  1.00 75.55      L21B C
ATOM   7589  C   PRO  211      91.139  56.737  -6.872  1.00 80.46      L21B C
ATOM   7590  O   PRO  211      91.237  57.751  -7.560  1.00 79.95      L21B O
ATOM   7591  N   THR  212      92.110  56.302  -6.078  1.00 85.46      L21B N
ATOM   7592  CA  THR  212      93.399  56.973  -6.031  1.00 91.28      L21B C
ATOM   7593  CB  THR  212      94.165  56.604  -4.734  1.00 93.31      L21B C
ATOM   7594  OG1 THR  212      93.322  56.861  -3.602  1.00 94.30      L21B O
ATOM   7595  CG2 THR  212      95.445  57.441  -4.596  1.00 94.22      L21B C
ATOM   7596  C   THR  212      94.220  56.575  -7.260  1.00 92.75      L21B C
ATOM   7597  O   THR  212      95.341  56.047  -7.083  1.00 96.55      L21B O
ATOM   7598  OXT THR  212      93.725  56.799  -8.390  1.00 95.26      L21B O
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TER    7599      THR   212                                                L21B
ATOM   7600  OB  GLU     1      55.064   6.583 -21.347  1.00 73.12      H21B C
ATOM   7601  CG  GLU     1      55.535   7.444 -22.516  1.00 78.67      H21B C
ATOM   7602  CD  GLU     1      56.988   7.870 -22.365  1.00 81.81      H21B C
ATOM   7603  OE1 GLU     1      57.708   7.220 -21.575  1.00 83.96      H21B O
ATOM   7604  OE2 GLU     1      57.407   8.852 -23.024  1.00 83.08      H21B O
ATOM   7605  C   GLU     1      53.453   8.324 -20.546  1.00 66.92      H21B C
ATOM   7606  O   GLU     1      53.014   9.132 -21.373  1.00 64.83      H21B O
ATOM   7607  N   GLU     1      52.655   6.390 -21.940  1.00 69.73      H21B N
ATOM   7608  CA  GLU     1      53.623   6.841 -20.892  1.00 69.78      H21B C
ATOM   7609  N   VAL     2      53.807   8.669 -19.313  1.00 63.58      H21B N
ATOM   7610  CA  VAL     2      53.524   9.995 -18.788  1.00 61.52      H21B C
ATOM   7611  CB  VAL     2      53.746  10.065 -17.280  1.00 63.16      H21B C
ATOM   7612  CG1 VAL     2      53.593  11.493 -16.815  1.00 63.16      H21B C
ATOM   7613  CG2 VAL     2      52.763   9.151 -16.571  1.00 63.55      H21B C
ATOM   7614  C   VAL     2      54.417  11.027 -19.429  1.00 59.20      H21B C
ATOM   7615  O   VAL     2      55.621  10.999 -19.259  1.00 57.87      H21B O
ATOM   7616  N   GLN     3      53.806  11.958 -20.142  1.00 58.46      H21B N
ATOM   7617  CA  GLN     3      54.511  12.768 -21.111  1.00 57.91      H21B C
ATOM   7618  CB  GLN     3      54.405  12.090 -22.477  1.00 59.52      H21B C
ATOM   7619  CG  GLN     3      55.250  12.691 -23.581  1.00 64.49      H21B C
ATOM   7620  CD  GLN     3      55.179  11.867 -24.857  1.00 66.57      H21B C
ATOM   7621  OE1 GLN     3      55.033  12.406 -25.963  1.00 67.65      H21B O
ATOM   7622  NE2 GLN     3      55.277  10.548 -24.709  1.00 66.83      H21B N
ATOM   7623  C   GLN     3      53.894  14.164 -21.146  1.00 57.21      H21B C
ATOM   7624  O   GLN     3      52.762  14.345 -21.599  1.00 57.22      H21B O
ATOM   7625  N   LEU     4      54.644  15.147 -20.659  1.00 55.99      H21B N
ATOM   7626  CA  LEU     4      54.177  16.523 -20.595  1.00 54.53      H21B C
ATOM   7627  CB  LEU     4      54.485  17.099 -19.224  1.00 52.38      H21B C
ATOM   7628  CG  LEU     4      53.635  16.536 -18.091  1.00 50.58      H21B C
ATOM   7629  CD1 LEU     4      54.188  16.982 -16.749  1.00 49.08      H21B C
ATOM   7630  CD2 LEU     4      52.213  17.010 -18.278  1.00 51.45      H21B C
ATOM   7631  C   LEU     4      54.858  17.359 -21.661  1.00 55.60      H21B C
ATOM   7632  O   LEU     4      56.076  17.298 -21.806  1.00 56.73      H21B O
ATOM   7633  N   VAL     5      54.075  18.141 -22.401  1.00 56.76      H21B N
ATOM   7634  CA  VAL     5      54.585  18.838 -23.582  1.00 59.30      H21B C
ATOM   7635  CB  VAL     5      54.087  18.158 -24.883  1.00 57.84      H21B C
ATOM   7636  CG1 VAL     5      54.545  18.953 -26.102  1.00 57.36      H21B C
ATOM   7637  CG2 VAL     5      54.604  16.735 -24.959  1.00 54.25      H21B C
ATOM   7638  C   VAL     5      54.212  20.323 -23.642  1.00 62.11      H21B C
ATOM   7639  O   VAL     5      53.066  20.677 -23.925  1.00 60.98      H21B O
ATOM   7640  N   GLN     6      55.193  21.189 -23.406  1.00 66.23      H21B N
ATOM   7641  CA  GLN     6      54.936  22.619 -23.271  1.00 72.12      H21B C
ATOM   7642  CB  GLN     6      55.978  23.242 -22.340  1.00 71.68      H21B C
ATOM   7643  CG  GLN     6      56.117  22.517 -21.013  1.00 74.65      H21B C
ATOM   7644  CD  GLN     6      57.118  23.171 -20.085  1.00 76.41      H21B C
ATOM   7645  OE1 GLN     6      57.615  22.547 -19.148  1.00 76.12      H21B O
ATOM   7646  NE2 GLN     6      57.419  24.434 -20.342  1.00 83.40      H21B N
ATOM   7647  C   GLN     6      54.900  23.380 -24.600  1.00 75.46      H21B C
ATOM   7648  O   GLN     6      55.484  22.953 -25.593  1.00 76.72      H21B O
ATOM   7649  N   SER     7      54.205  24.514 -24.603  1.00 79.41      H21B N
ATOM   7650  CA  SER     7      54.059  25.333 -25.795  1.00 81.97      H21B C
ATOM   7651  CB  SER     7      53.273  26.603 -25.468  1.00 81.36      H21B C
ATOM   7652  OG  SER     7      53.846  27.308 -24.380  1.00 76.16      H21B O
ATOM   7653  C   SER     7      55.413  25.709 -26.365  1.00 85.28      H21B C
ATOM   7654  O   SER     7      56.456  25.356 -25.817  1.00 86.88      H21B O
ATOM   7655  N   GLY     8      55.390  26.435 -27.472  1.00 88.03      H21B N
ATOM   7656  CA  GLY     8      56.628  26.845 -28.098  1.00 91.52      H21B C
ATOM   7657  C   GLY     8      57.347  27.965 -27.373  1.00 93.23      H21B C
ATOM   7658  O   GLY     8      56.899  28.455 -26.338  1.00 91.33      H21B O
ATOM   7659  N   ALA     9      58.481  28.371 -27.925  1.00 97.66      H21B N
ATOM   7660  CA  ALA     9      59.244  29.459 -27.350  1.00101.03      H21B C
ATOM   7661  CB  ALA     9      60.687  29.377 -27.793  1.00105.95      H21B C
ATOM   7662  C   ALA     9      58.633  30.774 -27.791  1.00101.99      H21B C
ATOM   7663  O   ALA     9      58.185  30.914 -28.930  1.00101.78      H21B O
ATOM   7664  N   GLU    10      58.612  31.734 -26.874  1.00101.94      H21B N
ATOM   7665  CA  GLU    10      57.969  33.011 -27.124  1.00102.88      H21B C
ATOM   7666  CB  GLU    10      56.567  33.016 -26.505  1.00106.55      H21B C
ATOM   7667  CG  GLU    10      55.579  32.157 -27.275  1.00112.96      H21B C
ATOM   7668  CD  GLU    10      54.401  31.728 -26.439  1.00116.32      H21B C
ATOM   7669  OE1 GLU    10      54.359  32.110 -25.253  1.00119.31      H21B O
ATOM   7670  OE2 GLU    10      53.522  31.011 -26.963  1.00119.07      H21B O
ATOM   7671  C   GLU    10      58.779  34.190 -26.607  1.00100.78      H21B C
ATOM   7672  O   GLU    10      59.532  34.061 -25.637  1.00 99.72      H21B O
ATOM   7673  N   VAL    11      58.614  35.335 -27.274  1.00 97.98      H21B N
ATOM   7674  CA  VAL    11      59.360  36.559 -26.964  1.00 94.70      H21B C
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ATOM   7675  CB  VAL   11      60.409  36.848 -28.061  1.00 94.26      H21B C
ATOM   7676  CG1 VAL   11      61.190  38.097 -27.722  1.00 94.82      H21B C
ATOM   7677  CG2 VAL   11      61.355  35.668 -28.192  1.00 93.77      H21B C
ATOM   7678  C   VAL   11      58.456  37.797 -26.799  1.00 90.98      H21B C
ATOM   7679  O   VAL   11      57.662  38.115 -27.678  1.00 90.99      H21B O
ATOM   7680  N   LYS   12      58.596  38.492 -25.669  1.00 86.92      H21B N
ATOM   7681  CA  LYS   12      57.657  39.540 -25.256  1.00 82.85      H21B C
ATOM   7682  CB  LYS   12      56.660  38.978 -24.225  1.00 83.38      H21B C
ATOM   7683  CG  LYS   12      55.735  37.868 -24.752  1.00 83.33      H21B C
ATOM   7684  CD  LYS   12      54.466  38.433 -25.388  1.00 84.59      H21B C
ATOM   7685  CE  LYS   12      53.732  37.413 -26.250  1.00 85.71      H21B C
ATOM   7686  NZ  LYS   12      52.793  38.089 -27.202  1.00 88.33      H21B N
ATOM   7687  C   LYS   12      58.358  40.772 -24.668  1.00 79.74      H21B C
ATOM   7688  O   LYS   12      59.426  40.666 -24.070  1.00 79.34      H21B O
ATOM   7689  N   LYS   13      57.744  41.939 -24.850  1.00 76.15      H21B N
ATOM   7690  CA  LYS   13      58.303  43.198 -24.367  1.00 73.12      H21B C
ATOM   7691  CB  LYS   13      57.899  44.356 -25.282  1.00 72.80      H21B C
ATOM   7692  CG  LYS   13      58.547  44.332 -26.644  1.00 72.33      H21B C
ATOM   7693  CD  LYS   13      58.100  45.521 -27.465  0.50 75.00      H21B C
ATOM   7694  CE  LYS   13      58.853  45.603 -28.778  1.00 76.22      H21B C
ATOM   7695  NZ  LYS   13      58.325  46.709 -29.614  1.00 79.14      H21B N
ATOM   7696  C   LYS   13      57.778  43.466 -22.975  1.00 69.85      H21B C
ATOM   7697  O   LYS   13      56.745  42.947 -22.589  1.00 70.18      H21B O
ATOM   7698  N   PRO   14      58.480  44.295 -22.201  1.00 66.74      H21B N
ATOM   7699  CD  PRO   14      59.782  44.935 -22.443  1.00 64.63      H21B C
ATOM   7700  CA  PRO   14      57.962  44.600 -20.867  1.00 65.80      H21B C
ATOM   7701  CB  PRO   14      58.893  45.707 -20.372  1.00 63.65      H21B C
ATOM   7702  CG  PRO   14      60.177  45.416 -21.067  1.00 63.20      H21B C
ATOM   7703  C   PRO   14      56.506  45.031 -20.935  1.00 64.83      H21B C
ATOM   7704  O   PRO   14      56.044  45.494 -21.975  1.00 65.82      H21B O
ATOM   7705  N   GLY   15      55.781  44.850 -19.836  1.00 63.25      H21B N
ATOM   7706  CA  GLY   15      54.388  45.265 -19.789  1.00 60.95      H21B C
ATOM   7707  C   GLY   15      53.404  44.306 -20.443  1.00 60.18      H21B C
ATOM   7708  O   GLY   15      52.208  44.384 -20.176  1.00 60.15      H21B O
ATOM   7709  N   ALA   16      53.892  43.402 -21.293  1.00 58.86      H21B N
ATOM   7710  CA  ALA   16      53.016  42.476 -22.013  1.00 57.51      H21B C
ATOM   7711  CB  ALA   16      53.634  42.097 -23.352  1.00 56.81      H21B C
ATOM   7712  C   ALA   16      52.686  41.203 -21.234  1.00 56.45      H21B C
ATOM   7713  O   ALA   16      52.973  41.096 -20.034  1.00 54.45      H21B O
ATOM   7714  N   SER   17      52.079  40.246 -21.944  1.00 55.06      H21B N
ATOM   7715  CA  SER   17      51.619  38.989 -21.355  1.00 54.13      H21B C
ATOM   7716  CB  SER   17      50.116  39.050 -21.119  1.00 52.63      H21B C
ATOM   7717  OG  SER   17      49.819  40.066 -20.184  1.00 52.89      H21B O
ATOM   7718  C   SER   17      51.942  37.752 -22.186  1.00 54.17      H21B C
ATOM   7719  O   SER   17      51.745  37.745 -23.401  1.00 54.06      H21B O
ATOM   7720  N   VAL   18      52.440  36.709 -21.522  1.00 54.75      H21B N
ATOM   7721  CA  VAL   18      52.538  35.380 -22.122  1.00 55.21      H21B C
ATOM   7722  CB  VAL   18      53.959  34.781 -22.042  1.00 54.56      H21B C
ATOM   7723  CG1 VAL   18      54.718  35.067 -23.313  1.00 55.71      H21B C
ATOM   7724  CG2 VAL   18      54.693  35.337 -20.836  1.00 53.13      H21B C
ATOM   7725  C   VAL   18      51.615  34.373 -21.477  1.00 54.61      H21B C
ATOM   7726  O   VAL   18      51.310  34.455 -20.288  1.00 52.93      H21B O
ATOM   7727  N   LYS   19      51.190  33.416 -22.292  1.00 55.58      H21B N
ATOM   7728  CA  LYS   19      50.429  32.270 -21.834  1.00 56.30      H21B C
ATOM   7729  CB  LYS   19      48.989  32.340 -22.368  1.00 57.21      H21B C
ATOM   7730  CG  LYS   19      48.025  31.342 -21.704  1.00 59.70      H21B C
ATOM   7731  CD  LYS   19      46.632  31.357 -22.314  1.00 64.05      H21B C
ATOM   7732  CE  LYS   19      45.643  30.623 -21.438  1.00 67.96      H21B C
ATOM   7733  NZ  LYS   19      44.243  30.905 -21.843  1.00 77.89      H21B N
ATOM   7734  C   LYS   19      51.130  31.015 -22.354  1.00 55.43      H21B C
ATOM   7735  O   LYS   19      51.354  30.885 -23.555  1.00 56.86      H21B O
ATOM   7736  N   VAL   20      51.471  30.107 -21.439  1.00 53.48      H21B N
ATOM   7737  CA  VAL   20      52.219  28.892 -21.745  1.00 51.91      H21B C
ATOM   7738  CB  VAL   20      53.465  28.788 -20.822  1.00 52.28      H21B C
ATOM   7739  CG1 VAL   20      54.133  27.415 -20.967  1.00 51.32      H21B C
ATOM   7740  CG2 VAL   20      54.454  29.897 -21.164  1.00 49.80      H21B C
ATOM   7741  C   VAL   20      51.361  27.633 -21.571  1.00 51.11      H21B C
ATOM   7742  O   VAL   20      50.687  27.479 -20.554  1.00 50.92      H21B O
ATOM   7743  N   SER   21      51.405  26.736 -22.557  1.00 49.84      H21B N
ATOM   7744  CA  SER   21      50.636  25.490 -22.538  1.00 49.48      H21B C
ATOM   7745  CB  SER   21      50.209  25.094 -23.954  1.00 47.62      H21B C
ATOM   7746  OG  SER   21      49.534  26.136 -24.630  1.00 45.69      H21B O
ATOM   7747  C   SER   21      51.426  24.323 -21.961  1.00 51.01      H21B C
ATOM   7748  O   SER   21      52.644  24.292 -22.061  1.00 53.07      H21B O
ATOM   7749  N   CYS   22      50.714  23.354 -21.388  1.00 52.61      H21B N
ATOM   7750  CA  CYS   22      51.288  22.089 -20.914  1.00 53.01      H21B C
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ATOM   7751  C   CYS  22      50.295  20.978 -21.235  1.00 54.06      H21B C
ATOM   7752  O   CYS  22      49.179  20.977 -20.726  1.00 53.18      H21B O
ATOM   7753  CB  CYS  22      51.486  22.139 -19.402  1.00 52.59      H21B C
ATOM   7754  SG  CYS  22      52.311  20.704 -18.625  1.00 50.07      H21B S
ATOM   7755  N   LYS  23      50.689  20.031 -22.069  1.00 56.00      H21B N
ATOM   7756  CA  LYS  23      49.755  19.003 -22.496  1.00 58.44      H21B C
ATOM   7757  CB  LYS  23      49.727  18.903 -24.020  1.00 59.90      H21B C
ATOM   7758  CG  LYS  23      48.465  19.442 -24.661  1.00 63.71      H21B C
ATOM   7759  CD  LYS  23      47.553  18.301 -25.077  1.00 71.14      H21B C
ATOM   7760  CE  LYS  23      48.115  17.544 -26.270  1.00 75.82      H21B C
ATOM   7761  NZ  LYS  23      47.296  16.349 -26.613  1.00 81.04      H21B N
ATOM   7762  C   LYS  23      50.155  17.674 -21.911  1.00 58.80      H21B C
ATOM   7763  O   LYS  23      51.310  17.284 -22.004  1.00 59.77      H21B O
ATOM   7764  N   ALA  24      49.194  16.979 -21.312  1.00 59.37      H21B N
ATOM   7765  CA  ALA  24      49.479  15.720 -20.637  1.00 60.19      H21B C
ATOM   7766  CB  ALA  24      48.990  15.780 -19.204  1.00 61.61      H21B C
ATOM   7767  C   ALA  24      48.871  14.510 -21.342  1.00 60.44      H21B C
ATOM   7768  O   ALA  24      47.842  14.600 -22.013  1.00 61.03      H21B O
ATOM   7769  N   SER  25      49.523  13.370 -21.180  1.00 59.33      H21B N
ATOM   7770  CA  SER  25      49.035  12.142 -21.759  1.00 57.93      H21B C
ATOM   7771  CB  SER  25      49.605  11.971 -23.154  1.00 56.24      H21B C
ATOM   7772  OG  SER  25      50.986  11.688 -23.076  1.00 52.37      H21B O
ATOM   7773  C   SER  25      49.511  11.003 -20.879  1.00 59.09      H21B C
ATOM   7774  O   SER  25      50.546  11.117 -20.221  1.00 60.76      H21B O
ATOM   7775  N   GLY  26      48.754   9.910 -20.862  1.00 57.10      H21B N
ATOM   7776  CA  GLY  26      49.281   8.663 -20.341  1.00 54.77      H21B C
ATOM   7777  C   GLY  26      49.093   8.453 -18.853  1.00 52.65      H21B C
ATOM   7778  O   GLY  26      49.749   7.593 -18.240  1.00 51.57      H21B O
ATOM   7779  N   TYR  27      48.203   9.244 -18.264  1.00 50.72      H21B N
ATOM   7780  CA  TYR  27      47.809   9.047 -16.883  1.00 49.62      H21B C
ATOM   7781  CB  TYR  27      48.888   9.581 -15.934  1.00 48.68      H21B C
ATOM   7782  CG  TYR  27      48.984  11.088 -15.900  1.00 48.27      H21B C
ATOM   7783  CD1 TYR  27      49.651  11.789 -16.898  1.00 48.25      H21B C
ATOM   7784  CE1 TYR  27      49.720  13.174 -16.875  1.00 47.93      H21B C
ATOM   7785  CD2 TYR  27      48.392  11.813 -14.878  1.00 46.65      H21B C
ATOM   7786  CE2 TYR  27      48.455  13.188 -14.852  1.00 46.51      H21B C
ATOM   7787  CZ  TYR  27      49.118  13.863 -15.850  1.00 46.31      H21B C
ATOM   7788  OH  TYR  27      49.157  15.232 -15.823  1.00 45.63      H21B O
ATOM   7789  C   TYR  27      46.495   9.770 -16.661  1.00 49.46      H21B C
ATOM   7790  O   TYR  27      46.059  10.543 -17.509  1.00 50.28      H21B O
ATOM   7791  N   THR  28      45.865   9.495 -15.523  1.00 49.35      H21B N
ATOM   7792  CA  THR  28      44.567  10.061 -15.186  1.00 47.73      H21B C
ATOM   7793  CB  THR  28      43.865   9.233 -14.114  1.00 47.99      H21B C
ATOM   7794  OG1 THR  28      43.650   7.904 -14.603  1.00 45.83      H21B O
ATOM   7795  CG2 THR  28      42.531   9.866 -13.753  1.00 46.00      H21B C
ATOM   7796  C   THR  28      44.731  11.456 -14.647  1.00 47.40      H21B C
ATOM   7797  O   THR  28      45.152  11.646 -13.509  1.00 46.65      H21B O
ATOM   7798  N   LEU  29      44.369  12.428 -15.470  1.00 48.24      H21B N
ATOM   7799  CA  LEU  29      44.654  13.828 -15.203  1.00 48.79      H21B C
ATOM   7800  CB  LEU  29      44.044  14.687 -16.310  1.00 51.03      H21B C
ATOM   7801  CG  LEU  29      44.477  14.391 -17.750  1.00 51.75      H21B C
ATOM   7802  CD1 LEU  29      43.533  15.076 -18.741  1.00 51.69      H21B C
ATOM   7803  CD2 LEU  29      45.909  14.871 -17.940  1.00 54.49      H21B C
ATOM   7804  C   LEU  29      44.112  14.280 -13.856  1.00 47.85      H21B C
ATOM   7805  O   LEU  29      44.683  15.164 -13.214  1.00 46.44      H21B O
ATOM   7806  N   THR  30      43.004  13.681 -13.435  1.00 48.64      H21B N
ATOM   7807  CA  THR  30      42.347  14.122 -12.219  1.00 50.74      H21B C
ATOM   7808  CB  THR  30      40.859  13.743 -12.211  1.00 49.03      H21B C
ATOM   7809  OG1 THR  30      40.722  12.334 -12.389  1.00 50.94      H21B O
ATOM   7810  CG2 THR  30      40.133  14.451 -13.317  1.00 47.98      H21B C
ATOM   7811  C   THR  30      43.046  13.619 -11.027  1.00 50.98      H21B C
ATOM   7812  O   THR  30      42.609  13.668  -9.889  1.00 50.89      H21B O
ATOM   7813  N   SER  31      44.160  12.854 -11.305  1.00 53.19      H21B N
ATOM   7814  CA  SER  31      44.955  12.233 -10.263  1.00 55.07      H21B C
ATOM   7815  CB  SER  31      45.561  10.934 -10.764  1.00 56.35      H21B C
ATOM   7816  OG  SER  31      44.999   9.858 -10.045  1.00 56.82      H21B O
ATOM   7817  C   SER  31      46.063  13.126  -9.751  1.00 55.17      H21B C
ATOM   7818  O   SER  31      46.625  12.875  -8.692  1.00 53.88      H21B O
ATOM   7819  N   TYR  32      46.375  14.179 -10.493  1.00 55.54      H21B N
ATOM   7820  CA  TYR  32      47.539  14.940 -10.144  1.00 55.48      H21B C
ATOM   7821  CB  TYR  32      48.660  14.586 -11.095  1.00 60.47      H21B C
ATOM   7822  CG  TYR  32      49.372  13.363 -10.654  1.00 64.82      H21B C
ATOM   7823  CD1 TYR  32      49.287  12.192 -11.386  1.00 67.41      H21B C
ATOM   7824  CE1 TYR  32      49.872  11.059 -10.943  1.00 69.47      H21B C
ATOM   7825  CD2 TYR  32      50.075  13.358  -9.465  1.00 66.41      H21B C
ATOM   7826  CE2 TYR  32      50.664  12.236  -9.020  1.00 68.99      H21B C
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ATOM   7827  CZ  TYR    32      50.561  11.082  -9.761  1.00 70.12      H21B C
ATOM   7828  OH  TYR    32      51.172   9.941  -9.328  1.00 72.42      H21B O
ATOM   7829  C   TYR    32      47.440  16.435 -10.011  1.00 53.36      H21B C
ATOM   7830  O   TYR    32      47.609  16.964  -8.907  1.00 53.86      H21B O
ATOM   7831  N   GLY    33      47.201  17.123 -11.123  1.00 50.34      H21B N
ATOM   7832  CA  GLY    33      47.374  18.580 -11.157  1.00 46.78      H21B C
ATOM   7833  C   GLY    33      48.767  19.025 -11.621  1.00 44.17      H21B C
ATOM   7834  O   GLY    33      49.703  18.211 -11.676  1.00 45.33      H21B O
ATOM   7835  N   ILE    34      48.933  20.307 -11.936  1.00 38.23      H21B N
ATOM   7836  CA  ILE    34      50.162  20.762 -12.580  1.00 34.77      H21B C
ATOM   7837  CB  ILE    34      49.857  21.401 -13.943  1.00 32.05      H21B C
ATOM   7838  CG2 ILE    34      51.147  21.870 -14.562  1.00 33.10      H21B C
ATOM   7839  CG1 ILE    34      49.265  20.307 -14.838  1.00 32.22      H21B C
ATOM   7840  CD1 ILE    34      50.188  19.133 -15.056  1.00 32.13      H21B C
ATOM   7841  C   ILE    34      50.946  21.742 -11.623  1.00 32.84      H21B C
ATOM   7842  O   ILE    34      50.397  22.732 -11.165  1.00 32.86      H21B O
ATOM   7843  N   SER    35      52.205  21.486 -11.280  1.00 30.04      H21B N
ATOM   7844  CA  SER    35      53.030  22.582 -10.782  1.00 28.79      H21B C
ATOM   7845  CB  SER    35      54.135  22.083  -9.875  1.00 30.88      H21B C
ATOM   7846  OG  SER    35      53.665  21.644  -8.608  1.00 33.45      H21B O
ATOM   7847  C   SER    35      53.660  23.332 -11.940  1.00 29.43      H21B C
ATOM   7848  O   SER    35      53.982  22.785 -12.991  1.00 27.88      H21B O
ATOM   7849  N   TRP    36      53.810  24.630 -11.739  1.00 32.51      H21B N
ATOM   7850  CA  TRP    36      54.652  25.453 -12.620  1.00 33.33      H21B C
ATOM   7851  CB  TRP    36      53.929  26.727 -13.099  1.00 30.43      H21B C
ATOM   7852  CG  TRP    36      52.810  26.363 -13.964  1.00 27.81      H21B C
ATOM   7853  CD2 TRP    36      52.851  26.161 -15.378  1.00 25.61      H21B C
ATOM   7854  CE2 TRP    36      51.585  25.696 -15.765  1.00 25.73      H21B C
ATOM   7855  CE3 TRP    36      53.836  26.334 -16.346  1.00 24.81      H21B C
ATOM   7856  CD1 TRP    36      51.565  26.032 -13.563  1.00 25.80      H21B C
ATOM   7857  NE1 TRP    36      50.817  25.625 -14.636  1.00 25.45      H21B N
ATOM   7858  CZ2 TRP    36      51.277  25.391 -17.077  1.00 26.10      H21B C
ATOM   7859  CZ3 TRP    36      53.543  26.040 -17.636  1.00 28.16      H21B C
ATOM   7860  CH2 TRP    36      52.266  25.572 -18.002  1.00 31.41      H21B C
ATOM   7861  C   TRP    36      55.935  25.862 -11.890  1.00 34.86      H21B C
ATOM   7862  O   TRP    36      55.868  26.388 -10.769  1.00 34.06      H21B O
ATOM   7863  N   VAL    37      57.085  25.579 -12.510  1.00 35.54      H21B N
ATOM   7864  CA  VAL    37      58.381  26.082 -12.061  1.00 39.02      H21B C
ATOM   7865  CB  VAL    37      59.361  24.915 -11.867  1.00 39.94      H21B C
ATOM   7866  CG1 VAL    37      60.629  25.398 -11.198  1.00 40.01      H21B C
ATOM   7867  CG2 VAL    37      58.711  23.828 -11.077  1.00 41.21      H21B C
ATOM   7868  C   VAL    37      58.958  27.033 -13.111  1.00 40.36      H21B C
ATOM   7869  O   VAL    37      58.892  26.747 -14.306  1.00 38.98      H21B O
ATOM   7870  N   ARG    38      59.511  28.164 -12.674  1.00 42.91      H21B N
ATOM   7871  CA  ARG    38      60.403  28.931 -13.548  1.00 43.59      H21B C
ATOM   7872  CB  ARG    38      59.987  30.401 -13.614  1.00 43.48      H21B C
ATOM   7873  CG  ARG    38      59.939  31.085 -12.277  1.00 47.20      H21B C
ATOM   7874  CD  ARG    38      60.930  32.217 -12.199  1.00 51.45      H21B C
ATOM   7875  NE  ARG    38      60.691  33.069 -11.037  1.00 57.33      H21B N
ATOM   7876  CZ  ARG    38      60.035  34.226 -11.083  1.00 60.44      H21B C
ATOM   7877  NH1 ARG    38      59.550  34.676 -12.236  1.00 61.49      H21B N
ATOM   7878  NH2 ARG    38      59.861  34.932  -9.974  1.00 62.75      H21B N
ATOM   7879  C   ARG    38      61.835  28.832 -13.061  1.00 43.70      H21B C
ATOM   7880  O   ARG    38      62.080  28.633 -11.868  1.00 42.17      H21B O
ATOM   7881  N   GLN    39      62.773  28.960 -14.001  1.00 44.87      H21B N
ATOM   7882  CA  GLN    39      64.213  29.023 -13.709  1.00 45.12      H21B C
ATOM   7883  CB  GLN    39      64.818  27.653 -13.955  1.00 41.61      H21B C
ATOM   7884  CG  GLN    39      66.312  27.567 -13.851  1.00 38.69      H21B C
ATOM   7885  CD  GLN    39      66.784  26.155 -14.143  1.00 38.75      H21B C
ATOM   7886  OE1 GLN    39      66.623  25.651 -15.267  1.00 38.09      H21B O
ATOM   7887  NE2 GLN    39      67.349  25.496 -13.131  1.00 34.80      H21B N
ATOM   7888  C   GLN    39      64.898  30.069 -14.607  1.00 46.70      H21B C
ATOM   7889  O   GLN    39      65.002  29.872 -15.813  1.00 48.11      H21B O
ATOM   7890  N   ALA    40      65.354  31.178 -14.032  1.00 48.34      H21B N
ATOM   7891  CA  ALA    40      66.033  32.219 -14.821  1.00 50.36      H21B C
ATOM   7892  CB  ALA    40      66.068  33.524 -14.048  1.00 48.17      H21B C
ATOM   7893  C   ALA    40      67.449  31.802 -15.175  1.00 51.42      H21B C
ATOM   7894  O   ALA    40      68.057  31.015 -14.469  1.00 51.99      H21B O
ATOM   7895  N   PRO    41      68.008  32.353 -16.260  1.00 53.51      H21B N
ATOM   7896  CD  PRO    41      67.494  33.481 -17.055  1.00 53.56      H21B C
ATOM   7897  CA  PRO    41      69.325  31.898 -16.732  1.00 54.44      H21B C
ATOM   7898  CB  PRO    41      69.695  32.911 -17.811  1.00 53.16      H21B C
ATOM   7899  CG  PRO    41      68.385  33.467 -18.267  1.00 54.68      H21B C
ATOM   7900  C   PRO    41      70.356  31.864 -15.607  1.00 54.76      H21B C
ATOM   7901  O   PRO    41      70.466  32.808 -14.833  1.00 54.35      H21B O
ATOM   7902  N   GLY    42      71.092  30.761 -15.517  1.00 54.62      H21B N
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ATOM   7903  CA  GLY    42      72.067  30.602 -14.458  1.00 55.36      H21B C
ATOM   7904  C   GLY    42      71.510  30.605 -13.039  1.00 57.03      H21B C
ATOM   7905  O   GLY    42      72.285  30.695 -12.087  1.00 57.54      H21B O
ATOM   7906  N   GLN    43      70.190  30.499 -12.882  1.00 57.38      H21B N
ATOM   7907  CA  GLN    43      69.556  30.603 -11.564  1.00 57.89      H21B C
ATOM   7908  CB  GLN    43      68.480  31.688 -11.585  1.00 62.24      H21B C
ATOM   7909  CG  GLN    43      69.024  33.096 -11.723  1.00 70.32      H21B C
ATOM   7910  CD  GLN    43      69.786  33.529 -10.490  1.00 74.85      H21B C
ATOM   7911  OE1 GLN    43      71.014  33.621 -10.506  1.00 78.51      H21B O
ATOM   7912  NE2 GLN    43      69.059  33.790  -9.404  1.00 79.48      H21B N
ATOM   7913  C   GLN    43      68.935  29.301 -11.058  1.00 55.77      H21B C
ATOM   7914  O   GLN    43      68.991  28.269 -11.725  1.00 54.47      H21B O
ATOM   7915  N   GLY    44      68.333  29.358  -9.873  1.00 53.17      H21B N
ATOM   7916  CA  GLY    44      67.761  28.153  -9.291  1.00 51.51      H21B C
ATOM   7917  C   GLY    44      66.307  27.935  -9.655  1.00 49.20      H21B C
ATOM   7918  O   GLY    44      65.633  28.875 -10.067  1.00 50.15      H21B O
ATOM   7919  N   LEU    45      65.812  26.709  -9.512  1.00 46.56      H21B N
ATOM   7920  CA  LEU    45      64.408  26.437  -9.803  1.00 45.04      H21B C
ATOM   7921  CB  LEU    45      64.111  24.944  -9.692  1.00 41.43      H21B C
ATOM   7922  CG  LEU    45      64.932  23.987 -10.558  1.00 40.15      H21B C
ATOM   7923  CD1 LEU    45      64.566  22.569 -10.200  1.00 37.42      H21B C
ATOM   7924  CD2 LEU    45      64.680  24.240 -12.030  1.00 39.47      H21B C
ATOM   7925  C   LEU    45      63.515  27.195  -8.819  1.00 45.66      H21B C
ATOM   7926  O   LEU    45      63.787  27.230  -7.621  1.00 44.89      H21B O
ATOM   7927  N   GLU    46      62.452  27.808  -9.332  1.00 45.83      H21B N
ATOM   7928  CA  GLU    46      61.463  28.445  -8.482  1.00 45.79      H21B C
ATOM   7929  CB  GLU    46      61.532  29.950  -8.679  1.00 47.33      H21B C
ATOM   7930  CG  GLU    46      61.516  30.727  -7.380  1.00 52.65      H21B C
ATOM   7931  CD  GLU    46      62.013  32.152  -7.546  1.00 56.81      H21B C
ATOM   7932  OE1 GLU    46      61.922  32.929  -6.566  1.00 59.52      H21B O
ATOM   7933  OE2 GLU    46      62.496  32.492  -8.654  1.00 58.17      H21B O
ATOM   7934  C   GLU    46      60.043  27.937  -8.760  1.00 45.68      H21B C
ATOM   7935  O   GLU    46      59.580  27.906  -9.909  1.00 44.98      H21B O
ATOM   7936  N   TRP    47      59.344  27.539  -7.705  1.00 44.68      H21B N
ATOM   7937  CA  TRP    47      57.964  27.103  -7.860  1.00 43.70      H21B C
ATOM   7938  CB  TRP    47      57.612  26.090  -6.780  1.00 46.89      H21B C
ATOM   7939  CG  TRP    47      56.206  25.542  -6.845  1.00 50.28      H21B C
ATOM   7940  CD2 TRP    47      55.053  26.055  -6.164  1.00 51.83      H21B C
ATOM   7941  CE2 TRP    47      53.994  25.161  -6.407  1.00 51.66      H21B C
ATOM   7942  CE3 TRP    47      54.818  27.181  -5.367  1.00 53.50      H21B C
ATOM   7943  CD1 TRP    47      55.800  24.395  -7.470  1.00 47.94      H21B C
ATOM   7944  NE1 TRP    47      54.475  24.159  -7.207  1.00 50.10      H21B N
ATOM   7945  CZ2 TRP    47      52.721  25.358  -5.882  1.00 53.80      H21B C
ATOM   7946  CZ3 TRP    47      53.556  27.375  -4.849  1.00 54.46      H21B C
ATOM   7947  CH2 TRP    47      52.524  26.469  -5.105  1.00 54.87      H21B C
ATOM   7948  C   TRP    47      57.025  28.294  -7.776  1.00 41.76      H21B C
ATOM   7949  O   TRP    47      57.174  29.156  -6.914  1.00 40.43      H21B O
ATOM   7950  N   MET    48      56.054  28.319  -8.683  1.00 40.99      H21B N
ATOM   7951  CA  MET    48      55.148  29.449  -8.853  1.00 39.09      H21B C
ATOM   7952  CB  MET    48      55.137  29.903 -10.300  1.00 39.03      H21B C
ATOM   7953  CG  MET    48      56.506  30.115 -10.893  1.00 43.44      H21B C
ATOM   7954  SD  MET    48      56.347  30.353 -12.653  1.00 43.36      H21B S
ATOM   7955  CE  MET    48      56.015  32.072 -12.708  1.00 46.47      H21B C
ATOM   7956  C   MET    48      53.735  29.074  -8.480  1.00 38.15      H21B C
ATOM   7957  O   MET    48      52.944  29.929  -8.102  1.00 38.64      H21B O
ATOM   7958  N   GLY    49      53.401  27.799  -8.620  1.00 37.00      H21B N
ATOM   7959  CA  GLY    49      52.054  27.391  -8.300  1.00 35.84      H21B C
ATOM   7960  C   GLY    49      51.577  26.060  -8.833  1.00 35.29      H21B C
ATOM   7961  O   GLY    49      52.330  25.301  -9.442  1.00 33.34      H21B O
ATOM   7962  N   TRP    50      50.300  25.788  -8.584  1.00 35.95      H21B N
ATOM   7963  CA  TRP    50      49.687  24.522  -8.948  1.00 38.80      H21B C
ATOM   7964  CB  TRP    50      49.799  23.571  -7.736  1.00 40.20      H21B C
ATOM   7965  CG  TRP    50      49.008  22.290  -7.818  1.00 27.92      H21B C
ATOM   7966  CD2 TRP    50      47.607  22.144  -7.581  1.00 30.40      H21B C
ATOM   7967  CE2 TRP    50      47.296  20.770  -7.682  1.00 33.19      H21B C
ATOM   7968  CE3 TRP    50      46.572  23.046  -7.278  1.00 30.96      H21B C
ATOM   7969  CD1 TRP    50      49.491  21.028  -8.057  1.00 29.31      H21B C
ATOM   7970  NE1 TRP    50      48.471  20.108  -7.977  1.00 32.07      H21B N
ATOM   7971  CZ2 TRP    50      46.003  20.271  -7.493  1.00 31.29      H21B C
ATOM   7972  CZ3 TRP    50      45.290  22.553  -7.086  1.00 33.31      H21B C
ATOM   7973  CH2 TRP    50      45.015  21.178  -7.192  1.00 32.81      H21B C
ATOM   7974  C   TRP    50      48.221  24.759  -9.387  1.00 41.34      H21B C
ATOM   7975  O   TRP    50      47.544  25.645  -8.864  1.00 40.95      H21B O
ATOM   7976  N   VAL    51      47.766  23.996 -10.382  1.00 43.51      H21B N
ATOM   7977  CA  VAL    51      46.375  24.028 -10.841  1.00 44.71      H21B C
ATOM   7978  CB  VAL    51      46.245  24.400 -12.333  1.00 44.92      H21B C
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ATOM   7979  CG1 VAL    51      45.148  25.428 -12.516  1.00 41.58      H21B C
ATOM   7980  CG2 VAL    51      47.561  24.873 -12.877  1.00 42.14      H21B C
ATOM   7981  C   VAL    51      45.813  22.622 -10.730  1.00 46.39      H21B C
ATOM   7982  O   VAL    51      46.561  21.650 -10.773  1.00 47.07      H21B O
ATOM   7983  N   SER    52      44.491  22.520 -10.621  1.00 46.89      H21B N
ATOM   7984  CA  SER    52      43.800  21.235 -10.530  1.00 47.12      H21B C
ATOM   7985  CB  SER    52      42.812  21.250  -9.373  1.00 46.23      H21B C
ATOM   7986  OG  SER    52      42.020  20.080  -9.375  1.00 43.24      H21B O
ATOM   7987  C   SER    52      43.046  20.895 -11.802  1.00 48.99      H21B C
ATOM   7988  O   SER    52      42.243  21.684 -12.297  1.00 47.79      H21B O
ATOM   7989  N   PHE    53      43.297  19.708 -12.329  1.00 51.06      H21B N
ATOM   7990  CA  PHE    53      42.536  19.241 -13.468  1.00 51.62      H21B C
ATOM   7991  CB  PHE    53      43.238  18.050 -14.110  1.00 49.15      H21B C
ATOM   7992  CG  PHE    53      43.992  18.401 -15.352  1.00 47.00      H21B C
ATOM   7993  CD1 PHE    53      45.359  18.196 -15.433  1.00 45.85      H21B C
ATOM   7994  CD2 PHE    53      43.330  18.930 -16.444  1.00 45.21      H21B C
ATOM   7995  CE1 PHE    53      46.046  18.511 -16.582  1.00 44.88      H21B C
ATOM   7996  CE2 PHE    53      44.015  19.250 -17.599  1.00 44.11      H21B C
ATOM   7997  CZ  PHE    53      45.372  19.038 -17.668  1.00 43.18      H21B C
ATOM   7998  C   PHE    53      41.158  18.839 -12.986  1.00 52.99      H21B C
ATOM   7999  O   PHE    53      40.161  19.146 -13.620  1.00 51.55      H21B O
ATOM   8000  N   TYR    54      41.119  18.160 -11.848  1.00 55.64      H21B N
ATOM   8001  CA  TYR    54      39.872  17.785 -11.202  1.00 58.44      H21B C
ATOM   8002  CB  TYR    54      40.162  17.329  -9.777  1.00 54.87      H21B C
ATOM   8003  CG  TYR    54      38.949  16.897  -9.001  1.00 51.36      H21B C
ATOM   8004  CD1 TYR    54      38.655  17.477  -7.774  1.00 48.87      H21B C
ATOM   8005  CE1 TYR    54      37.555  17.104  -7.058  1.00 49.07      H21B C
ATOM   8006  CD2 TYR    54      38.096  15.917  -9.494  1.00 47.97      H21B C
ATOM   8007  CE2 TYR    54      36.990  15.535  -8.786  1.00 47.93      H21B C
ATOM   8008  CZ  TYR    54      36.720  16.132  -7.563  1.00 49.35      H21B C
ATOM   8009  OH  TYR    54      35.618  15.755  -6.836  1.00 48.60      H21B O
ATOM   8010  C   TYR    54      38.835  18.912 -11.192  1.00 62.29      H21B C
ATOM   8011  O   TYR    54      37.946  18.949 -12.039  1.00 63.00      H21B O
ATOM   8012  N   ASN    55      38.923  19.827 -10.236  1.00 67.02      H21B N
ATOM   8013  CA  ASN    55      38.057  20.994 -10.293  1.00 71.09      H21B C
ATOM   8014  CB  ASN    55      37.730  21.518  -8.894  1.00 68.84      H21B C
ATOM   8015  CG  ASN    55      38.841  21.307  -7.921  1.00 67.93      H21B C
ATOM   8016  OD1 ASN    55      38.616  20.676  -6.797  1.00 67.67      H21B O
ATOM   8017  ND2 ASN    55      40.054  21.613  -8.338  1.00 69.63      H21B N
ATOM   8018  C   ASN    55      38.717  22.079 -11.118  1.00 74.37      H21B C
ATOM   8019  O   ASN    55      38.675  22.051 -12.347  1.00 75.61      H21B O
ATOM   8020  N   GLY    56      39.336  23.034 -10.449  1.00 76.48      H21B N
ATOM   8021  CA  GLY    56      40.142  23.993 -11.163  1.00 78.11      H21B C
ATOM   8022  C   GLY    56      40.842  24.836 -10.142  1.00 80.92      H21B C
ATOM   8023  O   GLY    56      41.260  25.945 -10.426  1.00 81.59      H21B O
ATOM   8024  N   ASN    57      40.959  24.307  -8.935  1.00 83.69      H21B N
ATOM   8025  CA  ASN    57      41.543  25.062  -7.846  1.00 84.39      H21B C
ATOM   8026  CB  ASN    57      41.483  24.249  -6.559  1.00 88.07      H21B C
ATOM   8027  CG  ASN    57      40.069  23.968  -6.121  1.00 90.83      H21B C
ATOM   8028  OD1 ASN    57      39.142  24.677  -6.501  1.00 92.97      H21B O
ATOM   8029  ND2 ASN    57      39.892  22.927  -5.319  1.00 92.57      H21B N
ATOM   8030  C   ASN    57      42.978  25.426  -8.172  1.00 81.95      H21B C
ATOM   8031  O   ASN    57      43.665  24.701  -8.883  1.00 82.38      H21B O
ATOM   8032  N   THR    58      43.422  26.566  -7.663  1.00 77.98      H21B N
ATOM   8033  CA  THR    58      44.770  27.034  -7.924  1.00 74.72      H21B C
ATOM   8034  CB  THR    58      44.751  28.353  -8.715  1.00 76.41      H21B C
ATOM   8035  OG1 THR    58      43.878  29.282  -8.070  1.00 77.32      H21B O
ATOM   8036  CG2 THR    58      44.265  28.121 -10.131  1.00 77.44      H21B C
ATOM   8037  C   THR    58      45.492  27.240  -6.609  1.00 71.58      H21B C
ATOM   8038  O   THR    58      44.864  27.368  -5.569  1.00 70.30      H21B O
ATOM   8039  N   ASN    59      46.817  27.242  -6.649  1.00 69.82      H21B N
ATOM   8040  CA  ASN    59      47.611  27.653  -5.500  1.00 68.25      H21B C
ATOM   8041  CB  ASN    59      48.109  26.428  -4.740  1.00 65.64      H21B C
ATOM   8042  CG  ASN    59      48.845  26.787  -3.473  1.00 64.96      H21B C
ATOM   8043  OD1 ASN    59      48.733  27.899  -2.966  1.00 65.43      H21B O
ATOM   8044  ND2 ASN    59      49.609  25.840  -2.953  1.00 65.84      H21B N
ATOM   8045  C   ASN    59      48.780  28.460  -6.034  1.00 68.81      H21B C
ATOM   8046  O   ASN    59      49.547  27.972  -6.851  1.00 68.36      H21B O
ATOM   8047  N   TYR    60      48.899  29.704  -5.588  1.00 70.01      H21B N
ATOM   8048  CA  TYR    60      49.915  30.607  -6.115  1.00 70.19      H21B C
ATOM   8049  CB  TYR    60      49.286  31.940  -6.547  1.00 64.48      H21B C
ATOM   8050  CG  TYR    60      48.377  31.831  -7.747  1.00 56.84      H21B C
ATOM   8051  CD1 TYR    60      48.894  31.719  -9.029  1.00 54.39      H21B C
ATOM   8052  CE1 TYR    60      48.061  31.542 -10.120  1.00 49.77      H21B C
ATOM   8053  CD2 TYR    60      47.005  31.776  -7.594  1.00 52.06      H21B C
ATOM   8054  CE2 TYR    60      46.176  31.602  -8.672  1.00 47.45      H21B C
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ATOM   8055  CZ  TYR    60      46.706  31.481  -9.930  1.00 47.20      H21B C
ATOM   8056  OH  TYR    60      45.873  31.262 -10.999  1.00 45.04      H21B O
ATOM   8057  C   TYR    60      51.011  30.876  -5.101  1.00 73.82      H21B C
ATOM   8058  O   TYR    60      50.773  30.873  -3.899  1.00 74.53      H21B O
ATOM   8059  N   ALA    61      52.216  31.105  -5.608  1.00 79.37      H21B N
ATOM   8060  CA  ALA    61      53.339  31.530  -4.791  1.00 86.15      H21B C
ATOM   8061  CB  ALA    61      54.629  31.211  -5.508  1.00 79.56      H21B C
ATOM   8062  C   ALA    61      53.236  33.031  -4.518  1.00 93.30      H21B C
ATOM   8063  O   ALA    61      52.501  33.743  -5.194  1.00 94.02      H21B O
ATOM   8064  N   GLN    62      53.972  33.509  -3.523  1.00102.91      H21B N
ATOM   8065  CA  GLN    62      53.978  34.930  -3.199  1.00111.50      H21B C
ATOM   8066  CB  GLN    62      55.022  35.221  -2.123  1.00113.10      H21B C
ATOM   8067  CG  GLN    62      54.598  36.265  -1.105  1.00114.99      H21B C
ATOM   8068  CD  GLN    62      53.632  35.705  -0.080  1.00115.65      H21B C
ATOM   8069  OE1 GLN    62      53.671  36.073   1.096  1.00115.70      H21B O
ATOM   8070  NE2 GLN    62      52.756  34.805  -0.521  1.00116.17      H21B N
ATOM   8071  C   GLN    62      54.282  35.778  -4.426  1.00115.83      H21B C
ATOM   8072  O   GLN    62      53.402  36.445  -4.953  1.00118.69      H21B O
ATOM   8073  N   LYS    63      55.532  35.745  -4.877  1.00120.29      H21B N
ATOM   8074  CA  LYS    63      55.975  36.610  -5.965  1.00123.83      H21B C
ATOM   8075  CB  LYS    63      57.238  36.042  -6.623  1.00126.62      H21B C
ATOM   8076  CG  LYS    63      58.518  36.271  -5.828  1.00129.64      H21B C
ATOM   8077  CD  LYS    63      59.760  36.088  -6.696  1.00132.38      H21B C
ATOM   8078  CE  LYS    63      61.026  36.494  -5.951  1.00134.60      H21B C
ATOM   8079  NZ  LYS    63      62.235  36.420  -6.816  1.00135.68      H21B N
ATOM   8080  C   LYS    63      54.892  36.781  -7.018  1.00125.33      H21B C
ATOM   8081  O   LYS    63      54.362  37.876  -7.210  1.00125.28      H21B O
ATOM   8082  N   LEU    64      54.558  35.691  -7.695  1.00127.68      H21B N
ATOM   8083  CA  LEU    64      53.576  35.745  -8.764  1.00130.37      H21B C
ATOM   8084  CB  LEU    64      53.597  34.444  -9.568  1.00132.98      H21B C
ATOM   8085  CG  LEU    64      54.806  34.276 -10.494  1.00136.29      H21B C
ATOM   8086  CD1 LEU    64      54.818  35.403 -11.518  1.00138.41      H21B C
ATOM   8087  CD2 LEU    64      56.100  34.275  -9.685  1.00138.56      H21B C
ATOM   8088  C   LEU    64      52.187  35.983  -8.198  1.00130.44      H21B C
ATOM   8089  O   LEU    64      51.278  35.176  -8.400  1.00130.29      H21B O
ATOM   8090  N   GLN    65      52.027  37.100  -7.489  1.00129.19      H21B N
ATOM   8091  CA  GLN    65      50.749  37.424  -6.870  1.00127.17      H21B C
ATOM   8092  CB  GLN    65      50.948  37.952  -5.451  1.00131.48      H21B C
ATOM   8093  CG  GLN    65      49.744  37.693  -4.556  1.00137.16      H21B C
ATOM   8094  CD  GLN    65      49.179  36.287  -4.742  1.00140.29      H21B C
ATOM   8095  OE1 GLN    65      47.964  36.082  -4.713  1.00142.41      H21B O
ATOM   8096  NE2 GLN    65      50.064  35.313  -4.936  1.00142.71      H21B N
ATOM   8097  C   GLN    65      49.958  38.437  -7.676  1.00121.86      H21B C
ATOM   8098  O   GLN    65      50.422  39.546  -7.930  1.00123.39      H21B O
ATOM   8099  N   GLY    66      48.750  38.041  -8.059  1.00114.88      H21B N
ATOM   8100  CA  GLY    66      47.966  38.820  -8.993  1.00104.98      H21B C
ATOM   8101  C   GLY    66      48.484  38.624 -10.400  1.00 96.86      H21B C
ATOM   8102  O   GLY    66      47.711  38.410 -11.334  1.00 97.14      H21B O
ATOM   8103  N   ARG    67      49.806  38.681 -10.538  1.00 88.00      H21B N
ATOM   8104  CA  ARG    67      50.467  38.666 -11.838  1.00 78.28      H21B C
ATOM   8105  CB  ARG    67      51.980  38.806 -11.662  1.00 79.16      H21B C
ATOM   8106  CG  ARG    67      52.424  40.093 -11.021  1.00 78.42      H21B C
ATOM   8107  CD  ARG    67      53.594  40.703 -11.775  1.00 76.25      H21B C
ATOM   8108  NE  ARG    67      54.808  39.901 -11.670  1.00 70.42      H21B N
ATOM   8109  CZ  ARG    67      55.549  39.532 -12.711  1.00 67.25      H21B C
ATOM   8110  NH1 ARG    67      55.197  39.891 -13.941  1.00 63.82      H21B N
ATOM   8111  NH2 ARG    67      56.643  38.804 -12.522  1.00 62.44      H21B N
ATOM   8112  C   ARG    67      50.183  37.376 -12.587  1.00 72.34      H21B C
ATOM   8113  O   ARG    67      50.251  37.328 -13.818  1.00 67.98      H21B O
ATOM   8114  N   GLY    68      49.879  36.327 -11.831  1.00 67.36      H21B N
ATOM   8115  CA  GLY    68      49.791  35.003 -12.411  1.00 61.33      H21B C
ATOM   8116  C   GLY    68      48.384  34.459 -12.429  1.00 57.15      H21B C
ATOM   8117  O   GLY    68      47.615  34.658 -11.487  1.00 54.96      H21B O
ATOM   8118  N   THR    69      48.058  33.773 -13.519  1.00 52.86      H21B N
ATOM   8119  CA  THR    69      46.802  33.040 -13.640  1.00 49.43      H21B C
ATOM   8120  CB  THR    69      45.755  33.817 -14.466  1.00 47.25      H21B C
ATOM   8121  OG1 THR    69      45.372  35.009 -13.771  1.00 46.17      H21B O
ATOM   8122  CG2 THR    69      44.536  32.955 -14.699  1.00 43.40      H21B C
ATOM   8123  C   THR    69      47.002  31.698 -14.325  1.00 46.56      H21B C
ATOM   8124  O   THR    69      47.148  31.639 -15.539  1.00 46.53      H21B O
ATOM   8125  N   MET    70      46.997  30.618 -13.558  1.00 44.10      H21B N
ATOM   8126  CA  MET    70      46.975  29.311 -14.185  1.00 42.39      H21B C
ATOM   8127  CB  MET    70      47.990  28.360 -13.551  1.00 41.28      H21B C
ATOM   8128  CG  MET    70      48.171  28.515 -12.084  1.00 40.21      H21B C
ATOM   8129  SD  MET    70      49.850  28.066 -11.687  1.00 39.87      H21B S
ATOM   8130  CE  MET    70      49.867  28.560 -10.019  1.00 44.15      H21B C
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ATOM   8131  C   MET   70      45.613  28.668 -14.181  1.00 40.27      H21B C
ATOM   8132  O   MET   70      44.851  28.780 -13.230  1.00 40.48      H21B O
ATOM   8133  N   THR   71      45.329  27.981 -15.273  1.00 40.53      H21B N
ATOM   8134  CA  THR   71      44.026  27.413 -15.512  1.00 41.99      H21B C
ATOM   8135  CB  THR   71      43.219  28.301 -16.470  1.00 41.68      H21B C
ATOM   8136  OG1 THR   71      43.930  28.448 -17.710  1.00 40.43      H21B O
ATOM   8137  CG2 THR   71      42.978  29.653 -15.841  1.00 39.62      H21B C
ATOM   8138  C   THR   71      44.154  26.021 -16.112  1.00 43.75      H21B C
ATOM   8139  O   THR   71      45.248  25.511 -16.296  1.00 41.50      H21B O
ATOM   8140  N   THR   72      43.015  25.422 -16.422  1.00 48.73      H21B N
ATOM   8141  CA  THR   72      42.960  24.041 -16.873  1.00 53.25      H21B C
ATOM   8142  CB  THR   72      42.802  23.107 -15.652  1.00 52.17      H21B C
ATOM   8143  OG1 THR   72      44.095  22.675 -15.216  1.00 53.13      H21B O
ATOM   8144  CG2 THR   72      41.942  21.918 -15.981  1.00 53.23      H21B C
ATOM   8145  C   THR   72      41.803  23.831 -17.865  1.00 55.94      H21B C
ATOM   8146  O   THR   72      40.699  24.344 -17.663  1.00 56.70      H21B O
ATOM   8147  N   ASP   73      42.083  23.113 -18.954  1.00 57.69      H21B N
ATOM   8148  CA  ASP   73      41.046  22.553 -19.811  1.00 58.33      H21B C
ATOM   8149  CB  ASP   73      41.142  23.100 -21.238  1.00 61.28      H21B C
ATOM   8150  CG  ASP   73      40.250  22.338 -22.218  1.00 63.71      H21B C
ATOM   8151  OD1 ASP   73      40.367  22.558 -23.442  1.00 64.93      H21B O
ATOM   8152  OD2 ASP   73      39.429  21.513 -21.768  1.00 64.36      H21B O
ATOM   8153  C   ASP   73      41.216  21.048 -19.849  1.00 57.27      H21B C
ATOM   8154  O   ASP   73      42.090  20.523 -20.529  1.00 58.48      H21B O
ATOM   8155  N   PRO   74      40.375  20.327 -19.110  1.00 56.40      H21B N
ATOM   8156  CD  PRO   74      39.411  20.780 -18.097  1.00 55.90      H21B C
ATOM   8157  CA  PRO   74      40.518  18.875 -19.105  1.00 56.74      H21B C
ATOM   8158  CB  PRO   74      39.543  18.424 -18.015  1.00 55.19      H21B C
ATOM   8159  CG  PRO   74      39.364  19.611 -17.157  1.00 53.64      H21B C
ATOM   8160  C   PRO   74      40.207  18.253 -20.460  1.00 56.88      H21B C
ATOM   8161  O   PRO   74      40.761  17.220 -20.816  1.00 57.34      H21B O
ATOM   8162  N   SER   75      39.320  18.874 -21.221  1.00 57.92      H21B N
ATOM   8163  CA  SER   75      38.854  18.220 -22.428  1.00 58.36      H21B C
ATOM   8164  CB  SER   75      37.682  18.978 -23.039  1.00 57.07      H21B C
ATOM   8165  OG  SER   75      38.122  20.173 -23.637  1.00 57.34      H21B O
ATOM   8166  C   SER   75      39.989  18.122 -23.426  1.00 58.17      H21B C
ATOM   8167  O   SER   75      40.015  17.212 -24.251  1.00 60.52      H21B O
ATOM   8168  N   THR   76      40.934  19.055 -23.342  1.00 56.65      H21B N
ATOM   8169  CA  THR   76      42.126  19.003 -24.184  1.00 54.54      H21B C
ATOM   8170  CB  THR   76      42.418  20.352 -24.850  1.00 55.46      H21B C
ATOM   8171  OG1 THR   76      42.689  21.336 -23.846  1.00 57.17      H21B O
ATOM   8172  CG2 THR   76      41.239  20.791 -25.679  1.00 55.89      H21B C
ATOM   8173  C   THR   76      43.378  18.586 -23.419  1.00 52.21      H21B C
ATOM   8174  O   THR   76      44.479  18.660 -23.955  1.00 51.31      H21B O
ATOM   8175  N   SER   77      43.205  18.150 -22.175  1.00 50.10      H21B N
ATOM   8176  CA  SER   77      44.314  17.651 -21.369  1.00 48.00      H21B C
ATOM   8177  CB  SER   77      44.801  16.302 -21.905  1.00 48.42      H21B C
ATOM   8178  OG  SER   77      44.105  15.938 -23.088  1.00 50.53      H21B O
ATOM   8179  C   SER   77      45.451  18.660 -21.383  1.00 46.50      H21B C
ATOM   8180  O   SER   77      46.607  18.312 -21.612  1.00 45.85      H21B O
ATOM   8181  N   THR   78      45.100  19.916 -21.128  1.00 44.26      H21B N
ATOM   8182  CA  THR   78      46.027  21.026 -21.253  1.00 42.27      H21B C
ATOM   8183  CB  THR   78      45.777  21.774 -22.577  1.00 43.01      H21B C
ATOM   8184  OG1 THR   78      46.092  20.906 -23.666  1.00 42.58      H21B O
ATOM   8185  CG2 THR   78      46.641  23.019 -22.676  1.00 41.56      H21B C
ATOM   8186  C   THR   78      45.909  22.007 -20.084  1.00 41.93      H21B C
ATOM   8187  O   THR   78      44.816  22.455 -19.739  1.00 41.80      H21B O
ATOM   8188  N   ALA   79      47.046  22.336 -19.477  1.00 41.30      H21B N
ATOM   8189  CA  ALA   79      47.082  23.328 -18.415  1.00 43.41      H21B C
ATOM   8190  CB  ALA   79      47.823  22.778 -17.197  1.00 39.93      H21B C
ATOM   8191  C   ALA   79      47.791  24.559 -18.944  1.00 46.13      H21B C
ATOM   8192  O   ALA   79      48.574  24.466 -19.899  1.00 46.64      H21B O
ATOM   8193  N   TYR   80      47.521  25.711 -18.337  1.00 47.82      H21B N
ATOM   8194  CA  TYR   80      48.086  26.950 -18.843  1.00 50.58      H21B C
ATOM   8195  CB  TYR   80      47.013  27.771 -19.561  1.00 51.23      H21B C
ATOM   8196  CG  TYR   80      46.386  27.079 -20.743  1.00 51.15      H21B C
ATOM   8197  CD1 TYR   80      45.311  26.214 -20.572  1.00 51.00      H21B C
ATOM   8198  CE1 TYR   80      44.719  25.589 -21.650  1.00 52.73      H21B C
ATOM   8199  CD2 TYR   80      46.856  27.303 -22.034  1.00 51.74      H21B C
ATOM   8200  CE2 TYR   80      46.268  26.687 -23.126  1.00 53.17      H21B C
ATOM   8201  CZ  TYR   80      45.200  25.828 -22.928  1.00 55.28      H21B C
ATOM   8202  OH  TYR   80      44.619  25.186 -24.006  1.00 58.32      H21B O
ATOM   8203  C   TYR   80      48.731  27.812 -17.772  1.00 51.42      H21B C
ATOM   8204  O   TYR   80      48.313  27.831 -16.614  1.00 50.53      H21B O
ATOM   8205  N   MET   81      49.755  28.537 -18.186  1.00 53.11      H21B N
ATOM   8206  CA  MET   81      50.388  29.514 -17.331  1.00 56.14      H21B C
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ATOM   8207  CB  MET    81      51.833  29.080 -17.073  1.00 57.65      H21B C
ATOM   8208  CG  MET    81      52.753  30.176 -16.565  1.00 58.66      H21B C
ATOM   8209  SD  MET    81      52.254  30.897 -14.991  1.00 57.87      H21B S
ATOM   8210  CE  MET    81      53.575  32.092 -14.801  1.00 54.87      H21B C
ATOM   8211  C   MET    81      50.336  30.872 -18.029  1.00 58.01      H21B C
ATOM   8212  O   MET    81      51.015  31.089 -19.033  1.00 56.65      H21B O
ATOM   8213  N   GLU    82      49.514  31.781 -17.514  1.00 61.28      H21B N
ATOM   8214  CA  GLU    82      49.511  33.149 -18.018  1.00 65.13      H21B C
ATOM   8215  CB  GLU    82      48.100  33.623 -18.342  1.00 68.11      H21B C
ATOM   8216  CG  GLU    82      48.049  35.126 -18.580  1.00 75.01      H21B C
ATOM   8217  CD  GLU    82      46.699  35.614 -19.041  1.00 78.54      H21B C
ATOM   8218  OE1 GLU    82      46.537  36.847 -19.191  1.00 80.65      H21B O
ATOM   8219  OE2 GLU    82      45.802  34.770 -19.253  1.00 85.32      H21B O
ATOM   8220  C   GLU    82      50.126  34.125 -17.035  1.00 65.78      H21B C
ATOM   8221  O   GLU    82      49.752  34.156 -15.866  1.00 64.06      H21B O
ATOM   8222  N   LEU    83      51.063  34.928 -17.532  1.00 68.70      H21B N
ATOM   8223  CA  LEU    83      51.749  35.920 -16.718  1.00 72.23      H21B C
ATOM   8224  CB  LEU    83      53.227  35.574 -16.610  1.00 69.70      H21B C
ATOM   8225  CG  LEU    83      53.852  35.648 -15.223  1.00 67.44      H21B C
ATOM   8226  CD1 LEU    83      55.336  35.825 -15.388  1.00 66.44      H21B C
ATOM   8227  CD2 LEU    83      53.273  36.788 -14.430  1.00 65.57      H21B C
ATOM   8228  C   LEU    83      51.601  37.309 -17.330  1.00 75.95      H21B C
ATOM   8229  O   LEU    83      51.904  37.512 -18.504  1.00 75.65      H21B O
ATOM   8230  N   ARG    84      51.147  38.261 -16.520  1.00 80.89      H21B N
ATOM   8231  CA  ARG    84      50.822  39.601 -16.992  1.00 84.83      H21B C
ATOM   8232  CB  ARG    84      49.499  40.064 -16.380  1.00 90.56      H21B C
ATOM   8233  CG  ARG    84      48.344  39.114 -16.614  1.00100.33      H21B C
ATOM   8234  CD  ARG    84      47.315  39.202 -15.499  1.00109.68      H21B C
ATOM   8235  NE  ARG    84      46.461  38.020 -15.475  1.00120.80      H21B N
ATOM   8236  CZ  ARG    84      45.558  37.737 -16.408  1.00127.57      H21B C
ATOM   8237  NH1 ARG    84      44.822  36.638 -16.309  1.00132.02      H21B N
ATOM   8238  NH2 ARG    84      45.389  38.556 -17.439  1.00132.38      H21B N
ATOM   8239  C   ARG    84      51.914  40.585 -16.612  1.00 83.70      H21B C
ATOM   8240  O   ARG    84      52.734  40.303 -15.739  1.00 84.67      H21B O
ATOM   8241  N   SER    85      51.916  41.741 -17.272  1.00 81.90      H21B N
ATOM   8242  CA  SER    85      52.799  42.840 -16.896  1.00 79.14      H21B C
ATOM   8243  CB  SER    85      52.363  43.449 -15.557  1.00 78.89      H21B C
ATOM   8244  OG  SER    85      51.028  43.919 -15.602  1.00 75.63      H21B O
ATOM   8245  C   SER    85      54.221  42.330 -16.772  1.00 77.66      H21B C
ATOM   8246  O   SER    85      54.863  42.496 -15.733  1.00 77.42      H21B O
ATOM   8247  N   LEU    86      54.704  41.701 -17.838  1.00 75.99      H21B N
ATOM   8248  CA  LEU    86      56.022  41.085 -17.836  1.00 74.39      H21B C
ATOM   8249  CB  LEU    86      56.258  40.346 -19.157  1.00 71.64      H21B C
ATOM   8250  CG  LEU    86      55.451  39.062 -19.373  1.00 69.27      H21B C
ATOM   8251  CD1 LEU    86      55.543  38.610 -20.823  1.00 67.28      H21B C
ATOM   8252  CD2 LEU    86      55.969  37.989 -18.435  1.00 67.92      H21B C
ATOM   8253  C   LEU    86      57.126  42.111 -17.616  1.00 74.83      H21B C
ATOM   8254  O   LEU    86      57.262  43.060 -18.378  1.00 73.84      H21B O
ATOM   8255  N   ARG    87      57.898  41.916 -16.554  1.00 76.20      H21B N
ATOM   8256  CA  ARG    87      59.179  42.585 -16.402  1.00 77.36      H21B C
ATOM   8257  CB  ARG    87      59.621  42.614 -14.939  1.00 78.84      H21B C
ATOM   8258  CG  ARG    87      58.580  43.085 -13.960  1.00 81.26      H21B C
ATOM   8259  CD  ARG    87      59.149  43.110 -12.555  1.00 83.79      H21B C
ATOM   8260  NE  ARG    87      58.087  43.040 -11.560  1.00 86.95      H21B N
ATOM   8261  CZ  ARG    87      57.869  41.994 -10.775  1.00 89.17      H21B C
ATOM   8262  NH1 ARG    87      56.870  42.022  -9.901  1.00 90.14      H21B N
ATOM   8263  NH2 ARG    87      58.655  40.925 -10.858  1.00 90.65      H21B N
ATOM   8264  C   ARG    87      60.208  41.797 -17.182  1.00 77.52      H21B C
ATOM   8265  O   ARG    87      59.898  40.774 -17.787  1.00 78.03      H21B O
ATOM   8266  N   SER    88      61.442  42.278 -17.143  1.00 77.89      H21B N
ATOM   8267  CA  SER    88      62.579  41.526 -17.647  1.00 76.92      H21B C
ATOM   8268  CB  SER    88      63.693  42.497 -18.027  1.00 76.95      H21B C
ATOM   8269  OG  SER    88      63.875  43.471 -17.016  1.00 76.63      H21B O
ATOM   8270  C   SER    88      63.076  40.519 -16.600  1.00 76.23      H21B C
ATOM   8271  O   SER    88      63.913  39.673 -16.902  1.00 75.13      H21B O
ATOM   8272  N   ASP    89      62.549  40.617 -15.377  1.00 76.04      H21B N
ATOM   8273  CA  ASP    89      62.825  39.646 -14.307  1.00 74.28      H21B C
ATOM   8274  CB  ASP    89      62.387  40.196 -12.941  1.00 78.49      H21B C
ATOM   8275  CG  ASP    89      62.941  41.586 -12.649  1.00 82.47      H21B C
ATOM   8276  OD1 ASP    89      62.171  42.451 -12.167  1.00 84.19      H21B O
ATOM   8277  OD2 ASP    89      64.143  41.816 -12.890  1.00 84.83      H21B O
ATOM   8278  C   ASP    89      62.073  38.334 -14.552  1.00 70.50      H21B C
ATOM   8279  O   ASP    89      62.232  37.372 -13.807  1.00 69.35      H21B O
ATOM   8280  N   ASP    90      61.240  38.321 -15.589  1.00 66.65      H21B N
ATOM   8281  CA  ASP    90      60.393  37.176 -15.916  1.00 61.73      H21B C
ATOM   8282  CB  ASP    90      59.011  37.644 -16.368  1.00 61.77      H21B C
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ATOM   8283  CG  ASP    90      58.318  38.488 -15.329  1.00 61.61      H21B C
ATOM   8284  OD1 ASP    90      58.704  38.408 -14.145  1.00 62.01      H21B O
ATOM   8285  OD2 ASP    90      57.385  39.235 -15.696  1.00 62.33      H21B O
ATOM   8286  C   ASP    90      61.026  36.382 -17.033  1.00 57.28      H21B C
ATOM   8287  O   ASP    90      60.504  35.356 -17.451  1.00 57.29      H21B O
ATOM   8288  N   THR    91      62.145  36.873 -17.533  1.00 53.21      H21B N
ATOM   8289  CA  THR    91      62.907  36.114 -18.499  1.00 50.91      H21B C
ATOM   8290  CB  THR    91      64.019  36.957 -19.130  1.00 51.25      H21B C
ATOM   8291  OG1 THR    91      63.440  37.861 -20.079  1.00 52.40      H21B O
ATOM   8292  CG2 THR    91      65.025  36.065 -19.834  1.00 49.86      H21B C
ATOM   8293  C   THR    91      63.526  34.900 -17.832  1.00 47.55      H21B C
ATOM   8294  O   THR    91      64.316  35.024 -16.892  1.00 46.34      H21B O
ATOM   8295  N   ALA    92      63.156  33.726 -18.334  1.00 43.20      H21B N
ATOM   8296  CA  ALA    92      63.539  32.482 -17.700  1.00 40.41      H21B C
ATOM   8297  CB  ALA    92      63.041  32.459 -16.267  1.00 40.90      H21B C
ATOM   8298  C   ALA    92      62.990  31.292 -18.456  1.00 37.21      H21B C
ATOM   8299  O   ALA    92      62.242  31.430 -19.415  1.00 38.12      H21B O
ATOM   8300  N   VAL    93      63.380  30.109 -18.027  1.00 34.64      H21B N
ATOM   8301  CA  VAL    93      62.733  28.919 -18.513  1.00 32.99      H21B C
ATOM   8302  CB  VAL    93      63.750  27.793 -18.666  1.00 30.83      H21B C
ATOM   8303  CG1 VAL    93      63.044  26.455 -18.927  1.00 26.30      H21B C
ATOM   8304  CG2 VAL    93      64.672  28.142 -19.822  1.00 25.24      H21B C
ATOM   8305  C   VAL    93      61.595  28.521 -17.574  1.00 31.86      H21B C
ATOM   8306  O   VAL    93      61.729  28.533 -16.348  1.00 29.01      H21B O
ATOM   8307  N   TYR    94      60.458  28.214 -18.180  1.00 32.50      H21B N
ATOM   8308  CA  TYR    94      59.251  27.901 -17.441  1.00 34.02      H21B C
ATOM   8309  CB  TYR    94      58.111  28.813 -17.900  1.00 31.77      H21B C
ATOM   8310  CG  TYR    94      58.243  30.236 -17.393  1.00 30.96      H21B C
ATOM   8311  CD1 TYR    94      57.570  30.650 -16.240  1.00 29.84      H21B C
ATOM   8312  CE1 TYR    94      57.696  31.944 -15.754  1.00 27.78      H21B C
ATOM   8313  CD2 TYR    94      59.047  31.164 -18.049  1.00 28.53      H21B C
ATOM   8314  CE2 TYR    94      59.179  32.472 -17.564  1.00 28.29      H21B C
ATOM   8315  CZ  TYR    94      58.503  32.852 -16.412  1.00 28.76      H21B C
ATOM   8316  OH  TYR    94      58.659  34.126 -15.884  1.00 29.65      H21B O
ATOM   8317  C   TYR    94      58.907  26.443 -17.681  1.00 33.72      H21B C
ATOM   8318  O   TYR    94      58.750  26.005 -18.821  1.00 34.33      H21B O
ATOM   8319  N   TYR    95      58.824  25.694 -16.591  1.00 35.12      H21B N
ATOM   8320  CA  TYR    95      58.441  24.289 -16.634  1.00 35.54      H21B C
ATOM   8321  CB  TYR    95      59.442  23.437 -15.862  1.00 37.61      H21B C
ATOM   8322  CG  TYR    95      60.852  23.484 -16.396  1.00 39.36      H21B C
ATOM   8323  CD1 TYR    95      61.187  22.835 -17.578  1.00 40.65      H21B C
ATOM   8324  CE1 TYR    95      62.476  22.841 -18.055  1.00 40.50      H21B C
ATOM   8325  CD2 TYR    95      61.856  24.146 -15.703  1.00 39.61      H21B C
ATOM   8326  CE2 TYR    95      63.155  24.155 -16.173  1.00 40.63      H21B C
ATOM   8327  CZ  TYR    95      63.460  23.502 -17.351  1.00 40.83      H21B C
ATOM   8328  OH  TYR    95      64.750  23.515 -17.834  1.00 41.60      H21B O
ATOM   8329  C   TYR    95      57.066  24.073 -16.024  1.00 34.62      H21B C
ATOM   8330  O   TYR    95      56.654  24.772 -15.087  1.00 29.49      H21B O
ATOM   8331  N   CYS    96      56.358  23.095 -16.570  1.00 36.69      H21B N
ATOM   8332  CA  CYS    96      55.310  22.427 -15.825  1.00 39.27      H21B C
ATOM   8333  C   CYS    96      55.813  21.042 -15.431  1.00 40.20      H21B C
ATOM   8334  O   CYS    96      56.535  20.390 -16.182  1.00 38.24      H21B O
ATOM   8335  CB  CYS    96      54.026  22.325 -16.657  1.00 38.82      H21B C
ATOM   8336  SG  CYS    96      54.180  21.440 -18.241  1.00 47.25      H21B S
ATOM   8337  N   ALA    97      55.453  20.620 -14.229  1.00 42.79      H21B N
ATOM   8338  CA  ALA    97      55.651  19.248 -13.802  1.00 45.99      H21B C
ATOM   8339  CB  ALA    97      56.701  19.189 -12.703  1.00 47.19      H21B C
ATOM   8340  C   ALA    97      54.320  18.724 -13.281  1.00 48.49      H21B C
ATOM   8341  O   ALA    97      53.412  19.504 -12.983  1.00 46.95      H21B O
ATOM   8342  N   ARG    98      54.207  17.401 -13.187  1.00 51.41      H21B N
ATOM   8343  CA  ARG    98      53.047  16.760 -12.588  1.00 53.37      H21B C
ATOM   8344  CB  ARG    98      53.046  15.269 -12.932  1.00 51.25      H21B C
ATOM   8345  CG  ARG    98      51.663  14.667 -13.187  1.00 49.60      H21B C
ATOM   8346  CD  ARG    98      51.753  13.294 -13.836  1.00 45.00      H21B C
ATOM   8347  NE  ARG    98      52.484  12.364 -12.984  1.00 44.76      H21B N
ATOM   8348  CZ  ARG    98      52.421  11.041 -13.085  1.00 43.23      H21B C
ATOM   8349  NH1 ARG    98      51.657  10.481 -14.007  1.00 44.12      H21B N
ATOM   8350  NH2 ARG    98      53.126  10.279 -12.262  1.00 42.06      H21B N
ATOM   8351  C   ARG    98      53.123  16.958 -11.075  1.00 56.47      H21B C
ATOM   8352  O   ARG    98      53.021  16.006 -10.315  1.00 57.42      H21B O
ATOM   8353  N   GLY    99      53.323  18.208 -10.653  1.00 59.29      H21B N
ATOM   8354  CA  GLY    99      53.343  18.538  -9.238  1.00 59.64      H21B C
ATOM   8355  C   GLY    99      52.179  17.874  -8.542  1.00 60.62      H21B C
ATOM   8356  O   GLY    99      51.325  17.298  -9.215  1.00 60.20      H21B O
ATOM   8357  N   TYR   100      52.130  17.940  -7.213  1.00 62.03      H21B N
ATOM   8358  CA  TYR   100      53.124  18.643  -6.419  1.00 62.23      H21B C
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ATOM   8359  CB  TYR   100      52.626  18.808  -4.981  1.00 63.21      H21B C
ATOM   8360  CG  TYR   100      51.617  19.916  -4.795  1.00 66.99      H21B C
ATOM   8361  CD1 TYR   100      50.253  19.662  -4.865  1.00 69.20      H21B C
ATOM   8362  CE1 TYR   100      49.326  20.676  -4.719  1.00 69.60      H21B C
ATOM   8363  CD2 TYR   100      52.026  21.219  -4.570  1.00 67.54      H21B C
ATOM   8364  CE2 TYR   100      51.105  22.236  -4.423  1.00 70.27      H21B C
ATOM   8365  CZ  TYR   100      49.757  21.960  -4.499  1.00 69.88      H21B C
ATOM   8366  OH  TYR   100      48.846  22.975  -4.357  1.00 68.94      H21B O
ATOM   8367  C   TYR   100      54.467  17.916  -6.412  1.00 61.98      H21B C
ATOM   8368  O   TYR   100      55.521  18.539  -6.522  1.00 63.49      H21B O
ATOM   8369  N   GLY   101      54.432  16.595  -6.292  1.00 59.53      H21B N
ATOM   8370  CA  GLY   101      55.663  15.839  -6.187  1.00 55.46      H21B C
ATOM   8371  C   GLY   101      56.609  16.052  -7.351  1.00 52.91      H21B C
ATOM   8372  O   GLY   101      57.785  15.688  -7.281  1.00 52.82      H21B O
ATOM   8373  N   MET   102      56.099  16.640  -8.427  1.00 50.64      H21B N
ATOM   8374  CA  MET   102      56.917  16.931  -9.603  1.00 49.55      H21B C
ATOM   8375  CB  MET   102      57.841  18.118  -9.334  1.00 45.02      H21B C
ATOM   8376  CG  MET   102      57.128  19.441  -9.291  1.00 40.44      H21B C
ATOM   8377  SD  MET   102      57.934  20.655  -8.250  1.00 32.66      H21B S
ATOM   8378  CE  MET   102      57.347  22.083  -8.992  1.00 41.41      H21B C
ATOM   8379  C   MET   102      57.756  15.726  -9.985  1.00 50.20      H21B C
ATOM   8380  O   MET   102      58.982  15.751  -9.873  1.00 50.67      H21B O
ATOM   8381  N   ASP   103      57.084  14.672 -10.433  1.00 49.62      H21B N
ATOM   8382  CA  ASP   103      57.756  13.426 -10.757  1.00 47.87      H21B C
ATOM   8383  CB  ASP   103      56.997  12.238 -10.138  1.00 49.38      H21B C
ATOM   8384  CG  ASP   103      55.590  12.042 -10.729  1.00 50.48      H21B C
ATOM   8385  OD1 ASP   103      55.088  12.937 -11.449  1.00 49.77      H21B O
ATOM   8386  OD2 ASP   103      54.986  10.978 -10.469  1.00 49.50      H21B O
ATOM   8387  C   ASP   103      57.890  13.258 -12.272  1.00 45.83      H21B C
ATOM   8388  O   ASP   103      58.508  12.306 -12.745  1.00 44.29      H21B O
ATOM   8389  N   VAL   104      57.314  14.187 -13.029  1.00 42.93      H21B N
ATOM   8390  CA  VAL   104      57.535  14.224 -14.471  1.00 42.13      H21B C
ATOM   8391  CB  VAL   104      56.421  13.493 -15.264  1.00 42.15      H21B C
ATOM   8392  CG1 VAL   104      56.696  13.618 -16.763  1.00 38.31      H21B C
ATOM   8393  CG2 VAL   104      56.334  12.035 -14.846  1.00 39.07      H21B C
ATOM   8394  C   VAL   104      57.539  15.659 -14.941  1.00 41.93      H21B C
ATOM   8395  O   VAL   104      56.611  16.399 -14.645  1.00 43.60      H21B O
ATOM   8396  N   TRP   105      58.562  16.053 -15.692  1.00 41.64      H21B N
ATOM   8397  CA  TRP   105      58.631  17.418 -16.202  1.00 41.31      H21B C
ATOM   8398  CB  TRP   105      59.989  18.032 -15.860  1.00 37.08      H21B C
ATOM   8399  CG  TRP   105      60.236  18.127 -14.398  1.00 32.86      H21B C
ATOM   8400  CD2 TRP   105      60.423  19.323 -13.638  1.00 30.58      H21B C
ATOM   8401  CE2 TRP   105      60.619  18.936 -12.293  1.00 31.25      H21B C
ATOM   8402  CE3 TRP   105      60.449  20.685 -13.960  1.00 29.62      H21B C
ATOM   8403  CD1 TRP   105      60.327  17.087 -13.508  1.00 32.58      H21B C
ATOM   8404  NE1 TRP   105      60.557  17.566 -12.242  1.00 30.54      H21B N
ATOM   8405  CZ2 TRP   105      60.831  19.869 -11.268  1.00 28.75      H21B C
ATOM   8406  CZ3 TRP   105      60.663  21.619 -12.934  1.00 28.89      H21B C
ATOM   8407  CH2 TRP   105      60.849  21.203 -11.610  1.00 29.51      H21B C
ATOM   8408  C   TRP   105      58.393  17.515 -17.706  1.00 43.85      H21B C
ATOM   8409  O   TRP   105      58.790  16.633 -18.462  1.00 43.17      H21B O
ATOM   8410  N   GLY   106      57.749  18.599 -18.134  1.00 48.37      H21B N
ATOM   8411  CA  GLY   106      57.772  18.971 -19.539  1.00 54.07      H21B C
ATOM   8412  C   GLY   106      59.143  19.475 -19.961  1.00 58.35      H21B C
ATOM   8413  O   GLY   106      60.004  19.725 -19.114  1.00 58.59      H21B O
ATOM   8414  N   GLN   107      59.353  19.626 -21.266  1.00 62.46      H21B N
ATOM   8415  CA  GLN   107      60.664  20.000 -21.785  1.00 66.67      H21B C
ATOM   8416  CB  GLN   107      60.642  20.044 -23.311  1.00 71.29      H21B C
ATOM   8417  CG  GLN   107      59.401  19.460 -23.927  1.00 81.85      H21B C
ATOM   8418  CD  GLN   107      58.376  20.516 -24.264  1.00 87.17      H21B C
ATOM   8419  OE1 GLN   107      58.653  21.711 -24.192  1.00 91.77      H21B O
ATOM   8420  NE2 GLN   107      57.180  20.082 -24.639  1.00 91.59      H21B N
ATOM   8421  C   GLN   107      61.091  21.363 -21.256  1.00 66.02      H21B C
ATOM   8422  O   GLN   107      62.263  21.604 -20.971  1.00 65.68      H21B O
ATOM   8423  N   GLY   108      60.117  22.251 -21.121  1.00 65.26      H21B N
ATOM   8424  CA  GLY   108      60.402  23.622 -20.760  1.00 61.19      H21B C
ATOM   8425  C   GLY   108      59.968  24.539 -21.879  1.00 58.99      H21B C
ATOM   8426  O   GLY   108      59.909  24.135 -23.038  1.00 55.89      H21B O
ATOM   8427  N   THR   109      59.638  25.775 -21.534  1.00 59.00      H21B N
ATOM   8428  CA  THR   109      59.567  26.817 -22.540  1.00 59.69      H21B C
ATOM   8429  CB  THR   109      58.099  27.114 -22.949  1.00 59.41      H21B C
ATOM   8430  OG1 THR   109      57.978  28.493 -23.313  1.00 59.56      H21B O
ATOM   8431  CG2 THR   109      57.140  26.775 -21.832  1.00 58.39      H21B C
ATOM   8432  C   THR   109      60.267  28.078 -22.067  1.00 57.64      H21B C
ATOM   8433  O   THR   109      60.179  28.452 -20.903  1.00 57.85      H21B O
ATOM   8434  N   THR   110      61.002  28.703 -22.977  1.00 56.06      H21B N
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ATOM   8435  CA  THR   110      61.757  29.905 -22.648  1.00 53.48      H21B C
ATOM   8436  CB  THR   110      63.106  29.974 -23.417  1.00 52.52      H21B C
ATOM   8437  OG1 THR   110      62.861  30.153 -24.814  1.00 51.93      H21B O
ATOM   8438  CG2 THR   110      63.896  28.701 -23.226  1.00 51.81      H21B C
ATOM   8439  C   THR   110      60.929  31.117 -23.025  1.00 51.15      H21B C
ATOM   8440  O   THR   110      60.340  31.170 -24.098  1.00 49.63      H21B O
ATOM   8441  N   VAL   111      60.883  32.086 -22.130  1.00 50.86      H21B N
ATOM   8442  CA  VAL   111      60.165  33.318 -22.382  1.00 53.09      H21B C
ATOM   8443  CB  VAL   111      59.024  33.498 -21.381  1.00 53.31      H21B C
ATOM   8444  CG1 VAL   111      58.293  34.802 -21.651  1.00 52.94      H21B C
ATOM   8445  CG2 VAL   111      58.086  32.321 -21.466  1.00 51.89      H21B C
ATOM   8446  C   VAL   111      61.127  34.481 -22.230  1.00 54.49      H21B C
ATOM   8447  O   VAL   111      61.634  34.722 -21.139  1.00 54.27      H21B O
ATOM   8448  N   THR   112      61.372  35.203 -23.320  1.00 56.71      H21B N
ATOM   8449  CA  THR   112      62.345  36.289 -23.315  1.00 59.52      H21B C
ATOM   8450  CB  THR   112      63.131  36.334 -24.627  1.00 60.00      H21B C
ATOM   8451  OG1 THR   112      63.445  35.001 -25.044  1.00 60.97      H21B O
ATOM   8452  CG2 THR   112      64.422  37.103 -24.437  1.00 59.68      H21B C
ATOM   8453  C   THR   112      61.640  37.622 -23.164  1.00 61.20      H21B C
ATOM   8454  O   THR   112      60.806  37.975 -23.981  1.00 62.07      H21B O
ATOM   8455  N   VAL   113      61.979  38.368 -22.123  1.00 63.91      H21B N
ATOM   8456  CA  VAL   113      61.405  39.692 -21.939  1.00 65.56      H21B C
ATOM   8457  CB  VAL   113      60.719  39.822 -20.578  1.00 65.05      H21B C
ATOM   8458  CG1 VAL   113      60.127  41.202 -20.438  1.00 63.91      H21B C
ATOM   8459  CG2 VAL   113      59.654  38.771 -20.434  1.00 64.43      H21B C
ATOM   8460  C   VAL   113      62.457  40.794 -22.039  1.00 67.13      H21B C
ATOM   8461  O   VAL   113      63.230  41.015 -21.106  1.00 68.10      H21B O
ATOM   8462  N   SER   114      62.468  41.492 -23.172  1.00 68.21      H21B N
ATOM   8463  CA  SER   114      63.407  42.588 -23.403  1.00 68.04      H21B C
ATOM   8464  CB  SER   114      64.642  42.068 -24.137  1.00 67.93      H21B C
ATOM   8465  OG  SER   114      65.609  43.086 -24.290  1.00 68.98      H21B O
ATOM   8466  C   SER   114      62.795  43.725 -24.219  1.00 68.45      H21B C
ATOM   8467  O   SER   114      61.954  43.509 -25.090  1.00 67.57      H21B O
ATOM   8468  N   SER   115      63.229  44.945 -23.938  1.00 69.36      H21B N
ATOM   8469  CA  SER   115      63.224  45.975 -24.961  1.00 69.76      H21B C
ATOM   8470  CB  SER   115      63.221  47.352 -24.320  1.00 70.24      H21B C
ATOM   8471  OG  SER   115      62.095  47.490 -23.474  1.00 72.93      H21B O
ATOM   8472  C   SER   115      64.512  45.746 -25.731  1.00 69.56      H21B C
ATOM   8473  O   SER   115      64.797  44.616 -26.131  1.00 71.87      H21B O
ATOM   8474  N   ALA   116      65.296  46.793 -25.933  1.00 67.88      H21B N
ATOM   8475  CA  ALA   116      66.653  46.616 -26.425  1.00 66.75      H21B C
ATOM   8476  CB  ALA   116      67.319  45.449 -25.711  1.00 67.94      H21B C
ATOM   8477  C   ALA   116      66.718  46.398 -27.925  1.00 66.13      H21B C
ATOM   8478  O   ALA   116      66.907  45.274 -28.387  1.00 65.19      H21B O
ATOM   8479  N   SER   117      66.569  47.484 -28.679  1.00 65.90      H21B N
ATOM   8480  CA  SER   117      66.897  47.492 -30.100  1.00 64.00      H21B C
ATOM   8481  CB  SER   117      66.318  48.732 -30.787  1.00 64.23      H21B C
ATOM   8482  OG  SER   117      66.935  48.950 -32.050  1.00 63.88      H21B O
ATOM   8483  C   SER   117      68.403  47.524 -30.222  1.00 62.28      H21B C
ATOM   8484  O   SER   117      69.087  48.047 -29.347  1.00 62.29      H21B O
ATOM   8485  N   THR   118      68.910  46.974 -31.316  1.00 60.02      H21B N
ATOM   8486  CA  THR   118      70.340  46.961 -31.579  1.00 58.31      H21B C
ATOM   8487  CB  THR   118      70.612  46.584 -33.050  1.00 58.63      H21B C
ATOM   8488  OG1 THR   118      71.785  47.270 -33.508  1.00 59.03      H21B O
ATOM   8489  CG2 THR   118      69.411  46.933 -33.926  1.00 55.99      H21B C
ATOM   8490  C   THR   118      71.048  48.280 -31.262  1.00 56.38      H21B C
ATOM   8491  O   THR   118      70.704  49.334 -31.794  1.00 57.50      H21B O
ATOM   8492  N   LYS   119      72.034  48.206 -30.376  1.00 53.48      H21B N
ATOM   8493  CA  LYS   119      72.906  49.337 -30.091  1.00 49.19      H21B C
ATOM   8494  CB  LYS   119      72.642  49.885 -28.693  1.00 47.24      H21B C
ATOM   8495  CG  LYS   119      73.647  50.952 -28.288  1.00 48.71      H21B C
ATOM   8496  CD  LYS   119      73.521  51.380 -26.829  1.00 47.72      H21B C
ATOM   8497  CE  LYS   119      74.717  52.241 -26.443  1.00 47.92      H21B C
ATOM   8498  NZ  LYS   119      75.119  52.084 -25.016  1.00 50.67      H21B N
ATOM   8499  C   LYS   119      74.375  48.922 -30.200  1.00 47.21      H21B C
ATOM   8500  O   LYS   119      74.735  47.761 -29.963  1.00 45.25      H21B O
ATOM   8501  N   GLY   120      75.220  49.879 -30.570  1.00 44.35      H21B N
ATOM   8502  CA  GLY   120      76.646  49.612 -30.652  1.00 40.92      H21B C
ATOM   8503  C   GLY   120      77.327  49.892 -29.326  1.00 38.58      H21B C
ATOM   8504  O   GLY   120      76.802  50.621 -28.474  1.00 36.46      H21B O
ATOM   8505  N   PRO   121      78.506  49.310 -29.114  1.00 36.94      H21B N
ATOM   8506  CD  PRO   121      79.208  48.416 -30.042  1.00 35.27      H21B C
ATOM   8507  CA  PRO   121      79.171  49.390 -27.813  1.00 36.50      H21B C
ATOM   8508  CB  PRO   121      80.132  48.220 -27.839  1.00 34.63      H21B C
ATOM   8509  CG  PRO   121      80.452  48.077 -29.285  1.00 36.69      H21B C
ATOM   8510  C   PRO   121      79.893  50.716 -27.558  1.00 37.73      H21B C
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ATOM   8511  O   PRO  121      80.301  51.421 -28.487  1.00 35.85      H21B O
ATOM   8512  N   SER  122      80.049  51.047 -26.284  1.00 37.92      H21B N
ATOM   8513  CA  SER  122      81.072  51.998 -25.896  1.00 38.88      H21B C
ATOM   8514  CB  SER  122      80.612  52.828 -24.689  1.00 39.55      H21B C
ATOM   8515  OG  SER  122      79.510  53.676 -25.028  1.00 41.26      H21B O
ATOM   8516  C   SER  122      82.326  51.205 -25.552  1.00 38.81      H21B C
ATOM   8517  O   SER  122      82.277  50.222 -24.805  1.00 39.06      H21B O
ATOM   8518  N   VAL  123      83.451  51.617 -26.114  1.00 37.51      H21B N
ATOM   8519  CA  VAL  123      84.702  50.973 -25.770  1.00 36.15      H21B C
ATOM   8520  CB  VAL  123      85.525  50.643 -27.024  1.00 34.75      H21B C
ATOM   8521  CG1 VAL  123      86.827  49.988 -26.615  1.00 32.49      H21B C
ATOM   8522  CG2 VAL  123      84.713  49.721 -27.955  1.00 31.95      H21B C
ATOM   8523  C   VAL  123      85.529  51.854 -24.851  1.00 35.96      H21B C
ATOM   8524  O   VAL  123      85.920  52.952 -25.230  1.00 37.71      H21B O
ATOM   8525  N   PHE  124      85.776  51.362 -23.639  1.00 34.63      H21B N
ATOM   8526  CA  PHE  124      86.671  52.014 -22.688  1.00 33.18      H21B C
ATOM   8527  CB  PHE  124      86.008  52.139 -21.319  1.00 29.98      H21B C
ATOM   8528  CG  PHE  124      84.785  53.005 -21.322  1.00 30.01      H21B C
ATOM   8529  CD1 PHE  124      84.885  54.368 -21.582  1.00 30.04      H21B C
ATOM   8530  CD2 PHE  124      83.534  52.471 -21.043  1.00 28.86      H21B C
ATOM   8531  CE1 PHE  124      83.757  55.186 -21.560  1.00 28.02      H21B C
ATOM   8532  CE2 PHE  124      82.411  53.279 -21.022  1.00 27.96      H21B C
ATOM   8533  CZ  PHE  124      82.526  54.639 -21.279  1.00 28.14      H21B C
ATOM   8534  C   PHE  124      87.966  51.242 -22.531  1.00 33.73      H21B C
ATOM   8535  O   PHE  124      87.992  50.015 -22.596  1.00 34.72      H21B O
ATOM   8536  N   PRO  125      89.068  51.960 -22.314  1.00 35.69      H21B N
ATOM   8537  CD  PRO  125      89.202  53.424 -22.358  1.00 34.56      H21B C
ATOM   8538  CA  PRO  125      90.351  51.311 -22.040  1.00 36.09      H21B C
ATOM   8539  CB  PRO  125      91.357  52.406 -22.339  1.00 34.77      H21B C
ATOM   8540  CG  PRO  125      90.632  53.632 -21.941  1.00 34.99      H21B C
ATOM   8541  C   PRO  125      90.474  50.812 -20.598  1.00 36.79      H21B C
ATOM   8542  O   PRO  125      90.186  51.553 -19.645  1.00 36.75      H21B O
ATOM   8543  N   LEU  126      90.911  49.565 -20.449  1.00 35.96      H21B N
ATOM   8544  CA  LEU  126      91.369  49.067 -19.155  1.00 38.37      H21B C
ATOM   8545  CB  LEU  126      90.912  47.623 -18.959  1.00 36.93      H21B C
ATOM   8546  CG  LEU  126      89.403  47.434 -19.047  1.00 32.95      H21B C
ATOM   8547  CD1 LEU  126      89.058  46.042 -18.690  1.00 30.64      H21B C
ATOM   8548  CD2 LEU  126      88.714  48.406 -18.114  1.00 33.63      H21B C
ATOM   8549  C   LEU  126      92.897  49.153 -18.994  1.00 41.03      H21B C
ATOM   8550  O   LEU  126      93.620  48.185 -19.232  1.00 40.02      H21B O
ATOM   8551  N   ALA  127      93.367  50.322 -18.568  1.00 44.45      H21B N
ATOM   8552  CA  ALA  127      94.789  50.605 -18.423  1.00 48.51      H21B C
ATOM   8553  CB  ALA  127      94.968  52.014 -17.860  1.00 46.45      H21B C
ATOM   8554  C   ALA  127      95.527  49.596 -17.546  1.00 51.79      H21B C
ATOM   8555  O   ALA  127      94.996  49.107 -16.552  1.00 50.48      H21B O
ATOM   8556  N   PRO  128      96.778  49.295 -17.902  1.00 55.84      H21B N
ATOM   8557  CD  PRO  128      97.532  49.968 -18.971  1.00 58.16      H21B C
ATOM   8558  CA  PRO  128      97.578  48.269 -17.232  1.00 61.96      H21B C
ATOM   8559  CB  PRO  128      98.792  48.112 -18.141  1.00 61.86      H21B C
ATOM   8560  CG  PRO  128      98.925  49.434 -18.799  1.00 59.88      H21B C
ATOM   8561  C   PRO  128      97.966  48.684 -15.818  1.00 66.51      H21B C
ATOM   8562  O   PRO  128      98.194  49.867 -15.551  1.00 67.81      H21B O
ATOM   8563  N   SER  129      98.041  47.697 -14.926  1.00 71.65      H21B N
ATOM   8564  CA  SER  129      98.268  47.930 -13.496  1.00 76.44      H21B C
ATOM   8565  CB  SER  129      98.539  46.594 -12.778  1.00 76.87      H21B C
ATOM   8566  OG  SER  129      97.610  45.589 -13.171  1.00 74.76      H21B O
ATOM   8567  C   SER  129      99.444  48.875 -13.269  1.00 80.19      H21B C
ATOM   8568  O   SER  129     100.343  48.974 -14.109  1.00 81.78      H21B O
ATOM   8569  N   GLY  135     104.091  39.639 -12.189  1.00119.83      H21B N
ATOM   8570  CA  GLY  135     105.514  39.440 -12.396  1.00119.78      H21B C
ATOM   8571  C   GLY  135     106.108  40.387 -13.424  1.00119.10      H21B C
ATOM   8572  O   GLY  135     106.017  41.611 -13.283  1.00120.34      H21B O
ATOM   8573  N   GLY  136     106.728  39.823 -14.458  1.00116.59      H21B N
ATOM   8574  CA  GLY  136     107.222  40.634 -15.557  1.00111.70      H21B C
ATOM   8575  C   GLY  136     106.106  40.963 -16.526  1.00107.95      H21B C
ATOM   8576  O   GLY  136     106.272  41.764 -17.444  1.00108.38      H21B O
ATOM   8577  N   THR  137     104.957  40.334 -16.313  1.00103.31      H21B N
ATOM   8578  CA  THR  137     103.792  40.520 -17.166  1.00 97.68      H21B C
ATOM   8579  CB  THR  137     103.096  39.171 -17.453  1.00 98.99      H21B C
ATOM   8580  OG1 THR  137     102.819  38.504 -16.215  1.00100.61      H21B O
ATOM   8581  CG2 THR  137     103.979  38.281 -18.314  1.00 99.84      H21B C
ATOM   8582  C   THR  137     102.783  41.458 -16.512  1.00 92.44      H21B C
ATOM   8583  O   THR  137     102.556  41.394 -15.303  1.00 91.69      H21B O
ATOM   8584  N   ALA  138     102.188  42.335 -17.316  1.00 86.07      H21B N
ATOM   8585  CA  ALA  138     101.075  43.159 -16.861  1.00 79.49      H21B C
ATOM   8586  CB  ALA  138     101.440  44.631 -16.949  1.00 79.63      H21B C
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ATOM   8587  C   ALA  138      99.834  42.871 -17.704  1.00 74.95      H21B C
ATOM   8588  O   ALA  138      99.937  42.528 -18.884  1.00 73.87      H21B O
ATOM   8589  N   ALA  139      98.663  43.000 -17.087  1.00 68.95      H21B N
ATOM   8590  CA  ALA  139      97.404  42.783 -17.784  1.00 63.72      H21B C
ATOM   8591  CB  ALA  139      96.487  41.900 -16.944  1.00 63.98      H21B C
ATOM   8592  C   ALA  139      96.716  44.116 -18.104  1.00 60.62      H21B C
ATOM   8593  O   ALA  139      96.707  45.045 -17.289  1.00 59.40      H21B O
ATOM   8594  N   LEU  140      96.154  44.197 -19.306  1.00 55.59      H21B N
ATOM   8595  CA  LEU  140      95.516  45.408 -19.795  1.00 51.29      H21B C
ATOM   8596  CB  LEU  140      96.535  46.309 -20.494  1.00 50.60      H21B C
ATOM   8597  CG  LEU  140      97.142  45.763 -21.799  1.00 50.41      H21B C
ATOM   8598  CD1 LEU  140      96.234  46.074 -22.988  1.00 49.55      H21B C
ATOM   8599  CD2 LEU  140      98.506  46.382 -22.014  1.00 49.82      H21B C
ATOM   8600  C   LEU  140      94.501  44.927 -20.803  1.00 49.58      H21B C
ATOM   8601  O   LEU  140      94.544  43.765 -21.210  1.00 48.85      H21B O
ATOM   8602  N   GLY  141      93.600  45.809 -21.219  1.00 47.47      H21B N
ATOM   8603  CA  GLY  141      92.610  45.412 -22.201  1.00 47.38      H21B C
ATOM   8604  C   GLY  141      91.581  46.490 -22.427  1.00 47.35      H21B C
ATOM   8605  O   GLY  141      91.850  47.658 -22.143  1.00 47.11      H21B O
ATOM   8606  N   CYS  142      90.406  46.127 -22.931  1.00 47.21      H21B N
ATOM   8607  CA  CYS  142      89.360  47.128 -23.024  1.00 47.00      H21B C
ATOM   8608  C   CYS  142      87.936  46.620 -22.781  1.00 43.01      H21B C
ATOM   8609  O   CYS  142      87.595  45.495 -23.137  1.00 42.06      H21B O
ATOM   8610  CB  CYS  142      89.459  47.844 -24.371  1.00 50.45      H21B C
ATOM   8611  SG  CYS  142      88.981  46.864 -25.834  1.00 59.47      H21B S
ATOM   8612  N   LEU  143      87.119  47.474 -22.161  1.00 38.35      H21B N
ATOM   8613  CA  LEU  143      85.750  47.140 -21.766  1.00 34.87      H21B C
ATOM   8614  CB  LEU  143      85.345  47.929 -20.513  1.00 35.86      H21B C
ATOM   8615  CG  LEU  143      83.870  47.892 -20.078  1.00 33.46      H21B C
ATOM   8616  CD1 LEU  143      83.550  46.536 -19.524  1.00 32.08      H21B C
ATOM   8617  CD2 LEU  143      83.602  48.964 -19.035  1.00 31.98      H21B C
ATOM   8618  C   LEU  143      84.787  47.485 -22.886  1.00 34.34      H21B C
ATOM   8619  O   LEU  143      84.717  48.643 -23.302  1.00 31.63      H21B O
ATOM   8620  N   VAL  144      84.047  46.488 -23.372  1.00 34.47      H21B N
ATOM   8621  CA  VAL  144      83.069  46.720 -24.419  1.00 35.66      H21B C
ATOM   8622  CB  VAL  144      83.162  45.660 -25.533  1.00 34.46      H21B C
ATOM   8623  CG1 VAL  144      82.203  46.022 -26.664  1.00 35.75      H21B C
ATOM   8624  CG2 VAL  144      84.594  45.556 -26.042  1.00 33.25      H21B C
ATOM   8625  C   VAL  144      81.683  46.690 -23.783  1.00 38.56      H21B C
ATOM   8626  O   VAL  144      81.134  45.630 -23.460  1.00 38.50      H21B O
ATOM   8627  N   LYS  145      81.120  47.879 -23.620  1.00 38.89      H21B N
ATOM   8628  CA  LYS  145      79.997  48.085 -22.734  1.00 39.76      H21B C
ATOM   8629  CB  LYS  145      80.342  49.261 -21.801  1.00 41.45      H21B C
ATOM   8630  CG  LYS  145      79.385  49.497 -20.642  1.00 41.62      H21B C
ATOM   8631  CD  LYS  145      80.143  49.965 -19.415  1.00 44.51      H21B C
ATOM   8632  CE  LYS  145      79.965  51.461 -19.152  1.00 47.71      H21B C
ATOM   8633  NZ  LYS  145      78.551  51.812 -18.786  1.00 52.84      H21B N
ATOM   8634  C   LYS  145      78.725  48.367 -23.548  1.00 40.29      H21B C
ATOM   8635  O   LYS  145      78.764  49.101 -24.546  1.00 37.61      H21B O
ATOM   8636  N   ASP  146      77.620  47.746 -23.124  1.00 41.41      H21B N
ATOM   8637  CA  ASP  146      76.255  48.156 -23.480  1.00 42.07      H21B C
ATOM   8638  CB  ASP  146      75.997  49.578 -23.000  1.00 44.43      H21B C
ATOM   8639  CG  ASP  146      75.619  49.625 -21.539  1.00 49.82      H21B C
ATOM   8640  OD1 ASP  146      75.227  48.554 -21.020  1.00 49.91      H21B O
ATOM   8641  OD2 ASP  146      75.708  50.715 -20.915  1.00 52.78      H21B O
ATOM   8642  C   ASP  146      75.812  48.059 -24.933  1.00 40.90      H21B C
ATOM   8643  O   ASP  146      75.271  49.021 -25.475  1.00 40.19      H21B O
ATOM   8644  N   TYR  147      76.012  46.895 -25.545  1.00 39.23      H21B N
ATOM   8645  CA  TYR  147      75.641  46.677 -26.939  1.00 40.02      H21B C
ATOM   8646  CB  TYR  147      76.847  46.185 -27.733  1.00 38.54      H21B C
ATOM   8647  CG  TYR  147      77.413  44.872 -27.241  1.00 38.05      H21B C
ATOM   8648  CD1 TYR  147      77.005  43.660 -27.799  1.00 36.64      H21B C
ATOM   8649  CE1 TYR  147      77.585  42.457 -27.406  1.00 35.87      H21B C
ATOM   8650  CD2 TYR  147      78.408  44.842 -26.261  1.00 37.18      H21B C
ATOM   8651  CE2 TYR  147      78.989  43.646 -25.863  1.00 36.14      H21B C
ATOM   8652  CZ  TYR  147      78.583  42.458 -26.445  1.00 35.73      H21B C
ATOM   8653  OH  TYR  147      79.240  41.290 -26.130  1.00 33.54      H21B O
ATOM   8654  C   TYR  147      74.501  45.662 -27.090  1.00 41.03      H21B C
ATOM   8655  O   TYR  147      74.063  45.055 -26.121  1.00 39.81      H21B O
ATOM   8656  N   PHE  148      74.038  45.476 -28.321  1.00 42.76      H21B N
ATOM   8657  CA  PHE  148      72.928  44.579 -28.589  1.00 44.38      H21B C
ATOM   8658  CB  PHE  148      71.654  45.101 -27.916  1.00 47.16      H21B C
ATOM   8659  CG  PHE  148      70.495  44.148 -28.013  1.00 51.60      H21B C
ATOM   8660  CD1 PHE  148      69.885  43.904 -29.231  1.00 52.66      H21B C
ATOM   8661  CD2 PHE  148      70.046  43.464 -26.893  1.00 53.38      H21B C
ATOM   8662  CE1 PHE  148      68.861  43.002 -29.336  1.00 53.99      H21B C
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ATOM   8663  CE2 PHE   148      69.021  42.561 -26.992  1.00 54.37      H21B C
ATOM   8664  CZ  PHE   148      68.426  42.329 -28.218  1.00 55.54      H21B C
ATOM   8665  C   PHE   148      72.690  44.474 -30.087  1.00 43.47      H21B C
ATOM   8666  O   PHE   148      72.795  45.454 -30.809  1.00 43.90      H21B O
ATOM   8667  N   PRO   149      72.363  43.279 -30.577  1.00 43.17      H21B N
ATOM   8668  CD  PRO   149      71.924  43.094 -31.969  1.00 43.32      H21B C
ATOM   8669  CA  PRO   149      72.424  42.006 -29.863  1.00 44.42      H21B C
ATOM   8670  CB  PRO   149      71.744  41.033 -30.818  1.00 42.48      H21B C
ATOM   8671  CG  PRO   149      72.009  41.596 -32.152  1.00 41.77      H21B C
ATOM   8672  C   PRO   149      73.874  41.627 -29.635  1.00 44.50      H21B C
ATOM   8673  O   PRO   149      74.748  42.479 -29.653  1.00 46.03      H21B O
ATOM   8674  N   GLU   150      74.116  40.342 -29.413  1.00 44.54      H21B N
ATOM   8675  CA  GLU   150      75.421  39.764 -29.666  1.00 45.89      H21B C
ATOM   8676  CB  GLU   150      75.544  38.405 -28.977  1.00 46.98      H21B C
ATOM   8677  CG  GLU   150      75.465  38.451 -27.465  1.00 48.70      H21B C
ATOM   8678  CD  GLU   150      76.823  38.263 -26.821  1.00 49.44      H21B C
ATOM   8679  OE1 GLU   150      77.643  39.205 -26.905  1.00 48.89      H21B O
ATOM   8680  OE2 GLU   150      77.073  37.178 -26.241  1.00 48.71      H21B O
ATOM   8681  C   GLU   150      75.507  39.577 -31.172  1.00 44.93      H21B C
ATOM   8682  O   GLU   150      74.527  39.787 -31.886  1.00 45.62      H21B O
ATOM   8683  N   PRO   151      76.680  39.188 -31.676  1.00 42.42      H21B N
ATOM   8684  CD  PRO   151      76.847  38.646 -33.034  1.00 40.81      H21B C
ATOM   8685  CA  PRO   151      77.943  39.164 -30.940  1.00 42.36      H21B C
ATOM   8686  CB  PRO   151      78.688  38.010 -31.588  1.00 40.92      H21B C
ATOM   8687  CG  PRO   151      78.279  38.116 -33.025  1.00 39.66      H21B C
ATOM   8688  C   PRO   151      78.701  40.488 -31.104  1.00 42.20      H21B C
ATOM   8689  O   PRO   151      78.303  41.359 -31.874  1.00 43.34      H21B O
ATOM   8690  N   VAL   152      79.796  40.624 -30.374  1.00 40.37      H21B N
ATOM   8691  CA  VAL   152      80.856  41.531 -30.763  1.00 39.57      H21B C
ATOM   8692  CB  VAL   152      81.098  42.617 -29.689  1.00 39.88      H21B C
ATOM   8693  CG1 VAL   152      79.869  43.496 -29.558  1.00 40.66      H21B C
ATOM   8694  CG2 VAL   152      81.432  41.967 -28.355  1.00 38.69      H21B C
ATOM   8695  C   VAL   152      82.112  40.689 -30.924  1.00 40.03      H21B C
ATOM   8696  O   VAL   152      82.258  39.655 -30.288  1.00 38.78      H21B O
ATOM   8697  N   THR   153      83.018  41.120 -31.780  1.00 40.25      H21B N
ATOM   8698  CA  THR   153      84.279  40.431 -31.900  1.00 42.37      H21B C
ATOM   8699  CB  THR   153      84.458  39.941 -33.336  1.00 44.70      H21B C
ATOM   8700  OG1 THR   153      84.342  41.054 -34.231  1.00 49.27      H21B O
ATOM   8701  CG2 THR   153      83.372  38.916 -33.683  1.00 46.36      H21B C
ATOM   8702  C   THR   153      85.399  41.392 -31.504  1.00 43.32      H21B C
ATOM   8703  O   THR   153      85.273  42.599 -31.688  1.00 44.64      H21B O
ATOM   8704  N   VAL   154      86.485  40.871 -30.941  1.00 43.33      H21B N
ATOM   8705  CA  VAL   154      87.588  41.733 -30.518  1.00 42.48      H21B C
ATOM   8706  CB  VAL   154      87.654  41.842 -28.985  1.00 40.02      H21B C
ATOM   8707  CG1 VAL   154      88.723  42.809 -28.587  1.00 39.30      H21B C
ATOM   8708  CG2 VAL   154      86.336  42.294 -28.452  1.00 39.07      H21B C
ATOM   8709  C   VAL   154      88.944  41.253 -31.038  1.00 44.23      H21B C
ATOM   8710  O   VAL   154      89.225  40.061 -31.084  1.00 45.14      H21B O
ATOM   8711  N   SER   155      89.779  42.199 -31.436  1.00 46.16      H21B N
ATOM   8712  CA  SER   155      91.058  41.898 -32.053  1.00 49.14      H21B C
ATOM   8713  CB  SER   155      90.953  42.124 -33.568  1.00 50.30      H21B C
ATOM   8714  OG  SER   155      92.205  42.456 -34.151  1.00 52.89      H21B O
ATOM   8715  C   SER   155      92.100  42.830 -31.429  1.00 51.51      H21B C
ATOM   8716  O   SER   155      91.742  43.875 -30.870  1.00 52.21      H21B O
ATOM   8717  N   TRP   156      93.377  42.461 -31.508  1.00 51.65      H21B N
ATOM   8718  CA  TRP   156      94.425  43.269 -30.892  1.00 53.25      H21B C
ATOM   8719  CB  TRP   156      94.963  42.580 -29.636  1.00 51.30      H21B C
ATOM   8720  CG  TRP   156      94.025  42.624 -28.473  1.00 48.72      H21B C
ATOM   8721  CD2 TRP   156      93.881  43.691 -27.533  1.00 46.26      H21B C
ATOM   8722  CE2 TRP   156      92.913  43.291 -26.592  1.00 45.15      H21B C
ATOM   8723  CE3 TRP   156      94.477  44.943 -27.396  1.00 43.80      H21B C
ATOM   8724  CD1 TRP   156      93.157  41.645 -28.077  1.00 47.86      H21B C
ATOM   8725  NE1 TRP   156      92.486  42.039 -26.945  1.00 45.30      H21B N
ATOM   8726  CZ2 TRP   156      92.529  44.102 -25.526  1.00 44.56      H21B C
ATOM   8727  CZ3 TRP   156      94.097  45.745 -26.340  1.00 43.81      H21B C
ATOM   8728  CH2 TRP   156      93.129  45.323 -25.416  1.00 42.87      H21B C
ATOM   8729  C   TRP   156      95.583  43.578 -31.828  1.00 56.08      H21B C
ATOM   8730  O   TRP   156      96.211  42.682 -32.388  1.00 55.96      H21B O
ATOM   8731  N   ASN   157      95.878  44.861 -31.977  1.00 60.61      H21B N
ATOM   8732  CA  ASN   157      96.870  45.289 -32.952  1.00 64.41      H21B C
ATOM   8733  CB  ASN   157      98.284  44.890 -32.479  1.00 65.66      H21B C
ATOM   8734  CG  ASN   157      98.725  45.647 -31.206  1.00 66.78      H21B C
ATOM   8735  OD1 ASN   157      98.165  46.697 -30.860  1.00 66.57      H21B O
ATOM   8736  ND2 ASN   157      99.731  45.110 -30.515  1.00 63.54      H21B N
ATOM   8737  C   ASN   157      96.515  44.625 -34.287  1.00 66.14      H21B C
ATOM   8738  O   ASN   157      97.389  44.180 -35.031  1.00 66.39      H21B O
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ATOM   8739  N   SER  158      95.211  44.546 -34.558  1.00 67.60      H21B N
ATOM   8740  CA  SER  158      94.688  44.092 -35.843  1.00 68.35      H21B C
ATOM   8741  CB  SER  158      95.198  44.984 -36.968  1.00 68.48      H21B C
ATOM   8742  OG  SER  158      94.876  46.343 -36.726  1.00 69.06      H21B O
ATOM   8743  C   SER  158      95.052  42.655 -36.141  1.00 69.01      H21B C
ATOM   8744  O   SER  158      94.963  42.206 -37.282  1.00 69.30      H21B O
ATOM   8745  N   GLY  159      95.464  41.938 -35.103  1.00 69.36      H21B N
ATOM   8746  CA  GLY  159      95.792  40.535 -35.255  1.00 69.78      H21B C
ATOM   8747  C   GLY  159      97.208  40.233 -34.819  1.00 70.12      H21B C
ATOM   8748  O   GLY  159      97.498  39.142 -34.331  1.00 69.25      H21B O
ATOM   8749  N   ALA  160      98.093  41.208 -34.985  1.00 71.83      H21B N
ATOM   8750  CA  ALA  160      99.512  41.003 -34.717  1.00 73.36      H21B C
ATOM   8751  CB  ALA  160     100.271  42.319 -34.871  1.00 73.77      H21B C
ATOM   8752  C   ALA  160      99.731  40.430 -33.320  1.00 74.01      H21B C
ATOM   8753  O   ALA  160     100.796  39.892 -33.022  1.00 75.23      H21B O
ATOM   8754  N   LEU  161      98.716  40.545 -32.469  1.00 73.29      H21B N
ATOM   8755  CA  LEU  161      98.813  40.076 -31.093  1.00 71.19      H21B C
ATOM   8756  CB  LEU  161      98.713  41.266 -30.135  1.00 71.12      H21B C
ATOM   8757  CG  LEU  161      99.008  41.016 -28.659  1.00 70.62      H21B C
ATOM   8758  CD1 LEU  161     100.385  40.403 -28.515  1.00 72.48      H21B C
ATOM   8759  CD2 LEU  161      98.927  42.322 -27.897  1.00 69.65      H21B C
ATOM   8760  C   LEU  161      97.703  39.073 -30.793  1.00 70.20      H21B C
ATOM   8761  O   LEU  161      96.541  39.445 -30.655  1.00 69.33      H21B O
ATOM   8762  N   THR  162      98.063  37.798 -30.699  1.00 69.03      H21B N
ATOM   8763  CA  THR  162      97.090  36.767 -30.376  1.00 69.44      H21B C
ATOM   8764  CB  THR  162      97.079  35.661 -31.444  1.00 70.49      H21B C
ATOM   8765  OG1 THR  162      98.431  35.293 -31.752  1.00 72.93      H21B O
ATOM   8766  CG2 THR  162      96.367  36.127 -32.697  1.00 69.81      H21B C
ATOM   8767  C   THR  162      97.417  36.143 -29.024  1.00 69.37      H21B C
ATOM   8768  O   THR  162      96.528  35.857 -28.220  1.00 68.44      H21B O
ATOM   8769  N   SER  163      98.703  35.943 -28.773  1.00 68.77      H21B N
ATOM   8770  CA  SER  163      99.130  35.291 -27.550  1.00 68.52      H21B C
ATOM   8771  CB  SER  163     100.646  35.102 -27.577  1.00 68.30      H21B C
ATOM   8772  OG  SER  163     101.019  33.934 -26.874  1.00 69.26      H21B O
ATOM   8773  C   SER  163      98.720  36.107 -26.319  1.00 68.15      H21B C
ATOM   8774  O   SER  163      98.861  37.332 -26.300  1.00 69.61      H21B O
ATOM   8775  N   GLY  164      98.203  35.419 -25.300  1.00 66.79      H21B N
ATOM   8776  CA  GLY  164      97.882  36.063 -24.033  1.00 63.70      H21B C
ATOM   8777  C   GLY  164      96.541  36.773 -24.043  1.00 61.53      H21B C
ATOM   8778  O   GLY  164      96.106  37.351 -23.038  1.00 59.15      H21B O
ATOM   8779  N   VAL  165      95.891  36.733 -25.200  1.00 60.65      H21B N
ATOM   8780  CA  VAL  165      94.596  37.360 -25.381  1.00 61.21      H21B C
ATOM   8781  CB  VAL  165      94.320  37.626 -26.867  1.00 60.77      H21B C
ATOM   8782  CG1 VAL  165      92.857  38.002 -27.064  1.00 61.36      H21B C
ATOM   8783  CG2 VAL  165      95.225  38.729 -27.365  1.00 61.58      H21B C
ATOM   8784  C   VAL  165      93.490  36.473 -24.842  1.00 61.45      H21B C
ATOM   8785  O   VAL  165      93.344  35.331 -25.271  1.00 62.50      H21B O
ATOM   8786  N   HIS  166      92.716  37.007 -23.903  1.00 61.14      H21B N
ATOM   8787  CA  HIS  166      91.531  36.329 -23.404  1.00 61.29      H21B C
ATOM   8788  CB  HIS  166      91.764  35.867 -21.964  1.00 65.46      H21B C
ATOM   8789  CG  HIS  166      91.226  34.498 -21.675  1.00 72.03      H21B C
ATOM   8790  CD2 HIS  166      91.738  33.267 -21.923  1.00 74.47      H21B C
ATOM   8791  ND1 HIS  166      90.012  34.289 -21.051  1.00 74.85      H21B N
ATOM   8792  CE1 HIS  166      89.800  32.990 -20.926  1.00 76.37      H21B C
ATOM   8793  NE2 HIS  166      90.832  32.347 -21.447  1.00 76.73      H21B N
ATOM   8794  C   HIS  166      90.306  37.248 -23.481  1.00 58.26      H21B C
ATOM   8795  O   HIS  166      90.182  38.217 -22.728  1.00 57.57      H21B O
ATOM   8796  N   THR  167      89.406  36.931 -24.404  1.00 54.96      H21B N
ATOM   8797  CA  THR  167      88.176  37.690 -24.574  1.00 52.16      H21B C
ATOM   8798  CB  THR  167      87.855  37.872 -26.068  1.00 52.06      H21B C
ATOM   8799  OG1 THR  167      89.000  38.412 -26.743  1.00 52.50      H21B O
ATOM   8800  CG2 THR  167      86.705  38.836 -26.241  1.00 50.49      H21B C
ATOM   8801  C   THR  167      86.994  36.988 -23.883  1.00 49.74      H21B C
ATOM   8802  O   THR  167      86.444  36.017 -24.396  1.00 49.74      H21B O
ATOM   8803  N   PHE  168      86.605  37.506 -22.722  1.00 46.79      H21B N
ATOM   8804  CA  PHE  168      85.631  36.862 -21.854  1.00 44.69      H21B C
ATOM   8805  CB  PHE  168      85.690  37.497 -20.469  1.00 42.76      H21B C
ATOM   8806  CG  PHE  168      86.901  37.114 -19.682  1.00 41.19      H21B C
ATOM   8807  CD1 PHE  168      88.035  37.895 -19.707  1.00 40.51      H21B C
ATOM   8808  CD2 PHE  168      86.896  35.976 -18.896  1.00 42.33      H21B C
ATOM   8809  CE1 PHE  168      89.141  37.551 -18.962  1.00 40.25      H21B C
ATOM   8810  CE2 PHE  168      88.003  35.625 -18.148  1.00 41.44      H21B C
ATOM   8811  CZ  PHE  168      89.122  36.411 -18.180  1.00 40.34      H21B C
ATOM   8812  C   PHE  168      84.199  36.929 -22.373  1.00 43.71      H21B C
ATOM   8813  O   PHE  168      83.829  37.859 -23.077  1.00 45.06      H21B O
ATOM   8814  N   PRO  169      83.367  35.942 -22.014  1.00 42.61      H21B N
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ATOM	8815	CD	PRO	169	83.682	34.799	-21.144	1.00	41.27	H21B	C
ATOM	8816	CA	PRO	169	81.976	35.900	-22.477	1.00	42.04	H21B	C
ATOM	8817	CB	PRO	169	81.472	34.548	-21.975	1.00	40.17	H21B	C
ATOM	8818	CG	PRO	169	82.714	33.768	-21.619	1.00	39.86	H21B	C
ATOM	8819	C	PRO	169	81.201	37.061	-21.867	1.00	41.81	H21B	C
ATOM	8820	O	PRO	169	81.511	37.493	-20.757	1.00	40.70	H21B	O
ATOM	8821	N	ALA	170	80.201	37.562	-22.587	1.00	41.85	H21B	N
ATOM	8822	CA	ALA	170	79.512	38.785	-22.184	1.00	43.47	H21B	C
ATOM	8823	CB	ALA	170	78.573	39.232	-23.285	1.00	42.59	H21B	C
ATOM	8824	C	ALA	170	78.735	38.618	-20.879	1.00	45.43	H21B	C
ATOM	8825	O	ALA	170	78.762	37.571	-20.255	1.00	46.51	H21B	O
ATOM	8826	N	VAL	171	78.049	39.668	-20.465	1.00	47.55	H21B	N
ATOM	8827	CA	VAL	171	77.162	39.592	-19.323	1.00	50.15	H21B	C
ATOM	8828	CB	VAL	171	77.784	40.306	-18.077	1.00	49.03	H21B	C
ATOM	8829	CG1	VAL	171	77.820	41.807	-18.287	1.00	46.37	H21B	C
ATOM	8830	CG2	VAL	171	77.008	39.969	-16.817	1.00	47.04	H21B	C
ATOM	8831	C	VAL	171	75.921	40.335	-19.788	1.00	54.76	H21B	C
ATOM	8832	O	VAL	171	76.035	41.327	-20.502	1.00	54.18	H21B	O
ATOM	8833	N	LEU	172	74.742	39.841	-19.423	1.00	60.45	H21B	N
ATOM	8834	CA	LEU	172	73.507	40.542	-19.741	1.00	66.51	H21B	C
ATOM	8835	CB	LEU	172	72.403	39.548	-20.094	1.00	69.05	H21B	C
ATOM	8836	CG	LEU	172	72.171	39.445	-21.602	1.00	71.42	H21B	C
ATOM	8837	CD1	LEU	172	71.366	38.195	-21.963	1.00	70.43	H21B	C
ATOM	8838	CD2	LEU	172	71.459	40.714	-22.037	1.00	71.49	H21B	C
ATOM	8839	C	LEU	172	73.088	41.398	-18.564	1.00	69.67	H21B	C
ATOM	8840	O	LEU	172	73.104	40.942	-17.423	1.00	70.14	H21B	O
ATOM	8841	N	GLN	173	72.719	42.640	-18.851	1.00	73.42	H21B	N
ATOM	8842	CA	GLN	173	72.562	43.655	-17.823	1.00	77.15	H21B	C
ATOM	8843	CB	GLN	173	72.867	45.038	-18.404	1.00	79.75	H21B	C
ATOM	8844	CG	GLN	173	74.312	45.248	-18.827	1.00	83.55	H21B	C
ATOM	8845	CD	GLN	173	75.172	45.782	-17.698	1.00	85.89	H21B	C
ATOM	8846	OE1	GLN	173	74.729	45.856	-16.548	1.00	86.83	H21B	O
ATOM	8847	NE2	GLN	173	76.410	46.159	-18.020	1.00	87.03	H21B	N
ATOM	8848	C	GLN	173	71.166	43.671	-17.216	1.00	78.60	H21B	C
ATOM	8849	O	GLN	173	70.259	42.965	-17.669	1.00	79.70	H21B	O
ATOM	8850	N	SER	174	71.007	44.489	-16.181	1.00	79.33	H21B	N
ATOM	8851	CA	SER	174	69.697	44.771	-15.607	1.00	79.53	H21B	C
ATOM	8852	CB	SER	174	69.846	45.630	-14.344	1.00	81.67	H21B	C
ATOM	8853	OG	SER	174	70.638	46.782	-14.601	1.00	84.74	H21B	O
ATOM	8854	C	SER	174	68.839	45.509	-16.629	1.00	78.22	H21B	C
ATOM	8855	O	SER	174	67.741	45.967	-16.306	1.00	77.64	H21B	O
ATOM	8856	N	SER	175	69.352	45.623	-17.855	1.00	76.80	H21B	N
ATOM	8857	CA	SER	175	68.678	46.367	-18.916	1.00	74.39	H21B	C
ATOM	8858	CB	SER	175	69.153	47.826	-18.914	1.00	74.01	H21B	C
ATOM	8859	OG	SER	175	70.555	47.916	-19.114	1.00	75.07	H21B	O
ATOM	8860	C	SER	175	68.868	45.759	-20.311	1.00	72.21	H21B	C
ATOM	8861	O	SER	175	69.097	46.469	-21.280	1.00	71.78	H21B	O
ATOM	8862	N	GLY	176	68.769	44.442	-20.415	1.00	70.15	H21B	N
ATOM	8863	CA	GLY	176	68.736	43.815	-21.728	1.00	66.86	H21B	C
ATOM	8864	C	GLY	176	70.029	43.889	-22.527	1.00	65.20	H21B	C
ATOM	8865	O	GLY	176	70.210	43.127	-23.485	1.00	64.83	H21B	O
ATOM	8866	N	LEU	177	70.931	44.791	-22.137	1.00	62.07	H21B	N
ATOM	8867	CA	LEU	177	72.141	45.050	-22.913	1.00	57.24	H21B	C
ATOM	8868	CB	LEU	177	72.404	46.530	-22.838	1.00	57.31	H21B	C
ATOM	8869	CG	LEU	177	71.501	47.426	-23.584	1.00	57.05	H21B	C
ATOM	8870	CD1	LEU	177	71.938	48.874	-23.476	1.00	57.20	H21B	C
ATOM	8871	CD2	LEU	177	71.436	46.997	-25.032	1.00	57.12	H21B	C
ATOM	8872	C	LEU	177	73.357	44.217	-22.511	1.00	54.53	H21B	C
ATOM	8873	O	LEU	177	73.426	43.700	-21.405	1.00	54.83	H21B	O
ATOM	8874	N	TYR	178	74.312	44.093	-23.428	1.00	51.17	H21B	N
ATOM	8875	CA	TYR	178	75.484	43.241	-23.240	1.00	48.45	H21B	C
ATOM	8876	CB	TYR	178	75.706	42.398	-24.488	1.00	50.40	H21B	C
ATOM	8877	CG	TYR	178	74.710	41.284	-24.665	1.00	54.91	H21B	C
ATOM	8878	CD1	TYR	178	75.084	39.954	-24.470	1.00	56.36	H21B	C
ATOM	8879	CE1	TYR	178	74.190	38.922	-24.662	1.00	55.89	H21B	C
ATOM	8880	CD2	TYR	178	73.400	41.547	-25.057	1.00	56.50	H21B	C
ATOM	8881	CE2	TYR	178	72.499	40.513	-25.251	1.00	57.35	H21B	C
ATOM	8882	CZ	TYR	178	72.904	39.206	-25.052	1.00	56.84	H21B	C
ATOM	8883	OH	TYR	178	72.014	38.179	-25.249	1.00	58.94	H21B	O
ATOM	8884	C	TYR	178	76.790	43.988	-22.922	1.00	45.48	H21B	C
ATOM	8885	O	TYR	178	76.992	45.136	-23.331	1.00	42.53	H21B	O
ATOM	8886	N	SER	179	77.678	43.312	-22.203	1.00	42.07	H21B	N
ATOM	8887	CA	SER	179	79.030	43.814	-21.992	1.00	41.79	H21B	C
ATOM	8888	CB	SER	179	79.101	44.678	-20.721	1.00	40.35	H21B	C
ATOM	8889	OG	SER	179	78.480	45.938	-20.886	1.00	39.39	H21B	O
ATOM	8890	C	SER	179	80.025	42.657	-21.868	1.00	40.86	H21B	C

ATOM	8891	O	SER	179	79.888	41.808	-20.998	1.00	41.44	H21B	O
ATOM	8892	N	LEU	180	81.025	42.615	-22.740	1.00	39.48	H21B	N
ATOM	8893	CA	LEU	180	82.168	41.752	-22.495	1.00	38.55	H21B	C
ATOM	8894	CB	LEU	180	82.356	40.749	-23.644	1.00	38.54	H21B	C
ATOM	8895	CG	LEU	180	82.521	41.258	-25.080	1.00	39.73	H21B	C
ATOM	8896	CD1	LEU	180	83.775	42.126	-25.212	1.00	40.85	H21B	C
ATOM	8897	CD2	LEU	180	82.624	40.063	-26.010	1.00	39.95	H21B	C
ATOM	8898	C	LEU	180	83.403	42.628	-22.334	1.00	38.44	H21B	C
ATOM	8899	O	LEU	180	83.321	43.848	-22.463	1.00	38.53	H21B	O
ATOM	8900	N	SER	181	84.540	42.011	-22.033	1.00	37.53	H21B	N
ATOM	8901	CA	SER	181	85.803	42.731	-22.001	1.00	37.68	H21B	C
ATOM	8902	CB	SER	181	86.069	43.318	-20.607	1.00	37.05	H21B	C
ATOM	8903	OG	SER	181	85.701	42.431	-19.566	1.00	37.89	H21B	O
ATOM	8904	C	SER	181	86.931	41.809	-22.408	1.00	38.52	H21B	C
ATOM	8905	O	SER	181	86.928	40.639	-22.067	1.00	38.90	H21B	O
ATOM	8906	N	SER	182	87.881	42.334	-23.174	1.00	39.83	H21B	N
ATOM	8907	CA	SER	182	88.983	41.528	-23.659	1.00	39.81	H21B	C
ATOM	8908	CB	SER	182	89.139	41.681	-25.166	1.00	39.45	H21B	C
ATOM	8909	OG	SER	182	90.164	40.820	-25.641	1.00	41.46	H21B	O
ATOM	8910	C	SER	182	90.262	41.955	-22.964	1.00	40.20	H21B	C
ATOM	8911	O	SER	182	90.544	43.149	-22.828	1.00	38.40	H21B	O
ATOM	8912	N	VAL	183	91.022	40.967	-22.507	1.00	40.90	H21B	N
ATOM	8913	CA	VAL	183	92.246	41.226	-21.767	1.00	42.48	H21B	C
ATOM	8914	CB	VAL	183	92.213	40.579	-20.356	1.00	41.50	H21B	C
ATOM	8915	CG1	VAL	183	93.340	39.579	-20.229	1.00	41.59	H21B	C
ATOM	8916	CG2	VAL	183	92.335	41.636	-19.273	1.00	40.00	H21B	C
ATOM	8917	C	VAL	183	93.384	40.596	-22.544	1.00	43.84	H21B	C
ATOM	8918	O	VAL	183	93.262	39.464	-23.007	1.00	43.94	H21B	O
ATOM	8919	N	VAL	184	94.486	41.321	-22.694	1.00	44.89	H21B	N
ATOM	8920	CA	VAL	184	95.714	40.683	-23.142	1.00	47.11	H21B	C
ATOM	8921	CB	VAL	184	96.204	41.307	-24.477	1.00	45.90	H21B	C
ATOM	8922	CG1	VAL	184	96.653	42.736	-24.269	1.00	45.60	H21B	C
ATOM	8923	CG2	VAL	184	97.322	40.469	-25.050	1.00	46.59	H21B	C
ATOM	8924	C	VAL	184	96.786	40.813	-22.053	1.00	49.55	H21B	C
ATOM	8925	O	VAL	184	96.842	41.827	-21.353	1.00	48.61	H21B	O
ATOM	8926	N	THR	185	97.604	39.775	-21.887	1.00	51.80	H21B	N
ATOM	8927	CA	THR	185	98.753	39.851	-20.986	1.00	54.28	H21B	C
ATOM	8928	CB	THR	185	98.912	38.581	-20.125	1.00	55.24	H21B	C
ATOM	8929	OG1	THR	185	99.015	37.434	-20.977	1.00	57.32	H21B	O
ATOM	8930	CG2	THR	185	97.728	38.409	-19.188	1.00	56.03	H21B	C
ATOM	8931	C	THR	185	100.028	40.027	-21.791	1.00	57.09	H21B	C
ATOM	8932	O	THR	185	100.246	39.332	-22.793	1.00	56.47	H21B	O
ATOM	8933	N	VAL	186	100.863	40.961	-21.340	1.00	60.48	H21B	N
ATOM	8934	CA	VAL	186	102.081	41.354	-22.045	1.00	63.14	H21B	C
ATOM	8935	CB	VAL	186	101.834	42.577	-22.931	1.00	63.28	H21B	C
ATOM	8936	CG1	VAL	186	100.857	42.232	-24.023	1.00	63.21	H21B	C
ATOM	8937	CG2	VAL	186	101.306	43.725	-22.082	1.00	63.11	H21B	C
ATOM	8938	C	VAL	186	103.187	41.719	-21.059	1.00	67.74	H21B	C
ATOM	8939	O	VAL	186	102.917	42.082	-19.909	1.00	67.37	H21B	O
ATOM	8940	N	PRO	187	104.453	41.639	-21.505	1.00	72.12	H21B	N
ATOM	8941	CD	PRO	187	104.854	41.036	-22.788	1.00	71.71	H21B	C
ATOM	8942	CA	PRO	187	105.618	42.055	-20.718	1.00	73.03	H21B	C
ATOM	8943	CB	PRO	187	106.789	41.807	-21.659	1.00	72.51	H21B	C
ATOM	8944	CG	PRO	187	106.303	40.741	-22.576	1.00	71.49	H21B	C
ATOM	8945	C	PRO	187	105.524	43.514	-20.312	1.00	76.95	H21B	C
ATOM	8946	O	PRO	187	105.373	44.389	-21.165	1.00	78.69	H21B	O
ATOM	8947	N	SER	188	105.619	43.767	-19.009	1.00	80.50	H21B	N
ATOM	8948	CA	SER	188	105.532	45.124	-18.464	1.00	82.60	H21B	C
ATOM	8949	CB	SER	188	105.372	45.064	-16.946	1.00	82.32	H21B	C
ATOM	8950	OG	SER	188	106.304	44.157	-16.387	1.00	81.51	H21B	O
ATOM	8951	C	SER	188	106.756	45.969	-18.819	1.00	83.71	H21B	C
ATOM	8952	O	SER	188	106.805	47.170	-18.543	1.00	84.16	H21B	O
ATOM	8953	N	SER	189	107.746	45.321	-19.420	1.00	84.52	H21B	N
ATOM	8954	CA	SER	189	108.873	46.014	-20.027	1.00	85.15	H21B	C
ATOM	8955	CB	SER	189	109.991	45.015	-20.320	1.00	84.73	H21B	C
ATOM	8956	OG	SER	189	109.521	43.969	-21.154	1.00	84.01	H21B	O
ATOM	8957	C	SER	189	108.413	46.671	-21.328	1.00	86.33	H21B	C
ATOM	8958	O	SER	189	108.841	47.777	-21.675	1.00	85.19	H21B	O
ATOM	8959	N	SER	190	107.530	45.982	-22.042	1.00	87.47	H21B	N
ATOM	8960	CA	SER	190	107.110	46.423	-23.361	1.00	88.98	H21B	C
ATOM	8961	CB	SER	190	106.444	45.262	-24.102	1.00	89.63	H21B	C
ATOM	8962	OG	SER	190	106.386	45.516	-25.496	1.00	90.22	H21B	O
ATOM	8963	C	SER	190	106.156	47.615	-23.282	1.00	88.95	H21B	C
ATOM	8964	O	SER	190	105.962	48.334	-24.264	1.00	89.75	H21B	O
ATOM	8965	N	LEU	191	105.562	47.826	-22.113	1.00	88.06	H21B	N
ATOM	8966	CA	LEU	191	104.711	48.990	-21.912	1.00	87.45	H21B	C

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ATOM   8967  CB  LEU   191     104.184  49.024 -20.474  1.00 86.50      H21B C
ATOM   8968  CG  LEU   191     103.411  47.794 -19.980  1.00 84.98      H21B C
ATOM   8969  CD1 LEU   191     103.043  47.973 -18.519  1.00 83.28      H21B C
ATOM   8970  CD2 LEU   191     102.159  47.595 -20.819  1.00 84.31      H21B C
ATOM   8971  C   LEU   191     105.534  50.241 -22.192  1.00 88.10      H21B C
ATOM   8972  O   LEU   191     106.747  50.257 -21.974  1.00 88.98      H21B O
ATOM   8973  N   GLY   192     104.882  51.287 -22.686  1.00 88.53      H21B N
ATOM   8974  CA  GLY   192     105.593  52.525 -22.956  1.00 89.47      H21B C
ATOM   8975  C   GLY   192     106.283  52.577 -24.314  1.00 90.10      H21B C
ATOM   8976  O   GLY   192     106.279  53.623 -24.971  1.00 90.36      H21B O
ATOM   8977  N   THR   193     106.877  51.462 -24.739  1.00 89.29      H21B N
ATOM   8978  CA  THR   193     107.531  51.396 -26.043  1.00 88.82      H21B C
ATOM   8979  CB  THR   193     108.976  50.904 -25.928  1.00 88.80      H21B C
ATOM   8980  OG1 THR   193     108.990  49.601 -25.336  1.00 89.49      H21B O
ATOM   8981  CG2 THR   193     109.791  51.861 -25.080  1.00 88.67      H21B C
ATOM   8982  C   THR   193     106.795  50.468 -26.992  1.00 88.03      H21B C
ATOM   8983  O   THR   193     107.223  50.261 -28.129  1.00 88.87      H21B O
ATOM   8984  N   GLN   194     105.695  49.895 -26.515  1.00 88.58      H21B N
ATOM   8985  CA  GLN   194     104.795  49.153 -27.387  1.00 87.54      H21B C
ATOM   8986  CB  GLN   194     104.771  47.671 -27.010  1.00 87.84      H21B C
ATOM   8987  CG  GLN   194     103.816  46.833 -27.853  1.00 87.81      H21B C
ATOM   8988  CD  GLN   194     104.129  46.880 -29.333  1.00 87.69      H21B C
ATOM   8989  OE1 GLN   194     104.621  47.885 -29.844  1.00 88.63      H21B O
ATOM   8990  NE2 GLN   194     103.844  45.788 -30.033  1.00 88.04      H21B N
ATOM   8991  C   GLN   194     103.381  49.712 -27.337  1.00 86.21      H21B C
ATOM   8992  O   GLN   194     102.820  49.944 -26.264  1.00 85.43      H21B O
ATOM   8993  N   THR   195     102.813  49.934 -28.514  1.00 84.94      H21B N
ATOM   8994  CA  THR   195     101.447  50.408 -28.602  1.00 83.65      H21B C
ATOM   8995  CB  THR   195     101.205  51.195 -29.918  1.00 84.72      H21B C
ATOM   8996  OG1 THR   195     101.519  50.364 -31.043  1.00 84.98      H21B O
ATOM   8997  CG2 THR   195     102.077  52.443 -29.959  1.00 85.59      H21B C
ATOM   8998  C   THR   195     100.521  49.204 -28.547  1.00 80.85      H21B C
ATOM   8999  O   THR   195     100.749  48.207 -29.227  1.00 79.88      H21B O
ATOM   9000  N   TYR   196      99.487  49.289 -27.719  1.00 78.25      H21B N
ATOM   9001  CA  TYR   196      98.466  48.256 -27.700  1.00 76.01      H21B C
ATOM   9002  CB  TYR   196      98.425  47.577 -26.326  1.00 77.54      H21B C
ATOM   9003  CG  TYR   196      99.661  46.749 -26.046  1.00 79.65      H21B C
ATOM   9004  CD1 TYR   196     100.738  47.281 -25.342  1.00 81.11      H21B C
ATOM   9005  CE1 TYR   196     101.898  46.540 -25.132  1.00 81.59      H21B C
ATOM   9006  CD2 TYR   196      99.773  45.449 -26.528  1.00 80.81      H21B C
ATOM   9007  CE2 TYR   196     100.927  44.700 -26.323  1.00 81.26      H21B C
ATOM   9008  CZ  TYR   196     101.987  45.251 -25.628  1.00 81.91      H21B C
ATOM   9009  OH  TYR   196     103.143  44.523 -25.455  1.00 81.33      H21B O
ATOM   9010  C   TYR   196      97.109  48.849 -28.059  1.00 73.03      H21B C
ATOM   9011  O   TYR   196      96.591  49.728 -27.362  1.00 72.39      H21B O
ATOM   9012  N   ILE   197      96.553  48.374 -29.170  1.00 68.83      H21B N
ATOM   9013  CA  ILE   197      95.256  48.838 -29.634  1.00 64.42      H21B C
ATOM   9014  CB  ILE   197      95.323  49.442 -31.055  1.00 63.76      H21B C
ATOM   9015  CG2 ILE   197      93.937  49.836 -31.516  1.00 59.55      H21B C
ATOM   9016  CG1 ILE   197      96.218  50.676 -31.064  1.00 63.89      H21B C
ATOM   9017  CD1 ILE   197      96.068  51.503 -32.321  1.00 66.94      H21B C
ATOM   9018  C   ILE   197      94.266  47.698 -29.670  1.00 61.31      H21B C
ATOM   9019  O   ILE   197      94.502  46.665 -30.296  1.00 61.15      H21B O
ATOM   9020  N   CYS   198      93.148  47.904 -28.990  1.00 57.57      H21B N
ATOM   9021  CA  CYS   198      92.029  46.985 -29.052  1.00 52.01      H21B C
ATOM   9022  C   CYS   198      91.039  47.522 -30.085  1.00 47.01      H21B C
ATOM   9023  O   CYS   198      90.588  48.659 -30.004  1.00 44.41      H21B O
ATOM   9024  CB  CYS   198      91.394  46.878 -27.665  1.00 52.51      H21B C
ATOM   9025  SG  CYS   198      89.863  47.823 -27.408  1.00 57.12      H21B S
ATOM   9026  N   ASN   199      90.735  46.713 -31.088  1.00 44.35      H21B N
ATOM   9027  CA  ASN   199      89.775  47.131 -32.096  1.00 42.24      H21B C
ATOM   9028  CB  ASN   199      90.440  47.172 -33.472  1.00 41.57      H21B C
ATOM   9029  CG  ASN   199      91.497  46.126 -33.630  1.00 39.25      H21B C
ATOM   9030  OD1 ASN   199      91.202  44.937 -33.614  1.00 37.00      H21B O
ATOM   9031  ND2 ASN   199      92.743  46.560 -33.779  1.00 37.76      H21B N
ATOM   9032  C   ASN   199      88.541  46.228 -32.110  1.00 40.58      H21B C
ATOM   9033  O   ASN   199      88.638  45.017 -32.287  1.00 39.56      H21B O
ATOM   9034  N   VAL   200      87.384  46.855 -31.919  1.00 39.66      H21B N
ATOM   9035  CA  VAL   200      86.151  46.167 -31.582  1.00 40.25      H21B C
ATOM   9036  CB  VAL   200      85.492  46.801 -30.333  1.00 38.62      H21B C
ATOM   9037  CG1 VAL   200      84.361  45.920 -29.842  1.00 37.80      H21B C
ATOM   9038  CG2 VAL   200      86.530  47.024 -29.237  1.00 35.75      H21B C
ATOM   9039  C   VAL   200      85.173  46.252 -32.748  1.00 41.51      H21B C
ATOM   9040  O   VAL   200      85.173  47.212 -33.515  1.00 41.66      H21B O
ATOM   9041  N   ASN   201      84.338  45.239 -32.887  1.00 43.17      H21B N
ATOM   9042  CA  ASN   201      83.464  45.187 -34.034  1.00 46.02      H21B C
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ATOM   9043  CB  ASN   201      84.057  44.284 -35.117  1.00 50.18      H21B C
ATOM   9044  CG  ASN   201      84.087  44.956 -36.485  1.00 53.80      H21B C
ATOM   9045  OD1 ASN   201      83.056  45.422 -36.991  1.00 55.38      H21B O
ATOM   9046  ND2 ASN   201      85.273  45.011 -37.089  1.00 55.20      H21B N
ATOM   9047  C   ASN   201      82.092  44.686 -33.657  1.00 46.12      H21B C
ATOM   9048  O   ASN   201      81.946  43.641 -33.017  1.00 45.78      H21B O
ATOM   9049  N   HIS   202      81.082  45.440 -34.065  1.00 46.00      H21B N
ATOM   9050  CA  HIS   202      79.718  45.037 -33.829  1.00 46.36      H21B C
ATOM   9051  CB  HIS   202      79.070  45.968 -32.815  1.00 44.13      H21B C
ATOM   9052  CG  HIS   202      77.789  45.438 -32.269  1.00 42.81      H21B C
ATOM   9053  CD2 HIS   202      77.351  44.168 -32.119  1.00 40.00      H21B C
ATOM   9054  ND1 HIS   202      76.762  46.252 -31.849  1.00 43.89      H21B N
ATOM   9055  CE1 HIS   202      75.743  45.504 -31.468  1.00 42.77      H21B C
ATOM   9056  NE2 HIS   202      76.076  44.237 -31.623  1.00 40.89      H21B N
ATOM   9057  C   HIS   202      78.942  45.067 -35.135  1.00 48.29      H21B C
ATOM   9058  O   HIS   202      78.308  46.067 -35.461  1.00 48.19      H21B O
ATOM   9059  N   LYS   203      78.987  43.966 -35.880  1.00 50.80      H21B N
ATOM   9060  CA  LYS   203      78.283  43.915 -37.151  1.00 53.50      H21B C
ATOM   9061  CB  LYS   203      78.396  42.522 -37.777  1.00 54.50      H21B C
ATOM   9062  CG  LYS   203      79.807  42.159 -38.238  1.00 59.30      H21B C
ATOM   9063  CD  LYS   203      80.169  40.691 -37.915  1.00 63.35      H21B C
ATOM   9064  CE  LYS   203      79.219  39.692 -38.598  1.00 66.63      H21B C
ATOM   9065  NZ  LYS   203      79.247  38.318 -38.006  1.00 69.31      H21B N
ATOM   9066  C   LYS   203      76.815  44.301 -36.977  1.00 52.84      H21B C
ATOM   9067  O   LYS   203      76.298  45.132 -37.715  1.00 53.76      H21B O
ATOM   9068  N   PRO   204      76.129  43.728 -35.979  1.00 51.43      H21B N
ATOM   9069  CD  PRO   204      76.587  42.808 -34.926  1.00 50.05      H21B C
ATOM   9070  CA  PRO   204      74.703  44.037 -35.850  1.00 51.46      H21B C
ATOM   9071  CB  PRO   204      74.311  43.393 -34.523  1.00 48.83      H21B C
ATOM   9072  CG  PRO   204      75.294  42.308 -34.338  1.00 48.98      H21B C
ATOM   9073  C   PRO   204      74.460  45.533 -35.845  1.00 51.76      H21B C
ATOM   9074  O   PRO   204      73.498  46.021 -36.431  1.00 52.73      H21B O
ATOM   9075  N   SER   205      75.354  46.259 -35.187  1.00 52.99      H21B N
ATOM   9076  CA  SER   205      75.254  47.708 -35.096  1.00 53.35      H21B C
ATOM   9077  CB  SER   205      75.774  48.181 -33.744  1.00 53.63      H21B C
ATOM   9078  OG  SER   205      75.929  49.585 -33.733  1.00 54.28      H21B O
ATOM   9079  C   SER   205      76.016  48.431 -36.203  1.00 53.57      H21B C
ATOM   9080  O   SER   205      75.883  49.642 -36.354  1.00 53.69      H21B O
ATOM   9081  N   ASN   206      76.810  47.696 -36.975  1.00 54.02      H21B N
ATOM   9082  CA  ASN   206      77.655  48.315 -37.986  1.00 54.69      H21B C
ATOM   9083  CB  ASN   206      76.796  48.942 -39.077  1.00 52.53      H21B C
ATOM   9084  CG  ASN   206      76.283  47.923 -40.060  1.00 52.56      H21B C
ATOM   9085  OD1 ASN   206      75.156  48.030 -40.540  1.00 51.22      H21B O
ATOM   9086  ND2 ASN   206      77.112  46.921 -40.373  1.00 51.22      H21B N
ATOM   9087  C   ASN   206      78.556  49.373 -37.350  1.00 55.72      H21B C
ATOM   9088  O   ASN   206      78.725  50.477 -37.867  1.00 56.49      H21B O
ATOM   9089  N   THR   207      79.129  49.011 -36.211  1.00 56.16      H21B N
ATOM   9090  CA  THR   207      79.997  49.892 -35.464  1.00 55.85      H21B C
ATOM   9091  CB  THR   207      79.431  50.120 -34.056  1.00 55.15      H21B C
ATOM   9092  OG1 THR   207      78.105  50.651 -34.155  1.00 52.50      H21B O
ATOM   9093  CG2 THR   207      80.305  51.085 -33.282  1.00 53.38      H21B C
ATOM   9094  C   THR   207      81.368  49.238 -35.354  1.00 57.06      H21B C
ATOM   9095  O   THR   207      81.475  48.067 -35.000  1.00 56.67      H21B O
ATOM   9096  N   LYS   208      82.414  49.986 -35.675  1.00 58.26      H21B N
ATOM   9097  CA  LYS   208      83.765  49.517 -35.437  1.00 59.43      H21B C
ATOM   9098  CB  LYS   208      84.490  49.263 -36.756  1.00 62.37      H21B C
ATOM   9099  CG  LYS   208      85.846  48.597 -36.585  1.00 68.71      H21B C
ATOM   9100  CD  LYS   208      86.654  48.621 -37.877  1.00 76.76      H21B C
ATOM   9101  CE  LYS   208      86.050  47.703 -38.944  1.00 85.44      H21B C
ATOM   9102  NZ  LYS   208      84.704  48.149 -39.439  1.00 94.28      H21B N
ATOM   9103  C   LYS   208      84.519  50.558 -34.629  1.00 57.52      H21B C
ATOM   9104  O   LYS   208      84.623  51.721 -35.036  1.00 58.10      H21B O
ATOM   9105  N   VAL   209      85.029  50.136 -33.476  1.00 54.42      H21B N
ATOM   9106  CA  VAL   209      85.791  51.019 -32.604  1.00 50.92      H21B C
ATOM   9107  CB  VAL   209      85.220  51.053 -31.177  1.00 48.81      H21B C
ATOM   9108  CG1 VAL   209      86.086  51.946 -30.313  1.00 48.17      H21B C
ATOM   9109  CG2 VAL   209      83.807  51.556 -31.188  1.00 46.41      H21B C
ATOM   9110  C   VAL   209      87.236  50.578 -32.491  1.00 49.62      H21B C
ATOM   9111  O   VAL   209      87.520  49.408 -32.274  1.00 48.07      H21B O
ATOM   9112  N   ASP   210      88.156  51.521 -32.626  1.00 50.05      H21B N
ATOM   9113  CA  ASP   210      89.555  51.238 -32.331  1.00 51.15      H21B C
ATOM   9114  CB  ASP   210      90.434  51.542 -33.551  1.00 53.47      H21B C
ATOM   9115  CG  ASP   210      90.104  50.655 -34.753  1.00 57.01      H21B C
ATOM   9116  OD1 ASP   210      89.597  49.530 -34.558  1.00 57.51      H21B O
ATOM   9117  OD2 ASP   210      90.351  51.085 -35.903  1.00 60.50      H21B O
ATOM   9118  C   ASP   210      89.999  52.068 -31.128  1.00 49.56      H21B C
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ATOM   9119  O   ASP 210      89.709  53.260 -31.033  1.00 47.85      H21B O
ATOM   9120  N   LYS 211      90.688  51.424 -30.197  1.00 48.61      H21B N
ATOM   9121  CA  LYS 211      91.069  52.086 -28.969  1.00 50.47      H21B C
ATOM   9122  CB  LYS 211      90.181  51.613 -27.828  1.00 48.97      H21B C
ATOM   9123  CG  LYS 211      90.526  52.218 -26.501  1.00 48.07      H21B C
ATOM   9124  CD  LYS 211      90.131  53.664 -26.478  1.00 51.05      H21B C
ATOM   9125  CE  LYS 211      88.626  53.799 -26.489  1.00 53.67      H21B C
ATOM   9126  NZ  LYS 211      88.191  55.222 -26.529  1.00 57.46      H21B N
ATOM   9127  C   LYS 211      92.518  51.784 -28.647  1.00 52.98      H21B C
ATOM   9128  O   LYS 211      92.950  50.644 -28.718  1.00 52.69      H21B O
ATOM   9129  N   LYS 212      93.276  52.816 -28.304  1.00 56.95      H21B N
ATOM   9130  CA  LYS 212      94.631  52.615 -27.835  1.00 59.46      H21B C
ATOM   9131  CB  LYS 212      95.538  53.764 -28.284  1.00 61.77      H21B C
ATOM   9132  CG  LYS 212      96.779  53.957 -27.410  1.00 65.33      H21B C
ATOM   9133  CD  LYS 212      97.512  55.244 -27.743  1.00 70.28      H21B C
ATOM   9134  CE  LYS 212      98.557  55.037 -28.832  1.00 74.23      H21B C
ATOM   9135  NZ  LYS 212      99.718  54.250 -28.339  1.00 79.11      H21B N
ATOM   9136  C   LYS 212      94.581  52.557 -26.326  1.00 60.12      H21B C
ATOM   9137  O   LYS 212      94.098  53.481 -25.670  1.00 59.63      H21B O
ATOM   9138  N   VAL 213      95.070  51.458 -25.774  1.00 62.31      H21B N
ATOM   9139  CA  VAL 213      95.155  51.335 -24.329  1.00 65.85      H21B C
ATOM   9140  CB  VAL 213      94.782  49.926 -23.860  1.00 64.74      H21B C
ATOM   9141  CG1 VAL 213      94.776  49.873 -22.337  1.00 62.94      H21B C
ATOM   9142  CG2 VAL 213      93.427  49.548 -24.422  1.00 63.10      H21B C
ATOM   9143  C   VAL 213      96.564  51.650 -23.875  1.00 68.37      H21B C
ATOM   9144  O   VAL 213      97.507  50.939 -24.208  1.00 66.85      H21B O
ATOM   9145  N   GLU 214      96.688  52.733 -23.120  1.00 73.64      H21B N
ATOM   9146  CA  GLU 214      97.975  53.188 -22.615  1.00 79.85      H21B C
ATOM   9147  CB  GLU 214      98.196  54.657 -22.980  1.00 80.72      H21B C
ATOM   9148  CG  GLU 214      98.423  54.944 -24.465  1.00 83.09      H21B C
ATOM   9149  CD  GLU 214      98.686  56.425 -24.731  1.00 84.01      H21B C
ATOM   9150  OE1 GLU 214      99.337  56.750 -25.748  1.00 84.45      H21B O
ATOM   9151  OE2 GLU 214      98.239  57.266 -23.916  1.00 84.49      H21B O
ATOM   9152  C   GLU 214      98.009  53.041 -21.100  1.00 83.59      H21B C
ATOM   9153  O   GLU 214      96.981  52.814 -20.463  1.00 83.78      H21B O
ATOM   9154  N   PRO 215      99.199  53.164 -20.505  1.00 86.98      H21B N
ATOM   9155  CD  PRO 215     100.495  52.995 -21.181  1.00 87.43      H21B C
ATOM   9156  CA  PRO 215      99.349  53.266 -19.049  1.00 91.94      H21B C
ATOM   9157  CB  PRO 215     100.858  53.356 -18.852  1.00 90.16      H21B C
ATOM   9158  CG  PRO 215     101.419  52.646 -20.051  1.00 88.99      H21B C
ATOM   9159  C   PRO 215      98.620  54.468 -18.466  1.00 96.26      H21B C
ATOM   9160  O   PRO 215      98.343  55.433 -19.167  1.00 96.59      H21B O
ATOM   9161  N   LYS 216      98.315  54.402 -17.177  1.00102.66      H21B N
ATOM   9162  CA  LYS 216      97.499  55.419 -16.521  1.00109.92      H21B C
ATOM   9163  CB  LYS 216      97.121  54.949 -15.109  1.00114.68      H21B C
ATOM   9164  CG  LYS 216      96.487  53.556 -15.053  1.00120.32      H21B C
ATOM   9165  CD  LYS 216      96.274  53.079 -13.613  1.00126.26      H21B C
ATOM   9166  CE  LYS 216      95.639  51.689 -13.570  1.00131.57      H21B C
ATOM   9167  NZ  LYS 216      95.637  51.080 -12.205  1.00135.66      H21B N
ATOM   9168  C   LYS 216      98.209  56.776 -16.448  1.00111.87      H21B C
ATOM   9169  O   LYS 216      98.520  57.236 -15.326  1.00113.12      H21B O
ATOM   9170  OXT LYS 216      98.451  57.368 -17.519  1.00114.65      H21B O
TER    9171       LYS 216                                             H21B
END
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TABLE 35.4

ATOM	1	CB	TRP	453	-29.096	34.072	11.297	1.00	62.90	A	C
ATOM	2	CG	TRP	453	-28.288	34.936	10.370	1.00	64.93	A	C
ATOM	3	CD2	TRP	453	-27.397	34.492	9.339	1.00	66.49	A	C
ATOM	4	CE2	TRP	453	-26.843	35.642	8.742	1.00	67.05	A	C
ATOM	5	CE3	TRP	453	-27.013	33.233	8.865	1.00	67.25	A	C
ATOM	6	CD1	TRP	453	-28.241	36.301	10.353	1.00	65.59	A	C
ATOM	7	NE1	TRP	453	-27.375	36.733	9.378	1.00	65.65	A	N
ATOM	8	CZ2	TRP	453	-25.924	35.571	7.694	1.00	67.23	A	C
ATOM	9	CZ3	TRP	453	-26.100	33.165	7.824	1.00	67.18	A	C
ATOM	10	CH2	TRP	453	-25.566	34.327	7.252	1.00	67.55	A	C
ATOM	11	C	TRP	453	-27.142	33.092	12.528	1.00	59.11	A	C
ATOM	12	O	TRP	453	-26.061	33.624	12.777	1.00	59.88	A	O
ATOM	13	N	TRP	453	-28.207	35.189	13.329	1.00	61.20	A	N
ATOM	14	CA	TRP	453	-28.450	33.869	12.675	1.00	60.89	A	C
ATOM	15	N	GLN	454	-27.248	31.827	12.137	1.00	55.81	A	N
ATOM	16	CA	GLN	454	-26.072	30.981	11.962	1.00	52.32	A	C
ATOM	17	CB	GLN	454	-26.117	29.821	12.962	1.00	54.31	A	C
ATOM	18	CG	GLN	454	-25.914	28.448	12.353	1.00	57.49	A	C
ATOM	19	CD	GLN	454	-24.637	27.773	12.827	1.00	59.72	A	C
ATOM	20	OE1	GLN	454	-23.617	28.430	13.055	1.00	60.00	A	O
ATOM	21	NE2	GLN	454	-24.688	26.450	12.973	1.00	59.65	A	N
ATOM	22	C	GLN	454	-26.011	30.456	10.530	1.00	47.73	A	C
ATOM	23	O	GLN	454	-27.041	30.329	9.867	1.00	46.06	A	O
ATOM	24	N	LEU	455	-24.808	30.158	10.049	1.00	43.78	A	N
ATOM	25	CA	LEU	455	-24.651	29.736	8.658	1.00	41.00	A	C
ATOM	26	CB	LEU	455	-23.386	30.356	8.063	1.00	41.51	A	C
ATOM	27	CG	LEU	455	-23.082	30.107	6.583	1.00	41.00	A	C
ATOM	28	CD1	LEU	455	-24.272	30.468	5.716	1.00	39.45	A	C
ATOM	29	CD2	LEU	455	-21.874	30.934	6.191	1.00	42.57	A	C
ATOM	30	C	LEU	455	-24.602	28.217	8.531	1.00	38.55	A	C
ATOM	31	O	LEU	455	-23.744	27.562	9.122	1.00	38.78	A	O
ATOM	32	N	PHE	456	-25.540	27.665	7.766	1.00	34.53	A	N
ATOM	33	CA	PHE	456	-25.605	26.226	7.524	1.00	32.19	A	C
ATOM	34	CB	PHE	456	-27.039	25.715	7.711	1.00	30.96	A	C
ATOM	35	CG	PHE	456	-27.517	25.742	9.135	1.00	30.83	A	C
ATOM	36	CD1	PHE	456	-27.303	24.657	9.973	1.00	29.93	A	C
ATOM	37	CD2	PHE	456	-28.186	26.850	9.636	1.00	30.09	A	C
ATOM	38	CE1	PHE	456	-27.749	24.675	11.291	1.00	30.85	A	C
ATOM	39	CE2	PHE	456	-28.635	26.877	10.952	1.00	31.15	A	C
ATOM	40	CZ	PHE	456	-28.417	25.789	11.781	1.00	29.38	A	C
ATOM	41	C	PHE	456	-25.154	25.915	6.100	1.00	30.99	A	C
ATOM	42	O	PHE	456	-25.649	26.508	5.148	1.00	30.86	A	O
ATOM	43	N	CYS	457	-24.222	24.981	5.957	1.00	30.33	A	N
ATOM	44	CA	CYS	457	-23.812	24.518	4.636	1.00	29.83	A	C
ATOM	45	C	CYS	457	-23.729	23.002	4.598	1.00	29.43	A	C
ATOM	46	O	CYS	457	-23.477	22.357	5.617	1.00	30.24	A	O
ATOM	47	CB	CYS	457	-22.442	25.067	4.260	1.00	30.42	A	C
ATOM	48	SG	CYS	457	-22.270	26.868	4.126	1.00	33.48	A	S
ATOM	49	N	ARG	458	-23.932	22.440	3.412	1.00	28.73	A	N
ATOM	50	CA	ARG	458	-23.732	21.013	3.192	1.00	27.98	A	C
ATOM	51	CB	ARG	458	-25.082	20.295	3.128	1.00	27.46	A	C
ATOM	52	CG	ARG	458	-25.982	20.816	2.021	1.00	27.71	A	C
ATOM	53	CD	ARG	458	-27.371	20.197	2.092	1.00	28.19	A	C
ATOM	54	NE	ARG	458	-28.104	20.474	0.864	1.00	28.26	A	N
ATOM	55	CZ	ARG	458	-28.211	19.619	-0.150	1.00	28.99	A	C
ATOM	56	NH1	ARG	458	-28.891	19.965	-1.235	1.00	28.33	A	N
ATOM	57	NH2	ARG	458	-27.651	18.416	-0.073	1.00	26.24	A	N
ATOM	58	C	ARG	458	-22.969	20.809	1.889	1.00	26.26	A	C
ATOM	59	O	ARG	458	-22.870	21.721	1.074	1.00	25.98	A	O
ATOM	60	N	THR	459	-22.425	19.612	1.705	1.00	24.49	A	N
ATOM	61	CA	THR	459	-21.655	19.296	0.520	1.00	24.90	A	C
ATOM	62	CB	THR	459	-20.334	18.588	0.898	1.00	25.10	A	C
ATOM	63	OG1	THR	459	-19.493	19.502	1.608	1.00	25.42	A	O
ATOM	64	CG2	THR	459	-19.609	18.094	-0.344	1.00	22.93	A	C
ATOM	65	C	THR	459	-22.476	18.383	-0.379	1.00	26.20	A	C
ATOM	66	O	THR	459	-23.093	17.428	0.093	1.00	24.85	A	O
ATOM	67	N	VAL	460	-22.493	18.690	-1.672	1.00	26.69	A	N
ATOM	68	CA	VAL	460	-23.209	17.869	-2.637	1.00	27.29	A	C
ATOM	69	CB	VAL	460	-24.313	18.676	-3.343	1.00	29.26	A	C
ATOM	70	CG1	VAL	460	-24.981	17.823	-4.419	1.00	26.28	A	C
ATOM	71	CG2	VAL	460	-25.339	19.146	-2.317	1.00	30.35	A	C
ATOM	72	C	VAL	460	-22.264	17.321	-3.693	1.00	27.26	A	C
ATOM	73	O	VAL	460	-21.661	18.078	-4.451	1.00	29.73	A	O
ATOM	74	N	TRP	461	-22.140	16.001	-3.740	1.00	26.25	A	N

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ATOM     75  CA  TRP   461     -21.368  15.351  -4.788  1.00 26.76      A    C
ATOM     76  CB  TRP   461     -20.764  14.038  -4.268  1.00 24.57      A    C
ATOM     77  CG  TRP   461     -19.606  14.235  -3.334  1.00 24.74      A    C
ATOM     78  CD2 TRP   461     -18.218  13.984  -3.613  1.00 24.43      A    C
ATOM     79  CE2 TRP   461     -17.494  14.304  -2.447  1.00 25.75      A    C
ATOM     80  CE3 TRP   461     -17.520  13.521  -4.736  1.00 26.44      A    C
ATOM     81  CD1 TRP   461     -19.660  14.684  -2.046  1.00 23.71      A    C
ATOM     82  NE1 TRP   461     -18.397  14.729  -1.507  1.00 25.76      A    N
ATOM     83  CZ2 TRP   461     -16.103  14.176  -2.368  1.00 28.09      A    C
ATOM     84  CZ3 TRP   461     -16.135  13.392  -4.657  1.00 27.37      A    C
ATOM     85  CH2 TRP   461     -15.444  13.719  -3.480  1.00 26.66      A    C
ATOM     86  C   TRP   461     -22.264  15.076  -5.998  1.00 27.35      A    C
ATOM     87  O   TRP   461     -23.406  14.629  -5.850  1.00 26.99      A    O
ATOM     88  N   SER   462     -21.745  15.364  -7.189  1.00 25.48      A    N
ATOM     89  CA  SER   462     -22.428  15.031  -8.432  1.00 25.19      A    C
ATOM     90  CB  SER   462     -21.732  15.698  -9.626  1.00 24.60      A    C
ATOM     91  OG  SER   462     -20.463  15.099  -9.866  1.00 24.13      A    O
ATOM     92  C   SER   462     -22.384  13.528  -8.627  1.00 24.96      A    C
ATOM     93  O   SER   462     -21.641  12.835  -7.947  1.00 26.57      A    O
ATOM     94  N   ALA   463     -23.176  13.030  -9.566  1.00 26.08      A    N
ATOM     95  CA  ALA   463     -22.921  11.714 -10.139  1.00 27.25      A    C
ATOM     96  CB  ALA   463     -24.042  11.351 -11.106  1.00 24.94      A    C
ATOM     97  C   ALA   463     -21.578  11.774 -10.883  1.00 26.16      A    C
ATOM     98  O   ALA   463     -21.131  12.849 -11.278  1.00 25.40      A    O
ATOM     99  N   HIS   464     -20.939  10.625 -11.066  1.00 26.74      A    N
ATOM    100  CA  HIS   464     -19.723  10.539 -11.873  1.00 29.13      A    C
ATOM    101  CB  HIS   464     -19.297   9.073 -11.991  1.00 30.10      A    C
ATOM    102  CG  HIS   464     -17.948   8.876 -12.610  1.00 31.56      A    C
ATOM    103  CD2 HIS   464     -16.713   8.823 -12.055  1.00 30.82      A    C
ATOM    104  ND1 HIS   464     -17.772   8.653 -13.959  1.00 32.23      A    N
ATOM    105  CE1 HIS   464     -16.488   8.466 -14.208  1.00 31.96      A    C
ATOM    106  NE2 HIS   464     -15.824   8.565 -13.068  1.00 31.64      A    N
ATOM    107  C   HIS   464     -19.977  11.125 -13.272  1.00 29.91      A    C
ATOM    108  O   HIS   464     -21.080  11.013 -13.804  1.00 32.59      A    O
ATOM    109  N   SER   465     -18.959  11.738 -13.870  1.00 28.42      A    N
ATOM    110  CA  SER   465     -19.143  12.475 -15.126  1.00 27.55      A    C
ATOM    111  CB  SER   465     -17.936  13.372 -15.405  1.00 26.69      A    C
ATOM    112  OG  SER   465     -16.774  12.600 -15.663  1.00 26.28      A    O
ATOM    113  C   SER   465     -19.348  11.580 -16.337  1.00 27.89      A    C
ATOM    114  O   SER   465     -19.898  12.012 -17.351  1.00 30.42      A    O
ATOM    115  N   GLY   466     -18.890  10.340 -16.247  1.00 24.79      A    N
ATOM    116  CA  GLY   466     -18.715   9.563 -17.457  1.00 22.23      A    C
ATOM    117  C   GLY   466     -17.259   9.660 -17.871  1.00 25.15      A    C
ATOM    118  O   GLY   466     -16.583  10.644 -17.547  1.00 22.33      A    O
ATOM    119  N   PRO   467     -16.741   8.642 -18.574  1.00 25.86      A    N
ATOM    120  CD  PRO   467     -17.448   7.387 -18.894  1.00 26.23      A    C
ATOM    121  CA  PRO   467     -15.304   8.535 -18.831  1.00 24.52      A    C
ATOM    122  CB  PRO   467     -15.095   7.030 -18.999  1.00 25.31      A    C
ATOM    123  CG  PRO   467     -16.374   6.532 -19.524  1.00 24.11      A    C
ATOM    124  C   PRO   467     -14.787   9.321 -20.029  1.00 24.85      A    C
ATOM    125  O   PRO   467     -13.596   9.522 -20.152  1.00 25.40      A    O
ATOM    126  N   THR   468     -15.660   9.762 -20.923  1.00 24.02      A    N
ATOM    127  CA  THR   468     -15.174  10.270 -22.201  1.00 23.19      A    C
ATOM    128  CB  THR   468     -16.305  10.341 -23.242  1.00 23.48      A    C
ATOM    129  OG1 THR   468     -17.407  11.085 -22.706  1.00 23.61      A    O
ATOM    130  CG2 THR   468     -16.766   8.946 -23.608  1.00 22.59      A    C
ATOM    131  C   THR   468     -14.503  11.640 -22.087  1.00 24.90      A    C
ATOM    132  O   THR   468     -14.594  12.310 -21.060  1.00 23.76      A    O
ATOM    133  N   ARG   469     -13.832  12.044 -23.159  1.00 23.46      A    N
ATOM    134  CA  ARG   469     -12.874  13.140 -23.115  1.00 24.59      A    C
ATOM    135  CB  ARG   469     -12.178  13.250 -24.476  1.00 25.09      A    C
ATOM    136  CG  ARG   469     -11.111  14.318 -24.558  1.00 26.25      A    C
ATOM    137  CD  ARG   469     -10.439  14.323 -25.935  1.00 27.32      A    C
ATOM    138  NE  ARG   469      -9.440  15.382 -26.040  1.00 27.17      A    N
ATOM    139  CZ  ARG   469      -8.581  15.512 -27.046  1.00 25.09      A    C
ATOM    140  NH1 ARG   469      -7.711  16.514 -27.044  1.00 24.35      A    N
ATOM    141  NH2 ARG   469      -8.591  14.644 -28.047  1.00 23.06      A    N
ATOM    142  C   ARG   469     -13.495  14.487 -22.736  1.00 25.75      A    C
ATOM    143  O   ARG   469     -12.891  15.276 -22.009  1.00 23.47      A    O
ATOM    144  N   MET   470     -14.698  14.759 -23.230  1.00 24.77      A    N
ATOM    145  CA  MET   470     -15.328  16.047 -22.965  1.00 24.49      A    C
ATOM    146  CB  MET   470     -15.904  16.621 -24.262  1.00 24.57      A    C
ATOM    147  CG  MET   470     -15.563  18.079 -24.508  1.00 30.96      A    C
ATOM    148  SD  MET   470     -13.788  18.409 -24.583  1.00 31.07      A    S
ATOM    149  CE  MET   470     -13.271  17.434 -25.959  1.00 31.18      A    C
ATOM    150  C   MET   470     -16.425  15.926 -21.908  1.00 24.68      A    C
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ATOM    151  O   MET   470     -17.226  16.848 -21.717  1.00 23.62      A    O
ATOM    152  N   ALA   471     -16.459  14.789 -21.220  1.00 24.16      A    N
ATOM    153  CA  ALA   471     -17.508  14.528 -20.238  1.00 24.20      A    C
ATOM    154  CB  ALA   471     -17.371  13.119 -19.687  1.00 20.75      A    C
ATOM    155  C   ALA   471     -17.452  15.537 -19.096  1.00 25.48      A    C
ATOM    156  O   ALA   471     -16.372  15.975 -18.693  1.00 27.11      A    O
ATOM    157  N   THR   472     -18.617  15.906 -18.578  1.00 24.57      A    N
ATOM    158  CA  THR   472     -18.675  16.811 -17.437  1.00 24.16      A    C
ATOM    159  CB  THR   472     -19.181  18.220 -17.836  1.00 24.01      A    C
ATOM    160  OG1 THR   472     -20.548  18.134 -18.257  1.00 21.91      A    O
ATOM    161  CG2 THR   472     -18.344  18.795 -18.970  1.00 18.59      A    C
ATOM    162  C   THR   472     -19.620  16.260 -16.386  1.00 25.66      A    C
ATOM    163  O   THR   472     -20.508  15.455 -16.690  1.00 24.53      A    O
ATOM    164  N   ALA   473     -19.415  16.700 -15.147  1.00 25.24      A    N
ATOM    165  CA  ALA   473     -20.319  16.393 -14.048  1.00 24.92      A    C
ATOM    166  CB  ALA   473     -19.560  15.661 -12.946  1.00 24.30      A    C
ATOM    167  C   ALA   473     -20.900  17.699 -13.513  1.00 25.27      A    C
ATOM    168  O   ALA   473     -20.241  18.741 -13.556  1.00 24.13      A    O
ATOM    169  N   ILE   474     -22.131  17.638 -13.012  1.00 24.89      A    N
ATOM    170  CA  ILE   474     -22.819  18.815 -12.489  1.00 25.52      A    C
ATOM    171  CB  ILE   474     -24.026  19.182 -13.365  1.00 29.25      A    C
ATOM    172  CG2 ILE   474     -24.883  20.220 -12.662  1.00 32.52      A    C
ATOM    173  CG1 ILE   474     -23.558  19.713 -14.721  1.00 33.35      A    C
ATOM    174  CD1 ILE   474     -22.979  21.105 -14.661  1.00 33.21      A    C
ATOM    175  C   ILE   474     -23.336  18.571 -11.069  1.00 26.48      A    C
ATOM    176  O   ILE   474     -24.055  17.603 -10.823  1.00 26.94      A    O
ATOM    177  N   ALA   475     -22.974  19.451 -10.140  1.00 25.26      A    N
ATOM    178  CA  ALA   475     -23.580  19.452  -8.811  1.00 25.92      A    C
ATOM    179  CB  ALA   475     -22.496  19.409  -7.732  1.00 23.88      A    C
ATOM    180  C   ALA   475     -24.459  20.695  -8.635  1.00 26.91      A    C
ATOM    181  O   ALA   475     -24.063  21.808  -8.998  1.00 25.49      A    O
ATOM    182  N   ARG   476     -25.655  20.490  -8.091  1.00 28.07      A    N
ATOM    183  CA  ARG   476     -26.642  21.558  -7.932  1.00 28.54      A    C
ATOM    184  CB  ARG   476     -27.883  21.299  -8.792  1.00 30.18      A    C
ATOM    185  CG  ARG   476     -27.672  21.375 -10.272  1.00 35.86      A    C
ATOM    186  CD  ARG   476     -29.005  21.304 -11.004  1.00 38.60      A    C
ATOM    187  NE  ARG   476     -28.838  21.409 -12.452  1.00 41.19      A    N
ATOM    188  CZ  ARG   476     -28.492  20.389 -13.231  1.00 43.32      A    C
ATOM    189  NH1 ARG   476     -28.277  19.191 -12.698  1.00 43.42      A    N
ATOM    190  NH2 ARG   476     -28.356  20.562 -14.538  1.00 44.05      A    N
ATOM    191  C   ARG   476     -27.093  21.615  -6.488  1.00 28.17      A    C
ATOM    192  O   ARG   476     -27.077  20.608  -5.783  1.00 27.75      A    O
ATOM    193  N   CYS   477     -27.519  22.797  -6.064  1.00 29.00      A    N
ATOM    194  CA  CYS   477     -28.103  22.979  -4.745  1.00 29.17      A    C
ATOM    195  C   CYS   477     -29.620  22.952  -4.848  1.00 30.65      A    C
ATOM    196  O   CYS   477     -30.173  23.016  -5.947  1.00 31.16      A    O
ATOM    197  CB  CYS   477     -27.664  24.316  -4.165  1.00 27.94      A    C
ATOM    198  SG  CYS   477     -25.866  24.479  -3.959  1.00 30.04      A    S
ATOM    199  N   ALA   478     -30.283  22.869  -3.699  1.00 30.57      A    N
ATOM    200  CA  ALA   478     -31.742  22.922  -3.643  1.00 31.86      A    C
ATOM    201  CB  ALA   478     -32.228  22.484  -2.260  1.00 29.25      A    C
ATOM    202  C   ALA   478     -32.240  24.333  -3.949  1.00 32.09      A    C
ATOM    203  O   ALA   478     -31.494  25.306  -3.847  1.00 31.96      A    O
ATOM    204  N   PRO   479     -33.521  24.461  -4.322  1.00 32.78      A    N
ATOM    205  CD  PRO   479     -34.502  23.383  -4.550  1.00 31.44      A    C
ATOM    206  CA  PRO   479     -34.054  25.778  -4.691  1.00 32.40      A    C
ATOM    207  CB  PRO   479     -35.515  25.486  -5.061  1.00 32.72      A    C
ATOM    208  CG  PRO   479     -35.513  24.037  -5.458  1.00 31.86      A    C
ATOM    209  C   PRO   479     -33.933  26.834  -3.592  1.00 32.60      A    C
ATOM    210  O   PRO   479     -33.819  28.022  -3.884  1.00 32.64      A    O
ATOM    211  N   ASP   480     -33.954  26.416  -2.331  1.00 31.47      A    N
ATOM    212  CA  ASP   480     -33.879  27.384  -1.241  1.00 33.46      A    C
ATOM    213  CB  ASP   480     -34.749  26.936  -0.057  1.00 35.16      A    C
ATOM    214  CG  ASP   480     -34.300  25.610   0.537  1.00 36.44      A    C
ATOM    215  OD1 ASP   480     -34.828  25.224   1.598  1.00 40.34      A    O
ATOM    216  OD2 ASP   480     -33.419  24.954  -0.053  1.00 38.40      A    O
ATOM    217  C   ASP   480     -32.449  27.623  -0.765  1.00 33.71      A    C
ATOM    218  O   ASP   480     -32.222  28.295   0.246  1.00 35.25      A    O
ATOM    219  N   GLU   481     -31.482  27.076  -1.492  1.00 32.10      A    N
ATOM    220  CA  GLU   481     -30.087  27.191  -1.086  1.00 30.90      A    C
ATOM    221  CB  GLU   481     -29.439  25.804  -1.016  1.00 30.17      A    C
ATOM    222  CG  GLU   481     -30.038  24.890   0.041  1.00 30.22      A    C
ATOM    223  CD  GLU   481     -29.692  23.425  -0.191  1.00 31.87      A    C
ATOM    224  OE1 GLU   481     -29.930  22.602   0.717  1.00 30.65      A    O
ATOM    225  OE2 GLU   481     -29.186  23.093  -1.285  1.00 33.77      A    O
ATOM    226  C   GLU   481     -29.292  28.073  -2.031  1.00 29.33      A    C
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ATOM    227  O   GLU   481     -29.723  28.354  -3.146  1.00 30.27      A    O
ATOM    228  N   GLU   482     -28.127  28.510  -1.569  1.00 29.51      A    N
ATOM    229  CA  GLU   482     -27.176  29.220  -2.411  1.00 29.75      A    C
ATOM    230  CB  GLU   482     -26.813  30.561  -1.777  1.00 30.90      A    C
ATOM    231  CG  GLU   482     -27.965  31.542  -1.682  1.00 35.54      A    C
ATOM    232  CD  GLU   482     -28.316  32.148  -3.031  1.00 38.85      A    C
ATOM    233  OE1 GLU   482     -27.426  32.200  -3.911  1.00 39.06      A    O
ATOM    234  OE2 GLU   482     -29.479  32.570  -3.212  1.00 39.52      A    O
ATOM    235  C   GLU   482     -25.915  28.372  -2.558  1.00 29.76      A    C
ATOM    236  O   GLU   482     -25.357  27.904  -1.569  1.00 28.44      A    O
ATOM    237  N   LEU   483     -25.464  28.176  -3.790  1.00 27.98      A    N
ATOM    238  CA  LEU   483     -24.174  27.546  -4.015  1.00 28.33      A    C
ATOM    239  CB  LEU   483     -24.101  26.996  -5.442  1.00 27.55      A    C
ATOM    240  CG  LEU   483     -22.796  26.296  -5.842  1.00 29.33      A    C
ATOM    241  CD1 LEU   483     -23.075  25.330  -6.976  1.00 29.06      A    C
ATOM    242  CD2 LEU   483     -21.746  27.326  -6.253  1.00 28.65      A    C
ATOM    243  C   LEU   483     -23.078  28.585  -3.793  1.00 28.13      A    C
ATOM    244  O   LEU   483     -22.993  29.566  -4.522  1.00 32.20      A    O
ATOM    245  N   LEU   484     -22.240  28.367  -2.789  1.00 27.32      A    N
ATOM    246  CA  LEU   484     -21.214  29.341  -2.437  1.00 26.58      A    C
ATOM    247  CB  LEU   484     -21.172  29.546  -0.916  1.00 25.53      A    C
ATOM    248  CG  LEU   484     -22.427  30.163  -0.281  1.00 27.36      A    C
ATOM    249  CD1 LEU   484     -22.065  30.781   1.072  1.00 27.53      A    C
ATOM    250  CD2 LEU   484     -23.013  31.234  -1.196  1.00 27.97      A    C
ATOM    251  C   LEU   484     -19.829  28.953  -2.943  1.00 27.62      A    C
ATOM    252  O   LEU   484     -18.926  29.793  -2.994  1.00 28.95      A    O
ATOM    253  N   SER   485     -19.655  27.688  -3.317  1.00 25.70      A    N
ATOM    254  CA  SER   485     -18.391  27.254  -3.908  1.00 27.74      A    C
ATOM    255  CB  SER   485     -17.304  27.106  -2.836  1.00 26.49      A    C
ATOM    256  OG  SER   485     -17.504  25.926  -2.068  1.00 28.56      A    O
ATOM    257  C   SER   485     -18.522  25.934  -4.664  1.00 27.50      A    C
ATOM    258  O   SER   485     -19.569  25.281  -4.643  1.00 25.09      A    O
ATOM    259  N   CYS   486     -17.432  25.548  -5.315  1.00 26.66      A    N
ATOM    260  CA  CYS   486     -17.457  24.490  -6.311  1.00 26.00      A    C
ATOM    261  C   CYS   486     -16.047  23.934  -6.434  1.00 26.11      A    C
ATOM    262  O   CYS   486     -15.097  24.677  -6.679  1.00 25.83      A    O
ATOM    263  CB  CYS   486     -17.932  25.090  -7.632  1.00 25.13      A    C
ATOM    264  SG  CYS   486     -17.774  24.120  -9.162  1.00 31.94      A    S
ATOM    265  N   SER   487     -15.912  22.628  -6.243  1.00 25.42      A    N
ATOM    266  CA  SER   487     -14.626  21.964  -6.403  1.00 24.57      A    C
ATOM    267  CB  SER   487     -14.031  21.612  -5.029  1.00 24.64      A    C
ATOM    268  OG  SER   487     -14.849  20.692  -4.324  1.00 27.88      A    O
ATOM    269  C   SER   487     -14.828  20.704  -7.237  1.00 23.91      A    C
ATOM    270  O   SER   487     -15.923  20.455  -7.734  1.00 22.75      A    O
ATOM    271  N   SER   488     -13.776  19.916  -7.408  1.00 24.08      A    N
ATOM    272  CA  SER   488     -13.883  18.706  -8.209  1.00 25.08      A    C
ATOM    273  CB  SER   488     -13.664  19.015  -9.698  1.00 27.25      A    C
ATOM    274  OG  SER   488     -12.416  19.654  -9.931  1.00 25.16      A    O
ATOM    275  C   SER   488     -12.864  17.693  -7.743  1.00 27.25      A    C
ATOM    276  O   SER   488     -11.940  18.027  -6.997  1.00 28.14      A    O
ATOM    277  N   PHE   489     -13.033  16.453  -8.182  1.00 26.11      A    N
ATOM    278  CA  PHE   489     -12.152  15.383  -7.757  1.00 26.68      A    C
ATOM    279  CB  PHE   489     -12.680  14.754  -6.462  1.00 26.29      A    C
ATOM    280  CG  PHE   489     -11.886  13.566  -5.992  1.00 28.17      A    C
ATOM    281  CD1 PHE   489     -10.732  13.738  -5.241  1.00 28.05      A    C
ATOM    282  CD2 PHE   489     -12.303  12.272  -6.291  1.00 28.40      A    C
ATOM    283  CE1 PHE   489     -10.004  12.637  -4.793  1.00 29.69      A    C
ATOM    284  CE2 PHE   489     -11.585  11.172  -5.849  1.00 29.13      A    C
ATOM    285  CZ  PHE   489     -10.434  11.354  -5.099  1.00 30.06      A    C
ATOM    286  C   PHE   489     -12.028  14.323  -8.837  1.00 27.40      A    C
ATOM    287  O   PHE   489     -13.009  13.951  -9.471  1.00 28.37      A    O
ATOM    288  N   SER   490     -10.807  13.846  -9.041  1.00 28.47      A    N
ATOM    289  CA  SER   490     -10.561  12.663  -9.853  1.00 30.35      A    C
ATOM    290  CB  SER   490      -9.863  13.056 -11.158  1.00 30.98      A    C
ATOM    291  OG  SER   490      -9.257  11.933 -11.776  1.00 33.81      A    O
ATOM    292  C   SER   490      -9.672  11.708  -9.066  1.00 31.89      A    C
ATOM    293  O   SER   490      -8.751  12.144  -8.378  1.00 32.56      A    O
ATOM    294  N   ARG   491      -9.941  10.412  -9.171  1.00 33.39      A    N
ATOM    295  CA  ARG   491      -9.110   9.402  -8.515  1.00 36.40      A    C
ATOM    296  CB  ARG   491      -9.709   8.007  -8.719  1.00 38.90      A    C
ATOM    297  CG  ARG   491     -11.109   7.848  -8.166  1.00 45.81      A    C
ATOM    298  CD  ARG   491     -11.268   6.514  -7.458  1.00 51.82      A    C
ATOM    299  NE  ARG   491     -12.322   6.565  -6.448  1.00 56.98      A    N
ATOM    300  CZ  ARG   491     -13.621   6.548  -6.731  1.00 58.08      A    C
ATOM    301  NH1 ARG   491     -14.518   6.597  -5.753  1.00 58.30      A    N
ATOM    302  NH2 ARG   491     -14.020   6.486  -7.995  1.00 57.04      A    N
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ATOM    303  C   ARG   491      -7.677   9.407  -9.050  1.00 35.22      A    C
ATOM    304  O   ARG   491      -6.731   9.192  -8.305  1.00 35.69      A    O
ATOM    305  N   SER   492      -7.533   9.654 -10.347  1.00 35.60      A    N
ATOM    306  CA  SER   492      -6.246   9.550 -11.024  1.00 34.80      A    C
ATOM    307  CB  SER   492      -6.459   9.060 -12.452  1.00 36.97      A    C
ATOM    308  OG  SER   492      -7.242   9.997 -13.179  1.00 38.02      A    O
ATOM    309  C   SER   492      -5.512  10.887 -11.068  1.00 35.54      A    C
ATOM    310  O   SER   492      -4.295  10.932 -11.258  1.00 36.68      A    O
ATOM    311  N   GLY   493      -6.254  11.977 -10.909  1.00 34.28      A    N
ATOM    312  CA  GLY   493      -5.654  13.292 -11.037  1.00 32.69      A    C
ATOM    313  C   GLY   493      -5.739  13.824 -12.456  1.00 31.91      A    C
ATOM    314  O   GLY   493      -5.287  14.929 -12.739  1.00 32.88      A    O
ATOM    315  N   LYS   494      -6.316  13.039 -13.357  1.00 31.63      A    N
ATOM    316  CA  LYS   494      -6.506  13.492 -14.728  1.00 32.29      A    C
ATOM    317  CB  LYS   494      -6.442  12.304 -15.693  1.00 34.42      A    C
ATOM    318  CG  LYS   494      -5.080  11.622 -15.758  1.00 38.00      A    C
ATOM    319  CD  LYS   494      -5.032  10.628 -16.909  1.00 39.72      A    C
ATOM    320  CE  LYS   494      -3.648  10.029 -17.078  1.00 42.01      A    C
ATOM    321  NZ  LYS   494      -3.585   9.143 -18.279  1.00 43.69      A    N
ATOM    322  C   LYS   494      -7.844  14.213 -14.870  1.00 29.91      A    C
ATOM    323  O   LYS   494      -8.870  13.604 -15.169  1.00 28.53      A    O
ATOM    324  N   ARG   495      -7.815  15.523 -14.659  1.00 29.57      A    N
ATOM    325  CA  ARG   495      -9.025  16.341 -14.669  1.00 28.53      A    C
ATOM    326  CB  ARG   495      -9.558  16.492 -13.243  1.00 29.81      A    C
ATOM    327  CG  ARG   495      -8.560  17.157 -12.306  1.00 31.40      A    C
ATOM    328  CD  ARG   495      -9.082  17.315 -10.881  1.00 32.62      A    C
ATOM    329  NE  ARG   495      -8.125  18.089 -10.096  1.00 34.88      A    N
ATOM    330  CZ  ARG   495      -8.424  19.180  -9.400  1.00 34.47      A    C
ATOM    331  NH1 ARG   495      -7.472  19.812  -8.733  1.00 35.19      A    N
ATOM    332  NH2 ARG   495      -9.672  19.633  -9.355  1.00 34.89      A    N
ATOM    333  C   ARG   495      -8.661  17.710 -15.225  1.00 27.43      A    C
ATOM    334  O   ARG   495      -7.486  18.039 -15.349  1.00 27.93      A    O
ATOM    335  N   ARG   496      -9.660  18.513 -15.564  1.00 26.03      A    N
ATOM    336  CA  ARG   496      -9.376  19.877 -15.972  1.00 24.54      A    C
ATOM    337  CB  ARG   496      -9.765  20.076 -17.436  1.00 24.73      A    C
ATOM    338  CG  ARG   496      -8.775  19.429 -18.396  1.00 26.29      A    C
ATOM    339  CD  ARG   496      -9.064  19.789 -19.843  1.00 25.61      A    C
ATOM    340  NE  ARG   496     -10.117  18.950 -20.405  1.00 24.57      A    N
ATOM    341  CZ  ARG   496     -10.601  19.095 -21.632  1.00 23.57      A    C
ATOM    342  NH1 ARG   496     -10.130  20.048 -22.429  1.00 20.57      A    N
ATOM    343  NH2 ARG   496     -11.555  18.287 -22.060  1.00 23.67      A    N
ATOM    344  C   ARG   496     -10.052  20.921 -15.084  1.00 25.65      A    C
ATOM    345  O   ARG   496     -10.266  22.062 -15.502  1.00 23.54      A    O
ATOM    346  N   GLY   497     -10.375  20.531 -13.853  1.00 25.25      A    N
ATOM    347  CA  GLY   497     -10.908  21.491 -12.900  1.00 25.50      A    C
ATOM    348  C   GLY   497     -12.406  21.709 -13.026  1.00 25.22      A    C
ATOM    349  O   GLY   497     -13.126  20.844 -13.524  1.00 24.92      A    O
ATOM    350  N   GLU   498     -12.880  22.867 -12.574  1.00 25.28      A    N
ATOM    351  CA  GLU   498     -14.313  23.099 -12.473  1.00 25.14      A    C
ATOM    352  CB  GLU   498     -14.840  22.589 -11.122  1.00 25.47      A    C
ATOM    353  CG  GLU   498     -14.497  23.467  -9.924  1.00 24.08      A    C
ATOM    354  CD  GLU   498     -13.039  23.373  -9.504  1.00 27.78      A    C
ATOM    355  OE1 GLU   498     -12.453  22.265  -9.526  1.00 27.03      A    O
ATOM    356  OE2 GLU   498     -12.477  24.419  -9.139  1.00 30.19      A    O
ATOM    357  C   GLU   498     -14.669  24.567 -12.640  1.00 26.49      A    C
ATOM    358  O   GLU   498     -13.817  25.445 -12.521  1.00 26.93      A    O
ATOM    359  N   ARG   499     -15.943  24.823 -12.911  1.00 27.54      A    N
ATOM    360  CA  ARG   499     -16.427  26.180 -13.122  1.00 27.29      A    C
ATOM    361  CB  ARG   499     -16.477  26.493 -14.615  1.00 27.26      A    C
ATOM    362  CG  ARG   499     -16.999  27.879 -14.936  1.00 31.36      A    C
ATOM    363  CD  ARG   499     -17.028  28.135 -16.438  1.00 29.97      A    C
ATOM    364  NE  ARG   499     -17.154  29.562 -16.717  1.00 33.21      A    N
ATOM    365  CZ  ARG   499     -16.543  30.183 -17.721  1.00 31.95      A    C
ATOM    366  NH1 ARG   499     -16.710  31.487 -17.898  1.00 33.98      A    N
ATOM    367  NH2 ARG   499     -15.767  29.498 -18.549  1.00 30.18      A    N
ATOM    368  C   ARG   499     -17.818  26.349 -12.526  1.00 28.46      A    C
ATOM    369  O   ARG   499     -18.652  25.445 -12.608  1.00 27.86      A    O
ATOM    370  N   MET   500     -18.062  27.507 -11.921  1.00 28.00      A    N
ATOM    371  CA  MET   500     -19.405  27.868 -11.492  1.00 28.68      A    C
ATOM    372  CB  MET   500     -19.339  28.774 -10.259  1.00 29.36      A    C
ATOM    373  CG  MET   500     -18.800  28.062  -9.033  1.00 31.43      A    C
ATOM    374  SD  MET   500     -18.596  29.123  -7.594  1.00 35.65      A    S
ATOM    375  CE  MET   500     -17.061  29.997  -8.031  1.00 30.13      A    C
ATOM    376  C   MET   500     -20.118  28.578 -12.633  1.00 28.47      A    C
ATOM    377  O   MET   500     -19.589  29.527 -13.213  1.00 27.47      A    O
ATOM    378  N   GLU   501     -21.314  28.106 -12.965  1.00 29.85      A    N
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ATOM    379  CA  GLU   501     -22.059  28.656 -14.093  1.00 32.67      A    C
ATOM    380  CB  GLU   501     -22.169  27.617 -15.222  1.00 30.99      A    C
ATOM    381  CG  GLU   501     -20.816  27.155 -15.771  1.00 32.75      A    C
ATOM    382  CD  GLU   501     -20.922  26.113 -16.886  1.00 32.72      A    C
ATOM    383  OE1 GLU   501     -21.997  25.503 -17.053  1.00 28.58      A    O
ATOM    384  OE2 GLU   501     -19.917  25.907 -17.603  1.00 36.53      A    O
ATOM    385  C   GLU   501     -23.446  29.079 -13.633  1.00 35.03      A    C
ATOM    386  O   GLU   501     -24.072  28.399 -12.823  1.00 33.00      A    O
ATOM    387  N   ALA   502     -23.919  30.210 -14.145  1.00 39.39      A    N
ATOM    388  CA  ALA   502     -25.231  30.722 -13.766  1.00 42.89      A    C
ATOM    389  CB  ALA   502     -25.258  32.240 -13.896  1.00 41.99      A    C
ATOM    390  C   ALA   502     -26.305  30.104 -14.647  1.00 46.27      A    C
ATOM    391  O   ALA   502     -26.267  30.243 -15.868  1.00 47.27      A    O
ATOM    392  N   GLN   503     -27.251  29.414 -14.015  1.00 49.35      A    N
ATOM    393  CA  GLN   503     -28.385  28.808 -14.707  1.00 53.23      A    C
ATOM    394  CB  GLN   503     -28.315  27.281 -14.636  1.00 53.57      A    C
ATOM    395  CG  GLN   503     -27.190  26.633 -15.425  1.00 56.31      A    C
ATOM    396  CD  GLN   503     -27.325  25.113 -15.471  1.00 56.99      A    C
ATOM    397  OE1 GLN   503     -28.338  24.555 -15.041  1.00 57.97      A    O
ATOM    398  NE2 GLN   503     -26.305  24.441 -15.991  1.00 56.50      A    N
ATOM    399  C   GLN   503     -29.683  29.256 -14.048  1.00 55.13      A    C
ATOM    400  O   GLN   503     -29.988  28.845 -12.927  1.00 56.06      A    O
ATOM    401  N   GLY   504     -30.449  30.091 -14.744  1.00 56.99      A    N
ATOM    402  CA  GLY   504     -31.757  30.473 -14.244  1.00 56.78      A    C
ATOM    403  C   GLY   504     -31.704  31.096 -12.863  1.00 56.53      A    C
ATOM    404  O   GLY   504     -32.463  30.711 -11.973  1.00 57.57      A    O
ATOM    405  N   GLY   505     -30.799  32.055 -12.680  1.00 55.20      A    N
ATOM    406  CA  GLY   505     -30.714  32.760 -11.416  1.00 53.52      A    C
ATOM    407  C   GLY   505     -29.851  32.064 -10.381  1.00 53.23      A    C
ATOM    408  O   GLY   505     -29.381  32.699  -9.437  1.00 53.62      A    O
ATOM    409  N   LYS   506     -29.638  30.762 -10.548  1.00 50.84      A    N
ATOM    410  CA  LYS   506     -28.827  29.998  -9.604  1.00 49.53      A    C
ATOM    411  CB  LYS   506     -29.567  28.722  -9.183  1.00 51.74      A    C
ATOM    412  CG  LYS   506     -30.536  28.905  -8.019  1.00 54.83      A    C
ATOM    413  CD  LYS   506     -31.603  29.954  -8.324  1.00 59.19      A    C
ATOM    414  CE  LYS   506     -32.608  30.090  -7.177  1.00 59.58      A    C
ATOM    415  NZ  LYS   506     -33.329  28.812  -6.895  1.00 58.69      A    N
ATOM    416  C   LYS   506     -27.464  29.627 -10.190  1.00 46.30      A    C
ATOM    417  O   LYS   506     -27.327  29.456 -11.397  1.00 46.29      A    O
ATOM    418  N   LEU   507     -26.459  29.504  -9.329  1.00 41.66      A    N
ATOM    419  CA  LEU   507     -25.161  28.990  -9.749  1.00 36.60      A    C
ATOM    420  CB  LEU   507     -24.042  29.594  -8.901  1.00 34.82      A    C
ATOM    421  CG  LEU   507     -23.892  31.114  -8.941  1.00 35.36      A    C
ATOM    422  CD1 LEU   507     -22.759  31.532  -8.016  1.00 35.13      A    C
ATOM    423  CD2 LEU   507     -23.625  31.576 -10.363  1.00 30.11      A    C
ATOM    424  C   LEU   507     -25.127  27.473  -9.623  1.00 34.72      A    C
ATOM    425  O   LEU   507     -25.691  26.910  -8.657  1.00 33.92      A    O
ATOM    426  N   VAL   508     -24.524  26.812 -10.607  1.00 31.44      A    N
ATOM    427  CA  VAL   508     -24.238  25.392 -10.492  1.00 29.22      A    C
ATOM    428  CB  VAL   508     -25.036  24.556 -11.505  1.00 31.89      A    C
ATOM    429  CG1 VAL   508     -26.513  24.933 -11.437  1.00 30.81      A    C
ATOM    430  CG2 VAL   508     -24.469  24.744 -12.895  1.00 29.53      A    C
ATOM    431  C   VAL   508     -22.761  25.132 -10.724  1.00 28.89      A    C
ATOM    432  O   VAL   508     -22.026  26.009 -11.192  1.00 26.64      A    O
ATOM    433  N   CYS   509     -22.344  23.913 -10.401  1.00 26.50      A    N
ATOM    434  CA  CYS   509     -20.943  23.519 -10.431  1.00 26.63      A    C
ATOM    435  C   CYS   509     -20.710  22.485 -11.523  1.00 26.83      A    C
ATOM    436  O   CYS   509     -21.233  21.372 -11.446  1.00 27.38      A    O
ATOM    437  CB  CYS   509     -20.572  22.924  -9.076  1.00 27.35      A    C
ATOM    438  SG  CYS   509     -18.841  22.425  -8.815  1.00 29.68      A    S
ATOM    439  N   ARG   510     -19.921  22.847 -12.530  1.00 26.66      A    N
ATOM    440  CA  ARG   510     -19.559  21.915 -13.600  1.00 25.83      A    C
ATOM    441  CB  ARG   510     -19.838  22.546 -14.973  1.00 24.38      A    C
ATOM    442  CG  ARG   510     -19.589  21.595 -16.148  1.00 26.06      A    C
ATOM    443  CD  ARG   510     -19.798  22.278 -17.504  1.00 24.80      A    C
ATOM    444  NE  ARG   510     -21.158  22.782 -17.661  1.00 25.36      A    N
ATOM    445  CZ  ARG   510     -22.193  22.037 -18.049  1.00 28.22      A    C
ATOM    446  NH1 ARG   510     -23.397  22.581 -18.166  1.00 26.65      A    N
ATOM    447  NH2 ARG   510     -22.028  20.747 -18.320  1.00 24.05      A    N
ATOM    448  C   ARG   510     -18.084  21.521 -13.518  1.00 25.64      A    C
ATOM    449  O   ARG   510     -17.207  22.389 -13.530  1.00 27.40      A    O
ATOM    450  N   ALA   511     -17.812  20.220 -13.439  1.00 24.50      A    N
ATOM    451  CA  ALA   511     -16.436  19.712 -13.462  1.00 23.75      A    C
ATOM    452  CB  ALA   511     -16.218  18.716 -12.321  1.00 23.65      A    C
ATOM    453  C   ALA   511     -16.131  19.033 -14.794  1.00 24.97      A    C
ATOM    454  O   ALA   511     -17.015  18.413 -15.387  1.00 23.22      A    O
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ATOM    455  N   HIS   512     -14.872  19.135 -15.233  1.00 24.16      A    N
ATOM    456  CA  HIS   512     -14.435  18.679 -16.559  1.00 24.87      A    C
ATOM    457  CB  HIS   512     -13.774  19.837 -17.323  1.00 25.25      A    C
ATOM    458  CG  HIS   512     -14.711  20.953 -17.660  1.00 25.82      A    C
ATOM    459  CD2 HIS   512     -14.982  22.115 -17.021  1.00 26.89      A    C
ATOM    460  ND1 HIS   512     -15.507  20.939 -18.785  1.00 27.87      A    N
ATOM    461  CE1 HIS   512     -16.229  22.046 -18.825  1.00 28.93      A    C
ATOM    462  NE2 HIS   512     -15.929  22.776 -17.766  1.00 30.14      A    N
ATOM    463  C   HIS   512     -13.440  17.518 -16.503  1.00 25.39      A    C
ATOM    464  O   HIS   512     -12.452  17.570 -15.762  1.00 24.51      A    O
ATOM    465  N   ASN   513     -13.688  16.491 -17.313  1.00 23.79      A    N
ATOM    466  CA  ASN   513     -12.757  15.378 -17.459  1.00 24.72      A    C
ATOM    467  CB  ASN   513     -13.447  14.196 -18.155  1.00 21.47      A    C
ATOM    468  CG  ASN   513     -12.676  12.888 -17.998  1.00 25.11      A    C
ATOM    469  OD1 ASN   513     -11.857  12.740 -17.087  1.00 26.09      A    O
ATOM    470  ND2 ASN   513     -12.943  11.927 -18.887  1.00 20.32      A    N
ATOM    471  C   ASN   513     -11.558  15.839 -18.285  1.00 23.06      A    C
ATOM    472  O   ASN   513     -11.572  16.930 -18.843  1.00 23.76      A    O
ATOM    473  N   ALA   514     -10.523  15.010 -18.359  1.00 23.46      A    N
ATOM    474  CA  ALA   514      -9.342  15.331 -19.170  1.00 24.44      A    C
ATOM    475  CB  ALA   514      -8.116  15.505 -18.268  1.00 21.96      A    C
ATOM    476  C   ALA   514      -9.066  14.247 -20.203  1.00 24.29      A    C
ATOM    477  O   ALA   514      -9.501  13.107 -20.041  1.00 24.54      A    O
ATOM    478  N   PHE   515      -8.340  14.608 -21.261  1.00 25.86      A    N
ATOM    479  CA  PHE   515      -7.806  13.632 -22.218  1.00 25.71      A    C
ATOM    480  CB  PHE   515      -6.761  14.313 -23.116  1.00 27.88      A    C
ATOM    481  CG  PHE   515      -6.124  13.397 -24.134  1.00 28.53      A    C
ATOM    482  CD1 PHE   515      -6.818  13.006 -25.270  1.00 29.33      A    C
ATOM    483  CD2 PHE   515      -4.827  12.936 -23.956  1.00 31.17      A    C
ATOM    484  CE1 PHE   515      -6.234  12.171 -26.216  1.00 30.11      A    C
ATOM    485  CE2 PHE   515      -4.231  12.098 -24.897  1.00 32.87      A    C
ATOM    486  CZ  PHE   515      -4.936  11.714 -26.030  1.00 30.57      A    C
ATOM    487  C   PHE   515      -7.170  12.473 -21.446  1.00 25.67      A    C
ATOM    488  O   PHE   515      -6.333  12.692 -20.570  1.00 25.88      A    O
ATOM    489  N   GLY   516      -7.589  11.247 -21.745  1.00 25.26      A    N
ATOM    490  CA  GLY   516      -6.993  10.089 -21.095  1.00 25.89      A    C
ATOM    491  C   GLY   516      -7.556   9.730 -19.725  1.00 28.02      A    C
ATOM    492  O   GLY   516      -7.309   8.634 -19.228  1.00 29.41      A    O
ATOM    493  N   GLY   517      -8.310  10.641 -19.112  1.00 27.77      A    N
ATOM    494  CA  GLY   517      -8.722  10.457 -17.727  1.00 28.63      A    C
ATOM    495  C   GLY   517      -9.944   9.571 -17.617  1.00 29.61      A    C
ATOM    496  O   GLY   517     -10.494   9.178 -18.629  1.00 29.48      A    O
ATOM    497  N   GLU   518     -10.385   9.246 -16.408  1.00 28.88      A    N
ATOM    498  CA  GLU   518     -11.479   8.292 -16.278  1.00 30.08      A    C
ATOM    499  CB  GLU   518     -11.096   7.152 -15.325  1.00 33.06      A    C
ATOM    500  CG  GLU   518     -10.923   7.562 -13.883  1.00 41.55      A    C
ATOM    501  CD  GLU   518      -9.561   8.160 -13.614  1.00 48.72      A    C
ATOM    502  OE1 GLU   518      -8.719   8.183 -14.546  1.00 50.19      A    O
ATOM    503  OE2 GLU   518      -9.332   8.607 -12.467  1.00 53.68      A    O
ATOM    504  C   GLU   518     -12.779   8.931 -15.818  1.00 28.72      A    C
ATOM    505  O   GLU   518     -13.720   8.233 -15.443  1.00 29.04      A    O
ATOM    506  N   GLY   519     -12.836  10.257 -15.862  1.00 26.44      A    N
ATOM    507  CA  GLY   519     -14.033  10.951 -15.430  1.00 25.94      A    C
ATOM    508  C   GLY   519     -13.808  11.639 -14.099  1.00 25.37      A    C
ATOM    509  O   GLY   519     -12.808  11.386 -13.429  1.00 24.07      A    O
ATOM    510  N   VAL   520     -14.738  12.507 -13.715  1.00 24.24      A    N
ATOM    511  CA  VAL   520     -14.545  13.374 -12.557  1.00 24.20      A    C
ATOM    512  CB  VAL   520     -14.047  14.770 -12.988  1.00 25.35      A    C
ATOM    513  CG1 VAL   520     -12.750  14.643 -13.781  1.00 23.64      A    C
ATOM    514  CG2 VAL   520     -15.114  15.454 -13.824  1.00 21.19      A    C
ATOM    515  C   VAL   520     -15.852  13.561 -11.800  1.00 24.80      A    C
ATOM    516  O   VAL   520     -16.920  13.207 -12.302  1.00 25.48      A    O
ATOM    517  N   TYR   521     -15.754  14.121 -10.596  1.00 23.24      A    N
ATOM    518  CA  TYR   521     -16.922  14.530  -9.821  1.00 23.59      A    C
ATOM    519  CB  TYR   521     -16.902  13.893  -8.430  1.00 24.30      A    C
ATOM    520  CG  TYR   521     -16.985  12.392  -8.457  1.00 26.30      A    C
ATOM    521  CD1 TYR   521     -15.833  11.620  -8.533  1.00 25.03      A    C
ATOM    522  CE1 TYR   521     -15.901  10.243  -8.597  1.00 29.75      A    C
ATOM    523  CD2 TYR   521     -18.216  11.746  -8.440  1.00 23.79      A    C
ATOM    524  CE2 TYR   521     -18.296  10.368  -8.505  1.00 28.16      A    C
ATOM    525  CZ  TYR   521     -17.133   9.624  -8.587  1.00 30.47      A    C
ATOM    526  OH  TYR   521     -17.196   8.259  -8.694  1.00 35.15      A    O
ATOM    527  C   TYR   521     -16.945  16.037  -9.650  1.00 23.08      A    C
ATOM    528  O   TYR   521     -15.898  16.666  -9.504  1.00 22.54      A    O
ATOM    529  N   ALA   522     -18.142  16.611  -9.660  1.00 23.04      A    N
ATOM    530  CA  ALA   522     -18.314  18.005  -9.283  1.00 22.28      A    C
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ATOM    531  CB  ALA   522     -19.336  18.685 -10.201  1.00 23.30      A    C
ATOM    532  C   ALA   522     -18.803  18.016  -7.843  1.00 23.52      A    C
ATOM    533  O   ALA   522     -19.561  17.132  -7.431  1.00 22.23      A    O
ATOM    534  N   ILE   523     -18.358  19.004  -7.074  1.00 23.74      A    N
ATOM    535  CA  ILE   523     -18.714  19.078  -5.665  1.00 23.56      A    C
ATOM    536  CB  ILE   523     -17.537  18.639  -4.779  1.00 26.23      A    C
ATOM    537  CG2 ILE   523     -17.959  18.618  -3.306  1.00 21.95      A    C
ATOM    538  CG1 ILE   523     -17.083  17.237  -5.193  1.00 24.93      A    C
ATOM    539  CD1 ILE   523     -15.614  17.139  -5.452  1.00 24.24      A    C
ATOM    540  C   ILE   523     -19.112  20.500  -5.303  1.00 26.24      A    C
ATOM    541  O   ILE   523     -18.282  21.408  -5.298  1.00 28.35      A    O
ATOM    542  N   ALA   524     -20.393  20.690  -5.013  1.00 24.72      A    N
ATOM    543  CA  ALA   524     -20.902  22.004  -4.658  1.00 24.50      A    C
ATOM    544  CB  ALA   524     -22.279  22.214  -5.295  1.00 23.50      A    C
ATOM    545  C   ALA   524     -20.999  22.141  -3.140  1.00 24.77      A    C
ATOM    546  O   ALA   524     -21.254  21.170  -2.435  1.00 26.27      A    O
ATOM    547  N   ARG   525     -20.784  23.352  -2.646  1.00 26.08      A    N
ATOM    548  CA  ARG   525     -21.076  23.684  -1.259  1.00 25.79      A    C
ATOM    549  CB  ARG   525     -19.931  24.513  -0.678  1.00 26.10      A    C
ATOM    550  CG  ARG   525     -19.969  24.682   0.825  1.00 27.35      A    C
ATOM    551  CD  ARG   525     -19.724  23.367   1.575  1.00 26.86      A    C
ATOM    552  NE  ARG   525     -19.572  23.634   3.001  1.00 26.08      A    N
ATOM    553  CZ  ARG   525     -19.631  22.720   3.963  1.00 28.22      A    C
ATOM    554  NH1 ARG   525     -19.480  23.097   5.225  1.00 24.72      A    N
ATOM    555  NH2 ARG   525     -19.838  21.436   3.671  1.00 28.59      A    N
ATOM    556  C   ARG   525     -22.374  24.492  -1.259  1.00 26.55      A    C
ATOM    557  O   ARG   525     -22.457  25.555  -1.877  1.00 27.38      A    O
ATOM    558  N   CYS   526     -23.393  23.973  -0.585  1.00 27.37      A    N
ATOM    559  CA  CYS   526     -24.727  24.557  -0.635  1.00 27.20      A    C
ATOM    560  C   CYS   526     -25.155  25.053   0.749  1.00 28.67      A    C
ATOM    561  O   CYS   526     -25.183  24.282   1.709  1.00 27.15      A    O
ATOM    562  CB  CYS   526     -25.722  23.512  -1.124  1.00 28.62      A    C
ATOM    563  SG  CYS   526     -25.404  22.884  -2.801  1.00 28.40      A    S
ATOM    564  N   CYS   527     -25.496  26.333   0.850  1.00 27.75      A    N
ATOM    565  CA  CYS   527     -25.764  26.923   2.153  1.00 30.68      A    C
ATOM    566  C   CYS   527     -27.111  27.637   2.230  1.00 30.03      A    C
ATOM    567  O   CYS   527     -27.672  28.049   1.213  1.00 29.33      A    O
ATOM    568  CB  CYS   527     -24.653  27.907   2.519  1.00 30.20      A    C
ATOM    569  SG  CYS   527     -22.945  27.315   2.258  1.00 34.88      A    S
ATOM    570  N   LEU   528     -27.626  27.777   3.448  1.00 31.45      A    N
ATOM    571  CA  LEU   528     -28.801  28.604   3.688  1.00 33.88      A    C
ATOM    572  CB  LEU   528     -29.615  28.058   4.862  1.00 31.33      A    C
ATOM    573  CG  LEU   528     -30.103  26.630   4.633  1.00 31.88      A    C
ATOM    574  CD1 LEU   528     -30.984  26.179   5.793  1.00 28.76      A    C
ATOM    575  CD2 LEU   528     -30.858  26.576   3.317  1.00 29.32      A    C
ATOM    576  C   LEU   528     -28.361  30.026   3.979  1.00 36.14      A    C
ATOM    577  O   LEU   528     -27.762  30.311   5.014  1.00 36.49      A    O
ATOM    578  N   LEU   529     -28.652  30.914   3.042  1.00 40.20      A    N
ATOM    579  CA  LEU   529     -28.208  32.292   3.135  1.00 45.03      A    C
ATOM    580  CB  LEU   529     -26.956  32.491   2.272  1.00 43.96      A    C
ATOM    581  CG  LEU   529     -26.141  33.767   2.499  1.00 45.14      A    C
ATOM    582  CD1 LEU   529     -25.404  33.667   3.822  1.00 45.34      A    C
ATOM    583  CD2 LEU   529     -25.146  33.956   1.368  1.00 45.47      A    C
ATOM    584  C   LEU   529     -29.348  33.181   2.638  1.00 48.10      A    C
ATOM    585  O   LEU   529     -29.527  33.369   1.431  1.00 48.79      A    O
ATOM    586  N   PRO   530     -30.144  33.726   3.570  1.00 48.91      A    N
ATOM    587  CD  PRO   530     -29.972  33.547   5.021  1.00 48.98      A    C
ATOM    588  CA  PRO   530     -31.254  34.638   3.259  1.00 50.42      A    C
ATOM    589  CB  PRO   530     -31.955  34.819   4.603  1.00 49.24      A    C
ATOM    590  CG  PRO   530     -30.872  34.610   5.604  1.00 50.12      A    C
ATOM    591  C   PRO   530     -30.773  35.972   2.687  1.00 49.69      A    C
ATOM    592  O   PRO   530     -29.729  36.488   3.080  1.00 48.57      A    O
ATOM    593  N   GLN   531     -31.544  36.527   1.759  1.00 52.50      A    N
ATOM    594  CA  GLN   531     -31.225  37.830   1.182  1.00 54.58      A    C
ATOM    595  CB  GLN   531     -31.325  38.929   2.247  1.00 57.89      A    C
ATOM    596  CG  GLN   531     -32.722  39.160   2.811  1.00 62.33      A    C
ATOM    597  CD  GLN   531     -32.754  40.294   3.831  1.00 65.19      A    C
ATOM    598  OE1 GLN   531     -32.220  41.381   3.590  1.00 66.51      A    O
ATOM    599  NE2 GLN   531     -33.378  40.043   4.978  1.00 64.89      A    N
ATOM    600  C   GLN   531     -29.820  37.843   0.586  1.00 53.73      A    C
ATOM    601  O   GLN   531     -29.021  38.734   0.882  1.00 54.88      A    O
ATOM    602  N   ALA   532     -29.518  36.858  -0.250  1.00 50.22      A    N
ATOM    603  CA  ALA   532     -28.234  36.828  -0.933  1.00 48.20      A    C
ATOM    604  CB  ALA   532     -27.514  35.525  -0.626  1.00 46.39      A    C
ATOM    605  C   ALA   532     -28.404  36.991  -2.443  1.00 47.17      A    C
ATOM    606  O   ALA   532     -29.324  36.432  -3.043  1.00 45.39      A    O
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ATOM    607  N   ASN   533     -27.520  37.772  -3.052  1.00 46.61      A    N
ATOM    608  CA  ASN   533     -27.359  37.735  -4.500  1.00 45.24      A    C
ATOM    609  CB  ASN   533     -27.689  39.097  -5.130  1.00 48.68      A    C
ATOM    610  CG  ASN   533     -27.745  39.039  -6.657  1.00 52.76      A    C
ATOM    611  OD1 ASN   533     -27.459  38.001  -7.259  1.00 55.43      A    O
ATOM    612  ND2 ASN   533     -28.117  40.154  -7.286  1.00 54.40      A    N
ATOM    613  C   ASN   533     -25.910  37.372  -4.781  1.00 42.36      A    C
ATOM    614  O   ASN   533     -25.004  38.166  -4.540  1.00 41.25      A    O
ATOM    615  N   CYS   534     -25.694  36.162  -5.278  1.00 40.03      A    N
ATOM    616  CA  CYS   534     -24.346  35.706  -5.572  1.00 38.85      A    C
ATOM    617  C   CYS   534     -24.139  35.637  -7.076  1.00 38.60      A    C
ATOM    618  O   CYS   534     -25.031  35.215  -7.812  1.00 38.28      A    O
ATOM    619  CB  CYS   534     -24.104  34.322  -4.980  1.00 38.71      A    C
ATOM    620  SG  CYS   534     -24.229  34.143  -3.170  1.00 39.44      A    S
ATOM    621  N   SER   535     -22.956  36.040  -7.525  1.00 35.57      A    N
ATOM    622  CA  SER   535     -22.642  36.029  -8.943  1.00 35.87      A    C
ATOM    623  CB  SER   535     -22.807  37.438  -9.523  1.00 34.93      A    C
ATOM    624  OG  SER   535     -22.034  38.375  -8.795  1.00 37.31      A    O
ATOM    625  C   SER   535     -21.220  35.528  -9.168  1.00 34.87      A    C
ATOM    626  O   SER   535     -20.364  35.643  -8.287  1.00 34.76      A    O
ATOM    627  N   VAL   536     -20.975  34.966 -10.347  1.00 32.81      A    N
ATOM    628  CA  VAL   536     -19.659  34.444 -10.681  1.00 32.05      A    C
ATOM    629  CB  VAL   536     -19.752  33.158 -11.524  1.00 33.02      A    C
ATOM    630  CG1 VAL   536     -18.353  32.630 -11.817  1.00 31.62      A    C
ATOM    631  CG2 VAL   536     -20.555  32.116 -10.787  1.00 37.22      A    C
ATOM    632  C   VAL   536     -18.871  35.465 -11.479  1.00 32.35      A    C
ATOM    633  O   VAL   536     -19.395  36.083 -12.406  1.00 32.68      A    O
ATOM    634  N   HIS   537     -17.603  35.631 -11.129  1.00 32.15      A    N
ATOM    635  CA  HIS   537     -16.743  36.550 -11.854  1.00 32.62      A    C
ATOM    636  CB  HIS   537     -16.430  37.757 -10.974  1.00 32.90      A    C
ATOM    637  CG  HIS   537     -17.657  38.492 -10.534  1.00 36.48      A    C
ATOM    638  CD2 HIS   537     -18.536  38.238  -9.534  1.00 39.02      A    C
ATOM    639  ND1 HIS   537     -18.153  39.587 -11.211  1.00 39.06      A    N
ATOM    640  CE1 HIS   537     -19.285  39.973 -10.650  1.00 40.38      A    C
ATOM    641  NE2 HIS   537     -19.540  39.171  -9.630  1.00 41.12      A    N
ATOM    642  C   HIS   537     -15.482  35.829 -12.268  1.00 31.22      A    C
ATOM    643  O   HIS   537     -14.825  35.197 -11.446  1.00 32.87      A    O
ATOM    644  N   THR   538     -15.160  35.923 -13.553  1.00 30.02      A    N
ATOM    645  CA  THR   538     -14.130  35.092 -14.162  1.00 28.85      A    C
ATOM    646  CB  THR   538     -14.745  34.189 -15.259  1.00 30.20      A    C
ATOM    647  OG1 THR   538     -15.658  33.264 -14.652  1.00 29.62      A    O
ATOM    648  CG2 THR   538     -13.658  33.417 -16.011  1.00 26.80      A    C
ATOM    649  C   THR   538     -13.027  35.934 -14.786  1.00 27.41      A    C
ATOM    650  O   THR   538     -13.290  37.001 -15.330  1.00 26.94      A    O
ATOM    651  N   ALA   539     -11.793  35.453 -14.704  1.00 26.67      A    N
ATOM    652  CA  ALA   539     -10.713  36.017 -15.502  1.00 26.84      A    C
ATOM    653  CB  ALA   539      -9.722  36.757 -14.617  1.00 27.43      A    C
ATOM    654  C   ALA   539     -10.010  34.901 -16.258  1.00 27.34      A    C
ATOM    655  O   ALA   539      -9.800  33.806 -15.727  1.00 29.03      A    O
ATOM    656  N   PRO   540      -9.645  35.168 -17.518  1.00 25.34      A    N
ATOM    657  CD  PRO   540      -9.835  36.482 -18.162  1.00 26.40      A    C
ATOM    658  CA  PRO   540      -9.058  34.190 -18.434  1.00 25.56      A    C
ATOM    659  CB  PRO   540      -9.227  34.841 -19.804  1.00 25.71      A    C
ATOM    660  CG  PRO   540      -9.165  36.312 -19.505  1.00 27.40      A    C
ATOM    661  C   PRO   540      -7.595  33.907 -18.126  1.00 27.52      A    C
ATOM    662  O   PRO   540      -6.923  34.699 -17.463  1.00 27.04      A    O
ATOM    663  N   PRO   541      -7.075  32.783 -18.636  1.00 27.53      A    N
ATOM    664  CD  PRO   541      -7.795  31.778 -19.435  1.00 28.31      A    C
ATOM    665  CA  PRO   541      -5.650  32.469 -18.522  1.00 29.94      A    C
ATOM    666  CB  PRO   541      -5.458  31.318 -19.510  1.00 26.51      A    C
ATOM    667  CG  PRO   541      -6.794  30.658 -19.562  1.00 28.03      A    C
ATOM    668  C   PRO   541      -4.788  33.676 -18.875  1.00 32.23      A    C
ATOM    669  O   PRO   541      -5.053  34.382 -19.848  1.00 30.35      A    O
ATOM    670  N   ALA   542      -3.759  33.910 -18.070  1.00 35.43      A    N
ATOM    671  CA  ALA   542      -2.864  35.034 -18.290  1.00 39.68      A    C
ATOM    672  CB  ALA   542      -3.169  36.142 -17.298  1.00 41.32      A    C
ATOM    673  C   ALA   542      -1.429  34.572 -18.125  1.00 41.82      A    C
ATOM    674  O   ALA   542      -1.077  33.982 -17.108  1.00 43.83      A    O
ATOM    675  N   GLU   543      -0.603  34.838 -19.125  1.00 44.83      A    N
ATOM    676  CA  GLU   543       0.816  34.523 -19.033  1.00 48.00      A    C
ATOM    677  CB  GLU   543       1.461  34.625 -20.418  1.00 52.56      A    C
ATOM    678  CG  GLU   543       2.492  33.547 -20.715  1.00 57.96      A    C
ATOM    679  CD  GLU   543       1.858  32.227 -21.120  1.00 61.23      A    C
ATOM    680  OE1 GLU   543       1.824  31.303 -20.277  1.00 62.99      A    O
ATOM    681  OE2 GLU   543       1.396  32.113 -22.283  1.00 63.66      A    O
ATOM    682  C   GLU   543       1.463  35.530 -18.082  1.00 47.94      A    C
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ATOM    683  O   GLU   543       2.038  36.525 -18.520  1.00 49.36      A    O
ATOM    684  N   ALA   544       1.359  35.274 -16.781  1.00 45.85      A    N
ATOM    685  CA  ALA   544       1.827  36.223 -15.780  1.00 43.79      A    C
ATOM    686  CB  ALA   544       0.678  37.126 -15.346  1.00 42.96      A    C
ATOM    687  C   ALA   544       2.418  35.508 -14.564  1.00 44.20      A    C
ATOM    688  O   ALA   544       1.998  34.402 -14.210  1.00 42.82      A    O
ATOM    689  N   SER   545       3.392  36.148 -13.926  1.00 43.11      A    N
ATOM    690  CA  SER   545       4.054  35.566 -12.767  1.00 42.80      A    C
ATOM    691  CB  SER   545       5.198  36.469 -12.297  1.00 44.06      A    C
ATOM    692  OG  SER   545       6.381  36.204 -13.028  1.00 49.18      A    O
ATOM    693  C   SER   545       3.082  35.350 -11.621  1.00 40.42      A    C
ATOM    694  O   SER   545       3.159  34.344 -10.917  1.00 41.59      A    O
ATOM    695  N   MET   546       2.171  36.297 -11.435  1.00 37.26      A    N
ATOM    696  CA  MET   546       1.224  36.224 -10.337  1.00 37.49      A    C
ATOM    697  CB  MET   546       0.851  37.631  -9.869  1.00 36.04      A    C
ATOM    698  CG  MET   546       1.973  38.326  -9.119  1.00 37.71      A    C
ATOM    699  SD  MET   546       1.626  40.049  -8.851  1.00 38.80      A    S
ATOM    700  CE  MET   546       2.137  40.710 -10.406  1.00 47.70      A    C
ATOM    701  C   MET   546      -0.028  35.451 -10.718  1.00 37.16      A    C
ATOM    702  O   MET   546      -1.076  35.596 -10.083  1.00 37.19      A    O
ATOM    703  N   GLY   547       0.091  34.630 -11.757  1.00 36.79      A    N
ATOM    704  CA  GLY   547      -0.994  33.747 -12.136  1.00 35.87      A    C
ATOM    705  C   GLY   547      -2.212  34.481 -12.659  1.00 36.73      A    C
ATOM    706  O   GLY   547      -2.119  35.599 -13.168  1.00 36.47      A    O
ATOM    707  N   THR   548      -3.366  33.840 -12.536  1.00 35.43      A    N
ATOM    708  CA  THR   548      -4.617  34.407 -13.017  1.00 35.74      A    C
ATOM    709  CB  THR   548      -5.403  33.365 -13.841  1.00 37.13      A    C
ATOM    710  OG1 THR   548      -4.580  32.892 -14.914  1.00 39.98      A    O
ATOM    711  CG2 THR   548      -6.664  33.976 -14.416  1.00 38.25      A    C
ATOM    712  C   THR   548      -5.442  34.817 -11.808  1.00 34.88      A    C
ATOM    713  O   THR   548      -5.597  34.041 -10.866  1.00 36.02      A    O
ATOM    714  N   ARG   549      -5.971  36.033 -11.831  1.00 33.45      A    N
ATOM    715  CA  ARG   549      -6.562  36.607 -10.632  1.00 34.68      A    C
ATOM    716  CB  ARG   549      -5.574  37.580  -9.975  1.00 34.64      A    C
ATOM    717  CG  ARG   549      -4.158  37.005  -9.856  1.00 37.46      A    C
ATOM    718  CD  ARG   549      -3.155  38.035  -9.350  1.00 35.98      A    C
ATOM    719  NE  ARG   549      -3.201  38.159  -7.901  1.00 35.43      A    N
ATOM    720  CZ  ARG   549      -2.454  37.440  -7.069  1.00 34.21      A    C
ATOM    721  NH1 ARG   549      -2.566  37.618  -5.760  1.00 30.15      A    N
ATOM    722  NH2 ARG   549      -1.593  36.545  -7.546  1.00 32.58      A    N
ATOM    723  C   ARG   549      -7.859  37.329 -10.947  1.00 34.29      A    C
ATOM    724  O   ARG   549      -7.990  37.968 -11.986  1.00 34.98      A    O
ATOM    725  N   VAL   550      -8.818  37.221 -10.040  1.00 33.25      A    N
ATOM    726  CA  VAL   550     -10.054  37.970 -10.147  1.00 33.71      A    C
ATOM    727  CB  VAL   550     -11.130  37.161 -10.914  1.00 33.51      A    C
ATOM    728  CG1 VAL   550     -11.547  35.936 -10.103  1.00 30.56      A    C
ATOM    729  CG2 VAL   550     -12.329  38.043 -11.222  1.00 32.51      A    C
ATOM    730  C   VAL   550     -10.530  38.244  -8.727  1.00 36.76      A    C
ATOM    731  O   VAL   550     -10.261  37.460  -7.812  1.00 36.18      A    O
ATOM    732  N   HIS   551     -11.222  39.358  -8.530  1.00 38.93      A    N
ATOM    733  CA  HIS   551     -11.719  39.683  -7.200  1.00 43.33      A    C
ATOM    734  CB  HIS   551     -10.779  40.675  -6.506  1.00 46.64      A    C
ATOM    735  CG  HIS   551     -10.856  42.067  -7.050  1.00 52.46      A    C
ATOM    736  CD2 HIS   551     -10.404  42.597  -8.212  1.00 54.58      A    C
ATOM    737  ND1 HIS   551     -11.460  43.101  -6.365  1.00 54.96      A    N
ATOM    738  CE1 HIS   551     -11.377  44.208  -7.082  1.00 56.62      A    C
ATOM    739  NE2 HIS   551     -10.741  43.930  -8.207  1.00 57.00      A    N
ATOM    740  C   HIS   551     -13.128  40.256  -7.262  1.00 43.49      A    C
ATOM    741  O   HIS   551     -13.585  40.702  -8.313  1.00 42.88      A    O
ATOM    742  N   CYS   552     -13.815  40.229  -6.128  1.00 44.34      A    N
ATOM    743  CA  CYS   552     -15.158  40.771  -6.045  1.00 47.65      A    C
ATOM    744  C   CYS   552     -15.074  42.292  -5.943  1.00 51.08      A    C
ATOM    745  O   CYS   552     -14.612  42.830  -4.937  1.00 50.38      A    O
ATOM    746  CB  CYS   552     -15.875  40.193  -4.824  1.00 44.92      A    C
ATOM    747  SG  CYS   552     -15.982  38.371  -4.816  1.00 44.45      A    S
ATOM    748  N   HIS   553     -15.518  42.979  -6.990  1.00 55.41      A    N
ATOM    749  CA  HIS   553     -15.266  44.410  -7.120  1.00 60.41      A    C
ATOM    750  CB  HIS   553     -14.882  44.748  -8.565  1.00 63.13      A    C
ATOM    751  CG  HIS   553     -15.927  44.379  -9.572  1.00 68.52      A    C
ATOM    752  CD2 HIS   553     -16.781  45.144 -10.294  1.00 70.80      A    C
ATOM    753  ND1 HIS   553     -16.187  43.073  -9.932  1.00 71.80      A    N
ATOM    754  CE1 HIS   553     -17.155  43.049 -10.831  1.00 71.80      A    C
ATOM    755  NE2 HIS   553     -17.533  44.293 -11.068  1.00 72.09      A    N
ATOM    756  C   HIS   553     -16.442  45.282  -6.685  1.00 60.90      A    C
ATOM    757  O   HIS   553     -16.292  46.491  -6.524  1.00 62.29      A    O
ATOM    758  N   GLN   554     -17.609  44.677  -6.497  1.00 61.02      A    N
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ATOM    759  CA  GLN   554     -18.792  45.442  -6.120  1.00 61.69      A    C
ATOM    760  CB  GLN   554     -20.062  44.695  -6.527  1.00 62.40      A    C
ATOM    761  CG  GLN   554     -20.209  44.514  -8.030  1.00 64.68      A    C
ATOM    762  CD  GLN   554     -21.646  44.278  -8.452  1.00 66.14      A    C
ATOM    763  OE1 GLN   554     -22.313  43.371  -7.952  1.00 66.12      A    O
ATOM    764  NE2 GLN   554     -22.133  45.100  -9.379  1.00 66.69      A    N
ATOM    765  C   GLN   554     -18.831  45.747  -4.628  1.00 61.75      A    C
ATOM    766  O   GLN   554     -18.285  45.002  -3.813  1.00 61.46      A    O
ATOM    767  N   GLN   555     -19.488  46.850  -4.281  1.00 61.73      A    N
ATOM    768  CA  GLN   555     -19.517  47.338  -2.907  1.00 61.26      A    C
ATOM    769  CB  GLN   555     -20.092  48.756  -2.869  1.00 64.04      A    C
ATOM    770  CG  GLN   555     -19.903  49.452  -1.534  1.00 67.91      A    C
ATOM    771  CD  GLN   555     -18.440  49.524  -1.132  1.00 71.48      A    C
ATOM    772  OE1 GLN   555     -17.700  50.401  -1.586  1.00 73.49      A    O
ATOM    773  NE2 GLN   555     -18.012  48.595  -0.279  1.00 71.05      A    N
ATOM    774  C   GLN   555     -20.336  46.437  -1.984  1.00 59.47      A    C
ATOM    775  O   GLN   555     -21.532  46.235  -2.199  1.00 58.75      A    O
ATOM    776  N   GLY   556     -19.683  45.907  -0.953  1.00 57.33      A    N
ATOM    777  CA  GLY   556     -20.378  45.081   0.019  1.00 55.06      A    C
ATOM    778  C   GLY   556     -20.304  43.591  -0.268  1.00 53.72      A    C
ATOM    779  O   GLY   556     -20.687  42.770   0.566  1.00 53.32      A    O
ATOM    780  N   HIS   557     -19.809  43.233  -1.446  1.00 51.42      A    N
ATOM    781  CA  HIS   557     -19.772  41.834  -1.841  1.00 49.50      A    C
ATOM    782  CB  HIS   557     -19.655  41.721  -3.367  1.00 51.95      A    C
ATOM    783  CG  HIS   557     -20.930  42.035  -4.092  1.00 56.53      A    C
ATOM    784  CD2 HIS   557     -21.535  41.427  -5.141  1.00 56.76      A    C
ATOM    785  ND1 HIS   557     -21.752  43.085  -3.733  1.00 57.67      A    N
ATOM    786  CE1 HIS   557     -22.807  43.110  -4.528  1.00 55.39      A    C
ATOM    787  NE2 HIS   557     -22.700  42.115  -5.391  1.00 58.10      A    N
ATOM    788  C   HIS   557     -18.633  41.089  -1.151  1.00 45.78      A    C
ATOM    789  O   HIS   557     -17.560  41.647  -0.920  1.00 45.56      A    O
ATOM    790  N   VAL   558     -18.892  39.829  -0.812  1.00 41.36      A    N
ATOM    791  CA  VAL   558     -17.934  38.982  -0.108  1.00 37.02      A    C
ATOM    792  CB  VAL   558     -18.500  38.521   1.244  1.00 37.11      A    C
ATOM    793  CG1 VAL   558     -17.454  37.737   1.998  1.00 37.46      A    C
ATOM    794  CG2 VAL   558     -18.964  39.721   2.050  1.00 39.30      A    C
ATOM    795  C   VAL   558     -17.628  37.736  -0.930  1.00 35.05      A    C
ATOM    796  O   VAL   558     -18.538  37.112  -1.475  1.00 34.44      A    O
ATOM    797  N   LEU   559     -16.351  37.373  -1.009  1.00 33.22      A    N
ATOM    798  CA  LEU   559     -15.934  36.175  -1.732  1.00 31.06      A    C
ATOM    799  CB  LEU   559     -14.440  36.250  -2.039  1.00 29.73      A    C
ATOM    800  CG  LEU   559     -13.822  34.965  -2.596  1.00 29.66      A    C
ATOM    801  CD1 LEU   559     -14.328  34.731  -4.025  1.00 27.03      A    C
ATOM    802  CD2 LEU   559     -12.300  35.079  -2.575  1.00 28.88      A    C
ATOM    803  C   LEU   559     -16.220  34.929  -0.892  1.00 31.01      A    C
ATOM    804  O   LEU   559     -15.760  34.829   0.245  1.00 31.89      A    O
ATOM    805  N   THR   560     -16.976  33.984  -1.447  1.00 28.29      A    N
ATOM    806  CA  THR   560     -17.335  32.777  -0.707  1.00 27.77      A    C
ATOM    807  CB  THR   560     -18.862  32.532  -0.728  1.00 27.96      A    C
ATOM    808  OG1 THR   560     -19.302  32.333  -2.080  1.00 29.50      A    O
ATOM    809  CG2 THR   560     -19.602  33.721  -0.117  1.00 24.95      A    C
ATOM    810  C   THR   560     -16.649  31.518  -1.234  1.00 27.81      A    C
ATOM    811  O   THR   560     -16.592  30.505  -0.544  1.00 27.32      A    O
ATOM    812  N   GLY   561     -16.149  31.579  -2.465  1.00 28.51      A    N
ATOM    813  CA  GLY   561     -15.529  30.415  -3.067  1.00 26.65      A    C
ATOM    814  C   GLY   561     -14.728  30.746  -4.312  1.00 28.04      A    C
ATOM    815  O   GLY   561     -15.044  31.696  -5.035  1.00 29.85      A    O
ATOM    816  N   CYS   562     -13.684  29.960  -4.555  1.00 26.78      A    N
ATOM    817  CA  CYS   562     -12.860  30.109  -5.745  1.00 28.08      A    C
ATOM    818  C   CYS   562     -12.868  28.802  -6.535  1.00 28.91      A    C
ATOM    819  O   CYS   562     -12.698  27.725  -5.965  1.00 29.35      A    O
ATOM    820  CB  CYS   562     -11.414  30.444  -5.358  1.00 29.31      A    C
ATOM    821  SG  CYS   562     -11.177  32.043  -4.523  1.00 30.67      A    S
ATOM    822  N   SER   563     -13.060  28.903  -7.845  1.00 27.63      A    N
ATOM    823  CA  SER   563     -12.994  27.741  -8.721  1.00 27.76      A    C
ATOM    824  CB  SER   563     -14.372  27.442  -9.321  1.00 25.87      A    C
ATOM    825  OG  SER   563     -15.267  26.932  -8.341  1.00 29.15      A    O
ATOM    826  C   SER   563     -11.991  27.990  -9.839  1.00 27.56      A    C
ATOM    827  O   SER   563     -11.673  29.137 -10.157  1.00 28.34      A    O
ATOM    828  N   SER   564     -11.506  26.912 -10.440  1.00 26.93      A    N
ATOM    829  CA  SER   564     -10.543  27.014 -11.531  1.00 28.10      A    C
ATOM    830  CB  SER   564      -9.121  27.017 -10.960  1.00 28.82      A    C
ATOM    831  OG  SER   564      -8.171  27.330 -11.957  1.00 33.70      A    O
ATOM    832  C   SER   564     -10.697  25.845 -12.502  1.00 26.61      A    C
ATOM    833  O   SER   564     -10.767  24.693 -12.083  1.00 27.41      A    O
ATOM    834  N   HIS   565     -10.744  26.139 -13.797  1.00 26.38      A    N
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ATOM    835  CA  HIS   565     -10.647  25.098 -14.814  1.00 24.40      A    C
ATOM    836  CB  HIS   565     -11.995  24.894 -15.510  1.00 23.27      A    C
ATOM    837  CG  HIS   565     -12.271  25.882 -16.603  1.00 24.99      A    C
ATOM    838  CD2 HIS   565     -12.022  25.831 -17.934  1.00 25.30      A    C
ATOM    839  ND1 HIS   565     -12.881  27.096 -16.375  1.00 25.00      A    N
ATOM    840  CE1 HIS   565     -12.994  27.752 -17.517  1.00 26.14      A    C
ATOM    841  NE2 HIS   565     -12.480  27.006 -18.479  1.00 24.33      A    N
ATOM    842  C   HIS   565      -9.589  25.467 -15.850  1.00 24.48      A    C
ATOM    843  O   HIS   565      -9.282  26.640 -16.046  1.00 25.19      A    O
ATOM    844  N   TRP   566      -9.029  24.463 -16.511  1.00 24.82      A    N
ATOM    845  CA  TRP   566      -8.034  24.710 -17.547  1.00 25.49      A    C
ATOM    846  CB  TRP   566      -6.618  24.503 -16.989  1.00 23.96      A    C
ATOM    847  CG  TRP   566      -6.428  23.185 -16.298  1.00 24.74      A    C
ATOM    848  CD2 TRP   566      -6.786  22.864 -14.944  1.00 24.42      A    C
ATOM    849  CE2 TRP   566      -6.434  21.519 -14.729  1.00 23.95      A    C
ATOM    850  CE3 TRP   566      -7.369  23.586 -13.896  1.00 25.18      A    C
ATOM    851  CD1 TRP   566      -5.886  22.051 -16.828  1.00 23.05      A    C
ATOM    852  NE1 TRP   566      -5.887  21.046 -15.894  1.00 23.71      A    N
ATOM    853  CZ2 TRP   566      -6.648  20.877 -13.508  1.00 25.70      A    C
ATOM    854  CZ3 TRP   566      -7.578  22.951 -12.688  1.00 24.87      A    C
ATOM    855  CH2 TRP   566      -7.220  21.608 -12.503  1.00 25.51      A    C
ATOM    856  C   TRP   566      -8.267  23.796 -18.746  1.00 25.03      A    C
ATOM    857  O   TRP   566      -8.687  22.645 -18.595  1.00 24.33      A    O
ATOM    858  N   GLU   567      -7.990  24.318 -19.934  1.00 24.46      A    N
ATOM    859  CA  GLU   567      -8.361  23.645 -21.170  1.00 25.18      A    C
ATOM    860  CB  GLU   567      -8.547  24.685 -22.274  1.00 24.46      A    C
ATOM    861  CG  GLU   567      -8.779  24.101 -23.646  1.00 24.68      A    C
ATOM    862  CD  GLU   567      -9.514  25.058 -24.557  1.00 25.49      A    C
ATOM    863  OE1 GLU   567     -10.669  25.420 -24.233  1.00 26.55      A    O
ATOM    864  OE2 GLU   567      -8.939  25.449 -25.596  1.00 28.35      A    O
ATOM    865  C   GLU   567      -7.333  22.608 -21.604  1.00 26.33      A    C
ATOM    866  O   GLU   567      -7.683  21.517 -22.072  1.00 25.57      A    O
ATOM    867  N   VAL   568      -6.063  22.953 -21.453  1.00 26.79      A    N
ATOM    868  CA  VAL   568      -4.991  22.112 -21.955  1.00 29.50      A    C
ATOM    869  CB  VAL   568      -3.874  22.974 -22.571  1.00 28.46      A    C
ATOM    870  CG1 VAL   568      -2.660  22.109 -22.901  1.00 28.66      A    C
ATOM    871  CG2 VAL   568      -4.402  23.646 -23.842  1.00 24.39      A    C
ATOM    872  C   VAL   568      -4.428  21.223 -20.853  1.00 32.48      A    C
ATOM    873  O   VAL   568      -3.985  21.710 -19.818  1.00 31.74      A    O
ATOM    874  N   GLU   569      -4.465  19.914 -21.080  1.00 36.29      A    N
ATOM    875  CA  GLU   569      -4.108  18.953 -20.047  1.00 41.85      A    C
ATOM    876  CB  GLU   569      -4.360  17.520 -20.538  1.00 36.99      A    C
ATOM    877  CG  GLU   569      -5.837  17.124 -20.604  1.00 31.58      A    C
ATOM    878  CD  GLU   569      -6.534  17.605 -21.868  1.00 30.99      A    C
ATOM    879  OE1 GLU   569      -7.728  17.286 -22.050  1.00 31.44      A    O
ATOM    880  OE2 GLU   569      -5.893  18.299 -22.687  1.00 28.21      A    O
ATOM    881  C   GLU   569      -2.653  19.114 -19.603  1.00 48.32      A    C
ATOM    882  O   GLU   569      -1.767  19.363 -20.424  1.00 47.22      A    O
ATOM    883  N   ASP   570      -2.439  18.957 -18.295  1.00 55.96      A    N
ATOM    884  CA  ASP   570      -1.185  19.278 -17.608  1.00 63.65      A    C
ATOM    885  CB  ASP   570       0.027  18.894 -18.464  1.00 66.25      A    C
ATOM    886  CG  ASP   570       0.493  17.475 -18.207  1.00 69.76      A    C
ATOM    887  OD1 ASP   570       0.436  17.039 -17.035  1.00 72.02      A    O
ATOM    888  OD2 ASP   570       0.915  16.798 -19.172  1.00 69.99      A    O
ATOM    889  C   ASP   570      -1.107  20.757 -17.236  1.00 67.13      A    C
ATOM    890  O   ASP   570      -0.426  21.538 -17.903  1.00 68.92      A    O
ATOM    891  N   LEU   571      -1.802  21.130 -16.161  1.00 69.67      A    N
ATOM    892  CA  LEU   571      -1.922  22.532 -15.760  1.00 72.16      A    C
ATOM    893  CB  LEU   571      -2.990  22.691 -14.664  1.00 71.90      A    C
ATOM    894  CG  LEU   571      -2.707  22.122 -13.265  1.00 71.59      A    C
ATOM    895  CD1 LEU   571      -3.756  22.633 -12.289  1.00 70.61      A    C
ATOM    896  CD2 LEU   571      -2.709  20.598 -13.304  1.00 72.02      A    C
ATOM    897  C   LEU   571      -0.591  23.094 -15.259  1.00 73.82      A    C
ATOM    898  O   LEU   571      -0.554  24.116 -14.566  1.00 75.52      A    O
ATOM    899  N   PRO   577       4.838  27.651 -10.922  1.00 57.36      A    N
ATOM    900  CD  PRO   577       6.223  27.162 -11.012  1.00 59.08      A    C
ATOM    901  CA  PRO   577       4.535  28.186  -9.587  1.00 56.03      A    C
ATOM    902  CB  PRO   577       5.864  28.077  -8.836  1.00 56.74      A    C
ATOM    903  CG  PRO   577       6.621  27.019  -9.563  1.00 58.87      A    C
ATOM    904  C   PRO   577       4.057  29.627  -9.671  1.00 53.74      A    C
ATOM    905  O   PRO   577       4.654  30.445 -10.365  1.00 55.21      A    O
ATOM    906  N   VAL   578       2.983  29.931  -8.956  1.00 49.44      A    N
ATOM    907  CA  VAL   578       2.397  31.260  -9.002  1.00 46.13      A    C
ATOM    908  CB  VAL   578       0.859  31.179  -8.897  1.00 45.53      A    C
ATOM    909  CG1 VAL   578       0.260  32.570  -8.892  1.00 45.46      A    C
ATOM    910  CG2 VAL   578       0.308  30.370 -10.057  1.00 47.41      A    C
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ATOM    911  C   VAL   578       2.933  32.114  -7.859  1.00 43.82      A    C
ATOM    912  O   VAL   578       2.958  31.671  -6.707  1.00 43.37      A    O
ATOM    913  N   LEU   579       3.365  33.331  -8.180  1.00 38.55      A    N
ATOM    914  CA  LEU   579       3.764  34.284  -7.150  1.00 36.58      A    C
ATOM    915  CB  LEU   579       4.586  35.429  -7.753  1.00 35.06      A    C
ATOM    916  CG  LEU   579       5.926  35.086  -8.408  1.00 35.00      A    C
ATOM    917  CD1 LEU   579       6.668  36.379  -8.730  1.00 34.09      A    C
ATOM    918  CD2 LEU   579       6.758  34.216  -7.480  1.00 32.12      A    C
ATOM    919  C   LEU   579       2.524  34.856  -6.475  1.00 35.58      A    C
ATOM    920  O   LEU   579       1.469  34.997  -7.101  1.00 35.13      A    O
ATOM    921  N   ARG   580       2.658  35.196  -5.198  1.00 35.00      A    N
ATOM    922  CA  ARG   580       1.539  35.720  -4.426  1.00 36.48      A    C
ATOM    923  CB  ARG   580       1.224  37.154  -4.863  1.00 36.24      A    C
ATOM    924  CG  ARG   580       2.223  38.184  -4.355  1.00 38.98      A    C
ATOM    925  CD  ARG   580       2.105  39.507  -5.099  1.00 38.47      A    C
ATOM    926  NE  ARG   580       2.801  40.584  -4.399  1.00 38.42      A    N
ATOM    927  CZ  ARG   580       3.303  41.662  -4.993  1.00 38.63      A    C
ATOM    928  NH1 ARG   580       3.917  42.593  -4.272  1.00 35.39      A    N
ATOM    929  NH2 ARG   580       3.196  41.807  -6.311  1.00 35.21      A    N
ATOM    930  C   ARG   580       0.299  34.839  -4.585  1.00 36.61      A    C
ATOM    931  O   ARG   580      -0.789  35.326  -4.901  1.00 36.06      A    O
ATOM    932  N   PRO   581       0.453  33.528  -4.351  1.00 37.30      A    N
ATOM    933  CD  PRO   581       1.638  32.925  -3.717  1.00 36.85      A    C
ATOM    934  CA  PRO   581      -0.611  32.544  -4.578  1.00 39.33      A    C
ATOM    935  CB  PRO   581       0.051  31.211  -4.242  1.00 40.31      A    C
ATOM    936  CG  PRO   581       1.147  31.569  -3.284  1.00 39.56      A    C
ATOM    937  C   PRO   581      -1.849  32.795  -3.724  1.00 40.74      A    C
ATOM    938  O   PRO   581      -2.935  32.322  -4.048  1.00 41.03      A    O
ATOM    939  N   ARG   582      -1.684  33.538  -2.633  1.00 41.34      A    N
ATOM    940  CA  ARG   582      -2.821  33.930  -1.809  1.00 43.35      A    C
ATOM    941  CB  ARG   582      -2.391  34.085  -0.346  1.00 44.09      A    C
ATOM    942  CG  ARG   582      -1.819  32.811   0.257  1.00 48.45      A    C
ATOM    943  CD  ARG   582      -0.916  33.106   1.448  1.00 52.04      A    C
ATOM    944  NE  ARG   582      -0.250  31.901   1.942  1.00 55.65      A    N
ATOM    945  CZ  ARG   582       0.862  31.392   1.419  1.00 58.45      A    C
ATOM    946  NH1 ARG   582       1.396  30.291   1.935  1.00 59.76      A    N
ATOM    947  NH2 ARG   582       1.442  31.979   0.378  1.00 59.55      A    N
ATOM    948  C   ARG   582      -3.411  35.243  -2.315  1.00 44.15      A    C
ATOM    949  O   ARG   582      -2.684  36.190  -2.612  1.00 45.85      A    O
ATOM    950  N   GLY   583      -4.731  35.301  -2.412  1.00 43.03      A    N
ATOM    951  CA  GLY   583      -5.352  36.495  -2.937  1.00 44.47      A    C
ATOM    952  C   GLY   583      -5.441  37.589  -1.896  1.00 47.04      A    C
ATOM    953  O   GLY   583      -5.458  37.320  -0.696  1.00 47.38      A    O
ATOM    954  N   GLN   584      -5.491  38.833  -2.355  1.00 47.91      A    N
ATOM    955  CA  GLN   584      -5.956  39.915  -1.513  1.00 50.00      A    C
ATOM    956  CB  GLN   584      -5.981  41.225  -2.302  1.00 54.25      A    C
ATOM    957  CG  GLN   584      -4.604  41.807  -2.565  1.00 60.44      A    C
ATOM    958  CD  GLN   584      -3.902  42.226  -1.286  1.00 65.17      A    C
ATOM    959  OE1 GLN   584      -4.440  43.010  -0.499  1.00 68.24      A    O
ATOM    960  NE2 GLN   584      -2.697  41.703  -1.068  1.00 66.44      A    N
ATOM    961  C   GLN   584      -7.361  39.548  -1.057  1.00 49.41      A    C
ATOM    962  O   GLN   584      -8.005  38.675  -1.640  1.00 48.96      A    O
ATOM    963  N   PRO   585      -7.858  40.204  -0.002  1.00 49.04      A    N
ATOM    964  CD  PRO   585      -7.320  41.365   0.727  1.00 48.81      A    C
ATOM    965  CA  PRO   585      -9.199  39.834   0.453  1.00 47.47      A    C
ATOM    966  CB  PRO   585      -9.475  40.806   1.602  1.00 48.68      A    C
ATOM    967  CG  PRO   585      -8.538  41.942   1.384  1.00 49.88      A    C
ATOM    968  C   PRO   585     -10.210  39.955  -0.677  1.00 45.46      A    C
ATOM    969  O   PRO   585     -10.194  40.922  -1.441  1.00 45.47      A    O
ATOM    970  N   ASN   586     -11.073  38.953  -0.784  1.00 42.42      A    N
ATOM    971  CA  ASN   586     -12.076  38.905  -1.836  1.00 39.41      A    C
ATOM    972  CB  ASN   586     -12.928  40.174  -1.818  1.00 39.50      A    C
ATOM    973  CG  ASN   586     -13.822  40.255  -0.595  1.00 43.25      A    C
ATOM    974  OD1 ASN   586     -14.272  39.236  -0.072  1.00 39.56      A    O
ATOM    975  ND2 ASN   586     -14.080  41.473  -0.128  1.00 44.78      A    N
ATOM    976  C   ASN   586     -11.481  38.706  -3.226  1.00 36.56      A    C
ATOM    977  O   ASN   586     -12.121  39.032  -4.222  1.00 37.07      A    O
ATOM    978  N   GLN   587     -10.266  38.183  -3.287  1.00 34.14      A    N
ATOM    979  CA  GLN   587      -9.629  37.824  -4.563  1.00 33.02      A    C
ATOM    980  CB  GLN   587      -8.337  38.631  -4.731  1.00 34.05      A    C
ATOM    981  CG  GLN   587      -7.551  38.349  -6.005  1.00 34.28      A    C
ATOM    982  CD  GLN   587      -6.272  39.189  -6.099  1.00 37.58      A    C
ATOM    983  OE1 GLN   587      -6.142  40.061  -6.966  1.00 36.03      A    O
ATOM    984  NE2 GLN   587      -5.328  38.927  -5.204  1.00 34.03      A    N
ATOM    985  C   GLN   587      -9.316  36.325  -4.660  1.00 32.64      A    C
ATOM    986  O   GLN   587      -8.899  35.712  -3.679  1.00 31.40      A    O
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ATOM    987  N   CYS   588      -9.523  35.750  -5.847  1.00 29.94      A    N
ATOM    988  CA  CYS   588      -9.140  34.368  -6.138  1.00 29.32      A    C
ATOM    989  C   CYS   588      -7.901  34.316  -7.018  1.00 28.74      A    C
ATOM    990  O   CYS   588      -7.692  35.196  -7.850  1.00 30.51      A    O
ATOM    991  CB  CYS   588     -10.256  33.644  -6.880  1.00 28.96      A    C
ATOM    992  SG  CYS   588     -11.771  33.388  -5.926  1.00 34.01      A    S
ATOM    993  N   VAL   589      -7.106  33.263  -6.866  1.00 27.74      A    N
ATOM    994  CA  VAL   589      -5.879  33.123  -7.644  1.00 28.08      A    C
ATOM    995  CB  VAL   589      -4.644  33.387  -6.752  1.00 28.90      A    C
ATOM    996  CG1 VAL   589      -3.364  33.266  -7.568  1.00 28.12      A    C
ATOM    997  CG2 VAL   589      -4.750  34.780  -6.123  1.00 29.60      A    C
ATOM    998  C   VAL   589      -5.761  31.728  -8.273  1.00 29.72      A    C
ATOM    999  O   VAL   589      -5.867  30.718  -7.580  1.00 29.20      A    O
ATOM   1000  N   GLY   590      -5.540  31.679  -9.585  1.00 29.03      A    N
ATOM   1001  CA  GLY   590      -5.365  30.401 -10.255  1.00 29.41      A    C
ATOM   1002  C   GLY   590      -4.067  30.331 -11.037  1.00 30.32      A    C
ATOM   1003  O   GLY   590      -3.875  31.335 -11.179  1.00 31.55      A    O
ATOM   1004  N   HIS   591      -3.732  29.151 -11.549  1.00 31.69      A    N
ATOM   1005  CA  HIS   591      -2.535  28.989 -12.367  1.00 34.04      A    C
ATOM   1006  CB  HIS   591      -2.343  27.516 -12.727  1.00 36.27      A    C
ATOM   1007  CG  HIS   591      -2.045  26.649 -11.546  1.00 44.11      A    C
ATOM   1008  CD2 HIS   591      -2.848  25.856 -10.796  1.00 46.70      A    C
ATOM   1009  ND1 HIS   591      -0.788  26.561 -10.986  1.00 46.49      A    N
ATOM   1010  CE1 HIS   591      -0.830  25.754  -9.940  1.00 46.26      A    C
ATOM   1011  NE2 HIS   591      -2.069  25.313  -9.803  1.00 49.17      A    N
ATOM   1012  C   HIS   591      -2.614  29.830 -13.640  1.00 34.21      A    C
ATOM   1013  O   HIS   591      -3.699  30.211 -14.077  1.00 32.59      A    O
ATOM   1014  N   ARG   592      -1.464  30.125 -14.234  1.00 33.67      A    N
ATOM   1015  CA  ARG   592      -1.443  31.006 -15.393  1.00 36.88      A    C
ATOM   1016  CB  ARG   592      -0.004  31.430 -15.716  1.00 39.26      A    C
ATOM   1017  CG  ARG   592       0.995  30.303 -15.710  1.00 45.79      A    C
ATOM   1018  CD  ARG   592       2.258  30.690 -16.456  1.00 50.53      A    C
ATOM   1019  NE  ARG   592       3.009  31.749 -15.787  1.00 53.30      A    N
ATOM   1020  CZ  ARG   592       3.575  32.773 -16.420  1.00 55.13      A    C
ATOM   1021  NH1 ARG   592       4.247  33.693 -15.740  1.00 55.24      A    N
ATOM   1022  NH2 ARG   592       3.465  32.879 -17.738  1.00 57.78      A    N
ATOM   1023  C   ARG   592      -2.112  30.410 -16.634  1.00 35.10      A    C
ATOM   1024  O   ARG   592      -2.566  31.149 -17.506  1.00 35.39      A    O
ATOM   1025  N   GLU   593      -2.192  29.084 -16.709  1.00 34.62      A    N
ATOM   1026  CA  GLU   593      -2.877  28.428 -17.823  1.00 33.78      A    C
ATOM   1027  CB  GLU   593      -2.181  27.120 -18.189  1.00 36.33      A    C
ATOM   1028  CG  GLU   593      -0.983  27.299 -19.098  1.00 46.12      A    C
ATOM   1029  CD  GLU   593       0.219  27.845 -18.362  1.00 52.97      A    C
ATOM   1030  OE1 GLU   593       0.553  27.303 -17.284  1.00 55.60      A    O
ATOM   1031  OE2 GLU   593       0.828  28.818 -18.861  1.00 57.96      A    O
ATOM   1032  C   GLU   593      -4.350  28.135 -17.558  1.00 32.11      A    C
ATOM   1033  O   GLU   593      -5.019  27.525 -18.388  1.00 29.63      A    O
ATOM   1034  N   ALA   594      -4.855  28.559 -16.406  1.00 28.88      A    N
ATOM   1035  CA  ALA   594      -6.236  28.267 -16.046  1.00 28.39      A    C
ATOM   1036  CB  ALA   594      -6.283  27.571 -14.687  1.00 26.56      A    C
ATOM   1037  C   ALA   594      -7.092  29.528 -16.018  1.00 28.13      A    C
ATOM   1038  O   ALA   594      -6.584  30.637 -15.843  1.00 28.06      A    O
ATOM   1039  N   SER   595      -8.396  29.358 -16.197  1.00 25.46      A    N
ATOM   1040  CA  SER   595      -9.335  30.416 -15.844  1.00 25.16      A    C
ATOM   1041  CB  SER   595     -10.656  30.226 -16.588  1.00 24.79      A    C
ATOM   1042  OG  SER   595     -10.490  30.436 -17.976  1.00 27.76      A    O
ATOM   1043  C   SER   595      -9.580  30.372 -14.338  1.00 24.98      A    C
ATOM   1044  O   SER   595      -9.479  29.312 -13.720  1.00 24.48      A    O
ATOM   1045  N   ILE   596      -9.891  31.524 -13.752  1.00 25.25      A    N
ATOM   1046  CA  ILE   596     -10.212  31.601 -12.333  1.00 25.69      A    C
ATOM   1047  CB  ILE   596      -9.165  32.467 -11.577  1.00 27.15      A    C
ATOM   1048  CG2 ILE   596      -9.259  33.923 -12.023  1.00 29.13      A    C
ATOM   1049  CG1 ILE   596      -9.375  32.349 -10.067  1.00 26.07      A    C
ATOM   1050  CD1 ILE   596      -8.902  31.029  -9.490  1.00 24.57      A    C
ATOM   1051  C   ILE   596     -11.619  32.192 -12.154  1.00 24.88      A    C
ATOM   1052  O   ILE   596     -12.005  33.135 -12.849  1.00 24.39      A    O
ATOM   1053  N   HIS   597     -12.385  31.613 -11.235  1.00 23.51      A    N
ATOM   1054  CA  HIS   597     -13.795  31.956 -11.062  1.00 22.95      A    C
ATOM   1055  CB  HIS   597     -14.689  30.798 -11.522  1.00 23.62      A    C
ATOM   1056  CG  HIS   597     -14.325  30.238 -12.863  1.00 22.44      A    C
ATOM   1057  CD2 HIS   597     -13.774  29.053 -13.213  1.00 25.26      A    C
ATOM   1058  ND1 HIS   597     -14.579  30.904 -14.042  1.00 23.06      A    N
ATOM   1059  CE1 HIS   597     -14.206  30.152 -15.063  1.00 23.37      A    C
ATOM   1060  NE2 HIS   597     -13.715  29.022 -14.588  1.00 24.84      A    N
ATOM   1061  C   HIS   597     -14.076  32.224  -9.586  1.00 25.18      A    C
ATOM   1062  O   HIS   597     -13.767  31.394  -8.735  1.00 25.64      A    O
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ATOM   1063  N   ALA   598     -14.672  33.373  -9.289  1.00 25.15      A    N
ATOM   1064  CA  ALA   598     -14.969  33.749  -7.909  1.00 26.23      A    C
ATOM   1065  CB  ALA   598     -14.384  35.114  -7.606  1.00 23.97      A    C
ATOM   1066  C   ALA   598     -16.472  33.778  -7.680  1.00 27.18      A    C
ATOM   1067  O   ALA   598     -17.224  34.295  -8.508  1.00 27.20      A    O
ATOM   1068  N   SER   599     -16.907  33.223  -6.556  1.00 26.97      A    N
ATOM   1069  CA  SER   599     -18.286  33.395  -6.133  1.00 29.40      A    C
ATOM   1070  CB  SER   599     -18.775  32.168  -5.376  1.00 30.19      A    C
ATOM   1071  OG  SER   599     -20.054  32.417  -4.826  1.00 33.89      A    O
ATOM   1072  C   SER   599     -18.387  34.618  -5.232  1.00 30.33      A    C
ATOM   1073  O   SER   599     -17.861  34.629  -4.120  1.00 31.52      A    O
ATOM   1074  N   CYS   600     -19.064  35.646  -5.728  1.00 32.35      A    N
ATOM   1075  CA  CYS   600     -19.170  36.923  -5.036  1.00 35.21      A    C
ATOM   1076  C   CYS   600     -20.606  37.142  -4.591  1.00 36.07      A    C
ATOM   1077  O   CYS   600     -21.520  37.174  -5.413  1.00 35.83      A    O
ATOM   1078  CB  CYS   600     -18.744  38.055  -5.971  1.00 35.88      A    C
ATOM   1079  SG  CYS   600     -17.010  37.951  -6.513  1.00 39.28      A    S
ATOM   1080  N   CYS   601     -20.816  37.282  -3.289  1.00 38.47      A    N
ATOM   1081  CA  CYS   601     -22.175  37.408  -2.790  1.00 40.53      A    C
ATOM   1082  C   CYS   601     -22.415  38.723  -2.079  1.00 42.20      A    C
ATOM   1083  O   CYS   601     -21.636  39.141  -1.218  1.00 40.46      A    O
ATOM   1084  CB  CYS   601     -22.511  36.277  -1.832  1.00 40.11      A    C
ATOM   1085  SG  CYS   601     -22.373  34.577  -2.469  1.00 41.58      A    S
ATOM   1086  N   HIS   602     -23.510  39.369  -2.453  1.00 44.38      A    N
ATOM   1087  CA  HIS   602     -24.081  40.418  -1.637  1.00 47.94      A    C
ATOM   1088  CB  HIS   602     -24.858  41.399  -2.518  1.00 51.93      A    C
ATOM   1089  CG  HIS   602     -25.222  42.674  -1.826  1.00 58.56      A    C
ATOM   1090  CD2 HIS   602     -24.451  43.620  -1.237  1.00 60.81      A    C
ATOM   1091  ND1 HIS   602     -26.526  43.095  -1.678  1.00 60.77      A    N
ATOM   1092  CE1 HIS   602     -26.543  44.246  -1.028  1.00 62.59      A    C
ATOM   1093  NE2 HIS   602     -25.297  44.587  -0.749  1.00 62.85      A    N
ATOM   1094  C   HIS   602     -25.017  39.713  -0.661  1.00 48.23      A    C
ATOM   1095  O   HIS   602     -26.025  39.129  -1.063  1.00 45.88      A    O
ATOM   1096  N   ALA   603     -24.661  39.746   0.619  1.00 49.87      A    N
ATOM   1097  CA  ALA   603     -25.450  39.089   1.653  1.00 52.49      A    C
ATOM   1098  CB  ALA   603     -25.006  37.647   1.804  1.00 50.93      A    C
ATOM   1099  C   ALA   603     -25.287  39.829   2.975  1.00 53.94      A    C
ATOM   1100  O   ALA   603     -24.183  39.934   3.507  1.00 55.03      A    O
ATOM   1101  N   PRO   604     -26.394  40.340   3.530  1.00 55.31      A    N
ATOM   1102  CD  PRO   604     -27.775  40.052   3.105  1.00 56.68      A    C
ATOM   1103  CA  PRO   604     -26.349  41.196   4.721  1.00 55.19      A    C
ATOM   1104  CB  PRO   604     -27.808  41.600   4.931  1.00 56.90      A    C
ATOM   1105  CG  PRO   604     -28.600  40.510   4.278  1.00 57.49      A    C
ATOM   1106  C   PRO   604     -25.762  40.507   5.951  1.00 53.84      A    C
ATOM   1107  O   PRO   604     -26.291  39.504   6.429  1.00 53.50      A    O
ATOM   1108  N   GLY   605     -24.664  41.058   6.459  1.00 51.66      A    N
ATOM   1109  CA  GLY   605     -24.084  40.547   7.683  1.00 49.75      A    C
ATOM   1110  C   GLY   605     -23.181  39.350   7.466  1.00 49.32      A    C
ATOM   1111  O   GLY   605     -22.695  38.750   8.426  1.00 49.78      A    O
ATOM   1112  N   LEU   606     -22.951  38.991   6.209  1.00 47.32      A    N
ATOM   1113  CA  LEU   606     -22.048  37.888   5.914  1.00 45.14      A    C
ATOM   1114  CB  LEU   606     -22.319  37.334   4.516  1.00 46.10      A    C
ATOM   1115  CG  LEU   606     -21.432  36.150   4.128  1.00 46.60      A    C
ATOM   1116  CD1 LEU   606     -21.603  35.038   5.146  1.00 45.47      A    C
ATOM   1117  CD2 LEU   606     -21.794  35.665   2.731  1.00 47.85      A    C
ATOM   1118  C   LEU   606     -20.599  38.351   6.015  1.00 43.35      A    C
ATOM   1119  O   LEU   606     -20.226  39.381   5.460  1.00 42.73      A    O
ATOM   1120  N   GLU   607     -19.792  37.584   6.737  1.00 41.82      A    N
ATOM   1121  CA  GLU   607     -18.388  37.913   6.949  1.00 41.58      A    C
ATOM   1122  CB  GLU   607     -18.194  38.408   8.390  1.00 43.33      A    C
ATOM   1123  CG  GLU   607     -16.751  38.685   8.788  1.00 48.75      A    C
ATOM   1124  CD  GLU   607     -16.585  38.939  10.289  1.00 52.26      A    C
ATOM   1125  OE1 GLU   607     -17.466  38.528  11.082  1.00 53.24      A    O
ATOM   1126  OE2 GLU   607     -15.564  39.548  10.675  1.00 52.44      A    O
ATOM   1127  C   GLU   607     -17.541  36.659   6.699  1.00 40.02      A    C
ATOM   1128  O   GLU   607     -17.845  35.587   7.217  1.00 39.94      A    O
ATOM   1129  N   CYS   608     -16.483  36.786   5.905  1.00 38.50      A    N
ATOM   1130  CA  CYS   608     -15.634  35.635   5.616  1.00 38.05      A    C
ATOM   1131  C   CYS   608     -14.154  35.926   5.793  1.00 38.09      A    C
ATOM   1132  O   CYS   608     -13.724  37.076   5.752  1.00 38.32      A    O
ATOM   1133  CB  CYS   608     -15.861  35.139   4.192  1.00 36.34      A    C
ATOM   1134  SG  CYS   608     -17.584  34.811   3.711  1.00 38.54      A    S
ATOM   1135  N   LYS   609     -13.381  34.864   5.981  1.00 37.60      A    N
ATOM   1136  CA  LYS   609     -11.938  34.967   6.132  1.00 37.54      A    C
ATOM   1137  CB  LYS   609     -11.566  35.070   7.611  1.00 36.16      A    C
ATOM   1138  CG  LYS   609     -11.789  33.783   8.375  1.00 39.02      A    C
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ATOM	1139	CD	LYS	609	-11.435	33.924	9.849	1.00	42.30	A	C
ATOM	1140	CE	LYS	609	-11.847	32.676	10.614	1.00	42.81	A	C
ATOM	1141	NZ	LYS	609	-11.420	32.722	12.039	1.00	44.87	A	N
ATOM	1142	C	LYS	609	-11.309	33.710	5.538	1.00	37.29	A	C
ATOM	1143	O	LYS	609	-12.004	32.728	5.264	1.00	37.32	A	O
ATOM	1144	N	VAL	610	-9.995	33.742	5.351	1.00	36.83	A	N
ATOM	1145	CA	VAL	610	-9.284	32.629	4.743	1.00	36.94	A	C
ATOM	1146	CB	VAL	610	-8.484	33.091	3.511	1.00	37.85	A	C
ATOM	1147	CG1	VAL	610	-7.745	31.913	2.902	1.00	38.22	A	C
ATOM	1148	CG2	VAL	610	-9.423	33.721	2.488	1.00	37.85	A	C
ATOM	1149	C	VAL	610	-8.323	31.984	5.729	1.00	37.80	A	C
ATOM	1150	O	VAL	610	-7.599	32.675	6.446	1.00	38.87	A	O
ATOM	1151	N	LYS	611	-8.330	30.655	5.761	1.00	37.56	A	N
ATOM	1152	CA	LYS	611	-7.388	29.883	6.568	1.00	38.62	A	C
ATOM	1153	CB	LYS	611	-8.116	29.133	7.684	1.00	38.63	A	C
ATOM	1154	CG	LYS	611	-8.917	30.008	8.625	1.00	42.75	A	C
ATOM	1155	CD	LYS	611	-9.432	29.193	9.793	1.00	42.52	A	C
ATOM	1156	CE	LYS	611	-8.336	28.291	10.342	1.00	44.58	A	C
ATOM	1157	NZ	LYS	611	-8.782	27.548	11.559	1.00	48.18	A	N
ATOM	1158	C	LYS	611	-6.718	28.865	5.664	1.00	39.23	A	C
ATOM	1159	O	LYS	611	-7.365	28.288	4.791	1.00	39.27	A	O
ATOM	1160	N	GLU	612	-5.429	28.632	5.874	1.00	39.16	A	N
ATOM	1161	CA	GLU	612	-4.726	27.624	5.094	1.00	42.41	A	C
ATOM	1162	CB	GLU	612	-3.867	28.295	4.025	1.00	44.71	A	C
ATOM	1163	CG	GLU	612	-2.809	29.235	4.571	1.00	49.08	A	C
ATOM	1164	CD	GLU	612	-2.008	29.896	3.465	1.00	53.20	A	C
ATOM	1165	OE1	GLU	612	-2.592	30.697	2.699	1.00	54.07	A	O
ATOM	1166	OE2	GLU	612	-0.796	29.609	3.360	1.00	53.71	A	O
ATOM	1167	C	GLU	612	-3.853	26.736	5.970	1.00	42.48	A	C
ATOM	1168	O	GLU	612	-3.592	27.058	7.127	1.00	43.10	A	O
ATOM	1169	N	HIS	613	-3.414	25.613	5.415	1.00	41.33	A	N
ATOM	1170	CA	HIS	613	-2.438	24.758	6.074	1.00	42.62	A	C
ATOM	1171	CB	HIS	613	-3.140	23.783	7.025	1.00	44.30	A	C
ATOM	1172	CG	HIS	613	-2.222	22.782	7.658	1.00	44.52	A	C
ATOM	1173	CD2	HIS	613	-1.137	22.941	8.453	1.00	44.27	A	C
ATOM	1174	ND1	HIS	613	-2.381	21.421	7.498	1.00	45.05	A	N
ATOM	1175	CE1	HIS	613	-1.435	20.786	8.167	1.00	43.50	A	C
ATOM	1176	NE2	HIS	613	-0.667	21.685	8.755	1.00	44.92	A	N
ATOM	1177	C	HIS	613	-1.677	23.994	5.003	1.00	44.32	A	C
ATOM	1178	O	HIS	613	-2.277	23.299	4.183	1.00	44.33	A	O
ATOM	1179	N	GLY	614	-0.356	24.142	5.000	1.00	45.57	A	N
ATOM	1180	CA	GLY	614	0.460	23.465	4.010	1.00	46.92	A	C
ATOM	1181	C	GLY	614	1.295	22.352	4.609	1.00	48.04	A	C
ATOM	1182	O	GLY	614	1.660	22.402	5.779	1.00	48.09	A	O
ATOM	1183	N	ILE	615	1.594	21.342	3.801	1.00	50.58	A	N
ATOM	1184	CA	ILE	615	2.451	20.237	4.214	1.00	53.77	A	C
ATOM	1185	CB	ILE	615	1.643	18.935	4.407	1.00	55.20	A	C
ATOM	1186	CG2	ILE	615	2.575	17.774	4.701	1.00	54.79	A	C
ATOM	1187	CG1	ILE	615	0.648	19.109	5.551	1.00	55.97	A	C
ATOM	1188	CD1	ILE	615	1.290	19.599	6.820	1.00	58.58	A	C
ATOM	1189	C	ILE	615	3.505	19.996	3.143	1.00	55.43	A	C
ATOM	1190	O	ILE	615	3.203	20.011	1.950	1.00	55.42	A	O
ATOM	1191	N	PRO	616	4.762	19.778	3.559	1.00	57.06	A	N
ATOM	1192	CD	PRO	616	5.226	19.885	4.952	1.00	57.57	A	C
ATOM	1193	CA	PRO	616	5.878	19.575	2.627	1.00	58.08	A	C
ATOM	1194	CB	PRO	616	7.100	19.486	3.541	1.00	58.83	A	C
ATOM	1195	CG	PRO	616	6.684	20.196	4.792	1.00	58.90	A	C
ATOM	1196	C	PRO	616	5.717	18.322	1.768	1.00	58.68	A	C
ATOM	1197	O	PRO	616	5.755	18.392	0.540	1.00	59.30	A	O
ATOM	1198	N	ALA	617	5.539	17.177	2.420	1.00	58.67	A	N
ATOM	1199	CA	ALA	617	5.396	15.915	1.704	1.00	60.47	A	C
ATOM	1200	CB	ALA	617	6.581	15.005	2.006	1.00	60.68	A	C
ATOM	1201	C	ALA	617	4.097	15.227	2.095	1.00	60.64	A	C
ATOM	1202	O	ALA	617	4.096	14.273	2.876	1.00	60.93	A	O
ATOM	1203	N	PRO	618	2.971	15.696	1.541	1.00	60.54	A	N
ATOM	1204	CD	PRO	618	2.883	16.645	0.417	1.00	60.69	A	C
ATOM	1205	CA	PRO	618	1.660	15.169	1.929	1.00	60.60	A	C
ATOM	1206	CB	PRO	618	0.679	15.977	1.079	1.00	60.68	A	C
ATOM	1207	CG	PRO	618	1.485	16.442	-0.093	1.00	61.12	A	C
ATOM	1208	C	PRO	618	1.557	13.669	1.673	1.00	60.54	A	C
ATOM	1209	O	PRO	618	1.962	13.178	0.621	1.00	60.46	A	O
ATOM	1210	N	GLN	619	1.017	12.947	2.647	1.00	61.08	A	N
ATOM	1211	CA	GLN	619	0.972	11.493	2.585	1.00	62.08	A	C
ATOM	1212	CB	GLN	619	1.260	10.902	3.963	1.00	64.75	A	C
ATOM	1213	CG	GLN	619	2.736	10.820	4.294	1.00	69.13	A	C
ATOM	1214	CD	GLN	619	3.405	9.620	3.650	1.00	72.41	A	C

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ATOM   1215  OE1 GLN   619       4.597   9.381   3.844  1.00 75.41      A    O
ATOM   1216  NE2 GLN   619       2.636   8.855   2.879  1.00 73.28      A    N
ATOM   1217  C   GLN   619      -0.366  10.974   2.082  1.00 60.67      A    C
ATOM   1218  O   GLN   619      -0.690   9.795   2.248  1.00 61.39      A    O
ATOM   1219  N   GLY   620      -1.140  11.858   1.463  1.00 58.38      A    N
ATOM   1220  CA  GLY   620      -2.412  11.446   0.902  1.00 55.08      A    C
ATOM   1221  C   GLY   620      -3.476  12.526   0.959  1.00 52.97      A    C
ATOM   1222  O   GLY   620      -4.332  12.608   0.076  1.00 52.81      A    O
ATOM   1223  N   GLN   621      -3.433  13.353   1.998  1.00 49.05      A    N
ATOM   1224  CA  GLN   621      -4.334  14.486   2.084  1.00 46.66      A    C
ATOM   1225  CB  GLN   621      -5.715  14.040   2.571  1.00 48.24      A    C
ATOM   1226  CG  GLN   621      -5.783  13.670   4.038  1.00 50.90      A    C
ATOM   1227  CD  GLN   621      -7.187  13.283   4.471  1.00 52.62      A    C
ATOM   1228  OE1 GLN   621      -8.035  12.943   3.645  1.00 53.23      A    O
ATOM   1229  NE2 GLN   621      -7.440  13.338   5.774  1.00 53.97      A    N
ATOM   1230  C   GLN   621      -3.793  15.570   2.998  1.00 44.42      A    C
ATOM   1231  O   GLN   621      -2.991  15.310   3.894  1.00 43.84      A    O
ATOM   1232  N   VAL   622      -4.230  16.795   2.742  1.00 41.31      A    N
ATOM   1233  CA  VAL   622      -3.895  17.932   3.576  1.00 38.86      A    C
ATOM   1234  CB  VAL   622      -3.022  18.943   2.814  1.00 37.61      A    C
ATOM   1235  CG1 VAL   622      -2.539  20.029   3.759  1.00 39.32      A    C
ATOM   1236  CG2 VAL   622      -1.858  18.234   2.162  1.00 38.06      A    C
ATOM   1237  C   VAL   622      -5.216  18.597   3.935  1.00 38.57      A    C
ATOM   1238  O   VAL   622      -6.068  18.804   3.068  1.00 38.51      A    O
ATOM   1239  N   THR   623      -5.392  18.926   5.207  1.00 35.77      A    N
ATOM   1240  CA  THR   623      -6.656  19.479   5.664  1.00 33.79      A    C
ATOM   1241  CB  THR   623      -7.387  18.504   6.604  1.00 34.25      A    C
ATOM   1242  OG1 THR   623      -6.629  18.345   7.810  1.00 33.79      A    O
ATOM   1243  CG2 THR   623      -7.559  17.148   5.932  1.00 33.56      A    C
ATOM   1244  C   THR   623      -6.457  20.786   6.407  1.00 33.76      A    C
ATOM   1245  O   THR   623      -5.388  21.040   6.968  1.00 33.37      A    O
ATOM   1246  N   VAL   624      -7.495  21.615   6.401  1.00 32.04      A    N
ATOM   1247  CA  VAL   624      -7.583  22.747   7.310  1.00 31.45      A    C
ATOM   1248  CB  VAL   624      -6.990  24.032   6.674  1.00 32.64      A    C
ATOM   1249  CG1 VAL   624      -7.790  24.431   5.446  1.00 31.89      A    C
ATOM   1250  CG2 VAL   624      -6.973  25.161   7.699  1.00 31.91      A    C
ATOM   1251  C   VAL   624      -9.058  22.959   7.630  1.00 33.28      A    C
ATOM   1252  O   VAL   624      -9.925  22.714   6.789  1.00 32.07      A    O
ATOM   1253  N   ALA   625      -9.346  23.392   8.852  1.00 33.71      A    N
ATOM   1254  CA  ALA   625     -10.723  23.491   9.305  1.00 35.41      A    C
ATOM   1255  CB  ALA   625     -10.971  22.492  10.434  1.00 33.24      A    C
ATOM   1256  C   ALA   625     -11.060  24.900   9.768  1.00 37.55      A    C
ATOM   1257  O   ALA   625     -10.185  25.646  10.200  1.00 38.33      A    O
ATOM   1258  N   CYS   626     -12.335  25.262   9.661  1.00 39.75      A    N
ATOM   1259  CA  CYS   626     -12.840  26.477  10.286  1.00 41.17      A    C
ATOM   1260  C   CYS   626     -13.170  26.144  11.743  1.00 43.54      A    C
ATOM   1261  O   CYS   626     -13.781  25.110  12.024  1.00 44.60      A    O
ATOM   1262  CB  CYS   626     -14.110  26.959   9.575  1.00 39.19      A    C
ATOM   1263  SG  CYS   626     -13.999  27.167   7.766  1.00 41.43      A    S
ATOM   1264  N   GLU   627     -12.770  27.010  12.669  1.00 45.20      A    N
ATOM   1265  CA  GLU   627     -13.077  26.793  14.085  1.00 47.32      A    C
ATOM   1266  CB  GLU   627     -12.212  27.696  14.966  1.00 47.58      A    C
ATOM   1267  CG  GLU   627     -12.348  29.175  14.660  1.00 49.95      A    C
ATOM   1268  CD  GLU   627     -11.403  29.627  13.566  1.00 52.51      A    C
ATOM   1269  OE1 GLU   627     -10.990  30.804  13.592  1.00 55.13      A    O
ATOM   1270  OE2 GLU   627     -11.071  28.810  12.681  1.00 52.16      A    O
ATOM   1271  C   GLU   627     -14.551  27.072  14.362  1.00 47.35      A    C
ATOM   1272  O   GLU   627     -15.252  27.648  13.522  1.00 46.45      A    O
ATOM   1273  N   GLU   628     -15.022  26.665  15.538  1.00 47.46      A    N
ATOM   1274  CA  GLU   628     -16.438  26.788  15.857  1.00 48.46      A    C
ATOM   1275  CB  GLU   628     -16.740  26.197  17.243  1.00 53.92      A    C
ATOM   1276  CG  GLU   628     -16.323  27.076  18.415  1.00 59.91      A    C
ATOM   1277  CD  GLU   628     -16.746  26.502  19.765  1.00 63.80      A    C
ATOM   1278  OE1 GLU   628     -16.934  27.291  20.719  1.00 63.78      A    O
ATOM   1279  OE2 GLU   628     -16.889  25.264  19.873  1.00 66.46      A    O
ATOM   1280  C   GLU   628     -16.840  28.257  15.815  1.00 46.20      A    C
ATOM   1281  O   GLU   628     -16.043  29.141  16.132  1.00 46.72      A    O
ATOM   1282  N   GLY   629     -18.077  28.513  15.410  1.00 42.69      A    N
ATOM   1283  CA  GLY   629     -18.499  29.878  15.171  1.00 40.02      A    C
ATOM   1284  C   GLY   629     -18.411  30.246  13.700  1.00 39.40      A    C
ATOM   1285  O   GLY   629     -19.083  31.173  13.250  1.00 40.26      A    O
ATOM   1286  N   TRP   630     -17.582  29.526  12.947  1.00 37.59      A    N
ATOM   1287  CA  TRP   630     -17.418  29.790  11.515  1.00 35.74      A    C
ATOM   1288  CB  TRP   630     -15.966  30.140  11.191  1.00 33.84      A    C
ATOM   1289  CG  TRP   630     -15.498  31.408  11.807  1.00 36.54      A    C
ATOM   1290  CD2 TRP   630     -15.352  32.675  11.155  1.00 36.10      A    C
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ATOM   1291  CE2 TRP   630     -14.864  33.585  12.118  1.00 38.43      A    C
ATOM   1292  CE3 TRP   630     -15.584  33.130   9.853  1.00 36.07      A    C
ATOM   1293  CD1 TRP   630     -15.104  31.595  13.105  1.00 37.06      A    C
ATOM   1294  NE1 TRP   630     -14.721  32.900  13.297  1.00 36.65      A    N
ATOM   1295  CZ2 TRP   630     -14.607  34.923  11.818  1.00 36.82      A    C
ATOM   1296  CZ3 TRP   630     -15.328  34.455   9.556  1.00 37.24      A    C
ATOM   1297  CH2 TRP   630     -14.844  35.338  10.535  1.00 37.91      A    C
ATOM   1298  C   TRP   630     -17.821  28.570  10.701  1.00 35.13      A    C
ATOM   1299  O   TRP   630     -17.703  27.434  11.165  1.00 35.66      A    O
ATOM   1300  N   THR   631     -18.285  28.804   9.481  1.00 32.78      A    N
ATOM   1301  CA  THR   631     -18.715  27.711   8.626  1.00 31.14      A    C
ATOM   1302  CB  THR   631     -20.210  27.847   8.289  1.00 31.87      A    C
ATOM   1303  OG1 THR   631     -20.961  27.895   9.508  1.00 28.32      A    O
ATOM   1304  CG2 THR   631     -20.686  26.666   7.452  1.00 28.98      A    C
ATOM   1305  C   THR   631     -17.900  27.686   7.337  1.00 30.23      A    C
ATOM   1306  O   THR   631     -17.683  28.714   6.708  1.00 29.72      A    O
ATOM   1307  N   LEU   632     -17.443  26.502   6.956  1.00 29.92      A    N
ATOM   1308  CA  LEU   632     -16.662  26.336   5.739  1.00 28.80      A    C
ATOM   1309  CB  LEU   632     -16.011  24.954   5.746  1.00 27.20      A    C
ATOM   1310  CG  LEU   632     -15.186  24.597   4.510  1.00 30.81      A    C
ATOM   1311  CD1 LEU   632     -14.066  25.609   4.346  1.00 30.24      A    C
ATOM   1312  CD2 LEU   632     -14.624  23.184   4.657  1.00 31.20      A    C
ATOM   1313  C   LEU   632     -17.550  26.493   4.498  1.00 28.43      A    C
ATOM   1314  O   LEU   632     -18.496  25.727   4.309  1.00 28.53      A    O
ATOM   1315  N   THR   633     -17.256  27.486   3.660  1.00 26.00      A    N
ATOM   1316  CA  THR   633     -18.047  27.692   2.449  1.00 27.86      A    C
ATOM   1317  CB  THR   633     -18.520  29.168   2.306  1.00 27.33      A    C
ATOM   1318  OG1 THR   633     -17.382  30.024   2.156  1.00 25.74      A    O
ATOM   1319  CG2 THR   633     -19.320  29.598   3.530  1.00 23.90      A    C
ATOM   1320  C   THR   633     -17.271  27.309   1.188  1.00 29.32      A    C
ATOM   1321  O   THR   633     -17.864  26.985   0.160  1.00 29.29      A    O
ATOM   1322  N   GLY   634     -15.946  27.352   1.267  1.00 29.44      A    N
ATOM   1323  CA  GLY   634     -15.131  27.078   0.096  1.00 29.46      A    C
ATOM   1324  C   GLY   634     -13.850  26.344   0.437  1.00 29.50      A    C
ATOM   1325  O   GLY   634     -13.244  26.581   1.485  1.00 30.22      A    O
ATOM   1326  N   CYS   635     -13.432  25.453  -0.454  1.00 28.04      A    N
ATOM   1327  CA  CYS   635     -12.296  24.581  -0.190  1.00 28.82      A    C
ATOM   1328  C   CYS   635     -11.526  24.342  -1.477  1.00 29.22      A    C
ATOM   1329  O   CYS   635     -12.107  23.936  -2.477  1.00 29.78      A    O
ATOM   1330  CB  CYS   635     -12.798  23.252   0.372  1.00 27.17      A    C
ATOM   1331  SG  CYS   635     -11.559  21.934   0.583  1.00 31.95      A    S
ATOM   1332  N   SER   636     -10.222  24.593  -1.449  1.00 29.87      A    N
ATOM   1333  CA  SER   636      -9.392  24.423  -2.635  1.00 31.98      A    C
ATOM   1334  CB  SER   636      -9.557  25.623  -3.567  1.00 30.53      A    C
ATOM   1335  OG  SER   636      -9.043  26.792  -2.957  1.00 32.60      A    O
ATOM   1336  C   SER   636      -7.916  24.274  -2.276  1.00 33.39      A    C
ATOM   1337  O   SER   636      -7.526  24.394  -1.113  1.00 33.25      A    O
ATOM   1338  N   ALA   637      -7.099  24.011  -3.289  1.00 35.31      A    N
ATOM   1339  CA  ALA   637      -5.653  23.970  -3.117  1.00 37.57      A    C
ATOM   1340  CB  ALA   637      -5.063  22.828  -3.937  1.00 35.81      A    C
ATOM   1341  C   ALA   637      -5.036  25.296  -3.554  1.00 38.81      A    C
ATOM   1342  O   ALA   637      -5.468  25.899  -4.537  1.00 39.80      A    O
ATOM   1343  N   LEU   638      -4.036  25.746  -2.811  1.00 40.54      A    N
ATOM   1344  CA  LEU   638      -3.236  26.893  -3.216  1.00 41.27      A    C
ATOM   1345  CB  LEU   638      -2.161  27.171  -2.163  1.00 41.30      A    C
ATOM   1346  CG  LEU   638      -1.633  28.595  -1.974  1.00 43.45      A    C
ATOM   1347  CD1 LEU   638      -2.751  29.509  -1.502  1.00 43.30      A    C
ATOM   1348  CD2 LEU   638      -0.505  28.577  -0.952  1.00 44.32      A    C
ATOM   1349  C   LEU   638      -2.583  26.506  -4.536  1.00 43.36      A    C
ATOM   1350  O   LEU   638      -2.027  25.415  -4.665  1.00 43.89      A    O
ATOM   1351  N   PRO   639      -2.654  27.386  -5.541  1.00 44.54      A    N
ATOM   1352  CD  PRO   639      -3.210  28.751  -5.535  1.00 42.27      A    C
ATOM   1353  CA  PRO   639      -2.072  27.020  -6.838  1.00 45.55      A    C
ATOM   1354  CB  PRO   639      -2.528  28.144  -7.764  1.00 44.15      A    C
ATOM   1355  CG  PRO   639      -2.731  29.322  -6.849  1.00 44.46      A    C
ATOM   1356  C   PRO   639      -0.551  26.924  -6.745  1.00 48.67      A    C
ATOM   1357  O   PRO   639       0.128  27.928  -6.530  1.00 49.10      A    O
ATOM   1358  N   GLY   640      -0.021  25.714  -6.895  1.00 52.08      A    N
ATOM   1359  CA  GLY   640       1.417  25.517  -6.809  1.00 55.68      A    C
ATOM   1360  C   GLY   640       1.913  24.533  -7.849  1.00 58.10      A    C
ATOM   1361  O   GLY   640       1.201  24.233  -8.802  1.00 58.79      A    O
ATOM   1362  N   THR   641       3.129  24.025  -7.679  1.00 61.04      A    N
ATOM   1363  CA  THR   641       3.606  22.939  -8.532  1.00 64.61      A    C
ATOM   1364  CB  THR   641       5.134  22.731  -8.392  1.00 65.87      A    C
ATOM   1365  OG1 THR   641       5.489  22.650  -7.006  1.00 67.03      A    O
ATOM   1366  CG2 THR   641       5.888  23.877  -9.048  1.00 66.73      A    C
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ATOM   1367  C   THR   641       2.883  21.653  -8.152  1.00 66.27      A    C
ATOM   1368  O   THR   641       3.336  20.548  -8.461  1.00 67.72      A    O
ATOM   1369  N   SER   642       1.740  21.824  -7.492  1.00 67.29      A    N
ATOM   1370  CA  SER   642       0.977  20.732  -6.901  1.00 66.68      A    C
ATOM   1371  CB  SER   642      -0.227  21.301  -6.138  1.00 68.50      A    C
ATOM   1372  OG  SER   642       0.163  22.269  -5.174  1.00 65.92      A    O
ATOM   1373  C   SER   642       0.490  19.719  -7.927  1.00 66.01      A    C
ATOM   1374  O   SER   642       0.079  20.081  -9.031  1.00 66.17      A    O
ATOM   1375  N   HIS   643       0.507  18.445  -7.548  1.00 65.59      A    N
ATOM   1376  CA  HIS   643      -0.260  17.419  -8.250  1.00 63.85      A    C
ATOM   1377  CB  HIS   643       0.565  16.134  -8.445  1.00 68.67      A    C
ATOM   1378  CG  HIS   643       2.049  16.339  -8.398  1.00 73.45      A    C
ATOM   1379  CD2 HIS   643       2.830  17.348  -8.853  1.00 75.15      A    C
ATOM   1380  ND1 HIS   643       2.905  15.423  -7.823  1.00 75.08      A    N
ATOM   1381  CE1 HIS   643       4.148  15.859  -7.925  1.00 76.32      A    C
ATOM   1382  NE2 HIS   643       4.130  17.025  -8.546  1.00 76.69      A    N
ATOM   1383  C   HIS   643      -1.461  17.099  -7.368  1.00 59.62      A    C
ATOM   1384  O   HIS   643      -1.347  16.321  -6.418  1.00 61.00      A    O
ATOM   1385  N   VAL   644      -2.606  17.697  -7.672  1.00 53.01      A    N
ATOM   1386  CA  VAL   644      -3.775  17.546  -6.813  1.00 47.88      A    C
ATOM   1387  CB  VAL   644      -4.312  18.931  -6.395  1.00 48.74      A    C
ATOM   1388  CG1 VAL   644      -5.721  18.813  -5.835  1.00 46.18      A    C
ATOM   1389  CG2 VAL   644      -3.379  19.535  -5.359  1.00 46.52      A    C
ATOM   1390  C   VAL   644      -4.894  16.729  -7.456  1.00 44.30      A    C
ATOM   1391  O   VAL   644      -5.270  16.973  -8.605  1.00 41.49      A    O
ATOM   1392  N   LEU   645      -5.416  15.756  -6.711  1.00 39.48      A    N
ATOM   1393  CA  LEU   645      -6.514  14.921  -7.198  1.00 35.69      A    C
ATOM   1394  CB  LEU   645      -6.636  13.646  -6.359  1.00 34.50      A    C
ATOM   1395  CG  LEU   645      -5.395  12.749  -6.310  1.00 33.63      A    C
ATOM   1396  CD1 LEU   645      -5.668  11.520  -5.449  1.00 29.01      A    C
ATOM   1397  CD2 LEU   645      -5.014  12.337  -7.722  1.00 32.25      A    C
ATOM   1398  C   LEU   645      -7.820  15.699  -7.120  1.00 33.69      A    C
ATOM   1399  O   LEU   645      -8.741  15.482  -7.911  1.00 33.63      A    O
ATOM   1400  N   GLY   646      -7.889  16.615  -6.164  1.00 30.73      A    N
ATOM   1401  CA  GLY   646      -9.099  17.387  -5.980  1.00 29.59      A    C
ATOM   1402  C   GLY   646      -9.210  17.891  -4.559  1.00 29.58      A    C
ATOM   1403  O   GLY   646      -8.271  17.766  -3.767  1.00 29.34      A    O
ATOM   1404  N   ALA   647     -10.362  18.466  -4.238  1.00 27.24      A    N
ATOM   1405  CA  ALA   647     -10.592  19.040  -2.927  1.00 26.99      A    C
ATOM   1406  CB  ALA   647     -10.043  20.453  -2.878  1.00 23.24      A    C
ATOM   1407  C   ALA   647     -12.091  19.044  -2.660  1.00 27.99      A    C
ATOM   1408  O   ALA   647     -12.898  19.142  -3.590  1.00 26.92      A    O
ATOM   1409  N   TYR   648     -12.462  18.923  -1.391  1.00 26.93      A    N
ATOM   1410  CA  TYR   648     -13.865  18.949  -1.011  1.00 28.19      A    C
ATOM   1411  CB  TYR   648     -14.535  17.621  -1.373  1.00 27.10      A    C
ATOM   1412  CG  TYR   648     -13.744  16.397  -0.960  1.00 29.81      A    C
ATOM   1413  CD1 TYR   648     -13.822  15.899   0.336  1.00 28.83      A    C
ATOM   1414  CE1 TYR   648     -13.090  14.788   0.725  1.00 30.03      A    C
ATOM   1415  CD2 TYR   648     -12.912  15.747  -1.862  1.00 29.05      A    C
ATOM   1416  CE2 TYR   648     -12.179  14.635  -1.486  1.00 32.30      A    C
ATOM   1417  CZ  TYR   648     -12.272  14.160  -0.187  1.00 31.60      A    C
ATOM   1418  OH  TYR   648     -11.544  13.057   0.195  1.00 33.44      A    O
ATOM   1419  C   TYR   648     -14.021  19.210   0.477  1.00 30.20      A    C
ATOM   1420  O   TYR   648     -13.126  18.905   1.272  1.00 30.34      A    O
ATOM   1421  N   ALA   649     -15.166  19.768   0.851  1.00 28.57      A    N
ATOM   1422  CA  ALA   649     -15.455  20.020   2.251  1.00 29.86      A    C
ATOM   1423  CB  ALA   649     -16.394  21.210   2.380  1.00 28.48      A    C
ATOM   1424  C   ALA   649     -16.083  18.783   2.884  1.00 32.30      A    C
ATOM   1425  O   ALA   649     -16.920  18.115   2.276  1.00 33.16      A    O
ATOM   1426  N   VAL   650     -15.653  18.469   4.101  1.00 33.30      A    N
ATOM   1427  CA  VAL   650     -16.356  17.509   4.939  1.00 33.70      A    C
ATOM   1428  CB  VAL   650     -15.487  16.269   5.234  1.00 34.68      A    C
ATOM   1429  CG1 VAL   650     -16.177  15.385   6.257  1.00 35.80      A    C
ATOM   1430  CG2 VAL   650     -15.251  15.487   3.954  1.00 33.77      A    C
ATOM   1431  C   VAL   650     -16.683  18.220   6.242  1.00 33.54      A    C
ATOM   1432  O   VAL   650     -15.784  18.580   7.005  1.00 34.44      A    O
ATOM   1433  N   ASP   651     -17.970  18.436   6.484  1.00 32.32      A    N
ATOM   1434  CA  ASP   651     -18.396  19.313   7.560  1.00 33.07      A    C
ATOM   1435  CB  ASP   651     -18.048  18.692   8.914  1.00 34.55      A    C
ATOM   1436  CG  ASP   651     -18.678  17.316   9.101  1.00 37.44      A    C
ATOM   1437  OD1 ASP   651     -19.927  17.218   9.110  1.00 36.19      A    O
ATOM   1438  OD2 ASP   651     -17.925  16.328   9.235  1.00 39.42      A    O
ATOM   1439  C   ASP   651     -17.696  20.658   7.382  1.00 33.97      A    C
ATOM   1440  O   ASP   651     -17.885  21.319   6.360  1.00 34.64      A    O
ATOM   1441  N   ASN   652     -16.884  21.060   8.357  1.00 32.48      A    N
ATOM   1442  CA  ASN   652     -16.196  22.346   8.273  1.00 32.00      A    C
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ATOM   1443  CB  ASN   652     -16.502  23.197   9.511  1.00 32.46      A    C
ATOM   1444  CG  ASN   652     -17.927  23.747   9.500  1.00 35.25      A    C
ATOM   1445  OD1 ASN   652     -18.527  23.930   8.433  1.00 32.30      A    O
ATOM   1446  ND2 ASN   652     -18.474  24.010  10.687  1.00 33.95      A    N
ATOM   1447  C   ASN   652     -14.693  22.176   8.102  1.00 31.47      A    C
ATOM   1448  O   ASN   652     -13.908  23.057   8.451  1.00 30.18      A    O
ATOM   1449  N   THR   653     -14.309  21.033   7.544  1.00 29.69      A    N
ATOM   1450  CA  THR   653     -12.921  20.746   7.225  1.00 28.78      A    C
ATOM   1451  CB  THR   653     -12.510  19.372   7.809  1.00 27.92      A    C
ATOM   1452  OG1 THR   653     -12.589  19.425   9.237  1.00 29.55      A    O
ATOM   1453  CG2 THR   653     -11.095  19.004   7.395  1.00 28.75      A    C
ATOM   1454  C   THR   653     -12.712  20.726   5.707  1.00 28.67      A    C
ATOM   1455  O   THR   653     -13.452  20.061   4.978  1.00 29.58      A    O
ATOM   1456  N   CYS   654     -11.705  21.454   5.236  1.00 28.76      A    N
ATOM   1457  CA  CYS   654     -11.330  21.421   3.827  1.00 27.76      A    C
ATOM   1458  C   CYS   654     -10.320  20.324   3.598  1.00 28.97      A    C
ATOM   1459  O   CYS   654      -9.254  20.321   4.215  1.00 31.24      A    O
ATOM   1460  CB  CYS   654     -10.709  22.750   3.399  1.00 28.74      A    C
ATOM   1461  SG  CYS   654     -10.081  22.798   1.685  1.00 32.18      A    S
ATOM   1462  N   VAL   655     -10.649  19.402   2.700  1.00 28.34      A    N
ATOM   1463  CA  VAL   655      -9.761  18.299   2.378  1.00 28.21      A    C
ATOM   1464  CB  VAL   655     -10.500  16.944   2.459  1.00 28.71      A    C
ATOM   1465  CG1 VAL   655      -9.552  15.821   2.103  1.00 28.30      A    C
ATOM   1466  CG2 VAL   655     -11.058  16.734   3.851  1.00 29.74      A    C
ATOM   1467  C   VAL   655      -9.187  18.455   0.971  1.00 29.61      A    C
ATOM   1468  O   VAL   655      -9.929  18.520  -0.011  1.00 29.60      A    O
ATOM   1469  N   VAL   656      -7.864  18.520   0.877  1.00 29.42      A    N
ATOM   1470  CA  VAL   656      -7.199  18.432  -0.411  1.00 29.18      A    C
ATOM   1471  CB  VAL   656      -6.140  19.550  -0.579  1.00 29.20      A    C
ATOM   1472  CG1 VAL   656      -5.368  19.352  -1.887  1.00 23.76      A    C
ATOM   1473  CG2 VAL   656      -6.827  20.915  -0.575  1.00 25.50      A    C
ATOM   1474  C   VAL   656      -6.526  17.075  -0.522  1.00 30.76      A    C
ATOM   1475  O   VAL   656      -5.797  16.663   0.376  1.00 29.50      A    O
ATOM   1476  N   ARG   657      -6.787  16.378  -1.625  1.00 33.71      A    N
ATOM   1477  CA  ARG   657      -6.201  15.062  -1.863  1.00 35.25      A    C
ATOM   1478  CB  ARG   657      -7.260  14.100  -2.409  1.00 35.66      A    C
ATOM   1479  CG  ARG   657      -8.393  13.791  -1.438  1.00 37.10      A    C
ATOM   1480  CD  ARG   657      -8.058  12.606  -0.541  1.00 37.52      A    C
ATOM   1481  NE  ARG   657      -7.781  11.393  -1.310  1.00 36.14      A    N
ATOM   1482  CZ  ARG   657      -8.701  10.489  -1.637  1.00 39.36      A    C
ATOM   1483  NH1 ARG   657      -8.358   9.416  -2.339  1.00 39.27      A    N
ATOM   1484  NH2 ARG   657      -9.966  10.655  -1.264  1.00 38.06      A    N
ATOM   1485  C   ARG   657      -5.048  15.166  -2.848  1.00 37.95      A    C
ATOM   1486  O   ARG   657      -5.200  15.678  -3.960  1.00 37.69      A    O
ATOM   1487  N   SER   658      -3.888  14.673  -2.433  1.00 41.37      A    N
ATOM   1488  CA  SER   658      -2.685  14.742  -3.254  1.00 44.93      A    C
ATOM   1489  CB  SER   658      -1.502  15.192  -2.397  1.00 45.44      A    C
ATOM   1490  OG  SER   658      -1.412  14.401  -1.223  1.00 50.04      A    O
ATOM   1491  C   SER   658      -2.392  13.378  -3.858  1.00 46.73      A    C
ATOM   1492  O   SER   658      -2.864  12.358  -3.357  1.00 45.37      A    O
ATOM   1493  N   ARG   659      -1.617  13.356  -4.936  1.00 50.91      A    N
ATOM   1494  CA  ARG   659      -1.109  12.095  -5.457  1.00 56.06      A    C
ATOM   1495  CB  ARG   659      -0.700  12.236  -6.920  1.00 60.94      A    C
ATOM   1496  CG  ARG   659       0.009  11.001  -7.473  1.00 66.87      A    C
ATOM   1497  CD  ARG   659      -0.801   9.724  -7.244  1.00 70.25      A    C
ATOM   1498  NE  ARG   659      -0.729   9.250  -5.863  1.00 73.87      A    N
ATOM   1499  CZ  ARG   659      -1.329   8.148  -5.418  1.00 75.97      A    C
ATOM   1500  NH1 ARG   659      -2.049   7.400  -6.247  1.00 76.24      A    N
ATOM   1501  NH2 ARG   659      -1.212   7.792  -4.144  1.00 76.11      A    N
ATOM   1502  C   ARG   659       0.091  11.633  -4.643  1.00 57.33      A    C
ATOM   1503  O   ARG   659      -0.051  10.845  -3.709  1.00 59.34      A    O
ATOM   1504  N   GLU   670       5.942  19.173  -5.234  1.00 59.80      A    N
ATOM   1505  CA  GLU   670       6.378  19.330  -3.843  1.00 59.19      A    C
ATOM   1506  CB  GLU   670       7.326  20.523  -3.727  1.00 60.88      A    C
ATOM   1507  CG  GLU   670       8.652  20.204  -3.026  1.00 65.12      A    C
ATOM   1508  CD  GLU   670       8.570  20.256  -1.508  1.00 68.07      A    C
ATOM   1509  OE1 GLU   670       7.987  21.220  -0.974  1.00 68.75      A    O
ATOM   1510  OE2 GLU   670       9.101  19.341  -0.839  1.00 70.08      A    O
ATOM   1511  C   GLU   670       5.291  19.476  -2.765  1.00 57.96      A    C
ATOM   1512  O   GLU   670       4.489  18.567  -2.544  1.00 56.88      A    O
ATOM   1513  N   ALA   671       5.273  20.628  -2.089  1.00 55.53      A    N
ATOM   1514  CA  ALA   671       4.300  20.874  -1.013  1.00 52.14      A    C
ATOM   1515  CB  ALA   671       4.729  22.075  -0.160  1.00 50.35      A    C
ATOM   1516  C   ALA   671       2.878  21.110  -1.532  1.00 50.50      A    C
ATOM   1517  O   ALA   671       2.685  21.547  -2.660  1.00 49.31      A    O
ATOM   1518  N   VAL   672       1.892  20.811  -0.689  1.00 48.22      A    N
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ATOM   1519  CA  VAL   672       0.485  21.038  -0.993  1.00 46.40      A    C
ATOM   1520  CB  VAL   672      -0.264  19.700  -1.175  1.00 46.60      A    C
ATOM   1521  CG1 VAL   672      -1.765  19.925  -1.123  1.00 45.09      A    C
ATOM   1522  CG2 VAL   672       0.127  19.068  -2.500  1.00 45.84      A    C
ATOM   1523  C   VAL   672      -0.164  21.812   0.140  1.00 44.64      A    C
ATOM   1524  O   VAL   672      -0.012  21.458   1.308  1.00 44.63      A    O
ATOM   1525  N   THR   673      -0.876  22.878  -0.205  1.00 42.97      A    N
ATOM   1526  CA  THR   673      -1.535  23.695   0.803  1.00 42.42      A    C
ATOM   1527  CB  THR   673      -1.014  25.151   0.769  1.00 41.83      A    C
ATOM   1528  OG1 THR   673       0.405  25.156   0.961  1.00 43.57      A    O
ATOM   1529  CG2 THR   673      -1.654  25.968   1.874  1.00 43.03      A    C
ATOM   1530  C   THR   673      -3.052  23.693   0.609  1.00 40.73      A    C
ATOM   1531  O   THR   673      -3.554  23.941  -0.491  1.00 40.35      A    O
ATOM   1532  N   ALA   674      -3.775  23.401   1.684  1.00 37.65      A    N
ATOM   1533  CA  ALA   674      -5.228  23.453   1.651  1.00 36.09      A    C
ATOM   1534  CB  ALA   674      -5.809  22.404   2.589  1.00 32.65      A    C
ATOM   1535  C   ALA   674      -5.700  24.848   2.054  1.00 35.70      A    C
ATOM   1536  O   ALA   674      -5.196  25.433   3.014  1.00 35.06      A    O
ATOM   1537  N   VAL   675      -6.669  25.378   1.313  1.00 33.36      A    N
ATOM   1538  CA  VAL   675      -7.155  26.732   1.554  1.00 31.12      A    C
ATOM   1539  CB  VAL   675      -6.893  27.646   0.344  1.00 32.06      A    C
ATOM   1540  CG1 VAL   675      -7.362  29.057   0.648  1.00 30.35      A    C
ATOM   1541  CG2 VAL   675      -5.423  27.625  -0.015  1.00 31.11      A    C
ATOM   1542  C   VAL   675      -8.652  26.721   1.814  1.00 32.06      A    C
ATOM   1543  O   VAL   675      -9.436  26.300   0.962  1.00 33.53      A    O
ATOM   1544  N   ALA   676      -9.046  27.191   2.991  1.00 31.39      A    N
ATOM   1545  CA  ALA   676     -10.446  27.210   3.369  1.00 31.28      A    C
ATOM   1546  CB  ALA   676     -10.631  26.537   4.731  1.00 30.37      A    C
ATOM   1547  C   ALA   676     -10.965  28.638   3.425  1.00 32.21      A    C
ATOM   1548  O   ALA   676     -10.342  29.513   4.026  1.00 32.29      A    O
ATOM   1549  N   ILE   677     -12.112  28.871   2.799  1.00 30.97      A    N
ATOM   1550  CA  ILE   677     -12.853  30.098   3.046  1.00 31.18      A    C
ATOM   1551  CB  ILE   677     -13.511  30.625   1.755  1.00 30.16      A    C
ATOM   1552  CG2 ILE   677     -14.405  31.810   2.071  1.00 28.01      A    C
ATOM   1553  CG1 ILE   677     -12.424  31.014   0.749  1.00 29.28      A    C
ATOM   1554  CD1 ILE   677     -12.949  31.276  -0.653  1.00 31.33      A    C
ATOM   1555  C   ILE   677     -13.925  29.812   4.094  1.00 32.12      A    C
ATOM   1556  O   ILE   677     -14.759  28.916   3.925  1.00 29.55      A    O
ATOM   1557  N   CYS   678     -13.880  30.576   5.181  1.00 33.00      A    N
ATOM   1558  CA  CYS   678     -14.780  30.386   6.316  1.00 34.13      A    C
ATOM   1559  C   CYS   678     -15.661  31.617   6.478  1.00 33.65      A    C
ATOM   1560  O   CYS   678     -15.180  32.744   6.384  1.00 33.63      A    O
ATOM   1561  CB  CYS   678     -13.966  30.179   7.594  1.00 35.39      A    C
ATOM   1562  SG  CYS   678     -12.794  28.787   7.523  1.00 37.76      A    S
ATOM   1563  N   CYS   679     -16.949  31.412   6.718  1.00 33.05      A    N
ATOM   1564  CA  CYS   679     -17.842  32.547   6.892  1.00 36.46      A    C
ATOM   1565  C   CYS   679     -18.750  32.419   8.113  1.00 39.97      A    C
ATOM   1566  O   CYS   679     -18.970  31.324   8.639  1.00 37.89      A    O
ATOM   1567  CB  CYS   679     -18.712  32.734   5.658  1.00 36.27      A    C
ATOM   1568  SG  CYS   679     -17.854  32.830   4.054  1.00 36.27      A    S
ATOM   1569  N   ARG   680     -19.282  33.556   8.547  1.00 43.34      A    N
ATOM   1570  CA  ARG   680     -20.257  33.591   9.630  1.00 48.28      A    C
ATOM   1571  CB  ARG   680     -19.547  33.641  10.983  1.00 47.10      A    C
ATOM   1572  CG  ARG   680     -18.769  34.919  11.195  1.00 50.25      A    C
ATOM   1573  CD  ARG   680     -18.249  35.029  12.610  1.00 50.93      A    C
ATOM   1574  NE  ARG   680     -17.474  36.250  12.786  1.00 53.09      A    N
ATOM   1575  CZ  ARG   680     -16.716  36.504  13.846  1.00 53.83      A    C
ATOM   1576  NH1 ARG   680     -16.043  37.645  13.919  1.00 54.26      A    N
ATOM   1577  NH2 ARG   680     -16.629  35.617  14.829  1.00 53.05      A    N
ATOM   1578  C   ARG   680     -21.146  34.820   9.479  1.00 49.82      A    C
ATOM   1579  O   ARG   680     -20.924  35.657   8.600  1.00 49.14      A    O
ATOM   1580  N   SER   681     -22.152  34.917  10.339  1.00 53.31      A    N
ATOM   1581  CA  SER   681     -23.029  36.083  10.378  1.00 58.05      A    C
ATOM   1582  CB  SER   681     -24.423  35.673  10.837  1.00 58.41      A    C
ATOM   1583  OG  SER   681     -24.392  35.272  12.195  1.00 60.93      A    O
ATOM   1584  C   SER   681     -22.486  37.127  11.346  1.00 59.77      A    C
ATOM   1585  O   SER   681     -21.714  36.806  12.248  1.00 60.55      A    O
ATOM   1586  N   ARG   682     -22.896  38.375  11.154  1.00 62.76      A    N
ATOM   1587  CA  ARG   682     -22.616  39.429  12.122  1.00 66.03      A    C
ATOM   1588  CB  ARG   682     -21.647  40.448  11.528  1.00 68.33      A    C
ATOM   1589  CG  ARG   682     -20.316  39.874  11.085  1.00 71.05      A    C
ATOM   1590  CD  ARG   682     -19.510  40.941  10.368  1.00 73.61      A    C
ATOM   1591  NE  ARG   682     -20.383  41.861   9.641  1.00 76.11      A    N
ATOM   1592  CZ  ARG   682     -19.966  42.729   8.723  1.00 76.34      A    C
ATOM   1593  NH1 ARG   682     -20.839  43.523   8.117  1.00 74.95      A    N
ATOM   1594  NH2 ARG   682     -18.678  42.801   8.409  1.00 77.06      A    N
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ATOM   1595  C   ARG   682      -23.909  40.138  12.514  1.00 66.85      A    C
ATOM   1596  O   ARG   682      -24.947  39.865  11.873  1.00 67.78      A    O
ATOM   1597  OXT ARG   682      -23.867  40.966  13.448  1.00 68.38      A    O
TER    1598      ARG   682                                               A
ATOM   3134  CB  ALA    -2      -22.599  17.038 -50.667  1.00 30.97      L1   C
ATOM   3135  C   ALA    -2      -20.735  18.001 -49.294  1.00 31.14      L1   C
ATOM   3136  O   ALA    -2      -19.631  17.468 -49.156  1.00 30.83      L1   O
ATOM   3137  N   ALA    -2      -20.438  17.670 -51.719  1.00 31.98      L1   N
ATOM   3138  CA  ALA    -2      -21.422  18.028 -50.655  1.00 32.78      L1   C
ATOM   3139  N   LEU    -1      -21.382  18.582 -48.290  1.00 30.72      L1   N
ATOM   3140  CA  LEU    -1      -20.904  18.443 -46.923  1.00 31.40      L1   C
ATOM   3141  CB  LEU    -1      -20.620  19.810 -46.300  1.00 30.42      L1   C
ATOM   3142  CG  LEU    -1      -20.062  19.746 -44.872  1.00 32.48      L1   C
ATOM   3143  CD1 LEU    -1      -18.735  19.007 -44.874  1.00 32.58      L1   C
ATOM   3144  CD2 LEU    -1      -19.875  21.146 -44.314  1.00 31.47      L1   C
ATOM   3145  C   LEU    -1      -21.939  17.701 -46.090  1.00 32.90      L1   C
ATOM   3146  O   LEU    -1      -23.076  18.154 -45.930  1.00 31.46      L1   O
ATOM   3147  N   GLN     1      -21.543  16.551 -45.565  1.00 32.45      L1   N
ATOM   3148  CA  GLN     1      -22.440  15.769 -44.739  1.00 36.30      L1   C
ATOM   3149  CB  GLN     1      -22.229  14.271 -45.006  1.00 39.00      L1   C
ATOM   3150  CG  GLN     1      -23.162  13.354 -44.229  1.00 46.21      L1   C
ATOM   3151  CD  GLN     1      -23.290  11.970 -44.857  1.00 51.70      L1   C
ATOM   3152  OE1 GLN     1      -24.374  11.383 -44.867  1.00 55.34      L1   O
ATOM   3153  NE2 GLN     1      -22.185  11.444 -45.384  1.00 50.89      L1   N
ATOM   3154  C   GLN     1      -22.148  16.113 -43.282  1.00 36.73      L1   C
ATOM   3155  O   GLN     1      -21.038  15.904 -42.801  1.00 39.33      L1   O
ATOM   3156  N   SER     2      -23.143  16.660 -42.592  1.00 34.09      L1   N
ATOM   3157  CA  SER     2      -22.984  17.045 -41.200  1.00 32.26      L1   C
ATOM   3158  CB  SER     2      -23.663  18.394 -40.951  1.00 33.35      L1   C
ATOM   3159  OG  SER     2      -22.995  19.426 -41.660  1.00 36.24      L1   O
ATOM   3160  C   SER     2      -23.579  15.975 -40.290  1.00 31.17      L1   C
ATOM   3161  O   SER     2      -24.551  15.314 -40.651  1.00 29.93      L1   O
ATOM   3162  N   VAL     3      -22.990  15.806 -39.111  1.00 29.66      L1   N
ATOM   3163  CA  VAL     3      -23.345  14.692 -38.242  1.00 27.57      L1   C
ATOM   3164  CB  VAL     3      -22.173  14.325 -37.304  1.00 26.77      L1   C
ATOM   3165  CG1 VAL     3      -22.604  13.235 -36.328  1.00 20.64      L1   C
ATOM   3166  CG2 VAL     3      -20.970  13.859 -38.135  1.00 24.71      L1   C
ATOM   3167  C   VAL     3      -24.573  14.995 -37.403  1.00 28.78      L1   C
ATOM   3168  O   VAL     3      -25.360  14.102 -37.110  1.00 29.90      L1   O
ATOM   3169  N   LEU     4      -24.736  16.256 -37.013  1.00 28.16      L1   N
ATOM   3170  CA  LEU     4      -25.909  16.661 -36.239  1.00 28.07      L1   C
ATOM   3171  CB  LEU     4      -25.540  17.780 -35.261  1.00 25.10      L1   C
ATOM   3172  CG  LEU     4      -24.396  17.481 -34.289  1.00 27.51      L1   C
ATOM   3173  CD1 LEU     4      -24.060  18.739 -33.483  1.00 27.28      L1   C
ATOM   3174  CD2 LEU     4      -24.793  16.341 -33.367  1.00 27.59      L1   C
ATOM   3175  C   LEU     4      -26.996  17.146 -37.190  1.00 26.91      L1   C
ATOM   3176  O   LEU     4      -26.701  17.620 -38.282  1.00 25.87      L1   O
ATOM   3177  N   THR     5      -28.251  17.031 -36.777  1.00 27.45      L1   N
ATOM   3178  CA  THR     5      -29.365  17.405 -37.650  1.00 29.69      L1   C
ATOM   3179  CB  THR     5      -30.419  16.281 -37.709  1.00 30.65      L1   C
ATOM   3180  OG1 THR     5      -29.813  15.091 -38.227  1.00 32.80      L1   O
ATOM   3181  CG2 THR     5      -31.584  16.682 -38.615  1.00 30.25      L1   C
ATOM   3182  C   THR     5      -30.062  18.693 -37.216  1.00 29.39      L1   C
ATOM   3183  O   THR     5      -30.597  18.773 -36.113  1.00 28.74      L1   O
ATOM   3184  N   GLN     6      -30.048  19.696 -38.093  1.00 27.91      L1   N
ATOM   3185  CA  GLN     6      -30.806  20.929 -37.885  1.00 28.70      L1   C
ATOM   3186  CB  GLN     6      -29.889  22.159 -37.866  1.00 26.66      L1   C
ATOM   3187  CG  GLN     6      -28.913  22.258 -36.721  1.00 24.05      L1   C
ATOM   3188  CD  GLN     6      -28.010  23.468 -36.872  1.00 25.09      L1   C
ATOM   3189  OE1 GLN     6      -26.843  23.341 -37.243  1.00 27.47      L1   O
ATOM   3190  NE2 GLN     6      -28.548  24.652 -36.597  1.00 22.42      L1   N
ATOM   3191  C   GLN     6      -31.770  21.109 -39.044  1.00 28.56      L1   C
ATOM   3192  O   GLN     6      -31.475  20.719 -40.170  1.00 27.37      L1   O
ATOM   3193  N   PRO     7      -32.921  21.745 -38.789  1.00 30.02      L1   N
ATOM   3194  CD  PRO     7      -33.416  22.218 -37.483  1.00 30.00      L1   C
ATOM   3195  CA  PRO     7      -33.756  22.206 -39.904  1.00 29.84      L1   C
ATOM   3196  CB  PRO     7      -34.944  22.876 -39.215  1.00 30.46      L1   C
ATOM   3197  CG  PRO     7      -34.421  23.278 -37.859  1.00 31.08      L1   C
ATOM   3198  C   PRO     7      -32.961  23.186 -40.761  1.00 30.79      L1   C
ATOM   3199  O   PRO     7      -32.206  24.007 -40.242  1.00 30.95      L1   O
ATOM   3200  N   PRO     8      -33.113  23.100 -42.087  1.00 29.99      L1   N
ATOM   3201  CD  PRO     8      -33.901  22.060 -42.767  1.00 29.60      L1   C
ATOM   3202  CA  PRO     8      -32.380  23.946 -43.036  1.00 30.24      L1   C
ATOM   3203  CB  PRO     8      -32.755  23.369 -44.404  1.00 30.10      L1   C
ATOM   3204  CG  PRO     8      -33.254  21.990 -44.113  1.00 31.61      L1   C
ATOM   3205  C   PRO     8      -32.747  25.428 -42.928  1.00 31.73      L1   C
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ATOM   3206  O   PRO     8     -31.943  26.308 -43.258  1.00 31.46      L1   O
ATOM   3207  N   SER     9     -33.960  25.707 -42.467  1.00 30.72      L1   N
ATOM   3208  CA  SER     9     -34.446  27.080 -42.450  1.00 32.53      L1   C
ATOM   3209  CB  SER     9     -35.043  27.456 -43.808  1.00 35.18      L1   C
ATOM   3210  OG  SER     9     -36.252  26.743 -44.038  1.00 39.78      L1   O
ATOM   3211  C   SER     9     -35.494  27.298 -41.382  1.00 32.20      L1   C
ATOM   3212  O   SER     9     -36.151  26.359 -40.933  1.00 32.14      L1   O
ATOM   3213  N   ALA    10     -35.637  28.556 -40.982  1.00 32.09      L1   N
ATOM   3214  CA  ALA    10     -36.668  28.970 -40.049  1.00 30.01      L1   C
ATOM   3215  CB  ALA    10     -36.196  28.744 -38.611  1.00 29.34      L1   C
ATOM   3216  C   ALA    10     -36.943  30.453 -40.294  1.00 31.41      L1   C
ATOM   3217  O   ALA    10     -36.162  31.139 -40.958  1.00 31.03      L1   O
ATOM   3218  N   SER    11     -38.058  30.946 -39.770  1.00 30.95      L1   N
ATOM   3219  CA  SER    11     -38.366  32.364 -39.876  1.00 33.18      L1   C
ATOM   3220  CB  SER    11     -39.055  32.667 -41.205  1.00 31.83      L1   C
ATOM   3221  OG  SER    11     -40.290  31.985 -41.292  1.00 34.06      L1   O
ATOM   3222  C   SER    11     -39.255  32.810 -38.731  1.00 33.60      L1   C
ATOM   3223  O   SER    11     -40.016  32.023 -38.175  1.00 34.21      L1   O
ATOM   3224  N   GLY    12     -39.143  34.084 -38.379  1.00 33.24      L1   N
ATOM   3225  CA  GLY    12     -39.983  34.638 -37.344  1.00 34.06      L1   C
ATOM   3226  C   GLY    12     -40.221  36.111 -37.594  1.00 34.78      L1   C
ATOM   3227  O   GLY    12     -39.510  36.738 -38.377  1.00 35.73      L1   O
ATOM   3228  N   THR    13     -41.224  36.664 -36.926  1.00 35.43      L1   N
ATOM   3229  CA  THR    13     -41.545  38.077 -37.055  1.00 35.76      L1   C
ATOM   3230  CB  THR    13     -43.057  38.299 -36.915  1.00 37.03      L1   C
ATOM   3231  OG1 THR    13     -43.732  37.621 -37.980  1.00 40.63      L1   O
ATOM   3232  CG2 THR    13     -43.389  39.781 -36.959  1.00 38.80      L1   C
ATOM   3233  C   THR    13     -40.834  38.834 -35.949  1.00 34.79      L1   C
ATOM   3234  O   THR    13     -40.768  38.358 -34.817  1.00 33.06      L1   O
ATOM   3235  N   PRO    14     -40.295  40.024 -36.261  1.00 33.81      L1   N
ATOM   3236  CD  PRO    14     -40.243  40.682 -37.579  1.00 33.59      L1   C
ATOM   3237  CA  PRO    14     -39.678  40.845 -35.213  1.00 35.17      L1   C
ATOM   3238  CB  PRO    14     -39.510  42.210 -35.881  1.00 35.17      L1   C
ATOM   3239  CG  PRO    14     -39.338  41.880 -37.339  1.00 32.62      L1   C
ATOM   3240  C   PRO    14     -40.562  40.904 -33.970  1.00 37.04      L1   C
ATOM   3241  O   PRO    14     -41.788  40.939 -34.076  1.00 36.13      L1   O
ATOM   3242  N   GLY    15     -39.933  40.880 -32.798  1.00 37.53      L1   N
ATOM   3243  CA  GLY    15     -40.684  40.885 -31.554  1.00 37.21      L1   C
ATOM   3244  C   GLY    15     -41.091  39.514 -31.044  1.00 37.71      L1   C
ATOM   3245  O   GLY    15     -41.394  39.364 -29.864  1.00 39.10      L1   O
ATOM   3246  N   GLN    16     -41.101  38.515 -31.918  1.00 38.28      L1   N
ATOM   3247  CA  GLN    16     -41.542  37.184 -31.514  1.00 39.95      L1   C
ATOM   3248  CB  GLN    16     -42.263  36.488 -32.673  1.00 42.22      L1   C
ATOM   3249  CG  GLN    16     -43.514  37.217 -33.166  1.00 49.57      L1   C
ATOM   3250  CD  GLN    16     -44.608  37.318 -32.108  1.00 53.19      L1   C
ATOM   3251  OE1 GLN    16     -45.367  36.371 -31.889  1.00 55.70      L1   O
ATOM   3252  NE2 GLN    16     -44.693  38.472 -31.449  1.00 53.13      L1   N
ATOM   3253  C   GLN    16     -40.393  36.301 -31.017  1.00 39.30      L1   C
ATOM   3254  O   GLN    16     -39.232  36.715 -30.998  1.00 36.61      L1   O
ATOM   3255  N   ARG    17     -40.743  35.085 -30.608  1.00 39.04      L1   N
ATOM   3256  CA  ARG    17     -39.782  34.093 -30.142  1.00 40.32      L1   C
ATOM   3257  CB  ARG    17     -40.225  33.551 -28.779  1.00 40.01      L1   C
ATOM   3258  CG  ARG    17     -39.384  32.406 -28.239  1.00 43.34      L1   C
ATOM   3259  CD  ARG    17     -39.923  31.930 -26.898  1.00 41.71      L1   C
ATOM   3260  NE  ARG    17     -39.086  30.906 -26.277  1.00 43.61      L1   N
ATOM   3261  CZ  ARG    17     -39.155  29.610 -26.565  1.00 43.68      L1   C
ATOM   3262  NH1 ARG    17     -40.019  29.178 -27.471  1.00 44.38      L1   N
ATOM   3263  NH2 ARG    17     -38.376  28.740 -25.933  1.00 43.92      L1   N
ATOM   3264  C   ARG    17     -39.697  32.949 -31.148  1.00 39.25      L1   C
ATOM   3265  O   ARG    17     -40.722  32.397 -31.548  1.00 40.22      L1   O
ATOM   3266  N   VAL    18     -38.480  32.601 -31.562  1.00 38.21      L1   N
ATOM   3267  CA  VAL    18     -38.269  31.428 -32.410  1.00 36.63      L1   C
ATOM   3268  CB  VAL    18     -37.688  31.814 -33.790  1.00 37.42      L1   C
ATOM   3269  CG1 VAL    18     -38.642  32.740 -34.510  1.00 37.93      L1   C
ATOM   3270  CG2 VAL    18     -36.331  32.479 -33.624  1.00 37.49      L1   C
ATOM   3271  C   VAL    18     -37.320  30.430 -31.755  1.00 36.06      L1   C
ATOM   3272  O   VAL    18     -36.493  30.800 -30.918  1.00 37.18      L1   O
ATOM   3273  N   THR    19     -37.447  29.161 -32.127  1.00 34.19      L1   N
ATOM   3274  CA  THR    19     -36.532  28.148 -31.635  1.00 33.85      L1   C
ATOM   3275  CB  THR    19     -37.231  27.152 -30.699  1.00 34.73      L1   C
ATOM   3276  OG1 THR    19     -38.256  26.459 -31.422  1.00 36.14      L1   O
ATOM   3277  CG2 THR    19     -37.839  27.875 -29.510  1.00 36.05      L1   C
ATOM   3278  C   THR    19     -35.932  27.367 -32.788  1.00 33.77      L1   C
ATOM   3279  O   THR    19     -36.581  27.136 -33.804  1.00 34.31      L1   O
ATOM   3280  N   ILE    20     -34.680  26.964 -32.627  1.00 32.67      L1   N
ATOM   3281  CA  ILE    20     -34.013  26.136 -33.619  1.00 30.93      L1   C
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ATOM	3282	CB	ILE	20	-32.856	26.902	-34.267	1.00	30.89	L1	C
ATOM	3283	CG2	ILE	20	-32.090	25.991	-35.202	1.00	32.28	L1	C
ATOM	3284	CG1	ILE	20	-33.410	28.121	-35.010	1.00	31.02	L1	C
ATOM	3285	CD1	ILE	20	-32.348	29.096	-35.461	1.00	29.96	L1	C
ATOM	3286	C	ILE	20	-33.479	24.902	-32.912	1.00	29.12	L1	C
ATOM	3287	O	ILE	20	-32.825	25.007	-31.877	1.00	27.11	L1	O
ATOM	3288	N	SER	21	-33.772	23.731	-33.460	1.00	28.13	L1	N
ATOM	3289	CA	SER	21	-33.418	22.493	-32.791	1.00	27.04	L1	C
ATOM	3290	CB	SER	21	-34.577	21.503	-32.857	1.00	28.51	L1	C
ATOM	3291	OG	SER	21	-34.646	20.916	-34.144	1.00	36.31	L1	O
ATOM	3292	C	SER	21	-32.183	21.856	-33.404	1.00	27.38	L1	C
ATOM	3293	O	SER	21	-31.850	22.087	-34.576	1.00	26.13	L1	O
ATOM	3294	N	CYS	22	-31.514	21.040	-32.599	1.00	26.91	L1	N
ATOM	3295	CA	CYS	22	-30.300	20.349	-33.012	1.00	28.14	L1	C
ATOM	3296	C	CYS	22	-30.378	18.953	-32.407	1.00	27.39	L1	C
ATOM	3297	O	CYS	22	-30.482	18.805	-31.191	1.00	26.72	L1	O
ATOM	3298	CB	CYS	22	-29.078	21.109	-32.470	1.00	28.78	L1	C
ATOM	3299	SG	CYS	22	-27.428	20.365	-32.716	1.00	33.98	L1	S
ATOM	3300	N	SER	23	-30.349	17.926	-33.246	1.00	26.59	L1	N
ATOM	3301	CA	SER	23	-30.434	16.574	-32.722	1.00	29.26	L1	C
ATOM	3302	CB	SER	23	-31.691	15.875	-33.244	1.00	30.37	L1	C
ATOM	3303	OG	SER	23	-31.715	15.896	-34.653	1.00	40.04	L1	O
ATOM	3304	C	SER	23	-29.202	15.750	-33.053	1.00	27.58	L1	C
ATOM	3305	O	SER	23	-28.689	15.783	-34.171	1.00	25.14	L1	O
ATOM	3306	N	GLY	24	-28.734	15.017	-32.049	1.00	27.83	L1	N
ATOM	3307	CA	GLY	24	-27.548	14.204	-32.195	1.00	28.19	L1	C
ATOM	3308	C	GLY	24	-27.832	12.832	-31.632	1.00	29.00	L1	C
ATOM	3309	O	GLY	24	-28.902	12.268	-31.867	1.00	29.40	L1	O
ATOM	3310	N	SER	25	-26.886	12.291	-30.878	1.00	26.84	L1	N
ATOM	3311	CA	SER	25	-27.045	10.942	-30.372	1.00	27.01	L1	C
ATOM	3312	CB	SER	25	-26.747	9.938	-31.478	1.00	26.36	L1	C
ATOM	3313	OG	SER	25	-25.400	10.061	-31.879	1.00	26.75	L1	O
ATOM	3314	C	SER	25	-26.113	10.705	-29.209	1.00	26.30	L1	C
ATOM	3315	O	SER	25	-25.447	11.629	-28.746	1.00	27.16	L1	O
ATOM	3316	N	SER	26	-26.066	9.462	-28.742	1.00	25.37	L1	N
ATOM	3317	CA	SER	26	-25.345	9.137	-27.522	1.00	25.52	L1	C
ATOM	3318	CB	SER	26	-25.588	7.678	-27.143	1.00	26.57	L1	C
ATOM	3319	OG	SER	26	-25.038	6.804	-28.119	1.00	32.72	L1	O
ATOM	3320	C	SER	26	-23.847	9.390	-27.673	1.00	25.31	L1	C
ATOM	3321	O	SER	26	-23.153	9.613	-26.688	1.00	26.31	L1	O
ATOM	3322	N	SER	27	-23.347	9.362	-28.904	1.00	25.12	L1	N
ATOM	3323	CA	SER	27	-21.916	9.549	-29.129	1.00	23.38	L1	C
ATOM	3324	CB	SER	27	-21.495	8.948	-30.472	1.00	22.37	L1	C
ATOM	3325	OG	SER	27	-21.943	9.746	-31.551	1.00	24.28	L1	O
ATOM	3326	C	SER	27	-21.521	11.023	-29.085	1.00	24.26	L1	C
ATOM	3327	O	SER	27	-20.342	11.342	-28.959	1.00	23.93	L1	O
ATOM	3328	N	ASN	28	-22.497	11.926	-29.195	1.00	23.53	L1	N
ATOM	3329	CA	ASN	28	-22.197	13.345	-29.020	1.00	23.52	L1	C
ATOM	3330	CB	ASN	28	-22.302	14.111	-30.361	1.00	21.17	L1	C
ATOM	3331	CG	ASN	28	-23.467	13.656	-31.228	1.00	23.34	L1	C
ATOM	3332	OD1	ASN	28	-24.629	13.895	-30.908	1.00	23.92	L1	O
ATOM	3333	ND2	ASN	28	-23.154	13.009	-32.345	1.00	21.27	L1	N
ATOM	3334	C	ASN	28	-23.028	14.033	-27.937	1.00	23.95	L1	C
ATOM	3335	O	ASN	28	-22.578	14.156	-26.799	1.00	25.36	L1	O
ATOM	3336	N	ILE	29	-24.230	14.486	-28.267	1.00	24.55	L1	N
ATOM	3337	CA	ILE	29	-24.995	15.263	-27.300	1.00	24.06	L1	C
ATOM	3338	CB	ILE	29	-26.300	15.799	-27.929	1.00	26.12	L1	C
ATOM	3339	CG2	ILE	29	-27.128	16.554	-26.883	1.00	21.66	L1	C
ATOM	3340	CG1	ILE	29	-25.951	16.716	-29.103	1.00	21.86	L1	C
ATOM	3341	CD1	ILE	29	-27.146	17.356	-29.766	1.00	26.70	L1	C
ATOM	3342	C	ILE	29	-25.313	14.456	-26.040	1.00	25.43	L1	C
ATOM	3343	O	ILE	29	-25.448	15.021	-24.956	1.00	24.34	L1	O
ATOM	3344	N	GLY	30	-25.401	13.134	-26.170	1.00	25.71	L1	N
ATOM	3345	CA	GLY	30	-25.640	12.301	-25.000	1.00	25.17	L1	C
ATOM	3346	C	GLY	30	-24.619	12.467	-23.875	1.00	28.74	L1	C
ATOM	3347	O	GLY	30	-24.952	12.284	-22.702	1.00	28.98	L1	O
ATOM	3348	N	SER	31	-23.376	12.806	-24.217	1.00	26.07	L1	N
ATOM	3349	CA	SER	31	-22.342	13.023	-23.207	1.00	25.52	L1	C
ATOM	3350	CB	SER	31	-21.104	12.176	-23.511	1.00	25.93	L1	C
ATOM	3351	OG	SER	31	-21.357	10.799	-23.314	1.00	27.84	L1	O
ATOM	3352	C	SER	31	-21.905	14.484	-23.092	1.00	26.39	L1	C
ATOM	3353	O	SER	31	-21.490	14.925	-22.022	1.00	25.97	L1	O
ATOM	3354	N	ASN	32	-21.991	15.227	-24.192	1.00	25.12	L1	N
ATOM	3355	CA	ASN	32	-21.240	16.471	-24.316	1.00	25.92	L1	C
ATOM	3356	CB	ASN	32	-20.279	16.342	-25.494	1.00	22.78	L1	C
ATOM	3357	CG	ASN	32	-19.380	15.133	-25.362	1.00	25.36	L1	C

ATOM	3358	OD1	ASN	32	-18.840	14.864	-24.284	1.00	24.71	L1	O
ATOM	3359	ND2	ASN	32	-19.220	14.387	-26.451	1.00	23.92	L1	N
ATOM	3360	C	ASN	32	-22.087	17.742	-24.445	1.00	26.14	L1	C
ATOM	3361	O	ASN	32	-23.258	17.697	-24.833	1.00	26.61	L1	O
ATOM	3362	N	THR	33	-21.477	18.873	-24.106	1.00	25.64	L1	N
ATOM	3363	CA	THR	33	-22.153	20.167	-24.125	1.00	25.44	L1	C
ATOM	3364	CB	THR	33	-21.272	21.272	-23.484	1.00	26.18	L1	C
ATOM	3365	OG1	THR	33	-20.054	21.407	-24.231	1.00	27.99	L1	O
ATOM	3366	CG2	THR	33	-20.937	20.925	-22.040	1.00	25.80	L1	C
ATOM	3367	C	THR	33	-22.478	20.589	-25.553	1.00	24.65	L1	C
ATOM	3368	O	THR	33	-21.778	20.220	-26.492	1.00	23.55	L1	O
ATOM	3369	N	VAL	34	-23.542	21.367	-25.713	1.00	22.97	L1	N
ATOM	3370	CA	VAL	34	-23.901	21.889	-27.024	1.00	20.22	L1	C
ATOM	3371	CB	VAL	34	-25.422	21.766	-27.273	1.00	20.79	L1	C
ATOM	3372	CG1	VAL	34	-25.765	22.309	-28.640	1.00	17.13	L1	C
ATOM	3373	CG2	VAL	34	-25.849	20.303	-27.162	1.00	18.86	L1	C
ATOM	3374	C	VAL	34	-23.497	23.349	-27.126	1.00	20.72	L1	C
ATOM	3375	O	VAL	34	-23.588	24.092	-26.153	1.00	20.35	L1	O
ATOM	3376	N	ASN	35	-23.062	23.760	-28.311	1.00	21.11	L1	N
ATOM	3377	CA	ASN	35	-22.553	25.104	-28.515	1.00	20.37	L1	C
ATOM	3378	CB	ASN	35	-21.028	25.068	-28.673	1.00	20.30	L1	C
ATOM	3379	CG	ASN	35	-20.345	24.360	-27.513	1.00	20.02	L1	C
ATOM	3380	OD1	ASN	35	-20.061	24.968	-26.477	1.00	20.69	L1	O
ATOM	3381	ND2	ASN	35	-20.092	23.069	-27.675	1.00	18.74	L1	N
ATOM	3382	C	ASN	35	-23.190	25.664	-29.765	1.00	23.63	L1	C
ATOM	3383	O	ASN	35	-23.363	24.943	-30.750	1.00	25.23	L1	O
ATOM	3384	N	TRP	36	-23.535	26.950	-29.728	1.00	24.33	L1	N
ATOM	3385	CA	TRP	36	-24.249	27.581	-30.828	1.00	23.53	L1	C
ATOM	3386	CB	TRP	36	-25.637	28.023	-30.361	1.00	24.65	L1	C
ATOM	3387	CG	TRP	36	-26.565	26.884	-30.058	1.00	23.30	L1	C
ATOM	3388	CD2	TRP	36	-27.466	26.254	-30.973	1.00	21.45	L1	C
ATOM	3389	CE2	TRP	36	-28.163	25.261	-30.257	1.00	22.89	L1	C
ATOM	3390	CE3	TRP	36	-27.754	26.436	-32.330	1.00	24.38	L1	C
ATOM	3391	CD1	TRP	36	-26.744	26.264	-28.853	1.00	22.85	L1	C
ATOM	3392	NE1	TRP	36	-27.706	25.284	-28.965	1.00	22.07	L1	N
ATOM	3393	CZ2	TRP	36	-29.131	24.452	-30.853	1.00	23.37	L1	C
ATOM	3394	CZ3	TRP	36	-28.718	25.630	-32.920	1.00	24.94	L1	C
ATOM	3395	CH2	TRP	36	-29.393	24.652	-32.181	1.00	23.94	L1	C
ATOM	3396	C	TRP	36	-23.492	28.783	-31.393	1.00	24.76	L1	C
ATOM	3397	O	TRP	36	-22.862	29.539	-30.651	1.00	24.29	L1	O
ATOM	3398	N	TYR	37	-23.580	28.950	-32.710	1.00	23.23	L1	N
ATOM	3399	CA	TYR	37	-22.890	30.017	-33.419	1.00	25.41	L1	C
ATOM	3400	CB	TYR	37	-21.710	29.444	-34.223	1.00	23.97	L1	C
ATOM	3401	CG	TYR	37	-20.751	28.674	-33.348	1.00	24.70	L1	C
ATOM	3402	CD1	TYR	37	-21.020	27.357	-32.993	1.00	23.14	L1	C
ATOM	3403	CE1	TYR	37	-20.220	26.679	-32.095	1.00	24.44	L1	C
ATOM	3404	CD2	TYR	37	-19.639	29.291	-32.789	1.00	21.55	L1	C
ATOM	3405	CE2	TYR	37	-18.826	28.618	-31.887	1.00	23.33	L1	C
ATOM	3406	CZ	TYR	37	-19.127	27.312	-31.541	1.00	23.69	L1	C
ATOM	3407	OH	TYR	37	-18.359	26.638	-30.618	1.00	20.32	L1	O
ATOM	3408	C	TYR	37	-23.845	30.737	-34.360	1.00	26.66	L1	C
ATOM	3409	O	TYR	37	-24.741	30.131	-34.946	1.00	27.88	L1	O
ATOM	3410	N	GLN	38	-23.640	32.040	-34.492	1.00	26.80	L1	N
ATOM	3411	CA	GLN	38	-24.395	32.858	-35.423	1.00	28.14	L1	C
ATOM	3412	CB	GLN	38	-24.986	34.053	-34.674	1.00	29.55	L1	C
ATOM	3413	CG	GLN	38	-25.560	35.128	-35.559	1.00	29.51	L1	C
ATOM	3414	CD	GLN	38	-25.992	36.335	-34.764	1.00	32.32	L1	C
ATOM	3415	OE1	GLN	38	-25.167	37.159	-34.364	1.00	32.45	L1	O
ATOM	3416	NE2	GLN	38	-27.292	36.449	-34.526	1.00	31.43	L1	N
ATOM	3417	C	GLN	38	-23.428	33.339	-36.497	1.00	27.48	L1	C
ATOM	3418	O	GLN	38	-22.313	33.746	-36.181	1.00	27.04	L1	O
ATOM	3419	N	GLN	39	-23.839	33.284	-37.760	1.00	27.49	L1	N
ATOM	3420	CA	GLN	39	-22.977	33.751	-38.838	1.00	28.01	L1	C
ATOM	3421	CB	GLN	39	-22.396	32.569	-39.630	1.00	28.56	L1	C
ATOM	3422	CG	GLN	39	-21.371	33.003	-40.684	1.00	25.91	L1	C
ATOM	3423	CD	GLN	39	-20.727	31.842	-41.412	1.00	26.00	L1	C
ATOM	3424	OE1	GLN	39	-21.345	30.791	-41.602	1.00	25.62	L1	O
ATOM	3425	NE2	GLN	39	-19.476	32.028	-41.832	1.00	22.22	L1	N
ATOM	3426	C	GLN	39	-23.697	34.682	-39.800	1.00	29.88	L1	C
ATOM	3427	O	GLN	39	-24.718	34.318	-40.387	1.00	28.22	L1	O
ATOM	3428	N	LEU	40	-23.143	35.881	-39.957	1.00	32.04	L1	N
ATOM	3429	CA	LEU	40	-23.601	36.838	-40.957	1.00	34.34	L1	C
ATOM	3430	CB	LEU	40	-23.473	38.267	-40.418	1.00	33.92	L1	C
ATOM	3431	CG	LEU	40	-24.221	38.579	-39.117	1.00	37.04	L1	C
ATOM	3432	CD1	LEU	40	-23.977	40.031	-38.729	1.00	38.63	L1	C
ATOM	3433	CD2	LEU	40	-25.711	38.328	-39.297	1.00	37.35	L1	C

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ATOM   3434  C   LEU   40     -22.765  36.686 -42.226  1.00 36.40      L1   C
ATOM   3435  O   LEU   40     -21.658  36.142 -42.194  1.00 35.68      L1   O
ATOM   3436  N   PRO   41     -23.285  37.167 -43.366  1.00 38.85      L1   N
ATOM   3437  CD  PRO   41     -24.597  37.818 -43.534  1.00 38.64      L1   C
ATOM   3438  CA  PRO   41     -22.597  36.981 -44.651  1.00 38.77      L1   C
ATOM   3439  CB  PRO   41     -23.531  37.655 -45.659  1.00 40.02      L1   C
ATOM   3440  CG  PRO   41     -24.884  37.619 -45.000  1.00 38.99      L1   C
ATOM   3441  C   PRO   41     -21.197  37.590 -44.668  1.00 38.74      L1   C
ATOM   3442  O   PRO   41     -20.978  38.690 -44.162  1.00 38.74      L1   O
ATOM   3443  N   GLY   42     -20.248  36.856 -45.236  1.00 38.68      L1   N
ATOM   3444  CA  GLY   42     -18.901  37.375 -45.382  1.00 40.14      L1   C
ATOM   3445  C   GLY   42     -18.135  37.520 -44.080  1.00 40.18      L1   C
ATOM   3446  O   GLY   42     -17.077  38.147 -44.042  1.00 41.17      L1   O
ATOM   3447  N   THR   43     -18.655  36.941 -43.007  1.00 37.47      L1   N
ATOM   3448  CA  THR   43     -17.972  37.040 -41.732  1.00 36.70      L1   C
ATOM   3449  CB  THR   43     -18.660  38.090 -40.841  1.00 38.93      L1   C
ATOM   3450  OG1 THR   43     -17.927  38.239 -39.619  1.00 44.97      L1   O
ATOM   3451  CG2 THR   43     -20.073  37.673 -40.529  1.00 40.28      L1   C
ATOM   3452  C   THR   43     -17.898  35.692 -41.007  1.00 33.58      L1   C
ATOM   3453  O   THR   43     -18.693  34.787 -41.268  1.00 32.26      L1   O
ATOM   3454  N   ALA   44     -16.924  35.556 -40.114  1.00 29.37      L1   N
ATOM   3455  CA  ALA   44     -16.748  34.320 -39.368  1.00 29.17      L1   C
ATOM   3456  CB  ALA   44     -15.469  34.376 -38.542  1.00 24.98      L1   C
ATOM   3457  C   ALA   44     -17.944  34.104 -38.453  1.00 28.88      L1   C
ATOM   3458  O   ALA   44     -18.588  35.057 -38.022  1.00 28.39      L1   O
ATOM   3459  N   PRO   45     -18.249  32.839 -38.139  1.00 28.41      L1   N
ATOM   3460  CD  PRO   45     -17.620  31.619 -38.670  1.00 26.65      L1   C
ATOM   3461  CA  PRO   45     -19.282  32.538 -37.143  1.00 27.06      L1   C
ATOM   3462  CB  PRO   45     -19.254  31.013 -37.036  1.00 25.83      L1   C
ATOM   3463  CG  PRO   45     -18.623  30.552 -38.310  1.00 27.18      L1   C
ATOM   3464  C   PRO   45     -18.922  33.199 -35.818  1.00 28.11      L1   C
ATOM   3465  O   PRO   45     -17.745  33.412 -35.520  1.00 29.04      L1   O
ATOM   3466  N   LYS   46     -19.934  33.527 -35.029  1.00 27.77      L1   N
ATOM   3467  CA  LYS   46     -19.718  34.126 -33.720  1.00 29.30      L1   C
ATOM   3468  CB  LYS   46     -20.324  35.529 -33.693  1.00 30.36      L1   C
ATOM   3469  CG  LYS   46     -20.595  36.080 -32.309  1.00 35.69      L1   C
ATOM   3470  CD  LYS   46     -21.447  37.342 -32.398  1.00 38.46      L1   C
ATOM   3471  CE  LYS   46     -21.611  38.008 -31.040  1.00 41.17      L1   C
ATOM   3472  NZ  LYS   46     -22.445  39.245 -31.131  1.00 43.66      L1   N
ATOM   3473  C   LYS   46     -20.366  33.249 -32.654  1.00 27.70      L1   C
ATOM   3474  O   LYS   46     -21.527  32.869 -32.784  1.00 28.72      L1   O
ATOM   3475  N   LEU   47     -19.611  32.927 -31.609  1.00 25.77      L1   N
ATOM   3476  CA  LEU   47     -20.108  32.072 -30.530  1.00 26.00      L1   C
ATOM   3477  CB  LEU   47     -18.971  31.738 -29.556  1.00 22.43      L1   C
ATOM   3478  CG  LEU   47     -19.368  30.964 -28.294  1.00 24.81      L1   C
ATOM   3479  CD1 LEU   47     -19.762  29.550 -28.680  1.00 20.56      L1   C
ATOM   3480  CD2 LEU   47     -18.207  30.937 -27.298  1.00 22.63      L1   C
ATOM   3481  C   LEU   47     -21.233  32.770 -29.772  1.00 26.08      L1   C
ATOM   3482  O   LEU   47     -21.056  33.895 -29.312  1.00 25.15      L1   O
ATOM   3483  N   LEU   48     -22.376  32.097 -29.642  1.00 26.83      L1   N
ATOM   3484  CA  LEU   48     -23.540  32.643 -28.934  1.00 28.45      L1   C
ATOM   3485  CB  LEU   48     -24.814  32.464 -29.763  1.00 30.33      L1   C
ATOM   3486  CG  LEU   48     -25.021  33.275 -31.042  1.00 35.48      L1   C
ATOM   3487  CD1 LEU   48     -26.383  32.926 -31.613  1.00 35.81      L1   C
ATOM   3488  CD2 LEU   48     -24.938  34.774 -30.756  1.00 34.18      L1   C
ATOM   3489  C   LEU   48     -23.757  31.955 -27.589  1.00 27.72      L1   C
ATOM   3490  O   LEU   48     -24.070  32.598 -26.588  1.00 27.24      L1   O
ATOM   3491  N   ILE   49     -23.608  30.636 -27.590  1.00 27.22      L1   N
ATOM   3492  CA  ILE   49     -23.803  29.810 -26.428  1.00 23.99      L1   C
ATOM   3493  CB  ILE   49     -25.270  29.109 -26.590  1.00 23.11      L1   C
ATOM   3494  CG2 ILE   49     -25.514  28.171 -25.427  1.00 23.24      L1   C
ATOM   3495  CG1 ILE   49     -26.387  30.153 -26.717  1.00 25.09      L1   C
ATOM   3496  CD1 ILE   49     -26.846  30.741 -25.395  1.00 26.07      L1   C
ATOM   3497  C   ILE   49     -22.830  28.738 -26.293  1.00 24.62      L1   C
ATOM   3498  O   ILE   49     -22.472  28.077 -27.272  1.00 25.83      L1   O
ATOM   3499  N   TYR   50     -22.314  28.558 -25.085  1.00 25.53      L1   N
ATOM   3500  CA  TYR   50     -21.466  27.403 -24.811  1.00 26.67      L1   C
ATOM   3501  CB  TYR   50     -20.003  27.833 -24.628  1.00 25.70      L1   C
ATOM   3502  CG  TYR   50     -19.748  28.688 -23.408  1.00 26.62      L1   C
ATOM   3503  CD1 TYR   50     -19.874  30.068 -23.466  1.00 26.79      L1   C
ATOM   3504  CE1 TYR   50     -19.645  30.858 -22.343  1.00 28.90      L1   C
ATOM   3505  CD2 TYR   50     -19.383  28.110 -22.196  1.00 26.72      L1   C
ATOM   3506  CE2 TYR   50     -19.153  28.882 -21.071  1.00 28.12      L1   C
ATOM   3507  CZ  TYR   50     -19.288  30.258 -21.148  1.00 30.76      L1   C
ATOM   3508  OH  TYR   50     -19.085  31.030 -20.026  1.00 29.96      L1   O
ATOM   3509  C   TYR   50     -21.976  26.697 -23.563  1.00 25.84      L1   C
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ATOM	3510	O	TYR	50	-22.784	27.251	-22.821	1.00	24.72	L1	O
ATOM	3511	N	SER	51	-21.518	25.470	-23.343	1.00	26.19	L1	N
ATOM	3512	CA	SER	51	-21.971	24.686	-22.198	1.00	26.90	L1	C
ATOM	3513	CB	SER	51	-21.436	25.284	-20.896	1.00	27.94	L1	C
ATOM	3514	OG	SER	51	-20.133	24.813	-20.626	1.00	31.95	L1	O
ATOM	3515	C	SER	51	-23.494	24.628	-22.129	1.00	24.91	L1	C
ATOM	3516	O	SER	51	-24.080	24.808	-21.060	1.00	22.95	L1	O
ATOM	3517	N	ASN	52	-24.122	24.384	-23.275	1.00	24.45	L1	N
ATOM	3518	CA	ASN	52	-25.571	24.212	-23.351	1.00	25.66	L1	C
ATOM	3519	CB	ASN	52	-26.051	23.237	-22.270	1.00	24.57	L1	C
ATOM	3520	CG	ASN	52	-25.501	21.835	-22.463	1.00	25.41	L1	C
ATOM	3521	OD1	ASN	52	-25.299	21.383	-23.589	1.00	26.77	L1	O
ATOM	3522	ND2	ASN	52	-25.251	21.143	-21.361	1.00	25.99	L1	N
ATOM	3523	C	ASN	52	-26.350	25.520	-23.224	1.00	24.22	L1	C
ATOM	3524	O	ASN	52	-27.273	25.767	-23.999	1.00	24.09	L1	O
ATOM	3525	N	ASN	53	-25.982	26.351	-22.251	1.00	25.54	L1	N
ATOM	3526	CA	ASN	53	-26.821	27.482	-21.875	1.00	27.46	L1	C
ATOM	3527	CB	ASN	53	-27.830	27.038	-20.811	1.00	27.71	L1	C
ATOM	3528	CG	ASN	53	-27.154	26.453	-19.570	1.00	33.27	L1	C
ATOM	3529	OD1	ASN	53	-26.023	26.819	-19.226	1.00	32.11	L1	O
ATOM	3530	ND2	ASN	53	-27.846	25.537	-18.894	1.00	32.47	L1	N
ATOM	3531	C	ASN	53	-26.064	28.712	-21.374	1.00	28.20	L1	C
ATOM	3532	O	ASN	53	-26.664	29.601	-20.766	1.00	26.89	L1	O
ATOM	3533	N	GLN	54	-24.758	28.772	-21.625	1.00	27.94	L1	N
ATOM	3534	CA	GLN	54	-23.949	29.907	-21.182	1.00	26.56	L1	C
ATOM	3535	CB	GLN	54	-22.620	29.410	-20.599	1.00	27.00	L1	C
ATOM	3536	CG	GLN	54	-22.772	28.528	-19.362	1.00	26.24	L1	C
ATOM	3537	CD	GLN	54	-23.496	29.231	-18.222	1.00	29.64	L1	C
ATOM	3538	OE1	GLN	54	-22.973	30.174	-17.630	1.00	28.40	L1	O
ATOM	3539	NE2	GLN	54	-24.708	28.774	-17.914	1.00	26.55	L1	N
ATOM	3540	C	GLN	54	-23.670	30.923	-22.294	1.00	28.39	L1	C
ATOM	3541	O	GLN	54	-23.343	30.557	-23.424	1.00	27.97	L1	O
ATOM	3542	N	ARG	55	-23.797	32.205	-21.971	1.00	27.97	L1	N
ATOM	3543	CA	ARG	55	-23.526	33.247	-22.952	1.00	31.26	L1	C
ATOM	3544	CB	ARG	55	-24.612	34.327	-22.897	1.00	31.77	L1	C
ATOM	3545	CG	ARG	55	-25.921	33.916	-23.551	1.00	35.02	L1	C
ATOM	3546	CD	ARG	55	-27.002	34.966	-23.340	1.00	37.92	L1	C
ATOM	3547	NE	ARG	55	-27.466	34.997	-21.956	1.00	40.09	L1	N
ATOM	3548	CZ	ARG	55	-27.186	35.969	-21.093	1.00	42.88	L1	C
ATOM	3549	NH1	ARG	55	-26.442	37.003	-21.471	1.00	42.75	L1	N
ATOM	3550	NH2	ARG	55	-27.642	35.900	-19.845	1.00	42.31	L1	N
ATOM	3551	C	ARG	55	-22.160	33.892	-22.755	1.00	31.34	L1	C
ATOM	3552	O	ARG	55	-21.785	34.257	-21.647	1.00	30.63	L1	O
ATOM	3553	N	PRO	56	-21.396	34.041	-23.842	1.00	32.93	L1	N
ATOM	3554	CD	PRO	56	-21.569	33.417	-25.162	1.00	30.73	L1	C
ATOM	3555	CA	PRO	56	-20.205	34.891	-23.787	1.00	33.81	L1	C
ATOM	3556	CB	PRO	56	-19.605	34.772	-25.188	1.00	33.09	L1	C
ATOM	3557	CG	PRO	56	-20.196	33.528	-25.754	1.00	33.61	L1	C
ATOM	3558	C	PRO	56	-20.632	36.322	-23.485	1.00	36.31	L1	C
ATOM	3559	O	PRO	56	-21.763	36.721	-23.787	1.00	36.25	L1	O
ATOM	3560	N	SER	57	-19.731	37.084	-22.891	1.00	37.50	L1	N
ATOM	3561	CA	SER	57	-19.946	38.507	-22.683	1.00	41.08	L1	C
ATOM	3562	CB	SER	57	-18.687	39.139	-22.080	1.00	42.30	L1	C
ATOM	3563	OG	SER	57	-18.841	40.537	-21.919	1.00	40.81	L1	O
ATOM	3564	C	SER	57	-20.259	39.151	-24.033	1.00	42.62	L1	C
ATOM	3565	O	SER	57	-19.583	38.883	-25.028	1.00	41.85	L1	O
ATOM	3566	N	GLY	58	-21.290	39.990	-24.068	1.00	43.59	L1	N
ATOM	3567	CA	GLY	58	-21.639	40.665	-25.308	1.00	44.33	L1	C
ATOM	3568	C	GLY	58	-22.635	39.902	-26.160	1.00	44.22	L1	C
ATOM	3569	O	GLY	58	-22.762	40.154	-27.362	1.00	46.95	L1	O
ATOM	3570	N	VAL	59	-23.329	38.949	-25.545	1.00	41.84	L1	N
ATOM	3571	CA	VAL	59	-24.456	38.284	-26.187	1.00	38.40	L1	C
ATOM	3572	CB	VAL	59	-24.213	36.759	-26.312	1.00	37.83	L1	C
ATOM	3573	CG1	VAL	59	-25.456	36.073	-26.854	1.00	35.01	L1	C
ATOM	3574	CG2	VAL	59	-23.019	36.494	-27.223	1.00	34.54	L1	C
ATOM	3575	C	VAL	59	-25.710	38.523	-25.353	1.00	37.89	L1	C
ATOM	3576	O	VAL	59	-25.750	38.195	-24.172	1.00	36.97	L1	O
ATOM	3577	N	PRO	60	-26.753	39.099	-25.967	1.00	38.59	L1	N
ATOM	3578	CD	PRO	60	-26.783	39.481	-27.387	1.00	38.95	L1	C
ATOM	3579	CA	PRO	60	-28.000	39.460	-25.281	1.00	39.12	L1	C
ATOM	3580	CB	PRO	60	-28.852	40.083	-26.384	1.00	38.57	L1	C
ATOM	3581	CG	PRO	60	-27.881	40.495	-27.428	1.00	40.32	L1	C
ATOM	3582	C	PRO	60	-28.694	38.251	-24.657	1.00	39.92	L1	C
ATOM	3583	O	PRO	60	-28.709	37.162	-25.238	1.00	40.29	L1	O
ATOM	3584	N	ASP	61	-29.281	38.449	-23.482	1.00	38.79	L1	N
ATOM	3585	CA	ASP	61	-29.944	37.362	-22.780	1.00	39.81	L1	C

ATOM	3586	CB	ASP	61	-30.266	37.774	-21.339	1.00	46.00	L1	C
ATOM	3587	CG	ASP	61	-31.094	39.054	-21.258	1.00	50.76	L1	C
ATOM	3588	OD1	ASP	61	-31.389	39.489	-20.123	1.00	53.85	L1	O
ATOM	3589	OD2	ASP	61	-31.446	39.624	-22.317	1.00	52.25	L1	O
ATOM	3590	C	ASP	61	-31.213	36.868	-23.467	1.00	37.81	L1	C
ATOM	3591	O	ASP	61	-31.837	35.917	-23.004	1.00	39.35	L1	O
ATOM	3592	N	ARG	62	-31.601	37.495	-24.571	1.00	36.57	L1	N
ATOM	3593	CA	ARG	62	-32.735	36.985	-25.332	1.00	35.73	L1	C
ATOM	3594	CB	ARG	62	-33.302	38.073	-26.248	1.00	36.98	L1	C
ATOM	3595	CG	ARG	62	-32.312	38.621	-27.243	1.00	40.00	L1	C
ATOM	3596	CD	ARG	62	-32.835	39.889	-27.898	1.00	40.98	L1	C
ATOM	3597	NE	ARG	62	-31.860	40.430	-28.836	1.00	38.44	L1	N
ATOM	3598	CZ	ARG	62	-31.799	40.077	-30.111	1.00	36.96	L1	C
ATOM	3599	NH1	ARG	62	-32.664	39.193	-30.580	1.00	37.84	L1	N
ATOM	3600	NH2	ARG	62	-30.870	40.589	-30.905	1.00	36.03	L1	N
ATOM	3601	C	ARG	62	-32.342	35.752	-26.153	1.00	34.56	L1	C
ATOM	3602	O	ARG	62	-33.199	35.055	-26.687	1.00	32.53	L1	O
ATOM	3603	N	PHE	63	-31.041	35.489	-26.248	1.00	34.16	L1	N
ATOM	3604	CA	PHE	63	-30.561	34.192	-26.738	1.00	33.82	L1	C
ATOM	3605	CB	PHE	63	-29.210	34.345	-27.436	1.00	31.07	L1	C
ATOM	3606	CG	PHE	63	-29.272	35.142	-28.706	1.00	31.60	L1	C
ATOM	3607	CD1	PHE	63	-29.616	34.535	-29.903	1.00	30.71	L1	C
ATOM	3608	CD2	PHE	63	-28.977	36.496	-28.703	1.00	31.31	L1	C
ATOM	3609	CE1	PHE	63	-29.664	35.262	-31.074	1.00	32.32	L1	C
ATOM	3610	CE2	PHE	63	-29.023	37.231	-29.875	1.00	33.14	L1	C
ATOM	3611	CZ	PHE	63	-29.367	36.612	-31.062	1.00	33.35	L1	C
ATOM	3612	C	PHE	63	-30.404	33.236	-25.564	1.00	33.18	L1	C
ATOM	3613	O	PHE	63	-29.770	33.576	-24.568	1.00	34.53	L1	O
ATOM	3614	N	SER	64	-30.982	32.046	-25.676	1.00	31.90	L1	N
ATOM	3615	CA	SER	64	-30.829	31.042	-24.633	1.00	31.99	L1	C
ATOM	3616	CB	SER	64	-31.984	31.139	-23.627	1.00	33.55	L1	C
ATOM	3617	OG	SER	64	-33.196	30.648	-24.176	1.00	34.69	L1	O
ATOM	3618	C	SER	64	-30.777	29.638	-25.226	1.00	30.65	L1	C
ATOM	3619	O	SER	64	-31.400	29.360	-26.247	1.00	32.44	L1	O
ATOM	3620	N	GLY	65	-30.029	28.750	-24.581	1.00	30.08	L1	N
ATOM	3621	CA	GLY	65	-29.917	27.396	-25.086	1.00	28.03	L1	C
ATOM	3622	C	GLY	65	-30.418	26.371	-24.097	1.00	28.41	L1	C
ATOM	3623	O	GLY	65	-30.419	26.609	-22.893	1.00	28.15	L1	O
ATOM	3624	N	SER	66	-30.854	25.223	-24.603	1.00	29.78	L1	N
ATOM	3625	CA	SER	66	-31.205	24.113	-23.730	1.00	29.50	L1	C
ATOM	3626	CB	SER	66	-32.708	24.111	-23.439	1.00	30.04	L1	C
ATOM	3627	OG	SER	66	-33.461	23.989	-24.632	1.00	33.58	L1	O
ATOM	3628	C	SER	66	-30.792	22.782	-24.337	1.00	29.31	L1	C
ATOM	3629	O	SER	66	-30.605	22.661	-25.551	1.00	28.96	L1	O
ATOM	3630	N	LYS	67	-30.636	21.790	-23.472	1.00	29.49	L1	N
ATOM	3631	CA	LYS	67	-30.268	20.448	-23.886	1.00	31.34	L1	C
ATOM	3632	CB	LYS	67	-28.789	20.182	-23.587	1.00	29.79	L1	C
ATOM	3633	CG	LYS	67	-28.422	18.700	-23.598	1.00	30.92	L1	C
ATOM	3634	CD	LYS	67	-26.945	18.480	-23.271	1.00	31.59	L1	C
ATOM	3635	CE	LYS	67	-26.665	17.014	-22.974	1.00	31.18	L1	C
ATOM	3636	NZ	LYS	67	-25.207	16.754	-22.808	1.00	32.99	L1	N
ATOM	3637	C	LYS	67	-31.123	19.451	-23.119	1.00	32.11	L1	C
ATOM	3638	O	LYS	67	-31.355	19.614	-21.921	1.00	33.29	L1	O
ATOM	3639	N	SER	68	-31.591	18.420	-23.808	1.00	32.41	L1	N
ATOM	3640	CA	SER	68	-32.289	17.334	-23.137	1.00	33.97	L1	C
ATOM	3641	CB	SER	68	-33.788	17.624	-23.098	1.00	35.12	L1	C
ATOM	3642	OG	SER	68	-34.481	16.551	-22.492	1.00	39.24	L1	O
ATOM	3643	C	SER	68	-32.038	16.010	-23.851	1.00	32.19	L1	C
ATOM	3644	O	SER	68	-32.356	15.861	-25.027	1.00	31.42	L1	O
ATOM	3645	N	GLY	69	-31.470	15.046	-23.134	1.00	31.82	L1	N
ATOM	3646	CA	GLY	69	-31.165	13.768	-23.750	1.00	30.65	L1	C
ATOM	3647	C	GLY	69	-30.148	13.927	-24.865	1.00	31.46	L1	C
ATOM	3648	O	GLY	69	-29.026	14.364	-24.623	1.00	31.98	L1	O
ATOM	3649	N	THR	70	-30.531	13.572	-26.087	1.00	30.21	L1	N
ATOM	3650	CA	THR	70	-29.627	13.705	-27.221	1.00	29.74	L1	C
ATOM	3651	CB	THR	70	-29.517	12.384	-28.011	1.00	31.03	L1	C
ATOM	3652	OG1	THR	70	-30.813	11.994	-28.472	1.00	32.35	L1	O
ATOM	3653	CG2	THR	70	-28.934	11.269	-27.126	1.00	26.28	L1	C
ATOM	3654	C	THR	70	-30.040	14.820	-28.179	1.00	29.23	L1	C
ATOM	3655	O	THR	70	-29.630	14.837	-29.336	1.00	28.90	L1	O
ATOM	3656	N	SER	71	-30.841	15.759	-27.687	1.00	28.99	L1	N
ATOM	3657	CA	SER	71	-31.244	16.908	-28.486	1.00	27.69	L1	C
ATOM	3658	CB	SER	71	-32.719	16.795	-28.860	1.00	29.27	L1	C
ATOM	3659	OG	SER	71	-32.874	15.835	-29.886	1.00	38.45	L1	O
ATOM	3660	C	SER	71	-31.003	18.208	-27.745	1.00	24.99	L1	C
ATOM	3661	O	SER	71	-30.824	18.208	-26.535	1.00	25.72	L1	O

ATOM	3662	N	ALA	72	-30.995	19.315	-28.480	1.00	23.94	L1	N
ATOM	3663	CA	ALA	72	-30.769	20.626	-27.884	1.00	26.16	L1	C
ATOM	3664	CB	ALA	72	-29.272	20.976	-27.903	1.00	22.79	L1	C
ATOM	3665	C	ALA	72	-31.555	21.669	-28.654	1.00	26.58	L1	C
ATOM	3666	O	ALA	72	-32.062	21.396	-29.742	1.00	26.75	L1	O
ATOM	3667	N	SER	73	-31.645	22.872	-28.102	1.00	28.14	L1	N
ATOM	3668	CA	SER	73	-32.438	23.910	-28.745	1.00	29.91	L1	C
ATOM	3669	CB	SER	73	-33.900	23.788	-28.305	1.00	30.18	L1	C
ATOM	3670	OG	SER	73	-34.699	24.715	-29.010	1.00	37.35	L1	O
ATOM	3671	C	SER	73	-31.926	25.313	-28.453	1.00	28.74	L1	C
ATOM	3672	O	SER	73	-31.501	25.613	-27.337	1.00	30.53	L1	O
ATOM	3673	N	LEU	74	-31.955	26.161	-29.472	1.00	29.11	L1	N
ATOM	3674	CA	LEU	74	-31.628	27.574	-29.321	1.00	31.29	L1	C
ATOM	3675	CB	LEU	74	-30.694	28.029	-30.449	1.00	28.58	L1	C
ATOM	3676	CG	LEU	74	-30.226	29.483	-30.412	1.00	29.48	L1	C
ATOM	3677	CD1	LEU	74	-29.347	29.689	-29.201	1.00	29.08	L1	C
ATOM	3678	CD2	LEU	74	-29.464	29.827	-31.694	1.00	28.84	L1	C
ATOM	3679	C	LEU	74	-32.931	28.366	-29.384	1.00	32.05	L1	C
ATOM	3680	O	LEU	74	-33.707	28.221	-30.327	1.00	31.82	L1	O
ATOM	3681	N	ALA	75	-33.176	29.194	-28.376	1.00	31.88	L1	N
ATOM	3682	CA	ALA	75	-34.368	30.028	-28.374	1.00	32.47	L1	C
ATOM	3683	CB	ALA	75	-35.148	29.833	-27.077	1.00	32.36	L1	C
ATOM	3684	C	ALA	75	-33.952	31.479	-28.520	1.00	32.90	L1	C
ATOM	3685	O	ALA	75	-33.072	31.955	-27.803	1.00	33.92	L1	O
ATOM	3686	N	ILE	76	-34.575	32.175	-29.462	1.00	33.40	L1	N
ATOM	3687	CA	ILE	76	-34.331	33.597	-29.635	1.00	34.46	L1	C
ATOM	3688	CB	ILE	76	-33.824	33.906	-31.058	1.00	34.25	L1	C
ATOM	3689	CG2	ILE	76	-33.403	35.370	-31.155	1.00	32.73	L1	C
ATOM	3690	CG1	ILE	76	-32.626	33.014	-31.393	1.00	35.86	L1	C
ATOM	3691	CD1	ILE	76	-32.170	33.122	-32.847	1.00	35.68	L1	C
ATOM	3692	C	ILE	76	-35.648	34.331	-29.410	1.00	36.20	L1	C
ATOM	3693	O	ILE	76	-36.586	34.210	-30.204	1.00	35.91	L1	O
ATOM	3694	N	SER	77	-35.722	35.082	-28.319	1.00	36.94	L1	N
ATOM	3695	CA	SER	77	-36.907	35.878	-28.038	1.00	38.93	L1	C
ATOM	3696	CB	SER	77	-37.273	35.777	-26.555	1.00	39.60	L1	C
ATOM	3697	OG	SER	77	-36.215	36.242	-25.739	1.00	42.81	L1	O
ATOM	3698	C	SER	77	-36.655	37.331	-28.424	1.00	39.07	L1	C
ATOM	3699	O	SER	77	-35.503	37.756	-28.574	1.00	37.94	L1	O
ATOM	3700	N	GLY	78	-37.733	38.089	-28.600	1.00	37.57	L1	N
ATOM	3701	CA	GLY	78	-37.585	39.485	-28.966	1.00	35.79	L1	C
ATOM	3702	C	GLY	78	-36.839	39.619	-30.276	1.00	35.60	L1	C
ATOM	3703	O	GLY	78	-35.946	40.458	-30.423	1.00	33.98	L1	O
ATOM	3704	N	LEU	79	-37.212	38.781	-31.236	1.00	35.78	L1	N
ATOM	3705	CA	LEU	79	-36.506	38.703	-32.506	1.00	35.73	L1	C
ATOM	3706	CB	LEU	79	-37.290	37.812	-33.468	1.00	36.94	L1	C
ATOM	3707	CG	LEU	79	-36.602	37.311	-34.738	1.00	37.13	L1	C
ATOM	3708	CD1	LEU	79	-35.333	36.537	-34.392	1.00	34.89	L1	C
ATOM	3709	CD2	LEU	79	-37.583	36.423	-35.490	1.00	37.56	L1	C
ATOM	3710	C	LEU	79	-36.301	40.082	-33.124	1.00	36.83	L1	C
ATOM	3711	O	LEU	79	-37.241	40.872	-33.244	1.00	36.62	L1	O
ATOM	3712	N	GLN	80	-35.064	40.362	-33.519	1.00	37.09	L1	N
ATOM	3713	CA	GLN	80	-34.739	41.593	-34.221	1.00	38.15	L1	C
ATOM	3714	CB	GLN	80	-33.738	42.405	-33.403	1.00	42.19	L1	C
ATOM	3715	CG	GLN	80	-34.200	42.669	-31.979	1.00	49.70	L1	C
ATOM	3716	CD	GLN	80	-33.230	43.539	-31.213	1.00	56.13	L1	C
ATOM	3717	OE1	GLN	80	-32.224	43.999	-31.764	1.00	58.59	L1	O
ATOM	3718	NE2	GLN	80	-33.520	43.772	-29.932	1.00	58.28	L1	N
ATOM	3719	C	GLN	80	-34.154	41.270	-35.592	1.00	38.49	L1	C
ATOM	3720	O	GLN	80	-33.627	40.177	-35.811	1.00	36.48	L1	O
ATOM	3721	N	SER	81	-34.243	42.220	-36.516	1.00	36.90	L1	N
ATOM	3722	CA	SER	81	-33.800	41.975	-37.881	1.00	38.24	L1	C
ATOM	3723	CB	SER	81	-34.135	43.181	-38.771	1.00	37.34	L1	C
ATOM	3724	OG	SER	81	-33.438	44.336	-38.346	1.00	41.79	L1	O
ATOM	3725	C	SER	81	-32.298	41.661	-37.959	1.00	38.41	L1	C
ATOM	3726	O	SER	81	-31.857	40.977	-38.876	1.00	39.41	L1	O
ATOM	3727	N	GLU	82	-31.518	42.156	-37.001	1.00	37.61	L1	N
ATOM	3728	CA	GLU	82	-30.085	41.857	-36.964	1.00	37.59	L1	C
ATOM	3729	CB	GLU	82	-29.382	42.649	-35.855	1.00	41.17	L1	C
ATOM	3730	CG	GLU	82	-29.697	44.126	-35.838	1.00	50.54	L1	C
ATOM	3731	CD	GLU	82	-31.072	44.404	-35.270	1.00	53.17	L1	C
ATOM	3732	OE1	GLU	82	-31.236	44.292	-34.036	1.00	57.55	L1	O
ATOM	3733	OE2	GLU	82	-31.986	44.727	-36.057	1.00	56.15	L1	O
ATOM	3734	C	GLU	82	-29.860	40.370	-36.706	1.00	34.12	L1	C
ATOM	3735	O	GLU	82	-28.765	39.862	-36.899	1.00	31.06	L1	O
ATOM	3736	N	ASP	83	-30.898	39.681	-36.250	1.00	31.15	L1	N
ATOM	3737	CA	ASP	83	-30.779	38.265	-35.941	1.00	30.79	L1	C

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ATOM   3738  CB  ASP   83     -31.877  37.843 -34.962  1.00 30.65      L1   C
ATOM   3739  CG  ASP   83     -31.797  38.587 -33.644  1.00 32.56      L1   C
ATOM   3740  OD1 ASP   83     -30.697  39.067 -33.289  1.00 34.97      L1   O
ATOM   3741  OD2 ASP   83     -32.834  38.692 -32.961  1.00 31.83      L1   O
ATOM   3742  C   ASP   83     -30.846  37.395 -37.194  1.00 29.63      L1   C
ATOM   3743  O   ASP   83     -30.649  36.188 -37.113  1.00 28.37      L1   O
ATOM   3744  N   GLU   84     -31.121  38.002 -38.348  1.00 29.00      L1   N
ATOM   3745  CA  GLU   84     -31.189  37.234 -39.588  1.00 30.03      L1   C
ATOM   3746  CB  GLU   84     -31.730  38.088 -40.741  1.00 30.22      L1   C
ATOM   3747  CG  GLU   84     -31.723  37.358 -42.089  1.00 31.01      L1   C
ATOM   3748  CD  GLU   84     -32.799  37.846 -43.050  1.00 35.95      L1   C
ATOM   3749  OE1 GLU   84     -32.515  37.939 -44.265  1.00 37.52      L1   O
ATOM   3750  OE2 GLU   84     -33.930  38.132 -42.598  1.00 36.88      L1   O
ATOM   3751  C   GLU   84     -29.800  36.714 -39.948  1.00 30.69      L1   C
ATOM   3752  O   GLU   84     -28.873  37.496 -40.160  1.00 31.59      L1   O
ATOM   3753  N   ALA   85     -29.663  35.394 -40.021  1.00 29.97      L1   N
ATOM   3754  CA  ALA   85     -28.345  34.780 -40.110  1.00 28.04      L1   C
ATOM   3755  CB  ALA   85     -27.499  35.206 -38.915  1.00 28.83      L1   C
ATOM   3756  C   ALA   85     -28.440  33.265 -40.153  1.00 27.69      L1   C
ATOM   3757  O   ALA   85     -29.522  32.692 -39.988  1.00 24.83      L1   O
ATOM   3758  N   ASP   86     -27.293  32.625 -40.377  1.00 28.83      L1   N
ATOM   3759  CA  ASP   86     -27.171  31.173 -40.263  1.00 27.36      L1   C
ATOM   3760  CB  ASP   86     -26.098  30.653 -41.224  1.00 30.73      L1   C
ATOM   3761  CG  ASP   86     -26.494  30.794 -42.683  1.00 32.29      L1   C
ATOM   3762  OD1 ASP   86     -27.659  30.501 -43.017  1.00 36.16      L1   O
ATOM   3763  OD2 ASP   86     -25.639  31.194 -43.497  1.00 33.33      L1   O
ATOM   3764  C   ASP   86     -26.775  30.813 -38.835  1.00 26.79      L1   C
ATOM   3765  O   ASP   86     -25.899  31.451 -38.245  1.00 26.47      L1   O
ATOM   3766  N   TYR   87     -27.419  29.789 -38.286  1.00 25.78      L1   N
ATOM   3767  CA  TYR   87     -27.113  29.317 -36.946  1.00 26.34      L1   C
ATOM   3768  CB  TYR   87     -28.346  29.450 -36.043  1.00 26.23      L1   C
ATOM   3769  CG  TYR   87     -28.702  30.888 -35.752  1.00 25.63      L1   C
ATOM   3770  CD1 TYR   87     -29.451  31.636 -36.656  1.00 26.80      L1   C
ATOM   3771  CE1 TYR   87     -29.708  32.973 -36.434  1.00 27.24      L1   C
ATOM   3772  CD2 TYR   87     -28.228  31.520 -34.612  1.00 25.13      L1   C
ATOM   3773  CE2 TYR   87     -28.479  32.856 -34.378  1.00 26.81      L1   C
ATOM   3774  CZ  TYR   87     -29.216  33.579 -35.292  1.00 27.91      L1   C
ATOM   3775  OH  TYR   87     -29.448  34.914 -35.069  1.00 28.02      L1   O
ATOM   3776  C   TYR   87     -26.642  27.870 -36.983  1.00 27.87      L1   C
ATOM   3777  O   TYR   87     -27.238  27.031 -37.662  1.00 28.78      L1   O
ATOM   3778  N   TYR   88     -25.568  27.592 -36.247  1.00 26.11      L1   N
ATOM   3779  CA  TYR   88     -24.955  26.271 -36.230  1.00 23.64      L1   C
ATOM   3780  CB  TYR   88     -23.557  26.328 -36.853  1.00 22.55      L1   C
ATOM   3781  CG  TYR   88     -23.532  26.659 -38.331  1.00 22.94      L1   C
ATOM   3782  CD1 TYR   88     -23.269  27.953 -38.768  1.00 23.36      L1   C
ATOM   3783  CE1 TYR   88     -23.236  28.260 -40.115  1.00 22.59      L1   C
ATOM   3784  CD2 TYR   88     -23.763  25.679 -39.286  1.00 22.71      L1   C
ATOM   3785  CE2 TYR   88     -23.732  25.977 -40.645  1.00 22.56      L1   C
ATOM   3786  CZ  TYR   88     -23.467  27.270 -41.051  1.00 23.14      L1   C
ATOM   3787  OH  TYR   88     -23.430  27.581 -42.399  1.00 23.24      L1   O
ATOM   3788  C   TYR   88     -24.835  25.759 -34.803  1.00 24.19      L1   C
ATOM   3789  O   TYR   88     -24.509  26.525 -33.887  1.00 23.05      L1   O
ATOM   3790  N   CYS   89     -25.087  24.466 -34.613  1.00 23.34      L1   N
ATOM   3791  CA  CYS   89     -24.701  23.808 -33.371  1.00 24.76      L1   C
ATOM   3792  C   CYS   89     -23.428  22.986 -33.551  1.00 24.26      L1   C
ATOM   3793  O   CYS   89     -23.127  22.513 -34.644  1.00 24.97      L1   O
ATOM   3794  CB  CYS   89     -25.829  22.909 -32.844  1.00 24.95      L1   C
ATOM   3795  SG  CYS   89     -26.482  21.652 -33.993  1.00 29.02      L1   S
ATOM   3796  N   ALA   90     -22.684  22.824 -32.467  1.00 22.28      L1   N
ATOM   3797  CA  ALA   90     -21.419  22.115 -32.511  1.00 21.93      L1   C
ATOM   3798  CB  ALA   90     -20.279  23.103 -32.663  1.00 21.85      L1   C
ATOM   3799  C   ALA   90     -21.262  21.320 -31.229  1.00 22.39      L1   C
ATOM   3800  O   ALA   90     -21.575  21.813 -30.139  1.00 23.97      L1   O
ATOM   3801  N   VAL   91     -20.790  20.087 -31.365  1.00 21.91      L1   N
ATOM   3802  CA  VAL   91     -20.699  19.158 -30.241  1.00 22.14      L1   C
ATOM   3803  CB  VAL   91     -21.992  18.315 -30.112  1.00 22.24      L1   C
ATOM   3804  CG1 VAL   91     -21.908  17.407 -28.898  1.00 18.74      L1   C
ATOM   3805  CG2 VAL   91     -23.211  19.238 -30.019  1.00 19.37      L1   C
ATOM   3806  C   VAL   91     -19.530  18.209 -30.487  1.00 23.31      L1   C
ATOM   3807  O   VAL   91     -19.298  17.798 -31.619  1.00 24.62      L1   O
ATOM   3808  N   TRP   92     -18.792  17.864 -29.436  1.00 22.18      L1   N
ATOM   3809  CA  TRP   92     -17.749  16.857 -29.566  1.00 21.63      L1   C
ATOM   3810  CB  TRP   92     -16.850  16.862 -28.331  1.00 20.72      L1   C
ATOM   3811  CG  TRP   92     -15.604  16.033 -28.480  1.00 24.87      L1   C
ATOM   3812  CD2 TRP   92     -14.339  16.463 -29.011  1.00 23.79      L1   C
ATOM   3813  CE2 TRP   92     -13.453  15.372 -28.914  1.00 24.92      L1   C
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ATOM	3814	CE3	TRP	92	-13.874	17.663	-29.556	1.00	24.00	L1	C
ATOM	3815	CD1	TRP	92	-15.435	14.735	-28.100	1.00	24.13	L1	C
ATOM	3816	NE1	TRP	92	-14.147	14.331	-28.356	1.00	26.35	L1	N
ATOM	3817	CZ2	TRP	92	-12.125	15.443	-29.339	1.00	25.22	L1	C
ATOM	3818	CZ3	TRP	92	-12.581	17.734	-29.981	1.00	27.22	L1	C
ATOM	3819	CH2	TRP	92	-11.694	16.629	-29.868	1.00	26.53	L1	C
ATOM	3820	C	TRP	92	-18.378	15.470	-29.739	1.00	22.42	L1	C
ATOM	3821	O	TRP	92	-19.333	15.120	-29.053	1.00	21.21	L1	O
ATOM	3822	N	ASP	93	-17.838	14.684	-30.661	1.00	22.63	L1	N
ATOM	3823	CA	ASP	93	-18.303	13.316	-30.847	1.00	22.79	L1	C
ATOM	3824	CB	ASP	93	-18.629	13.054	-32.312	1.00	22.67	L1	C
ATOM	3825	CG	ASP	93	-19.345	11.732	-32.517	1.00	24.52	L1	C
ATOM	3826	OD1	ASP	93	-20.593	11.733	-32.528	1.00	24.55	L1	O
ATOM	3827	OD2	ASP	93	-18.662	10.695	-32.662	1.00	24.78	L1	O
ATOM	3828	C	ASP	93	-17.234	12.338	-30.393	1.00	23.59	L1	C
ATOM	3829	O	ASP	93	-16.085	12.430	-30.815	1.00	21.15	L1	O
ATOM	3830	N	ASP	94	-17.622	11.396	-29.541	1.00	22.22	L1	N
ATOM	3831	CA	ASP	94	-16.672	10.480	-28.931	1.00	23.56	L1	C
ATOM	3832	CB	ASP	94	-17.222	10.010	-27.579	1.00	22.41	L1	C
ATOM	3833	CG	ASP	94	-17.423	11.162	-26.610	1.00	25.03	L1	C
ATOM	3834	OD1	ASP	94	-16.432	11.869	-26.310	1.00	24.18	L1	O
ATOM	3835	OD2	ASP	94	-18.567	11.367	-26.153	1.00	23.82	L1	O
ATOM	3836	C	ASP	94	-16.291	9.279	-29.803	1.00	22.66	L1	C
ATOM	3837	O	ASP	94	-15.337	8.576	-29.498	1.00	23.33	L1	O
ATOM	3838	N	SER	95	-17.027	9.043	-30.885	1.00	23.45	L1	N
ATOM	3839	CA	SER	95	-16.663	7.979	-31.821	1.00	22.75	L1	C
ATOM	3840	CB	SER	95	-17.913	7.348	-32.444	1.00	21.53	L1	C
ATOM	3841	OG	SER	95	-18.715	6.725	-31.450	1.00	19.78	L1	O
ATOM	3842	C	SER	95	-15.762	8.524	-32.918	1.00	24.04	L1	C
ATOM	3843	O	SER	95	-14.806	7.867	-33.329	1.00	24.34	L1	O
ATOM	3844	N	LEU	96	-16.060	9.736	-33.374	1.00	22.90	L1	N
ATOM	3845	CA	LEU	96	-15.206	10.427	-34.328	1.00	23.33	L1	C
ATOM	3846	CB	LEU	96	-16.022	11.480	-35.076	1.00	25.43	L1	C
ATOM	3847	CG	LEU	96	-17.131	10.906	-35.962	1.00	26.90	L1	C
ATOM	3848	CD1	LEU	96	-18.143	11.991	-36.285	1.00	23.29	L1	C
ATOM	3849	CD2	LEU	96	-16.522	10.341	-37.226	1.00	27.63	L1	C
ATOM	3850	C	LEU	96	-14.008	11.093	-33.644	1.00	23.40	L1	C
ATOM	3851	O	LEU	96	-13.029	11.442	-34.300	1.00	22.31	L1	O
ATOM	3852	N	ASN	97	-14.094	11.267	-32.327	1.00	21.76	L1	N
ATOM	3853	CA	ASN	97	-13.032	11.919	-31.564	1.00	23.62	L1	C
ATOM	3854	CB	ASN	97	-11.765	11.064	-31.554	1.00	23.22	L1	C
ATOM	3855	CG	ASN	97	-11.961	9.742	-30.897	1.00	25.94	L1	C
ATOM	3856	OD1	ASN	97	-12.281	9.698	-29.701	1.00	24.63	L1	O
ATOM	3857	ND2	ASN	97	-11.794	8.651	-31.637	1.00	27.24	L1	N
ATOM	3858	C	ASN	97	-12.719	13.326	-32.054	1.00	24.97	L1	C
ATOM	3859	O	ASN	97	-11.554	13.677	-32.290	1.00	24.92	L1	O
ATOM	3860	N	GLY	98	-13.764	14.134	-32.199	1.00	21.90	L1	N
ATOM	3861	CA	GLY	98	-13.575	15.504	-32.632	1.00	23.01	L1	C
ATOM	3862	C	GLY	98	-14.900	16.223	-32.796	1.00	22.94	L1	C
ATOM	3863	O	GLY	98	-15.961	15.660	-32.516	1.00	19.38	L1	O
ATOM	3864	N	TRP	99	-14.835	17.465	-33.258	1.00	21.48	L1	N
ATOM	3865	CA	TRP	99	-16.017	18.309	-33.354	1.00	21.80	L1	C
ATOM	3866	CB	TRP	99	-15.602	19.784	-33.485	1.00	20.41	L1	C
ATOM	3867	CG	TRP	99	-15.097	20.349	-32.201	1.00	21.29	L1	C
ATOM	3868	CD2	TRP	99	-15.887	20.731	-31.070	1.00	20.83	L1	C
ATOM	3869	CE2	TRP	99	-14.993	21.130	-30.052	1.00	21.78	L1	C
ATOM	3870	CE3	TRP	99	-17.264	20.773	-30.819	1.00	21.01	L1	C
ATOM	3871	CD1	TRP	99	-13.792	20.533	-31.838	1.00	20.69	L1	C
ATOM	3872	NE1	TRP	99	-13.721	21.000	-30.548	1.00	18.42	L1	N
ATOM	3873	CZ2	TRP	99	-15.429	21.565	-28.797	1.00	21.72	L1	C
ATOM	3874	CZ3	TRP	99	-17.701	21.207	-29.572	1.00	24.75	L1	C
ATOM	3875	CH2	TRP	99	-16.782	21.597	-28.575	1.00	23.40	L1	C
ATOM	3876	C	TRP	99	-16.896	17.915	-34.530	1.00	21.60	L1	C
ATOM	3877	O	TRP	99	-16.411	17.673	-35.640	1.00	20.77	L1	O
ATOM	3878	N	VAL	100	-18.196	17.843	-34.280	1.00	21.15	L1	N
ATOM	3879	CA	VAL	100	-19.155	17.745	-35.366	1.00	20.99	L1	C
ATOM	3880	CB	VAL	100	-19.959	16.412	-35.299	1.00	21.63	L1	C
ATOM	3881	CG1	VAL	100	-19.013	15.237	-35.493	1.00	18.33	L1	C
ATOM	3882	CG2	VAL	100	-20.685	16.283	-33.965	1.00	20.08	L1	C
ATOM	3883	C	VAL	100	-20.094	18.945	-35.310	1.00	22.73	L1	C
ATOM	3884	O	VAL	100	-20.206	19.604	-34.271	1.00	21.19	L1	O
ATOM	3885	N	PHE	101	-20.731	19.244	-36.439	1.00	23.17	L1	N
ATOM	3886	CA	PHE	101	-21.625	20.393	-36.556	1.00	24.04	L1	C
ATOM	3887	CB	PHE	101	-21.052	21.421	-37.538	1.00	23.68	L1	C
ATOM	3888	CG	PHE	101	-19.847	22.175	-37.015	1.00	25.65	L1	C
ATOM	3889	CD1	PHE	101	-18.558	21.762	-37.330	1.00	25.40	L1	C

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ATOM   3890  CD2 PHE   101     -20.005  23.323 -36.253  1.00 25.22      L1   C
ATOM   3891  CE1 PHE   101     -17.449  22.484 -36.900  1.00 27.16      L1   C
ATOM   3892  CE2 PHE   101     -18.899  24.053 -35.818  1.00 26.53      L1   C
ATOM   3893  CZ  PHE   101     -17.621  23.632 -36.144  1.00 25.47      L1   C
ATOM   3894  C   PHE   101     -22.995  19.944 -37.060  1.00 24.46      L1   C
ATOM   3895  O   PHE   101     -23.134  18.856 -37.620  1.00 25.98      L1   O
ATOM   3896  N   GLY   102     -24.006  20.782 -36.856  1.00 24.12      L1   N
ATOM   3897  CA  GLY   102     -25.261  20.601 -37.560  1.00 23.71      L1   C
ATOM   3898  C   GLY   102     -25.108  21.230 -38.930  1.00 24.83      L1   C
ATOM   3899  O   GLY   102     -24.139  21.951 -39.169  1.00 25.59      L1   O
ATOM   3900  N   GLY   103     -26.049  20.974 -39.831  1.00 24.86      L1   N
ATOM   3901  CA  GLY   103     -25.923  21.487 -41.188  1.00 23.32      L1   C
ATOM   3902  C   GLY   103     -26.218  22.973 -41.301  1.00 23.92      L1   C
ATOM   3903  O   GLY   103     -26.063  23.565 -42.365  1.00 23.16      L1   O
ATOM   3904  N   GLY   104     -26.637  23.590 -40.205  1.00 24.09      L1   N
ATOM   3905  CA  GLY   104     -26.940  25.010 -40.247  1.00 25.44      L1   C
ATOM   3906  C   GLY   104     -28.406  25.294 -40.535  1.00 26.46      L1   C
ATOM   3907  O   GLY   104     -29.032  24.613 -41.348  1.00 25.23      L1   O
ATOM   3908  N   THR   105     -28.951  26.296 -39.850  1.00 27.15      L1   N
ATOM   3909  CA  THR   105     -30.313  26.762 -40.080  1.00 27.11      L1   C
ATOM   3910  CB  THR   105     -31.161  26.699 -38.801  1.00 26.94      L1   C
ATOM   3911  OG1 THR   105     -31.235  25.348 -38.335  1.00 25.27      L1   O
ATOM   3912  CG2 THR   105     -32.574  27.229 -39.074  1.00 27.18      L1   C
ATOM   3913  C   THR   105     -30.275  28.220 -40.511  1.00 28.67      L1   C
ATOM   3914  O   THR   105     -29.776  29.072 -39.770  1.00 28.70      L1   O
ATOM   3915  N   LYS   106     -30.811  28.511 -41.693  1.00 27.94      L1   N
ATOM   3916  CA  LYS   106     -30.958  29.895 -42.132  1.00 27.78      L1   C
ATOM   3917  CB  LYS   106     -31.076  29.959 -43.660  1.00 30.73      L1   C
ATOM   3918  CG  LYS   106     -31.292  31.370 -44.224  1.00 34.36      L1   C
ATOM   3919  CD  LYS   106     -31.505  31.327 -45.743  1.00 40.73      L1   C
ATOM   3920  CE  LYS   106     -32.204  32.589 -46.269  1.00 43.20      L1   C
ATOM   3921  NZ  LYS   106     -31.412  33.828 -45.992  1.00 43.79      L1   N
ATOM   3922  C   LYS   106     -32.204  30.506 -41.492  1.00 28.25      L1   C
ATOM   3923  O   LYS   106     -33.324  30.102 -41.790  1.00 27.19      L1   O
ATOM   3924  N   LEU   107     -32.002  31.476 -40.606  1.00 28.39      L1   N
ATOM   3925  CA  LEU   107     -33.115  32.167 -39.965  1.00 29.11      L1   C
ATOM   3926  CB  LEU   107     -32.759  32.529 -38.524  1.00 27.84      L1   C
ATOM   3927  CG  LEU   107     -33.833  33.302 -37.750  1.00 30.76      L1   C
ATOM   3928  CD1 LEU   107     -35.095  32.460 -37.646  1.00 28.82      L1   C
ATOM   3929  CD2 LEU   107     -33.320  33.665 -36.364  1.00 29.24      L1   C
ATOM   3930  C   LEU   107     -33.438  33.440 -40.731  1.00 29.95      L1   C
ATOM   3931  O   LEU   107     -32.583  34.313 -40.881  1.00 30.14      L1   O
ATOM   3932  N   THR   108     -34.668  33.544 -41.222  1.00 29.24      L1   N
ATOM   3933  CA  THR   108     -35.124  34.781 -41.835  1.00 29.21      L1   C
ATOM   3934  CB  THR   108     -35.949  34.519 -43.118  1.00 30.64      L1   C
ATOM   3935  OG1 THR   108     -35.126  33.888 -44.107  1.00 30.87      L1   O
ATOM   3936  CG2 THR   108     -36.482  35.837 -43.686  1.00 30.92      L1   C
ATOM   3937  C   THR   108     -35.993  35.535 -40.840  1.00 29.75      L1   C
ATOM   3938  O   THR   108     -36.920  34.973 -40.257  1.00 29.93      L1   O
ATOM   3939  N   VAL   109     -35.685  36.808 -40.637  1.00 30.82      L1   N
ATOM   3940  CA  VAL   109     -36.549  37.666 -39.837  1.00 31.06      L1   C
ATOM   3941  CB  VAL   109     -35.720  38.675 -39.012  1.00 29.89      L1   C
ATOM   3942  CG1 VAL   109     -36.636  39.619 -38.253  1.00 31.03      L1   C
ATOM   3943  CG2 VAL   109     -34.827  37.922 -38.043  1.00 30.10      L1   C
ATOM   3944  C   VAL   109     -37.469  38.404 -40.803  1.00 30.26      L1   C
ATOM   3945  O   VAL   109     -37.037  39.294 -41.537  1.00 28.33      L1   O
ATOM   3946  N   LEU   110     -38.737  38.007 -40.808  1.00 29.80      L1   N
ATOM   3947  CA  LEU   110     -39.672  38.430 -41.844  1.00 30.88      L1   C
ATOM   3948  CB  LEU   110     -41.047  37.813 -41.571  1.00 29.96      L1   C
ATOM   3949  CG  LEU   110     -41.036  36.279 -41.536  1.00 29.38      L1   C
ATOM   3950  CD1 LEU   110     -42.338  35.739 -40.950  1.00 28.86      L1   C
ATOM   3951  CD2 LEU   110     -40.828  35.757 -42.938  1.00 22.82      L1   C
ATOM   3952  C   LEU   110     -39.779  39.950 -41.934  1.00 30.13      L1   C
ATOM   3953  O   LEU   110     -40.124  40.616 -40.962  1.00 30.33      L1   O
ATOM   3954  N   GLY   111     -39.468  40.486 -43.110  1.00 31.70      L1   N
ATOM   3955  CA  GLY   111     -39.525  41.923 -43.327  1.00 32.20      L1   C
ATOM   3956  C   GLY   111     -40.510  42.285 -44.428  1.00 33.43      L1   C
ATOM   3957  O   GLY   111     -40.689  43.461 -44.768  1.00 35.10      L1   O
ATOM   3958  N   GLN   112     -41.150  41.266 -44.990  1.00 33.16      L1   N
ATOM   3959  CA  GLN   112     -42.217  41.465 -45.964  1.00 36.01      L1   C
ATOM   3960  CB  GLN   112     -41.627  41.718 -47.354  1.00 37.16      L1   C
ATOM   3961  CG  GLN   112     -40.857  40.542 -47.931  1.00 37.31      L1   C
ATOM   3962  CD  GLN   112     -40.276  40.863 -49.291  1.00 38.17      L1   C
ATOM   3963  OE1 GLN   112     -40.967  40.775 -50.306  1.00 40.76      L1   O
ATOM   3964  NE2 GLN   112     -39.004  41.243 -49.320  1.00 35.15      L1   N
ATOM   3965  C   GLN   112     -43.097  40.219 -45.984  1.00 36.50      L1   C
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ATOM	3966	O	GLN	112	-42.783	39.228	-45.333	1.00	35.56	L1	O
ATOM	3967	N	PRO	113	-44.224	40.263	-46.714	1.00	37.98	L1	N
ATOM	3968	CD	PRO	113	-44.796	41.426	-47.415	1.00	38.44	L1	C
ATOM	3969	CA	PRO	113	-45.108	39.091	-46.789	1.00	38.56	L1	C
ATOM	3970	CB	PRO	113	-46.269	39.566	-47.669	1.00	38.32	L1	C
ATOM	3971	CG	PRO	113	-46.248	41.057	-47.550	1.00	38.11	L1	C
ATOM	3972	C	PRO	113	-44.401	37.887	-47.403	1.00	39.28	L1	C
ATOM	3973	O	PRO	113	-43.503	38.037	-48.330	1.00	38.01	L1	O
ATOM	3974	N	LYS	114	-44.698	36.697	-46.891	1.00	39.70	L1	N
ATOM	3975	CA	LYS	114	-44.209	35.472	-47.515	1.00	43.10	L1	C
ATOM	3976	CB	LYS	114	-44.646	34.243	-46.708	1.00	41.99	L1	C
ATOM	3977	CG	LYS	114	-44.031	34.182	-45.310	1.00	44.06	L1	C
ATOM	3978	CD	LYS	114	-44.233	32.820	-44.646	1.00	44.49	L1	C
ATOM	3979	CE	LYS	114	-43.571	32.774	-43.276	1.00	46.38	L1	C
ATOM	3980	NZ	LYS	114	-43.874	31.518	-42.529	1.00	47.56	L1	N
ATOM	3981	C	LYS	114	-44.754	35.387	-48.937	1.00	44.55	L1	C
ATOM	3982	O	LYS	114	-45.889	35.782	-49.192	1.00	45.57	L1	O
ATOM	3983	N	ALA	115	-43.936	34.889	-49.861	1.00	45.01	L1	N
ATOM	3984	CA	ALA	115	-44.316	34.830	-51.267	1.00	45.57	L1	C
ATOM	3985	CB	ALA	115	-43.751	36.038	-52.015	1.00	43.32	L1	C
ATOM	3986	C	ALA	115	-43.831	33.540	-51.920	1.00	47.27	L1	C
ATOM	3987	O	ALA	115	-42.635	33.236	-51.918	1.00	47.27	L1	O
ATOM	3988	N	ALA	116	-44.774	32.789	-52.478	1.00	47.90	L1	N
ATOM	3989	CA	ALA	116	-44.469	31.572	-53.213	1.00	48.80	L1	C
ATOM	3990	CB	ALA	116	-45.758	30.823	-53.533	1.00	49.49	L1	C
ATOM	3991	C	ALA	116	-43.724	31.904	-54.500	1.00	49.45	L1	C
ATOM	3992	O	ALA	116	-43.989	32.922	-55.139	1.00	49.52	L1	O
ATOM	3993	N	PRO	117	-42.779	31.038	-54.892	1.00	50.68	L1	N
ATOM	3994	CD	PRO	117	-42.512	29.768	-54.194	1.00	50.62	L1	C
ATOM	3995	CA	PRO	117	-41.904	31.218	-56.055	1.00	52.06	L1	C
ATOM	3996	CB	PRO	117	-40.870	30.110	-55.897	1.00	52.12	L1	C
ATOM	3997	CG	PRO	117	-41.594	29.043	-55.141	1.00	51.66	L1	C
ATOM	3998	C	PRO	117	-42.639	31.105	-57.385	1.00	53.83	L1	C
ATOM	3999	O	PRO	117	-43.584	30.327	-57.518	1.00	53.40	L1	O
ATOM	4000	N	SER	118	-42.193	31.884	-58.366	1.00	56.36	L1	N
ATOM	4001	CA	SER	118	-42.597	31.679	-59.755	1.00	59.62	L1	C
ATOM	4002	CB	SER	118	-42.568	33.003	-60.526	1.00	60.11	L1	C
ATOM	4003	OG	SER	118	-43.416	33.970	-59.935	1.00	62.95	L1	O
ATOM	4004	C	SER	118	-41.621	30.703	-60.406	1.00	60.93	L1	C
ATOM	4005	O	SER	118	-40.406	30.854	-60.272	1.00	61.04	L1	O
ATOM	4006	N	VAL	119	-42.152	29.708	-61.110	1.00	62.70	L1	N
ATOM	4007	CA	VAL	119	-41.313	28.719	-61.779	1.00	64.16	L1	C
ATOM	4008	CB	VAL	119	-41.591	27.300	-61.236	1.00	63.53	L1	C
ATOM	4009	CG1	VAL	119	-40.704	26.287	-61.945	1.00	63.21	L1	C
ATOM	4010	CG2	VAL	119	-41.344	27.263	-59.737	1.00	64.05	L1	C
ATOM	4011	C	VAL	119	-41.522	28.715	-63.293	1.00	65.49	L1	C
ATOM	4012	O	VAL	119	-42.626	28.471	-63.779	1.00	65.80	L1	O
ATOM	4013	N	THR	120	-40.450	28.987	-64.032	1.00	67.43	L1	N
ATOM	4014	CA	THR	120	-40.470	28.914	-65.490	1.00	68.73	L1	C
ATOM	4015	CB	THR	120	-40.051	30.262	-66.117	1.00	68.82	L1	C
ATOM	4016	OG1	THR	120	-40.923	31.298	-65.648	1.00	68.76	L1	O
ATOM	4017	CG2	THR	120	-40.125	30.192	-67.636	1.00	68.58	L1	C
ATOM	4018	C	THR	120	-39.510	27.820	-65.971	1.00	69.56	L1	C
ATOM	4019	O	THR	120	-38.316	27.856	-65.669	1.00	69.35	L1	O
ATOM	4020	N	LEU	121	-40.036	26.853	-66.720	1.00	70.38	L1	N
ATOM	4021	CA	LEU	121	-39.257	25.683	-67.122	1.00	70.76	L1	C
ATOM	4022	CB	LEU	121	-39.920	24.411	-66.594	1.00	70.00	L1	C
ATOM	4023	CG	LEU	121	-39.202	23.105	-66.926	1.00	69.81	L1	C
ATOM	4024	CD1	LEU	121	-37.857	23.072	-66.219	1.00	69.72	L1	C
ATOM	4025	CD2	LEU	121	-40.061	21.928	-66.496	1.00	70.35	L1	C
ATOM	4026	C	LEU	121	-39.087	25.565	-68.637	1.00	71.13	L1	C
ATOM	4027	O	LEU	121	-40.058	25.372	-69.368	1.00	70.75	L1	O
ATOM	4028	N	PHE	122	-37.846	25.663	-69.101	1.00	71.79	L1	N
ATOM	4029	CA	PHE	122	-37.547	25.537	-70.523	1.00	73.09	L1	C
ATOM	4030	CB	PHE	122	-36.521	26.591	-70.944	1.00	72.54	L1	C
ATOM	4031	CG	PHE	122	-37.013	28.004	-70.821	1.00	72.59	L1	C
ATOM	4032	CD1	PHE	122	-36.678	28.777	-69.722	1.00	72.06	L1	C
ATOM	4033	CD2	PHE	122	-37.803	28.564	-71.813	1.00	72.99	L1	C
ATOM	4034	CE1	PHE	122	-37.121	30.084	-69.613	1.00	73.02	L1	C
ATOM	4035	CE2	PHE	122	-38.250	29.873	-71.711	1.00	72.95	L1	C
ATOM	4036	CZ	PHE	122	-37.908	30.633	-70.610	1.00	72.77	L1	C
ATOM	4037	C	PHE	122	-37.009	24.149	-70.864	1.00	74.17	L1	C
ATOM	4038	O	PHE	122	-36.059	23.669	-70.243	1.00	74.07	L1	O
ATOM	4039	N	PRO	123	-37.618	23.485	-71.861	1.00	74.99	L1	N
ATOM	4040	CD	PRO	123	-38.879	23.887	-72.509	1.00	75.22	L1	C
ATOM	4041	CA	PRO	123	-37.079	22.243	-72.430	1.00	74.86	L1	C

ATOM	4042	CB	PRO	123	-38.219	21.728	-73.305	1.00	74.82	L1	C
ATOM	4043	CG	PRO	123	-38.970	22.961	-73.693	1.00	75.30	L1	C
ATOM	4044	C	PRO	123	-35.808	22.515	-73.238	1.00	74.68	L1	C
ATOM	4045	O	PRO	123	-35.491	23.666	-73.542	1.00	74.38	L1	O
ATOM	4046	N	PRO	124	-35.061	21.456	-73.591	1.00	75.00	L1	N
ATOM	4047	CD	PRO	124	-35.268	20.052	-73.198	1.00	74.86	L1	C
ATOM	4048	CA	PRO	124	-33.841	21.625	-74.390	1.00	74.74	L1	C
ATOM	4049	CB	PRO	124	-33.309	20.199	-74.545	1.00	74.36	L1	C
ATOM	4050	CG	PRO	124	-33.908	19.446	-73.404	1.00	74.67	L1	C
ATOM	4051	C	PRO	124	-34.141	22.267	-75.742	1.00	74.62	L1	C
ATOM	4052	O	PRO	124	-35.135	21.934	-76.387	1.00	74.35	L1	O
ATOM	4053	N	SER	125	-33.284	23.189	-76.167	1.00	74.96	L1	N
ATOM	4054	CA	SER	125	-33.440	23.813	-77.475	1.00	75.14	L1	C
ATOM	4055	CB	SER	125	-32.533	25.043	-77.591	1.00	73.75	L1	C
ATOM	4056	OG	SER	125	-31.171	24.709	-77.395	1.00	72.15	L1	O
ATOM	4057	C	SER	125	-33.095	22.801	-78.563	1.00	76.50	L1	C
ATOM	4058	O	SER	125	-32.531	21.743	-78.282	1.00	76.63	L1	O
ATOM	4059	N	SER	126	-33.446	23.120	-79.804	1.00	77.77	L1	N
ATOM	4060	CA	SER	126	-33.135	22.240	-80.924	1.00	78.66	L1	C
ATOM	4061	CB	SER	126	-33.941	22.651	-82.160	1.00	79.08	L1	C
ATOM	4062	OG	SER	126	-35.334	22.551	-81.917	1.00	79.83	L1	O
ATOM	4063	C	SER	126	-31.641	22.294	-81.235	1.00	78.82	L1	C
ATOM	4064	O	SER	126	-31.039	21.288	-81.610	1.00	78.71	L1	O
ATOM	4065	N	GLU	127	-31.051	23.474	-81.063	1.00	79.12	L1	N
ATOM	4066	CA	GLU	127	-29.627	23.677	-81.312	1.00	80.10	L1	C
ATOM	4067	CB	GLU	127	-29.247	25.144	-81.071	1.00	80.24	L1	C
ATOM	4068	CG	GLU	127	-29.826	26.127	-82.078	1.00	81.24	L1	C
ATOM	4069	CD	GLU	127	-31.272	26.496	-81.787	1.00	82.17	L1	C
ATOM	4070	OE1	GLU	127	-31.928	25.788	-80.993	1.00	82.13	L1	O
ATOM	4071	OE2	GLU	127	-31.754	27.499	-82.356	1.00	81.88	L1	O
ATOM	4072	C	GLU	127	-28.763	22.786	-80.424	1.00	80.55	L1	C
ATOM	4073	O	GLU	127	-27.606	22.516	-80.742	1.00	80.88	L1	O
ATOM	4074	N	GLU	128	-29.326	22.335	-79.308	1.00	81.34	L1	N
ATOM	4075	CA	GLU	128	-28.570	21.553	-78.339	1.00	82.00	L1	C
ATOM	4076	CB	GLU	128	-28.967	21.953	-76.911	1.00	82.28	L1	C
ATOM	4077	CG	GLU	128	-28.026	21.427	-75.834	1.00	81.82	L1	C
ATOM	4078	CD	GLU	128	-28.481	21.778	-74.430	1.00	81.09	L1	C
ATOM	4079	OE1	GLU	128	-27.643	21.730	-73.505	1.00	79.98	L1	O
ATOM	4080	OE2	GLU	128	-29.674	22.099	-74.251	1.00	81.48	L1	O
ATOM	4081	C	GLU	128	-28.794	20.057	-78.539	1.00	82.27	L1	C
ATOM	4082	O	GLU	128	-27.845	19.273	-78.510	1.00	81.56	L1	O
ATOM	4083	N	LEU	129	-30.047	19.664	-78.746	1.00	82.77	L1	N
ATOM	4084	CA	LEU	129	-30.367	18.262	-78.984	1.00	83.70	L1	C
ATOM	4085	CB	LEU	129	-31.860	18.099	-79.282	1.00	83.20	L1	C
ATOM	4086	CG	LEU	129	-32.808	18.383	-78.116	1.00	83.28	L1	C
ATOM	4087	CD1	LEU	129	-34.253	18.198	-78.557	1.00	82.64	L1	C
ATOM	4088	CD2	LEU	129	-32.476	17.449	-76.965	1.00	83.46	L1	C
ATOM	4089	C	LEU	129	-29.544	17.733	-80.154	1.00	84.66	L1	C
ATOM	4090	O	LEU	129	-29.147	16.567	-80.172	1.00	84.88	L1	O
ATOM	4091	N	GLN	130	-29.286	18.602	-81.127	1.00	85.33	L1	N
ATOM	4092	CA	GLN	130	-28.463	18.242	-82.275	1.00	86.17	L1	C
ATOM	4093	CB	GLN	130	-28.619	19.281	-83.390	1.00	86.49	L1	C
ATOM	4094	CG	GLN	130	-28.015	20.639	-83.063	1.00	86.88	L1	C
ATOM	4095	CD	GLN	130	-28.181	21.640	-84.191	1.00	87.75	L1	C
ATOM	4096	OE1	GLN	130	-27.319	22.496	-84.409	1.00	87.18	L1	O
ATOM	4097	NE2	GLN	130	-29.292	21.540	-84.915	1.00	86.91	L1	N
ATOM	4098	C	GLN	130	-26.997	18.152	-81.863	1.00	86.31	L1	C
ATOM	4099	O	GLN	130	-26.260	17.289	-82.342	1.00	87.49	L1	O
ATOM	4100	N	ALA	131	-26.581	19.046	-80.971	1.00	85.27	L1	N
ATOM	4101	CA	ALA	131	-25.210	19.055	-80.478	1.00	84.10	L1	C
ATOM	4102	CB	ALA	131	-24.918	20.373	-79.770	1.00	83.24	L1	C
ATOM	4103	C	ALA	131	-24.971	17.882	-79.528	1.00	83.51	L1	C
ATOM	4104	O	ALA	131	-23.918	17.792	-78.894	1.00	83.48	L1	O
ATOM	4105	N	ASN	132	-25.959	16.994	-79.433	1.00	82.74	L1	N
ATOM	4106	CA	ASN	132	-25.842	15.767	-78.649	1.00	82.64	L1	C
ATOM	4107	CB	ASN	132	-24.526	15.054	-78.983	1.00	82.71	L1	C
ATOM	4108	CG	ASN	132	-24.503	13.611	-78.508	1.00	82.60	L1	C
ATOM	4109	OD1	ASN	132	-25.549	12.972	-78.367	1.00	81.48	L1	O
ATOM	4110	ND2	ASN	132	-23.304	13.090	-78.261	1.00	81.72	L1	N
ATOM	4111	C	ASN	132	-25.920	16.049	-77.146	1.00	82.67	L1	C
ATOM	4112	O	ASN	132	-25.553	15.209	-76.322	1.00	82.72	L1	O
ATOM	4113	N	LYS	133	-26.400	17.241	-76.801	1.00	82.45	L1	N
ATOM	4114	CA	LYS	133	-26.575	17.639	-75.408	1.00	81.60	L1	C
ATOM	4115	CB	LYS	133	-25.705	18.861	-75.100	1.00	81.98	L1	C
ATOM	4116	CG	LYS	133	-24.278	18.755	-75.616	1.00	82.82	L1	C
ATOM	4117	CD	LYS	133	-23.379	18.012	-74.641	1.00	84.05	L1	C

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ATOM   4118  CE  LYS   133     -22.792  18.960 -73.604  1.00 85.60      L1   C
ATOM   4119  NZ  LYS   133     -21.971  20.039 -74.232  1.00 85.73      L1   N
ATOM   4120  C   LYS   133     -28.044  17.979 -75.156  1.00 81.04      L1   C
ATOM   4121  O   LYS   133     -28.780  18.307 -76.087  1.00 81.66      L1   O
ATOM   4122  N   ALA   134     -28.469  17.900 -73.900  1.00 79.38      L1   N
ATOM   4123  CA  ALA   134     -29.825  18.294 -73.537  1.00 77.97      L1   C
ATOM   4124  CB  ALA   134     -30.769  17.107 -73.687  1.00 77.34      L1   C
ATOM   4125  C   ALA   134     -29.882  18.839 -72.110  1.00 77.09      L1   C
ATOM   4126  O   ALA   134     -29.587  18.126 -71.147  1.00 76.45      L1   O
ATOM   4127  N   THR   135     -30.262  20.107 -71.983  1.00 75.44      L1   N
ATOM   4128  CA  THR   135     -30.332  20.765 -70.682  1.00 73.38      L1   C
ATOM   4129  CB  THR   135     -29.333  21.937 -70.590  1.00 72.52      L1   C
ATOM   4130  OG1 THR   135     -28.001  21.455 -70.806  1.00 71.97      L1   O
ATOM   4131  CG2 THR   135     -29.414  22.596 -69.221  1.00 71.78      L1   C
ATOM   4132  C   THR   135     -31.724  21.320 -70.404  1.00 72.81      L1   C
ATOM   4133  O   THR   135     -32.213  22.186 -71.131  1.00 72.54      L1   O
ATOM   4134  N   LEU   136     -32.358  20.824 -69.346  1.00 71.97      L1   N
ATOM   4135  CA  LEU   136     -33.573  21.446 -68.833  1.00 71.28      L1   C
ATOM   4136  CB  LEU   136     -34.423  20.421 -68.076  1.00 70.86      L1   C
ATOM   4137  CG  LEU   136     -35.265  19.454 -68.911  1.00 70.08      L1   C
ATOM   4138  CD1 LEU   136     -34.370  18.679 -69.858  1.00 70.16      L1   C
ATOM   4139  CD2 LEU   136     -36.016  18.507 -67.992  1.00 69.02      L1   C
ATOM   4140  C   LEU   136     -33.208  22.601 -67.901  1.00 71.09      L1   C
ATOM   4141  O   LEU   136     -32.394  22.441 -66.987  1.00 71.21      L1   O
ATOM   4142  N   VAL   137     -33.808  23.764 -68.141  1.00 69.73      L1   N
ATOM   4143  CA  VAL   137     -33.551  24.943 -67.321  1.00 67.62      L1   C
ATOM   4144  CB  VAL   137     -33.154  26.153 -68.190  1.00 67.33      L1   C
ATOM   4145  CG1 VAL   137     -32.927  27.370 -67.311  1.00 65.83      L1   C
ATOM   4146  CG2 VAL   137     -31.901  25.830 -68.989  1.00 67.16      L1   C
ATOM   4147  C   VAL   137     -34.783  25.317 -66.509  1.00 66.67      L1   C
ATOM   4148  O   VAL   137     -35.820  25.671 -67.067  1.00 66.72      L1   O
ATOM   4149  N   CYS   138     -34.662  25.230 -65.188  1.00 65.71      L1   N
ATOM   4150  CA  CYS   138     -35.747  25.611 -64.290  1.00 63.89      L1   C
ATOM   4151  C   CYS   138     -35.386  26.893 -63.559  1.00 62.65      L1   C
ATOM   4152  O   CYS   138     -34.407  26.939 -62.812  1.00 62.18      L1   O
ATOM   4153  CB  CYS   138     -36.011  24.507 -63.268  1.00 64.16      L1   C
ATOM   4154  SG  CYS   138     -37.540  24.736 -62.308  1.00 64.86      L1   S
ATOM   4155  N   LEU   139     -36.182  27.933 -63.775  1.00 61.05      L1   N
ATOM   4156  CA  LEU   139     -35.925  29.224 -63.156  1.00 59.25      L1   C
ATOM   4157  CB  LEU   139     -35.920  30.313 -64.223  1.00 59.69      L1   C
ATOM   4158  CG  LEU   139     -34.894  30.039 -65.322  1.00 60.35      L1   C
ATOM   4159  CD1 LEU   139     -34.880  31.188 -66.302  1.00 61.08      L1   C
ATOM   4160  CD2 LEU   139     -33.518  29.850 -64.700  1.00 60.90      L1   C
ATOM   4161  C   LEU   139     -36.957  29.540 -62.078  1.00 57.87      L1   C
ATOM   4162  O   LEU   139     -38.167  29.469 -62.317  1.00 57.52      L1   O
ATOM   4163  N   ILE   140     -36.462  29.888 -60.893  1.00 55.44      L1   N
ATOM   4164  CA  ILE   140     -37.306  30.104 -59.723  1.00 52.95      L1   C
ATOM   4165  CB  ILE   140     -36.927  29.123 -58.601  1.00 52.69      L1   C
ATOM   4166  CG2 ILE   140     -38.000  29.115 -57.525  1.00 50.12      L1   C
ATOM   4167  CG1 ILE   140     -36.764  27.717 -59.185  1.00 51.28      L1   C
ATOM   4168  CD1 ILE   140     -35.977  26.778 -58.301  1.00 51.25      L1   C
ATOM   4169  C   ILE   140     -37.097  31.531 -59.230  1.00 52.13      L1   C
ATOM   4170  O   ILE   140     -35.985  31.912 -58.874  1.00 51.51      L1   O
ATOM   4171  N   SER   141     -38.165  32.321 -59.214  1.00 52.05      L1   N
ATOM   4172  CA  SER   141     -38.045  33.739 -58.892  1.00 51.89      L1   C
ATOM   4173  CB  SER   141     -38.136  34.569 -60.176  1.00 52.46      L1   C
ATOM   4174  OG  SER   141     -39.277  34.202 -60.934  1.00 55.32      L1   O
ATOM   4175  C   SER   141     -39.086  34.235 -57.889  1.00 50.72      L1   C
ATOM   4176  O   SER   141     -40.109  33.583 -57.661  1.00 50.55      L1   O
ATOM   4177  N   ASP   142     -38.802  35.387 -57.285  1.00 48.87      L1   N
ATOM   4178  CA  ASP   142     -39.777  36.103 -56.470  1.00 47.69      L1   C
ATOM   4179  CB  ASP   142     -40.938  36.587 -57.342  1.00 48.13      L1   C
ATOM   4180  CG  ASP   142     -40.522  37.671 -58.310  1.00 47.87      L1   C
ATOM   4181  OD1 ASP   142     -40.884  37.580 -59.500  1.00 49.98      L1   O
ATOM   4182  OD2 ASP   142     -39.832  38.615 -57.879  1.00 48.34      L1   O
ATOM   4183  C   ASP   142     -40.339  35.310 -55.298  1.00 46.46      L1   C
ATOM   4184  O   ASP   142     -41.513  35.458 -54.958  1.00 47.09      L1   O
ATOM   4185  N   PHE   143     -39.521  34.467 -54.678  1.00 45.07      L1   N
ATOM   4186  CA  PHE   143     -39.969  33.806 -53.462  1.00 43.57      L1   C
ATOM   4187  CB  PHE   143     -39.638  32.304 -53.483  1.00 45.13      L1   C
ATOM   4188  CG  PHE   143     -38.186  31.989 -53.738  1.00 46.48      L1   C
ATOM   4189  CD1 PHE   143     -37.335  31.673 -52.686  1.00 46.51      L1   C
ATOM   4190  CD2 PHE   143     -37.684  31.963 -55.032  1.00 46.16      L1   C
ATOM   4191  CE1 PHE   143     -36.010  31.334 -52.919  1.00 45.94      L1   C
ATOM   4192  CE2 PHE   143     -36.360  31.626 -55.273  1.00 46.80      L1   C
ATOM   4193  CZ  PHE   143     -35.522  31.310 -54.217  1.00 46.20      L1   C
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ATOM   4194  C   PHE   143     -39.377  34.465 -52.230  1.00 41.63      L1   C
ATOM   4195  O   PHE   143     -38.354  35.150 -52.311  1.00 39.49      L1   O
ATOM   4196  N   TYR   144     -40.050  34.280 -51.098  1.00 39.96      L1   N
ATOM   4197  CA  TYR   144     -39.596  34.030 -49.827  1.00 40.17      L1   C
ATOM   4198  CB  TYR   144     -39.838  36.343 -49.777  1.00 37.41      L1   C
ATOM   4199  CG  TYR   144     -39.404  36.963 -48.468  1.00 36.24      L1   C
ATOM   4200  CD1 TYR   144     -40.278  37.034 -47.386  1.00 33.84      L1   C
ATOM   4201  CE1 TYR   144     -39.873  37.566 -46.176  1.00 32.19      L1   C
ATOM   4202  CD2 TYR   144     -38.111  37.445 -48.300  1.00 34.26      L1   C
ATOM   4203  CE2 TYR   144     -37.698  37.978 -47.095  1.00 32.36      L1   C
ATOM   4204  CZ  TYR   144     -38.582  38.038 -46.038  1.00 31.01      L1   C
ATOM   4205  OH  TYR   144     -38.180  38.588 -44.844  1.00 32.81      L1   O
ATOM   4206  C   TYR   144     -40.340  34.158 -48.680  1.00 40.52      L1   C
ATOM   4207  O   TYR   144     -41.560  34.011 -48.728  1.00 42.86      L1   O
ATOM   4208  N   PRO   145     -39.615  33.740 -47.633  1.00 40.04      L1   N
ATOM   4209  CD  PRO   145     -40.238  33.151 -46.436  1.00 39.62      L1   C
ATOM   4210  CA  PRO   145     -38.156  33.852 -47.500  1.00 41.17      L1   C
ATOM   4211  CB  PRO   145     -37.875  33.348 -46.080  1.00 40.22      L1   C
ATOM   4212  CG  PRO   145     -39.073  32.561 -45.692  1.00 39.21      L1   C
ATOM   4213  C   PRO   145     -37.354  33.086 -48.555  1.00 43.02      L1   C
ATOM   4214  O   PRO   145     -37.912  32.315 -49.340  1.00 42.96      L1   O
ATOM   4215  N   GLY   146     -36.041  33.310 -48.569  1.00 44.40      L1   N
ATOM   4216  CA  GLY   146     -35.200  32.759 -49.618  1.00 46.46      L1   C
ATOM   4217  C   GLY   146     -34.687  31.351 -49.356  1.00 48.24      L1   C
ATOM   4218  O   GLY   146     -33.481  31.137 -49.233  1.00 47.83      L1   O
ATOM   4219  N   ALA   147     -35.603  30.389 -49.279  1.00 49.04      L1   N
ATOM   4220  CA  ALA   147     -35.233  28.992 -49.084  1.00 50.76      L1   C
ATOM   4221  CB  ALA   147     -35.225  28.657 -47.597  1.00 48.27      L1   C
ATOM   4222  C   ALA   147     -36.193  28.063 -49.825  1.00 52.27      L1   C
ATOM   4223  O   ALA   147     -37.394  28.034 -49.544  1.00 54.27      L1   O
ATOM   4224  N   VAL   148     -35.658  27.302 -50.773  1.00 52.98      L1   N
ATOM   4225  CA  VAL   148     -36.448  26.317 -51.504  1.00 53.70      L1   C
ATOM   4226  CB  VAL   148     -36.780  26.809 -52.935  1.00 52.32      L1   C
ATOM   4227  CG1 VAL   148     -37.632  28.061 -52.872  1.00 51.32      L1   C
ATOM   4228  CG2 VAL   148     -35.495  27.084 -53.698  1.00 50.98      L1   C
ATOM   4229  C   VAL   148     -35.704  24.990 -51.615  1.00 54.30      L1   C
ATOM   4230  O   VAL   148     -34.493  24.927 -51.415  1.00 53.75      L1   O
ATOM   4231  N   THR   149     -36.442  23.930 -51.928  1.00 55.67      L1   N
ATOM   4232  CA  THR   149     -35.835  22.680 -52.369  1.00 56.75      L1   C
ATOM   4233  CB  THR   149     -36.112  21.540 -51.373  1.00 56.63      L1   C
ATOM   4234  OG1 THR   149     -37.519  21.462 -51.115  1.00 55.86      L1   O
ATOM   4235  CG2 THR   149     -35.364  21.780 -50.071  1.00 55.91      L1   C
ATOM   4236  C   THR   149     -36.392  22.294 -53.734  1.00 58.18      L1   C
ATOM   4237  O   THR   149     -37.597  22.400 -53.977  1.00 58.23      L1   O
ATOM   4238  N   VAL   150     -35.510  21.849 -54.623  1.00 59.58      L1   N
ATOM   4239  CA  VAL   150     -35.909  21.510 -55.983  1.00 62.22      L1   C
ATOM   4240  CB  VAL   150     -35.024  22.235 -57.015  1.00 62.03      L1   C
ATOM   4241  CG1 VAL   150     -35.498  21.913 -58.422  1.00 62.67      L1   C
ATOM   4242  CG2 VAL   150     -35.063  23.732 -56.768  1.00 62.69      L1   C
ATOM   4243  C   VAL   150     -35.833  20.008 -56.248  1.00 63.79      L1   C
ATOM   4244  O   VAL   150     -34.762  19.401 -56.166  1.00 64.56      L1   O
ATOM   4245  N   ALA   151     -36.978  19.414 -56.566  1.00 64.58      L1   N
ATOM   4246  CA  ALA   151     -37.034  18.014 -56.964  1.00 65.49      L1   C
ATOM   4247  CB  ALA   151     -38.086  17.282 -56.144  1.00 64.19      L1   C
ATOM   4248  C   ALA   151     -37.364  17.913 -58.450  1.00 66.84      L1   C
ATOM   4249  O   ALA   151     -38.276  18.580 -58.939  1.00 67.66      L1   O
ATOM   4250  N   TRP   152     -36.613  17.084 -59.167  1.00 67.84      L1   N
ATOM   4251  CA  TRP   152     -36.882  16.842 -60.579  1.00 69.16      L1   C
ATOM   4252  CB  TRP   152     -35.579  16.851 -61.380  1.00 67.01      L1   C
ATOM   4253  CG  TRP   152     -34.968  18.213 -61.549  1.00 64.07      L1   C
ATOM   4254  CD2 TRP   152     -35.151  19.103 -62.658  1.00 62.91      L1   C
ATOM   4255  CE2 TRP   152     -34.345  20.235 -62.422  1.00 62.02      L1   C
ATOM   4256  CE3 TRP   152     -35.916  19.050 -63.829  1.00 62.20      L1   C
ATOM   4257  CD1 TRP   152     -34.088  18.827 -60.707  1.00 62.61      L1   C
ATOM   4258  NE1 TRP   152     -33.707  20.042 -61.225  1.00 61.78      L1   N
ATOM   4259  CZ2 TRP   152     -34.281  21.304 -63.313  1.00 61.34      L1   C
ATOM   4260  CZ3 TRP   152     -35.850  20.113 -64.714  1.00 61.43      L1   C
ATOM   4261  CH2 TRP   152     -35.038  21.224 -64.451  1.00 61.91      L1   C
ATOM   4262  C   TRP   152     -37.579  15.500 -60.751  1.00 71.64      L1   C
ATOM   4263  O   TRP   152     -37.339  14.564 -59.985  1.00 71.81      L1   O
ATOM   4264  N   LYS   153     -38.440  15.407 -61.759  1.00 73.83      L1   N
ATOM   4265  CA  LYS   153     -39.267  14.221 -61.939  1.00 76.53      L1   C
ATOM   4266  CB  LYS   153     -40.708  14.532 -61.520  1.00 77.51      L1   C
ATOM   4267  CG  LYS   153     -41.453  13.361 -60.900  1.00 78.87      L1   C
ATOM   4268  CD  LYS   153     -41.399  13.407 -59.376  1.00 79.45      L1   C
ATOM   4269  CE  LYS   153     -39.981  13.219 -58.855  1.00 80.16      L1   C
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ATOM	4270	NZ	LYS	153	-39.927	13.194	-57.366	1.00	79.77	L1	N
ATOM	4271	C	LYS	153	-39.243	13.737	-63.389	1.00	77.70	L1	C
ATOM	4272	O	LYS	153	-39.608	14.475	-64.304	1.00	77.96	L1	O
ATOM	4273	N	ALA	154	-38.809	12.495	-63.590	1.00	79.25	L1	N
ATOM	4274	CA	ALA	154	-38.815	11.880	-64.917	1.00	80.39	L1	C
ATOM	4275	CB	ALA	154	-37.575	11.013	-65.098	1.00	79.94	L1	C
ATOM	4276	C	ALA	154	-40.075	11.035	-65.085	1.00	81.09	L1	C
ATOM	4277	O	ALA	154	-40.234	10.008	-64.426	1.00	80.56	L1	O
ATOM	4278	N	ASP	155	-40.962	11.470	-65.976	1.00	82.58	L1	N
ATOM	4279	CA	ASP	155	-42.319	10.933	-66.034	1.00	84.19	L1	C
ATOM	4280	CB	ASP	155	-42.302	9.446	-66.414	1.00	85.20	L1	C
ATOM	4281	CG	ASP	155	-42.034	9.222	-67.893	1.00	86.13	L1	C
ATOM	4282	OD1	ASP	155	-42.635	9.939	-68.723	1.00	85.84	L1	O
ATOM	4283	OD2	ASP	155	-41.225	8.326	-68.225	1.00	86.10	L1	O
ATOM	4284	C	ASP	155	-42.981	11.105	-64.673	1.00	84.53	L1	C
ATOM	4285	O	ASP	155	-43.575	12.144	-64.388	1.00	84.90	L1	O
ATOM	4286	N	SER	156	-42.866	10.081	-63.834	1.00	84.79	L1	N
ATOM	4287	CA	SER	156	-43.383	10.136	-62.472	1.00	84.82	L1	C
ATOM	4288	CB	SER	156	-44.545	9.154	-62.310	1.00	85.15	L1	C
ATOM	4289	OG	SER	156	-45.527	9.358	-63.312	1.00	85.88	L1	O
ATOM	4290	C	SER	156	-42.270	9.777	-61.495	1.00	84.72	L1	C
ATOM	4291	O	SER	156	-42.374	10.029	-60.295	1.00	84.40	L1	O
ATOM	4292	N	SER	157	-41.205	9.182	-62.023	1.00	84.73	L1	N
ATOM	4293	CA	SER	157	-40.101	8.708	-61.198	1.00	84.69	L1	C
ATOM	4294	CB	SER	157	-39.301	7.635	-61.946	1.00	84.54	L1	C
ATOM	4295	OG	SER	157	-40.079	6.469	-62.163	1.00	84.37	L1	O
ATOM	4296	C	SER	157	-39.169	9.843	-60.796	1.00	84.54	L1	C
ATOM	4297	O	SER	157	-38.933	10.776	-61.565	1.00	83.92	L1	O
ATOM	4298	N	PRO	158	-38.631	9.773	-59.571	1.00	84.61	L1	N
ATOM	4299	CD	PRO	158	-39.055	8.832	-58.521	1.00	84.52	L1	C
ATOM	4300	CA	PRO	158	-37.633	10.728	-59.079	1.00	84.62	L1	C
ATOM	4301	CB	PRO	158	-37.393	10.286	-57.635	1.00	84.58	L1	C
ATOM	4302	CG	PRO	158	-38.625	9.522	-57.263	1.00	84.46	L1	C
ATOM	4303	C	PRO	158	-36.352	10.680	-59.909	1.00	84.68	L1	C
ATOM	4304	O	PRO	158	-35.918	9.607	-60.333	1.00	84.90	L1	O
ATOM	4305	N	VAL	159	-35.753	11.844	-60.141	1.00	84.33	L1	N
ATOM	4306	CA	VAL	159	-34.479	11.919	-60.847	1.00	83.83	L1	C
ATOM	4307	CB	VAL	159	-34.463	13.086	-61.860	1.00	84.04	L1	C
ATOM	4308	CG1	VAL	159	-33.084	13.210	-62.492	1.00	83.62	L1	C
ATOM	4309	CG2	VAL	159	-35.516	12.857	-62.931	1.00	83.67	L1	C
ATOM	4310	C	VAL	159	-33.328	12.110	-59.864	1.00	83.32	L1	C
ATOM	4311	O	VAL	159	-33.329	13.042	-59.060	1.00	83.25	L1	O
ATOM	4312	N	LYS	160	-32.345	11.221	-59.937	1.00	82.48	L1	N
ATOM	4313	CA	LYS	160	-31.199	11.284	-59.040	1.00	81.67	L1	C
ATOM	4314	CB	LYS	160	-30.533	9.905	-58.947	1.00	81.53	L1	C
ATOM	4315	CG	LYS	160	-31.408	8.839	-58.305	1.00	81.07	L1	C
ATOM	4316	CD	LYS	160	-31.257	7.491	-58.997	1.00	80.52	L1	C
ATOM	4317	CE	LYS	160	-32.249	6.480	-58.440	1.00	80.19	L1	C
ATOM	4318	NZ	LYS	160	-32.296	5.224	-59.240	1.00	80.25	L1	N
ATOM	4319	C	LYS	160	-30.180	12.328	-59.499	1.00	80.59	L1	C
ATOM	4320	O	LYS	160	-30.155	13.449	-58.990	1.00	80.48	L1	O
ATOM	4321	N	ALA	161	-29.352	11.958	-60.472	1.00	78.88	L1	N
ATOM	4322	CA	ALA	161	-28.194	12.764	-60.839	1.00	76.98	L1	C
ATOM	4323	CB	ALA	161	-27.015	11.855	-61.163	1.00	77.49	L1	C
ATOM	4324	C	ALA	161	-28.471	13.694	-62.014	1.00	75.33	L1	C
ATOM	4325	O	ALA	161	-29.578	13.719	-62.554	1.00	74.70	L1	O
ATOM	4326	N	GLY	162	-27.450	14.455	-62.400	1.00	73.68	L1	N
ATOM	4327	CA	GLY	162	-27.592	15.404	-63.490	1.00	71.48	L1	C
ATOM	4328	C	GLY	162	-28.113	16.746	-63.013	1.00	70.15	L1	C
ATOM	4329	O	GLY	162	-28.263	17.681	-63.802	1.00	69.01	L1	O
ATOM	4330	N	VAL	163	-28.383	16.839	-61.713	1.00	69.08	L1	N
ATOM	4331	CA	VAL	163	-29.013	18.020	-61.132	1.00	67.56	L1	C
ATOM	4332	CB	VAL	163	-29.963	17.627	-59.978	1.00	67.21	L1	C
ATOM	4333	CG1	VAL	163	-30.631	18.868	-59.406	1.00	66.89	L1	C
ATOM	4334	CG2	VAL	163	-31.006	16.642	-60.477	1.00	66.02	L1	C
ATOM	4335	C	VAL	163	-27.986	19.019	-60.601	1.00	66.76	L1	C
ATOM	4336	O	VAL	163	-27.092	18.659	-59.846	1.00	65.41	L1	O
ATOM	4337	N	GLU	164	-28.139	20.277	-61.000	1.00	66.54	L1	N
ATOM	4338	CA	GLU	164	-27.250	21.345	-60.551	1.00	66.37	L1	C
ATOM	4339	CB	GLU	164	-26.269	21.708	-61.667	1.00	67.88	L1	C
ATOM	4340	CG	GLU	164	-24.859	21.983	-61.192	1.00	68.70	L1	C
ATOM	4341	CD	GLU	164	-24.073	20.712	-60.971	1.00	70.76	L1	C
ATOM	4342	OE1	GLU	164	-23.934	20.298	-59.801	1.00	71.18	L1	O
ATOM	4343	OE2	GLU	164	-23.597	20.127	-61.969	1.00	71.13	L1	O
ATOM	4344	C	GLU	164	-28.076	22.577	-60.174	1.00	64.94	L1	C
ATOM	4345	O	GLU	164	-28.643	23.243	-61.041	1.00	64.32	L1	O

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ATOM   4346  N   THR   165     -28.140  22.876 -58.881  1.00 62.91      L1   N
ATOM   4347  CA  THR   165     -28.991  23.957 -58.391  1.00 61.09      L1   C
ATOM   4348  CB  THR   165     -30.086  23.418 -57.441  1.00 60.86      L1   C
ATOM   4349  OG1 THR   165     -30.918  22.485 -58.144  1.00 60.45      L1   O
ATOM   4350  CG2 THR   165     -30.947  24.560 -56.915  1.00 59.22      L1   C
ATOM   4351  C   THR   165     -28.192  25.025 -57.649  1.00 59.97      L1   C
ATOM   4352  O   THR   165     -27.360  24.714 -56.795  1.00 60.22      L1   O
ATOM   4353  N   THR   166     -28.455  26.286 -57.973  1.00 57.86      L1   N
ATOM   4354  CA  THR   166     -27.779  27.393 -57.312  1.00 57.52      L1   C
ATOM   4355  CB  THR   166     -27.989  28.715 -58.071  1.00 57.94      L1   C
ATOM   4356  OG1 THR   166     -29.386  29.037 -58.088  1.00 56.92      L1   O
ATOM   4357  CG2 THR   166     -27.471  28.599 -59.496  1.00 57.35      L1   C
ATOM   4358  C   THR   166     -28.276  27.600 -55.883  1.00 56.25      L1   C
ATOM   4359  O   THR   166     -29.303  27.052 -55.475  1.00 55.75      L1   O
ATOM   4360  N   THR   167     -27.536  28.400 -55.127  1.00 54.65      L1   N
ATOM   4361  CA  THR   167     -28.032  28.910 -53.860  1.00 54.27      L1   C
ATOM   4362  CB  THR   167     -26.877  29.393 -52.963  1.00 54.52      L1   C
ATOM   4363  OG1 THR   167     -26.172  30.451 -53.623  1.00 53.89      L1   O
ATOM   4364  CG2 THR   167     -25.909  28.250 -52.680  1.00 54.44      L1   C
ATOM   4365  C   THR   167     -28.949  30.089 -54.161  1.00 53.66      L1   C
ATOM   4366  O   THR   167     -28.730  30.829 -55.121  1.00 52.20      L1   O
ATOM   4367  N   PRO   168     -29.998  30.273 -53.351  1.00 54.17      L1   N
ATOM   4368  CD  PRO   168     -30.507  29.340 -52.331  1.00 53.81      L1   C
ATOM   4369  CA  PRO   168     -30.894  31.418 -53.544  1.00 53.90      L1   C
ATOM   4370  CB  PRO   168     -31.944  31.238 -52.450  1.00 53.59      L1   C
ATOM   4371  CG  PRO   168     -31.945  29.761 -52.185  1.00 54.66      L1   C
ATOM   4372  C   PRO   168     -30.153  32.748 -53.427  1.00 53.20      L1   C
ATOM   4373  O   PRO   168     -29.358  32.942 -52.511  1.00 54.37      L1   O
ATOM   4374  N   SER   169     -30.413  33.654 -54.363  1.00 52.86      L1   N
ATOM   4375  CA  SER   169     -29.782  34.971 -54.360  1.00 52.56      L1   C
ATOM   4376  CB  SER   169     -28.984  35.169 -55.647  1.00 51.77      L1   C
ATOM   4377  OG  SER   169     -29.775  34.846 -56.779  1.00 53.89      L1   O
ATOM   4378  C   SER   169     -30.821  36.083 -54.224  1.00 52.72      L1   C
ATOM   4379  O   SER   169     -31.943  35.962 -54.716  1.00 51.23      L1   O
ATOM   4380  N   LYS   170     -30.440  37.167 -53.556  1.00 53.95      L1   N
ATOM   4381  CA  LYS   170     -31.375  38.246 -53.266  1.00 55.76      L1   C
ATOM   4382  CB  LYS   170     -30.868  39.080 -52.086  1.00 57.25      L1   C
ATOM   4383  CG  LYS   170     -31.817  40.193 -51.657  1.00 59.84      L1   C
ATOM   4384  CD  LYS   170     -31.307  40.916 -50.417  1.00 60.88      L1   C
ATOM   4385  CE  LYS   170     -31.110  39.953 -49.254  1.00 63.31      L1   C
ATOM   4386  NZ  LYS   170     -32.375  39.253 -48.887  1.00 65.05      L1   N
ATOM   4387  C   LYS   170     -31.590  39.147 -54.478  1.00 55.79      L1   C
ATOM   4388  O   LYS   170     -30.646  39.745 -54.992  1.00 56.34      L1   O
ATOM   4389  N   GLN   171     -32.839  39.240 -54.927  1.00 55.98      L1   N
ATOM   4390  CA  GLN   171     -33.200  40.121 -56.032  1.00 56.08      L1   C
ATOM   4391  CB  GLN   171     -34.559  39.716 -56.610  1.00 54.88      L1   C
ATOM   4392  CG  GLN   171     -34.622  38.299 -57.149  1.00 56.49      L1   C
ATOM   4393  CD  GLN   171     -36.046  37.850 -57.439  1.00 56.91      L1   C
ATOM   4394  OE1 GLN   171     -36.271  36.805 -58.053  1.00 56.50      L1   O
ATOM   4395  NE2 GLN   171     -37.018  38.643 -56.995  1.00 56.59      L1   N
ATOM   4396  C   GLN   171     -33.263  41.575 -55.567  1.00 57.33      L1   C
ATOM   4397  O   GLN   171     -32.982  41.882 -54.407  1.00 58.32      L1   O
ATOM   4398  N   SER   172     -33.637  42.465 -56.481  1.00 57.40      L1   N
ATOM   4399  CA  SER   172     -33.736  43.889 -56.178  1.00 57.00      L1   C
ATOM   4400  CB  SER   172     -33.911  44.682 -57.473  1.00 57.43      L1   C
ATOM   4401  OG  SER   172     -34.948  44.123 -58.263  1.00 58.48      L1   O
ATOM   4402  C   SER   172     -34.900  44.183 -55.233  1.00 56.38      L1   C
ATOM   4403  O   SER   172     -34.776  45.001 -54.322  1.00 57.23      L1   O
ATOM   4404  N   ASN   173     -36.026  43.511 -55.454  1.00 54.81      L1   N
ATOM   4405  CA  ASN   173     -37.207  43.684 -54.613  1.00 52.81      L1   C
ATOM   4406  CB  ASN   173     -38.442  43.191 -55.356  1.00 53.01      L1   C
ATOM   4407  CG  ASN   173     -38.286  41.770 -55.848  1.00 54.06      L1   C
ATOM   4408  OD1 ASN   173     -37.432  41.027 -55.364  1.00 54.19      L1   O
ATOM   4409  ND2 ASN   173     -39.108  41.383 -56.816  1.00 54.46      L1   N
ATOM   4410  C   ASN   173     -37.062  42.919 -53.298  1.00 52.76      L1   C
ATOM   4411  O   ASN   173     -38.020  42.770 -52.540  1.00 51.62      L1   O
ATOM   4412  N   ASN   174     -35.858  42.423 -53.042  1.00 52.25      L1   N
ATOM   4413  CA  ASN   174     -35.560  41.753 -51.788  1.00 50.29      L1   C
ATOM   4414  CB  ASN   174     -35.930  42.661 -50.615  1.00 53.67      L1   C
ATOM   4415  CG  ASN   174     -34.994  43.848 -50.488  1.00 55.99      L1   C
ATOM   4416  OD1 ASN   174     -33.839  43.694 -50.091  1.00 58.37      L1   O
ATOM   4417  ND2 ASN   174     -35.483  45.037 -50.830  1.00 56.42      L1   N
ATOM   4418  C   ASN   174     -36.253  40.402 -51.654  1.00 48.00      L1   C
ATOM   4419  O   ASN   174     -36.163  39.755 -50.611  1.00 45.22      L1   O
ATOM   4420  N   LYS   175     -36.944  39.977 -52.709  1.00 46.20      L1   N
ATOM   4421  CA  LYS   175     -37.315  38.574 -52.844  1.00 45.58      L1   C
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ATOM   4422  CB  LYS   175     -38.486  38.410 -53.819  1.00 44.39      L1   C
ATOM   4423  CG  LYS   175     -39.801  39.024 -53.348  1.00 44.61      L1   C
ATOM   4424  CD  LYS   175     -40.991  38.369 -54.048  1.00 44.52      L1   C
ATOM   4425  CE  LYS   175     -42.325  38.951 -53.593  1.00 44.60      L1   C
ATOM   4426  NZ  LYS   175     -42.567  40.326 -54.117  1.00 43.73      L1   N
ATOM   4427  C   LYS   175     -36.095  37.812 -53.359  1.00 46.51      L1   C
ATOM   4428  O   LYS   175     -35.042  38.404 -53.590  1.00 46.31      L1   O
ATOM   4429  N   TYR   176     -36.229  36.503 -53.535  1.00 47.57      L1   N
ATOM   4430  CA  TYR   176     -35.092  35.680 -53.940  1.00 49.06      L1   C
ATOM   4431  CB  TYR   176     -34.766  34.665 -52.847  1.00 47.73      L1   C
ATOM   4432  CG  TYR   176     -34.098  35.252 -51.631  1.00 48.77      L1   C
ATOM   4433  CD1 TYR   176     -32.771  34.961 -51.345  1.00 49.57      L1   C
ATOM   4434  CE1 TYR   176     -32.160  35.455 -50.210  1.00 51.42      L1   C
ATOM   4435  CD2 TYR   176     -34.799  36.066 -50.743  1.00 48.10      L1   C
ATOM   4436  CE2 TYR   176     -34.193  36.568 -49.600  1.00 48.23      L1   C
ATOM   4437  CZ  TYR   176     -32.871  36.255 -49.339  1.00 50.96      L1   C
ATOM   4438  OH  TYR   176     -32.247  36.721 -48.202  1.00 51.85      L1   O
ATOM   4439  C   TYR   176     -35.296  34.936 -55.260  1.00 50.28      L1   C
ATOM   4440  O   TYR   176     -36.421  34.615 -55.648  1.00 50.45      L1   O
ATOM   4441  N   ALA   177     -34.186  34.658 -55.937  1.00 50.51      L1   N
ATOM   4442  CA  ALA   177     -34.201  33.888 -57.170  1.00 52.09      L1   C
ATOM   4443  CB  ALA   177     -33.844  34.784 -58.352  1.00 50.29      L1   C
ATOM   4444  C   ALA   177     -33.212  32.731 -57.074  1.00 53.43      L1   C
ATOM   4445  O   ALA   177     -32.246  32.784 -56.307  1.00 52.69      L1   O
ATOM   4446  N   ALA   178     -33.459  31.689 -57.861  1.00 54.62      L1   N
ATOM   4447  CA  ALA   178     -32.601  30.513 -57.875  1.00 55.83      L1   C
ATOM   4448  CB  ALA   178     -33.021  29.550 -56.773  1.00 55.75      L1   C
ATOM   4449  C   ALA   178     -32.690  29.825 -59.230  1.00 56.94      L1   C
ATOM   4450  O   ALA   178     -33.667  29.995 -59.958  1.00 56.28      L1   O
ATOM   4451  N   SER   179     -31.662  29.052 -59.563  1.00 58.47      L1   N
ATOM   4452  CA  SER   179     -31.650  28.281 -60.800  1.00 59.49      L1   C
ATOM   4453  CB  SER   179     -30.569  28.812 -61.742  1.00 58.95      L1   C
ATOM   4454  OG  SER   179     -30.847  30.142 -62.130  1.00 60.29      L1   O
ATOM   4455  C   SER   179     -31.395  26.805 -60.520  1.00 60.01      L1   C
ATOM   4456  O   SER   179     -30.626  26.455 -59.625  1.00 60.02      L1   O
ATOM   4457  N   SER   180     -32.050  25.943 -61.290  1.00 60.42      L1   N
ATOM   4458  CA  SER   180     -31.746  24.520 -61.264  1.00 61.35      L1   C
ATOM   4459  CB  SER   180     -32.836  23.755 -60.509  1.00 60.80      L1   C
ATOM   4460  OG  SER   180     -32.458  22.404 -60.306  1.00 59.95      L1   O
ATOM   4461  C   SER   180     -31.637  23.994 -62.690  1.00 62.30      L1   C
ATOM   4462  O   SER   180     -32.556  24.150 -63.493  1.00 61.67      L1   O
ATOM   4463  N   TYR   181     -30.505  23.376 -63.003  1.00 63.90      L1   N
ATOM   4464  CA  TYR   181     -30.283  22.822 -64.332  1.00 64.83      L1   C
ATOM   4465  CB  TYR   181     -28.971  23.354 -64.904  1.00 63.82      L1   C
ATOM   4466  CG  TYR   181     -28.972  24.844 -65.154  1.00 64.75      L1   C
ATOM   4467  CD1 TYR   181     -28.789  25.744 -64.112  1.00 64.15      L1   C
ATOM   4468  CE1 TYR   181     -28.756  27.107 -64.346  1.00 65.10      L1   C
ATOM   4469  CD2 TYR   181     -29.127  25.350 -66.439  1.00 63.88      L1   C
ATOM   4470  CE2 TYR   181     -29.096  26.708 -66.682  1.00 63.92      L1   C
ATOM   4471  CZ  TYR   181     -28.908  27.583 -65.634  1.00 65.26      L1   C
ATOM   4472  OH  TYR   181     -28.850  28.934 -65.881  1.00 65.71      L1   O
ATOM   4473  C   TYR   181     -30.244  21.302 -64.290  1.00 66.15      L1   C
ATOM   4474  O   TYR   181     -29.520  20.712 -63.486  1.00 66.57      L1   O
ATOM   4475  N   LEU   182     -31.030  20.668 -65.152  1.00 67.55      L1   N
ATOM   4476  CA  LEU   182     -30.982  19.217 -65.284  1.00 69.82      L1   C
ATOM   4477  CB  LEU   182     -32.388  18.618 -65.181  1.00 69.84      L1   C
ATOM   4478  CG  LEU   182     -32.470  17.116 -65.479  1.00 70.75      L1   C
ATOM   4479  CD1 LEU   182     -31.428  16.367 -64.653  1.00 70.11      L1   C
ATOM   4480  CD2 LEU   182     -33.870  16.602 -65.179  1.00 70.49      L1   C
ATOM   4481  C   LEU   182     -30.351  18.814 -66.613  1.00 70.96      L1   C
ATOM   4482  O   LEU   182     -30.884  19.117 -67.683  1.00 71.15      L1   O
ATOM   4483  N   SER   183     -29.214  18.131 -66.536  1.00 71.50      L1   N
ATOM   4484  CA  SER   183     -28.553  17.619 -67.728  1.00 71.86      L1   C
ATOM   4485  CB  SER   183     -27.033  17.734 -67.584  1.00 71.20      L1   C
ATOM   4486  OG  SER   183     -26.626  19.092 -67.550  1.00 70.56      L1   O
ATOM   4487  C   SER   183     -28.937  16.166 -67.964  1.00 72.44      L1   C
ATOM   4488  O   SER   183     -28.869  15.339 -67.057  1.00 72.17      L1   O
ATOM   4489  N   LEU   184     -29.353  15.865 -69.188  1.00 73.34      L1   N
ATOM   4490  CA  LEU   184     -29.605  14.489 -69.591  1.00 74.57      L1   C
ATOM   4491  CB  LEU   184     -31.074  14.125 -69.346  1.00 74.23      L1   C
ATOM   4492  CG  LEU   184     -32.149  15.157 -69.697  1.00 73.95      L1   C
ATOM   4493  CD1 LEU   184     -32.186  15.392 -71.194  1.00 74.22      L1   C
ATOM   4494  CD2 LEU   184     -33.496  14.657 -69.211  1.00 73.72      L1   C
ATOM   4495  C   LEU   184     -29.243  14.299 -71.060  1.00 75.82      L1   C
ATOM   4496  O   LEU   184     -29.060  15.273 -71.791  1.00 75.56      L1   O
ATOM   4497  N   THR   185     -29.124  13.047 -71.489  1.00 77.26      L1   N
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ATOM   4498  CA  THR   185     -28.787  12.766 -72.878  1.00 78.90      L1   C
ATOM   4499  CB  THR   185     -28.252  11.321 -73.056  1.00 79.12      L1   C
ATOM   4500  OG1 THR   185     -29.259  10.378 -72.664  1.00 79.12      L1   O
ATOM   4501  CG2 THR   185     -27.001  11.109 -72.210  1.00 78.20      L1   C
ATOM   4502  C   THR   185     -30.023  12.955 -73.748  1.00 79.76      L1   C
ATOM   4503  O   THR   185     -31.148  12.692 -73.313  1.00 79.50      L1   O
ATOM   4504  N   PRO   186     -29.828  13.420 -74.992  1.00 80.66      L1   N
ATOM   4505  CD  PRO   186     -28.522  13.709 -75.608  1.00 80.61      L1   C
ATOM   4506  CA  PRO   186     -30.939  13.701 -75.908  1.00 81.88      L1   C
ATOM   4507  CB  PRO   186     -30.240  14.073 -77.215  1.00 81.41      L1   C
ATOM   4508  CG  PRO   186     -28.889  14.557 -76.788  1.00 80.92      L1   C
ATOM   4509  C   PRO   186     -31.843  12.485 -76.068  1.00 83.38      L1   C
ATOM   4510  O   PRO   186     -33.050  12.612 -76.280  1.00 83.33      L1   O
ATOM   4511  N   GLU   187     -31.242  11.305 -75.955  1.00 84.89      L1   N
ATOM   4512  CA  GLU   187     -31.967  10.050 -76.094  1.00 86.26      L1   C
ATOM   4513  CB  GLU   187     -30.988   8.872 -76.050  1.00 86.44      L1   C
ATOM   4514  CG  GLU   187     -29.896   8.924 -77.117  1.00 87.62      L1   C
ATOM   4515  CD  GLU   187     -28.843   9.992 -76.843  1.00 88.12      L1   C
ATOM   4516  OE1 GLU   187     -28.057   9.823 -75.886  1.00 88.36      L1   O
ATOM   4517  OE2 GLU   187     -28.800  10.996 -77.587  1.00 87.80      L1   O
ATOM   4518  C   GLU   187     -33.007   9.900 -74.987  1.00 86.75      L1   C
ATOM   4519  O   GLU   187     -34.201   9.762 -75.260  1.00 87.11      L1   O
ATOM   4520  N   GLN   188     -32.551   9.938 -73.738  1.00 86.85      L1   N
ATOM   4521  CA  GLN   188     -33.449   9.776 -72.601  1.00 86.86      L1   C
ATOM   4522  CB  GLN   188     -32.645   9.606 -71.310  1.00 86.25      L1   C
ATOM   4523  CG  GLN   188     -31.628  10.699 -71.052  1.00 86.47      L1   C
ATOM   4524  CD  GLN   188     -30.730  10.378 -69.873  1.00 86.63      L1   C
ATOM   4525  OE1 GLN   188     -29.527  10.641 -69.901  1.00 86.47      L1   O
ATOM   4526  NE2 GLN   188     -31.312   9.802 -68.828  1.00 86.42      L1   N
ATOM   4527  C   GLN   188     -34.415  10.951 -72.466  1.00 87.03      L1   C
ATOM   4528  O   GLN   188     -35.455  10.836 -71.817  1.00 86.87      L1   O
ATOM   4529  N   TRP   189     -34.072  12.078 -73.085  1.00 86.98      L1   N
ATOM   4530  CA  TRP   189     -34.980  13.220 -73.142  1.00 87.11      L1   C
ATOM   4531  CB  TRP   189     -34.300  14.398 -73.847  1.00 87.05      L1   C
ATOM   4532  CG  TRP   189     -35.253  15.455 -74.337  1.00 87.49      L1   C
ATOM   4533  CD2 TRP   189     -36.222  16.172 -73.558  1.00 87.50      L1   C
ATOM   4534  CE2 TRP   189     -36.888  17.056 -74.430  1.00 87.06      L1   C
ATOM   4535  CE3 TRP   189     -36.590  16.151 -72.208  1.00 87.12      L1   C
ATOM   4536  CD1 TRP   189     -35.371  15.923 -75.615  1.00 87.14      L1   C
ATOM   4537  NE1 TRP   189     -36.351  16.885 -75.679  1.00 87.03      L1   N
ATOM   4538  CZ2 TRP   189     -37.900  17.911 -73.998  1.00 87.27      L1   C
ATOM   4539  CZ3 TRP   189     -37.597  17.000 -71.781  1.00 87.07      L1   C
ATOM   4540  CH2 TRP   189     -38.239  17.868 -72.673  1.00 87.37      L1   C
ATOM   4541  C   TRP   189     -36.256  12.839 -73.886  1.00 87.39      L1   C
ATOM   4542  O   TRP   189     -37.366  13.094 -73.416  1.00 86.90      L1   O
ATOM   4543  N   LYS   190     -36.086  12.216 -75.047  1.00 88.13      L1   N
ATOM   4544  CA  LYS   190     -37.209  11.865 -75.910  1.00 88.58      L1   C
ATOM   4545  CB  LYS   190     -36.711  11.657 -77.343  1.00 88.28      L1   C
ATOM   4546  CG  LYS   190     -35.788  12.759 -77.847  1.00 87.78      L1   C
ATOM   4547  CD  LYS   190     -36.571  13.950 -78.371  1.00 87.40      L1   C
ATOM   4548  CE  LYS   190     -37.307  13.602 -79.657  1.00 86.94      L1   C
ATOM   4549  NZ  LYS   190     -38.075  14.758 -80.198  1.00 86.22      L1   N
ATOM   4550  C   LYS   190     -37.917  10.600 -75.423  1.00 88.94      L1   C
ATOM   4551  O   LYS   190     -39.078  10.361 -75.756  1.00 88.76      L1   O
ATOM   4552  N   SER   191     -37.212   9.797 -74.631  1.00 89.21      L1   N
ATOM   4553  CA  SER   191     -37.719   8.494 -74.213  1.00 89.73      L1   C
ATOM   4554  CB  SER   191     -36.556   7.586 -73.801  1.00 90.11      L1   C
ATOM   4555  OG  SER   191     -35.907   8.073 -72.639  1.00 90.61      L1   O
ATOM   4556  C   SER   191     -38.728   8.582 -73.068  1.00 89.98      L1   C
ATOM   4557  O   SER   191     -39.285   7.567 -72.645  1.00 89.77      L1   O
ATOM   4558  N   HIS   192     -38.957   9.790 -72.561  1.00 90.04      L1   N
ATOM   4559  CA  HIS   192     -39.911   9.986 -71.474  1.00 89.79      L1   C
ATOM   4560  CB  HIS   192     -39.214  10.625 -70.268  1.00 89.58      L1   C
ATOM   4561  CG  HIS   192     -38.255   9.712 -69.569  1.00 89.63      L1   C
ATOM   4562  CD2 HIS   192     -36.938   9.472 -69.776  1.00 89.30      L1   C
ATOM   4563  ND1 HIS   192     -38.625   8.915 -68.506  1.00 89.88      L1   N
ATOM   4564  CE1 HIS   192     -37.578   8.225 -68.089  1.00 89.37      L1   C
ATOM   4565  NE2 HIS   192     -36.541   8.545 -68.843  1.00 89.09      L1   N
ATOM   4566  C   HIS   192     -41.093  10.850 -71.906  1.00 89.78      L1   C
ATOM   4567  O   HIS   192     -40.961  11.703 -72.785  1.00 89.72      L1   O
ATOM   4568  N   ARG   193     -42.246  10.620 -71.282  1.00 89.66      L1   N
ATOM   4569  CA  ARG   193     -43.456  11.380 -71.583  1.00 89.47      L1   C
ATOM   4570  CB  ARG   193     -44.650  10.806 -70.812  1.00 90.32      L1   C
ATOM   4571  CG  ARG   193     -45.282   9.593 -71.468  1.00 91.88      L1   C
ATOM   4572  CD  ARG   193     -45.834   9.961 -72.836  1.00 93.71      L1   C
ATOM   4573  NE  ARG   193     -46.143   8.787 -73.648  1.00 94.98      L1   N
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ATOM	4574	CZ	ARG	193	-46.579	8.844	-74.904	1.00	95.50	L1	C
ATOM	4575	NH1	ARG	193	-46.834	7.725	-75.570	1.00	95.39	L1	N
ATOM	4576	NH2	ARG	193	-46.761	10.020	-75.492	1.00	95.47	L1	N
ATOM	4577	C	ARG	193	-43.300	12.862	-71.251	1.00	88.54	L1	C
ATOM	4578	O	ARG	193	-43.756	13.724	-72.004	1.00	88.43	L1	O
ATOM	4579	N	SER	194	-42.658	13.150	-70.121	1.00	87.52	L1	N
ATOM	4580	CA	SER	194	-42.402	14.528	-69.707	1.00	85.86	L1	C
ATOM	4581	CB	SER	194	-43.723	15.262	-69.446	1.00	85.32	L1	C
ATOM	4582	OG	SER	194	-44.394	14.726	-68.318	1.00	83.93	L1	O
ATOM	4583	C	SER	194	-41.537	14.584	-68.450	1.00	84.95	L1	C
ATOM	4584	O	SER	194	-41.534	13.653	-67.643	1.00	84.18	L1	O
ATOM	4585	N	TYR	195	-40.803	15.684	-68.298	1.00	83.96	L1	N
ATOM	4586	CA	TYR	195	-40.034	15.947	-67.085	1.00	82.58	L1	C
ATOM	4587	CB	TYR	195	-38.584	16.284	-67.439	1.00	82.92	L1	C
ATOM	4588	CG	TYR	195	-37.729	15.078	-67.748	1.00	83.66	L1	C
ATOM	4589	CD1	TYR	195	-36.938	14.495	-66.765	1.00	83.87	L1	C
ATOM	4590	CE1	TYR	195	-36.145	13.398	-67.042	1.00	84.28	L1	C
ATOM	4591	CD2	TYR	195	-37.703	14.526	-69.023	1.00	83.62	L1	C
ATOM	4592	CE2	TYR	195	-36.911	13.428	-69.311	1.00	83.74	L1	C
ATOM	4593	CZ	TYR	195	-36.134	12.869	-68.317	1.00	84.20	L1	C
ATOM	4594	OH	TYR	195	-35.332	11.786	-68.600	1.00	84.08	L1	O
ATOM	4595	C	TYR	195	-40.646	17.101	-66.300	1.00	81.49	L1	C
ATOM	4596	O	TYR	195	-41.169	18.050	-66.886	1.00	81.16	L1	O
ATOM	4597	N	SER	196	-40.576	17.016	-64.973	1.00	80.07	L1	N
ATOM	4598	CA	SER	196	-41.152	18.040	-64.106	1.00	78.68	L1	C
ATOM	4599	CB	SER	196	-42.303	17.454	-63.284	1.00	79.05	L1	C
ATOM	4600	OG	SER	196	-43.396	17.099	-64.113	1.00	80.30	L1	O
ATOM	4601	C	SER	196	-40.127	18.660	-63.159	1.00	77.25	L1	C
ATOM	4602	O	SER	196	-39.337	17.952	-62.530	1.00	76.71	L1	O
ATOM	4603	N	CYS	197	-40.153	19.987	-63.063	1.00	75.59	L1	N
ATOM	4604	CA	CYS	197	-39.390	20.701	-62.046	1.00	73.09	L1	C
ATOM	4605	C	CYS	197	-40.296	21.060	-60.876	1.00	72.59	L1	C
ATOM	4606	O	CYS	197	-41.221	21.859	-61.020	1.00	72.65	L1	O
ATOM	4607	CB	CYS	197	-38.786	21.979	-62.628	1.00	71.16	L1	C
ATOM	4608	SG	CYS	197	-37.815	22.934	-61.416	1.00	69.32	L1	S
ATOM	4609	N	GLN	198	-40.027	20.466	-59.718	1.00	71.96	L1	N
ATOM	4610	CA	GLN	198	-40.866	20.671	-58.542	1.00	71.74	L1	C
ATOM	4611	CB	GLN	198	-41.277	19.321	-57.952	1.00	73.01	L1	C
ATOM	4612	CG	GLN	198	-41.917	18.376	-58.954	1.00	74.84	L1	C
ATOM	4613	CD	GLN	198	-42.165	16.997	-58.373	1.00	76.39	L1	C
ATOM	4614	OE1	GLN	198	-41.697	16.676	-57.277	1.00	75.67	L1	O
ATOM	4615	NE2	GLN	198	-42.903	16.171	-59.107	1.00	77.19	L1	N
ATOM	4616	C	GLN	198	-40.140	21.493	-57.479	1.00	70.00	L1	C
ATOM	4617	O	GLN	198	-39.145	21.045	-56.905	1.00	70.10	L1	O
ATOM	4618	N	VAL	199	-40.647	22.694	-57.217	1.00	67.33	L1	N
ATOM	4619	CA	VAL	199	-40.060	23.570	-56.210	1.00	64.63	L1	C
ATOM	4620	CB	VAL	199	-39.894	25.006	-56.750	1.00	63.62	L1	C
ATOM	4621	CG1	VAL	199	-39.232	25.880	-55.700	1.00	62.58	L1	C
ATOM	4622	CG2	VAL	199	-39.072	24.990	-58.029	1.00	62.84	L1	C
ATOM	4623	C	VAL	199	-40.942	23.618	-54.969	1.00	63.15	L1	C
ATOM	4624	O	VAL	199	-42.132	23.908	-55.059	1.00	62.70	L1	O
ATOM	4625	N	THR	200	-40.359	23.331	-53.810	1.00	62.14	L1	N
ATOM	4626	CA	THR	200	-41.100	23.436	-52.558	1.00	62.84	L1	C
ATOM	4627	CB	THR	200	-40.974	22.147	-51.712	1.00	63.21	L1	C
ATOM	4628	OG1	THR	200	-41.466	21.029	-52.461	1.00	63.77	L1	O
ATOM	4629	CG2	THR	200	-41.785	22.272	-50.429	1.00	62.81	L1	C
ATOM	4630	C	THR	200	-40.611	24.623	-51.732	1.00	62.22	L1	C
ATOM	4631	O	THR	200	-39.409	24.821	-51.559	1.00	62.43	L1	O
ATOM	4632	N	HIS	201	-41.558	25.409	-51.229	1.00	61.86	L1	N
ATOM	4633	CA	HIS	201	-41.251	26.622	-50.481	1.00	61.76	L1	C
ATOM	4634	CB	HIS	201	-41.224	27.826	-51.428	1.00	61.02	L1	C
ATOM	4635	CG	HIS	201	-41.179	29.150	-50.729	1.00	59.75	L1	C
ATOM	4636	CD2	HIS	201	-40.135	29.901	-50.306	1.00	59.40	L1	C
ATOM	4637	ND1	HIS	201	-42.316	29.863	-50.415	1.00	59.04	L1	N
ATOM	4638	CE1	HIS	201	-41.974	30.998	-49.830	1.00	58.23	L1	C
ATOM	4639	NE2	HIS	201	-40.657	31.046	-49.752	1.00	58.22	L1	N
ATOM	4640	C	HIS	201	-42.299	26.843	-49.398	1.00	62.27	L1	C
ATOM	4641	O	HIS	201	-43.491	26.934	-49.690	1.00	62.33	L1	O
ATOM	4642	N	GLU	203	-41.849	26.928	-48.149	1.00	63.08	L1	N
ATOM	4643	CA	GLU	203	-42.749	27.081	-47.009	1.00	64.31	L1	C
ATOM	4644	CB	GLU	203	-43.440	28.447	-47.050	1.00	64.10	L1	C
ATOM	4645	CG	GLU	203	-42.490	29.628	-46.933	1.00	64.37	L1	C
ATOM	4646	CD	GLU	203	-41.781	29.682	-45.593	1.00	66.35	L1	C
ATOM	4647	OE1	GLU	203	-40.533	29.779	-45.584	1.00	66.23	L1	O
ATOM	4648	OE2	GLU	203	-42.471	29.629	-44.549	1.00	66.40	L1	O
ATOM	4649	C	GLU	203	-43.800	25.978	-46.978	1.00	65.33	L1	C

ATOM	4650	O	GLU	203	-44.909	26.183	-46.490	1.00	65.44	L1	O
ATOM	4651	N	GLY	203	-43.448	24.810	-47.507	1.00	66.98	L1	N
ATOM	4652	CA	GLY	203	-44.328	23.660	-47.415	1.00	69.40	L1	C
ATOM	4653	C	GLY	203	-45.215	23.457	-48.631	1.00	71.29	L1	C
ATOM	4654	O	GLY	203	-45.869	22.422	-48.761	1.00	71.55	L1	O
ATOM	4655	N	SER	204	-45.240	24.440	-49.523	1.00	72.19	L1	N
ATOM	4656	CA	SER	204	-46.079	24.371	-50.711	1.00	73.00	L1	C
ATOM	4657	CB	SER	204	-46.931	25.638	-50.821	1.00	74.04	L1	C
ATOM	4658	OG	SER	204	-47.605	25.699	-52.068	1.00	75.31	L1	O
ATOM	4659	C	SER	204	-45.237	24.200	-51.974	1.00	73.53	L1	C
ATOM	4660	O	SER	204	-44.322	24.983	-52.231	1.00	74.09	L1	O
ATOM	4661	N	THR	205	-45.555	23.174	-52.760	1.00	73.50	L1	N
ATOM	4662	CA	THR	205	-44.801	22.864	-53.971	1.00	72.84	L1	C
ATOM	4663	CB	THR	205	-44.781	21.345	-54.243	1.00	73.16	L1	C
ATOM	4664	OG1	THR	205	-44.142	20.669	-53.153	1.00	72.97	L1	O
ATOM	4665	CG2	THR	205	-44.031	21.044	-55.533	1.00	73.27	L1	C
ATOM	4666	C	THR	205	-45.392	23.558	-55.193	1.00	72.59	L1	C
ATOM	4667	O	THR	205	-46.586	23.445	-55.465	1.00	72.07	L1	O
ATOM	4668	N	VAL	206	-44.548	24.280	-55.923	1.00	72.50	L1	N
ATOM	4669	CA	VAL	206	-44.934	24.850	-57.208	1.00	73.02	L1	C
ATOM	4670	CB	VAL	206	-44.562	26.341	-57.302	1.00	72.13	L1	C
ATOM	4671	CG1	VAL	206	-44.947	26.890	-58.666	1.00	70.61	L1	C
ATOM	4672	CG2	VAL	206	-45.257	27.117	-56.201	1.00	71.61	L1	C
ATOM	4673	C	VAL	206	-44.207	24.092	-58.310	1.00	74.77	L1	C
ATOM	4674	O	VAL	206	-42.994	23.893	-58.240	1.00	74.80	L1	O
ATOM	4675	N	GLU	207	-44.952	23.671	-59.326	1.00	76.39	L1	N
ATOM	4676	CA	GLU	207	-44.438	22.716	-60.298	1.00	77.74	L1	C
ATOM	4677	CB	GLU	207	-45.033	21.335	-60.011	1.00	78.83	L1	C
ATOM	4678	CG	GLU	207	-44.608	20.244	-60.973	1.00	81.28	L1	C
ATOM	4679	CD	GLU	207	-45.153	18.886	-60.574	1.00	82.80	L1	C
ATOM	4680	OE1	GLU	207	-45.501	18.710	-59.386	1.00	84.12	L1	O
ATOM	4681	OE2	GLU	207	-45.235	17.995	-61.445	1.00	83.95	L1	O
ATOM	4682	C	GLU	207	-44.744	23.126	-61.735	1.00	77.99	L1	C
ATOM	4683	O	GLU	207	-45.837	23.603	-62.040	1.00	77.18	L1	O
ATOM	4684	N	LYS	208	-43.765	22.941	-62.615	1.00	78.59	L1	N
ATOM	4685	CA	LYS	208	-43.973	23.134	-64.043	1.00	79.79	L1	C
ATOM	4686	CB	LYS	208	-43.289	24.418	-64.515	1.00	79.80	L1	C
ATOM	4687	CG	LYS	208	-43.911	25.692	-63.961	1.00	80.02	L1	C
ATOM	4688	CD	LYS	208	-45.400	25.764	-64.266	1.00	79.10	L1	C
ATOM	4689	CE	LYS	208	-45.956	27.147	-63.964	1.00	79.03	L1	C
ATOM	4690	NZ	LYS	208	-45.344	28.191	-64.835	1.00	77.78	L1	N
ATOM	4691	C	LYS	208	-43.433	21.941	-64.823	1.00	80.92	L1	C
ATOM	4692	O	LYS	208	-42.413	21.356	-64.454	1.00	80.76	L1	O
ATOM	4693	N	THR	209	-44.125	21.584	-65.900	1.00	82.27	L1	N
ATOM	4694	CA	THR	209	-43.779	20.402	-66.679	1.00	83.34	L1	C
ATOM	4695	CB	THR	209	-44.889	19.335	-66.582	1.00	83.02	L1	C
ATOM	4696	OG1	THR	209	-45.127	19.012	-65.207	1.00	82.81	L1	O
ATOM	4697	CG2	THR	209	-44.482	18.071	-67.327	1.00	82.62	L1	C
ATOM	4698	C	THR	209	-43.566	20.749	-68.151	1.00	85.00	L1	C
ATOM	4699	O	THR	209	-44.209	21.653	-68.687	1.00	85.03	L1	O
ATOM	4700	N	VAL	210	-42.652	20.029	-68.793	1.00	86.55	L1	N
ATOM	4701	CA	VAL	210	-42.468	20.124	-70.236	1.00	87.92	L1	C
ATOM	4702	CB	VAL	210	-41.190	20.918	-70.590	1.00	87.63	L1	C
ATOM	4703	CG1	VAL	210	-41.310	22.347	-70.085	1.00	87.34	L1	C
ATOM	4704	CG2	VAL	210	-39.970	20.240	-69.988	1.00	87.16	L1	C
ATOM	4705	C	VAL	210	-42.372	18.728	-70.848	1.00	89.44	L1	C
ATOM	4706	O	VAL	210	-41.880	17.794	-70.212	1.00	89.28	L1	O
ATOM	4707	N	ALA	211	-42.851	18.591	-72.081	1.00	91.11	L1	N
ATOM	4708	CA	ALA	211	-42.826	17.311	-72.782	1.00	92.64	L1	C
ATOM	4709	CB	ALA	211	-44.206	17.004	-73.356	1.00	92.29	L1	C
ATOM	4710	C	ALA	211	-41.785	17.331	-73.897	1.00	93.85	L1	C
ATOM	4711	O	ALA	211	-41.449	18.392	-74.424	1.00	94.16	L1	O
ATOM	4712	N	PRO	212	-41.265	16.151	-74.273	1.00	95.11	L1	N
ATOM	4713	CD	PRO	212	-41.685	14.830	-73.774	1.00	95.23	L1	C
ATOM	4714	CA	PRO	212	-40.227	16.043	-75.306	1.00	95.91	L1	C
ATOM	4715	CB	PRO	212	-39.882	14.553	-75.318	1.00	95.62	L1	C
ATOM	4716	CG	PRO	212	-41.095	13.880	-74.776	1.00	95.26	L1	C
ATOM	4717	C	PRO	212	-40.691	16.540	-76.672	1.00	96.68	L1	C
ATOM	4718	O	PRO	212	-39.878	16.917	-77.520	1.00	96.58	L1	O
ATOM	4719	N	THR	213	-42.005	16.543	-76.872	1.00	97.41	L1	N
ATOM	4720	CA	THR	213	-42.600	16.998	-78.123	1.00	98.13	L1	C
ATOM	4721	CB	THR	213	-44.081	16.575	-78.209	1.00	98.19	L1	C
ATOM	4722	OG1	THR	213	-44.792	17.076	-77.068	1.00	97.41	L1	O
ATOM	4723	CG2	THR	213	-44.195	15.056	-78.248	1.00	98.19	L1	C
ATOM	4724	C	THR	213	-42.504	18.516	-78.277	1.00	98.77	L1	C
ATOM	4725	O	THR	213	-41.782	19.146	-77.474	1.00	99.20	L1	O

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ATOM   4726  OXT THR   213     -43.142  19.057 -79.208  1.00 99.08      L1   O
TER    4727      THR   213                              L1
ATOM   4728  CB  GLN     1     -10.333  41.335 -26.837  1.00 58.55      H1   C
ATOM   4729  CG  GLN     1     -11.368  41.133 -27.931  1.00 61.95      H1   C
ATOM   4730  CD  GLN     1     -12.786  41.172 -27.394  1.00 64.59      H1   C
ATOM   4731  OE1 GLN     1     -13.725  40.692 -28.034  1.00 66.92      H1   O
ATOM   4732  NE2 GLN     1     -12.949  41.749 -26.209  1.00 65.52      H1   N
ATOM   4733  C   GLN     1      -8.067  40.598 -27.611  1.00 51.02      H1   C
ATOM   4734  O   GLN     1      -6.978  40.491 -27.047  1.00 50.88      H1   O
ATOM   4735  N   GLN     1      -9.151  42.604 -28.597  1.00 56.41      H1   N
ATOM   4736  CA  GLN     1      -8.969  41.799 -27.352  1.00 54.79      H1   C
ATOM   4737  N   VAL     2      -8.528  39.693 -28.465  1.00 46.33      H1   N
ATOM   4738  CA  VAL     2      -7.658  38.679 -29.042  1.00 40.97      H1   C
ATOM   4739  CB  VAL     2      -8.026  37.266 -28.541  1.00 40.66      H1   C
ATOM   4740  CG1 VAL     2      -7.187  36.223 -29.261  1.00 37.87      H1   C
ATOM   4741  CG2 VAL     2      -7.800  37.175 -27.041  1.00 39.09      H1   C
ATOM   4742  C   VAL     2      -7.831  38.737 -30.551  1.00 40.14      H1   C
ATOM   4743  O   VAL     2      -8.948  38.863 -31.050  1.00 39.38      H1   O
ATOM   4744  N   GLN     3      -6.730  38.672 -31.285  1.00 38.15      H1   N
ATOM   4745  CA  GLN     3      -6.836  38.657 -32.733  1.00 37.80      H1   C
ATOM   4746  CB  GLN     3      -6.288  39.951 -33.340  1.00 41.11      H1   C
ATOM   4747  CG  GLN     3      -6.598  40.078 -34.830  1.00 48.63      H1   C
ATOM   4748  CD  GLN     3      -5.966  41.299 -35.480  1.00 54.48      H1   C
ATOM   4749  OE1 GLN     3      -5.933  41.412 -36.710  1.00 56.23      H1   O
ATOM   4750  NE2 GLN     3      -5.461  42.221 -34.658  1.00 55.22      H1   N
ATOM   4751  C   GLN     3      -6.097  37.475 -33.315  1.00 34.51      H1   C
ATOM   4752  O   GLN     3      -4.950  37.209 -32.960  1.00 34.76      H1   O
ATOM   4753  N   LEU     4      -6.774  36.764 -34.206  1.00 31.16      H1   N
ATOM   4754  CA  LEU     4      -6.152  35.706 -34.983  1.00 30.62      H1   C
ATOM   4755  CB  LEU     4      -7.036  34.455 -34.966  1.00 26.45      H1   C
ATOM   4756  CG  LEU     4      -6.920  33.556 -33.728  1.00 27.80      H1   C
ATOM   4757  CD1 LEU     4      -7.145  34.374 -32.449  1.00 25.47      H1   C
ATOM   4758  CD2 LEU     4      -7.930  32.423 -33.831  1.00 26.22      H1   C
ATOM   4759  C   LEU     4      -5.943  36.200 -36.419  1.00 32.23      H1   C
ATOM   4760  O   LEU     4      -6.899  36.554 -37.120  1.00 31.55      H1   O
ATOM   4761  N   GLN     5      -4.683  36.229 -36.840  1.00 31.02      H1   N
ATOM   4762  CA  GLN     5      -4.307  36.758 -38.144  1.00 31.99      H1   C
ATOM   4763  CB  GLN     5      -3.168  37.772 -37.976  1.00 35.77      H1   C
ATOM   4764  CG  GLN     5      -2.719  38.446 -39.262  1.00 41.14      H1   C
ATOM   4765  CD  GLN     5      -3.830  39.250 -39.904  1.00 45.66      H1   C
ATOM   4766  OE1 GLN     5      -3.749  39.625 -41.073  1.00 48.29      H1   O
ATOM   4767  NE2 GLN     5      -4.881  39.519 -39.137  1.00 48.64      H1   N
ATOM   4768  C   GLN     5      -3.863  35.614 -39.050  1.00 29.87      H1   C
ATOM   4769  O   GLN     5      -2.974  34.842 -38.690  1.00 31.50      H1   O
ATOM   4770  N   GLN     6      -4.482  35.509 -40.221  1.00 28.29      H1   N
ATOM   4771  CA  GLN     6      -4.243  34.377 -41.117  1.00 30.55      H1   C
ATOM   4772  CB  GLN     6      -5.564  33.740 -41.540  1.00 29.81      H1   C
ATOM   4773  CG  GLN     6      -6.344  33.071 -40.433  1.00 30.43      H1   C
ATOM   4774  CD  GLN     6      -7.562  32.345 -40.972  1.00 28.63      H1   C
ATOM   4775  OE1 GLN     6      -8.691  32.631 -40.580  1.00 30.41      H1   O
ATOM   4776  NE2 GLN     6      -7.336  31.408 -41.882  1.00 26.47      H1   N
ATOM   4777  C   GLN     6      -3.511  34.800 -42.376  1.00 30.75      H1   C
ATOM   4778  O   GLN     6      -3.696  35.916 -42.858  1.00 32.04      H1   O
ATOM   4779  N   TRP     7      -2.695  33.904 -42.918  1.00 28.95      H1   N
ATOM   4780  CA  TRP     7      -2.211  34.076 -44.278  1.00 28.90      H1   C
ATOM   4781  CB  TRP     7      -0.966  34.976 -44.301  1.00 28.68      H1   C
ATOM   4782  CG  TRP     7       0.236  34.402 -43.598  1.00 27.66      H1   C
ATOM   4783  CD2 TRP     7       0.704  34.740 -42.286  1.00 25.98      H1   C
ATOM   4784  CE2 TRP     7       1.888  34.010 -42.063  1.00 27.69      H1   C
ATOM   4785  CE3 TRP     7       0.238  35.594 -41.282  1.00 26.98      H1   C
ATOM   4786  CD1 TRP     7       1.128  33.499 -44.101  1.00 27.44      H1   C
ATOM   4787  NE1 TRP     7       2.176  33.258 -43.185  1.00 28.36      H1   N
ATOM   4788  CZ2 TRP     7       2.612  34.109 -40.880  1.00 28.83      H1   C
ATOM   4789  CZ3 TRP     7       0.956  35.692 -40.111  1.00 28.53      H1   C
ATOM   4790  CH2 TRP     7       2.132  34.955 -39.917  1.00 30.19      H1   C
ATOM   4791  C   TRP     7      -1.899  32.733 -44.923  1.00 28.22      H1   C
ATOM   4792  O   TRP     7      -2.070  31.681 -44.307  1.00 28.99      H1   O
ATOM   4793  N   GLY     8      -1.432  32.783 -46.166  1.00 28.35      H1   N
ATOM   4794  CA  GLY     8      -1.234  31.575 -46.945  1.00 26.37      H1   C
ATOM   4795  C   GLY     8      -1.920  31.683 -48.295  1.00 27.33      H1   C
ATOM   4796  O   GLY     8      -2.925  32.386 -48.444  1.00 27.76      H1   O
ATOM   4797  N   ALA     9      -1.377  30.984 -49.282  1.00 27.19      H1   N
ATOM   4798  CA  ALA     9      -1.887  31.057 -50.646  1.00 28.12      H1   C
ATOM   4799  CB  ALA     9      -0.828  30.527 -51.619  1.00 27.62      H1   C
ATOM   4800  C   ALA     9      -3.179  30.253 -50.791  1.00 28.31      H1   C
ATOM   4801  O   ALA     9      -3.233  29.079 -50.431  1.00 28.12      H1   O
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ATOM	4802	N	GLY	10	-4.212	30.888	-51.330	1.00	29.12	H1	N
ATOM	4803	CA	GLY	10	-5.493	30.222	-51.479	1.00	30.10	H1	C
ATOM	4804	C	GLY	10	-5.820	29.691	-52.871	1.00	32.33	H1	C
ATOM	4805	O	GLY	10	-6.659	28.798	-53.005	1.00	32.38	H1	O
ATOM	4806	N	LEU	11	-5.177	30.221	-53.909	1.00	31.52	H1	N
ATOM	4807	CA	LEU	11	-5.494	29.799	-55.275	1.00	33.59	H1	C
ATOM	4808	CB	LEU	11	-5.308	30.959	-56.255	1.00	36.39	H1	C
ATOM	4809	CG	LEU	11	-5.584	30.620	-57.725	1.00	40.88	H1	C
ATOM	4810	CD1	LEU	11	-6.969	29.990	-57.844	1.00	43.73	H1	C
ATOM	4811	CD2	LEU	11	-5.491	31.877	-58.586	1.00	40.70	H1	C
ATOM	4812	C	LEU	11	-4.626	28.629	-55.697	1.00	34.18	H1	C
ATOM	4813	O	LEU	11	-3.429	28.787	-55.936	1.00	35.52	H1	O
ATOM	4814	N	LEU	12	-5.232	27.448	-55.785	1.00	34.06	H1	N
ATOM	4815	CA	LEU	12	-4.470	26.219	-55.974	1.00	33.09	H1	C
ATOM	4816	CB	LEU	12	-4.533	25.358	-54.711	1.00	34.00	H1	C
ATOM	4817	CG	LEU	12	-4.079	25.990	-53.395	1.00	35.02	H1	C
ATOM	4818	CD1	LEU	12	-4.057	24.931	-52.297	1.00	33.56	H1	C
ATOM	4819	CD2	LEU	12	-2.698	26.606	-53.580	1.00	36.04	H1	C
ATOM	4820	C	LEU	12	-4.979	25.401	-57.147	1.00	33.61	H1	C
ATOM	4821	O	LEU	12	-6.129	25.542	-57.566	1.00	34.76	H1	O
ATOM	4822	N	LYS	13	-4.115	24.542	-57.671	1.00	33.61	H1	N
ATOM	4823	CA	LYS	13	-4.533	23.537	-58.636	1.00	35.02	H1	C
ATOM	4824	CB	LYS	13	-3.558	23.486	-59.813	1.00	35.86	H1	C
ATOM	4825	CG	LYS	13	-3.544	24.732	-60.671	1.00	39.05	H1	C
ATOM	4826	CD	LYS	13	-2.763	24.491	-61.952	0.50	41.55	H1	C
ATOM	4827	CE	LYS	13	-2.408	25.800	-62.627	1.00	44.69	H1	C
ATOM	4828	NZ	LYS	13	-1.530	26.633	-61.750	1.00	48.57	H1	N
ATOM	4829	C	LYS	13	-4.584	22.171	-57.967	1.00	35.21	H1	C
ATOM	4830	O	LYS	13	-3.887	21.924	-56.983	1.00	37.52	H1	O
ATOM	4831	N	PRO	14	-5.405	21.258	-58.499	1.00	34.61	H1	N
ATOM	4832	CD	PRO	14	-6.318	21.454	-59.638	1.00	34.09	H1	C
ATOM	4833	CA	PRO	14	-5.476	19.894	-57.969	1.00	34.50	H1	C
ATOM	4834	CB	PRO	14	-6.368	19.169	-58.974	1.00	34.26	H1	C
ATOM	4835	CG	PRO	14	-7.214	20.257	-59.557	1.00	33.95	H1	C
ATOM	4836	C	PRO	14	-4.093	19.253	-57.854	1.00	34.86	H1	C
ATOM	4837	O	PRO	14	-3.222	19.493	-58.690	1.00	35.81	H1	O
ATOM	4838	N	SER	15	-3.914	18.455	-56.803	1.00	34.00	H1	N
ATOM	4839	CA	SER	15	-2.684	17.718	-56.515	1.00	34.80	H1	C
ATOM	4840	CB	SER	15	-1.999	17.245	-57.805	1.00	35.51	H1	C
ATOM	4841	OG	SER	15	-1.036	18.192	-58.234	1.00	38.11	H1	O
ATOM	4842	C	SER	15	-1.694	18.534	-55.684	1.00	34.51	H1	C
ATOM	4843	O	SER	15	-0.761	17.980	-55.099	1.00	35.24	H1	O
ATOM	4844	N	GLU	16	-1.901	19.845	-55.622	1.00	34.15	H1	N
ATOM	4845	CA	GLU	16	-1.018	20.705	-54.845	1.00	34.52	H1	C
ATOM	4846	CB	GLU	16	-1.183	22.164	-55.283	1.00	36.08	H1	C
ATOM	4847	CG	GLU	16	-0.552	22.478	-56.646	1.00	41.58	H1	C
ATOM	4848	CD	GLU	16	-0.918	23.864	-57.160	1.00	44.69	H1	C
ATOM	4849	OE1	GLU	16	-1.573	24.626	-56.423	1.00	44.93	H1	O
ATOM	4850	OE2	GLU	16	-0.555	24.193	-58.309	1.00	49.52	H1	O
ATOM	4851	C	GLU	16	-1.256	20.583	-53.338	1.00	34.67	H1	C
ATOM	4852	O	GLU	16	-2.252	20.008	-52.898	1.00	32.76	H1	O
ATOM	4853	N	THR	17	-0.319	21.120	-52.559	1.00	33.99	H1	N
ATOM	4854	CA	THR	17	-0.421	21.165	-51.104	1.00	32.35	H1	C
ATOM	4855	CB	THR	17	0.928	20.860	-50.447	1.00	31.91	H1	C
ATOM	4856	OG1	THR	17	1.289	19.505	-50.722	1.00	35.50	H1	O
ATOM	4857	CG2	THR	17	0.853	21.073	-48.940	1.00	31.69	H1	C
ATOM	4858	C	THR	17	-0.867	22.539	-50.627	1.00	31.96	H1	C
ATOM	4859	O	THR	17	-0.296	23.554	-51.024	1.00	31.96	H1	O
ATOM	4860	N	LEU	18	-1.889	22.565	-49.774	1.00	29.65	H1	N
ATOM	4861	CA	LEU	18	-2.355	23.812	-49.179	1.00	28.49	H1	C
ATOM	4862	CB	LEU	18	-3.841	23.720	-48.831	1.00	27.90	H1	C
ATOM	4863	CG	LEU	18	-4.387	24.865	-47.973	1.00	27.75	H1	C
ATOM	4864	CD1	LEU	18	-4.433	26.133	-48.808	1.00	28.48	H1	C
ATOM	4865	CD2	LEU	18	-5.784	24.520	-47.467	1.00	27.32	H1	C
ATOM	4866	C	LEU	18	-1.557	24.086	-47.914	1.00	29.31	H1	C
ATOM	4867	O	LEU	18	-1.324	23.183	-47.105	1.00	29.34	H1	O
ATOM	4868	N	SER	19	-1.138	25.332	-47.740	1.00	28.71	H1	N
ATOM	4869	CA	SER	19	-0.342	25.693	-46.580	1.00	28.95	H1	C
ATOM	4870	CB	SER	19	1.136	25.745	-46.969	1.00	29.72	H1	C
ATOM	4871	OG	SER	19	1.927	26.134	-45.861	1.00	37.84	H1	O
ATOM	4872	C	SER	19	-0.790	27.032	-46.000	1.00	28.99	H1	C
ATOM	4873	O	SER	19	-0.614	28.084	-46.611	1.00	30.84	H1	O
ATOM	4874	N	LEU	20	-1.384	26.985	-44.814	1.00	28.20	H1	N
ATOM	4875	CA	LEU	20	-1.932	28.179	-44.193	1.00	26.04	H1	C
ATOM	4876	CB	LEU	20	-3.447	28.056	-44.050	1.00	25.36	H1	C
ATOM	4877	CG	LEU	20	-4.265	27.889	-45.334	1.00	26.53	H1	C

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ATOM   4878  CD1 LEU   20      -5.744  27.831 -44.970  1.00 23.87      H1   C
ATOM   4879  CD2 LEU   20      -3.999  29.053 -46.288  1.00 25.15      H1   C
ATOM   4880  C   LEU   20      -1.308  28.376 -42.825  1.00 25.77      H1   C
ATOM   4881  O   LEU   20      -0.850  27.427 -42.197  1.00 26.14      H1   O
ATOM   4882  N   THR   21      -1.299  29.618 -42.365  1.00 26.29      H1   N
ATOM   4883  CA  THR   21      -0.685  29.947 -41.093  1.00 27.57      H1   C
ATOM   4884  CB  THR   21       0.691  30.616 -41.314  1.00 28.87      H1   C
ATOM   4885  OG1 THR   21       1.536  29.733 -42.072  1.00 30.29      H1   O
ATOM   4886  CG2 THR   21       1.348  30.928 -39.988  1.00 26.32      H1   C
ATOM   4887  C   THR   21      -1.599  30.897 -40.332  1.00 27.47      H1   C
ATOM   4888  O   THR   21      -2.294  31.715 -40.930  1.00 26.74      H1   O
ATOM   4889  N   CYS   22      -1.606  30.778 -39.011  1.00 28.77      H1   N
ATOM   4890  CA  CYS   22      -2.393  31.675 -38.175  1.00 29.72      H1   C
ATOM   4891  C   CYS   22      -1.469  32.192 -37.082  1.00 30.55      H1   C
ATOM   4892  O   CYS   22      -0.752  31.411 -36.456  1.00 30.73      H1   O
ATOM   4893  CB  CYS   22      -3.564  30.899 -37.565  1.00 30.30      H1   C
ATOM   4894  SG  CYS   22      -4.786  31.772 -36.524  1.00 31.82      H1   S
ATOM   4895  N   ALA   23      -1.481  33.504 -36.864  1.00 29.76      H1   N
ATOM   4896  CA  ALA   23      -0.725  34.101 -35.771  1.00 31.19      H1   C
ATOM   4897  CB  ALA   23       0.187  35.216 -36.298  1.00 31.06      H1   C
ATOM   4898  C   ALA   23      -1.684  34.662 -34.732  1.00 29.91      H1   C
ATOM   4899  O   ALA   23      -2.677  35.300 -35.075  1.00 29.39      H1   O
ATOM   4900  N   VAL   24      -1.375  34.429 -33.462  1.00 29.49      H1   N
ATOM   4901  CA  VAL   24      -2.243  34.855 -32.375  1.00 29.62      H1   C
ATOM   4902  CB  VAL   24      -2.415  33.717 -31.352  1.00 28.75      H1   C
ATOM   4903  CG1 VAL   24      -3.211  34.202 -30.149  1.00 28.80      H1   C
ATOM   4904  CG2 VAL   24      -3.112  32.543 -32.015  1.00 26.06      H1   C
ATOM   4905  C   VAL   24      -1.708  36.095 -31.663  1.00 31.01      H1   C
ATOM   4906  O   VAL   24      -0.566  36.118 -31.206  1.00 32.54      H1   O
ATOM   4907  N   TYR   25      -2.543  37.124 -31.567  1.00 31.55      H1   N
ATOM   4908  CA  TYR   25      -2.175  38.336 -30.848  1.00 32.59      H1   C
ATOM   4909  CB  TYR   25      -2.177  39.527 -31.807  1.00 34.98      H1   C
ATOM   4910  CG  TYR   25      -1.229  39.346 -32.968  1.00 36.34      H1   C
ATOM   4911  CD1 TYR   25      -1.704  39.045 -34.236  1.00 37.36      H1   C
ATOM   4912  CE1 TYR   25      -0.838  38.843 -35.297  1.00 40.15      H1   C
ATOM   4913  CD2 TYR   25       0.144  39.445 -32.788  1.00 38.49      H1   C
ATOM   4914  CE2 TYR   25       1.020  39.243 -33.843  1.00 40.50      H1   C
ATOM   4915  CZ  TYR   25       0.523  38.942 -35.093  1.00 40.26      H1   C
ATOM   4916  OH  TYR   25       1.389  38.726 -36.142  1.00 44.41      H1   O
ATOM   4917  C   TYR   25      -3.129  38.594 -29.689  1.00 32.47      H1   C
ATOM   4918  O   TYR   25      -4.289  38.942 -29.896  1.00 32.98      H1   O
ATOM   4919  N   GLY   26      -2.637  38.415 -28.468  1.00 31.79      H1   N
ATOM   4920  CA  GLY   26      -3.465  38.647 -27.296  1.00 32.41      H1   C
ATOM   4921  C   GLY   26      -3.828  37.380 -26.537  1.00 32.65      H1   C
ATOM   4922  O   GLY   26      -3.801  36.279 -27.093  1.00 31.74      H1   O
ATOM   4923  N   GLY   27      -4.167  37.539 -25.259  1.00 32.32      H1   N
ATOM   4924  CA  GLY   27      -4.515  36.399 -24.431  1.00 30.56      H1   C
ATOM   4925  C   GLY   27      -3.347  35.444 -24.311  1.00 31.96      H1   C
ATOM   4926  O   GLY   27      -2.258  35.738 -24.791  1.00 33.98      H1   O
ATOM   4927  N   SER   28      -3.570  34.297 -23.677  1.00 31.00      H1   N
ATOM   4928  CA  SER   28      -2.546  33.261 -23.580  1.00 29.78      H1   C
ATOM   4929  CB  SER   28      -2.782  32.397 -22.340  1.00 27.61      H1   C
ATOM   4930  OG  SER   28      -2.050  31.184 -22.424  1.00 29.12      H1   O
ATOM   4931  C   SER   28      -2.535  32.367 -24.815  1.00 29.77      H1   C
ATOM   4932  O   SER   28      -3.569  32.152 -25.442  1.00 31.74      H1   O
ATOM   4933  N   PHE   29      -1.365  31.836 -25.153  1.00 28.60      H1   N
ATOM   4934  CA  PHE   29      -1.246  30.952 -26.301  1.00 28.86      H1   C
ATOM   4935  CB  PHE   29       0.048  31.229 -27.077  1.00 28.06      H1   C
ATOM   4936  CG  PHE   29       0.202  30.383 -28.307  1.00 26.67      H1   C
ATOM   4937  CD1 PHE   29      -0.550  30.638 -29.436  1.00 28.66      H1   C
ATOM   4938  CD2 PHE   29       1.081  29.317 -28.327  1.00 27.57      H1   C
ATOM   4939  CE1 PHE   29      -0.431  29.841 -30.559  1.00 29.40      H1   C
ATOM   4940  CE2 PHE   29       1.205  28.518 -29.442  1.00 27.25      H1   C
ATOM   4941  CZ  PHE   29       0.452  28.778 -30.560  1.00 28.48      H1   C
ATOM   4942  C   PHE   29      -1.260  29.506 -25.851  1.00 28.91      H1   C
ATOM   4943  O   PHE   29      -1.893  28.653 -26.483  1.00 31.89      H1   O
ATOM   4944  N   SER   30      -0.566  29.226 -24.755  1.00 27.90      H1   N
ATOM   4945  CA  SER   30      -0.390  27.849 -24.311  1.00 29.14      H1   C
ATOM   4946  CB  SER   30       0.850  27.734 -23.418  1.00 31.78      H1   C
ATOM   4947  OG  SER   30       0.687  28.470 -22.221  1.00 36.30      H1   O
ATOM   4948  C   SER   30      -1.612  27.295 -23.574  1.00 27.44      H1   C
ATOM   4949  O   SER   30      -1.699  26.097 -23.340  1.00 26.60      H1   O
ATOM   4950  N   ALA   31      -2.558  28.163 -23.227  1.00 27.53      H1   N
ATOM   4951  CA  ALA   31      -3.706  27.760 -22.413  1.00 27.10      H1   C
ATOM   4952  CB  ALA   31      -4.175  28.937 -21.580  1.00 23.65      H1   C
ATOM   4953  C   ALA   31      -4.888  27.190 -23.211  1.00 27.46      H1   C
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ATOM   4954  O   ALA  31      -5.917  26.850 -22.633  1.00 28.90      H1   O
ATOM   4955  N   TYR  32      -4.743  27.091 -24.527  1.00 26.48      H1   N
ATOM   4956  CA  TYR  32      -5.864  26.726 -25.396  1.00 25.86      H1   C
ATOM   4957  CB  TYR  32      -6.393  27.962 -26.132  1.00 24.51      H1   C
ATOM   4958  CG  TYR  32      -6.878  29.066 -25.223  1.00 27.00      H1   C
ATOM   4959  CD1 TYR  32      -6.062  30.146 -24.912  1.00 27.70      H1   C
ATOM   4960  CE1 TYR  32      -6.510  31.167 -24.086  1.00 27.16      H1   C
ATOM   4961  CD2 TYR  32      -8.161  29.034 -24.681  1.00 25.53      H1   C
ATOM   4962  CE2 TYR  32      -8.616  30.045 -23.857  1.00 25.30      H1   C
ATOM   4963  CZ  TYR  32      -7.790  31.109 -23.564  1.00 26.69      H1   C
ATOM   4964  OH  TYR  32      -8.250  32.131 -22.771  1.00 28.27      H1   O
ATOM   4965  C   TYR  32      -5.486  25.688 -26.440  1.00 24.12      H1   C
ATOM   4966  O   TYR  32      -4.327  25.601 -26.856  1.00 24.09      H1   O
ATOM   4967  N   TYR  33      -6.471  24.917 -26.887  1.00 22.70      H1   N
ATOM   4968  CA  TYR  33      -6.348  24.241 -28.173  1.00 22.68      H1   C
ATOM   4969  CB  TYR  33      -7.374  23.116 -28.298  1.00 22.31      H1   C
ATOM   4970  CG  TYR  33      -7.016  21.898 -27.492  1.00 21.33      H1   C
ATOM   4971  CD1 TYR  33      -6.239  20.893 -28.042  1.00 20.79      H1   C
ATOM   4972  CE1 TYR  33      -5.886  19.783 -27.311  1.00 22.09      H1   C
ATOM   4973  CD2 TYR  33      -7.440  21.758 -26.173  1.00 22.16      H1   C
ATOM   4974  CE2 TYR  33      -7.092  20.642 -25.429  1.00 23.42      H1   C
ATOM   4975  CZ  TYR  33      -6.311  19.659 -26.010  1.00 23.95      H1   C
ATOM   4976  OH  TYR  33      -5.941  18.549 -25.298  1.00 23.94      H1   O
ATOM   4977  C   TYR  33      -6.554  25.244 -29.299  1.00 22.45      H1   C
ATOM   4978  O   TYR  33      -7.292  26.218 -29.155  1.00 22.76      H1   O
ATOM   4979  N   TRP  34      -5.887  25.002 -30.417  1.00 22.98      H1   N
ATOM   4980  CA  TRP  34      -6.012  25.859 -31.582  1.00 23.60      H1   C
ATOM   4981  CB  TRP  34      -4.650  26.490 -31.885  1.00 23.80      H1   C
ATOM   4982  CG  TRP  34      -4.247  27.427 -30.790  1.00 23.49      H1   C
ATOM   4983  CD2 TRP  34      -4.745  28.749 -30.577  1.00 22.41      H1   C
ATOM   4984  CE2 TRP  34      -4.165  29.227 -29.385  1.00 25.04      H1   C
ATOM   4985  CE3 TRP  34      -5.629  29.578 -31.280  1.00 23.05      H1   C
ATOM   4986  CD1 TRP  34      -3.402  27.165 -29.752  1.00 24.44      H1   C
ATOM   4987  NE1 TRP  34      -3.347  28.240 -28.900  1.00 23.72      H1   N
ATOM   4988  CZ2 TRP  34      -4.442  30.495 -28.879  1.00 21.78      H1   C
ATOM   4989  CZ3 TRP  34      -5.900  30.833 -30.777  1.00 21.48      H1   C
ATOM   4990  CH2 TRP  34      -5.309  31.280 -29.587  1.00 22.95      H1   C
ATOM   4991  C   TRP  34      -6.530  25.048 -32.766  1.00 21.45      H1   C
ATOM   4992  O   TRP  34      -5.988  23.987 -33.070  1.00 21.93      H1   O
ATOM   4993  N   ASN  35      -7.578  25.556 -33.420  1.00 19.88      H1   N
ATOM   4994  CA  ASN  35      -8.355  24.785 -34.392  1.00 20.46      H1   C
ATOM   4995  CB  ASN  35      -9.841  24.803 -34.021  1.00 20.89      H1   C
ATOM   4996  CG  ASN  35     -10.101  24.285 -32.627  1.00 23.23      H1   C
ATOM   4997  OD1 ASN  35     -10.302  23.087 -32.428  1.00 24.03      H1   O
ATOM   4998  ND2 ASN  35     -10.099  25.185 -31.650  1.00 22.85      H1   N
ATOM   4999  C   ASN  35      -8.236  25.315 -35.810  1.00 20.87      H1   C
ATOM   5000  O   ASN  35      -8.048  26.505 -36.019  1.00 23.07      H1   O
ATOM   5001  N   TRP  36      -8.377  24.418 -36.782  1.00 22.36      H1   N
ATOM   5002  CA  TRP  36      -8.702  24.801 -38.152  1.00 22.05      H1   C
ATOM   5003  CB  TRP  36      -7.658  24.246 -39.121  1.00 22.90      H1   C
ATOM   5004  CG  TRP  36      -6.318  24.938 -39.025  1.00 25.04      H1   C
ATOM   5005  CD2 TRP  36      -5.956  26.193 -39.619  1.00 22.07      H1   C
ATOM   5006  CE2 TRP  36      -4.612  26.439 -39.284  1.00 24.03      H1   C
ATOM   5007  CE3 TRP  36      -6.639  27.129 -40.403  1.00 22.32      H1   C
ATOM   5008  CD1 TRP  36      -5.208  24.490 -38.370  1.00 20.33      H1   C
ATOM   5009  NE1 TRP  36      -4.179  25.385 -38.521  1.00 23.82      H1   N
ATOM   5010  CZ2 TRP  36      -3.933  27.587 -39.706  1.00 23.84      H1   C
ATOM   5011  CZ3 TRP  36      -5.968  28.266 -40.821  1.00 24.39      H1   C
ATOM   5012  CH2 TRP  36      -4.627  28.486 -40.471  1.00 24.85      H1   C
ATOM   5013  C   TRP  36     -10.087  24.261 -38.517  1.00 22.80      H1   C
ATOM   5014  O   TRP  36     -10.434  23.122 -38.193  1.00 21.90      H1   O
ATOM   5015  N   ILE  37     -10.871  25.094 -39.187  1.00 22.71      H1   N
ATOM   5016  CA  ILE  37     -12.244  24.773 -39.554  1.00 21.18      H1   C
ATOM   5017  CB  ILE  37     -13.232  25.478 -38.591  1.00 22.23      H1   C
ATOM   5018  CG2 ILE  37     -14.669  25.262 -39.039  1.00 20.06      H1   C
ATOM   5019  CG1 ILE  37     -13.024  24.950 -37.171  1.00 21.40      H1   C
ATOM   5020  CD1 ILE  37     -13.704  25.776 -36.102  1.00 20.52      H1   C
ATOM   5021  C   ILE  37     -12.445  25.310 -40.966  1.00 23.40      H1   C
ATOM   5022  O   ILE  37     -11.964  26.395 -41.285  1.00 23.50      H1   O
ATOM   5023  N   ARG  38     -13.134  24.571 -41.827  1.00 22.17      H1   N
ATOM   5024  CA  ARG  38     -13.417  25.120 -43.151  1.00 23.30      H1   C
ATOM   5025  CB  ARG  38     -12.680  24.328 -44.237  1.00 22.10      H1   C
ATOM   5026  CG  ARG  38     -13.113  22.886 -44.372  1.00 24.22      H1   C
ATOM   5027  CD  ARG  38     -12.272  22.174 -45.418  1.00 23.06      H1   C
ATOM   5028  NE  ARG  38     -12.792  20.840 -45.694  1.00 23.01      H1   N
ATOM   5029  CZ  ARG  38     -12.283  20.000 -46.587  1.00 22.72      H1   C
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ATOM	5030	NH1	ARG	38	-11.212	20.337	-47.310	1.00	18.87	H1	N
ATOM	5031	NH2	ARG	38	-12.875	18.832	-46.782	1.00	20.60	H1	N
ATOM	5032	C	ARG	38	-14.906	25.185	-43.469	1.00	23.88	H1	C
ATOM	5033	O	ARG	38	-15.720	24.482	-42.863	1.00	23.72	H1	O
ATOM	5034	N	GLN	39	-15.256	26.055	-44.411	1.00	24.72	H1	N
ATOM	5035	CA	GLN	39	-16.638	26.222	-44.829	1.00	24.67	H1	C
ATOM	5036	CB	GLN	39	-17.224	27.509	-44.238	1.00	25.79	H1	C
ATOM	5037	CG	GLN	39	-18.684	27.793	-44.635	1.00	27.90	H1	C
ATOM	5038	CD	GLN	39	-19.318	28.866	-43.757	1.00	29.29	H1	C
ATOM	5039	OE1	GLN	39	-18.788	29.966	-43.633	1.00	32.06	H1	O
ATOM	5040	NE2	GLN	39	-20.447	28.545	-43.141	1.00	28.85	H1	N
ATOM	5041	C	GLN	39	-16.690	26.288	-46.345	1.00	25.97	H1	C
ATOM	5042	O	GLN	39	-16.266	27.270	-46.947	1.00	25.21	H1	O
ATOM	5043	N	PRO	40	-17.213	25.233	-46.981	1.00	26.39	H1	N
ATOM	5044	CD	PRO	40	-17.750	24.017	-46.345	1.00	28.52	H1	C
ATOM	5045	CA	PRO	40	-17.377	25.213	-48.433	1.00	29.08	H1	C
ATOM	5046	CB	PRO	40	-17.779	23.768	-48.731	1.00	27.66	H1	C
ATOM	5047	CG	PRO	40	-18.437	23.304	-47.485	1.00	28.54	H1	C
ATOM	5048	C	PRO	40	-18.448	26.219	-48.836	1.00	32.62	H1	C
ATOM	5049	O	PRO	40	-19.359	26.516	-48.061	1.00	32.18	H1	O
ATOM	5050	N	PRO	41	-18.345	26.766	-50.053	1.00	36.33	H1	N
ATOM	5051	CD	PRO	41	-17.406	26.340	-51.105	1.00	37.35	H1	C
ATOM	5052	CA	PRO	41	-19.204	27.872	-50.494	1.00	37.78	H1	C
ATOM	5053	CB	PRO	41	-18.839	28.044	-51.969	1.00	39.57	H1	C
ATOM	5054	CG	PRO	41	-17.449	27.484	-52.074	1.00	40.10	H1	C
ATOM	5055	C	PRO	41	-20.687	27.568	-50.307	1.00	38.36	H1	C
ATOM	5056	O	PRO	41	-21.165	26.510	-50.704	1.00	37.50	H1	O
ATOM	5057	N	GLY	42	-21.406	28.497	-49.682	1.00	40.85	H1	N
ATOM	5058	CA	GLY	42	-22.837	28.332	-49.499	1.00	43.07	H1	C
ATOM	5059	C	GLY	42	-23.227	27.130	-48.653	1.00	44.02	H1	C
ATOM	5060	O	GLY	42	-24.360	26.657	-48.727	1.00	44.58	H1	O
ATOM	5061	N	LYS	43	-22.297	26.634	-47.842	1.00	42.75	H1	N
ATOM	5062	CA	LYS	43	-22.551	25.440	-47.046	1.00	41.76	H1	C
ATOM	5063	CB	LYS	43	-21.778	24.254	-47.627	1.00	44.31	H1	C
ATOM	5064	CG	LYS	43	-22.140	23.962	-49.074	1.00	48.96	H1	C
ATOM	5065	CD	LYS	43	-22.254	22.468	-49.324	1.00	51.38	H1	C
ATOM	5066	CE	LYS	43	-23.015	22.187	-50.617	1.00	54.57	H1	C
ATOM	5067	NZ	LYS	43	-22.406	22.886	-51.784	1.00	54.57	H1	N
ATOM	5068	C	LYS	43	-22.171	25.647	-45.590	1.00	37.71	H1	C
ATOM	5069	O	LYS	43	-21.889	26.766	-45.175	1.00	40.41	H1	O
ATOM	5070	N	GLY	44	-22.160	24.562	-44.823	1.00	32.46	H1	N
ATOM	5071	CA	GLY	44	-21.927	24.654	-43.393	1.00	28.89	H1	C
ATOM	5072	C	GLY	44	-20.473	24.575	-42.949	1.00	28.33	H1	C
ATOM	5073	O	GLY	44	-19.570	25.015	-43.654	1.00	28.78	H1	O
ATOM	5074	N	LEU	45	-20.251	24.004	-41.770	1.00	26.37	H1	N
ATOM	5075	CA	LEU	45	-18.938	24.020	-41.139	1.00	24.45	H1	C
ATOM	5076	CB	LEU	45	-19.024	24.701	-39.766	1.00	23.45	H1	C
ATOM	5077	CG	LEU	45	-19.498	26.154	-39.752	1.00	24.17	H1	C
ATOM	5078	CD1	LEU	45	-19.692	26.636	-38.323	1.00	20.49	H1	C
ATOM	5079	CD2	LEU	45	-18.471	27.012	-40.474	1.00	23.06	H1	C
ATOM	5080	C	LEU	45	-18.379	22.616	-40.965	1.00	22.98	H1	C
ATOM	5081	O	LEU	45	-19.101	21.689	-40.600	1.00	21.96	H1	O
ATOM	5082	N	GLU	46	-17.081	22.475	-41.208	1.00	22.29	H1	N
ATOM	5083	CA	GLU	46	-16.397	21.199	-41.044	1.00	22.11	H1	C
ATOM	5084	CB	GLU	46	-16.060	20.609	-42.416	1.00	21.07	H1	C
ATOM	5085	CG	GLU	46	-15.392	19.253	-42.342	1.00	23.85	H1	C
ATOM	5086	CD	GLU	46	-15.117	18.646	-43.711	1.00	26.12	H1	C
ATOM	5087	OE1	GLU	46	-15.068	19.390	-44.714	1.00	25.34	H1	O
ATOM	5088	OE2	GLU	46	-14.947	17.414	-43.779	1.00	28.37	H1	O
ATOM	5089	C	GLU	46	-15.112	21.411	-40.245	1.00	22.69	H1	C
ATOM	5090	O	GLU	46	-14.246	22.188	-40.643	1.00	22.85	H1	O
ATOM	5091	N	TRP	47	-14.994	20.727	-39.113	1.00	23.32	H1	N
ATOM	5092	CA	TRP	47	-13.810	20.853	-38.271	1.00	24.50	H1	C
ATOM	5093	CB	TRP	47	-14.124	20.356	-36.862	1.00	23.99	H1	C
ATOM	5094	CG	TRP	47	-12.985	20.457	-35.899	1.00	23.86	H1	C
ATOM	5095	CD2	TRP	47	-12.225	19.369	-35.357	1.00	23.15	H1	C
ATOM	5096	CE2	TRP	47	-11.346	19.912	-34.401	1.00	22.05	H1	C
ATOM	5097	CE3	TRP	47	-12.209	17.986	-35.584	1.00	22.33	H1	C
ATOM	5098	CD1	TRP	47	-12.536	21.586	-35.276	1.00	23.05	H1	C
ATOM	5099	NE1	TRP	47	-11.555	21.268	-34.373	1.00	23.21	H1	N
ATOM	5100	CZ2	TRP	47	-10.461	19.124	-33.665	1.00	20.75	H1	C
ATOM	5101	CZ3	TRP	47	-11.326	17.200	-34.852	1.00	20.87	H1	C
ATOM	5102	CH2	TRP	47	-10.467	17.774	-33.903	1.00	23.70	H1	C
ATOM	5103	C	TRP	47	-12.687	20.020	-38.872	1.00	24.58	H1	C
ATOM	5104	O	TRP	47	-12.877	18.849	-39.176	1.00	23.77	H1	O
ATOM	5105	N	ILE	48	-11.519	20.626	-39.042	1.00	24.78	H1	N

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ATOM   5106  CA  ILE    48     -10.389  19.929 -39.647  1.00 24.00      H1   C
ATOM   5107  CB  ILE    48      -9.526  20.913 -40.481  1.00 24.53      H1   C
ATOM   5108  CG2 ILE    48      -8.285  20.196 -41.038  1.00 24.01      H1   C
ATOM   5109  CG1 ILE    48     -10.383  21.502 -41.611  1.00 23.41      H1   C
ATOM   5110  CD1 ILE    48      -9.698  22.583 -42.429  1.00 24.34      H1   C
ATOM   5111  C   ILE    48      -9.530  19.259 -38.575  1.00 23.04      H1   C
ATOM   5112  O   ILE    48      -9.181  18.080 -38.689  1.00 21.34      H1   O
ATOM   5113  N   GLY    49      -9.207  20.011 -37.528  1.00 22.63      H1   N
ATOM   5114  CA  GLY    49      -8.387  19.476 -36.459  1.00 21.49      H1   C
ATOM   5115  C   GLY    49      -7.929  20.562 -35.502  1.00 22.86      H1   C
ATOM   5116  O   GLY    49      -8.258  21.733 -35.679  1.00 22.27      H1   O
ATOM   5117  N   GLU    50      -7.166  20.164 -34.490  1.00 22.34      H1   N
ATOM   5118  CA  GLU    50      -6.673  21.078 -33.469  1.00 23.19      H1   C
ATOM   5119  CB  GLU    50      -7.577  21.023 -32.234  1.00 23.22      H1   C
ATOM   5120  CG  GLU    50      -7.510  19.674 -31.530  1.00 20.57      H1   C
ATOM   5121  CD  GLU    50      -8.512  19.538 -30.396  1.00 23.89      H1   C
ATOM   5122  OE1 GLU    50      -9.378  20.428 -30.251  1.00 22.86      H1   O
ATOM   5123  OE2 GLU    50      -8.438  18.533 -29.656  1.00 20.61      H1   O
ATOM   5124  C   GLU    50      -5.257  20.664 -33.053  1.00 25.41      H1   C
ATOM   5125  O   GLU    50      -4.799  19.559 -33.349  1.00 26.68      H1   O
ATOM   5126  N   ILE    51      -4.577  21.549 -32.340  1.00 25.24      H1   N
ATOM   5127  CA  ILE    51      -3.275  21.223 -31.780  1.00 25.42      H1   C
ATOM   5128  CB  ILE    51      -2.151  21.625 -32.768  1.00 25.30      H1   C
ATOM   5129  CG2 ILE    51      -2.110  23.145 -32.919  1.00 20.77      H1   C
ATOM   5130  CG1 ILE    51      -0.805  21.091 -32.288  1.00 25.23      H1   C
ATOM   5131  CD1 ILE    51       0.268  21.133 -33.364  1.00 26.63      H1   C
ATOM   5132  C   ILE    51      -3.150  22.023 -30.492  1.00 25.89      H1   C
ATOM   5133  O   ILE    51      -3.846  23.019 -30.318  1.00 25.01      H1   O
ATOM   5134  N   ASN    52      -2.294  21.585 -29.575  1.00 27.77      H1   N
ATOM   5135  CA  ASN    52      -1.924  22.442 -28.456  1.00 28.51      H1   C
ATOM   5136  CB  ASN    52      -2.366  21.828 -27.119  1.00 31.18      H1   C
ATOM   5137  CG  ASN    52      -1.642  20.539 -26.785  1.00 33.23      H1   C
ATOM   5138  OD1 ASN    52      -0.514  20.307 -27.221  1.00 37.24      H1   O
ATOM   5139  ND2 ASN    52      -2.294  19.688 -25.999  1.00 34.97      H1   N
ATOM   5140  C   ASN    52      -0.423  22.706 -28.463  1.00 29.67      H1   C
ATOM   5141  O   ASN    52       0.293  22.205 -29.327  1.00 29.70      H1   O
ATOM   5142  N   HIS    53       0.047  23.497 -27.505  1.00 30.65      H1   N
ATOM   5143  CA  HIS    53       1.391  24.059 -27.565  1.00 34.02      H1   C
ATOM   5144  CB  HIS    53       1.597  25.058 -26.431  1.00 36.59      H1   C
ATOM   5145  CG  HIS    53       1.574  24.427 -25.077  1.00 41.39      H1   C
ATOM   5146  CD2 HIS    53       0.542  24.090 -24.268  1.00 42.73      H1   C
ATOM   5147  ND1 HIS    53       2.719  24.025 -24.425  1.00 44.09      H1   N
ATOM   5148  CE1 HIS    53       2.393  23.465 -23.273  1.00 43.74      H1   C
ATOM   5149  NE2 HIS    53       1.077  23.492 -23.154  1.00 44.27      H1   N
ATOM   5150  C   HIS    53       2.473  22.990 -27.475  1.00 34.37      H1   C
ATOM   5151  O   HIS    53       3.621  23.241 -27.819  1.00 34.24      H1   O
ATOM   5152  N   SER    54       2.115  21.805 -26.999  1.00 34.17      H1   N
ATOM   5153  CA  SER    54       3.103  20.752 -26.852  1.00 34.95      H1   C
ATOM   5154  CB  SER    54       2.728  19.819 -25.708  1.00 37.13      H1   C
ATOM   5155  OG  SER    54       1.673  18.955 -26.090  1.00 44.66      H1   O
ATOM   5156  C   SER    54       3.207  19.960 -28.140  1.00 35.47      H1   C
ATOM   5157  O   SER    54       4.033  19.058 -28.255  1.00 37.27      H1   O
ATOM   5158  N   GLY    55       2.360  20.291 -29.110  1.00 33.32      H1   N
ATOM   5159  CA  GLY    55       2.456  19.646 -30.407  1.00 33.10      H1   C
ATOM   5160  C   GLY    55       1.537  18.459 -30.649  1.00 30.70      H1   C
ATOM   5161  O   GLY    55       1.519  17.913 -31.744  1.00 31.90      H1   O
ATOM   5162  N   ARG    56       0.770  18.047 -29.647  1.00 31.34      H1   N
ATOM   5163  CA  ARG    56      -0.181  16.959 -29.860  1.00 31.52      H1   C
ATOM   5164  CB  ARG    56      -0.674  16.399 -28.520  1.00 33.02      H1   C
ATOM   5165  CG  ARG    56      -1.671  15.245 -28.675  1.00 39.22      H1   C
ATOM   5166  CD  ARG    56      -1.844  14.428 -27.393  1.00 41.40      H1   C
ATOM   5167  NE  ARG    56      -2.188  15.260 -26.239  1.00 44.28      H1   N
ATOM   5168  CZ  ARG    56      -3.363  15.858 -26.069  1.00 45.93      H1   C
ATOM   5169  NH1 ARG    56      -4.318  15.722 -26.983  1.00 45.10      H1   N
ATOM   5170  NH2 ARG    56      -3.584  16.588 -24.980  1.00 46.47      H1   N
ATOM   5171  C   ARG    56      -1.376  17.427 -30.702  1.00 29.42      H1   C
ATOM   5172  O   ARG    56      -1.861  18.544 -30.547  1.00 29.24      H1   O
ATOM   5173  N   THR    57      -1.844  16.560 -31.590  1.00 27.46      H1   N
ATOM   5174  CA  THR    57      -2.908  16.912 -32.518  1.00 28.02      H1   C
ATOM   5175  CB  THR    57      -2.401  16.915 -33.974  1.00 26.99      H1   C
ATOM   5176  OG1 THR    57      -1.806  15.646 -34.267  1.00 28.13      H1   O
ATOM   5177  CG2 THR    57      -1.384  18.032 -34.194  1.00 24.99      H1   C
ATOM   5178  C   THR    57      -4.097  15.956 -32.442  1.00 27.51      H1   C
ATOM   5179  O   THR    57      -3.968  14.814 -31.996  1.00 24.38      H1   O
ATOM   5180  N   ASP    58      -5.252  16.447 -32.882  1.00 26.49      H1   N
ATOM   5181  CA  ASP    58      -6.422  15.606 -33.141  1.00 26.22      H1   C
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ATOM	5182	CB	ASP	58	-7.491	15.830	-32.064	1.00	26.18	H1	C
ATOM	5183	CG	ASP	58	-6.973	15.585	-30.655	1.00	25.96	H1	C
ATOM	5184	OD1	ASP	58	-6.604	14.436	-30.349	1.00	28.58	H1	O
ATOM	5185	OD2	ASP	58	-6.941	16.538	-29.849	1.00	25.89	H1	O
ATOM	5186	C	ASP	58	-6.985	16.012	-34.506	1.00	25.72	H1	C
ATOM	5187	O	ASP	58	-7.107	17.200	-34.795	1.00	24.91	H1	O
ATOM	5188	N	TYR	59	-7.326	15.035	-35.339	1.00	25.29	H1	N
ATOM	5189	CA	TYR	59	-7.818	15.326	-36.681	1.00	25.55	H1	C
ATOM	5190	CB	TYR	59	-6.882	14.740	-37.743	1.00	24.54	H1	C
ATOM	5191	CG	TYR	59	-5.476	15.296	-37.709	1.00	27.17	H1	C
ATOM	5192	CD1	TYR	59	-5.247	16.653	-37.827	1.00	25.11	H1	C
ATOM	5193	CE1	TYR	59	-3.971	17.167	-37.782	1.00	28.62	H1	C
ATOM	5194	CD2	TYR	59	-4.378	14.458	-37.550	1.00	28.77	H1	C
ATOM	5195	CE2	TYR	59	-3.091	14.966	-37.510	1.00	29.01	H1	C
ATOM	5196	CZ	TYR	59	-2.895	16.324	-37.622	1.00	28.28	H1	C
ATOM	5197	OH	TYR	59	-1.624	16.855	-37.546	1.00	29.69	H1	O
ATOM	5198	C	TYR	59	-9.217	14.774	-36.912	1.00	26.93	H1	C
ATOM	5199	O	TYR	59	-9.610	13.775	-36.306	1.00	25.54	H1	O
ATOM	5200	N	ASN	60	-9.960	15.435	-37.795	1.00	25.64	H1	N
ATOM	5201	CA	ASN	60	-11.180	14.866	-38.339	1.00	27.20	H1	C
ATOM	5202	CB	ASN	60	-11.914	15.908	-39.193	1.00	27.87	H1	C
ATOM	5203	CG	ASN	60	-13.295	15.439	-39.636	1.00	29.92	H1	C
ATOM	5204	OD1	ASN	60	-13.552	14.242	-39.734	1.00	27.12	H1	O
ATOM	5205	ND2	ASN	60	-14.191	16.388	-39.906	1.00	27.79	H1	N
ATOM	5206	C	ASN	60	-10.766	13.682	-39.211	1.00	28.06	H1	C
ATOM	5207	O	ASN	60	-9.900	13.815	-40.076	1.00	26.75	H1	O
ATOM	5208	N	PRO	61	-11.383	12.509	-38.988	1.00	28.60	H1	N
ATOM	5209	CD	PRO	61	-12.379	12.272	-37.929	1.00	28.36	H1	C
ATOM	5210	CA	PRO	61	-11.103	11.288	-39.750	1.00	28.67	H1	C
ATOM	5211	CB	PRO	61	-12.156	10.300	-39.243	1.00	30.97	H1	C
ATOM	5212	CG	PRO	61	-12.471	10.774	-37.878	1.00	31.87	H1	C
ATOM	5213	C	PRO	61	-11.199	11.489	-41.254	1.00	29.49	H1	C
ATOM	5214	O	PRO	61	-10.505	10.830	-42.016	1.00	30.22	H1	O
ATOM	5215	N	SER	62	-12.064	12.402	-41.682	1.00	29.86	H1	N
ATOM	5216	CA	SER	62	-12.310	12.593	-43.105	1.00	30.55	H1	C
ATOM	5217	CB	SER	62	-13.540	13.472	-43.307	1.00	30.58	H1	C
ATOM	5218	OG	SER	62	-13.283	14.785	-42.854	1.00	31.57	H1	O
ATOM	5219	C	SER	62	-11.117	13.242	-43.799	1.00	31.98	H1	C
ATOM	5220	O	SER	62	-11.054	13.280	-45.026	1.00	32.06	H1	O
ATOM	5221	N	LEU	63	-10.180	13.763	-43.010	1.00	30.93	H1	N
ATOM	5222	CA	LEU	63	-9.039	14.489	-43.558	1.00	31.13	H1	C
ATOM	5223	CB	LEU	63	-9.206	15.987	-43.293	1.00	29.01	H1	C
ATOM	5224	CG	LEU	63	-10.329	16.656	-44.092	1.00	31.66	H1	C
ATOM	5225	CD1	LEU	63	-10.593	18.044	-43.535	1.00	28.33	H1	C
ATOM	5226	CD2	LEU	63	-9.939	16.731	-45.575	1.00	28.56	H1	C
ATOM	5227	C	LEU	63	-7.703	14.006	-42.988	1.00	30.49	H1	C
ATOM	5228	O	LEU	63	-6.640	14.416	-43.451	1.00	30.14	H1	O
ATOM	5229	N	LYS	64	-7.773	13.133	-41.989	1.00	30.11	H1	N
ATOM	5230	CA	LYS	64	-6.602	12.670	-41.257	1.00	32.57	H1	C
ATOM	5231	CB	LYS	64	-7.007	11.514	-40.337	1.00	36.89	H1	C
ATOM	5232	CG	LYS	64	-5.877	10.924	-39.511	1.00	40.28	H1	C
ATOM	5233	CD	LYS	64	-6.349	9.706	-38.706	1.00	44.49	H1	C
ATOM	5234	CE	LYS	64	-6.817	10.078	-37.290	1.00	48.41	H1	C
ATOM	5235	NZ	LYS	64	-8.133	10.801	-37.258	1.00	48.82	H1	N
ATOM	5236	C	LYS	64	-5.443	12.229	-42.153	1.00	34.71	H1	C
ATOM	5237	O	LYS	64	-4.282	12.504	-41.851	1.00	34.00	H1	O
ATOM	5238	N	SER	65	-5.753	11.547	-43.254	1.00	35.63	H1	N
ATOM	5239	CA	SER	65	-4.709	10.978	-44.099	1.00	38.29	H1	C
ATOM	5240	CB	SER	65	-5.299	9.924	-45.043	1.00	39.24	H1	C
ATOM	5241	OG	SER	65	-6.082	10.525	-46.065	1.00	44.72	H1	O
ATOM	5242	C	SER	65	-3.961	12.035	-44.916	1.00	39.01	H1	C
ATOM	5243	O	SER	65	-2.842	11.793	-45.378	1.00	39.10	H1	O
ATOM	5244	N	ARG	66	-4.573	13.204	-45.089	1.00	37.26	H1	N
ATOM	5245	CA	ARG	66	-3.992	14.240	-45.937	1.00	35.64	H1	C
ATOM	5246	CB	ARG	66	-5.018	14.691	-46.983	1.00	34.32	H1	C
ATOM	5247	CG	ARG	66	-5.552	13.575	-47.868	1.00	35.93	H1	C
ATOM	5248	CD	ARG	66	-6.867	13.993	-48.513	1.00	35.14	H1	C
ATOM	5249	NE	ARG	66	-6.689	15.244	-49.224	1.00	34.04	H1	N
ATOM	5250	CZ	ARG	66	-7.614	16.185	-49.375	1.00	32.60	H1	C
ATOM	5251	NH1	ARG	66	-8.833	16.046	-48.867	1.00	24.54	H1	N
ATOM	5252	NH2	ARG	66	-7.296	17.285	-50.040	1.00	30.85	H1	N
ATOM	5253	C	ARG	66	-3.495	15.462	-45.157	1.00	35.29	H1	C
ATOM	5254	O	ARG	66	-2.810	16.320	-45.713	1.00	36.20	H1	O
ATOM	5255	N	VAL	67	-3.848	15.549	-43.878	1.00	33.06	H1	N
ATOM	5256	CA	VAL	67	-3.663	16.787	-43.132	1.00	32.98	H1	C
ATOM	5257	CB	VAL	67	-4.944	17.169	-42.347	1.00	34.24	H1	C

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ATOM   5258  CG1 VAL    67      -4.725  18.471 -41.600  1.00 36.90      H1   C
ATOM   5259  CG2 VAL    67      -6.107  17.326 -43.297  1.00 38.06      H1   C
ATOM   5260  C   VAL    67      -2.513  16.725 -42.137  1.00 30.38      H1   C
ATOM   5261  O   VAL    67      -2.260  15.693 -41.525  1.00 29.73      H1   O
ATOM   5262  N   THR    68      -1.824  17.847 -41.978  1.00 29.72      H1   N
ATOM   5263  CA  THR    68      -0.890  18.021 -40.878  1.00 31.21      H1   C
ATOM   5264  CB  THR    68       0.581  17.868 -41.333  1.00 32.80      H1   C
ATOM   5265  OG1 THR    68       0.773  16.575 -41.917  1.00 36.87      H1   O
ATOM   5266  CG2 THR    68       1.521  18.003 -40.141  1.00 33.68      H1   C
ATOM   5267  C   THR    68      -1.067  19.413 -40.291  1.00 31.32      H1   C
ATOM   5268  O   THR    68      -1.042  20.420 -41.011  1.00 31.21      H1   O
ATOM   5269  N   ILE    69      -1.255  19.466 -38.980  1.00 27.82      H1   N
ATOM   5270  CA  ILE    69      -1.280  20.733 -38.274  1.00 26.11      H1   C
ATOM   5271  CB  ILE    69      -2.548  20.844 -37.404  1.00 22.50      H1   C
ATOM   5272  CG2 ILE    69      -2.490  22.088 -36.539  1.00 21.71      H1   C
ATOM   5273  CG1 ILE    69      -3.786  20.862 -38.309  1.00 22.12      H1   C
ATOM   5274  CD1 ILE    69      -5.111  20.739 -37.559  1.00 19.06      H1   C
ATOM   5275  C   ILE    69      -0.034  20.796 -37.399  1.00 26.90      H1   C
ATOM   5276  O   ILE    69       0.344  19.808 -36.763  1.00 26.17      H1   O
ATOM   5277  N   SER    70       0.620  21.950 -37.386  1.00 26.96      H1   N
ATOM   5278  CA  SER    70       1.825  22.112 -36.590  1.00 29.41      H1   C
ATOM   5279  CB  SER    70       3.066  22.082 -37.490  1.00 29.33      H1   C
ATOM   5280  OG  SER    70       2.970  23.059 -38.508  1.00 34.80      H1   O
ATOM   5281  C   SER    70       1.753  23.419 -35.829  1.00 29.15      H1   C
ATOM   5282  O   SER    70       0.891  24.257 -36.096  1.00 29.52      H1   O
ATOM   5283  N   VAL    71       2.647  23.588 -34.864  1.00 30.15      H1   N
ATOM   5284  CA  VAL    71       2.617  24.776 -34.029  1.00 30.65      H1   C
ATOM   5285  CB  VAL    71       1.970  24.461 -32.653  1.00 30.44      H1   C
ATOM   5286  CG1 VAL    71       2.819  23.450 -31.900  1.00 28.16      H1   C
ATOM   5287  CG2 VAL    71       1.804  25.744 -31.842  1.00 32.12      H1   C
ATOM   5288  C   VAL    71       4.026  25.320 -33.813  1.00 31.95      H1   C
ATOM   5289  O   VAL    71       4.994  24.564 -33.722  1.00 30.74      H1   O
ATOM   5290  N   ASP    72       4.134  26.639 -33.743  1.00 33.49      H1   N
ATOM   5291  CA  ASP    72       5.392  27.289 -33.405  1.00 35.48      H1   C
ATOM   5292  CB  ASP    72       5.876  28.143 -34.582  1.00 38.44      H1   C
ATOM   5293  CG  ASP    72       7.211  28.814 -34.310  1.00 40.69      H1   C
ATOM   5294  OD1 ASP    72       7.609  28.903 -33.131  1.00 41.96      H1   O
ATOM   5295  OD2 ASP    72       7.865  29.252 -35.279  1.00 43.43      H1   O
ATOM   5296  C   ASP    72       5.139  28.165 -32.187  1.00 35.14      H1   C
ATOM   5297  O   ASP    72       4.727  29.317 -32.316  1.00 33.42      H1   O
ATOM   5298  N   THR    73       5.376  27.614 -31.002  1.00 36.86      H1   N
ATOM   5299  CA  THR    73       5.012  28.306 -29.774  1.00 40.27      H1   C
ATOM   5300  CB  THR    73       5.158  27.390 -28.545  1.00 41.47      H1   C
ATOM   5301  OG1 THR    73       6.539  27.065 -28.356  1.00 44.90      H1   O
ATOM   5302  CG2 THR    73       4.360  26.103 -28.740  1.00 38.24      H1   C
ATOM   5303  C   THR    73       5.877  29.541 -29.569  1.00 40.62      H1   C
ATOM   5304  O   THR    73       5.410  30.547 -29.039  1.00 41.80      H1   O
ATOM   5305  N   SER    74       7.133  29.475 -29.998  1.00 41.71      H1   N
ATOM   5306  CA  SER    74       8.024  30.617 -29.833  1.00 42.42      H1   C
ATOM   5307  CB  SER    74       9.418  30.313 -30.403  1.00 42.23      H1   C
ATOM   5308  OG  SER    74       9.455  30.467 -31.811  1.00 45.59      H1   O
ATOM   5309  C   SER    74       7.423  31.820 -30.542  1.00 41.31      H1   C
ATOM   5310  O   SER    74       7.602  32.959 -30.111  1.00 43.56      H1   O
ATOM   5311  N   LYS    75       6.692  31.562 -31.622  1.00 40.07      H1   N
ATOM   5312  CA  LYS    75       6.081  32.635 -32.402  1.00 37.23      H1   C
ATOM   5313  CB  LYS    75       6.283  32.381 -33.894  1.00 40.70      H1   C
ATOM   5314  CG  LYS    75       7.640  32.786 -34.430  1.00 42.41      H1   C
ATOM   5315  CD  LYS    75       7.552  33.046 -35.925  1.00 45.61      H1   C
ATOM   5316  CE  LYS    75       8.933  33.131 -36.550  1.00 48.78      H1   C
ATOM   5317  NZ  LYS    75       9.706  31.882 -36.306  1.00 49.50      H1   N
ATOM   5318  C   LYS    75       4.591  32.813 -32.129  1.00 34.20      H1   C
ATOM   5319  O   LYS    75       3.958  33.725 -32.673  1.00 31.98      H1   O
ATOM   5320  N   LYS    76       4.031  31.939 -31.298  1.00 31.53      H1   N
ATOM   5321  CA  LYS    76       2.583  31.894 -31.099  1.00 30.16      H1   C
ATOM   5322  CB  LYS    76       2.129  33.120 -30.301  1.00 31.13      H1   C
ATOM   5323  CG  LYS    76       2.773  33.203 -28.914  1.00 32.98      H1   C
ATOM   5324  CD  LYS    76       2.476  34.526 -28.236  1.00 35.35      H1   C
ATOM   5325  CE  LYS    76       3.315  34.711 -26.976  1.00 42.53      H1   C
ATOM   5326  NZ  LYS    76       3.033  33.680 -25.934  1.00 45.06      H1   N
ATOM   5327  C   LYS    76       1.867  31.830 -32.451  1.00 28.30      H1   C
ATOM   5328  O   LYS    76       1.027  32.664 -32.773  1.00 26.79      H1   O
ATOM   5329  N   GLN    77       2.234  30.837 -33.248  1.00 28.03      H1   N
ATOM   5330  CA  GLN    77       1.610  30.613 -34.544  1.00 30.21      H1   C
ATOM   5331  CB  GLN    77       2.528  31.065 -35.685  1.00 29.25      H1   C
ATOM   5332  CG  GLN    77       2.730  32.565 -35.770  1.00 32.24      H1   C
ATOM   5333  CD  GLN    77       3.757  32.963 -36.834  1.00 31.77      H1   C
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ATOM   5334  OE1 GLN    77       4.144  34.126 -36.922  1.00 35.74      H1   O
ATOM   5335  NE2 GLN    77       4.198  32.002 -37.634  1.00 28.74      H1   N
ATOM   5336  C   GLN    77       1.353  29.124 -34.680  1.00 29.69      H1   C
ATOM   5337  O   GLN    77       2.085  28.307 -34.119  1.00 31.84      H1   O
ATOM   5338  N   PHE    78       0.313  28.767 -35.417  1.00 28.53      H1   N
ATOM   5339  CA  PHE    78       0.149  27.385 -35.817  1.00 28.21      H1   C
ATOM   5340  CB  PHE    78      -0.842  26.658 -34.888  1.00 28.77      H1   C
ATOM   5341  CG  PHE    78      -2.240  27.200 -34.928  1.00 28.71      H1   C
ATOM   5342  CD1 PHE    78      -3.272  26.444 -35.469  1.00 28.64      H1   C
ATOM   5343  CD2 PHE    78      -2.531  28.452 -34.412  1.00 28.33      H1   C
ATOM   5344  CE1 PHE    78      -4.568  26.928 -35.494  1.00 28.19      H1   C
ATOM   5345  CE2 PHE    78      -3.824  28.941 -34.434  1.00 28.05      H1   C
ATOM   5346  CZ  PHE    78      -4.844  28.177 -34.978  1.00 27.78      H1   C
ATOM   5347  C   PHE    78      -0.291  27.334 -37.267  1.00 26.74      H1   C
ATOM   5348  O   PHE    78      -0.746  28.329 -37.820  1.00 26.40      H1   O
ATOM   5349  N   SER    79      -0.121  26.173 -37.884  1.00 26.74      H1   N
ATOM   5350  CA  SER    79      -0.190  26.058 -39.332  1.00 28.66      H1   C
ATOM   5351  CB  SER    79       1.217  25.906 -39.911  1.00 26.94      H1   C
ATOM   5352  OG  SER    79       1.991  27.059 -39.670  1.00 37.18      H1   O
ATOM   5353  C   SER    79      -1.019  24.857 -39.751  1.00 27.22      H1   C
ATOM   5354  O   SER    79      -1.198  23.915 -38.982  1.00 28.89      H1   O
ATOM   5355  N   LEU    80      -1.492  24.899 -40.988  1.00 28.24      H1   N
ATOM   5356  CA  LEU    80      -2.221  23.797 -41.595  1.00 28.21      H1   C
ATOM   5357  CB  LEU    80      -3.657  24.228 -41.873  1.00 27.82      H1   C
ATOM   5358  CG  LEU    80      -4.499  23.249 -42.689  1.00 29.21      H1   C
ATOM   5359  CD1 LEU    80      -4.684  21.959 -41.906  1.00 25.31      H1   C
ATOM   5360  CD2 LEU    80      -5.843  23.888 -43.000  1.00 29.81      H1   C
ATOM   5361  C   LEU    80      -1.560  23.383 -42.910  1.00 29.34      H1   C
ATOM   5362  O   LEU    80      -1.283  24.224 -43.764  1.00 29.99      H1   O
ATOM   5363  N   LYS    81      -1.315  22.088 -43.072  1.00 30.34      H1   N
ATOM   5364  CA  LYS    81      -0.952  21.543 -44.372  1.00 32.50      H1   C
ATOM   5365  CB  LYS    81       0.433  20.880 -44.315  1.00 36.42      H1   C
ATOM   5366  CG  LYS    81       1.585  21.867 -44.454  1.00 43.74      H1   C
ATOM   5367  CD  LYS    81       2.892  21.145 -44.778  1.00 47.76      H1   C
ATOM   5368  CE  LYS    81       3.901  22.096 -45.405  1.00 48.81      H1   C
ATOM   5369  NZ  LYS    81       3.752  23.471 -44.859  1.00 49.14      H1   N
ATOM   5370  C   LYS    81      -1.976  20.514 -44.830  1.00 32.18      H1   C
ATOM   5371  O   LYS    81      -2.286  19.568 -44.102  1.00 32.47      H1   O
ATOM   5372  N   LEU    82      -2.493  20.694 -46.041  1.00 31.45      H1   N
ATOM   5373  CA  LEU    82      -3.388  19.708 -46.642  1.00 31.74      H1   C
ATOM   5374  CB  LEU    82      -4.777  20.322 -46.853  1.00 29.16      H1   C
ATOM   5375  CG  LEU    82      -5.901  19.419 -47.374  1.00 29.44      H1   C
ATOM   5376  CD1 LEU    82      -6.172  18.295 -46.380  1.00 26.17      H1   C
ATOM   5377  CD2 LEU    82      -7.163  20.251 -47.579  1.00 27.84      H1   C
ATOM   5378  C   LEU    82      -2.815  19.252 -47.980  1.00 31.38      H1   C
ATOM   5379  O   LEU    82      -2.719  20.049 -48.913  1.00 30.59      H1   O
ATOM   5380  N   ASN    83      -2.439  17.976 -48.068  1.00 33.35      H1   N
ATOM   5381  CA  ASN    83      -1.862  17.403 -49.294  1.00 34.64      H1   C
ATOM   5382  CB  ASN    83      -1.143  16.079 -48.999  1.00 37.82      H1   C
ATOM   5383  CG  ASN    83      -0.053  16.223 -47.975  1.00 43.06      H1   C
ATOM   5384  OD1 ASN    83       0.637  17.244 -47.924  1.00 45.32      H1   O
ATOM   5385  ND2 ASN    83       0.114  15.198 -47.141  1.00 42.01      H1   N
ATOM   5386  C   ASN    83      -2.907  17.112 -50.368  1.00 34.00      H1   C
ATOM   5387  O   ASN    83      -4.092  16.917 -50.066  1.00 31.80      H1   O
ATOM   5388  N   SER    84      -2.437  17.067 -51.616  1.00 32.98      H1   N
ATOM   5389  CA  SER    84      -3.175  16.485 -52.733  1.00 32.51      H1   C
ATOM   5390  CB  SER    84      -3.226  14.963 -52.596  1.00 33.13      H1   C
ATOM   5391  OG  SER    84      -1.930  14.422 -52.419  1.00 37.34      H1   O
ATOM   5392  C   SER    84      -4.594  17.013 -52.848  1.00 32.10      H1   C
ATOM   5393  O   SER    84      -5.544  16.235 -52.918  1.00 32.45      H1   O
ATOM   5394  N   VAL    85      -4.737  18.331 -52.870  1.00 30.78      H1   N
ATOM   5395  CA  VAL    85      -6.054  18.939 -52.891  1.00 31.67      H1   C
ATOM   5396  CB  VAL    85      -5.955  20.468 -52.767  1.00 31.43      H1   C
ATOM   5397  CG1 VAL    85      -5.392  20.841 -51.400  1.00 31.90      H1   C
ATOM   5398  CG2 VAL    85      -5.072  21.019 -53.869  1.00 31.67      H1   C
ATOM   5399  C   VAL    85      -6.815  18.589 -54.164  1.00 32.60      H1   C
ATOM   5400  O   VAL    85      -6.214  18.241 -55.180  1.00 34.22      H1   O
ATOM   5401  N   THR    86      -8.142  18.659 -54.090  1.00 31.22      H1   N
ATOM   5402  CA  THR    86      -8.995  18.603 -55.275  1.00 31.00      H1   C
ATOM   5403  CB  THR    86      -9.725  17.249 -55.400  1.00 31.98      H1   C
ATOM   5404  OG1 THR    86     -10.653  17.102 -54.316  1.00 34.46      H1   O
ATOM   5405  CG2 THR    86      -8.725  16.091 -55.368  1.00 32.26      H1   C
ATOM   5406  C   THR    86     -10.038  19.709 -55.152  1.00 30.48      H1   C
ATOM   5407  O   THR    86     -10.059  20.439 -54.162  1.00 30.01      H1   O
ATOM   5408  N   ALA    87     -10.904  19.836 -56.150  1.00 28.60      H1   N
ATOM   5409  CA  ALA    87     -11.891  20.908 -56.145  1.00 29.42      H1   C
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ATOM   5410  CB  ALA    87     -12.797  20.806 -57.386  1.00 25.14      H1   C
ATOM   5411  C   ALA    87     -12.734  20.876 -54.871  1.00 27.88      H1   C
ATOM   5412  O   ALA    87     -13.204  21.908 -54.415  1.00 29.57      H1   O
ATOM   5413  N   ALA    88     -12.915  19.689 -54.301  1.00 26.65      H1   N
ATOM   5414  CA  ALA    88     -13.753  19.515 -53.115  1.00 26.47      H1   C
ATOM   5415  CB  ALA    88     -14.045  18.023 -52.907  1.00 22.83      H1   C
ATOM   5416  C   ALA    88     -13.147  20.115 -51.834  1.00 27.55      H1   C
ATOM   5417  O   ALA    88     -13.818  20.204 -50.800  1.00 25.90      H1   O
ATOM   5418  N   ASP    89     -11.879  20.514 -51.896  1.00 26.38      H1   N
ATOM   5419  CA  ASP    89     -11.249  21.202 -50.773  1.00 26.18      H1   C
ATOM   5420  CB  ASP    89      -9.752  20.899 -50.735  1.00 24.35      H1   C
ATOM   5421  CG  ASP    89      -9.462  19.433 -50.473  1.00 25.95      H1   C
ATOM   5422  OD1 ASP    89      -8.997  18.882 -49.486  1.00 26.03      H1   O
ATOM   5423  OD2 ASP    89      -8.704  18.831 -51.262  1.00 25.18      H1   O
ATOM   5424  C   ASP    89     -11.461  22.712 -50.844  1.00 25.45      H1   C
ATOM   5425  O   ASP    89     -11.031  23.446 -49.960  1.00 25.56      H1   O
ATOM   5426  N   THR    90     -12.124  23.169 -51.901  1.00 24.03      H1   N
ATOM   5427  CA  THR    90     -12.500  24.572 -52.021  1.00 22.80      H1   C
ATOM   5428  CB  THR    90     -13.279  24.808 -53.327  1.00 23.31      H1   C
ATOM   5429  OG1 THR    90     -12.440  24.464 -54.434  1.00 23.59      H1   O
ATOM   5430  CG2 THR    90     -13.720  26.278 -53.458  1.00 22.37      H1   C
ATOM   5431  C   THR    90     -13.363  24.994 -50.833  1.00 23.24      H1   C
ATOM   5432  O   THR    90     -14.360  24.342 -50.512  1.00 23.24      H1   O
ATOM   5433  N   ALA    91     -12.975  26.083 -50.178  1.00 22.25      H1   N
ATOM   5434  CA  ALA    91     -13.673  26.534 -48.978  1.00 22.73      H1   C
ATOM   5435  CB  ALA    91     -13.718  25.406 -47.943  1.00 18.43      H1   C
ATOM   5436  C   ALA    91     -12.982  27.757 -48.378  1.00 24.17      H1   C
ATOM   5437  O   ALA    91     -11.855  28.094 -48.755  1.00 25.58      H1   O
ATOM   5438  N   VAL    92     -13.663  28.420 -47.447  1.00 23.43      H1   N
ATOM   5439  CA  VAL    92     -13.002  29.380 -46.575  1.00 23.10      H1   C
ATOM   5440  CB  VAL    92     -13.983  30.477 -46.087  1.00 24.14      H1   C
ATOM   5441  CG1 VAL    92     -13.263  31.430 -45.107  1.00 23.43      H1   C
ATOM   5442  CG2 VAL    92     -14.535  31.260 -47.281  1.00 23.45      H1   C
ATOM   5443  C   VAL    92     -12.436  28.627 -45.372  1.00 24.98      H1   C
ATOM   5444  O   VAL    92     -13.160  27.903 -44.675  1.00 26.01      H1   O
ATOM   5445  N   TYR    93     -11.137  28.779 -45.143  1.00 23.58      H1   N
ATOM   5446  CA  TYR    93     -10.479  28.124 -44.019  1.00 23.55      H1   C
ATOM   5447  CB  TYR    93      -9.135  27.537 -44.471  1.00 20.25      H1   C
ATOM   5448  CG  TYR    93      -9.272  26.302 -45.342  1.00 20.65      H1   C
ATOM   5449  CD1 TYR    93      -8.895  25.048 -44.871  1.00 19.71      H1   C
ATOM   5450  CE1 TYR    93      -9.047  23.911 -45.651  1.00 20.45      H1   C
ATOM   5451  CD2 TYR    93      -9.804  26.385 -46.620  1.00 21.99      H1   C
ATOM   5452  CE2 TYR    93      -9.964  25.250 -47.412  1.00 21.69      H1   C
ATOM   5453  CZ  TYR    93      -9.583  24.020 -46.918  1.00 21.47      H1   C
ATOM   5454  OH  TYR    93      -9.757  22.894 -47.687  1.00 24.56      H1   O
ATOM   5455  C   TYR    93     -10.263  29.110 -42.869  1.00 24.76      H1   C
ATOM   5456  O   TYR    93      -9.700  30.190 -43.064  1.00 26.10      H1   O
ATOM   5457  N   TYR    94     -10.719  28.735 -41.676  1.00 22.97      H1   N
ATOM   5458  CA  TYR    94     -10.583  29.576 -40.489  1.00 23.93      H1   C
ATOM   5459  CB  TYR    94     -11.946  29.821 -39.846  1.00 22.58      H1   C
ATOM   5460  CG  TYR    94     -12.952  30.548 -40.702  1.00 25.89      H1   C
ATOM   5461  CD1 TYR    94     -12.938  31.936 -40.801  1.00 25.89      H1   C
ATOM   5462  CE1 TYR    94     -13.908  32.614 -41.531  1.00 25.38      H1   C
ATOM   5463  CD2 TYR    94     -13.962  29.850 -41.365  1.00 26.81      H1   C
ATOM   5464  CE2 TYR    94     -14.935  30.516 -42.095  1.00 25.75      H1   C
ATOM   5465  CZ  TYR    94     -14.906  31.897 -42.172  1.00 27.48      H1   C
ATOM   5466  OH  TYR    94     -15.889  32.561 -42.874  1.00 28.03      H1   O
ATOM   5467  C   TYR    94      -9.691  28.909 -39.446  1.00 24.14      H1   C
ATOM   5468  O   TYR    94      -9.695  27.681 -39.310  1.00 22.75      H1   O
ATOM   5469  N   CYS    95      -8.938  29.716 -38.705  1.00 24.10      H1   N
ATOM   5470  CA  CYS    95      -8.466  29.294 -37.395  1.00 25.55      H1   C
ATOM   5471  C   CYS    95      -9.410  29.829 -36.338  1.00 25.49      H1   C
ATOM   5472  O   CYS    95     -10.104  30.819 -36.560  1.00 24.75      H1   O
ATOM   5473  CB  CYS    95      -7.034  29.781 -37.107  1.00 27.36      H1   C
ATOM   5474  SG  CYS    95      -6.561  31.491 -37.539  1.00 31.48      H1   S
ATOM   5475  N   ALA    96      -9.439  29.157 -35.192  1.00 24.09      H1   N
ATOM   5476  CA  ALA    96     -10.354  29.504 -34.117  1.00 22.01      H1   C
ATOM   5477  CB  ALA    96     -11.730  28.887 -34.380  1.00 20.76      H1   C
ATOM   5478  C   ALA    96      -9.782  28.977 -32.813  1.00 22.32      H1   C
ATOM   5479  O   ALA    96      -9.166  27.910 -32.775  1.00 22.28      H1   O
ATOM   5480  N   ARG    97      -9.997  29.723 -31.739  1.00 22.97      H1   N
ATOM   5481  CA  ARG    97      -9.439  29.355 -30.450  1.00 22.39      H1   C
ATOM   5482  CB  ARG    97      -9.076  30.614 -29.668  1.00 22.09      H1   C
ATOM   5483  CG  ARG    97      -8.510  30.352 -28.276  1.00 21.83      H1   C
ATOM   5484  CD  ARG    97      -8.172  31.672 -27.581  1.00 23.69      H1   C
ATOM   5485  NE  ARG    97      -9.342  32.258 -26.927  1.00 23.06      H1   N
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ATOM   5486  CZ  ARG    97      -9.273  33.201 -25.993  1.00 26.31      H1   C
ATOM   5487  NH1 ARG    97      -8.090  33.668 -25.613  1.00 26.46      H1   N
ATOM   5488  NH2 ARG    97     -10.379  33.662 -25.422  1.00 23.30      H1   N
ATOM   5489  C   ARG    97     -10.426  28.533 -29.636  1.00 24.53      H1   C
ATOM   5490  O   ARG    97     -11.632  28.801 -29.658  1.00 22.12      H1   O
ATOM   5491  N   GLY    98      -9.905  27.537 -28.921  1.00 22.25      H1   N
ATOM   5492  CA  GLY    98     -10.606  27.020 -27.760  1.00 23.81      H1   C
ATOM   5493  C   GLY    98     -11.259  25.658 -27.913  1.00 23.68      H1   C
ATOM   5494  O   GLY    98     -11.391  25.128 -29.023  1.00 23.49      H1   O
ATOM   5495  N   GLN    99     -11.668  25.095 -26.780  1.00 22.67      H1   N
ATOM   5496  CA  GLN    99     -12.293  23.780 -26.749  1.00 23.45      H1   C
ATOM   5497  CB  GLN    99     -11.253  22.706 -26.419  1.00 22.97      H1   C
ATOM   5498  CG  GLN    99     -11.774  21.269 -26.560  1.00 20.76      H1   C
ATOM   5499  CD  GLN    99     -10.690  20.232 -26.328  1.00 22.06      H1   C
ATOM   5500  OE1 GLN    99     -10.482  19.768 -25.202  1.00 22.02      H1   O
ATOM   5501  NE2 GLN    99      -9.992  19.860 -27.395  1.00 21.38      H1   N
ATOM   5502  C   GLN    99     -13.436  23.724 -25.725  1.00 24.22      H1   C
ATOM   5503  O   GLN    99     -14.550  23.329 -26.056  1.00 24.20      H1   O
ATOM   5504  N   LEU   100     -13.161  24.117 -24.485  1.00 23.64      H1   N
ATOM   5505  CA  LEU   100     -14.182  24.086 -23.448  1.00 24.07      H1   C
ATOM   5506  CB  LEU   100     -13.543  24.137 -22.055  1.00 23.74      H1   C
ATOM   5507  CG  LEU   100     -12.653  22.935 -21.697  1.00 23.65      H1   C
ATOM   5508  CD1 LEU   100     -12.218  23.039 -20.250  1.00 21.96      H1   C
ATOM   5509  CD2 LEU   100     -13.409  21.629 -21.918  1.00 22.57      H1   C
ATOM   5510  C   LEU   100     -15.133  25.257 -23.642  1.00 26.71      H1   C
ATOM   5511  O   LEU   100     -16.334  25.145 -23.386  1.00 27.08      H1   O
ATOM   5512  N   VAL   101     -14.591  26.380 -24.104  1.00 26.05      H1   N
ATOM   5513  CA  VAL   101     -15.413  27.446 -24.665  1.00 26.77      H1   C
ATOM   5514  CB  VAL   101     -15.146  28.799 -23.957  1.00 25.01      H1   C
ATOM   5515  CG1 VAL   101     -16.083  29.869 -24.501  1.00 25.38      H1   C
ATOM   5516  CG2 VAL   101     -15.361  28.646 -22.453  1.00 24.75      H1   C
ATOM   5517  C   VAL   101     -15.034  27.530 -26.141  1.00 26.59      H1   C
ATOM   5518  O   VAL   101     -14.149  28.290 -26.528  1.00 26.80      H1   O
ATOM   5519  N   PRO   102     -15.687  26.710 -26.980  1.00 25.65      H1   N
ATOM   5520  CD  PRO   102     -16.834  25.848 -26.646  1.00 23.49      H1   C
ATOM   5521  CA  PRO   102     -15.148  26.427 -28.315  1.00 23.89      H1   C
ATOM   5522  CB  PRO   102     -15.863  25.141 -28.729  1.00 23.18      H1   C
ATOM   5523  CG  PRO   102     -17.147  25.158 -27.954  1.00 25.40      H1   C
ATOM   5524  C   PRO   102     -15.315  27.542 -29.339  1.00 24.72      H1   C
ATOM   5525  O   PRO   102     -16.411  28.061 -29.550  1.00 24.05      H1   O
ATOM   5526  N   PHE   103     -14.206  27.900 -29.974  1.00 22.69      H1   N
ATOM   5527  CA  PHE   103     -14.223  28.841 -31.080  1.00 22.97      H1   C
ATOM   5528  CB  PHE   103     -15.059  28.262 -32.240  1.00 20.62      H1   C
ATOM   5529  CG  PHE   103     -14.912  26.755 -32.412  1.00 21.68      H1   C
ATOM   5530  CD1 PHE   103     -16.005  25.970 -32.759  1.00 17.84      H1   C
ATOM   5531  CD2 PHE   103     -13.693  26.125 -32.185  1.00 19.12      H1   C
ATOM   5532  CE1 PHE   103     -15.888  24.595 -32.868  1.00 19.25      H1   C
ATOM   5533  CE2 PHE   103     -13.565  24.745 -32.294  1.00 17.07      H1   C
ATOM   5534  CZ  PHE   103     -14.666  23.978 -32.634  1.00 18.07      H1   C
ATOM   5535  C   PHE   103     -14.781  30.195 -30.619  1.00 23.73      H1   C
ATOM   5536  O   PHE   103     -15.712  30.724 -31.217  1.00 24.85      H1   O
ATOM   5537  N   ASP   104     -14.215  30.752 -29.548  1.00 24.65      H1   N
ATOM   5538  CA  ASP   104     -14.669  32.050 -29.069  1.00 24.76      H1   C
ATOM   5539  CB  ASP   104     -14.406  32.217 -27.563  1.00 25.91      H1   C
ATOM   5540  CG  ASP   104     -12.945  32.005 -27.175  1.00 30.36      H1   C
ATOM   5541  OD1 ASP   104     -12.122  31.635 -28.036  1.00 31.12      H1   O
ATOM   5542  OD2 ASP   104     -12.619  32.212 -25.987  1.00 28.71      H1   O
ATOM   5543  C   ASP   104     -14.021  33.180 -29.856  1.00 27.29      H1   C
ATOM   5544  O   ASP   104     -14.586  34.262 -29.971  1.00 27.52      H1   O
ATOM   5545  N   TYR   105     -12.841  32.920 -30.411  1.00 26.93      H1   N
ATOM   5546  CA  TYR   105     -12.227  33.847 -31.355  1.00 26.35      H1   C
ATOM   5547  CB  TYR   105     -10.971  34.479 -30.743  1.00 25.73      H1   C
ATOM   5548  CG  TYR   105     -11.278  35.498 -29.675  1.00 28.28      H1   C
ATOM   5549  CD1 TYR   105     -11.345  35.133 -28.334  1.00 26.26      H1   C
ATOM   5550  CE1 TYR   105     -11.661  36.060 -27.359  1.00 27.70      H1   C
ATOM   5551  CD2 TYR   105     -11.534  36.825 -30.012  1.00 29.38      H1   C
ATOM   5552  CE2 TYR   105     -11.855  37.756 -29.044  1.00 29.45      H1   C
ATOM   5553  CZ  TYR   105     -11.918  37.368 -27.722  1.00 29.50      H1   C
ATOM   5554  OH  TYR   105     -12.255  38.299 -26.766  1.00 32.67      H1   O
ATOM   5555  C   TYR   105     -11.869  33.131 -32.657  1.00 25.43      H1   C
ATOM   5556  O   TYR   105     -11.426  31.982 -32.641  1.00 24.81      H1   O
ATOM   5557  N   TRP   106     -12.065  33.826 -33.774  1.00 24.15      H1   N
ATOM   5558  CA  TRP   106     -11.831  33.273 -35.106  1.00 25.25      H1   C
ATOM   5559  CB  TRP   106     -13.145  33.160 -35.879  1.00 21.93      H1   C
ATOM   5560  CG  TRP   106     -14.111  32.140 -35.362  1.00 22.83      H1   C
ATOM   5561  CD2 TRP   106     -14.605  30.995 -36.072  1.00 21.41      H1   C
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ATOM   5562  CE2 TRP   106     -15.565  30.379 -35.250  1.00 20.48      H1   C
ATOM   5563  CE3 TRP   106     -14.328  30.437 -37.326  1.00 20.75      H1   C
ATOM   5564  CD1 TRP   106     -14.767  32.164 -34.170  1.00 19.87      H1   C
ATOM   5565  NE1 TRP   106     -15.646  31.112 -34.095  1.00 22.85      H1   N
ATOM   5566  CZ2 TRP   106     -16.258  29.229 -35.637  1.00 20.26      H1   C
ATOM   5567  CZ3 TRP   106     -15.015  29.297 -37.711  1.00 20.27      H1   C
ATOM   5568  CH2 TRP   106     -15.970  28.705 -36.868  1.00 23.31      H1   C
ATOM   5569  C   TRP   106     -10.902  34.181 -35.906  1.00 27.18      H1   C
ATOM   5570  O   TRP   106     -10.923  35.400 -35.741  1.00 26.73      H1   O
ATOM   5571  N   GLY   107     -10.114  33.587 -36.797  1.00 27.12      H1   N
ATOM   5572  CA  GLY   107      -9.418  34.377 -37.797  1.00 27.10      H1   C
ATOM   5573  C   GLY   107     -10.379  34.963 -38.822  1.00 28.57      H1   C
ATOM   5574  O   GLY   107     -11.583  34.686 -38.790  1.00 27.48      H1   O
ATOM   5575  N   GLN   108      -9.852  35.771 -39.737  1.00 28.26      H1   N
ATOM   5576  CA  GLN   108     -10.680  36.424 -40.743  1.00 30.59      H1   C
ATOM   5577  CB  GLN   108      -9.980  37.684 -41.275  1.00 33.27      H1   C
ATOM   5578  CG  GLN   108      -8.939  37.432 -42.371  1.00 36.99      H1   C
ATOM   5579  CD  GLN   108      -7.566  37.046 -41.831  1.00 42.07      H1   C
ATOM   5580  OE1 GLN   108      -7.395  36.782 -40.633  1.00 42.16      H1   O
ATOM   5581  NE2 GLN   108      -6.577  37.012 -42.720  1.00 40.22      H1   N
ATOM   5582  C   GLN   108     -10.976  35.464 -41.895  1.00 30.47      H1   C
ATOM   5583  O   GLN   108     -11.868  35.710 -42.707  1.00 31.09      H1   O
ATOM   5584  N   GLY   109     -10.223  34.369 -41.953  1.00 29.81      H1   N
ATOM   5585  CA  GLY   109     -10.492  33.323 -42.923  1.00 29.09      H1   C
ATOM   5586  C   GLY   109      -9.718  33.498 -44.216  1.00 28.69      H1   C
ATOM   5587  O   GLY   109      -9.425  34.618 -44.619  1.00 28.50      H1   O
ATOM   5588  N   THR   110      -9.386  32.389 -44.866  1.00 28.50      H1   N
ATOM   5589  CA  THR   110      -8.682  32.431 -46.139  1.00 28.41      H1   C
ATOM   5590  CB  THR   110      -7.285  31.793 -46.021  1.00 29.01      H1   C
ATOM   5591  OG1 THR   110      -6.522  32.494 -45.034  1.00 32.03      H1   O
ATOM   5592  CG2 THR   110      -6.554  31.856 -47.356  1.00 29.38      H1   C
ATOM   5593  C   THR   110      -9.476  31.669 -47.193  1.00 29.43      H1   C
ATOM   5594  O   THR   110      -9.712  30.471 -47.053  1.00 27.15      H1   O
ATOM   5595  N   LEU   111      -9.890  32.364 -48.246  1.00 29.78      H1   N
ATOM   5596  CA  LEU   111     -10.616  31.723 -49.328  1.00 29.48      H1   C
ATOM   5597  CB  LEU   111     -11.234  32.766 -50.258  1.00 31.55      H1   C
ATOM   5598  CG  LEU   111     -11.853  32.182 -51.537  1.00 36.06      H1   C
ATOM   5599  CD1 LEU   111     -13.054  31.310 -51.180  1.00 35.44      H1   C
ATOM   5600  CD2 LEU   111     -12.272  33.308 -52.470  1.00 33.87      H1   C
ATOM   5601  C   LEU   111      -9.659  30.839 -50.109  1.00 30.13      H1   C
ATOM   5602  O   LEU   111      -8.661  31.308 -50.664  1.00 30.30      H1   O
ATOM   5603  N   VAL   112      -9.953  29.547 -50.129  1.00 28.78      H1   N
ATOM   5604  CA  VAL   112      -9.134  28.603 -50.862  1.00 27.19      H1   C
ATOM   5605  CB  VAL   112      -8.700  27.434 -49.964  1.00 27.32      H1   C
ATOM   5606  CG1 VAL   112      -7.986  26.383 -50.793  1.00 26.80      H1   C
ATOM   5607  CG2 VAL   112      -7.788  27.951 -48.857  1.00 25.55      H1   C
ATOM   5608  C   VAL   112      -9.934  28.071 -52.039  1.00 28.91      H1   C
ATOM   5609  O   VAL   112     -11.012  27.493 -51.867  1.00 26.37      H1   O
ATOM   5610  N   THR   113      -9.405  28.288 -53.237  1.00 28.59      H1   N
ATOM   5611  CA  THR   113     -10.076  27.865 -54.457  1.00 31.25      H1   C
ATOM   5612  CB  THR   113     -10.365  29.071 -55.379  1.00 30.80      H1   C
ATOM   5613  OG1 THR   113     -11.220  29.992 -54.691  1.00 33.89      H1   O
ATOM   5614  CG2 THR   113     -11.064  28.623 -56.658  1.00 32.74      H1   C
ATOM   5615  C   THR   113      -9.190  26.868 -55.183  1.00 31.51      H1   C
ATOM   5616  O   THR   113      -8.041  27.163 -55.506  1.00 32.27      H1   O
ATOM   5617  N   VAL   114      -9.721  25.674 -55.406  1.00 32.99      H1   N
ATOM   5618  CA  VAL   114      -8.990  24.643 -56.122  1.00 34.40      H1   C
ATOM   5619  CB  VAL   114      -8.865  23.356 -55.288  1.00 32.69      H1   C
ATOM   5620  CG1 VAL   114      -8.002  22.346 -56.026  1.00 30.00      H1   C
ATOM   5621  CG2 VAL   114      -8.294  23.671 -53.916  1.00 32.66      H1   C
ATOM   5622  C   VAL   114      -9.745  24.305 -57.401  1.00 37.41      H1   C
ATOM   5623  O   VAL   114     -10.881  23.828 -57.349  1.00 37.23      H1   O
ATOM   5624  N   SER   115      -9.119  24.564 -58.544  1.00 39.18      H1   N
ATOM   5625  CA  SER   115      -9.668  24.115 -59.813  1.00 44.39      H1   C
ATOM   5626  CB  SER   115     -10.785  25.059 -60.277  1.00 45.61      H1   C
ATOM   5627  OG  SER   115     -10.365  25.884 -61.347  1.00 45.98      H1   O
ATOM   5628  C   SER   115      -8.586  24.014 -60.881  1.00 46.47      H1   C
ATOM   5629  O   SER   115      -7.543  24.667 -60.796  1.00 45.73      H1   O
ATOM   5630  N   SER   116      -8.841  23.177 -61.880  1.00 49.79      H1   N
ATOM   5631  CA  SER   116      -7.891  22.949 -62.961  1.00 53.39      H1   C
ATOM   5632  CB  SER   116      -7.763  21.450 -63.230  1.00 52.69      H1   C
ATOM   5633  OG  SER   116      -9.040  20.871 -63.434  1.00 54.43      H1   O
ATOM   5634  C   SER   116      -8.348  23.658 -64.230  1.00 55.63      H1   C
ATOM   5635  O   SER   116      -7.679  23.595 -65.261  1.00 57.09      H1   O
ATOM   5636  N   ALA   117      -9.491  24.332 -64.146  1.00 57.80      H1   N
ATOM   5637  CA  ALA   117     -10.149  24.878 -65.329  1.00 60.79      H1   C
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ATOM   5638  CB  ALA   117     -11.572  25.305 -64.983  1.00 60.52      H1   C
ATOM   5639  C   ALA   117      -9.391  26.049 -65.946  1.00 62.07      H1   C
ATOM   5640  O   ALA   117      -8.832  26.890 -65.240  1.00 62.18      H1   O
ATOM   5641  N   SER   118      -9.380  26.095 -67.274  1.00 64.43      H1   N
ATOM   5642  CA  SER   118      -8.766  27.201 -67.996  1.00 65.80      H1   C
ATOM   5643  CB  SER   118      -7.636  26.692 -68.890  1.00 66.05      H1   C
ATOM   5644  OG  SER   118      -6.810  27.764 -69.309  1.00 68.87      H1   O
ATOM   5645  C   SER   118      -9.807  27.920 -68.845  1.00 65.94      H1   C
ATOM   5646  O   SER   118     -10.856  27.358 -69.164  1.00 65.51      H1   O
ATOM   5647  N   THR   119      -9.505  29.161 -69.211  1.00 66.97      H1   N
ATOM   5648  CA  THR   119     -10.462  30.027 -69.893  1.00 67.67      H1   C
ATOM   5649  CB  THR   119      -9.775  31.297 -70.430  1.00 67.99      H1   C
ATOM   5650  OG1 THR   119      -9.195  32.026 -69.339  1.00 68.98      H1   O
ATOM   5651  CG2 THR   119     -10.784  32.181 -71.146  1.00 67.51      H1   C
ATOM   5652  C   THR   119     -11.167  29.336 -71.054  1.00 67.83      H1   C
ATOM   5653  O   THR   119     -10.543  28.631 -71.848  1.00 67.72      H1   O
ATOM   5654  N   LYS   120     -12.477  29.546 -71.141  1.00 67.80      H1   N
ATOM   5655  CA  LYS   120     -13.282  28.985 -72.217  1.00 68.20      H1   C
ATOM   5656  CB  LYS   120     -13.685  27.545 -71.889  1.00 67.62      H1   C
ATOM   5657  CG  LYS   120     -14.595  26.908 -72.931  1.00 67.51      H1   C
ATOM   5658  CD  LYS   120     -15.178  25.590 -72.436  1.00 69.11      H1   C
ATOM   5659  CE  LYS   120     -16.168  25.004 -73.437  1.00 69.25      H1   C
ATOM   5660  NZ  LYS   120     -17.270  25.956 -73.782  1.00 70.85      H1   N
ATOM   5661  C   LYS   120     -14.537  29.824 -72.432  1.00 69.04      H1   C
ATOM   5662  O   LYS   120     -15.329  30.023 -71.508  1.00 69.38      H1   O
ATOM   5663  N   GLY   121     -14.715  30.308 -73.659  1.00 69.06      H1   N
ATOM   5664  CA  GLY   121     -15.886  31.102 -73.982  1.00 68.10      H1   C
ATOM   5665  C   GLY   121     -17.159  30.283 -73.922  1.00 68.13      H1   C
ATOM   5666  O   GLY   121     -17.129  29.061 -74.089  1.00 67.67      H1   O
ATOM   5667  N   PRO   122     -18.303  30.936 -73.678  1.00 67.96      H1   N
ATOM   5668  CD  PRO   122     -18.403  32.385 -73.433  1.00 67.50      H1   C
ATOM   5669  CA  PRO   122     -19.596  30.260 -73.523  1.00 68.40      H1   C
ATOM   5670  CB  PRO   122     -20.492  31.337 -72.925  1.00 68.31      H1   C
ATOM   5671  CG  PRO   122     -19.885  32.622 -73.381  1.00 68.44      H1   C
ATOM   5672  C   PRO   122     -20.173  29.701 -74.819  1.00 69.03      H1   C
ATOM   5673  O   PRO   122     -19.899  30.207 -75.906  1.00 69.06      H1   O
ATOM   5674  N   SER   123     -20.972  28.649 -74.688  1.00 69.50      H1   N
ATOM   5675  CA  SER   123     -21.822  28.187 -75.776  1.00 70.46      H1   C
ATOM   5676  CB  SER   123     -21.828  26.658 -75.823  1.00 70.16      H1   C
ATOM   5677  OG  SER   123     -22.886  26.177 -76.631  1.00 70.42      H1   O
ATOM   5678  C   SER   123     -23.238  28.708 -75.532  1.00 71.61      H1   C
ATOM   5679  O   SER   123     -23.796  28.518 -74.450  1.00 71.85      H1   O
ATOM   5680  N   VAL   124     -23.814  29.366 -76.536  1.00 71.93      H1   N
ATOM   5681  CA  VAL   124     -25.099  30.039 -76.368  1.00 71.46      H1   C
ATOM   5682  CB  VAL   124     -25.035  31.486 -76.898  1.00 70.79      H1   C
ATOM   5683  CG1 VAL   124     -26.356  32.191 -76.640  1.00 69.89      H1   C
ATOM   5684  CG2 VAL   124     -23.886  32.233 -76.236  1.00 68.40      H1   C
ATOM   5685  C   VAL   124     -26.244  29.313 -77.073  1.00 72.64      H1   C
ATOM   5686  O   VAL   124     -26.151  28.983 -78.255  1.00 73.01      H1   O
ATOM   5687  N   PHE   125     -27.324  29.071 -76.338  1.00 73.89      H1   N
ATOM   5688  CA  PHE   125     -28.510  28.431 -76.894  1.00 75.85      H1   C
ATOM   5689  CB  PHE   125     -28.723  27.062 -76.247  1.00 76.97      H1   C
ATOM   5690  CG  PHE   125     -27.641  26.071 -76.557  1.00 78.42      H1   C
ATOM   5691  CD1 PHE   125     -27.704  25.293 -77.702  1.00 78.83      H1   C
ATOM   5692  CD2 PHE   125     -26.560  25.916 -75.704  1.00 78.71      H1   C
ATOM   5693  CE1 PHE   125     -26.712  24.380 -77.992  1.00 79.49      H1   C
ATOM   5694  CE2 PHE   125     -25.564  25.004 -75.987  1.00 79.38      H1   C
ATOM   5695  CZ  PHE   125     -25.639  24.235 -77.134  1.00 80.12      H1   C
ATOM   5696  C   PHE   125     -29.749  29.292 -76.671  1.00 77.12      H1   C
ATOM   5697  O   PHE   125     -29.803  30.084 -75.731  1.00 76.96      H1   O
ATOM   5698  N   PRO   126     -30.760  29.151 -77.543  1.00 78.18      H1   N
ATOM   5699  CD  PRO   126     -30.705  28.371 -78.793  1.00 78.12      H1   C
ATOM   5700  CA  PRO   126     -32.048  29.838 -77.388  1.00 78.24      H1   C
ATOM   5701  CB  PRO   126     -32.641  29.793 -78.789  1.00 77.98      H1   C
ATOM   5702  CG  PRO   126     -32.099  28.522 -79.358  1.00 77.57      H1   C
ATOM   5703  C   PRO   126     -32.951  29.136 -76.376  1.00 78.94      H1   C
ATOM   5704  O   PRO   126     -33.020  27.909 -76.344  1.00 78.95      H1   O
ATOM   5705  N   LEU   127     -33.643  29.918 -75.555  1.00 80.12      H1   N
ATOM   5706  CA  LEU   127     -34.690  29.378 -74.695  1.00 81.26      H1   C
ATOM   5707  CB  LEU   127     -34.451  29.790 -73.239  1.00 80.50      H1   C
ATOM   5708  CG  LEU   127     -33.138  29.323 -72.604  1.00 79.91      H1   C
ATOM   5709  CD1 LEU   127     -33.050  29.821 -71.171  1.00 79.12      H1   C
ATOM   5710  CD2 LEU   127     -33.061  27.806 -72.647  1.00 79.76      H1   C
ATOM   5711  C   LEU   127     -36.039  29.905 -75.174  1.00 82.39      H1   C
ATOM   5712  O   LEU   127     -36.562  30.883 -74.639  1.00 82.90      H1   O
ATOM   5713  N   ALA   128     -36.595  29.247 -76.187  1.00 83.36      H1   N
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ATOM   5714  CA  ALA   128     -37.769  29.753 -76.891  1.00 84.32      H1   C
ATOM   5715  CB  ALA   128     -37.982  28.959 -78.174  1.00 84.16      H1   C
ATOM   5716  C   ALA   128     -39.031  29.711 -76.036  1.00 85.07      H1   C
ATOM   5717  O   ALA   128     -39.252  28.766 -75.275  1.00 85.35      H1   O
ATOM   5718  N   PRO   129     -39.884  30.740 -76.164  1.00 85.56      H1   N
ATOM   5719  CD  PRO   129     -39.691  31.851 -77.113  1.00 85.77      H1   C
ATOM   5720  CA  PRO   129     -41.121  30.890 -75.387  1.00 85.80      H1   C
ATOM   5721  CB  PRO   129     -41.604  32.293 -75.747  1.00 85.81      H1   C
ATOM   5722  CG  PRO   129     -41.019  32.555 -77.094  1.00 86.05      H1   C
ATOM   5723  C   PRO   129     -42.175  29.827 -75.697  1.00 86.17      H1   C
ATOM   5724  O   PRO   129     -42.318  29.391 -76.840  1.00 85.88      H1   O
ATOM   5725  N   SER   130     -42.911  29.421 -74.667  1.00 86.84      H1   N
ATOM   5726  CA  SER   130     -43.963  28.421 -74.816  1.00 87.43      H1   C
ATOM   5727  CB  SER   130     -44.385  27.890 -73.441  1.00 87.92      H1   C
ATOM   5728  OG  SER   130     -43.298  27.275 -72.771  1.00 88.97      H1   O
ATOM   5729  C   SER   130     -45.175  29.014 -75.531  1.00 87.44      H1   C
ATOM   5730  O   SER   130     -45.780  28.371 -76.391  1.00 87.06      H1   O
ATOM   5731  N   GLY   136     -50.193  34.409 -68.488  1.00 99.67      H1   N
ATOM   5732  CA  GLY   136     -51.016  34.123 -69.648  1.00 99.80      H1   C
ATOM   5733  C   GLY   136     -51.050  35.272 -70.638  1.00100.02      H1   C
ATOM   5734  O   GLY   136     -51.114  35.055 -71.850  1.00100.03      H1   O
ATOM   5735  N   GLY   137     -51.007  36.498 -70.123  1.00 99.85      H1   N
ATOM   5736  CA  GLY   137     -50.995  37.666 -70.986  1.00 99.13      H1   C
ATOM   5737  C   GLY   137     -49.608  37.961 -71.527  1.00 98.66      H1   C
ATOM   5738  O   GLY   137     -49.456  38.642 -72.542  1.00 98.56      H1   O
ATOM   5739  N   THR   138     -48.591  37.446 -70.842  1.00 98.05      H1   N
ATOM   5740  CA  THR   138     -47.209  37.602 -71.282  1.00 96.71      H1   C
ATOM   5741  CB  THR   138     -46.393  38.445 -70.280  1.00 96.65      H1   C
ATOM   5742  OG1 THR   138     -46.424  37.817 -68.991  1.00 96.53      H1   O
ATOM   5743  CG2 THR   138     -46.967  39.851 -70.175  1.00 95.98      H1   C
ATOM   5744  C   THR   138     -46.526  36.247 -71.444  1.00 95.56      H1   C
ATOM   5745  O   THR   138     -46.947  35.250 -70.853  1.00 95.36      H1   O
ATOM   5746  N   ALA   139     -45.472  36.220 -72.253  1.00 94.21      H1   N
ATOM   5747  CA  ALA   139     -44.699  35.022 -72.418  1.00 92.92      H1   C
ATOM   5748  CB  ALA   139     -44.780  34.497 -73.846  1.00 93.04      H1   C
ATOM   5749  C   ALA   139     -43.202  35.339 -72.095  1.00 91.71      H1   C
ATOM   5750  O   ALA   139     -42.713  36.430 -72.395  1.00 91.15      H1   O
ATOM   5751  N   ALA   140     -42.513  34.383 -71.480  1.00 90.37      H1   N
ATOM   5752  CA  ALA   140     -41.123  34.582 -71.092  1.00 88.70      H1   C
ATOM   5753  CB  ALA   140     -40.928  34.187 -69.634  1.00 89.09      H1   C
ATOM   5754  C   ALA   140     -40.178  33.783 -71.982  1.00 87.42      H1   C
ATOM   5755  O   ALA   140     -40.491  32.667 -72.400  1.00 86.82      H1   O
ATOM   5756  N   LEU   141     -39.020  34.368 -72.267  1.00 86.13      H1   N
ATOM   5757  CA  LEU   141     -38.002  33.722 -73.084  1.00 85.33      H1   C
ATOM   5758  CB  LEU   141     -38.303  33.941 -74.572  1.00 85.25      H1   C
ATOM   5759  CG  LEU   141     -38.499  35.376 -75.081  1.00 85.32      H1   C
ATOM   5760  CD1 LEU   141     -37.162  36.103 -75.165  1.00 84.61      H1   C
ATOM   5761  CD2 LEU   141     -39.154  35.334 -76.453  1.00 85.35      H1   C
ATOM   5762  C   LEU   141     -36.621  34.275 -72.741  1.00 85.01      H1   C
ATOM   5763  O   LEU   141     -36.496  35.396 -72.247  1.00 85.36      H1   O
ATOM   5764  N   GLY   142     -35.587  33.484 -73.001  1.00 83.90      H1   N
ATOM   5765  CA  GLY   142     -34.233  33.941 -72.752  1.00 83.19      H1   C
ATOM   5766  C   GLY   142     -33.231  33.175 -73.589  1.00 82.79      H1   C
ATOM   5767  O   GLY   142     -33.609  32.491 -74.539  1.00 83.09      H1   O
ATOM   5768  N   CYS   143     -31.952  33.289 -73.248  1.00 82.17      H1   N
ATOM   5769  CA  CYS   143     -30.940  32.465 -73.890  1.00 81.85      H1   C
ATOM   5770  C   CYS   143     -29.954  31.864 -72.882  1.00 80.13      H1   C
ATOM   5771  O   CYS   143     -29.537  32.518 -71.925  1.00 79.94      H1   O
ATOM   5772  CB  CYS   143     -30.204  33.270 -74.974  1.00 83.41      H1   C
ATOM   5773  SG  CYS   143     -28.939  34.449 -74.400  1.00 87.73      H1   S
ATOM   5774  N   LEU   144     -29.601  30.602 -73.109  1.00 78.24      H1   N
ATOM   5775  CA  LEU   144     -28.826  29.808 -72.161  1.00 75.92      H1   C
ATOM   5776  CB  LEU   144     -29.286  28.349 -72.233  1.00 75.88      H1   C
ATOM   5777  CG  LEU   144     -28.506  27.285 -71.462  1.00 76.15      H1   C
ATOM   5778  CD1 LEU   144     -28.543  27.594 -69.977  1.00 76.40      H1   C
ATOM   5779  CD2 LEU   144     -29.115  25.915 -71.739  1.00 75.59      H1   C
ATOM   5780  C   LEU   144     -27.329  29.895 -72.449  1.00 74.33      H1   C
ATOM   5781  O   LEU   144     -26.890  29.633 -73.566  1.00 74.67      H1   O
ATOM   5782  N   VAL   145     -26.549  30.260 -71.436  1.00 72.28      H1   N
ATOM   5783  CA  VAL   145     -25.103  30.401 -71.591  1.00 70.90      H1   C
ATOM   5784  CB  VAL   145     -24.623  31.746 -71.003  1.00 69.27      H1   C
ATOM   5785  CG1 VAL   145     -23.127  31.873 -71.147  1.00 69.00      H1   C
ATOM   5786  CG2 VAL   145     -25.319  32.894 -71.705  1.00 68.74      H1   C
ATOM   5787  C   VAL   145     -24.358  29.254 -70.899  1.00 71.16      H1   C
ATOM   5788  O   VAL   145     -24.173  29.269 -69.680  1.00 71.43      H1   O
ATOM   5789  N   LYS   146     -23.927  28.266 -71.684  1.00 70.62      H1   N
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ATOM   5790  CA  LYS   146     -23.358  27.031 -71.141  1.00 70.17      H1   C
ATOM   5791  CB  LYS   146     -23.870  25.819 -71.925  1.00 71.40      H1   C
ATOM   5792  CG  LYS   146     -25.286  25.393 -71.583  1.00 74.93      H1   C
ATOM   5793  CD  LYS   146     -25.822  24.395 -72.605  1.00 78.01      H1   C
ATOM   5794  CE  LYS   146     -24.948  23.146 -72.701  1.00 79.46      H1   C
ATOM   5795  NZ  LYS   146     -25.126  22.233 -71.534  1.00 80.50      H1   N
ATOM   5796  C   LYS   146     -21.831  26.988 -71.127  1.00 68.82      H1   C
ATOM   5797  O   LYS   146     -21.173  27.538 -72.011  1.00 68.87      H1   O
ATOM   5798  N   ASP   147     -21.286  26.330 -70.106  1.00 66.87      H1   N
ATOM   5799  CA  ASP   147     -19.906  25.850 -70.110  1.00 64.95      H1   C
ATOM   5800  CB  ASP   147     -19.759  24.720 -71.129  1.00 64.08      H1   C
ATOM   5801  CG  ASP   147     -20.730  23.582 -70.877  1.00 65.39      H1   C
ATOM   5802  OD1 ASP   147     -20.976  23.252 -69.697  1.00 64.41      H1   O
ATOM   5803  OD2 ASP   147     -21.250  23.015 -71.862  1.00 66.30      H1   O
ATOM   5804  C   ASP   147     -18.857  26.920 -70.390  1.00 64.29      H1   C
ATOM   5805  O   ASP   147     -18.121  26.829 -71.368  1.00 64.65      H1   O
ATOM   5806  N   TYR   148     -18.775  27.925 -69.527  1.00 63.41      H1   N
ATOM   5807  CA  TYR   148     -17.760  28.956 -69.687  1.00 63.64      H1   C
ATOM   5808  CB  TYR   148     -18.411  30.291 -70.061  1.00 63.45      H1   C
ATOM   5809  CG  TYR   148     -19.292  30.886 -68.984  1.00 62.08      H1   C
ATOM   5810  CD1 TYR   148     -18.787  31.813 -68.081  1.00 61.94      H1   C
ATOM   5811  CE1 TYR   148     -19.594  32.383 -67.112  1.00 61.63      H1   C
ATOM   5812  CD2 TYR   148     -20.632  30.539 -68.887  1.00 62.19      H1   C
ATOM   5813  CE2 TYR   148     -21.448  31.103 -67.920  1.00 62.61      H1   C
ATOM   5814  CZ  TYR   148     -20.923  32.024 -67.037  1.00 61.07      H1   C
ATOM   5815  OH  TYR   148     -21.731  32.594 -66.082  1.00 61.02      H1   O
ATOM   5816  C   TYR   148     -16.920  29.126 -68.425  1.00 64.03      H1   C
ATOM   5817  O   TYR   148     -17.296  28.664 -67.348  1.00 63.78      H1   O
ATOM   5818  N   PHE   149     -15.777  29.788 -68.572  1.00 64.42      H1   N
ATOM   5819  CA  PHE   149     -14.897  30.073 -67.446  1.00 65.06      H1   C
ATOM   5820  CB  PHE   149     -14.209  28.787 -66.972  1.00 63.77      H1   C
ATOM   5821  CG  PHE   149     -13.335  28.979 -65.765  1.00 62.58      H1   C
ATOM   5822  CD1 PHE   149     -12.012  29.371 -65.904  1.00 61.56      H1   C
ATOM   5823  CD2 PHE   149     -13.843  28.789 -64.489  1.00 62.08      H1   C
ATOM   5824  CE1 PHE   149     -11.210  29.576 -64.795  1.00 60.99      H1   C
ATOM   5825  CE2 PHE   149     -13.047  28.992 -63.373  1.00 62.34      H1   C
ATOM   5826  CZ  PHE   149     -11.727  29.386 -63.527  1.00 61.88      H1   C
ATOM   5827  C   PHE   149     -13.844  31.092 -67.865  1.00 66.30      H1   C
ATOM   5828  O   PHE   149     -13.365  31.069 -68.998  1.00 67.25      H1   O
ATOM   5829  N   PRO   150     -13.473  32.006 -66.955  1.00 67.74      H1   N
ATOM   5830  CD  PRO   150     -12.274  32.850 -67.107  1.00 67.53      H1   C
ATOM   5831  CA  PRO   150     -14.104  32.195 -65.645  1.00 68.63      H1   C
ATOM   5832  CB  PRO   150     -13.037  32.928 -64.841  1.00 67.62      H1   C
ATOM   5833  CG  PRO   150     -12.295  33.712 -65.866  1.00 67.46      H1   C
ATOM   5834  C   PRO   150     -15.389  33.010 -65.742  1.00 70.45      H1   C
ATOM   5835  O   PRO   150     -16.077  32.997 -66.764  1.00 71.47      H1   O
ATOM   5836  N   GLU   151     -15.700  33.719 -64.663  1.00 71.80      H1   N
ATOM   5837  CA  GLU   151     -16.743  34.734 -64.676  1.00 72.52      H1   C
ATOM   5838  CB  GLU   151     -17.550  34.681 -63.378  1.00 72.76      H1   C
ATOM   5839  CG  GLU   151     -18.290  33.376 -63.155  1.00 74.07      H1   C
ATOM   5840  CD  GLU   151     -19.458  33.535 -62.204  1.00 74.56      H1   C
ATOM   5841  OE1 GLU   151     -20.618  33.454 -62.665  1.00 74.92      H1   O
ATOM   5842  OE2 GLU   151     -19.216  33.745 -60.997  1.00 73.75      H1   O
ATOM   5843  C   GLU   151     -16.079  36.099 -64.806  1.00 73.24      H1   C
ATOM   5844  O   GLU   151     -14.868  36.227 -64.624  1.00 72.62      H1   O
ATOM   5845  N   PRO   152     -16.867  37.142 -65.113  1.00 74.62      H1   N
ATOM   5846  CD  PRO   152     -16.420  38.539 -64.959  1.00 74.59      H1   C
ATOM   5847  CA  PRO   152     -18.292  37.070 -65.452  1.00 75.10      H1   C
ATOM   5848  CB  PRO   152     -18.849  38.361 -64.867  1.00 75.76      H1   C
ATOM   5849  CG  PRO   152     -17.712  39.337 -65.029  1.00 75.41      H1   C
ATOM   5850  C   PRO   152     -18.556  36.968 -66.952  1.00 75.76      H1   C
ATOM   5851  O   PRO   152     -17.630  36.989 -67.761  1.00 75.62      H1   O
ATOM   5852  N   VAL   153     -19.830  36.854 -67.312  1.00 76.74      H1   N
ATOM   5853  CA  VAL   153     -20.266  37.125 -68.676  1.00 78.44      H1   C
ATOM   5854  CB  VAL   153     -20.722  35.836 -69.410  1.00 78.41      H1   C
ATOM   5855  CG1 VAL   153     -19.573  34.848 -69.489  1.00 78.22      H1   C
ATOM   5856  CG2 VAL   153     -21.909  35.219 -68.700  1.00 78.16      H1   C
ATOM   5857  C   VAL   153     -21.428  38.114 -68.651  1.00 79.27      H1   C
ATOM   5858  O   VAL   153     -22.353  37.981 -67.848  1.00 79.85      H1   O
ATOM   5859  N   THR   154     -21.367  39.111 -69.528  1.00 79.67      H1   N
ATOM   5860  CA  THR   154     -22.408  40.130 -69.613  1.00 79.68      H1   C
ATOM   5861  CB  THR   154     -21.808  41.505 -69.963  1.00 79.51      H1   C
ATOM   5862  OG1 THR   154     -21.139  41.422 -71.228  1.00 79.48      H1   O
ATOM   5863  CG2 THR   154     -20.811  41.945 -68.897  1.00 78.20      H1   C
ATOM   5864  C   THR   154     -23.417  39.754 -70.693  1.00 80.21      H1   C
ATOM   5865  O   THR   154     -23.034  39.413 -71.812  1.00 80.43      H1   O
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ATOM   5866  N   VAL  155     -24.703  39.813 -70.359  1.00 80.71      H1   N
ATOM   5867  CA  VAL  155     -25.752  39.518 -71.330  1.00 81.00      H1   C
ATOM   5868  CB  VAL  155     -26.580  38.284 -70.910  1.00 80.93      H1   C
ATOM   5869  CG1 VAL  155     -27.614  37.965 -71.979  1.00 80.65      H1   C
ATOM   5870  CG2 VAL  155     -25.666  37.094 -70.689  1.00 81.15      H1   C
ATOM   5871  C   VAL  155     -26.701  40.702 -71.503  1.00 81.78      H1   C
ATOM   5872  O   VAL  155     -27.351  41.139 -70.551  1.00 81.71      H1   O
ATOM   5873  N   SER  156     -26.772  41.216 -72.727  1.00 82.68      H1   N
ATOM   5874  CA  SER  156     -27.654  42.334 -73.046  1.00 83.16      H1   C
ATOM   5875  CB  SER  156     -26.847  43.473 -73.674  1.00 83.30      H1   C
ATOM   5876  OG  SER  156     -27.652  44.620 -73.878  1.00 84.63      H1   O
ATOM   5877  C   SER  156     -28.739  41.868 -74.014  1.00 83.16      H1   C
ATOM   5878  O   SER  156     -28.596  40.831 -74.658  1.00 82.89      H1   O
ATOM   5879  N   TRP  157     -29.825  42.627 -74.110  1.00 83.58      H1   N
ATOM   5880  CA  TRP  157     -30.919  42.259 -75.001  1.00 84.15      H1   C
ATOM   5881  CB  TRP  157     -32.180  41.952 -74.188  1.00 83.42      H1   C
ATOM   5882  CG  TRP  157     -32.120  40.627 -73.494  1.00 83.40      H1   C
ATOM   5883  CD2 TRP  157     -32.489  39.355 -74.042  1.00 83.53      H1   C
ATOM   5884  CE2 TRP  157     -32.248  38.386 -73.047  1.00 83.19      H1   C
ATOM   5885  CE3 TRP  157     -33.000  38.941 -75.277  1.00 83.42      H1   C
ATOM   5886  CD1 TRP  157     -31.684  40.385 -72.224  1.00 83.06      H1   C
ATOM   5887  NE1 TRP  157     -31.757  39.040 -71.947  1.00 82.93      H1   N
ATOM   5888  CZ2 TRP  157     -32.498  37.030 -73.250  1.00 83.39      H1   C
ATOM   5889  CZ3 TRP  157     -33.247  37.594 -75.476  1.00 83.62      H1   C
ATOM   5890  CH2 TRP  157     -32.997  36.655 -74.468  1.00 83.34      H1   C
ATOM   5891  C   TRP  157     -31.219  43.334 -76.041  1.00 84.76      H1   C
ATOM   5892  O   TRP  157     -31.410  44.505 -75.706  1.00 84.03      H1   O
ATOM   5893  N   ASN  158     -31.260  42.921 -77.306  1.00 85.58      H1   N
ATOM   5894  CA  ASN  158     -31.495  43.842 -78.411  1.00 86.37      H1   C
ATOM   5895  CB  ASN  158     -32.946  44.331 -78.389  1.00 86.40      H1   C
ATOM   5896  CG  ASN  158     -33.946  43.196 -78.553  1.00 87.21      H1   C
ATOM   5897  OD1 ASN  158     -33.691  42.224 -79.268  1.00 86.86      H1   O
ATOM   5898  ND2 ASN  158     -35.091  43.315 -77.889  1.00 86.97      H1   N
ATOM   5899  C   ASN  158     -30.536  45.026 -78.315  1.00 86.98      H1   C
ATOM   5900  O   ASN  158     -30.939  46.183 -78.439  1.00 86.86      H1   O
ATOM   5901  N   SER  159     -29.264  44.717 -78.077  1.00 87.36      H1   N
ATOM   5902  CA  SER  159     -28.203  45.716 -78.013  1.00 87.62      H1   C
ATOM   5903  CB  SER  159     -27.970  46.322 -79.398  1.00 87.30      H1   C
ATOM   5904  OG  SER  159     -27.479  45.341 -80.297  1.00 86.83      H1   O
ATOM   5905  C   SER  159     -28.471  46.822 -76.999  1.00 87.96      H1   C
ATOM   5906  O   SER  159     -27.791  47.849 -76.998  1.00 87.94      H1   O
ATOM   5907  N   GLY  160     -29.459  46.608 -76.136  1.00 88.50      H1   N
ATOM   5908  CA  GLY  160     -29.716  47.551 -75.062  1.00 88.84      H1   C
ATOM   5909  C   GLY  160     -31.108  48.150 -75.103  1.00 88.98      H1   C
ATOM   5910  O   GLY  160     -31.570  48.731 -74.120  1.00 88.61      H1   O
ATOM   5911  N   ALA  161     -31.779  48.006 -76.242  1.00 89.24      H1   N
ATOM   5912  CA  ALA  161     -33.105  48.585 -76.430  1.00 89.63      H1   C
ATOM   5913  CB  ALA  161     -33.569  48.373 -77.868  1.00 89.23      H1   C
ATOM   5914  C   ALA  161     -34.112  47.971 -75.461  1.00 89.76      H1   C
ATOM   5915  O   ALA  161     -35.083  48.619 -75.067  1.00 89.81      H1   O
ATOM   5916  N   LEU  162     -33.874  46.718 -75.084  1.00 89.21      H1   N
ATOM   5917  CA  LEU  162     -34.769  46.006 -74.182  1.00 88.42      H1   C
ATOM   5918  CB  LEU  162     -35.189  44.673 -74.806  1.00 87.77      H1   C
ATOM   5919  CG  LEU  162     -36.018  43.738 -73.923  1.00 87.40      H1   C
ATOM   5920  CD1 LEU  162     -37.267  44.456 -73.439  1.00 86.87      H1   C
ATOM   5921  CD2 LEU  162     -36.384  42.491 -74.710  1.00 86.79      H1   C
ATOM   5922  C   LEU  162     -34.106  45.753 -72.833  1.00 88.21      H1   C
ATOM   5923  O   LEU  162     -33.176  44.954 -72.727  1.00 88.43      H1   O
ATOM   5924  N   THR  163     -34.591  46.441 -71.804  1.00 87.73      H1   N
ATOM   5925  CA  THR  163     -34.069  46.270 -70.453  1.00 87.13      H1   C
ATOM   5926  CB  THR  163     -33.414  47.569 -69.937  1.00 87.76      H1   C
ATOM   5927  OG1 THR  163     -34.374  48.634 -69.962  1.00 88.69      H1   O
ATOM   5928  CG2 THR  163     -32.218  47.946 -70.804  1.00 87.58      H1   C
ATOM   5929  C   THR  163     -35.194  45.880 -69.503  1.00 86.16      H1   C
ATOM   5930  O   THR  163     -34.957  45.286 -68.449  1.00 85.76      H1   O
ATOM   5931  N   SER  164     -36.420  46.216 -69.888  1.00 84.83      H1   N
ATOM   5932  CA  SER  164     -37.584  45.961 -69.052  1.00 83.71      H1   C
ATOM   5933  CB  SER  164     -38.737  46.877 -69.475  1.00 84.14      H1   C
ATOM   5934  OG  SER  164     -39.855  46.731 -68.616  1.00 84.04      H1   O
ATOM   5935  C   SER  164     -38.016  44.500 -69.149  1.00 82.36      H1   C
ATOM   5936  O   SER  164     -38.185  43.962 -70.244  1.00 82.47      H1   O
ATOM   5937  N   GLY  165     -38.193  43.864 -67.995  1.00 80.97      H1   N
ATOM   5938  CA  GLY  165     -38.616  42.476 -67.972  1.00 79.30      H1   C
ATOM   5939  C   GLY  165     -37.452  41.504 -68.016  1.00 78.12      H1   C
ATOM   5940  O   GLY  165     -37.651  40.294 -68.120  1.00 77.82      H1   O
ATOM   5941  N   VAL  166     -36.233  42.030 -67.935  1.00 76.72      H1   N
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ATOM	5942	CA	VAL	166	-35.037	41.205	-68.040	1.00	75.81	H1	C
ATOM	5943	CB	VAL	166	-33.905	41.954	-68.770	1.00	75.90	H1	C
ATOM	5944	CG1	VAL	166	-32.673	41.070	-68.856	1.00	75.62	H1	C
ATOM	5945	CG2	VAL	166	-34.365	42.364	-70.161	1.00	75.64	H1	C
ATOM	5946	C	VAL	166	-34.526	40.775	-66.670	1.00	75.14	H1	C
ATOM	5947	O	VAL	166	-34.253	41.610	-65.809	1.00	75.37	H1	O
ATOM	5948	N	HIS	167	-34.396	39.466	-66.480	1.00	73.47	H1	N
ATOM	5949	CA	HIS	167	-33.935	38.911	-65.214	1.00	72.00	H1	C
ATOM	5950	CB	HIS	167	-35.057	38.090	-64.569	1.00	72.27	H1	C
ATOM	5951	CG	HIS	167	-34.777	37.683	-63.155	1.00	72.47	H1	C
ATOM	5952	CD2	HIS	167	-33.696	37.887	-62.363	1.00	72.15	H1	C
ATOM	5953	ND1	HIS	167	-35.684	36.981	-62.390	1.00	71.80	H1	N
ATOM	5954	CE1	HIS	167	-35.176	36.771	-61.189	1.00	71.72	H1	C
ATOM	5955	NE2	HIS	167	-33.971	37.311	-61.147	1.00	72.50	H1	N
ATOM	5956	C	HIS	167	-32.708	38.028	-65.438	1.00	70.88	H1	C
ATOM	5957	O	HIS	167	-32.783	37.008	-66.121	1.00	69.91	H1	O
ATOM	5958	N	THR	168	-31.581	38.424	-64.855	1.00	70.00	H1	N
ATOM	5959	CA	THR	168	-30.335	37.687	-65.027	1.00	68.86	H1	C
ATOM	5960	CB	THR	168	-29.156	38.650	-65.269	1.00	69.61	H1	C
ATOM	5961	OG1	THR	168	-29.449	39.489	-66.394	1.00	70.62	H1	O
ATOM	5962	CG2	THR	168	-27.878	37.867	-65.548	1.00	69.75	H1	C
ATOM	5963	C	THR	168	-30.026	36.826	-63.803	1.00	67.17	H1	C
ATOM	5964	O	THR	168	-29.731	37.344	-62.727	1.00	67.02	H1	O
ATOM	5965	N	PHE	169	-30.093	35.510	-63.979	1.00	64.86	H1	N
ATOM	5966	CA	PHE	169	-29.878	34.576	-62.881	1.00	62.90	H1	C
ATOM	5967	CB	PHE	169	-30.615	33.263	-63.150	1.00	62.05	H1	C
ATOM	5968	CG	PHE	169	-32.108	33.400	-63.176	1.00	62.68	H1	C
ATOM	5969	CD1	PHE	169	-32.748	33.972	-64.266	1.00	62.88	H1	C
ATOM	5970	CD2	PHE	169	-32.877	32.943	-62.118	1.00	62.49	H1	C
ATOM	5971	CE1	PHE	169	-34.128	34.086	-64.299	1.00	62.61	H1	C
ATOM	5972	CE2	PHE	169	-34.257	33.053	-62.144	1.00	62.55	H1	C
ATOM	5973	CZ	PHE	169	-34.884	33.625	-63.237	1.00	62.31	H1	C
ATOM	5974	C	PHE	169	-28.398	34.290	-62.683	1.00	61.96	H1	C
ATOM	5975	O	PHE	169	-27.663	34.061	-63.645	1.00	61.82	H1	O
ATOM	5976	N	PRO	170	-27.942	34.294	-61.421	1.00	60.72	H1	N
ATOM	5977	CD	PRO	170	-28.702	34.746	-60.244	1.00	60.03	H1	C
ATOM	5978	CA	PRO	170	-26.555	33.964	-61.078	1.00	59.89	H1	C
ATOM	5979	CB	PRO	170	-26.526	34.083	-59.554	1.00	59.29	H1	C
ATOM	5980	CG	PRO	170	-27.618	35.058	-59.246	1.00	59.11	H1	C
ATOM	5981	C	PRO	170	-26.132	32.576	-61.561	1.00	59.02	H1	C
ATOM	5982	O	PRO	170	-26.948	31.653	-61.647	1.00	57.43	H1	O
ATOM	5983	N	ALA	171	-24.847	32.440	-61.873	1.00	58.34	H1	N
ATOM	5984	CA	ALA	171	-24.329	31.229	-62.496	1.00	58.50	H1	C
ATOM	5985	CB	ALA	171	-22.955	31.503	-63.094	1.00	57.77	H1	C
ATOM	5986	C	ALA	171	-24.243	30.066	-61.517	1.00	58.57	H1	C
ATOM	5987	O	ALA	171	-24.030	30.259	-60.320	1.00	57.67	H1	O
ATOM	5988	N	VAL	172	-24.415	28.855	-62.035	1.00	58.56	H1	N
ATOM	5989	CA	VAL	172	-24.118	27.654	-61.268	1.00	59.16	H1	C
ATOM	5990	CB	VAL	172	-25.148	26.538	-61.553	1.00	58.86	H1	C
ATOM	5991	CG1	VAL	172	-25.070	26.114	-63.011	1.00	59.09	H1	C
ATOM	5992	CG2	VAL	172	-24.898	25.353	-60.632	1.00	60.06	H1	C
ATOM	5993	C	VAL	172	-22.721	27.157	-61.641	1.00	59.63	H1	C
ATOM	5994	O	VAL	172	-22.304	27.266	-62.796	1.00	59.35	H1	O
ATOM	5995	N	LEU	173	-21.990	26.631	-60.663	1.00	59.79	H1	N
ATOM	5996	CA	LEU	173	-20.688	26.032	-60.937	1.00	60.87	H1	C
ATOM	5997	CB	LEU	173	-19.698	26.348	-59.811	1.00	59.98	H1	C
ATOM	5998	CG	LEU	173	-18.205	26.413	-60.167	1.00	60.10	H1	C
ATOM	5999	CD1	LEU	173	-17.376	26.202	-58.905	1.00	57.51	H1	C
ATOM	6000	CD2	LEU	173	-17.858	25.356	-61.201	1.00	58.71	H1	C
ATOM	6001	C	LEU	173	-20.879	24.528	-61.041	1.00	61.60	H1	C
ATOM	6002	O	LEU	173	-21.379	23.897	-60.110	1.00	62.78	H1	O
ATOM	6003	N	GLN	174	-20.482	23.956	-62.172	1.00	62.51	H1	N
ATOM	6004	CA	GLN	174	-20.744	22.547	-62.446	1.00	63.92	H1	C
ATOM	6005	CB	GLN	174	-21.044	22.352	-63.929	1.00	65.18	H1	C
ATOM	6006	CG	GLN	174	-22.139	23.256	-64.450	1.00	67.24	H1	C
ATOM	6007	CD	GLN	174	-22.385	23.065	-65.928	1.00	69.04	H1	C
ATOM	6008	OE1	GLN	174	-23.070	22.125	-66.337	1.00	70.60	H1	O
ATOM	6009	NE2	GLN	174	-21.823	23.953	-66.741	1.00	68.01	H1	N
ATOM	6010	C	GLN	174	-19.565	21.672	-62.050	1.00	63.96	H1	C
ATOM	6011	O	GLN	174	-18.445	22.159	-61.895	1.00	64.54	H1	O
ATOM	6012	N	SER	175	-19.820	20.376	-61.896	1.00	63.64	H1	N
ATOM	6013	CA	SER	175	-18.777	19.438	-61.501	1.00	63.62	H1	C
ATOM	6014	CB	SER	175	-19.346	18.016	-61.407	1.00	64.57	H1	C
ATOM	6015	OG	SER	175	-19.975	17.626	-62.616	1.00	66.59	H1	O
ATOM	6016	C	SER	175	-17.610	19.464	-62.481	1.00	63.00	H1	C
ATOM	6017	O	SER	175	-16.541	18.926	-62.197	1.00	64.06	H1	O

ATOM	6018	N	SER	176	-17.816	20.095	-63.632	1.00	61.79		H1	N
ATOM	6019	CA	SER	176	-16.776	20.187	-64.650	1.00	60.10		H1	C
ATOM	6020	CB	SER	176	-17.399	20.418	-66.025	1.00	59.94		H1	C
ATOM	6021	OG	SER	176	-17.863	21.752	-66.150	1.00	60.80		H1	O
ATOM	6022	C	SER	176	-15.806	21.323	-64.355	1.00	59.39		H1	C
ATOM	6023	O	SER	176	-14.733	21.398	-64.951	1.00	59.39		H1	O
ATOM	6024	N	GLY	177	-16.191	22.211	-63.443	1.00	57.70		H1	N
ATOM	6025	CA	GLY	177	-15.384	23.389	-63.176	1.00	56.90		H1	C
ATOM	6026	C	GLY	177	-15.778	24.550	-64.068	1.00	56.87		H1	C
ATOM	6027	O	GLY	177	-15.157	25.614	-64.043	1.00	56.89		H1	O
ATOM	6028	N	LEU	178	-16.822	24.345	-64.860	1.00	56.61		H1	N
ATOM	6029	CA	LEU	178	-17.298	25.370	-65.778	1.00	57.28		H1	C
ATOM	6030	CB	LEU	178	-17.436	24.786	-67.188	1.00	57.13		H1	C
ATOM	6031	CG	LEU	178	-16.115	24.423	-67.872	1.00	56.25		H1	C
ATOM	6032	CD1	LEU	178	-16.387	23.798	-69.230	1.00	55.52		H1	C
ATOM	6033	CD2	LEU	178	-15.268	25.675	-68.017	1.00	56.22		H1	C
ATOM	6034	C	LEU	178	-18.631	25.959	-65.328	1.00	57.07		H1	C
ATOM	6035	O	LEU	178	-19.428	25.295	-64.665	1.00	55.79		H1	O
ATOM	6036	N	TYR	179	-18.864	27.213	-65.697	1.00	57.71		H1	N
ATOM	6037	CA	TYR	179	-20.064	27.921	-65.280	1.00	59.38		H1	C
ATOM	6038	CB	TYR	179	-19.725	29.378	-64.961	1.00	58.68		H1	C
ATOM	6039	CG	TYR	179	-18.831	29.558	-63.754	1.00	58.53		H1	C
ATOM	6040	CD1	TYR	179	-19.346	29.474	-62.465	1.00	58.26		H1	C
ATOM	6041	CE1	TYR	179	-18.540	29.680	-61.356	1.00	58.61		H1	C
ATOM	6042	CD2	TYR	179	-17.482	29.849	-63.904	1.00	57.93		H1	C
ATOM	6043	CE2	TYR	179	-16.667	30.058	-62.803	1.00	58.41		H1	C
ATOM	6044	CZ	TYR	179	-17.200	29.974	-61.531	1.00	58.60		H1	C
ATOM	6045	OH	TYR	179	-16.394	30.195	-60.438	1.00	56.00		H1	O
ATOM	6046	C	TYR	179	-21.174	27.877	-66.328	1.00	60.05		H1	C
ATOM	6047	O	TYR	179	-20.914	27.891	-67.531	1.00	59.38		H1	O
ATOM	6048	N	SER	180	-22.413	27.819	-65.852	1.00	61.78		H1	N
ATOM	6049	CA	SER	180	-23.589	27.990	-66.698	1.00	64.17		H1	C
ATOM	6050	CB	SER	180	-24.290	26.646	-66.917	1.00	63.61		H1	C
ATOM	6051	OG	SER	180	-23.501	25.775	-67.708	1.00	65.69		H1	O
ATOM	6052	C	SER	180	-24.554	28.960	-66.024	1.00	65.91		H1	C
ATOM	6053	O	SER	180	-24.671	28.979	-64.797	1.00	65.96		H1	O
ATOM	6054	N	HIS	181	-25.242	29.768	-66.823	1.00	67.43		H1	N
ATOM	6055	CA	HIS	181	-26.293	30.625	-66.290	1.00	69.34		H1	C
ATOM	6056	CB	HIS	181	-25.689	31.850	-65.606	1.00	70.05		H1	C
ATOM	6057	CG	HIS	181	-25.466	33.009	-66.524	1.00	70.38		H1	C
ATOM	6058	CD2	HIS	181	-24.413	33.336	-67.309	1.00	70.89		H1	C
ATOM	6059	ND1	HIS	181	-26.387	34.023	-66.676	1.00	70.55		H1	N
ATOM	6060	CE1	HIS	181	-25.909	34.927	-67.513	1.00	71.32		H1	C
ATOM	6061	NE2	HIS	181	-24.713	34.533	-67.911	1.00	71.85		H1	N
ATOM	6062	C	HIS	181	-27.264	31.075	-67.371	1.00	70.14		H1	C
ATOM	6063	O	HIS	181	-26.990	30.942	-68.564	1.00	69.87		H1	O
ATOM	6064	N	SER	182	-28.402	31.609	-66.941	1.00	71.10		H1	N
ATOM	6065	CA	SER	182	-29.458	32.010	-67.859	1.00	70.94		H1	C
ATOM	6066	CB	SER	182	-30.702	31.146	-67.636	1.00	70.38		H1	C
ATOM	6067	OG	SER	182	-30.418	29.774	-67.830	1.00	68.38		H1	O
ATOM	6068	C	SER	182	-29.827	33.479	-67.688	1.00	71.53		H1	C
ATOM	6069	O	SER	182	-29.763	34.023	-66.584	1.00	71.22		H1	O
ATOM	6070	N	SER	183	-30.205	34.114	-68.794	1.00	72.30		H1	N
ATOM	6071	CA	SER	183	-30.900	35.396	-68.757	1.00	72.41		H1	C
ATOM	6072	CB	SER	183	-30.111	36.456	-69.525	1.00	71.36		H1	C
ATOM	6073	OG	SER	183	-30.820	37.682	-69.569	1.00	69.72		H1	O
ATOM	6074	C	SER	183	-32.276	35.227	-69.388	1.00	73.63		H1	C
ATOM	6075	O	SER	183	-32.414	34.573	-70.421	1.00	73.54		H1	O
ATOM	6076	N	VAL	184	-33.294	35.809	-68.762	1.00	75.17		H1	N
ATOM	6077	CA	VAL	184	-34.658	35.702	-69.269	1.00	76.46		H1	C
ATOM	6078	CB	VAL	184	-35.491	34.712	-68.424	1.00	76.29		H1	C
ATOM	6079	CG1	VAL	184	-36.949	34.751	-68.852	1.00	76.19		H1	C
ATOM	6080	CG2	VAL	184	-34.941	33.311	-68.594	1.00	76.69		H1	C
ATOM	6081	C	VAL	184	-35.378	37.045	-69.300	1.00	77.45		H1	C
ATOM	6082	O	VAL	184	-35.375	37.793	-68.320	1.00	77.46		H1	O
ATOM	6083	N	VAL	185	-35.992	37.345	-70.439	1.00	78.31		H1	N
ATOM	6084	CA	VAL	185	-36.816	38.538	-70.572	1.00	78.79		H1	C
ATOM	6085	CB	VAL	185	-36.362	39.400	-71.778	1.00	78.96		H1	C
ATOM	6086	CG1	VAL	185	-36.360	38.566	-73.049	1.00	78.89		H1	C
ATOM	6087	CG2	VAL	185	-37.281	40.601	-71.931	1.00	79.45		H1	C
ATOM	6088	C	VAL	185	-38.281	38.151	-70.743	1.00	78.60		H1	C
ATOM	6089	O	VAL	185	-38.632	37.366	-71.624	1.00	77.83		H1	O
ATOM	6090	N	THR	186	-39.132	38.696	-69.880	1.00	79.19		H1	N
ATOM	6091	CA	THR	186	-40.571	38.499	-69.998	1.00	80.16		H1	C
ATOM	6092	CB	THR	186	-41.244	38.482	-68.609	1.00	80.16		H1	C
ATOM	6093	OG1	THR	186	-40.924	39.689	-67.907	1.00	80.54		H1	O

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ATOM   6094  CG2 THR   186     -40.761  37.290 -67.796  1.00 80.44      H1   C
ATOM   6095  C   THR   186     -41.172  39.627 -70.833  1.00 80.74      H1   C
ATOM   6096  O   THR   186     -41.107  40.798 -70.455  1.00 79.91      H1   O
ATOM   6097  N   VAL   187     -41.750  39.268 -71.975  1.00 82.12      H1   N
ATOM   6098  CA  VAL   187     -42.248  40.258 -72.923  1.00 83.45      H1   C
ATOM   6099  CB  VAL   187     -41.451  40.210 -74.246  1.00 83.13      H1   C
ATOM   6100  CG1 VAL   187     -39.975  40.456 -73.975  1.00 82.39      H1   C
ATOM   6101  CG2 VAL   187     -41.662  38.869 -74.933  1.00 82.96      H1   C
ATOM   6102  C   VAL   187     -43.726  40.048 -73.239  1.00 84.60      H1   C
ATOM   6103  O   VAL   187     -44.303  39.012 -72.906  1.00 83.97      H1   O
ATOM   6104  N   PRO   188     -44.357  41.041 -73.886  1.00 86.10      H1   N
ATOM   6105  CD  PRO   188     -43.802  42.380 -74.148  1.00 86.23      H1   C
ATOM   6106  CA  PRO   188     -45.756  40.939 -74.316  1.00 87.32      H1   C
ATOM   6107  CB  PRO   188     -46.050  42.313 -74.919  1.00 87.01      H1   C
ATOM   6108  CG  PRO   188     -45.028  43.220 -74.315  1.00 86.99      H1   C
ATOM   6109  C   PRO   188     -45.950  39.825 -75.339  1.00 88.92      H1   C
ATOM   6110  O   PRO   188     -45.245  39.772 -76.347  1.00 89.36      H1   O
ATOM   6111  N   SER   189     -46.908  38.940 -75.080  1.00 90.12      H1   N
ATOM   6112  CA  SER   189     -47.239  37.889 -76.033  1.00 91.74      H1   C
ATOM   6113  CB  SER   189     -48.429  37.067 -75.529  1.00 91.44      H1   C
ATOM   6114  OG  SER   189     -48.108  36.379 -74.333  1.00 91.54      H1   O
ATOM   6115  C   SER   189     -47.584  38.513 -77.382  1.00 93.06      H1   C
ATOM   6116  O   SER   189     -47.261  37.962 -78.436  1.00 93.63      H1   O
ATOM   6117  N   SER   190     -48.233  39.673 -77.334  1.00 94.02      H1   N
ATOM   6118  CA  SER   190     -48.654  40.386 -78.536  1.00 94.90      H1   C
ATOM   6119  CB  SER   190     -49.228  41.754 -78.159  1.00 94.23      H1   C
ATOM   6120  OG  SER   190     -50.258  41.630 -77.192  1.00 94.02      H1   O
ATOM   6121  C   SER   190     -47.498  40.582 -79.512  1.00 95.63      H1   C
ATOM   6122  O   SER   190     -47.679  40.497 -80.727  1.00 95.93      H1   O
ATOM   6123  N   SER   191     -46.311  40.845 -78.974  1.00 96.22      H1   N
ATOM   6124  CA  SER   191     -45.161  41.202 -79.796  1.00 96.93      H1   C
ATOM   6125  OB  SER   191     -44.210  42.100 -79.002  1.00 97.27      H1   C
ATOM   6126  OG  SER   191     -44.862  43.293 -78.596  1.00 97.43      H1   O
ATOM   6127  C   SER   191     -44.405  39.981 -80.314  1.00 97.19      H1   C
ATOM   6128  O   SER   191     -43.393  40.116 -81.003  1.00 96.64      H1   O
ATOM   6129  N   LEU   192     -44.895  38.792 -79.982  1.00 97.72      H1   N
ATOM   6130  CA  LEU   192     -44.297  37.561 -80.489  1.00 98.54      H1   C
ATOM   6131  CB  LEU   192     -44.718  36.367 -79.626  1.00 98.53      H1   C
ATOM   6132  CG  LEU   192     -44.246  36.361 -78.170  1.00 98.25      H1   C
ATOM   6133  CD1 LEU   192     -44.734  35.095 -77.483  1.00 98.23      H1   C
ATOM   6134  CD2 LEU   192     -42.727  36.446 -78.118  1.00 97.62      H1   C
ATOM   6135  C   LEU   192     -44.719  37.323 -81.935  1.00 98.79      H1   C
ATOM   6136  O   LEU   192     -45.907  37.187 -82.232  1.00 98.91      H1   O
ATOM   6137  N   GLY   193     -43.738  37.272 -82.831  1.00 98.91      H1   N
ATOM   6138  CA  GLY   193     -44.034  37.097 -84.241  1.00 99.03      H1   C
ATOM   6139  C   GLY   193     -43.844  38.374 -85.037  1.00 98.92      H1   C
ATOM   6140  O   GLY   193     -43.566  38.330 -86.236  1.00 98.95      H1   O
ATOM   6141  N   THR   194     -43.995  39.515 -84.373  1.00 98.94      H1   N
ATOM   6142  CA  THR   194     -43.785  40.808 -85.018  1.00 98.85      H1   C
ATOM   6143  CB  THR   194     -45.009  41.738 -84.838  1.00 98.60      H1   C
ATOM   6144  OG1 THR   194     -45.217  41.997 -83.444  1.00 98.43      H1   O
ATOM   6145  CG2 THR   194     -46.257  41.093 -85.424  1.00 98.49      H1   C
ATOM   6146  C   THR   194     -42.556  41.508 -84.446  1.00 98.74      H1   C
ATOM   6147  O   THR   194     -42.136  42.554 -84.944  1.00 98.84      H1   O
ATOM   6148  N   GLN   195     -41.986  40.928 -83.394  1.00 98.32      H1   N
ATOM   6149  CA  GLN   195     -40.809  41.500 -82.750  1.00 97.95      H1   C
ATOM   6150  CB  GLN   195     -41.181  42.046 -81.368  1.00 97.62      H1   C
ATOM   6151  CG  GLN   195     -40.009  42.597 -80.572  1.00 97.74      H1   C
ATOM   6152  CD  GLN   195     -39.320  43.757 -81.265  1.00 98.15      H1   C
ATOM   6153  OE1 GLN   195     -38.732  43.594 -82.335  1.00 98.48      H1   O
ATOM   6154  NE2 GLN   195     -39.387  44.936 -80.656  1.00 97.92      H1   N
ATOM   6155  C   GLN   195     -39.694  40.465 -82.615  1.00 97.59      H1   C
ATOM   6156  O   GLN   195     -39.919  39.356 -82.127  1.00 97.42      H1   O
ATOM   6157  N   THR   196     -38.493  40.832 -83.052  1.00 96.87      H1   N
ATOM   6158  CA  THR   196     -37.335  39.951 -82.936  1.00 96.09      H1   C
ATOM   6159  CB  THR   196     -36.406  40.071 -84.167  1.00 96.10      H1   C
ATOM   6160  OG1 THR   196     -35.943  41.421 -84.292  1.00 96.22      H1   O
ATOM   6161  CG2 THR   196     -37.146  39.672 -85.434  1.00 95.59      H1   C
ATOM   6162  C   THR   196     -36.526  40.270 -81.681  1.00 95.48      H1   C
ATOM   6163  O   THR   196     -36.262  41.434 -81.375  1.00 95.11      H1   O
ATOM   6164  N   TYR   197     -36.137  39.225 -80.958  1.00 94.60      H1   N
ATOM   6165  CA  TYR   197     -35.347  39.389 -79.745  1.00 93.36      H1   C
ATOM   6166  CB  TYR   197     -36.104  38.817 -78.545  1.00 93.41      H1   C
ATOM   6167  CG  TYR   197     -37.457  39.455 -78.317  1.00 92.83      H1   C
ATOM   6168  CD1 TYR   197     -38.628  38.773 -78.621  1.00 92.54      H1   C
ATOM   6169  CE1 TYR   197     -39.867  39.354 -78.419  1.00 92.56      H1   C
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ATOM   6170  CD2 TYR   197     -37.561  40.741 -77.801  1.00 92.54      H1   C
ATOM   6171  CE2 TYR   197     -38.794  41.331 -77.596  1.00 92.67      H1   C
ATOM   6172  CZ  TYR   197     -39.945  40.633 -77.907  1.00 92.86      H1   C
ATOM   6173  OH  TYR   197     -41.175  41.221 -77.710  1.00 92.25      H1   O
ATOM   6174  C   TYR   197     -34.000  38.690 -79.888  1.00 92.48      H1   C
ATOM   6175  O   TYR   197     -33.937  37.498 -80.190  1.00 92.42      H1   O
ATOM   6176  N   ILE   198     -32.925  39.441 -79.670  1.00 91.09      H1   N
ATOM   6177  CA  ILE   198     -31.577  38.901 -79.783  1.00 89.67      H1   C
ATOM   6178  CB  ILE   198     -30.846  39.489 -81.011  1.00 89.35      H1   C
ATOM   6179  CG2 ILE   198     -29.408  38.996 -81.045  1.00 89.56      H1   C
ATOM   6180  CG1 ILE   198     -31.579  39.091 -82.295  1.00 88.73      H1   C
ATOM   6181  CD1 ILE   198     -30.918  39.598 -83.560  1.00 87.61      H1   C
ATOM   6182  C   ILE   198     -30.760  39.201 -78.530  1.00 89.23      H1   C
ATOM   6183  O   ILE   198     -30.556  40.362 -78.170  1.00 88.50      H1   O
ATOM   6184  N   CYS   199     -30.292  38.145 -77.870  1.00 88.78      H1   N
ATOM   6185  CA  CYS   199     -29.523  38.287 -76.640  1.00 87.96      H1   C
ATOM   6186  C   CYS   199     -28.039  38.438 -76.971  1.00 87.57      H1   C
ATOM   6187  O   CYS   199     -27.521  37.774 -77.868  1.00 87.15      H1   O
ATOM   6188  CB  CYS   199     -29.752  37.069 -75.731  1.00 87.80      H1   C
ATOM   6189  SG  CYS   199     -28.628  35.661 -76.005  1.00 86.80      H1   S
ATOM   6190  N   ASN   200     -27.363  39.323 -76.247  1.00 87.80      H1   N
ATOM   6191  CA  ASN   200     -26.000  39.714 -76.593  1.00 87.93      H1   C
ATOM   6192  CB  ASN   200     -25.917  41.235 -76.743  1.00 88.63      H1   C
ATOM   6193  CG  ASN   200     -27.051  41.801 -77.577  1.00 89.36      H1   C
ATOM   6194  OD1 ASN   200     -27.673  42.794 -77.201  1.00 89.44      H1   O
ATOM   6195  ND2 ASN   200     -27.326  41.171 -78.716  1.00 89.25      H1   N
ATOM   6196  C   ASN   200     -24.999  39.254 -75.539  1.00 87.45      H1   C
ATOM   6197  O   ASN   200     -24.758  39.948 -74.550  1.00 87.13      H1   O
ATOM   6198  N   VAL   201     -24.408  38.086 -75.766  1.00 86.80      H1   N
ATOM   6199  CA  VAL   201     -23.438  37.526 -74.835  1.00 85.93      H1   C
ATOM   6200  CB  VAL   201     -23.413  35.987 -74.926  1.00 85.26      H1   C
ATOM   6201  CG1 VAL   201     -22.441  35.421 -73.907  1.00 85.22      H1   C
ATOM   6202  CG2 VAL   201     -24.809  35.433 -74.698  1.00 84.35      H1   C
ATOM   6203  C   VAL   201     -22.038  38.062 -75.121  1.00 85.88      H1   C
ATOM   6204  O   VAL   201     -21.636  38.188 -76.277  1.00 85.83      H1   O
ATOM   6205  N   ASN   202     -21.302  38.382 -74.062  1.00 86.08      H1   N
ATOM   6206  CA  ASN   202     -19.929  38.856 -74.196  1.00 86.41      H1   C
ATOM   6207  CB  ASN   202     -19.897  40.388 -74.196  1.00 86.12      H1   C
ATOM   6208  CG  ASN   202     -18.503  40.945 -74.437  1.00 86.38      H1   C
ATOM   6209  OD1 ASN   202     -17.597  40.226 -74.860  1.00 86.49      H1   O
ATOM   6210  ND2 ASN   202     -18.327  42.234 -74.168  1.00 86.34      H1   N
ATOM   6211  C   ASN   202     -19.063  38.322 -73.056  1.00 86.59      H1   C
ATOM   6212  O   ASN   202     -19.273  38.666 -71.893  1.00 86.68      H1   O
ATOM   6213  N   HIS   203     -18.089  37.483 -73.397  1.00 87.00      H1   N
ATOM   6214  CA  HIS   203     -17.166  36.934 -72.409  1.00 87.34      H1   C
ATOM   6215  CB  HIS   203     -17.108  35.409 -72.537  1.00 88.04      H1   C
ATOM   6216  CG  HIS   203     -16.304  34.741 -71.465  1.00 89.65      H1   C
ATOM   6217  CD2 HIS   203     -16.174  35.015 -70.145  1.00 90.06      H1   C
ATOM   6218  ND1 HIS   203     -15.509  33.641 -71.707  1.00 89.81      H1   N
ATOM   6219  CE1 HIS   203     -14.923  33.267 -70.583  1.00 89.71      H1   C
ATOM   6220  NE2 HIS   203     -15.310  34.084 -69.620  1.00 90.14      H1   N
ATOM   6221  C   HIS   203     -15.772  37.524 -72.603  1.00 87.37      H1   C
ATOM   6222  O   HIS   203     -14.946  36.961 -73.318  1.00 87.61      H1   O
ATOM   6223  N   LYS   204     -15.517  38.656 -71.953  1.00 87.39      H1   N
ATOM   6224  CA  LYS   204     -14.282  39.411 -72.154  1.00 87.67      H1   C
ATOM   6225  CB  LYS   204     -14.296  40.671 -71.281  1.00 87.25      H1   C
ATOM   6226  CG  LYS   204     -15.279  41.733 -71.759  1.00 88.05      H1   C
ATOM   6227  CD  LYS   204     -15.674  42.676 -70.637  1.00 88.66      H1   C
ATOM   6228  CE  LYS   204     -14.453  43.279 -69.965  1.00 89.19      H1   C
ATOM   6229  NZ  LYS   204     -14.820  44.035 -68.736  1.00 90.12      H1   N
ATOM   6230  C   LYS   204     -12.996  38.621 -71.903  1.00 87.80      H1   C
ATOM   6231  O   LYS   204     -11.983  38.852 -72.566  1.00 88.12      H1   O
ATOM   6232  N   PRO   205     -13.013  37.683 -70.940  1.00 87.81      H1   N
ATOM   6233  CD  PRO   205     -14.006  37.497 -69.868  1.00 87.68      H1   C
ATOM   6234  CA  PRO   205     -11.854  36.799 -70.763  1.00 87.47      H1   C
ATOM   6235  CB  PRO   205     -12.254  35.915 -69.583  1.00 87.20      H1   C
ATOM   6236  CG  PRO   205     -13.239  36.737 -68.825  1.00 86.89      H1   C
ATOM   6237  C   PRO   205     -11.542  35.975 -72.012  1.00 87.26      H1   C
ATOM   6238  O   PRO   205     -10.378  35.786 -72.369  1.00 86.44      H1   O
ATOM   6239  N   SER   206     -12.589  35.492 -72.673  1.00 87.15      H1   N
ATOM   6240  CA  SER   206     -12.432  34.595 -73.812  1.00 87.33      H1   C
ATOM   6241  CB  SER   206     -13.477  33.480 -73.738  1.00 86.84      H1   C
ATOM   6242  OG  SER   206     -13.303  32.540 -74.781  1.00 87.39      H1   O
ATOM   6243  C   SER   206     -12.559  35.326 -75.150  1.00 87.95      H1   C
ATOM   6244  O   SER   206     -12.301  34.749 -76.209  1.00 87.21      H1   O
ATOM   6245  N   ASN   207     -12.955  36.594 -75.095  1.00 88.68      H1   N
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ATOM   6246  CA  ASN   207     -13.253  37.365 -76.299  1.00 88.90      H1   C
ATOM   6247  CB  ASN   207     -11.971  37.647 -77.088  1.00 89.10      H1   C
ATOM   6248  CG  ASN   207     -11.180  38.808 -76.515  1.00 89.83      H1   C
ATOM   6249  OD1 ASN   207     -10.011  39.004 -76.850  1.00 90.22      H1   O
ATOM   6250  ND2 ASN   207     -11.816  39.587 -75.645  1.00 90.17      H1   N
ATOM   6251  C   ASN   207     -14.260  36.641 -77.185  1.00 88.67      H1   C
ATOM   6252  O   ASN   207     -14.192  36.717 -78.411  1.00 88.61      H1   O
ATOM   6253  N   THR   208     -15.189  35.933 -76.551  1.00 88.73      H1   N
ATOM   6254  CA  THR   208     -16.290  35.300 -77.264  1.00 88.95      H1   C
ATOM   6255  CB  THR   208     -16.700  33.971 -76.583  1.00 88.70      H1   C
ATOM   6256  OG1 THR   208     -15.581  33.075 -76.577  1.00 88.37      H1   O
ATOM   6257  CG2 THR   208     -17.850  33.323 -77.350  1.00 87.89      H1   C
ATOM   6258  C   THR   208     -17.492  36.238 -77.271  1.00 89.54      H1   C
ATOM   6259  O   THR   208     -18.310  36.221 -76.352  1.00 89.28      H1   O
ATOM   6260  N   LYS   209     -17.586  37.059 -78.312  1.00 90.21      H1   N
ATOM   6261  CA  LYS   209     -18.681  38.012 -78.445  1.00 90.36      H1   C
ATOM   6262  CB  LYS   209     -18.158  39.319 -79.049  1.00 90.46      H1   C
ATOM   6263  CG  LYS   209     -18.875  40.570 -78.568  1.00 91.30      H1   C
ATOM   6264  CD  LYS   209     -20.221  40.752 -79.256  1.00 92.32      H1   C
ATOM   6265  CE  LYS   209     -20.057  41.171 -80.713  1.00 92.31      H1   C
ATOM   6266  NZ  LYS   209     -21.372  41.397 -81.382  1.00 91.57      H1   N
ATOM   6267  C   LYS   209     -19.748  37.396 -79.347  1.00 90.48      H1   C
ATOM   6268  O   LYS   209     -19.498  37.130 -80.523  1.00 91.28      H1   O
ATOM   6269  N   VAL   210     -20.934  37.163 -78.790  1.00 90.31      H1   N
ATOM   6270  CA  VAL   210     -21.966  36.387 -79.474  1.00 90.33      H1   C
ATOM   6271  CB  VAL   210     -22.164  35.008 -78.792  1.00 90.01      H1   C
ATOM   6272  CG1 VAL   210     -23.304  34.246 -79.450  1.00 88.87      H1   C
ATOM   6273  CG2 VAL   210     -20.880  34.200 -78.879  1.00 89.86      H1   C
ATOM   6274  C   VAL   210     -23.311  37.106 -79.519  1.00 90.85      H1   C
ATOM   6275  O   VAL   210     -23.579  38.000 -78.715  1.00 90.85      H1   O
ATOM   6276  N   ASP   211     -24.146  36.715 -80.477  1.00 91.03      H1   N
ATOM   6277  CA  ASP   211     -25.536  37.151 -80.526  1.00 91.23      H1   C
ATOM   6278  CB  ASP   211     -25.690  38.294 -81.532  1.00 91.67      H1   C
ATOM   6279  CG  ASP   211     -24.793  39.472 -81.228  1.00 92.43      H1   C
ATOM   6280  OD1 ASP   211     -25.173  40.317 -80.391  1.00 92.57      H1   O
ATOM   6281  OD2 ASP   211     -23.699  39.552 -81.828  1.00 92.96      H1   O
ATOM   6282  C   ASP   211     -26.416  35.975 -80.940  1.00 91.08      H1   C
ATOM   6283  O   ASP   211     -26.143  35.305 -81.935  1.00 91.24      H1   O
ATOM   6284  N   LYS   212     -27.466  35.720 -80.168  1.00 90.73      H1   N
ATOM   6285  CA  LYS   212     -28.407  34.659 -80.501  1.00 90.77      H1   C
ATOM   6286  CB  LYS   212     -28.410  33.585 -79.410  1.00 90.14      H1   C
ATOM   6287  CG  LYS   212     -29.376  32.438 -79.673  1.00 89.67      H1   C
ATOM   6288  CD  LYS   212     -29.021  31.683 -80.949  1.00 89.85      H1   C
ATOM   6289  CE  LYS   212     -27.672  30.983 -80.831  1.00 89.83      H1   C
ATOM   6290  NZ  LYS   212     -27.417  30.051 -81.967  1.00 89.22      H1   N
ATOM   6291  C   LYS   212     -29.810  35.226 -80.668  1.00 91.33      H1   C
ATOM   6292  O   LYS   212     -30.237  36.091 -79.902  1.00 91.11      H1   O
ATOM   6293  N   LYS   213     -30.523  34.732 -81.674  1.00 91.62      H1   N
ATOM   6294  CA  LYS   213     -31.861  35.221 -81.977  1.00 92.18      H1   C
ATOM   6295  CB  LYS   213     -32.032  35.352 -83.494  1.00 92.88      H1   C
ATOM   6296  CG  LYS   213     -33.204  36.218 -83.933  1.00 93.56      H1   C
ATOM   6297  CD  LYS   213     -33.200  36.412 -85.446  1.00 93.60      H1   C
ATOM   6298  CE  LYS   213     -34.314  37.347 -85.896  1.00 93.91      H1   C
ATOM   6299  NZ  LYS   213     -34.365  37.479 -87.380  1.00 92.57      H1   N
ATOM   6300  C   LYS   213     -32.893  34.253 -81.415  1.00 92.17      H1   C
ATOM   6301  O   LYS   213     -32.933  33.085 -81.799  1.00 92.12      H1   O
ATOM   6302  N   VAL   214     -33.723  34.739 -80.501  1.00 92.74      H1   N
ATOM   6303  CA  VAL   214     -34.723  33.891 -79.869  1.00 93.76      H1   C
ATOM   6304  CB  VAL   214     -34.883  34.231 -78.370  1.00 93.57      H1   C
ATOM   6305  CG1 VAL   214     -35.835  33.241 -77.709  1.00 92.68      H1   C
ATOM   6306  CG2 VAL   214     -33.524  34.212 -77.684  1.00 93.40      H1   C
ATOM   6307  C   VAL   214     -36.075  34.044 -80.557  1.00 94.58      H1   C
ATOM   6308  O   VAL   214     -36.666  35.126 -80.559  1.00 94.67      H1   O
ATOM   6309  N   GLU   215     -36.556  32.952 -81.141  1.00 95.50      H1   N
ATOM   6310  CA  GLU   215     -37.829  32.956 -81.850  1.00 96.61      H1   C
ATOM   6311  CB  GLU   215     -37.591  32.750 -83.347  1.00 97.66      H1   C
ATOM   6312  CG  GLU   215     -36.659  33.777 -83.970  1.00 99.30      H1   C
ATOM   6313  CD  GLU   215     -36.464  33.561 -85.458  1.00100.39      H1   C
ATOM   6314  OE1 GLU   215     -35.937  34.475 -86.128  1.00100.64      H1   O
ATOM   6315  OE2 GLU   215     -36.841  32.477 -85.956  1.00101.04      H1   O
ATOM   6316  C   GLU   215     -38.735  31.853 -81.313  1.00 96.73      H1   C
ATOM   6317  O   GLU   215     -38.263  30.886 -80.717  1.00 96.90      H1   O
ATOM   6318  N   PRO   216     -40.053  31.989 -81.519  1.00 97.02      H1   N
ATOM   6319  CD  PRO   216     -40.692  33.162 -82.139  1.00 96.97      H1   C
ATOM   6320  CA  PRO   216     -41.045  31.022 -81.032  1.00 97.59      H1   C
ATOM   6321  CB  PRO   216     -42.376  31.602 -81.507  1.00 97.22      H1   C
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ATOM   6322  CG  PRO   216     -42.109  33.068 -81.652  1.00 97.02      H1   C
ATOM   6323  C   PRO   216     -40.819  29.603 -81.553  1.00 98.15      H1   C
ATOM   6324  O   PRO   216     -40.189  29.404 -82.591  1.00 97.93      H1   O
ATOM   6325  N   LYS   217     -41.343  28.624 -80.821  1.00 99.25      H1   N
ATOM   6326  CA  LYS   217     -41.133  27.213 -81.137  1.00100.54      H1   C
ATOM   6327  CB  LYS   217     -41.567  26.340 -79.954  1.00100.87      H1   C
ATOM   6328  CG  LYS   217     -40.832  26.634 -78.652  1.00101.66      H1   C
ATOM   6329  CD  LYS   217     -41.398  25.815 -77.496  1.00101.93      H1   C
ATOM   6330  CE  LYS   217     -40.737  26.185 -76.173  1.00102.31      H1   C
ATOM   6331  NZ  LYS   217     -41.369  25.499 -75.006  1.00101.76      H1   N
ATOM   6332  C   LYS   217     -41.892  26.784 -82.392  1.00100.93      H1   C
ATOM   6333  O   LYS   217     -42.770  25.901 -82.277  1.00101.08      H1   O
ATOM   6334  OXT LYS   217     -41.600  27.333 -83.476  1.00101.13      H1   O
TER    6335       LYS   217                                             H1
END
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WHAT IS CLAIMED IS:

1. A monoclonal antibody or fragment thereof that competes for binding to human PCSK9 with:
(a) an antibody comprising a heavy chain variable region of the amino acid sequence in SEQ ID NO: 49; and a light chain variable region of the amino acid sequence in SEQ ID NO: 23; or
(b) an antibody comprising a heavy chain variable region of the amino acid sequence in SEQ ID NO: 67; and a light chain variable region of the amino acid sequence in SEQ ID NO: 12,
and wherein said monoclonal antibody or fragment thereof is neutralizing in that an excess of said monoclonal antibody or fragment thereof is capable of reducing the quantity of human PCSK9 bound to LDLR in an *in vitro* competitive binding assay.

2. The monoclonal antibody or fragment thereof of claim 1, wherein the monoclonal antibody or fragment thereof is capable of reducing the quantity of human PCSK9 bound to LDLR by at least about 80% in an *in vitro* competitive binding assay.

3. An isolated monoclonal antibody, wherein, when bound to human PCSK9, the monoclonal antibody binds to at least one of the following residues: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, or S381 of SEQ ID NO:3, and wherein the monoclonal antibody blocks binding of human PCSK9 to LDLR.

4. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least S153.

5. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least I154.

6. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least P155.

7. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least T377.

8. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least R194.

9. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least D238.

10. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least A239.

11. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least I369.

12. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least S372.

13. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least D374.

14. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least C375.

15. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least C378.

16. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least F379.

17. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least V380.

18. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody binds to at least S381.

19. The isolated monoclonal antibody of claim 3, wherein the isolated monoclonal antibody binds to at least two of the following residues: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, or S381 of SEQ ID NO:3.

20. The isolated monoclonal antibody of claim 3, wherein the isolated monoclonal antibody binds to at least four of the following residues: S153, I154, P155, R194, D238, A239, I369, S372, D374, C375, T377, C378, F379, V380, or S381 of SEQ ID NO:3.

21. The isolated monoclonal antibody of claim 3, wherein the monoclonal antibody blocks binding of PCSK9 to LDLR by at least 80%.

22. The isolated monoclonal antibody of claim 19, wherein the monoclonal antibody blocks binding of PCSK9 to LDLR by at least 80%.

23. The isolated monoclonal antibody of claim 20, wherein the monoclonal antibody blocks binding of PCSK9 to LDLR by at least 80%.

24. A monoclonal antibody or fragment thereof that specifically binds to human PCSK9 and is neutralizing in that an excess of said antigen binding protein is capable of reducing the quantity of human PCSK9 bound to LDLR in an *in vitro* competitive binding assay, said monoclonal antibody or fragment thereof comprising a combination of a light chain variable domain and a heavy chain variable domain selected from the group of combinations consisting of:
a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 9 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 71;
a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 10 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 72;
a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 12 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 67;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 13 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 87;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 16 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 52;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 17 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 51;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 20 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 54;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 21 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 55;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 22 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 56;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 23, and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 49;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 24 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 57;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 26 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 50;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 30 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 64;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 31 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 62;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 33 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 65;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 35 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 79;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 36 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 80;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 37 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 76;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 38 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 77;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 39 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 78;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 40 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 83;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 461 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 459;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 465, and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 463;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 473 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO:471;

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 477 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO:475; and

a light chain variable domain having a sequence with at least 90% identity to SEQ ID NO: 485 and a heavy chain variable domain having a sequence with at least 90% identity to SEQ ID NO:483.

25. The monoclonal antibody or fragment thereof of Claim 24, wherein the light chain variable domain and the heavy chain variable domain are selected from the group of combinations consisting of: a light chain variable domain having the sequence set forth in SEQ ID NO: 10 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 72; a light chain variable domain having the sequence set forth in SEQ ID NO: 12 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 67; a light chain variable domain having the sequence set forth in SEQ ID NO: 17 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 51; a light chain variable domain having the sequence set forth in SEQ ID NO: 23, and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 49; a light chain variable domain having the sequence set forth in SEQ ID NO: 35 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 79; a light chain variable domain having the sequence set forth in SEQ ID NO: 461 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 459; and a light chain variable domain having the sequence set forth in SEQ ID NO: 465, and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 463; a light chain variable domain having the sequence set forth in SEQ ID NO: 485 and a heavy chain variable domain having the sequence set forth in SEQ ID NO:483, wherein the antigen binding protein specifically binds to the PCSK9.

26. A monoclonal antibody or fragment thereof, wherein said monoclonal antibody comprises:

(a) a light chain variable domain comprising the amino acid sequence of SEQ ID NO: 23, and a heavy chain variable domain comprising the amino acid sequence of SEQ ID NO: 49; or

(b) a light chain variable domain comprising the amino acid sequence of SEQ ID NO: 23, wherein the first residue is a glutamic acid instead of a glutamine, and a heavy chain variable domain comprising the amino acid sequence of SEQ ID NO: 49, wherein the first residue is a glutamic acid instead of a glutamine.

27. A monoclonal antibody or fragment thereof that specifically binds to human PCSK9 and is neutralizing in that an excess of said monoclonal antibody or fragment thereof is capable of reducing the quantity of human PCSK9 bound to LDLR in an *in vitro* competitive binding assay, said monoclonal antibody or fragment thereof comprising a heavy chain complementarity determining region (CDR) of the CDRH1 sequence in SEQ ID NO: 51, a CDRH2 of the CDRH2 sequence in SEQ ID NO: 51, and a CDRH3 of the CDRH3 sequence in SEQ ID NO: 51, and a light chain CDRL1 of the CDRL1 sequence in SEQ ID NO: 17, a CDRL2 of the CDRL2 sequence in SEQ ID NO: 17, and a CDRL3 of the CDRL3 sequence in SEQ ID NO: 17.

28. A monoclonal antibody or fragment thereof that specifically binds to human PCSK9 and is neutralizing in that an excess of said monoclonal antibody or fragment thereof is capable of reducing the quantity of human PCSK9 bound to LDLR in an *in vitro* competitive binding assay, said monoclonal antibody or fragment thereof comprising a heavy chain complementarity determining region (CDR) of the CDRH1 sequence in SEQ ID NO: 459, a CDRH2 of the CDRH2 sequence in SEQ ID NO: 459, and a CDRH3 of the CDRH3 sequence in SEQ ID NO: 459, and a light chain CDRL1 of the CDRL1 sequence in SEQ ID NO: 461, a CDRL2 of the CDRL2 sequence in SEQ ID NO: 461, and a CDRL3 of the CDRL3 sequence in SEQ ID NO: 461.

29. A monoclonal antibody or fragment thereof that specifically binds to human PCSK9 and is neutralizing in that an excess of said monoclonal antibody or fragment thereof is capable of reducing the quantity of human PCSK9 bound to LDLR in an *in vitro* competitive binding assay, said monoclonal antibody or fragment thereof comprising a heavy chain complementarity determining region (CDR) of the CDRH1 sequence in SEQ ID NO: 49, a CDRH2 of the CDRH2 sequence in SEQ ID NO: 49, and a CDRH3 of the CDRH3 sequence in SEQ ID NO: 49, and a light chain CDRL1 of the CDRL1 sequence in SEQ ID NO: 23, a CDRL2 of the CDRL2 sequence in SEQ ID NO: 23, and a CDRL3 of the CDRL3 sequence in SEQ ID NO: 23.

30. The monoclonal antibody or fragment thereof of claim 29, wherein the monoclonal antibody or fragment thereof comprises all six of the following amino acid sequences: SEQ ID NO: 308, SEQ ID NO: 175, SEQ ID NO: 180, SEQ ID NO: 158, SEQ ID NO: 162, and SEQ ID NO: 395.

31. The monoclonal antibody or fragment thereof of claim 29, wherein the monoclonal antibody or fragment thereof comprises all six of the following amino acid sequences: SEQ ID NO: 368, SEQ ID NO: 175, SEQ ID NO: 180, SEQ ID NO: 158, SEQ ID NO: 162, and SEQ ID NO: 395.

32. The monoclonal antibody or fragment thereof of claim 29, wherein the monoclonal antibody or fragment thereof comprises: a CDRH1 comprising SEQ ID NO: 308, a CDRH2 comprising SEQ ID NO: 175, and a CDRH3 comprising SEQ ID NO: 180, and a CDRL1 comprising SEQ ID NO: 158, a CDRL2 comprising SEQ ID NO: 162, and a CDRL3 comprising SEQ ID NO: 395.

33. The monoclonal antibody or fragment thereof of claim 29, wherein the monoclonal antibody or fragment thereof comprises: a CDRH1 comprising SEQ ID NO: 368, a CDRH2 comprising SEQ ID NO: 175, and a CDRH3 comprising SEQ ID NO: 180, and a CDRL1 comprising SEQ ID NO: 158, a CDRL2 comprising SEQ ID NO: 162, and a CDRL3 comprising SEQ ID NO: 395.

34. The monoclonal antibody or fragment thereof of claim 29 comprising a light chain variable domain comprising the amino acid sequence of SEQ ID NO:23, and a heavy chain variable domain comprising the amino acid sequence of SEQ ID NO:49.

35. The monoclonal antibody or fragment thereof of claim 34 wherein the first residue of the light chain variable domain is a glutamic acid instead of a glutamine, and wherein the first residue of the heavy chain variable domain is a glutamic acid instead of a glutamine.

36. The monoclonal antibody or fragment thereof of claim 35, further comprising a glycine residue at the C-terminal end of said light chain variable domain.

37. The monoclonal antibody or fragment thereof of claim 27, wherein the monoclonal antibody or fragment thereof comprises all six of the following amino acid sequences: SEQ ID NO: 368, SEQ ID NO: 174, SEQ ID NO: 180, SEQ ID NO: 158, SEQ ID NO: 162, and SEQ ID NO:164.

38. The monoclonal antibody or fragment thereof of claim 27, wherein the monoclonal antibody or fragment thereof comprises: a CDRH1 comprising SEQ ID NO: 368, a CDRH2 comprising SEQ ID NO: 174, and a CDRH3 comprising SEQ ID NO: 180, and a CDRL1 comprising SEQ ID NO: 158, a CDRL2 comprising SEQ ID NO: 162, and a CDRL3 comprising SEQ ID NO: 164.

39. The monoclonal antibody or fragment thereof of claim 37 comprising a light chain variable domain comprising the amino acid sequence of SEQ ID NO:17, and a heavy chain variable domain comprising the amino acid sequence of SEQ ID NO:51.

40. The monoclonal antibody or fragment thereof of claim 28, wherein the monoclonal antibody or fragment thereof comprises all six of the following amino acid sequences: GFTFSSYWMS, SIKQDGSEKYYVDSVKG, DLVLMVYDIDYYYYGMDV, RSSQSLLHSNGYNYLD, LGSNRAS and MQALQTPLT.

41. The monoclonal antibody or fragment thereof of claim 28, wherein the monoclonal antibody or fragment thereof comprises: a CDRH1 comprising GFTFSSYWMS, a CDRH2 comprising SIKQDGSEKYYVDSVKG, and a CDRH3 comprising DLVLMVYDIDYYYYGMDV, and a CDRL1 comprising RSSQSLLHSNGYNYLD, a CDRL2 comprising LGSNRAS, and a CDRL3 comprising MQALQTPLT.

42. The monoclonal antibody or fragment thereof of claim 28 comprising a light chain variable domain comprising the amino acid sequence of SEQ ID NO:461, and a heavy chain variable domain comprising the amino acid sequence of SEQ ID NO:459.

43. An isolated monoclonal antibody or fragment thereof that binds to a human PCSK9 protein comprising a heavy chain polypeptide comprising residue 1 to residue 115 of SEQ ID NO:298 and a light chain polypeptide comprising residue 1 to 110 of SEQ ID NO:297.

44. The monoclonal antibody or fragment thereof of any one of claims 24 to 43, further comprising:

(a) the light chain constant sequence of SEQ ID NO: 156;

(b) the light chain constant sequence of SEQ ID NO: 157;

(c) the heavy chain constant sequence of SEQ ID NO: 154;

(d) the heavy chain constant sequence of SEQ ID NO: 155;

(e) the light chain constant sequence of SEQ ID NO: 156 and the heavy chain constant sequence of SEQ ID NO: 154,

(f) the light chain constant sequence of SEQ ID NO: 157 and the heavy chain constant sequence of SEQ ID NO: 154;

(g) the light chain constant sequence of SEQ ID NO: 156 and the heavy chain constant sequence of SEQ ID NO: 155; or

(h) the light chain constant sequence of SEQ ID NO: 157 and the heavy chain constant sequence of SEQ ID NO: 155.

45. The monoclonal antibody or fragment thereof of any one of the preceding claims, wherein the monoclonal antibody or fragment thereof is an LDLR competitive neutralizing monoclonal antibody or fragment thereof.

46. The monoclonal antibody or fragment thereof of any one of the preceding claims, wherein the monoclonal antibody or fragment thereof has at least one of the following characteristics:

(a) binds to a PCSK9 variant that has a D374Y point mutation;

(b) is selected from the group consisting of a human antibody, a humanized antibody, a chimeric antibody, a multispecific antibody, a recombinant antibody, a single chain antibody, a diabody, a Fab fragment, an $F(ab)_2$ fragment, an IgG1 antibody, an IgG2 antibody, and IgG3 antibody, and an IgG4 antibody;

(c) binds to PCSK9 with a K_d that is smaller than 1 nM, is smaller than 100 pM, is smaller than 10 pM, or is smaller than 5 pM; or

(d) can block the binding of D374Y PCSK9 to LDLR with an IC_{50} of less than 200 pM.

47. The monoclonal antibody or fragment thereof of any one of the preceding claims that, when bound to human PCSK9:

(a) binds to PCSK9 with the same Kd as a reference antibody;

(b) inhibits binding of PCSK9 to LDLR with the same IC_{50} as said reference antibody; or

(c) cross-competes for binding with said reference antibody to PCSK9,

wherein said reference antibody comprises a combination of light chain and heavy chain variable domain sequences selected from the group consisting of a light chain variable domain having the sequence as set forth in SEQ ID NO: 10 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 72; a light chain variable domain having the sequence as set forth in SEQ ID NO: 12 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 67; a light chain variable domain having the sequence set forth in SEQ ID NO: 17 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 51; a light chain variable domain having the sequence set forth in SEQ ID NO: 23, and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 49; a light chain variable domain having the sequence set forth in SEQ ID NO: 35 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 79; a light chain variable domain having the sequence set forth in SEQ ID NO: 461 and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 459; a light chain variable domain having the sequence set forth in SEQ ID NO: 465, and a heavy chain variable domain having the sequence set forth in SEQ ID NO: 463; and a light chain variable domain having the sequence set forth in SEQ ID NO: 485 and a heavy chain variable domain having the sequence set forth in SEQ ID NO:483.

48. A nucleic acid encoding the monoclonal antibody or fragment thereof of any one of the preceding claims.

49. A recombinant expression vector comprising the nucleic acid molecule according to claim 48.

50. A host cell comprising the vector of claim 49.

51. A hybridoma capable of producing the monoclonal antibody or fragment thereof of claim 47.

52. A method of making the monoclonal antibody or fragment thereof of any one of claims 1-47, comprising the step of preparing said monoclonal antibody or fragment thereof from a host cell that secretes said monoclonal antibody or fragment thereof.

53. A monoclonal antibody or fragment thereof produced by the expression of recombinant DNA in the host cell of claim 50 or the hybridoma of claim 51.

54. The method of claim 52 or the monoclonal antibody or fragment thereof or antigen binding protein of claim 53, wherein the host cell is selected from the group consisting of Chinese hamster ovary (CHO) cells, HeLa cells, baby hamster kidney (BHK) cells, monkey kidney cells (COS), human hepatocellular carcinoma cells, and human epithelial kidney 293 cells.

55. A pharmaceutical composition comprising at least one monoclonal antibody or fragment thereof according to any one of claims 1 to 47, or 53 and a pharmaceutically acceptable excipient.

56. The pharmaceutical composition of claim 55, wherein the pharmaceutical composition is to be administered prior to, concurrent with, or subsequent to, the administration of at least one other therapeutic agent, wherein optionally the other therapeutic agent is a statin.

57. A kit for the treatment of cholesterol related disorders comprising the composition of claims 55 or 56.

58. Use of the monoclonal antibody or fragment thereof of any one of claims 1 to 47 or 53 in the manufacture of a medicament for treating or preventing a condition selected from hypercholesterolemia, cardiovascular disease, metabolic syndrome, diabetes, stroke, and dyslipidemia.

59. The monoclonal antibody or fragment thereof of any one of claims 27 to 29, wherein each CDR is defined in accordance with the Kabat definition, the Chothia definition, the combination of the Kabat definition and the Chothia definition, the AbM definition, or the contact definition of CDR.

60. The monoclonal antibody or fragment thereof of Claim 59, wherein each CDR is defined in accordance with the CDR definition of Kabat.

61. The monoclonal antibody or fragment thereof of Claim 59, wherein each CDR is defined in accordance with the CDR definition of Chothia.

62. The monoclonal antibody or fragment thereof of Claim 59, wherein each CDR is defined in accordance with the combination of the Kabat definition and the Chothia definition.

63. An isolated neutralizing human monoclonal antibody that binds to a PCSK9 protein comprising:
a heavy chain polypeptide comprising:
the amino acid sequence of SEQ ID NO: 180;
the amino acid sequence of SEQ ID NO: 175; and
the amino acid sequence of SEQ ID NO: 308; and
a light chain polypeptide comprising:
the amino acid sequence of SEQ ID NO: 395;
the amino acid sequence of SEQ ID NO: 162; and
the amino acid sequence of SEQ ID NO: 158.