P-CADHERIN ANTIBODIES

This application claims priority to U.S. Provisional Application No. 60/675,311 filed on April 26, 2005, which is incorporated herein by reference in its entirety.

Field of the Invention

The present invention relates to antibodies and antigen-binding portions thereof that bind to P-cadherin. The invention also relates to nucleic acid molecules encoding such antibodies and antigen-binding portions, methods of making P-cadherin antibodies and antigen-binding portions, compositions comprising these antibodies and antigen-binding portions and methods of using the antibodies, antigen-binding portions, and compositions.

Background of the Invention

Cadherins are a superfamily of transmembrane glycoproteins that regulate cell-cell adhesion during development and tissue homeostasis (Gumbiner *J. Cell. Biol.*, 148:399-404 (2000); Yagi, et al., *Genes Dev.*, 14:1169-1180 (2000)). The intracellular domains of cadherins interact with cytoplasmic proteins such as catenins and p120, which form the basis of cadherin attachment to the actin cytoskeleton. Cadherins have five extracellular Ca^{2+} binding domains and a small cytoplasmic domain that is highly conserved among the classical cadherins. Members of the classical cadherin family include P-cadherin, E-cadherin, and N-cadherin. Cellular adhesion molecules such as cadherins are considered to play a significant role in the cellular connections of cancer and metastatic cells (Furukawa, et al., *Microscopy Res. Technique* 38 (4):343-352 (1997)). P-cadherin expression in normal adult tissues is low and is restricted primarily to myoepithelial cells and the basal layers of stratified epithelium (Shimoyama, et al. *Cancer Res.* 49:2128-33 (1989)). P-cadherin is upregulated in inflammatory bowel diseases such as Crohn's disease and colitis (Hardy, et al., *Gut* 50:513-519 (2002)). A large body of evidence now also reveals that aberrant P-cadherin expression is associated with cell proliferation and with tumors of the colon, breast, lung, thyroid, and cervix (Gamallo, *Modern Pathology*, 14:650-654, (2001); and Stefansson, et al., *J. Clin. Oncol.* 22(7):1242-1252 (2004)). Human P-cadherin was reported to be the antigen recognized by the NCC-CAD-299 monoclonal antibody raised against a vulvar epidermoid carcinoma (Shimoyama, et al., *Cancer Res.*, 49:2128-2133 (1989)). Modulation of P-cadherin mediated adhesion and intracellular signaling is expected to result in decreased proliferation and survival of tumor cells *in vivo*. Accordingly, in view of the pivotal role that P-cadherin appears to possess in cell proliferation and solid tumor progression, it is desirable to generate antibodies to P-cadherin that can provide a therapeutic benefit to patients with a variety of cancers.

Summary of the Invention

In one aspect of the present invention is a P-cadherin antibody or antigen-binding portion thereof wherein the antibody or antigen-binding portion thereof has at least one of several functional properties as described below in A) thru K).

A) For example, in one embodiment the antibodies or antigen-binding portions thereof have a greater binding affinity for P-cadherin (K_D(P)) than for E-cadherin (K_D(E)). In one embodiment, the antibodies or antigen-binding portions thereof of the present invention have a K_D(E)/K_D(P) that is

greater than or equal to 1.5. In a further embodiment the antibodies or antigen-binding portions thereof of the present invention have a $K_D(E)/K_D(P)$ that is greater than or equal 2, greater than or equal to 3, greater than or equal to 5, greater than or equal to 10, greater than or equal to 20, greater than or equal to 50, greater than or equal to 100, greater than or equal to 200, greater than or equal to 500, or greater than or equal to 1000. Typically there is no upper limit on the value of $K_D(E)/K_D(P)$ because the $K_D(E)$ value can be very small, such as 0. For practical purposes, however, an upper limit of $K_D(E)/K_D(P)$ can be 1×10^6. Such K_D values for both P-cadherin and for E-cadherin can be measured by any technique known to those of skill in the art, such as by ELISA, RIA, flow cytometry, or surface plasmon resonance, such as BIACORE™.

B) In another embodiment, the antibody or portion thereof binds to P-cadherin with a K_D of 1000 nM or less as measured by surface plasmon resonance. In a further embodiment, the antibody or portion binds to P-cadherin with a K_D of less than 500 nM, less than 100 nM, less than 50 nM, less than 20 nM, less than 10 nM, less than 1 nM, less than 500 pM, or less than 100 pM, as measured by surface plasmon resonance. Typically, there is no lower limit on the value of K_D. For practical purposes, however, the lower limit can be assumed to be about 1 pM.

C) In another embodiment, the antibody or portion thereof has an off rate (k_{off}) for P-cadherin of less than or equal to 0.01 $^{s-1}$ as measured by surface plasmon resonance. For example, in certain embodiments the antibody or portion has a k_{off} for P-cadherin of less than 0.005 $^{s-1}$, less than 0.004 $^{s-1}$, less than 0.003 $^{s-1}$, less than 0.002 $^{s-1}$, or less than 0.001 $^{s-1}$. Typically, there is no lower limit for the value of k_{off}. For practical purposes, however, the lower limit can be assumed to be about $1x10^{-7}$ $^{s-1}$.

D) In another embodiment, the P-cadherin antibody or portion thereof has an IC_{50} of 100 nM or less as measured by a P-cadherin dependent cell adhesion assay. In a further embodiment, said IC_{50} is less than 50 nM, less than 40 nM, less than 20 nM, less than 10 nM, less than 1 nM, less than 500 pM, less than 200 pM, less than 100 pM, or less than 10 pM, as measured by a P-cadherin dependent cell adhesion assay. Typically, there is no lower limit for the value of IC_{50} as measured by a P-cadherin dependent cell adhesion assay. For practical purposes, however, the lower limit can be assumed to be about 1 pM.

E) In another embodiment, the P-cadherin antibody or portion thereof has an IC_{50} of 100 nM or less as measured by a P-cadherin dependent cell aggregation assay. In a further embodiment, said IC_{50} is less than 50 nM, less than 40 nM, less than 20 nM, less than 10 nM, less than 1 nM, less than 500 pM, less than 200 pM, less than 100 pM, or less than 1 pM, as measured by a P-cadherin dependent cell aggregation assay. Typically, there is no lower limit for the value of IC_{50} as measured by a P-cadherin dependent cell aggregation assay. For practical purposes, however, the lower limit can be assumed to be about 1 pM.

F) In another embodiment, the P-cadherin antibody or portion thereof increases spheroid disruption in a P-cadherin-dependent spheroid disruption assay by a factor of at least 2 as compared to a control sample with no IgG present. In a further embodiment, the P-cadherin antibody or portion thereof increases spheroid disruption in a P-cadherin-dependent spheroid disruption assay by a

factor of at least 3, at least 4, at least 6, at least 10, or at least 15 as compared to a control sample with no IgG present.

G) In another embodiment, the P-cadherin antibody or portion thereof competes for binding to P-cadherin with an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4.

H) In another embodiment, the P-cadherin antibody or portion thereof cross-competes for binding to P-cadherin with an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06129-1c4; and g-129-1c4.

I) In another embodiment, the P-cadherin antibody or portion thereof binds to the same epitope of P-cadherin as an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4.

J) In another embodiment, the P-cadherin antibody or portion thereof binds to P-cadherin with substantially the same K_D as an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4.

K) In another embodiment, the P-cadherin antibody or portion thereof binds to P-cadherin with substantially the same k_{off} as an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4.

A further aspect of the present invention is an antibody or antigen-binding portion thereof with at least one of the functional propertes described previously in A) thru K), comprising a V_H domain that is at least 90% identical in amino acid sequence to any one of SEQ ID NOs: 1 to 13 and 320 to 325. In one embodiment, said V_H domain is at least 91%, at least 93%, at least 95%, at least 97%, at least 99%, or 100% identical in amino acid sequence to any one of SEQ ID NOs: 1 to 12 and 320 to 325.

In a further embodiment, the antibody or portion thereof has at least one of the functional properties described previously in A) thru K), comprising a V_H domain that is any one of SEQ ID NOs: 1 to 13 and 320 to 325, or differs from any one of SEQ ID NOs: 1 to 13 and 320 to 325 by having at least one conservative amino acid substitution. For example, the V_H domain can differ by 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 conversative amino acid substitutions from any one of SEQ ID NOs: 1 to 13 and 320 to 325. In a further embodiment, any of these conservative amino acid substitutions can occur in the CDR1, CDR2, and/or CDR3 regions.

A further aspect of the present invention is an antibody or antigen-binding portion thereof with at least one of the functional properties described previously in A) thru K), comprising a V_L

domain that is at least 90% identical in amino acid sequence to any one of SEQ ID NOs: 14 to 23 and 326 to 331. In one embodiment, said V_L domain is at least 91%, at least 93%, at least 95%, at least 97%, at least 99%, or 100% identical in amino acid sequence to any one of SEQ ID NOs: 14 to 23 and 326 to 331.

In a further embodiment, the antibody or portion thereof has at least one of the functional properties described previously in A) thru K), and comprises a V_L domain that is any one of SEQ ID NOs: 14 to 23 and 326 to 331, or differs from any one of SEQ ID Nos: 14 to 23 and 326 to 331 by having at least one conservative amino acid substitution. For example, the V_L domain can differ by 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 converative amino acid substitutions from any one of SEQ ID NOs: 14 to 23 and 326 to 331. In a further embodiment, any of these conservative amino acid substitutions can occur in the CDR1, CDR2, and/or CDR3 regions.

Another aspect of the present invention is an antibody or antigen-binding portion thereof with at least one of the functional properties described previously in A) thru K), wherein the V_L and V_H domains are each at least 90% identical in amino acid sequence to the V_L and V_H domains, respectively, of any one of antibodies 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4. For example, the V_L and V_H domains are each at least 91%, 93%, 95%, 97%, 99% or 100% identical in amino acid sequences to the V_L and V_H domains, respectively, of any one of antibodies 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4.

In another aspect of the present invention is an antibody or antigen-binding portion thereof that is selected from the group consisting of: a) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 1, and a V_L domain as set forth in SEQ ID NO: 14; b) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 2, and a V_L domain as set forth in SEQ ID NO: 14; c) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 2 and a V_L domain as set forth in SEQ ID NO: 15; d) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 3, and a V_L domain as set forth in SEQ ID NO: 16; e) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 4 and a V_L domain as set forth in SEQ ID NO: 17; f) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 4 and a V_L domain as set forth in SEQ ID NO: 23; g) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 5 and a V_L domain as set forth in SEQ ID NO: 18; h) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 6 and a V_L domain as set forth in SEQ ID NO: 23; i) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 7 and a V_L domain as set forth in SEQ ID NO: 23; j) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 8 and a V_L domain as set forth in SEQ ID NO: 23; k) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 9 and a V_L domain as set forth in SEQ ID NO: 23; l) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 10 and a V_L domain as set forth in

SEQ ID NO: 19; m) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 11 and a V_L domain as set forth in SEQ ID NO: 20; n) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 12 and a V_L domain as set forth in SEQ ID NO: 21; o) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 13 and a V_L domain as set forth in SEQ ID NO: 22; p) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 320 and a V_L domain as set forth in SEQ ID NO: 326; q) an antibody or antigen-binding portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 321 and a V_L domain as set forth in SEQ ID NO: 327; r) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 322 and a V_L domain as set forth in SEQ ID NO: 328; s) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 323 and a V_L domain as set forth in SEQ ID NO: 329; t) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 324 and a V_L domain as set forth in SEQ ID NO: 330; and u) an antibody or portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 325, and a V_L domain as set forth in SEQ ID NO: 331.

In a further embodiment, for any of the antibodies or portions thereof as described above in groups a) to u) the V_H and/or V_L domains can differ from the specific SEQ ID NOs recited therein by at least one conservative amino acid substitution. For example, the V_H and/or V_L domains can differ from the recited SEQ ID NO by 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 conservative amino acid substitutions. In a further embodiment, any of these conservative amino acid substitutions can occur in the CDR1, CDR2, and/or CDR3 regions.

In another embodiment, the present invention provides an antibody or antigen-binding portion thereof with at least one of the functional properties described previously in A) thru K), comprising a V_H domain that is independently selected from any one of SEQ ID NOs: 1 to 13 and 320 to 325, or a sequence that differs from any one of SEQ ID NOs: 1 to 13 and 320 to 325 by at least one conservative amino acid substitution, and further comprises a V_L domain that is independently selected from any one of SEQ ID NOs: 14 to 23 and 326 to 331, or a sequence that differs from any one of SEQ ID NOs: 14 to 23 and 326 to 331 by at least one conservative amino acid substitution. For example, the V_H and V_L domains can each independently differ from any one of SEQ ID NOs: 1 to 13, 320 to 325, 14 to 23, and 326 to 331 by 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 conservative amino acid substitutions.

In a further embodiment, the present invention provides an antibody or antigen-binding portion thereof with at least one of the functional properties described previously in A) thru K), wherein said antibody or portion comprises a V_H CDR3 selected from any one of SEQ ID NOs: 26 to 37 and 91 to 256, or a sequence that differs from any one of SEQ ID NOs: 26 to 37 and 91 to 256 by at least one conservative amino acid substitution. For example, the V_H CDR3 can differ from any one of SEQ ID NOs: 26 to 37 and 91 to 256 by 1, 2, 3, or 4 conservative amino acid substitutions.

In a further embodiment, the present invention provides an antibody or antigen-binding portion thereof with at least one of the functional properties described previously in A) thru K), wherein said antibody or antigen-binding portion comprises a V_L CDR3 selected from any one of

SEQ ID NOs: 40 to 47 and 257 to 319, or a sequence that differs from any one of SEQ ID NOs: 40 to 47 and 257 to 319 by at least one conservative amino acid substitution. For example, the V_L CDR3 can differ from any one of SEQ ID NOs: 40 to 47 and 257 to 319 by 1, 2, 3, 4, or 5 conservative amino acid substitutions.

In a further embodiment, the present invention provides an antibody or antigen-binding portion thereof thereof, wherein said antibody or antigen-binding portion comprises: a V_H CDR1 as set forth in SEQ ID NO: 24, or a sequence that differs from SEQ ID NO: 24 by at least one conservative amino acid substitution; a V_H CDR2 as set forth in SEQ ID NO: 25 or a sequence that differs from SEQ ID NO: 25 by at least one conservative amino acid substitution; and a V_H CDR3 that is independently selected from any one of SEQ ID NOs: 26 to 37 and 91 to 256, or a sequence that differs from any one of SEQ ID NOs: 26 to 37 and 91 to 256 by at least one conservative amino acid substitution. For example, the V_H CDR1, CDR2, and CDR3 sequences mentioned above can each independently differ from the respective recited SEQ ID NOs by 1, 2, 3, 4 or 5 conservative amino acid substitutions.

In another embodiment, the present invention provides an antibody or antigen-binding portion thereof, wherein said antibody or antigen-binding portion comprises: a V_L CDR1 as set forth in SEQ ID NO: 38, or a sequence that differs from SEQ ID NO: 38 by at least one conservative amino acid substitution; a V_L CDR2 as set forth in SEQ ID NO: 39, or a sequence that differs from SEQ ID NO: 39 by at least one conservative amino acid substitution; and a V_L CDR3 that is independently selected from any one of SEQ ID NOs: 40 to 47 and 257 to 319, or a sequence that differs from any one of SEQ ID NOs: 40 to 47 and 257 to 319 by at least one conservative amino acid substitution. For example, the V_L CDR1, CDR2, and CDR3 sequences mentioned above can each independently differ from the respective recited SEQ ID NOs by 1, 2, 3, 4 or 5 conservative amino acid substitutions.

The present invention further provides an antibody or antigen-binding portion thereof, wherein said antibody or antigen-binding portion comprises: a V_H CDR1 as set forth in SEQ ID NO: 24; a V_H CDR2 as set forth in SEQ ID NO: 25; a V_H CDR3 selected from any one of SEQ ID NOs: 26 to 37 and 91 to 256; a V_L CDR1 as set forth in SEQ ID NO: 38; a V_L CDR2 as set forth in SEQ ID NO: 39; and a V_L CDR3 selected from any one of SEQ ID NOs: 40 to 47 and 257 to 319. In a further embodiment, the V_H and V_L CDR1, CDR2, and CDR3 sequences mentioned can also each independently differ from the specific SEQ ID NOs recited above by at least one conservative amino acid substitution. For example, the CDR1, CDR2, and CDR3 sequences can each independently differ by 1, 2, 3, 4, or 5 conservative amino acid substitutions from the respective specific SEQ ID NOs recited above.

The present invention further provides an antibody or antigen-binding portion thereof wherein said antibody or antigen-binding portion comprises the V_H and V_L CDR1, the V_H and V_L CDR2, and the V_H and V_L CDR3 as found in any of antibodies 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g129-1c4.

The present invention further provides an antibody or antigen-binding portion thereof, comprising a V_H domain selected from any one of SEQ ID NOs: 1 to 13 and 320 to 325, or differs from any one of SEQ ID NOs: 1 to 13 and 320 to 325 by 1 to 10 conservative amino acid substitutions.

The present invention further provides an antibody or antigen-binding portion thereof, comprising a V_L domain selected from any one of SEQ ID NOs: 14 to 23 and 326 to 331, or differs from any one of SEQ ID NOs: 14 to 23 and 326 to 331 by 1 to 10 conservative amino acid substitutions.

The present invention further provides an antibody or antigen-binding portion thereof, comprising a V_H domain that is independently selected from any one of SEQ ID NOs: 1 to 13 and 320 to 325, or a sequence that differs from any one of SEQ ID NOs: 1 to 13 and 320 to 325 by 1 to 10 conservative amino acid substitutions, and further comprising a V_L domain that is independently selected from any one of SEQ ID NOs: 14 to 23 and 326 to 331, or a sequence that differs from any one of SEQ ID NOs: 14 to 23 and 326 to 331 by 1 to 10 conservative amino acid substitutions.

The present invention further provides an antibody or antigen-binding portion thereof, wherein said antibody or antigen-binding portion comprises: a V_H CDR1 as set forth in SEQ ID NO: 24, or a sequence that differs from SEQ ID NO: 24 by 1 to 4 conservative amino acid substitutions; a V_H CDR2 as set forth in SEQ ID NO: 25, or a sequence that differs from SEQ ID NO: 25 by 1 to 4 conservative amino acid substitutions; and a V_H CDR3 that is selected from any one of SEQ ID NOs: 26 to 37 and 91 to 256, or a sequence that differs from any one of SEQ ID NOs: 26 to 37 and 91 to 256 by 1 to 4 conservative amino acid substitutions.

The present invention further provides an antibody or antigen-binding portion thereof, wherein said antibody or antigen-binding portion comprises: a V_L CDR1 as set forth in SEQ ID NO: 38, or a sequence that differs from SEQ ID NO: 38 by 1 to 4 conservative amino acid substitutions; a V_L CDR2 as set forth in SEQ ID NO: 39, or a sequence that differs from SEQ ID NO: 39 by 1 to 4 conservative amino acid substitutions; and a V_L CDR3 that is selected from any one of SEQ ID NOs: 40 to 47 and 257 to 319, or a sequence that differs from any one of SEQ ID NOs: 40 to 47 and 257 to 319 by 1 to 4 conservative amino acid substitutions.

The present invention further provides an antibody or antigen-binding portion thereof with at least one of the functional properties described previously in A) thru K), wherein said antibody or antigen-binding portion comprises: a V_H FR1 as set forth in SEQ ID NO: 48; a V_H FR2 as set forth in SEQ ID NO: 49; a V_H FR3 selected from any one of SEQ ID NOs: 50 to 55; a V_H FR4 selected from any one of SEQ ID NOs: 56 and 57; a V_L FR1 selected from any one of SEQ ID NOs: 58 and 59; a V_L FR2 selected from any one of SEQ ID NOs: 60 to 62; a V_L FR3 selected from any one of SEQ ID NOs: 63 to 66; and a V_L FR4 as set forth in SEQ ID NO: 67. In a further embodiment, the V_H and V_L FR1, FR2, FR3, and FR4 sequences mentioned can also each independently differ from the specific SEQ ID NOs. recited above by at least one conservative amino acid substitution. For example, the FR1, FR2, FR3, and FR4 sequences can each independently differ by 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 conservative amino acid subsitions from the respective specific SEQ ID NOs recited above. In a still

further embodiment, any of the FR1, FR2, FR3, and FR4 sequences can each independently be mutated to match the respective germline framework sequence.

In a further embodiment of the present invention is any of the antibodies described herein wherein the antibody is an IgG, an IgM, an IgE, an IgA, or an IgD molecule, or is derived therefrom. For example, the antibody can be an IgG_1 or IgG_2. For example, in one embodiment IgG is an IgG_1 wherein the heavy chain constant region comprises SEQ ID NO: 344 and wherein the light chain constant region comprises SEQ ID NO: 345, with the proviso that the C-terminal lysine residue of SEQ ID NO: 344 is optionally cleaved.

Another embodiment provides any of the antibodies or antigen-binding portions described above which is an Fab fragment, an $F(ab')_2$ fragment, an F_V fragment, a single chain Fv fragment, a single chain V_H fragment, a single chain V_L fragment, a humanized antibody, a chimeric antibody or a bispecific antibody.

In a further embodiment is a derivatized antibody or antigen-binding portion comprising any of the antibodies or portions thereof as described herein and at least one additional molecular entity. For example, the at least one additional molecular entity can be another antibody (e.g., a bispecific antibody or a diabody), a detection agent, a label, a cytotoxic agent, a pharmaceutical agent, and/or a protein or peptide that can mediate association of the antibody or antibody portion with another molecule (such as a streptavidin core region or a polyhistidine tag). For example, useful detection agents with which an antibody or antigen-binding portion of the invention may be derivatized include fluorescent compounds, including fluorescein, fluorescein isothiocyanate, rhodamine, 5-dimethylamine-1-naphthalenesulfonyl chloride, phycoerythrin, lanthanide phosphors, and the like. An antibody can also be labeled with enzymes that are useful for detection, such as horseradish peroxidase, β-galactosidase, luciferase, alkaline phosphatase, glucose oxidase, and the like. In a further embodiment the antibodies or portions thereof of the present invention can also be labeled with biotin, or with a predetermined polypeptide epitope recognized by a secondary reporter (e.g., leucine zipper pair sequences, binding sites for secondary antibodies, metal binding domains, epitope tags). In a still further embodiment of the present invention, any of the antibodies or portions thereof can also be derivatized with a chemical group such as polyethylene glycol (PEG), a methyl or ethyl group, or a carbohydrate group.

In some embodiments, the P-cadherin antibodies or antigen binding portions disclosed herein are attached to a solid support.

In some embodiments, the C-terminal lysine of the heavy chain of any of the P-cadherin antibodies of the invention is cleaved. In various embodiments of the invention, the heavy and light chains of the P-cadherin antibodies may optionally include a N-terminal signal sequence. For example the heavy chain signal sequence can be SEQ ID NO: 346, and the light chain signal sequence can be SEQ ID NO: 347.

In a further embodiment, the present invention relates to any of the antibodies or antigen-binding portions thereof as described herein wherein the antibodies or antigen-bindingn portions thereof are of human origin.

The present invention also provides a pharmaceutical composition comprising any of the the antibodies or antigen-binding portions thereof as described above and a pharmaceutically acceptable carrier.

In another embodiment, the invention relates to an isolated nucleic acid molecule comprising a nucleotide sequence that encodes any of the antibodies or antigen binding portions thereof as described herein. In one particular embodiment is an isolated nucleic acid molecule comprising a nucleotide sequence as set forth in any one of SEQ ID NOs: 68 to 90 and 332 to 343. The invention further relates to a vector comprising any of the the nucleic acid molecules described herein, wherein the vector optionally comprises an expression control sequence operably linked to the nucleic acid molecule.

Another embodiment provides a host cell comprising any one of the vectors described herein or comprising any one of the nucleic acid molecules described herein. The present invention also provides an isolated cell line that produces any of the antibodies or antigen-binding portions as described herein or that produces the heavy chain or light chain of any of said antibodies or said antigen-binding portions.

In another embodiment, the present invention relates to a method for producing a P-cadherin antibody or antigen-binding portion thereof, comprising culturing any of the host cells or cell lines described herein under suitable conditions and recovering said antibody or antigen-binding portion.

The present invention also relates to a non-human transgenic animal or transgenic plant comprising any of the the nucleic acids described herein, wherein the non-human transgenic animal or transgenic plant expresses said nucleic acid.

The present invention further provides a method for isolating an antibody or antigen-binding portion thereof that binds to P-cadherin, comprising the step of isolating the antibody from the non-human transgenic animal or transgenic plant as described herein.

The present invention also provides a method for treating abnormal cell growth in a mammal in need thereof, comprising the step of administering to said mammal any of the antibodies or antigen-binding portions thereof, or any of the pharmaceutical compositions, as described herein. The present invention further provides a method for treating abnormal cell growth in a mammal in need thereof with an antibody or antigen-binding portion thereof that binds to P-cadherin comprising the steps of administering to said mammal an effective amount of any of the nucleic acid molecules described herein under suitable conditions that allow expression of said nucleic acid molecules. In another embodiment, the method of treating abnormal cell growth further comprises administering an amount of one or more substances selected from anti-tumor agents, anti-angiogenesis agents, signal transduction inhibitors, and antiproliferative agents, which amounts are together effective in treating said abnormal cell growth. In particular embodiments, said abnormal cell growth is cancerous.

The present invention further provides a method for reducing P-cadherin-dependent cellular aggregation comprising contacting the cells with any of the antibodies or antigen-binding portions thereof described herein or any of the the pharmaceutical compositions described herein.

Another aspect of the present invention is an antibody or antigen-binding portion thereof comprising a heavy chain variable region amino acid sequence that utilizes a human V_H-3 family gene. In one embodiment, for example, the human V_H-3 family gene is V_H-3-23.

Another aspect of the present invention provides any of the antibodies or antigen-binding portions thereof as described herein, where said antibody or antigen-binding portion is a human antibody. In a further aspect, said antibody or antigen-binding portion is a human recombinant antibody.

The invention also provides a method for producing a P-cadherin antibody or antigen-binding portion thereof comprising the steps of synthesizing a library of human antibodies on phage, screening the library with P-cadherin, or an antigenic portion thereof, isolating phage that bind P-cadherin, and obtaining the antibody from the phage.

Definitions and General Techniques

Unless otherwise defined herein, scientific and technical terms used in connection with the present invention shall have the meanings that are commonly understood by those of ordinary skill in the art. Further, unless otherwise required by context, singular terms shall include pluralities and plural terms shall include the singular. Generally, nomenclature used in connection with, and techniques of, cell and tissue culture, molecular biology, immunology, microbiology, genetics and protein and nucleic acid chemistry and hybridization described herein are those well known and commonly used in the art.

The methods and techniques of the present invention are 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 unless otherwise indicated. See, e.g., Sambrook, et al. Molecular Cloning: A Laboratory Manual, 3rd ed., Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y. (2000); Ausubel, et al., Short Protocols in Molecular Biology: A Compendium of Methods from Current Protocols in Molecular Biology, Wiley, John & Sons, Inc. (2002); Harlow and Lane Using Antibodies: A Laboratory Manual, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y. (1998); and Coligan, et al., Short Protocols in Protein Science, Wiley, John & Sons, Inc. (2003), the disclosures of which are incorporated herein by reference. Enzymatic reactions and purification techniques are performed according to manufacturer's specifications, as commonly accomplished in the art or as described herein. The nomenclature used 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 are used for chemical syntheses, chemical analyses, pharmaceutical preparation, formulation, delivery, and treatment of patients.

The basic antibody structural unit is known to comprise a tetramer. Each tetramer is composed of two identical pairs of polypeptide chains, each pair having one "light" (about 25 kDa) and one "heavy" chain (about 50-70 kDa). The amino-terminal portion of each chain includes a variable region of about 100 to 120 or more amino acids primarily responsible for antigen recognition.

The carboxy-terminal portion of each chain defines a constant region primarily responsible for effector function. Human light chains are classified as kappa and lambda light chains. Heavy chains are classified as mu, delta, gamma, alpha, or epsilon, and define the antibody's isotype as IgM, IgD, IgG, IgA, and IgE, respectively. Within light and heavy chains, 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 3 or more amino acids. See generally, Fundamental Immunology, Ch. 7 (Paul, W., ed., 2nd ed. Raven Press, N.Y. (1989)) (incorporated herein by reference in its entirety for all purposes). The variable regions of each heavy/light chain pair (V_H and V_L) form the antibody binding site. Thus, an intact IgG antibody, for example, has two binding sites. Except in bifunctional or bispecific antibodies, the two binding sites are the same.

The variable regions of the heavy and light chains 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 term "variable" refers to the fact that certain portions of the variable domains differ extensively in sequence among antibodies and are used in the binding and specificity of each particular antibody for its particular antigen. The variability, however, is not evenly distributed throughout the variable domains of antibodies, but is concentrated in the CDRs, which are separated by the more highly conserved FRs. The CDRs from the two chains of each pair are aligned by the FRs, enabling binding to a specific epitope. From N-terminal to C-terminal, both light and heavy chains comprise the domains FR1, CDR1, FR2, CDR2, FR3, CDR3 and FR4. The assignment of amino acids to each domain is 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), the disclosures of which are herein incorporated by reference.

The following terms, unless otherwise indicated, shall be understood to have the following meanings:

Unless specifically indicated otherwise, the term "P-cadherin" refers to human P-cadherin, which is an integral membrane protein and a member of the classical cadherin family of transmembrane glycoproteins that regulate cell-cell adhesion. The cloning and sequence of human P-cadherin has been reported, e.g. Shimoyama, et al., J. *Cell Biol.* 109 (4 Pt 1), 1787-1794 (1989), the disclosure of which is herein incorporated by reference. The term P-cadherin is intended to include recombinant human P-cadherin and recombinant chimeric forms of P-cadherin, which can be prepared by standard recombinant expression methods or purchased commercially (R&D Systems 861-PC-100).

Unless specifically indicated otherwise, as used herein the term "E-cadherin" refers to human E-cadherin, which is an integral membrane protein and a member of the classical cadherin family of transmembrane glycoproteins that regulate cell-cell adhesion. E-cadherin is described, for example, in Takeichi, *Science,* 251: 1451-1455 (1991), the disclosure of which is herein incorporated by reference. The term E-cadherin is intended to include recombinant human E-cadherin and

recombinant chimeric forms of E-cadherin, which can be prepared by standard recombinant expression methods or purchased commercially (R&D 648-EC-100).

The term "polypeptide" encompasses native or artificial proteins, protein fragments and polypeptide analogs of a protein sequence. A polypeptide may be monomeric or polymeric.

The term "isolated protein", "isolated polypeptide" or "isolated antibody" is a protein, polypeptide or antibody that by virtue of its origin or source of derivation (1) is not associated with naturally associated components that accompany it in its native state, (2) is free of other proteins from the same species, (3) is expressed by a cell from a different species, or (4) does not occur in nature. Thus, a polypeptide that is chemically synthesized or synthesized in a cellular system different from the cell from which it naturally originates will be "isolated" from its naturally associated components. A protein may also be rendered substantially free of naturally associated components by isolation, using protein purification techniques well known in the art.

Examples of isolated antibodies include a P-cadherin antibody that has been affinity purified using P-cadherin, and a P-cadherin antibody that has been synthesized by a cell line *in vitro*.

A protein or polypeptide is "substantially pure," "substantially homogeneous," or "substantially purified" when at least about 60 to 75% of a sample exhibits a single species of polypeptide. The polypeptide or protein may be monomeric or multimeric. A substantially pure polypeptide or protein can typically comprise about 50%, 60%, 70%, 80% or 90% w/w of a protein sample, more usually about 95%, and preferably can be over 99% pure. Protein purity or homogeneity may be indicated by a number of means well known in the art, such as polyacrylamide gel electrophoresis of a protein sample, followed by visualizing a single polypeptide band upon staining the gel with a stain well known in the art. For certain purposes, higher resolution may be provided by using HPLC or other means well known in the art for purification.

The term "polypeptide fragment" as used herein refers to a polypeptide that has an amino-terminal and/or carboxy-terminal deletion, but where the remaining amino acid sequence is identical to the corresponding positions in the naturally-occurring sequence. In some embodiments, fragments are at least 5, 6, 8 or 10 amino acids long. In other embodiments, the fragments are at least 14, at least 20, at least 50, or at least 70, 80, 90, 100, 150 or 200 amino acids long.

The term "analog" or "polypeptide analog" as used herein refers to a polypeptide that comprises a segment that has substantial identity to some reference amino acid sequence and has substantially the same function or activity as the reference amino acid sequence. Typically, polypeptide analogs comprise a conservative amino acid substitution (or insertion or deletion) with respect to the reference sequence. Analogs can be at least 20 or 25 amino acids long, or can be at least 50, 60, 70, 80, 90, 100, 150 or 200 amino acids long or longer, and can often be as long as the full-length polypeptide. Some embodiments of the invention include polypeptide fragments or polypeptide analog antibodies with 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16 or 17 substitutions from the germline amino acid sequence. Fragments or analogs of antibodies or immunoglobulin molecules can be readily prepared by those of ordinary skill in the art following the teachings of this specification.

In certain embodiments, amino acid substitutions to a P-cadherin antibody or antigen-binding portion thereof are those which: (1) reduce susceptibility to proteolysis, (2) reduce susceptibility to oxidation, (3) alter binding affinity for forming protein complexes, and (4) confer or modify other physicochemical or functional properties of such analogs, but still retain specific binding to P-cadherin. Analogs can include various substitutions to the normally-occurring peptide sequence. For example, single or multiple amino acid substitutions, preferably conservative amino acid substitutions, may be made in the normally-occurring sequence, for example in the portion of the polypeptide outside the domain(s) forming intermolecular contacts. Amino acid substitutions can also be made in the domain(s) that form intermolecular contacts that can improve the activity of the polypeptide. A conservative amino acid substitution should not substantially change the structural characteristics of the parent sequence; e.g., a replacement amino acid should not alter the anti-parallel β-sheet that makes up the immunoglobulin binding domain that occurs in the parent sequence, or disrupt other types of secondary structure that characterizes the parent sequence. In general, glycine and proline would not be used in an anti-parallel β-sheet. 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 and J. Tooze, eds., Garland Publishing, New York, N.Y. (1991)); and Thornton, et al., *Nature*, 354:105 (1991), incorporated herein by reference.

As used herein, the term "antibody" is synonymous with immunoglobulin and is to be understood as commonly known in the art. In particular, the term antibody is not limited by any particular method of producing the antibody. For example, the term antibody includes, without limitation, recombinant antibodies, monoclonal antibodies, and polyclonal antibodies.

The term "antigen-binding portion" of an antibody (or simply "antibody portion"), as used herein, refers to one or more fragments of an antibody that retain the ability to specifically bind to an antigen (e.g., P-cadherin). It has been shown that the antigen-binding function of an antibody can be performed by fragments of a full-length antibody. Examples of binding fragments encompassed within the term "antigen-binding portion" of an antibody include (i) a Fab fragment, a monovalent fragment consisting of the V_L, V_H, C_L and C_H1 domains; (ii) a $F(ab')_2$ fragment, a bivalent fragment comprising two Fab fragments linked by a disulfide bridge at the hinge region; (iii) a Fd fragment consisting of the V_H and C_H1 domains; (iv) a Fv fragment consisting of the V_L and V_H domains of a single arm of an antibody, (v) a dAb fragment (Ward, et al., *Nature*, (1989) 341:544-546), which consists of a V_H domain; and (vi) an isolated complementarity determining region (CDR). Furthermore, although the two domains of the Fv fragment, V_L and V_H, are coded for by separate genes, they can be joined, using recombinant methods, by a synthetic linker that enables them to be made as a single protein chain in which the V_L and V_H regions pair to form monovalent molecules (known as single chain Fv (scFv)); see e.g., Bird, et al., *Science* (1988) 242:423-426 and Huston, et al., *Proc. Natl. Acad. Sci. USA,* 85:5879-5883 (1988)). Such single chain antibodies are also intended to be encompassed within the term "antigen-binding portion" of an antibody. Other forms of single chain antibodies, such as diabodies are also encompassed. Diabodies are bivalent, bispecific

antibodies in which V_H and V_L domains are expressed on a single polypeptide chain, but using a linker that is too short to allow for pairing between the two domains on the same chain, thereby forcing the domains to pair with complementary domains of another chain and creating two antigen binding sites (see e.g., Holliger, et al., *Proc. Natl. Acad. Sci. USA,* 90:6444-6448 (1993); Poljak, et al. *Structure,* 2:1121-1123 (1994)).

Still further, an antibody or antigen-binding portion thereof may be part of larger immunoadhesion molecules, formed by covalent or noncovalent association of the antibody or antibody portion with one or more other proteins or peptides. Examples of such immunoadhesion molecules include use of the streptavidin core region to make a tetrameric scFv molecule (Kipriyanov, et al., *Human Antibodies and Hybridomas,* 6:93-101 (1995)) and use of a cysteine residue, a marker peptide and a C-terminal polyhistidine tag to make bivalent and biotinylated scFv molecules (Kipriyanov, et al., *Mol. Immunol.*, 31:1047-1058 (1994)). Other examples include where one or more CDRs from an antibody are incorporated into a molecule either covalently or noncovalently to make it an immunoadhesin that specifically binds to an antigen of interest, such as P-cadherin. In such embodiments, the CDR(s) may be incorporated as part of a larger polypeptide chain, may be covalently linked to another polypeptide chain, or may be incorporated noncovalently. Antibody portions, such as Fab and $F(ab')_2$ fragments, can be prepared from whole antibodies using conventional techniques, such as papain or pepsin digestion, respectively, of whole antibodies. Moreover, antibodies, antibody portions and immunoadhesion molecules can be obtained using standard recombinant DNA techniques, as described herein.

Where an "antibody" is referred to herein with respect to the present invention, it should be understood that an antigen-binding portion thereof may also be used. An antigen-binding portion competes with the intact antibody for specific binding. See generally, Fundamental Immunology, Ch. 7 (Paul, W., ed., 2nd ed. Raven Press, N.Y. (1989)) (incorporated by reference in its entirety for all purposes). Antigen-binding portions may be produced by recombinant DNA techniques or by enzymatic or chemical cleavage of intact antibodies. In some embodiments, antigen-binding portions include Fab, Fab', $F(ab')_2$, Fd, Fv, dAb, and complementarity determining region (CDR) fragments, single-chain antibodies (scFv), chimeric antibodies, diabodies and polypeptides that contain at least a portion of an antibody that is sufficient to confer specific antigen binding to the polypeptide. In embodiments having one or more binding sites, the binding sites may be identical to one another or may be different.

As used herein, the term "human antibody" means any antibody in which the variable and constant domain sequences are human sequences. The term encompasses antibodies with sequences derived from human genes, but which have been changed, e.g. to decrease possible immunogenicity, increase affinity, eliminate cysteines that might cause undesirable folding, etc. The term also encompasses such antibodies produced recombinantly in non-human cells, which might impart glycosylation not typical of human cells. These antibodies may be prepared in a variety of ways, as described below.

The term "chimeric antibody" as used herein means an antibody that comprises regions from two or more different antibodies, including antibodies from different species. For example, one or more of the CDRs of a chimeric antibody can be derived from a human P-cadherin antibody. In one example, the CDRs from a human antibody can be combined with CDRs from a non-human antibody, such as mouse or rat. In another example, all of the CDRs can be derived from human P-cadherin antibodies. In another example, the CDRs from more than one human P-cadherin antibody can be combined in a chimeric antibody. For instance, a chimeric antibody may comprise a CDR1 from the light chain of a first human P-cadherin antibody, a CDR2 from the light chain of a second human P-cadherin antibody and a CDR3 from the light chain of a third human P-cadherin antibody, and CDRs from the heavy chain may be derived from one or more other P-cadherin antibodies. Further, the framework regions may be derived from one of the P-cadherin antibodies from which one or more of the CDRs are taken or from one or more different human antibodies. Further, the term "chimeric antibody" is intended to encompass any of the above mentioned combinations where the combinations involved human and non-human antibodies.

As used herein, the term "humanized antibody" refers to antibodies of non-human origin, wherein the amino acid residues that are characteristic of antibody sequences of the non-human species are replaced with residues found in the corresponding positions of human antibodies. This "humanization" process is thought to reduce the immunogenicity in humans of the resulting antibody. It will be appreciated that antibodies of non-human origin can be humanized using techniques well known in the art. See, e.g. Winter, et al., *Immunol. Today*, 14:43-46 (1993). The antibody of interest may be engineered by recombinant DNA techniques to substitute the CH1, CH2, CH3, hinge domains, and/or the framework domain with the corresponding human sequence. See, e.g. WO 92/02190, and U.S. Patent Nos. 5,530,101, 5,585,089, 5,693,761, 5,693,792, 5,714,350, and 5,777,085). The term "humanized antibody", as used herein, includes within its meaning, chimeric human antibodies and CDR-grafted antibodies. Chimeric human antibodies of the invention include the V_H and V_L of an antibody of a non-human species and the C_H and C_L domains of a human antibody. The CDR-transplanted antibodies of the invention result from the replacement of CDRs of the V_H and V_L of a human antibody with those of the V_H and V_L, respectively, of an antibody of an animal other than a human.

As used herein, the term "ELISA" refers to an enzyme-linked immunosorbent assay. This assay is well known to those of skill in the art. Examples of this assay can be found in Vaughan, T.J., et al., *Nat. Biotech.*, 14:309-314 (1996), as well as in Example 7 of the present application.

The term "surface plasmon resonance", as used herein, refers to an optical phenomenon that allows for the analysis of real-time biospecific interactions by detection of alterations in protein concentrations within a biosensor matrix, for example using the BIACORE™ system (Pharmacia Biosensor AB, Uppsala, Sweden and Piscataway, N.J.). For further descriptions, see Jonsson et al., *Ann. Biol. Clin.*, 51:19-26 (1993); Jonsson, et al., *Biotechniques,* 11:620-627 (1991); Jonsson, et al., *J. Mol. Recognit.*, 8:125-131 (1995); and Johnsson, et al., *Anal. Biochem.*, 198:268-277 (1991).

The term "K_D" refers to the binding affinity equilibrium constant of a particular antibody-antigen interaction. An antibody is said to specifically bind an antigen when the K_D is ≤ 1 mM, preferably ≤ 100 nM and most preferably ≤ 10 nM. A K_D binding affinity constant can be measured by surface plasmon resonance, for example using the BIACORE™ system as discussed in Example 6.

The term "k_{off}" refers to the dissociation rate constant of a particular antibody-antigen interaction. A k_{off} dissociation rate constant can be measured by surface plasmon resonance, for example using the BIACORE™ system as discussed in Example 6.

As used herein, the term "P-cadherin dependent cell adhesion assay" refers to an assay used to measure the ability of a P-cadherin antibody to block the adhesion of cells to a receptor P-cadherin that has been immobilized on a solid support. This type of assay can be carried out, for example, by immobilizing P-cadherin on a solid support, such as plastic. Cells over-expressing P-cadherin are then allowed to adhere to the solid support via P-cadherin-P-cadherin interactions. The level of adhesion can then be quantified with and without a P-cadherin antibody. Adhesion as a function of antibody concentration can then be used to determine an IC_{50} value. Example 3 provides further details of a P-cadherin-dependent cell adhesion assay that was used to measure IC_{50} values for P-cadherin antibodies.

As used herein, the term "P-cadherin dependent cell aggregation assay" refers to an assay for measuring the ability of a P-cadherin antibody to block aggregation of cells expressing P-cadherin on their surfaces. For example, this type of assay can use a cell line that over-expresses P-cadherin, wherein the cells are placed into suspension and allowed to form P-cadherin-dependent aggregates. The aggregation assay is then used to quantify the ability of a P-cadherin antibody to prevent this aggregation by measuring the size of cellular aggregates that result with and without the antibody. Cell aggregate size as a function of P-cadherin antibody concentration can then be used to determine an IC_{50} value. Example 4 provides further details of a P-cadherin-dependent aggregation assay that was used to measure IC_{50} values for several P-cadherin antibodies.

As used herein, the term "P-cadherin-dependent spheroid disruption assay" refers to an assay for measuring the ability of a P-cadherin antibody to disrupt pre-formed P-cadherin-dependent cellular aggregations. By measuring the size reduction of aggregates as a function of antibody concentration, an IC_{50} value can be determined. Example 5 provides further details of a P-cadherin-dependent spheroid disruption assay that was used to measure IC_{50} values for P-cadherin antibodies.

As used herein, the term "molecular selectivity" refers to the binding affinity of an antibody for a specific antigen being greater than for a related antigen. For example, the antibodies of the present invention can be selective for P-cadherin over E-cadherin, meaning that the binding affinity of the antibody for P-cadherin is at least 2 times greater, for example 4 times, or 10 times, or 50 times, or 100 times or more, than for E-cadherin. Such binding affinities can be measured using standard techniques known to those of skill in the art.

The term "epitope" includes any protein determinant capable of specific binding to an immunoglobulin or T-cell receptor or otherwise interacting with a molecule. Epitopic determinants generally consist of chemically active surface groupings of molecules such as amino acids or carbohydrate or sugar side chains and generally have specific three dimensional structural characteristics, as well as specific charge characteristics. An epitope may be "linear" or "conformational." In a linear epitope, all of the points of interaction between the protein and the interacting molecule (such as an antibody) occur linearally along the primary amino acid sequence of the protein. In a conformational epitope, the points of interaction occur across amino acid residues on the protein that are separated from one another. Once a desired epitope on an antigen is determined, it is possible to generate antibodies to that epitope, e.g., using the techniques described in the present invention. Alternatively, during the discovery process, the generation and characterization of antibodies may elucidate information about desirable epitopes. From this information, it is then possible to competitively screen antibodies for binding to the same epitope. An approach to achieve this is to conduct cross-competition studies to find antibodies that competitively bind with one another, i.e. the antibodies compete for binding to the antigen. A high throughput process for "binning" antibodies based upon their cross-competition is described in International Patent Publication No. WO 03/48731.

The term "compete", as used herein with regard to an antibody, means that a first antibody, or an antigen-binding portion thereof, competes for binding with a second antibody, or an antigen-binding portion thereof, where binding of the first antibody with its cognate epitope is detectably decreased in the presence of the second antibody compared to the binding of the first antibody in the absence of the second antibody. The alternative, where the binding of the second antibody to its epitope is also detectably decreased in the presence of the first antibody, can, but need not be the case. That is, a first antibody can inhibit the binding of a second antibody to its epitope without that second antibody inhibiting the binding of the first antibody to its respective epitope. However, where each antibody detectably inhibits the binding of the other antibody with its cognate epitope or ligand, whether to the same, greater, or lesser extent, the antibodies are said to "cross-compete" with each other for binding of their respective epitope(s). Both competing and cross-competing antibodies are encompassed by the present invention. Regardless of the mechanism by which such competition or cross-competition occurs (*e.g.*, steric hindrance, conformational change, or binding to a common epitope, or portion thereof, and the like), the skilled artisan would appreciate, based upon the teachings provided herein, that such competing and/or cross-competing antibodies are encompassed and can be useful for the methods disclosed herein.

As used herein, the term "utilizes" with reference to a particular gene means that the amino acid sequence of a particular region in an antibody was ultimately derived from that gene during B-cell maturation. For example, the phrase "a heavy chain variable region amino acid sequence that utilizes a human V_H-3 family gene" refers to the situation where the V_H region of the antibody was derived from the VH-3 family of gene segments during B-cell maturation. In human B-cells, there are more than 30 distinct functional heavy chain variable genes with which to generate antibodies. Use

of a particular heavy chain variable gene, therefore, is indicative of a preferred binding motif of the antibody-antigen interaction with respect to the combined properties of binding to the antigen and functional activity. As will be appreciated, gene utilization analysis provides only a limited overview of antibody structure. As human B-cells stocastically generate V-D-J heavy or V-J kappa light chain transcripts, there are a number of secondary processes that occur, including, without limitation, somatic hypermutation, n-additions, and CDR3 extensions. See, for example, Mendez et al. *Nature Genetics* 15:146-156 (1997).

As used herein, the twenty conventional 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)), incorporated herein by reference.

The term "polynucleotide" as referred to herein means a polymeric form of nucleotides of at least 10 bases in length, either ribonucleotides or deoxynucleotides or a modified form of either type of nucleotide. The term includes single and double stranded forms.

The term "isolated polynucleotide" as used herein means a polynucleotide of genomic, cDNA, or synthetic origin or some combination thereof, which by virtue of its origin the "isolated polynucleotide" (1) is not associated with all or a portion of polynucleotides with which the "isolated polynucleotide" is found in nature, (2) is operably linked to a polynucleotide to which it is not linked in nature, or (3) does not occur in nature as part of a larger sequence.

The term "naturally occurring nucleotides" as used herein includes deoxyribonucleotides and ribonucleotides. The term "modified nucleotides" as used herein includes nucleotides with modified or substituted sugar groups and the like. The term "oligonucleotide linkages" referred to herein includes oligonucleotides linkages such as phosphorothioate, phosphorodithioate, phosphoroselenoate, phosphorodiselenoate, phosphoroanilothioate, phoshoraniladate, phosphoroamidate, and the like. See e.g., LaPlanche, et al., *Nucl. Acids Res.*, 14:9081 (1986); Stec, et al., *J. Am. Chem. Soc.*, 106:6077 (1984); Stein, et al., *Nucl. Acids Res.*,16:3209 (1988); Zon, et al., *Anti-Cancer Drug Design,* 6:539 (1991); Zon, et al., Oligonucleotides and Analogues: A Practical Approach, pp. 87-108 (F. Eckstein, Ed., Oxford University Press, Oxford England (1991)); U.S. Patent No. 5,151,510; Uhlmann and Peyman, *Chemical Reviews,* 90:543 (1990), the disclosures of which are hereby incorporated by reference. An oligonucleotide can include a label for detection, if desired.

"Operably linked" sequences include both expression control sequences that are contiguous with the gene of interest and expression control sequences that act in *trans* or at a distance to control the gene of interest.

The term "expression control sequence" as used herein means polynucleotide sequences that are necessary to effect the expression and processing of coding sequences to which they are ligated. Expression control sequences include appropriate transcription initiation, termination, promoter and enhancer sequences; efficient RNA processing signals such as splicing and polyadenylation signals; sequences that stabilize cytoplasmic mRNA; sequences that enhance translation efficiency (i.e., Kozak consensus sequence); sequences that enhance protein stability;

and when desired, sequences that enhance protein secretion. The nature of such control sequences differs depending upon the host organism; in prokaryotes, such control sequences generally include promoter, ribosomal binding site, and transcription termination sequence; in eukaryotes, generally, such control sequences include promoters and transcription termination sequence. The term "control sequences" is intended to include, at a minimum, all components whose presence is essential for expression and processing, and can also include additional components whose presence is advantageous, for example, leader sequences and fusion partner sequences.

The term "vector", as used herein, means a nucleic acid molecule capable of transporting another nucleic acid to which it has been linked. In some embodiments, the vector is a plasmid, i.e., a circular double stranded piece of DNA into which additional DNA segments may be ligated. In some embodiments, the vector is a viral vector, wherein additional DNA segments may be ligated into the viral genome. In some embodiments, the vectors are capable of autonomous replication in a host cell into which they are introduced (e.g., bacterial vectors having a bacterial origin of replication and episomal mammalian vectors). In other embodiments, the vectors (e.g., non-episomal mammalian vectors) can be integrated into the genome of a host cell upon introduction into the host cell, and thereby are replicated along with the host genome. Moreover, certain vectors are capable of directing the expression of genes to which they are operatively linked. Such vectors are referred to herein as "recombinant expression vectors" (or simply, "expression vectors").

The term "recombinant host cell" (or simply "host cell"), as used herein, means a cell into which a recombinant expression vector has been introduced. It should be understood that "recombinant host cell" and "host cell" mean not only the particular subject cell but also the progeny of such a cell. Because certain modifications may occur in succeeding generations due to either mutation or environmental influences, such progeny may not, in fact, be identical to the parent cell, but are still included within the scope of the term "host cell" as used herein.

As used herein, the term "germline" refers to the nucleotide sequences of the antibody genes and gene segments as they are passed from parents to offspring via the germ cells. This germline sequence is distinguished from the nucleotide sequences encoding antibodies in mature B cells which have been altered by recombination and hypermutation events during the course of B cell maturation.

The term "percent sequence identity" in the context of nucleic acid sequences means the residues in two sequences that are the same when aligned for maximum correspondence. The length of sequence identity comparison may be over a stretch of at least about nine nucleotides, usually at least about 18 nucleotides, more usually at least about 24 nucleotides, typically at least about 28 nucleotides, more typically at least about 32 nucleotides, and preferably at least about 36, 48 or more nucleotides. There are a number of different algorithms known in the art which can be used to measure nucleotide sequence identity. For instance, polynucleotide sequences can be compared using FASTA, Gap or Bestfit, which are programs in Wisconsin Package Version 10.0, Genetics Computer Group (GCG), Madison, Wisconsin. FASTA, which includes, e.g., the programs FASTA2 and FASTA3, provides alignments and percent sequence identity of the regions of the best

overlap between the query and search sequences (Pearson, *Methods Enzymol.*, 183:63-98 (1990); Pearson, *Methods Mol. Biol.*, 132:185-219 (2000); Pearson, *Methods Enzymol.*, 266:227-258 (1996); Pearson, *J. Mol. Biol.* 276:71-84 (1998); incorporated herein by reference). Unless otherwise specified, default parameters for a particular program or algorithm are used. For instance, percent sequence identity between nucleic acid sequences can be determined using FASTA with its default parameters (a word size of 6 and the NOPAM factor for the scoring matrix) or using Gap with its default parameters as provided in GCG Version 6.1, incorporated herein by reference.

A reference to a nucleotide sequence encompasses its complement unless otherwise specified. Thus, a reference to a nucleic acid having a particular sequence should be understood to encompass its complementary strand, with its complementary sequence.

The term "substantial similarity" or "substantial sequence similarity," when referring to a nucleic acid or fragment thereof, means that when optimally aligned with appropriate nucleotide insertions or deletions with another nucleic acid (or its complementary strand), there is nucleotide sequence identity in at least about 85%, preferably at least about 90%, and more preferably at least about 95%, 96%, 97%, 98% or 99% of the nucleotide bases, as measured by any well-known algorithm of sequence identity, such as FASTA, BLAST or Gap, as discussed above.

The term "percent sequence identity" in the context of amino acid sequences means the residues in two sequences that are the same when aligned for maximum correspondence. The length of sequence identity comparison may be over a stretch of at least about five amino acids, usually at least about 20 amino acids, more usually at least about 30 amino acids, typically at least about 50 amino acids, more typically at least about 100 amino acids, and even more typically about 150, 200 or more amino acids. There are a number of different algorithms known in the art that can be used to measure amino acid sequence identity. For instance, amino acid sequences can be compared using FASTA, Gap or Bestfit, which are programs in Wisconsin Package Version 10.0, Genetics Computer Group (GCG), Madison, Wisconsin.

As applied to polypeptides, the term "substantial identity" or "substantial similarity" means that two amino acid sequences, when optimally aligned, such as by the programs GAP or BESTFIT using default gap weights as supplied with the programs, share at least 70%, 75% or 80% sequence identity, preferably at least 90% or 95% sequence identity, and more preferably at least 97%, 98% or 99% sequence identity. In certain embodiments, residue positions that are not identical differ by conservative amino acid substitutions.

As the ther is used herein, a "conservative amino acid substitution" is one in which an amino acid residue is substituted by another amino acid residue having a side chain R group with similar chemical properties (e.g., charge or hydrophobicity). In general, a conservative amino acid substitution will not substantially change the functional properties of a protein. In cases where two or more amino acid sequences differ from each other by conservative substitutions, the percent sequence identity may be adjusted upwards to correct for the conservative nature of the substitution. Means for making this adjustment are well-known to those of skill in the art. See, e.g., Pearson, *Methods Mol. Biol.*, 243:307-31 (1994). Examples of groups of amino acids that have side chains

with similar chemical properties include 1) aliphatic side chains: glycine, alanine, valine, leucine, and isoleucine; 2) aliphatic-hydroxyl side chains: serine and threonine; 3) amide-containing side chains: asparagine and glutamine; 4) aromatic side chains: phenylalanine, tyrosine, and tryptophan; 5) basic side chains: lysine, arginine, and histidine; 6) acidic side chains: aspartic acid and glutamic acid; and 7) sulfur-containing side chains: cysteine and methionine. For example, conservative amino acids substitution groups can be: valine-leucine-isoleucine, phenylalanine-tyrosine, lysine-arginine, alanine-valine, glutamate-aspartate, and asparagine-glutamine.

Alternatively, a conservative replacement is any change having a positive value in the PAM250 log-likelihood matrix disclosed in Gonnet et al., *Science* 256:1443-45 (1992), incorporated herein by reference. A "moderately conservative" replacement is any change having a nonnegative value in the PAM250 log-likelihood matrix.

Sequence identity for polypeptides is typically measured using sequence analysis software. Protein analysis software matches sequences using measures of similarity assigned to various substitutions, deletions and other modifications, including conservative amino acid substitutions. For instance, GCG contains programs such as "Gap" and "Bestfit" which can be used with default parameters as specified by the programs to determine sequence homology or sequence identity between closely related polypeptides, such as homologous polypeptides from different species of organisms or between a wild type protein and an analog thereof. See, e.g., GCG Version 6.1 (University of Wisconsin, WI). Polypeptide sequences also can be compared using FASTA using default or recommended parameters, see GCG Version 6.1. FASTA (e.g., FASTA2 and FASTA3) provides alignments and percent sequence identity of the regions of the best overlap between the query and search sequences (Pearson, *Methods Enzymol.,* 183:63-98 (1990); Pearson, *Methods Mol. Biol.,* 132:185-219 (2000)). Another preferred algorithm when comparing a sequence of the invention to a database containing a large number of sequences from different organisms is the computer program BLAST, especially blastp or tblastn, using default parameters as supplied with the programs. See, e.g., Altschul, et al., *J. Mol. Biol.,* 215:403-410 (1990); Altschul, et al., *Nucleic Acids Res.,* 25:3389-402 (1997).

The length of polypeptide sequences compared for homology will generally be at least about 16 amino acid residues, usually at least about 20 residues, more usually at least about 24 residues, typically at least about 28 residues, and preferably more than about 35 residues. When searching a database containing sequences from a large number of different organisms, it is preferable to compare amino acid sequences.

As used herein, the terms "label" or "labeled" refers to incorporation of another molecule in the antibody. In one embodiment, the label is a detectable marker, e.g., incorporation of a radiolabeled amino acid or attachment to a polypeptide of biotinyl 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 another embodiment, the label or marker can be therapeutic, e.g., a drug conjugate or toxin. Various methods of labeling polypeptides and glycoproteins are known in the art and may 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 markers, 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), magnetic agents, such as gadolinium chelates, toxins such as pertussis toxin, taxol, cytochalasin B, gramicidin D, ethidium bromide, emetine, mitomycin, etoposide, tenoposide, vincristine, vinblastine, colchicine, doxorubicin, daunorubicin, dihydroxy anthracin dione, mitoxantrone, mithramycin, actinomycin D, 1-dehydrotestosterone, glucocorticoids, procaine, tetracaine, lidocaine, propranolol, and puromycin and analogs or homologs thereof. In some embodiments, labels are attached by spacer arms of various lengths to reduce potential steric hindrance.

"Therapeutically effective amount" refers to that amount of the therapeutic agent being administered which will relieve to some extent one or more of the symptoms of the disorder being treated. In reference to the treatment of cancer, a therapeutically effective amount refers to that amount which has at least one of the following effects: reducing the size of the tumor; inhibiting (that is, slowing to some extent, preferably stopping) tumor metastasis; inhibiting to some extent (that is, slowing to some extent, preferably stopping) tumor growth, and relieving to some extent (or, preferably, eliminating) one or more symptoms associated with the cancer.

"Treat", "treating" and "treatment" refer to a method of alleviating or abrogating a biological disorder and/or its attendant symptoms. With regard to cancer, these terms simply mean that the life expectancy of an individual affected with a cancer will be increased or that one or more of the symptoms of the disease will be reduced.

"Contacting" refers to bringing an antibody or antigen binding portion thereof of the present invention and a target P-cadherin, or epitiope thereof, together in such a manner that the antibody can affect the biological activity of the P-cadherin. Such "contacting" can be accomplished "*in vitro*," i.e., in a test tube, a petri dish, or the like. In a test tube, contacting may involve only an antibody or antigen binding portion thereof and P-cadherin or epitope thereof or it may involve whole cells. Cells may also be maintained or grown in cell culture dishes and contacted with antibodies or antigen binding portions thereof in that environment. In this context, the ability of a particular antibody or antigen binding portion thereof to affect a P-cadherin-related disorder, i.e., the IC_{50} of the antibody, can be determined before use of the antibody *in vivo* with more complex living organisms is attempted. For cells outside the organism, multiple methods exist, and are well-known to those skilled in the art, to contact P-caderin with the antibodies or antigen-binding portions thereof.

"Abnormal cell growth", as used herein, unless otherwise indicated, refers to cell growth that is independent of normal regulatory mechanisms (e.g., loss of contact inhibition), including the abnormal growth of normal cells and the growth of abnormal cells. This includes, but is not limited to, the abnormal growth of: tumor cells (tumors) that proliferate by expressing a mutated tyrosine kinase or overexpression of a receptor tyrosine kinase; benign and malignant cells of other

proliferative diseases in which aberrant tyrosine kinase activation occurs; any tumors that proliferate by receptor tyrosine kinases; any tumors that proliferate by aberrant serine/threonine kinase activation; benign and malignant cells of other proliferative diseases in which aberrant serine/threonine kinase activation occurs; tumors, both benign and malignant, expressing an activated Ras oncogene; tumor cells, both benign and malignant, in which the Ras protein is activated as a result of oncogenic mutation in another gene; benign and malignant cells of other proliferative diseases in which aberrant Ras activation occurs. Examples of such benign proliferative diseases are psoriasis, benign prostatic hypertrophy, human papilloma virus (HPV), and restinosis. "Abnormal cell growth" also refers to and includes the abnormal growth of cells, both benign and malignant, resulting from activity of the enzyme farnesyl protein transferase.

The terms "abnormal cell growth" and "hyperproliferative disorder" are used interchangeably in this application.

"*In vitro*" refers to procedures performed in an artificial environment such as, e.g., without limitation, in a test tube or culture medium.

"*In vivo*" refers to procedures performed within a living organism such as, without limitation, a mouse, rat or rabbit.

Brief Description Of The Drawings

Figure 1 shows the amino acid and nucleic acid sequences of SEQ ID NOs: 1-347.

Detailed Description Of The Invention

Human P-cadherin Antibodies

This invention pertains to isolated human antibodies, or antigen-binding portions thereof, that bind to human P-cadherin. Preferably, the human antibodies are recombinant human P-cadherin antibodies that have greater affinity for P-cadherin than for E-cadherin. Various aspects of the invention relate to such antibodies and antigen-binding portions, and pharmaceutical compositions thereof, as well as nucleic acids, recombinant expression vectors and host cells for making such antibodies and antigen-binding portions. Methods of using the antibodies and antigen-binding portions of the present invention to detect human P-cadherin or to inhibit human P-cadherin activity, either *in vitro* or *in vivo*, are also encompassed by the invention.

The P-cadherin amino acid and nucleotide sequences from several species, including human, are known (see e.g. Accession No. NM_001793.3). Human P-cadherin, or antigenic portions thereof, can be prepared according to methods well known to those in the art, or can be purchased from commercial vendors (e.g. R&D Systems 861-PC-100).

In certain embodiments, antibodies of the present invention are IgGs designated as: 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4. As discussed in greater detail in Example 1, high throughput screening of a scFv phage display library was used to identify the 129-1c4 scFv, which was subsequently converted to an IgG. 129-1c4 represents the lead antibody identified during initial phage display screening and is the parent antibody of a lineage from which several other antibodies of the present invention were

derived. Several of such derived antibodies are designated as 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; and 200-h06 and represent optimized antibodies within the 129-1c4 parent lineage. The g-129-1c4 antibody is the germline version of the 129-1c4 parent antibody where certain amino acids in the framework regions of the V_H and V_L domains were mutated to match those in the germline framework regions. Any of the antibodies mentioned above can also be germlined so that the framework region sequences are identical to the germline framework sequences as in g-129-1c4. For example in one embodiment of the present invention, the antibodies g-194-b09, g-194-g09, g-196-g03, g-196-h02, g-194-e01, g-195-e11, g-200-h06, and g-194-e06 are the germlined versions of 194-b09, 194-g09, 196-g03, 196-h02, 194-e01, 195-e11, 200-h06, and 194-e06, respectively. The specific amino acids that were mutated to arrive at the germlined versions are apparent to those of skill in the art by comparing the sequences of a germlined vs. a non-germlined antibody. As discussed below, specific amino acid sequences of the antibodies of the present invention are described in Tables 1-3 and Figure 1.

Antibodies of the present invention were generated with a strong bias towards the utilization of the V_H3 gene family of heavy chain variable regions. In particular, the 129-1c4 parent antibody derived from the V_H3-23 variable gene segment. In human B-cells, there are more than 30 distinct functional heavy chain variable genes with which to generate antibodies. Bias, therefore, is indicative of a preferred binding motif of the antibody-antigen interaction with respect to the combined properties of binding to the antigen and functional activity.

As will be appreciated, gene utilization analysis provides only a limited overview of antibody structure. As human B-cells stocastically generate V-D-J heavy or V-J kappa light chain transcripts, there are a number of secondary processes that occur, including, without limitation, somatic hypermutation, n-additions, and CDR3 extensions. See, for example, Mendez et al. *Nature Genetics* 15:146-156 (1997) and U.S. published patent application 2003-0070185, filed Feb. 19, 2002. Accordingly, to further examine antibody structure of the present invention, predicted amino acid sequences of the antibodies were generated from the cDNAs obtained from the clones. In addition, N-terminal amino acid sequences were obtained through protein sequencing. Figure 1 provides nucleotide and amino acid sequences of the heavy and light chain variable regions of several of the antibodies of the present invention.

Each of the specific antibodies mentioned above can be described by their variable domain sequences of the heavy (V_H) and light (V_L) chains as indicated in Tables 1 and 2. The specific sequences referred to by these SEQ ID NOs. are shown in Figure 1. As indicated in Tables 1 and 2, the corresponding V_H and V_L amino acid and DNA sequences of the antibodies mentioned above are described by SEQ ID NOs: 1-23, 68-90, and 320-343.

Table 1

Human P-cadherin antibodies				
Antibody	Sequence Identifier (SEQ ID NO:)			
	V_H		V_L	
	Amino acid	DNA	Amino acid	DNA
194-e06	1	68	14	81
194-a02	2	69	14	81
194-b09	2	69	15	82
195-e11	3	70	16	83
194-g09	4	71	17	84
196-h02	4	71	23	90
194-e01	5	72	18	85
196-d10	6	73	23	90
196-g03	7	74	23	90
196-e06	8	75	23	90
195-a09	9	76	23	90
198-a09	10	77	19	86
200-h06	11	78	20	87
129-1c4	12	79	21	88

Table 2

Germlined Human P-cadherin antibodies				
Antibody	Sequence Identifier (SEQ ID NO:)			
	V_H		V_L	
	Amino acid	DNA	Amino acid	DNA
g-194-b09	320	332	326	338
g-194-g09	321	333	327	339
g-196-g03	322	334	328	340
g-196-h02	323	335	329	341
g-194-e01	324	336	330	342
g-194-e06	325	337	331	343
g-129-1c4	13	80	22	89

Additional antibodies and antigen-binding portions of the present invention can also be described as comprising the various CDR and FR sequences that make up the heavy and light chain variable regions of the antibodies indicated in Tables 1 and 2. Accordingly, SEQ ID NOs. that correspond to various CDR and FR sequences of antibodies of the present invention are indicated in Table 3. Furthermore, numerous randomized mutations in the heavy and light chain CDR3 regions of the 129-1c4 parent antibody were also performed, which yielded improved P-cadherin affinity ranging from 10- to 417-fold improvement as measured by an epitope competition assay (see Example 8). The SEQ ID NOs. of these mutated V_H and V_L CDR3 sequences (SEQ ID NOs: 91-256, and 257-319) are also included in Table 3 below.

Table 3

SEQ ID NOs:	Description
24	V_H CDR1
25	V_H CDR2
26-37, 91-256	V_H CDR3
38	V_L CDR1
39	V_L CDR2
40-47, 257-319	V_L CDR3
48	V_H FR1
49	V_H FR2
50-55	V_H FR3
56, 57	V_H FR4
58, 59	V_L FR1
60-62	V_L FR2
63-66	V_L FR3
67	V_L FR4

Methods of Producing Antibodies

Phage Display Libraries

The antibodies or antigen-binding portions of the present invention can be prepared according to several methods known in the art. For example, phage display techniques can be used to provide libraries containing a repertoire of antibodies with varying affinities for P-cadherin. These libraries can then be screened to identify and isolate antibodies with the desired affinity for P-cadherin.

For example, recombinant human P-cadherin antibodies of the present invention can be isolated by screening a recombinant combinatorial antibody library. Preferably the library is a scFv phage display library, generated using human V_L and V_H cDNAs prepared from mRNA isolated from human B cells. Methods for preparing and screening such libraries are known in the art. Kits for generating phage display libraries are commercially available (e.g., the Pharmacia Recombinant Phage Antibody System, catalog no. 27-9400-01; and the Stratagene SurfZAP™ phage display kit, catalog no. 240612). There are also other methods and reagents that can be used in generating and screening antibody display libraries (see, e.g., U.S. Patent No. 5,223,409; PCT Publication Nos. WO 92/18619, WO 91/17271, WO 92/20791, WO 92/15679, WO 93/01288, WO 92/01047, and WO 92/09690; Fuchs et al., *Bio/Technology* 9:1370-1372 (1991); Hay, et al., *Hum. Antibod. Hybridomas*, 3:81-85 (1992); Huse, et al., *Science*, 246:1275-1281 (1989); McCafferty et al., *Nature*, 348:552-554 (1990); Griffiths, et al., *EMBO J.*, 12:725-734 (1993); Hawkins, et al., *J. Mol. Biol.*, 226:889-896 (1992); Clackson, et al., *Nature* 352:624-628 (1991); Gram, et al., *Proc. Natl. Acad. Sci. USA*, 89:3576-3580 (1992); Garrad, et al., *Bio/Technology*, 9:1373-1377 (1991); Hoogenboom, et al., *Nuc. Acid Res.*, 19:4133-4137 (1991); Barbas, et al., *Proc. Natl. Acad. Sci. USA*, 88:7978-7982 (1991); and Griffiths, et al., *EMBO J.*, 13:3245-3260 (1994); which are all incorporated herein by reference).

Another method for preparing a library of antibodies for use in phage display techniques comprises the steps of immunizing a non-human animal comprising human immunoglobulin loci with P-cadherin or an antigenic portion thereof to create an immune response, extracting antibody-producing cells from the immunized animal; isolating RNA encoding heavy and light chains of

antibodies of the invention from the extracted cells, reverse transcribing the RNA to produce cDNA, amplifying the cDNA using primers, and inserting the cDNA into a phage display vector such that antibodies are expressed on the phage. For production of such repertoires, it is unnecessary to immortalize the B cells from the immunized animal. Rather, the primary B cells can be used directly as a source of DNA. The mixture of cDNAs obtained from B cells, e.g., derived from spleens, is used to prepare an expression library, for example, a phage display library transfected into *E.coli*. Ultimately, clones from the library are identified that produce binding affinities of a desired magnitude for the antigen and the DNA encoding the product responsible for such binding is recovered and manipulated for standard recombinant expression. Phage display libraries may also be constructed using previously manipulated nucleotide sequences and screened in a similar fashion. In general, the cDNAs encoding heavy and light chains are independently supplied or linked to form Fv analogs for production in the phage library. The phage library is then screened for the antibodies with the highest affinities for P-cadherin and the genetic material is recovered from the appropriate clone. Further rounds of screening can increase affinity of the original antibody isolated.

In one embodiment, to isolate and produce human P-cadherin antibodies with the desired characteristics, a human P-cadherin antibody as described herein is first used to select human heavy and light chain sequences having similar binding activity toward P-cadherin, using the epitope imprinting methods described in PCT Publication No. WO 93/06213, incorporated herein by reference. The antibody libraries used in this method are preferably scFv libraries prepared and screened as described in PCT Publication No. WO 92/01047, McCafferty, et al., *Nature,* 348:552-554 (1990); and Griffiths, et al., *EMBO J.* 12:725-734 (1993), all incorporated herein by reference. The scFv antibody libraries can be screened using human P-cadherin as the antigen. The phage library is screened for the antibodies with the highest affinities for P-cadherin and the genetic material recovered from the appropriate clone. Further rounds of screening can increase affinity of the original antibody isolated.

Once initial human V_L and V_H domains are selected, "mix and match" experiments can then be performed, in which different pairs of the initially selected V_L and V_H segments are screened for P-cadherin binding to select preferred V_L/V_H pair combinations. These mix and match experiments can also be performed after the V_H and V_L segments have been randomly mutated for optimized binding as described below. Additionally, to further improve the quality of the antibody, the V_L and V_H segments of the preferred V_L/V_H pair(s) can be randomly mutated, preferably within the CDR3 region of V_H and/or V_L, in a process analogous to the *in vivo* somatic mutation process responsible for affinity maturation of antibodies during a natural immune response. This *in vitro* affinity maturation can be accomplished, for example, by amplifying V_H and V_L domains using PCR primers complimentary to the V_H CDR3 or V_L CDR3, respectively, which primers have been "spiked" with a random mixture of the four nucleotide bases at certain positions such that the resultant PCR products encode V_H and V_L segments into which random mutations have been introduced into the V_H and/or V_L CDR3 regions. These randomly mutated V_H and V_L segments can be re-screened for binding to P-cadherin, and sequences that exhibit high affinity and a low off rate for P-cadherin can

be selected. As discussed previously, several V_H and V_L CDR3 sequences of the present invention that were randomly mutated and showed improved affinity are indicated by SEQ ID NOs: 91-256 and 257-319.

Following screening and isolation of a P-cadherin antibody of the invention from a recombinant immunoglobulin display library, nucleic acids encoding the selected antibody can be recovered from the display package (e.g., from the phage genome) and subcloned into other expression vectors by standard recombinant DNA techniques. If desired, the nucleic acid can further be manipulated to create other antibody forms of the invention, as described below. To express a recombinant human antibody isolated by screening of a combinatorial library, the DNA encoding the antibody is cloned into a recombinant expression vector and introduced into a host cell, as described below.

Immunization

In another embodiment, human P-cadherin antibodies can be produced by immunizing a non-human, transgenic animal comprising within its genome some or all of human immunoglobulin heavy chain and light chain loci with a P-cadherin antigen. For example, the non-human animal can be a XENOMOUSE™ animal. (Abgenix, Inc., Fremont, CA).

XENOMOUSE™ mice are engineered mouse strains that comprise large fragments of human immunoglobulin heavy chain and light chain loci and are deficient in mouse antibody production. See, e.g., Green, et al., *Nature Genetics*, 7:13-21 (1994) and U.S. Patent Nos. 5,916,771, 5,939,598, 5,985,615, 5,998,209, 6,075,181, 6,091,001, 6,114,598, 6,130,364, 6,162,963 and 6,150,584. See also WO 91/10741, WO 94/02602, WO 96/34096, WO 96/33735, WO 98/16654, WO 98/24893, WO 98/50433, WO 99/45031, WO 99/53049, WO 00/09560, and WO 00/037504.

The methods disclosed in these documents can be modified as described in U.S. Patent No. 5,994,619, which is hereby incorporated by reference. U.S. Patent No. 5,994,619 describes methods for producing novel cultured inner cell mass (CICM) cells and cell lines, derived from pigs and cows, and transgenic CICM cells into which heterologous DNA has been inserted. CICM transgenic cells can be used to produce cloned transgenic embryos, fetuses, and offspring. The '619 patent also describes methods of producing transgenic animals that are capable of transmitting the heterologous DNA to their progeny. Examples of non-human animals that can be used with these methods include rats, sheep, pigs, goats, cattle, chicken, and horses.

XENOMOUSE™ mice produce an adult-like human repertoire of fully human antibodies and generate antigen-specific human antibodies. In some embodiments, the XENOMOUSE™ mice contain approximately 80% of the human antibody V gene repertoire through introduction of megabase sized, germline configuration fragments of the human heavy chain loci and kappa light chain loci in yeast artificial chromosome (YAC). In other embodiments, XENOMOUSE™ mice further contain approximately all of the human lambda light chain locus. See Mendez, et al., *Nature Genetics*, 15:146-156 (1997), Green and Jakobovits, *J. Exp. Med.*, 188:483-495 (1998), and WO 98/24893, the disclosures of which are hereby incorporated by reference.

In some embodiments, the non-human animal comprising human immunoglobulin genes are animals that have a human immunoglobulin "minilocus". In the minilocus approach, an exogenous Ig locus is mimicked through the inclusion of 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 domain, and a second constant domain (preferably a gamma constant domain) are formed into a construct for insertion into an animal. This approach is described in U.S. Patent Nos. 5,545,807, 5,545,806, 5,569,825, 5,625,126, 5,633,425, 5,661,016, 5,770,429, 5,789,650, 5,814,318, 5,591,669, 5,612,205, 5,721,367, 5,789,215, and 5,643,763, hereby incorporated by reference.

Non-human animals as described above can then be immunized with a P-cadherin antigen as described below under conditions that permit antibody production. Antibody-producing cells are isolated from the animals, and nucleic acids encoding the heavy and light chains of P-cadherin antibody of interest are isolated from the isolated antibody-producing cells or from an immortalized cell line produced from such cells. These nucleic acids are subsequently engineered using techniques known to those of skill in the art and as described further below to reduce the amount of non-human sequence, i.e., to humanize the antibody to reduce the immune response in humans.

In some embodiments, the P-cadherin antigen can be isolated and/or purified P-cadherin. In some embodiments, the P-cadherin antigen is human P-cadherin. In other embodiments, the P-cadherin antigen can be a cell that expresses or over-expresses P-cadherin. In other embodiments, the P-cadherin antigen is a recombinant protein expressed from yeast, insect cells, bacteria such as *E.coli*, or other resources by recombinant technology. Immunization of animals can be by any method known in the art. See, e.g., Harlow and Lane, Antibodies: A Laboratory Manual, New York: Cold Spring Harbor Press (1990). Methods for immunizing non-human animals such as mice, rats, sheep, goats, pigs, cattle and horses are well known in the art. See, e.g., Harlow and Lane, supra, and U.S. Patent No. 5,994,619. For example, the P-cadherin antigen can be administered with an adjuvant to stimulate the immune response. Exemplary adjuvants include complete or incomplete Freund's adjuvant, RIBI (muramyl dipeptides) or ISCOM (immunostimulating complexes). Such adjuvants may protect the polypeptide from rapid dispersal by sequestering it in a local deposit, or they may contain substances that stimulate the host to secrete factors that are chemotactic for macrophages and other components of the immune system. Preferably, if a polypeptide is being administered, the immunization schedule will involve two or more administrations of the polypeptide, spread out over several weeks.

For example, following immunization of a transgenic animal as described above with P-cadherin, primary cells (e.g. spleen or peripheral blood B cells) can be isolated from the immunized transgenic animal and individual cells produing antibodies specific for the desired antigen can be identified. Polyadenylated mRNA from each individual cell is then isolated and reverse transcription polymerase chain reaction (RT-PCR) is performed using sense primers that anneal to variable region sequences (e.g., degenerate primers that recognize most or all of the FR1 regions of human heavy and light chain variable region genes and anti-sense primers that anneal to constant or joining region sequences). cDNAs of the heavy and light chain variable domains are then cloned and expressed in

any suitable host cell, e.g., a myeloma cell, as chimeric antibodies with respective immunoglobulin constant regions, such as the heavy chain and κ or λ constant domains. See Babcook, et al., *Proc. Natl. Acad. Sci. USA*, 93:7843-48, (1996), incorporated herein by reference. P-cadherin antibodies may then be identified and isolated as described herein.

Recombinant Methods of Producing Antibodies

An antibody, or antibody portion, of the invention can be prepared by recombinant expression of immunoglobulin light and heavy chain genes in a host cell. For example, to express an antibody recombinantly, a host cell is transfected with one or more recombinant expression vectors carrying DNA fragments encoding the immunoglobulin light and heavy chains of the antibody such that the light and heavy chains are expressed in the host cell and, preferably, secreted into the medium in which the host cells are cultured, from which medium the antibodies can be recovered. Standard recombinant DNA methodologies are used to obtain antibody heavy and light chain genes, to incorporate these genes into recombinant expression vectors and to introduce the vectors into host cells, such as those described in Sambrook, Fritsch and Maniatis (eds), Molecular Cloning: A Laboratory Manual, Second Edition, Cold Spring Harbor, N.Y., (1989), Ausubel, F. M., et al. (eds.) Current Protocols in Molecular Biology, Greene Publishing Associates, (1989) and in U.S. Patent No. 4,816,397, the disclosures of which are incorporated herein by reference.

Mutations and Modifications

To express the P-cadherin antibodies of the present invention, DNA fragments encoding V_H and V_L regions can first be obtained using any of the methods described above. Various mutations, deletions, and/or additions can also be introduced into the DNA sequences using standard methods known to those of skill in the art. For example, mutagenesis can be carried out using standard methods, such as PCR-mediated mutagenesis, in which the mutated nucleotides are incorporated into the PCR primers such that the PCR product contains the desired mutations or site-directed mutagenesis. One type of substitution, for example, that may be made is to change one or more cysteines in the antibody, which may be chemically reactive, to another residue, such as, without limitation, alanine or serine. For example, there can be a substitution of a non-canonical cysteine. The substitution can be made in a CDR or framework region of a variable domain or in the constant domain of an antibody. In some embodiments, the cysteine is canonical.

The antibodies may also be mutated in the variable domains of the heavy and/or light chains, e.g., to alter a binding property of the antibody. For example, a mutation may be made in one or more of the CDR regions to increase or decrease the K_D of the antibody for P-cadherin, to increase or decrease k_{off}, or to alter the binding specificity of the antibody. Techniques in site-directed mutagenesis are well-known in the art. See, e.g., Sambrook, et al. and Ausubel, et al., *supra*, which is incorporated herein by reference. For example, as discussed in greater detail in Example 8, numerous variant V_H and V_L CDR3 sequences of the 129-1c4 parent were made according to the procedures discussed above, and are indicated as SEQ ID NOs: 91 to 256 (V_H CDR3 variants) and SEQ ID NOs: 257 to 319 (V_L CDR3 variants) in Figure 1.

A mutation may also be made in a framework region or constant domain to increase the half-life of a P-cadherin antibody. See, e.g., PCT Publication No. WO 00/09560, incorporated herein by reference. A mutation in a framework region or constant domain can also be made to alter the immunogenicity of the antibody, to provide a site for covalent or non-covalent binding to another molecule, or to alter such properties as complement fixation, FcR binding and antibody-dependent cell-mediated cytotoxicity (ADCC). According to the invention, a single antibody may have mutations in any one or more of the CDRs or framework regions of the variable domain or in the constant domain.

In a process known as "germlining", certain amino acids in the V_H and V_L sequences can be mutated to match thoses found naturally in germline V_H and V_L sequences. In particular, the amino acid sequences of the framework regions in the V_H and V_L sequences can be mutated to match the germline sequences to reduce the risk of immunogenicity when the antibody is administered. Germline DNA sequences for human V_H and V_L genes are known in the art (see e.g., the "Vbase" human germline sequence database; see also Kabat, E. A., et al. (1991) Sequences of Proteins of Immunological Interest, Fifth Edition, U.S. Department of Health and Human Services, NIH Publication No. 91-3242; Tomlinson, et al. (1992) *J. Mol. Biol.* 227:776-798; and Cox, et al. *Eur. J. Immunol.* 24:827-836 (1994); the contents of each of which are expressly incorporated herein by reference).

Another type of amino acid substitution that may be made is to remove potential proteolytic sites in the antibody. Such sites may occur in a CDR or framework region of a variable domain or in the constant domain of an antibody. Substitution of cysteine residues and removal of proteolytic sites may decrease the risk of heterogeneity in the antibody product and thus increase its homogeneity. Another type of amino acid substitution is to eliminate asparagine-glycine pairs, which form potential deamidation sites, by altering one or both of the residues. In another example, the C-terminal lysine of the heavy chain of a P-cadherin antibody of the invention can be cleaved. In various embodiments of the invention, the heavy and light chains of the P-cadherin antibodies may optionally include a N-terminal signal sequence, such as those indicated by SEQ ID NOs: 346 and 347.

Once DNA fragments encoding the V_H and V_L segments of the present invention are obtained, these DNA fragments can be further manipulated by standard recombinant DNA techniques, for example to convert the variable region genes to full-length antibody chain genes, to Fab fragment genes, or to a scFv gene. In these manipulations, a V_L- or V_H-encoding DNA fragment is operatively linked to another DNA fragment encoding another protein, such as an antibody constant region or a flexible linker. The term "operatively linked", as used in this context, is intended to mean that the two DNA fragments are joined such that the amino acid sequences encoded by the two DNA fragments remain in-frame.

The isolated DNA encoding the V_H region can be converted to a full-length heavy chain gene by operatively linking the V_H-encoding DNA to another DNA molecule encoding heavy chain constant regions (CH1, CH2 and CH3). The sequences of human heavy chain constant region genes are

known in the art (see e.g., Kabat, E. A., et al. (1991) Sequences of Proteins of Immunological Interest, Fifth Ed., U.S. Department of Health and Human Services, NIH Publication No. 91-3242) and DNA fragments encompassing these regions can be obtained by standard PCR amplification. The heavy chain constant region can be an IgG1, IgG2, IgG3, IgG4, IgA, IgE, IgM or IgD constant region, but most preferably is an IgG1 or IgG2 constant region. The IgG1 constant region sequence can be any of the various alleles or allotypes known to occur among different individuals, such as Gm(1), Gm(2), Gm(3), and Gm(17). These allotypes represent naturally occurring amino acid substitution in the IgG1 constant regions. For example, the heavy chain IgG1 constant region can be SEQ ID NO: 344. For a Fab fragment heavy chain gene, the V_H-encoding DNA can be operatively linked to another DNA molecule encoding only the heavy chain CH1 constant region. The CH1 heavy chain constant region may be derived from any of the heavy chain genes.

The isolated DNA encoding the V_L region can be converted to a full-length light chain gene (as well as a Fab light chain gene) by operatively linking the V_L-encoding DNA to another DNA molecule encoding the light chain constant region, C_L. The sequences of human light chain constant region genes are known in the art (see e.g., Kabat, E. A., et al. (1991) Sequences of Proteins of Immunological Interest, Fifth Ed., U.S. Department of Health and Human Services, NIH Publication No. 91-3242) and DNA fragments encompassing these regions can be obtained by standard PCR amplification. The light chain constant region can be a kappa or lambda constant region. The kappa constant region may be any of the various alleles known to occur among different individuals, such as Inv(1), Inv(2), and Inv(3). The lambda constant region may be derived from any of the three lambda genes. For example the light chain IgG1 constant region can be SEQ ID NO: 347.

To create a scFv gene, the V_H- and V_L-encoding DNA fragments are operatively linked to another fragment encoding a flexible linker, e.g., encoding the amino acid sequence $(Gly_4\text{-}Ser)_3$, such that the V_H and V_L sequences can be expressed as a contiguous single-chain protein, with the V_L and V_H regions joined by the flexible linker (see e.g., Bird et al. *Science* 242:423-426 (1988); Huston et al. *Proc. Natl. Acad. Sci. USA* 85:5879-5883 (1988); McCafferty, et al., *Nature,* 348:552-554 (1990)). The single chain antibody may be monovalent, if only a single V_H and V_L are used, bivalent, if two V_H and V_L are used, or polyvalent, if more than two V_H and V_L are used. Bispecific or polyvalent antibodies may be generated that bind specifically to P-cadherin and to another molecule.

In another embodiment, a fusion antibody or immunoadhesin may be made that comprises all or a portion of a P-cadherin antibody of the invention linked to another polypeptide. In another embodiment, only the variable domains of the P-cadherin antibody are linked to the polypeptide. In another embodiment, the V_H domain of a P-cadherin antibody is linked to a first polypeptide, while the V_L domain of a P-cadherin antibody is linked to a second polypeptide that associates with the first polypeptide in a manner such that the V_H and V_L domains can interact with one another to form an antigen binding site. In another preferred embodiment, the V_H domain is separated from the V_L domain by a linker such that the V_H and V_L domains can interact with one another. The V_H-linker-V_L antibody is then linked to the polypeptide of interest. In addition, fusion antibodies can be created in which two (or more) single-chain antibodies are linked to one another. This is useful if one wants to

create a divalent or polyvalent antibody on a single polypeptide chain, or if one wants to create a bispecific antibody.

In other embodiments, other modified antibodies may be prepared using P-cadherin antibody encoding nucleic acid molecules. For instance, "Kappa bodies" (III, et al., *Protein Eng.* 10: 949-57 (1997)), "Minibodies" (Martin, et al., *EMBO J.*, 13: 5303-9 (1994)), "Diabodies" (Holliger, et al., *Proc. Natl. Acad. Sci. USA*, 90: 6444-6448 (1993)), or "Janusins" (Traunecker, et al., *EMBO J.*, 10:3655-3659 (1991) and Traunecker, et al., *Int. J. Cancer*, (Suppl.) 7:51-52 (1992)) may be prepared using standard molecular biological techniques following the teachings of the specification.

Bispecific antibodies or antigen-binding fragments can be produced by a variety of methods including fusion of hybridomas or linking of Fab' fragments. See, e.g., Songsivilai & Lachmann, *Clin. Exp. Immunol.*, 79:315-321 (1990), Kostelny, et al., *J. Immunol.*, 148:1547-1553 (1992). In addition, bispecific antibodies may be formed as "diabodies" or "Janusins." In some embodiments, the bispecific antibody binds to two different epitopes of P-cadherin. In some embodiments, the modified antibodies described above are prepared using one or more of the variable domains or CDR regions from a human P-cadherin antibody provided herein.

Vectors and Host Cells

To express the antibodies and antigen-binding portions of the invention, DNAs encoding partial or full-length light and heavy chains, obtained as described above, are inserted into expression vectors such that the genes are operatively linked to transcriptional and translational control sequences. In this context, the term "operatively linked" is intended to mean that an antibody gene is ligated into a vector such that transcriptional and translational control sequences within the vector serve their intended function of regulating the transcription and translation of the antibody gene. The expression vector and expression control sequences are chosen to be compatible with the expression host cell used. Expression vectors include plasmids, retroviruses, adenoviruses, adeno-associated viruses (AAV), plant viruses such as cauliflower mosaic virus, tobacco mosaic virus, cosmids, YACs, EBV derived episomes, and the like. The antibody gene is ligated into a vector such that transcriptional and translational control sequences within the vector serve their intended function of regulating the transcription and translation of the antibody gene. The expression vector and expression control sequences are chosen to be compatible with the expression host cell used. The antibody light chain gene and the antibody heavy chain gene can be inserted into separate vectors. In a preferred embodiment, both genes are inserted into the same expression vector. The antibody genes are inserted into the expression vector by standard methods (e.g., ligation of complementary restriction sites on the antibody gene fragment and vector, or blunt end ligation if no restriction sites are present).

A convenient vector is one that encodes a functionally complete human C_H or C_L immunoglobulin sequence, with appropriate restriction sites engineered so that any V_H or V_L sequence can easily be inserted and expressed, as described above. In such vectors, splicing usually occurs between the splice donor site in the inserted J region and the splice acceptor site preceding the human C domain, and also at the splice regions that occur within the human C_H exons.

Polyadenylation and transcription termination occur at native chromosomal sites downstream of the coding regions. The recombinant expression vector also can encode a signal peptide that facilitates secretion of the antibody chain from a host cell. The antibody chain gene may be cloned into the vector such that the signal peptide is linked in-frame to the amino terminus of the immunoglobulin chain. The signal peptide can be an immunoglobulin signal peptide or a heterologous signal peptide (i.e., a signal peptide from a non-immunoglobulin protein).

In addition to the antibody chain genes, the recombinant expression vectors of the invention carry regulatory sequences that control the expression of the antibody chain genes in a host cell. It will be appreciated by those skilled in the art that the design of the expression vector, including the selection of regulatory sequences may depend on such factors as the choice of the host cell to be transformed, the level of expression of protein desired, and so forth. Preferred regulatory sequences for mammalian host cell expression include viral elements that direct high levels of protein expression in mammalian cells, such as promoters and/or enhancers derived from retroviral LTRs, cytomegalovirus (CMV) (such as the CMV promoter/enhancer), Simian Virus 40 (SV40) (such as the SV40 promoter/enhancer), adenovirus, (e.g., the adenovirus major late promoter (AdMLP)), polyoma and strong mammalian promoters such as native immunoglobulin and actin promoters. For further description of viral regulatory elements, and sequences thereof, see e.g., U.S. Patent No. 5,168,062, U.S. Patent No. 4,510,245 and U.S. Patent No. 4,968,615. Methods for expressing antibodies in plants, including a description of promoters and vectors, as well as transformation of plants is known in the art. See, e.g., U. S. Patent No. 6,517,529, incorporated herein by reference. Methods of expressing polypeptides in bacterial cells or fungal cells, e.g., yeast cells, are also well known in the art.

In addition to the antibody chain genes and regulatory sequences, the recombinant expression vectors of the invention may carry additional sequences, such as sequences that regulate replication of the vector in host cells (e.g., origins of replication) and selectable marker genes. The selectable marker gene facilitates selection of host cells into which the vector has been introduced (see e.g., U.S. Patent Nos. 4,399,216, 4,634,665 and 5,179,017, incorporated herein by reference). For example, typically the selectable marker gene confers resistance to drugs, such as G418, hygromycin or methotrexate, on a host cell into which the vector has been introduced. Preferred selectable marker genes include the dihydrofolate reductase (DHFR) gene (for use in dhfr-host cells with methotrexate selection/amplification), the neomycin phosphotransferase gene (for G418 selection), and the glutamate synthetase gene.

Nucleic acid molecules encoding P-cadherin antibodies and vectors comprising these nucleic acid molecules can be used for transfection of a suitable mammalian, plant, bacterial or yeast host cell. Transformation can be by any known method for introducing polynucleotides into a host cell. Methods for introduction of 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. In addition, nucleic acid molecules may

be introduced into mammalian cells by viral vectors. Methods of transforming cells are well known in the art. See, e.g., U.S. Patent Nos. 4,399,216, 4,912,040, 4,740,461, and 4,959,455, incorporated herein by reference). Methods of transforming plant cells are well known in the art, including, e.g., Agrobacterium-mediated transformation, biolistic transformation, direct injection, electroporation and viral transformation. Methods of transforming bacterial and yeast cells are also well known in the art.

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). These include, for example, Chinese hamster ovary (CHO) cells, NSO cells, SP2 cells, HEK-293T cells, NIH-3T3 cells, HeLa cells, baby hamster kidney (BHK) cells, African green monkey kidney cells (COS), human hepatocellular carcinoma cells (e.g., Hep G2), A549 cells, and a number of other cell lines. Cell lines of particular preference are selected through determining which cell lines have high expression levels. Other cell lines that may be used are insect cell lines, such as Sf9 or Sf21 cells. When recombinant expression vectors encoding antibody genes are introduced into mammalian host cells, the antibodies are produced by culturing the host cells for a period of time sufficient to allow for expression of the antibody in the host cells or, more preferably, secretion of the antibody into the culture medium in which the host cells are grown. Antibodies can be recovered from the culture medium using standard protein purification methods. Plant host cells include, e.g., Nicotiana, Arabidopsis, duckweed, corn, wheat, potato, and so forth. Bacterial host cells include *E. coli* and *Streptomyces* species. Yeast host cells include *Schizosaccharomyces pombe*, *Saccharomyces cerevisiae* and *Pichia pastoris*.

Further, expression of antibodies of the invention from production cell lines can be enhanced using a number of known techniques. For example, the glutamine synthetase (the GS system) and DHFR gene expression systems are common approaches for enhancing expression under certain conditions. High expressing cell clones can be identified using conventional techniques, such as limited dilution cloning and Microdrop technology. The GS system is discussed in European Patent Nos. 0 216 846, 0 256 055, 0 323 997 and 0 338 841.

It is likely that antibodies expressed by different cell lines or in transgenic animals will have different glycosylation from each other. However, all antibodies encoded by the nucleic acid molecules provided herein, or comprising the amino acid sequences provided herein are part of the present invention, regardless of the glycosylation of the antibodies.

Transgenic Animals and Plants

P-cadherin antibodies of the invention also can be produced transgenically through the generation of a mammal or plant that is transgenic for the immunoglobulin heavy and light chain sequences of interest and production of the antibody in a recoverable form therefrom. In connection with the transgenic production in mammals, P-cadherin antibodies can be produced in, and recovered from, the milk of goats, cows, or other mammals. See, e.g., U.S. Patent Nos. 5,827,690, 5,756,687, 5,750,172, and 5,741,957, incorporated herein by reference. In some embodiments, non-human transgenic animals that comprise human immunoglobulin loci are immunized with P-cadherin

or an immunogenic portion thereof, as described above. Methods for making antibodies in plants are described, e.g., in U.S. patents 6,046,037 and 5,959,177, incorporated herein by reference.

In some embodiments, non-human transgenic animals or plants are produced by introducing one or more nucleic acid molecules encoding a P-cadherin antibody, or antigen binding portion thereof, of the invention into the animal or plant by standard transgenic techniques. See Hogan and United States Patent 6,417,429, *supra.* The transgenic cells used for making the transgenic animal can be embryonic stem cells or somatic cells or a fertilized egg. The transgenic non-human organisms can be chimeric, nonchimeric heterozygotes, and nonchimeric homozygotes. See, e.g., Hogan et al., Manipulating the Mouse Embryo: A Laboratory Manual, 2nd ed., Cold Spring Harbor Press (1999); Jackson, et al., Mouse Genetics and Transgenics: A Practical Approach, Oxford University Press (2000); and Pinkert, Transgenic Animal Technology: A Laboratory Handbook, Academic Press (1999), all incorporated herein by reference. In some embodiments, the transgenic non-human animals have a targeted disruption and replacement by a targeting construct that encodes a heavy chain and/or a light chain of interest. The P-cadherin antibodies may be made in any transgenic animal. In a preferred embodiment, the non-human animals are mice, rats, sheep, pigs, goats, cattle or horses. The non-human transgenic animal expresses said encoded polypeptides in blood, milk, urine, saliva, tears, mucus and other bodily fluids.

Class switching

The class (e.g. IgG, IgM, IgE, IgA, or IgD) and subclass (e.g. IgG1, IgG2, IgG3, or IgG4) of P-cadherin antibodies may be determined by any method known in the art. In general, the class and subclass of an antibody may be determined using antibodies that are specific for a particular class and subclass of antibody. Such antibodies are commercially available. The class and subclass can be determined by ELISA, or Western Blot as well as other techniques. Alternatively, the class and subclass may be determined by sequencing all or a portion of the constant domains of the heavy and/or light chains of the antibodies, comparing their amino acid sequences to the known amino acid sequences of various class and subclasses of immunoglobulins, and determining the class and subclass of the antibodies. The P-cadherin antibodies of the present invention can be an IgG, an IgM, an IgE, an IgA, or an IgD molecule. For example, the P-cadherin antibodies can be an IgG that is an IgG1, IgG2, IgG3, or an IgG4 subclass. In one embodiment, the P-cadherin antibodies can have a heavy chain constant region indicated by SEQ ID NO: 344 and a light chain constant region indicated by SEQ ID NO: 345.

One aspect of the invention provides a method for converting the class or subclass of a P-cadherin antibody to another class or subclass. In some embodiments, a nucleic acid molecule encoding a V_L or V_H that does not include sequences encoding C_L or C_H is isolated using methods well-known in the art. The nucleic acid molecule then is operatively linked to a nucleic acid sequence encoding a C_L or C_H from a desired immunoglobulin class or subclass. This can be achieved using a vector or nucleic acid molecule that comprises a C_L or C_H chain, as described above. For example, a P-cadherin antibody that was originally IgM can be class switched to an IgG. Further, the class switching may be used to convert one IgG subclass to another, e.g., from IgG1 to

IgG2. Another method for producing an antibody of the invention comprising a desired isotype comprises the steps of isolating a nucleic acid encoding a heavy chain of a P-cadherin antibody and a nucleic acid encoding a light chain of a P-cadherin antibody, isolating the sequence encoding the V_H region, ligating the V_H sequence to a sequence encoding a heavy chain constant domain of the desired isotype, expressing the light chain gene and the heavy chain construct in a cell, and collecting the P-cadherin antibody with the desired isotype.

Deimmunized Antibodies

In another aspect of the invention, the antibodies or antigen binding portions thereof may be deimmunized to reduce their immunogenicity using the techniques described in, e.g., PCT Publication Nos. WO98/52976 and WO00/34317 (incorporated herein by reference).

Derivatized and Labeled Antibodies

A P-cadherin antibody or antigen-binding portion of the invention can be derivatized or linked to another molecule (e.g., another peptide or protein). In general, the antibodies or portion thereof are derivatized such that the P-cadherin binding is not affected adversely by the derivatization or labeling. Accordingly, the antibodies and antibody portions of the invention are intended to include both intact and modified forms of the human P-cadherin antibodies described herein. For example, an antibody or antibody portion of the invention can be functionally linked (by chemical coupling, genetic fusion, noncovalent association or otherwise) to one or more other molecular entities, such as another antibody (e.g., a bispecific antibody or a diabody), a detection agent, a label, a cytotoxic agent, a pharmaceutical agent, and/or a protein or peptide that can mediate association of the antibody or antibody portion with another molecule (such as a streptavidin core region or a polyhistidine tag).

One type of derivatized antibody is produced by crosslinking two or more antibodies (of the same type or of different types, e.g., to create bispecific antibodies). Suitable crosslinkers include those that are heterobifunctional, having two distinctly reactive groups separated by an appropriate spacer (e.g., m-maleimidobenzoyl-N-hydroxysuccinimide ester) or homobifunctional (e.g., disuccinimidyl suberate). Such linkers are available from Pierce Chemical Company, Rockford, IL.

Another type of derivatized antibody is a labeled antibody. Useful detection agents with which an antibody or antigen-binding portion of the invention may be derivatized include fluorescent compounds, including fluorescein, fluorescein isothiocyanate, rhodamine, 5-dimethylamine-1-naphthalenesulfonyl chloride, phycoerythrin, lanthanide phosphors, and the like. An antibody can also be labeled with enzymes that are useful for detection, such as horseradish peroxidase, β-galactosidase, luciferase, alkaline phosphatase, glucose oxidase, and the like. When an antibody is labeled with a detectable enzyme, it is detected by adding additional reagents that the enzyme uses to produce a reaction product that can be discerned. For example, when the agent horseradish peroxidase is present, the addition of hydrogen peroxide and diaminobenzidine leads to a colored reaction product, which is detectable. An antibody can also be labeled with biotin, and detected through indirect measurement of avidin or streptavidin binding. An antibody can also be labeled with a predetermined polypeptide epitope recognized by a secondary reporter (e.g., leucine zipper pair

sequences, binding sites for secondary antibodies, metal binding domains, epitope tags). In some embodiments, labels are attached by spacer arms of various lengths to reduce potential steric hindrance. A P-cadherin antibody can also be derivatized with a chemical group such as polyethylene glycol (PEG), a methyl or ethyl group, or a carbohydrate group. These groups are useful to improve the biological characteristics of the antibody, e.g., to increase serum half-life.

Binding Affinity of P-cadherin Antibodies to P-cadherin

The binding affinity (K_D) and dissociation rate (k_{off}) of a P-cadherin antibody or antigen-binding portion thereof to P-cadherin can be determined by methods known in the art. The binding affinity can be measured by ELISAs, RIAs, flow cytometry, or surface plasmon resonance, such as BIACORE™. The dissociation rate can be measured by surface plasmon resonance. Preferably, the binding affinity and dissociation rate is measured by surface plasmon resonance. More preferably, the binding affinity and dissociation rate are measured using BIACORE™. One can determine whether an antibody has substantially the same K_D as a P-cadherin antibody by using methods known in the art. Such methods of determining K_D and k_{off} can be used during the initial screening stage, as well as during subsequent optimization stages.

Identification of P-cadherin Epitopes Recognized by P-cadherin Antibodies

The invention provides human P-cadherin antibodies that bind to P-cadherin and compete or cross-compete with and/or binds the same epitope as any of the antibodies as described in Tables 1 or 2. One can determine whether an antibody binds to the same epitope or cross competes for binding with a P-cadherin antibody of the present invention by using methods known in the art. In one embodiment, one allows the P-cadherin antibody of the invention to bind to P-cadherin under saturating conditions and then measures the ability of the test antibody to bind to P-cadherin. If the test antibody is able to bind to P-cadherin at the same time as the P-cadherin antibody, then the test antibody binds to a different epitope as the P-cadherin antibody. However, if the test antibody is not able to bind to P-cadherin at the same time, then the test antibody binds to the same epitope, an overlapping epitope, or an epitope that is in close proximity to the epitope bound by the human P-cadherin antibody. This experiment can be performed using ELISA, RIA, BIACORE™, or flow cytometry. In a preferred embodiment, the experiment is performed using ELISA.

Inhibition of P-cadherin Activity by P-cadherin Antibody

P-cadherin antibodies that inhibit P-cadherin activity can be identified using a number of assays. A cell aggregation assay, for example, provides a method of measuring P-cadherin-dependent cellular aggregation. This type of assay uses a cell line that over-expresses P-cadherin, wherein the cells are placed into suspension and allowed to form P-cadherin-dependent aggregates. The aggregation assay is then used to quantify the ability of a P-cadherin antibody to prevent this aggregation by measuring the size of cellular aggregates that result with and without the antibody. Cell aggregate size as a function of P-cadherin antibody concentration can then be used to determine an IC_{50} value. Example 4 provides further details of a P-cadherin-dependent aggregation assay that was used to measure IC_{50} values for several P-cadherin antibodies.

Cell adhesion assays can also be used to measure the ability of a P-cadherin antibody to block the adhesion of cells to a receptor P-cadherin that has been immobilized on a solid support. This type of assay can be carried out, for example, by immobilizing P-cadherin on a solid support, such as plastic. Cells over-expressing P-cadherin are then allowed to adhere to the solid support via P-cadherin-P-cadherin interactions. The level of adhesion can then be quantified with and without a P-cadherin antibody. Adhesion as a function of antibody concentration is then used to determine an IC_{50} value. Example 3 provides further details of a P-cadherin-dependent cell adhesion assay that was used to measure IC_{50} values for P-cadherin antibodies.

Inhibition of P-cadherin activity can also be measured using a P-cadherin dependent spheroid disruption assay. This type of assay measures the ability of a P-cadherin antibody to disrupt pre-formed P-cadherin-dependent cellular aggregations. By measuring the size reduction of aggregates as a function of antibody concentration, an IC_{50} value can be determined. Example 5 provides further details of a P-cadherin-dependent spheroid disruption assay that was used to measure IC_{50} values for P-cadherin antibodies. The methods and assays described above for determining inhibition of P-cadherin activity by various antibodies or antigen-binding portions thereof can be used during the initial screening stage, as well as during subsequent optimization stages.

Molecular Selectivity

The selectivity of the P-cadherin antibodies of the present invention over other cadherins, such as E-cadherin, can be determined using methods well known in the art. For example one can determine the selectivity using Western blot, flow cytometry, ELISA, immunoprecipitation or RIA. Example 7 provides further details of an ELISA assay that was used to measure the selectivity of specific antibodies to P-cadherin over E-cadherin. The methods and assays described above for determining the selectivity for P-cadherin of various antibodies or antigen-binding portions thereof can be used during the initial screening stage, as well as during subsequent optimization stages.

Pharmaceutical Compositions and Administration

This invention also relates to a pharmaceutical composition for the treatment of abnormal cell growth in a mammal, including a human, comprising an amount of a P-cadherin antibody or antigen binding portion thereof, as described herein, that is effective in treating abnormal cell growth, and a pharmaceutically acceptable carrier.

The antibodies and antigen-binding portions of the present invention can be incorporated into pharmaceutical compositions suitable for administration to a subject. Typically, the pharmaceutical composition comprises an antibody or antigen-binding portion of the invention and a pharmaceutically acceptable carrier. As used herein, "pharmaceutically acceptable carrier" means any and all solvents, dispersion media, coatings, antibacterial and antifungal agents, isotonic and absorption delaying agents, and the like that are physiologically compatible. Some examples of pharmaceutically acceptable carriers are water, saline, phosphate buffered saline, dextrose, glycerol, ethanol and the like, as well as combinations thereof. In many cases, it will be preferable to include isotonic agents, for example, sugars, polyalcohols such as mannitol, sorbitol, or sodium chloride in the composition. Additional examples of pharmaceutically acceptable substances are wetting agents

or minor amounts of auxiliary substances such as wetting or emulsifying agents, preservatives or buffers, which enhance the shelf life or effectiveness of the antibody.

The compositions of this invention may be in a variety of forms, for example, liquid, semi-solid and solid dosage forms, such as liquid solutions (e.g., injectable and infusible solutions), dispersions or suspensions, tablets, pills, powders, liposomes and suppositories. The preferred form depends on the intended mode of administration and therapeutic application. Typical preferred compositions are in the form of injectable or infusible solutions, such as compositions similar to those used for passive immunization of humans. The preferred mode of administration is parenteral (e.g., intravenous, subcutaneous, intraperitoneal, intramuscular). In a preferred embodiment, the antibody is administered by intravenous infusion or injection. In another preferred embodiment, the antibody is administered by intramuscular or subcutaneous injection. Formulations for injection may be presented in unit dosage form, e.g., in ampoules or in multi-dose containers, with or without an added preservative. The compositions may take such forms as suspensions, solutions, or emulsions in oily or aqueous vehicles, and may contain formulatory agents such as suspending, stabilizing and/or dispersing agents. Alternatively, the active ingredient may be in powder form for constitution with a suitable vehicle, e.g., sterile pyrogen-free water, before use.

Therapeutic compositions typically must be sterile and stable under the conditions of manufacture and storage. The composition can be formulated as a solution, microemulsion, dispersion, liposome, or other ordered structure suitable to high drug concentration. Sterile injectable solutions can be prepared by incorporating the P-cadherin antibody in the required amount in an appropriate solvent with one or a combination of ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the active compound into a sterile vehicle that contains a basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the preferred methods of preparation are vacuum drying and freeze-drying that yields a powder of the active ingredient plus any additional desired ingredient from a previously sterile-filtered solution thereof. The proper fluidity of a solution can be maintained, for example, by the use of a coating such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. Prolonged absorption of injectable compositions can be brought about by including in the composition an agent that delays absorption, for example, monostearate salts and gelatin.

The antibodies or antibody portions of the present invention can be administered by a variety of methods known in the art, although for many therapeutic applications, the preferred route/mode of administration is subcutaneous, intramuscular, or intravenous infusion. As will be appreciated by the skilled artisan, the route and/or mode of administration will vary depending upon the desired results.

In certain embodiments, the antibody compositions of the present invention may be prepared with a carrier that will protect the antibody against rapid release, such as a controlled release formulation, including implants, transdermal patches, and microencapsulated delivery systems. Biodegradable, biocompatible polymers can be used, such as ethylene vinyl acetate, polyanhydrides,

polyglycolic acid, collagen, polyorthoesters, and polylactic acid. Many methods for the preparation of such formulations are generally known to those skilled in the art. See, e.g., Sustained and Controlled Release Drug Delivery Systems, J. R. Robinson, ed., Marcel Dekker, Inc., New York, 1978, which is incorporated herein by reference.

Additional active compounds also can be incorporated into the compositions. In certain embodiments, an inhibitory P-cadherin antibody of the invention is co-formulated with and/or co-administered with one or more additional therapeutic agents. These agents include, without limitation, antibodies that bind other targets, anti-tumor agents, anti-angiogenesis agents, signal transduction inhibitors, anti-proliferative agents, chemotherapeutic agents, or peptide analogues that inhibit P-cadherin. Such combination therapies may require lower dosages of the inhibitory P-cadherin antibody as well as the co-administered agents, thus avoiding possible toxicities or complications associated with the various monotherapies.

The compositions of the invention may include a "therapeutically effective amount" or a "prophylactically effective amount" of an antibody or antigen-binding portion of the invention. A "therapeutically effective amount" refers to an amount effective, at dosages and for periods of time necessary, to achieve the desired therapeutic result. A therapeutically effective amount of the antibody or antigen-binding portion may vary according to factors such as the disease state, age, sex, and weight of the individual, and the ability of the antibody or antibody portion to elicit a desired response in the individual. A therapeutically effective amount is also one in which any toxic or detrimental effects of the antibody or antigen-binding portion are outweighed by the therapeutically beneficial effects. A "prophylactically effective amount" refers to an amount effective, at dosages and for periods of time necessary, to achieve the desired prophylactic result. Typically, since a prophylactic dose is used in subjects prior to or at an earlier stage of disease, the prophylactically effective amount may be less than the therapeutically effective amount.

Dosage regimens can be adjusted to provide the optimum desired response (e.g., a therapeutic or prophylactic response). For example, a single bolus can be administered, several divided doses can be administered over time or the dose can be proportionally reduced or increased as indicated by the exigencies of the therapeutic situation. It is especially advantageous to formulate parenteral compositions in dosage unit form for ease of administration and uniformity of dosage. Dosage unit form as used herein refers to physically discrete units suited as unitary dosages for the mammalian subjects to be treated; each unit containing a predetermined quantity of active compound calculated to produce the desired therapeutic effect in association with the required pharmaceutical carrier. The specification for the dosage unit forms of the invention are dictated by and directly dependent on (a) the unique characteristics of the P-cadherin antibody or portion thereof and the particular therapeutic or prophylactic effect to be achieved, and (b) the limitations inherent in the art of compounding such an antibody for the treatment of sensitivity in individuals.

An exemplary, non-limiting range for a therapeutically or prophylactically effective amount of an antibody or antibody portion of the invention is 0.025 to 50 mg/kg, more preferably 0.1 to 50 mg/kg, more preferably 0.1-25, 0.1 to 10 or 0.1 to 3 mg/kg. In some embodiments, a formulation

contains 5 mg/mL of antibody in a buffer of 20mM sodium citrate, pH 5.5, 140mM NaCl, and 0.2mg/mL polysorbate 80. It is to be noted that dosage values may vary with the type and severity of the condition to be alleviated. It is to be further understood that for any particular subject, specific dosage regimens should be adjusted over time according to the individual need and the professional judgment of the person administering or supervising the administration of the compositions, and that dosage ranges set forth herein are exemplary only and are not intended to limit the scope or practice of the claimed composition.

Another aspect of the present invention provides kits comprising a P-cadherin antibody or antigen-binding portion of the invention or a composition comprising such an antibody or portion. A kit may include, in addition to the antibody or composition, diagnostic or therapeutic agents. A kit can also include instructions for use in a diagnostic or therapeutic method. In a preferred embodiment, the kit includes the antibody or a composition comprising it and a diagnostic agent that can be used in a method described below. In another preferred embodiment, the kit includes the antibody or a composition comprising it and one or more therapeutic agents that can be used in a method described below.

Diagnostic Methods of Use

The P-cadherin antibodies or antigen-binding portions thereof can be used in diagnostic methods to detect P-cadherin in a biological sample *in vitro* or *in vivo*. For example, the P-cadherin antibodies can be used in a conventional immunoassay, including, without limitation, an ELISA, an RIA, flow cytometry, tissue immunohistochemistry, Western blot or immunoprecipitation. The P-cadherin antibodies of the invention can be used to detect P-cadherin from humans. The P-cadherin antibodies can also be used to detect P-cadherin from mice, rats, and cynomolgus monkeys.

The invention provides a method for detecting P-cadherin in a biological sample comprising contacting the biological sample with a P-cadherin antibody of the invention and detecting the bound antibody. In one embodiment, the P-cadherin antibody is directly labeled with a detectable label. In another embodiment, the P-cadherin antibody (the first antibody) is unlabeled and a second antibody or other molecule that can bind the P-cadherin antibody is labeled. As is well known to one of skill in the art, a second antibody is chosen that is able to specifically bind the particular species and class of the first antibody. For example, if the P-cadherin antibody is a human IgG, then the secondary antibody could be an anti-human-IgG. Other molecules that can bind to antibodies include, without limitation, Protein A and Protein G, both of which are available commercially, e.g., from Pierce Chemical Co.

Suitable labels for the antibody or secondary antibody have been discussed previously, and include various enzymes, prosthetic groups, fluorescent materials, luminescent materials and radioactive materials. Examples of suitable enzymes include horseradish peroxidase, alkaline phosphatase, β-galactosidase, or acetylcholinesterase; examples of suitable prosthetic group complexes include streptavidin/biotin and avidin/biotin; examples of suitable fluorescent materials include umbelliferone, fluorescein, fluorescein isothiocyanate, rhodamine, dichlorotriazinylamine

fluorescein, dansyl chloride or phycoerythrin; an example of a luminescent material includes luminol; and examples of suitable radioactive material include ^{125}I, ^{131}I, ^{35}S or ^{3}H.

In other embodiments, P-cadherin can be assayed in a biological sample by a competition immunoassay utilizing P-cadherin standards labeled with a detectable substance and an unlabeled P-cadherin antibody. In this assay, the biological sample, the labeled P-cadherin standards and the P-cadherin antibody are combined and the amount of labeled P-cadherin standard bound to the unlabeled antibody is determined. The amount of P-cadherin in the biological sample is inversely proportional to the amount of labeled P-cadherin standard bound to the P-cadherin antibody.

One can use the immunoassays disclosed above for a number of purposes. For example, the P-cadherin antibodies can be used to detect P-cadherin in cultured cells. In a preferred embodiment, the P-cadherin antibodies are used to determine the amount of P-cadherin produced by cells that have been treated with various compounds. This method can be used to identify compounds that modulate P-cadherin protein levels. According to this method, one sample of cells is treated with a test compound for a period of time while another sample is left untreated. If the total level of P-cadherin is to be measured, the cells are lysed and the total P-cadherin level is measured using one of the immunoassays described above. The total level of P-cadherin in the treated versus the untreated cells is compared to determine the effect of the test compound.

A preferred immunoassay for measuring total P-cadherin levels is flow cytometry or immunohistochemistry. Methods such as ELISA, RIA, flow cytometry, Western blot, immunohistochemistry, cell surface labeling of integral membrane proteins and immunoprecipitation are well known in the art. See, e.g., Harlow and Lane, *supra*. In addition, the immunoassays can be scaled up for high throughput screening in order to test a large number of compounds for either activation or inhibition of P-cadherin expression.

The P-cadherin antibodies of the invention also can be used to determine the levels of P-cadherin in a tissue or in cells derived from the tissue. In some embodiments, the tissue is a diseased tissue. In some embodiments of the method, a tissue or a biopsy thereof is excised from a patient. The tissue or biopsy is then used in an immunoassay to determine, e.g., total P-cadherin levels or localization of P-cadherin by the methods discussed above.

The antibodies of the present invention also can be used *in vivo* to identify tissues and organs that express P-cadherin. One advantage of using the human P-cadherin antibodies of the present invention is that they may safely be used *in vivo* without eliciting a substantial immune response to the antibody upon administration, unlike antibodies of non-human origin or with humanized or chimeric antibodies.

The method comprises the steps of administering a detectably labeled P-cadherin antibody or a composition comprising them to a patient in need of such a diagnostic test and subjecting the patient to imaging analysis to determine the location of the P-cadherin-expressing tissues. Imaging analysis is well known in the medical art, and includes, without limitation, x-ray analysis, magnetic resonance imaging (MRI) or computed tomography (CT). The antibody can be labeled with any agent suitable for *in vivo* imaging, for example a contrast agent, such as barium, which can be used

for x-ray analysis, or a magnetic contrast agent, such as a gadolinium chelate, which can be used for MRI or CT. Other labeling agents include, without limitation, radioisotopes, such as ^{99}Tc. In another embodiment, the P-cadherin antibody will be unlabeled and will be imaged by administering a second antibody or other molecule that is detectable and that can bind the P-cadherin antibody. In one embodiment, a biopsy is obtained from the patient to determine whether the tissue of interest expresses P-cadherin.

Therapeutic Methods of Use

In another embodiment, the invention provides a method for inhibiting P-cadherin activity by administering a P-cadherin antibody to a patient in need thereof. Any of the antibodies or antigen-binding portions thereof described herein may be used therapeutically. In a preferred embodiment, the P-cadherin antibody is a human, chimeric or humanized antibody. In another preferred embodiment, the P-cadherin is human and the patient is a human patient. Alternatively, the patient may be a mammal that expresses a P-cadherin that the P-cadherin antibody cross-reacts with. The antibody may be administered to a non-human mammal expressing P-cadherin with which the antibody cross-reacts (e.g. a rat, a mouse, or a cynomolgus monkey) for veterinary purposes or as an animal model of human disease. Such animal models may be useful for evaluating the therapeutic efficacy of antibodies of this invention.

In another embodiment, a P-cadherin antibody or antibody portion thereof may be administered to a patient who expresses inappropriately high levels of P-cadherin. The antibody may be administered once, but more preferably is administered multiple times. The antibody may be administered from three times daily to once every six months or longer. The administering may be on a schedule such as three times daily, twice daily, once daily, once every two days, once every three days, once weekly, once every two weeks, once every month, once every two months, once every three months and once every six months. The antibody may also be administered continuously via a minipump. The antibody may be administered via a mucosal, buccal, intranasal, inhalable, intravenous, subcutaneous, intramuscular, parenteral, or intratumor route. The antibody may be administered once, at least twice or for at least the period of time until the condition is treated, palliated or cured. The antibody generally will be administered for as long as the condition is present. The antibody will generally be administered as part of a pharmaceutical composition as described *supra*. The dosage of antibody will generally be in the range of 0.1 to 100 mg/kg, more preferably 0.5 to 50 mg/kg, more preferably 1 to 20 mg/kg, and even more preferably 1 to 10 mg/kg. The serum concentration of the antibody may be measured by any method known in the art.

This invention also relates to a method for the treatment of abnormal cell growth in a mammal, including a human, comprising administering to said mammal a therapeutically effective amount of a P-cadherin antibody or antigen binding portion thereof, as described herein, that is effective in treating abnormal cell growth.

In one embodiment of this method, the abnormal cell growth is cancer, including, but not limited to, mesothelioma, hepatobilliary (hepatic and billiary duct), a primary or secondary CNS tumor, a primary or secondary brain tumor, lung cancer (NSCLC and SCLC), bone cancer,

pancreatic cancer, skin cancer, cancer of the head or neck, cutaneous or intraocular melanoma, ovarian cancer, colon cancer, rectal cancer, cancer of the anal region, stomach cancer, gastrointestinal (gastric, colorectal, and duodenal), breast cancer, uterine cancer, carcinoma of the fallopian tubes, carcinoma of the endometrium, carcinoma of the cervix, carcinoma of the vagina, carcinoma of the vulva, Hodgkin's Disease, cancer of the esophagus, cancer of the small intestine, cancer of the endocrine system, cancer of the thyroid gland, cancer of the parathyroid gland, cancer of the adrenal gland, sarcoma of soft tissue, cancer of the urethra, cancer of the penis, prostate cancer, testicular cancer, chronic or acute leukemia, chronic myeloid leukemia, lymphocytic lymphomas, cancer of the bladder, cancer of the kidney or ureter, renal cell carcinoma, carcinoma of the renal pelvis, neoplasms of the central nervous system (CNS), primary CNS lymphoma, non hodgkins's lymphoma, spinal axis tumors, brain stem glioma, pituitary adenoma, adrenocortical cancer, gall bladder cancer, multiple myeloma, cholangiocarcinoma, fibrosarcoma, neuroblastoma, retinoblastoma, or a combination of one or more of the foregoing cancers.

In a preferred embodiment of the present invention the cancer is selected from lung cancer (NSCLC and SCLC), cancer of the head or neck, ovarian cancer, colon cancer, rectal cancer, cancer of the anal region, stomach cancer, breast cancer, cancer of the kidney or ureter, renal cell carcinoma, carcinoma of the renal pelvis, neoplasms of the central nervous system (CNS), primary CNS lymphoma, non hodgkins's lymphoma, spinal axis tumors, or a combination of one or more of the foregoing cancers.

In another preferred embodiment of the present invention the cancer is selected from lung cancer (NSCLC and SCLC), ovarian cancer, colon cancer, rectal cancer, cancer of the anal region, or a combination of one or more of the foregoing cancers.

In another embodiment of said method, said abnormal cell growth is a benign proliferative disease, including, but not limited to, psoriasis, benign prostatic hypertrophy or restinosis.

This invention also relates to a method for the treatment of abnormal cell growth in a mammal which comprises administering to said mammal an amount of a P-cadherin antibody or antigen binding portion thereof, as described herein, that is effective in treating abnormal cell growth in combination with an anti-tumor agent selected from the group consisting of mitotic inhibitors, alkylating agents, anti-metabolites, intercalating antibiotics, growth factor inhibitors, cell cycle inhibitors, enzymes, topoisomerase inhibitors, biological response modifiers, antibodies, cytotoxics, anti-hormones, and anti-androgens.

The invention also relates to a pharmaceutical composition for the treatment of abnormal cell growth in a mammal, including a human, which comprises an amount of a P-cadherin antibody or antigen binding portion thereof, as described herein, that is effective in treating abnormal cell growth in combination with a pharmaceutically acceptable carrier and an anti-tumor agent selected from the group consisting of mitotic inhibitors, alkylating agents, anti-metabolites, intercalating antibiotics, growth factor inhibitors, cell cycle inhibitors, enzymes, topoisomerase inhibitors, biological response modifiers, anti-hormones, and anti-androgens.

The invention also relates to a method for the treatment of a hyperproliferative disorder in a mammal which comprises administering to said mammal a therapeutically effective amount of a P-cadherin antibody or antigen binding portion thereof, as described herein, in combination with an anti-tumor agent selected from the group consisting antiproliferative agents, kinase inhibitors, angiogenesis inhibitors, growth factor inhibitors, cox-I inhibitors, cox-II inhibitors, mitotic inhibitors, alkylating agents, anti-metabolites, intercalating antibiotics, growth factor inhibitors, radiation, cell cycle inhibitors, enzymes, topoisomerase inhibitors, biological response modifiers, antibodies, cytotoxics, anti-hormones, statins, and anti-androgens.

In one embodiment of the present invention the anti-tumor agent used in conjunction with a P-cadherin antibody or antigen binding portion thereof, and pharmaceutical compositions described herein, is an anti-angiogenesis agent, kinase inhibitor, pan kinase inhibitor or growth factor inhibitor. Preferred pan kinase inhibitors include SU-11248, described in U.S. Patent No. 6,573,293 (Pfizer, Inc, NY, USA).

Anti-angiogenesis agents, include but are not limited to the following agents, such as EGF inhibitor, EGFR inhibitors, VEGF inhibitors, VEGFR inhibitors, TIE2 inhibitors, IGF1R inhibitors, COX-II (cyclooxygenase II) inhibitors, MMP-2 (matrix-metalloprotienase 2) inhibitors, and MMP-9 (matrix-metalloprotienase 9) inhibitors. Preferred VEGF inhibitors, include for example, Avastin (bevacizumab), an anti-VEGF monoclonal antibody of Genentech, Inc. of South San Francisco, California.

Additional VEGF inhibitors include CP-547,632 (Pfizer Inc., NY, USA), AG13736 (Pfizer Inc.), ZD-6474 (AstraZeneca), AEE788 (Novartis), AZD-2171), VEGF Trap (Regeneron,/Aventis), Vatalanib (also known as PTK-787, ZK-222584: Novartis & Schering AG), Macugen (pegaptanib octasodium, NX-1838, EYE-001, Pfizer Inc./Gilead/Eyetech), IM862 (Cytran Inc. of Kirkland, Washington, USA); and angiozyme, a synthetic ribozyme from Ribozyme (Boulder, Colorado) and Chiron (Emeryville, California) and combinations thereof. VEGF inhibitors useful in the practice of the present invention are disclosed in US Patent No. 6,534,524 and 6,235,764, both of which are incorporated in their entirety for all purposed. Particularly preferred VEGF inhibitors include CP-547,632, AG13736, Vatalanib, Macugen and combinations thereof.

Additional VEGF inhibitors are described in, for example in WO 99/24440 (published May 20, 1999), PCT International Application PCT/IB99/00797 (filed May 3, 1999), in WO 95/21613 (published August 17, 1995), WO 99/61422 (published December 2, 1999), United States Patent 6, 534,524 (discloses AG13736), United States Patent 5,834,504 (issued November 10, 1998), WO 98/50356 (published November 12, 1998), United States Patent 5,883,113 (issued March 16, 1999), United States Patent 5,886,020 (issued March 23, 1999), United States Patent 5,792,783 (issued August 11, 1998), U.S. Patent No. US 6,653,308 (issued November 25, 2003), WO 99/10349 (published March 4, 1999), WO 97/32856 (published September 12, 1997), WO 97/22596 (published June 26, 1997), WO 98/54093 (published December 3, 1998), WO 98/02438 (published January 22, 1998), WO 99/16755 (published April 8, 1999), and WO 98/02437 (published January 22, 1998), all of which are herein incorporated by reference in their entirety.

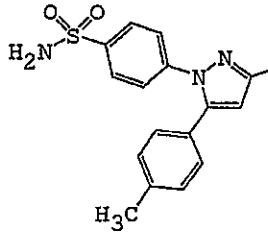
Other antiproliferative agents that may be used with the antibodies, or antigen-binding portions thereof, of the present invention include inhibitors of the enzyme farnesyl protein transferase and inhibitors of the receptor tyrosine kinase PDGFr, including the compounds disclosed and claimed in the following United States patent applications: 09/221946 (filed December 28, 1998); 09/454058 (filed December 2, 1999); 09/501163 (filed February 9, 2000); 09/539930 (filed March 31, 2000); 09/202796 (filed May 22, 1997); 09/384339 (filed August 26, 1999); and 09/383755 (filed August 26, 1999); and the compounds disclosed and claimed in the following United States provisional patent applications: 60/168207 (filed November 30, 1999); 60/170119 (filed December 10, 1999); 60/177718 (filed January 21, 2000); 60/168217 (filed November 30, 1999), and 60/200834 (filed May 1, 2000). Each of the foregoing patent applications and provisional patent applications is herein incorporated by reference in their entirety.

PDGRr inhibitors include but are not limited to those disclosed in WO01/40217, published July 7, 2001 and WO2004/020431, published March 11, 2004, the contents of which are incorporated in their entirety for all purposes. Preferred PDGFr inhibitors include Pfizer's CP-673,451 and CP-868,596 and its pharmaceutically acceptable salts.

Preferred GARF inhibitors include Pfizer's AG-2037 (pelitrexol and its pharmaceutically acceptable salts). GARF inhibitors useful in the practice of the present invention are disclosed in U.S. Patent No. 5,608,082 which is incorporated in its entirety for all purposes.

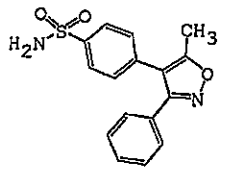
Examples of useful COX-II inhibitors which can be used in conjunction with a P-cadherin antibody or antigen binding portion thereof, as described herein, and pharmaceutical compositions described herein include CELEBREX™ (celecoxib), parecoxib, deracoxib, ABT-963, MK-663 (etoricoxib), COX-189 (Lumiracoxib), BMS 347070, RS 57067, NS-398, Bextra (valdecoxib), paracoxib, Vioxx (rofecoxib), SD-8381, 4-Methyl-2-(3,4-dimethylphenyl)-1-(4-sulfamoyl-phenyl)-1H-pyrrole, 2-(4-Ethoxyphenyl)-4-methyl-1-(4-sulfamoylphenyl)-1H-pyrrole, T-614, JTE-522, S-2474, SVT-2016, CT-3, SC-58125 and Arcoxia (etoricoxib). Additonally, COX-II inhibitors are disclosed in U.S. Patent Application Nos. 10/801,446 and 10/801,429, the contents of which are incorporated in their entirety for all purposes.

In one preferred embodiment the anti-tumor agent is celecoxib as disclosed in U.S. Patent No. 5,466,823, the contents of which are incorporated by reference in its entirety for all purposes. The structure for Celecoxib is shown below:



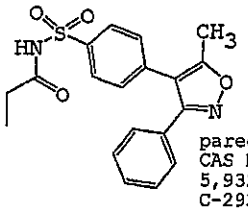
celecoxib
CAS No. 169590-42-5
5,466,823
C-2779
SC-58635

In one preferred embodiment the anti-tumor agent is valecoxib as disclosed in U.S. Patent No. 5,633,272, the contents of which are incorporated by reference in its entirety for all purposes. The structure for valdecoxib is shown below:



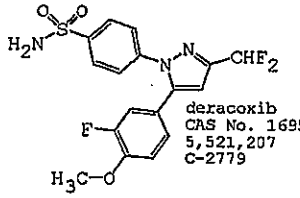
valdecoxib
CAS No. 181695-72-7
5,633,272
C-2865
SC-65872

In one preferred embodiment the anti-tumor agent is parecoxib as disclosed in U.S. Patent No. 5,932,598, the contents of which are incorporated by reference in its entirety for all purposes. The structure for paracoxib is shown below:



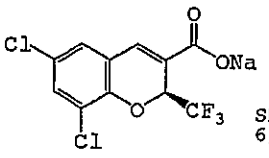
parecoxib
CAS No. 198470-84-7
5,932,598
C-2931

In one preferred embodiment the anti-tumor agent is deracoxib as disclosed in U.S. Patent No. 5,521,207, the contents of which are incorporated by reference in its entirety for all purposes. The structure for deracoxib is shown below:



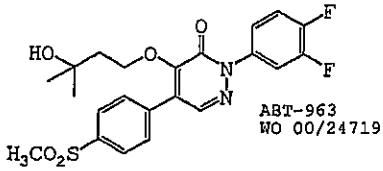
deracoxib
CAS No. 169590-41-4
5,521,207
C-2779

In one preferred embodiment the anti-tumor agent is SD-8381 as disclosed in U.S. Patent No. 6,034,256, the contents of which are incorporated by reference in its entirety for all purposes. The structure for SD-8381 is shown below:



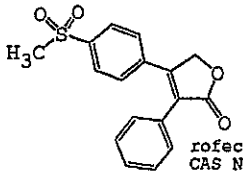
SD-8381
6,034,256
Ex. 175

In one preferred embodiment the anti-tumor agent is ABT-963 as disclosed in International Publication Number WO 2002/24719, the contents of which are incorporated by reference in its entirety for all purposes. The structure for ABT-963 is shown below:



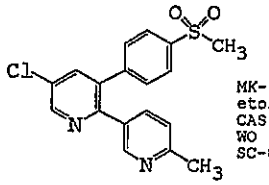
ABT-963
WO 00/24719

In one preferred embodiment the anti-tumor agent is rofecoxib as shown below:



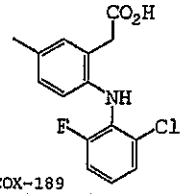
rofecoxib
CAS No. 162011-90-7

In one preferred embodiment the anti-tumor agent is MK-663 (etoricoxib) as disclosed in International Publication Number WO 1998/03484, the contents of which are incorporated by reference in its entirety for all purposes. The structure for etoricoxib is shown below:



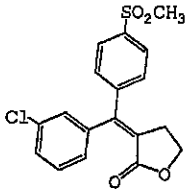
MK-663
etoricoxib
CAS No. 202409-33-4
WO 98/03484
SC-86218

In one preferred embodiment the anti-tumor agent is COX-189 (Lumiracoxib) as disclosed in International Publication Number WO 1999/11605, the contents of which are incorporated by reference in its entirety for all purposes. The structure for Lumiracoxib is shown below:



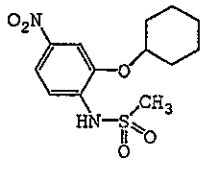
COX-189
Lumiracoxib
CAS No. 220991-20-8
Novartis
WO 99/11605

In one preferred embodiment the anti-tumor agent is BMS-347070 as disclosed in United States Patent No. 6,180,651, the contents of which are incorporated by reference in its entirety for all purposes. The structure for BMS-347070 is shown below:



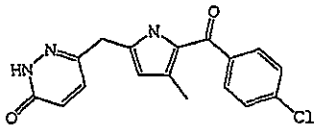
BMS 347070
CAS No. 197438-48-5
6,180,651

In one preferred embodiment the anti-tumor agent is NS-398 (CAS 123653-11-2). The structure for NS-398 (CAS 123653-11-2) is shown below:



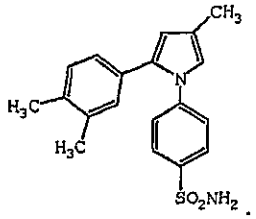
NS-398
CAS No. 123653-11-2

In one preferred embodiment the anti-tumor agent is RS 57067 (CAS 17932-91-3). The structure for RS-57067 (CAS 17932-91-3) is shown below:

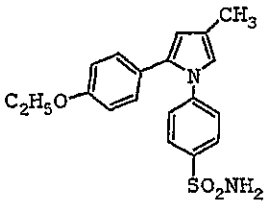


RS 57067
CAS No. 17932-91-3

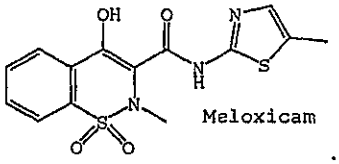
In one preferred embodiment the anti-tumor agent is 4-Methyl-2-(3,4-dimethylphenyl)-1-(4-sulfamoyl-phenyl)-1H-pyrrole. The structure for 4-Methyl-2-(3,4-dimethylphenyl)-1-(4-sulfamoyl-phenyl)-1H-pyrrole is shown below:



In one preferred embodiment the anti-tumor agent is 2-(4-Ethoxyphenyl)-4-methyl-1-(4-sulfamoylphenyl)-1H-pyrrole. The structure for 2-(4-Ethoxyphenyl)-4-methyl-1-(4-sulfamoylphenyl)-1H-pyrrole is shown below:



In one preferred embodiment the anti-tumor agent is meloxicam. The structure for meloxicam is shown below:



Meloxicam

Other useful inhibitors as anti-tumor agents used in conjunction with antibodies of the present invention and pharmaceutical compositions described herein include aspirin, and non-steroidal anti-inflammatory drugs (NSAIDs) which inhibit the enzyme that makes prostaglandins (cyclooxygenase I and II), resulting in lower levels of prostaglandins, include but are not limited to the following, Salsalate (Amigesic), Diflunisal (Dolobid), Ibuprofen (Motrin), Ketoprofen (Orudis), Nabumetone (Relafen), Piroxicam (Feldene), Naproxen (Aleve, Naprosyn), Diclofenac (Voltaren), Indomethacin (Indocin), Sulindac (Clinoril), Tolmetin (Tolectin), Etodolac (Lodine), Ketorolac (Toradol), Oxaprozin (Daypro) and combinations thereof. Preferred COX-I inhibitors include ibuprofen (Motrin), nuprin, naproxen (Aleve), indomethacin (Indocin), nabumetone (Relafen) and combinations thereof.

Targeted agents used in conjunction with a P-cadherin antibody or antigen binding portion thereof, as described herein, and pharmaceutical compositions thereof as described herein, include EGFr inhibitors such as Iressa (gefitinib, AstraZeneca), Tarceva (erlotinib or OSI-774, OSI Pharmaceuticals Inc.), Erbitux (cetuximab, Imclone Pharmaceuticals, Inc.), EMD-7200 (Merck AG), ABX-EGF (Amgen Inc. and Abgenix Inc.), HR3 (Cuban Government), IgA antibodies (University of Erlangen-Nuremberg), TP-38 (IVAX), EGFR fusion protein, EGF-vaccine, anti-EGFr immunoliposomes (Hermes Biosciences Inc.) and combinations thereof.

Preferred EGFr inhibitors include Iressa, Erbitux, Tarceva and combinations thereof.

The present invention also relates to anti-tumor agents selected from pan erb receptor inhibitors or ErbB2 receptor inhibitors, such as CP-724,714 (Pfizer, Inc.), CI-1033 (canertinib, Pfizer, Inc.), Herceptin (trastuzumab, Genentech Inc.), Omitarg (2C4, pertuzumab, Genentech Inc.), TAK-165 (Takeda), GW-572016 (Ionafarnib, GlaxoSmithKline), GW-282974 (GlaxoSmithKline), EKB-569 (Wyeth), PKI-166 (Novartis), dHER2 (HER2 Vaccine, Corixa and GlaxoSmithKline), APC8024 (HER2 Vaccine, Dendreon), anti-HER2/neu bispecific antibody (Decof Cancer Center), B7.her2.IgG3 (Agensys), AS HER2 (Research Institute for Rad Biology & Medicine), trifuntional bispecific antibodies (University of Munich) and mAB AR-209 (Aronex Pharmaceuticals Inc) and mAB 2B-1 (Chiron) and combinations thereof. Preferred erb selective anti-tumor agents include Herceptin, TAK-165, CP-724,714, ABX-EGF, HER3 and combinations thereof. Preferred pan erbb receptor inhibitors include GW572016, CI-1033, EKB-569, and Omitarg and combinations thereof.

Additional erbB2 inhibitors include those described in WO 98/02434 (published January 22, 1998), WO 99/35146 (published July 15, 1999), WO 99/35132 (published July 15, 1999), WO 98/02437 (published January 22, 1998), WO 97/13760 (published April 17, 1997), WO 95/19970 (published July 27, 1995), United States Patent 5,587,458 (issued December 24, 1996), and United States Patent 5,877,305 (issued March 2, 1999), each of which is herein incorporated by reference in its entirety. ErbB2 receptor inhibitors useful in the present invention are also described in United States Patent Nos. 6,465,449, and 6,284,764, and International Application No. WO 2001/98277 each of which are herein incorporated by reference in their entirety.

Additionally, other anti-tumor agents may be selected from the following agents, BAY-43-9006 (Onyx Pharmaceuticals Inc.), Genasense (augmerosen, Genta), Panitumumab (Abgenix/Amgen), Zevalin (Schering), Bexxar (Corixa/GlaxoSmithKline), Abarelix, Alimta, EPO 906 (Novartis), discodermolide (XAA-296), ABT-510 (Abbott), Neovastat (Aeterna), enzastaurin (Eli Lilly), Combrestatin A4P (Oxigene), ZD-6126 (AstraZeneca), flavopiridol (Aventis), CYC-202 (Cyclacel), AVE-8062 (Aventis), DMXAA (Roche/Antisoma), Thymitaq (Eximias), Temodar (temozolomide, Schering Plough) and Revilimd (Celegene) and combinations thereof.

Other anti-tumor agents may be selected from the following agents, CyPat (cyproterone acetate), Histerelin (histrelin acetate), Plenaixis (abarelix depot), Atrasentan (ABT-627), Satraplatin (JM-216), thalomid (Thalidomide), Theratope, Temilifene (DPPE), ABI-007 (paclitaxel), Evista (raloxifene), Atamestane (Biomed-777), Xyotax (polyglutamate paclitaxel), Targetin (bexarotine) and combinations thereof.

Additionally, other anti-tumor agents may be selected from the following agents, Trizaone (tirapazamine), Aposyn (exisulind), Nevastat (AE-941), Ceplene (histamine dihydrochloride), Orathecin (rubitecan), Virulizin, Gastrimmune (G17DT), DX-8951f (exatecan mesylate), Onconase (ranpirnase), BEC2 (mitumoab), Xcytrin (motexafin gadolinium) and combinations thereof.

Further anti-tumor agents may selected from the following agents, CeaVac (CEA), NeuTrexin (trimetresate glucuronate) and combinations thereof. Additional anti-tumor agents may selected from the following agents, OvaRex (oregovomab), Osidem (IDM-1), and combinations thereof. Additional anti-tumor agents may be selected from the following agents, Advexin (ING 201), Tirazone (tirapazamine), and combinations thereof. Additional anti-tumor agents may selected from the following agents, RSR13 (efaproxiral), Cotara (131i chTNT 1/b), NBI-3001 (IL-4) and combinations thereof. Additional anti-tumor agents may be selected from the following agents, Canvaxin, GMK vaccine, PEG Interon A, Taxoprexin (DHA/paclitaxel) and combinations thereof. Other preferred anti-tumor agents include Pfizer's MEK1/2 inhibitor PD325901, Array Biopharm's MEK inhibitor ARRY-142886, Bristol Myers' CDK2 inhibitor BMS-387,032, Pfizer's CDK inhibitor PD0332991 and AstraZeneca's AXD-5438 and combinations thereof. Additionally, mTOR inhibitors may also be utilized such as CCI-779 (Wyeth) and rapamycin derivatives RAD001 (Novartis) and AP-23573 (Ariad), HDAC inhibitors SAHA (Merck Inc./Aton Pharmaceuticals) and combinations thereof. Additional anti-tumor agents include aurora 2 inhibitor VX-680 (Vertex), Chk1/2 inhibitor XL844 (Exilixis).

The following cytotoxic agents, e.g., one or more selected from the group consisting of epirubicin (Ellence), docetaxel (Taxotere), paclitaxel, Zinecard (dexrazoxane), rituximab (Rituxan) imatinib mesylate (Gleevec), and combinations thereof, may be used in conjunction with a P-cadherin antibody or antigen binding portion thereof, as described herein, and pharmaceutical compositions thereof , as described herein.

The invention also contemplates the use of the antibodies and antigen-binding portions thereof of the present invention together with hormonal therapy, including but not limited to, exemestane (Aromasin, Pfizer Inc.), leuprorelin (Lupron or Leuplin, TAP/Abbott/Takeda), anastrozole (Arimidex, Astrazeneca), gosrelin (Zoladex, AstraZeneca), doxercalciferol, fadrozole, formestane, tamoxifen citrate (tamoxifen, Nolvadex, AstraZeneca), Casodex (AstraZeneca), Abarelix (Praecis), Trelstar, and combinations thereof.

The invention also relates to hormonal therapy agents such as anti-estrogens including, but not limited to fulvestrant, toremifene, raloxifene, lasofoxifene, letrozole (Femara, Novartis), anti-androgens such as bicalutamide, flutamide, mifepristone, nilutamide, Casodex®(4'-cyano-3-(4-fluorophenylsulphonyl)-2-hydroxy-2-methyl-3'-(trifluoromethyl) propionanilide, bicalutamide) and combinations thereof.

Further, the invention provides antibodies of the present invention alone or in combination with one or more supportive care products, e.g., a product selected from the group consisting of Filgrastim (Neupogen), ondansetron (Zofran), Fragmin, Procrit, Aloxi, Emend, or combinations thereof.

Particularly preferred cytotoxic agents include Camptosar, Erbitux, Iressa, Gleevec, Taxotere and combinations thereof.

The following topoisomerase I inhibitors may be utilized as anti-tumor agents camptothecin, irinotecan HCl (Camptosar), edotecarin, orathecin (Supergen), exatecan (Daiichi), BN-80915 (Roche) and combinations thereof. Particularly preferred toposimerase II inhibitors include epirubicin (Ellence).

The antibodies of the invention may be used with antitumor agents, alkylating agents, antimetabolites, antibiotics, plant-derived antitumor agents, camptothecin derivatives, tyrosine kinase inhibitors, other antibodies, interferons, and/or biological response modifiers.

Alkylating agents include, but are not limited to, nitrogen mustard N-oxide, cyclophosphamide, ifosfamide, melphalan, busulfan, mitobronitol, carboquone, thiotepa, ranimustine, nimustine, temozolomide, AMD-473, altretamine, AP-5280, apaziquone, brostallicin, bendamustine, carmustine, estramustine, fotemustine, glufosfamide, ifosfamide, KW-2170, mafosfamide, and mitolactol; platinum-coordinated alkylating compounds include but are not limited to, cisplatin, Paraplatin (carboplatin), eptaplatin, lobaplatin, nedaplatin, Eloxatin (oxaliplatin, Sanofi) or satrplatin and combinations thereof. Particularly preferred alkylating agents include Eloxatin (oxaliplatin).

Antimetabolites include but are not limited to, methotrexate, 6-mercaptopurine riboside, mercaptopurine, 5-fluorouracil (5-FU) alone or in combination with leucovorin, tegafur, UFT, doxifluridine, carmofur, cytarabine, cytarabine ocfosfate, enocitabine, S-1, Alimta (premetrexed disodium, LY231514, MTA), Gemzar (gemcitabine, Eli Lilly), fludarabin, 5-azacitidine, capecitabine, cladribine, clofarabine, decitabine, eflornithine, ethynylcytidine, cytosine arabinoside, hydroxyurea, TS-1, melphalan, nelarabine, nolatrexed, ocfosfate, disodium premetrexed, pentostatin, pelitrexol, raltitrexed, triapine, trimetrexate, vidarabine, vincristine, vinorelbine; or for example, one of the preferred anti-metabolites disclosed in European Patent Application No. 239362 such as N-(5-[N-(3,4-dihydro-2-methyl-4-oxoquinazolin-6-ylmethyl)-N-methylamino]-2-thenoyl)-L-glutamic acid and combinations thereof.

Antibiotics include intercalating antibiotics but are not limited to: aclarubicin, actinomycin D, amrubicin, annamycin, adriamycin, bleomycin, daunorubicin, doxorubicin, elsamitrucin, epirubicin, galarubicin, idarubicin, mitomycin C, nemorubicin, neocarzinostatin, peplomycin, pirarubicin, rebeccamycin, stimalamer, streptozocin, valrubicin, zinostatin and combinations thereof.

Plant derived anti-tumor substances include for example those selected from mitotic inhibitors, for example vinblastine, docetaxel (Taxotere), paclitaxel and combinations thereof.

Cytotoxic topoisomerase inhibiting agents include one or more agents selected from the group consisting of aclarubicn, amonafide, belotecan, camptothecin, 10-hydroxycamptothecin, 9-aminocamptothecin, diflomotecan, irinotecan HCl (Camptosar), edotecarin, epirubicin (Ellence), etoposide, exatecan, gimatecan, lurtotecan, mitoxantrone, pirarubicin, pixantrone, rubitecan, sobuzoxane, SN-38, tafluposide, topotecan, and combinations thereof.

Preferred cytotoxic topoisomerase inhibiting agents include one or more agents selected from the group consisting of camptothecin, 10-hydroxycamptothecin, 9-aminocamptothecin, irinotecan HCl (Camptosar), edotecarin, epirubicin (Ellence), etoposide, SN-38, topotecan, and combinations thereof.

Immunologicals include interferons and numerous other immune enhancing agents. Interferons include interferon alpha, interferon alpha-2a, interferon, alpha-2b, interferon beta, interferon gamma-1a, interferon gamma-1b (Actimmune), or interferon gamma-n1 and combinations thereof. Other agents include filgrastim, lentinan, sizofilan, TheraCys, ubenimex, WF-10, aldesleukin, alemtuzumab, BAM-002, dacarbazine, daclizumab, denileukin, gemtuzumab ozogamicin, ibritumomab, imiquimod, lenograstim, lentinan, melanoma vaccine (Corixa), molgramostim, OncoVAX-CL, sargramostim, tasonermin, tecleukin, thymalasin, tositumomab, Virulizin, Z-100, epratuzumab, mitumomab, oregovomab, pemtumomab (Y-muHMFG1), Provenge (Dendreon) and combinations thereof.

Biological response modifiers are agents that modify defense mechanisms of living organisms or biological responses, such as survival, growth, or differentiation of tissue cells to direct them to have anti-tumor activity. Such agents include krestin, lentinan, sizofiran, picibanil, ubenimex and combinations thereof.

Other anticancer agents include alitretinoin, ampligen, atrasentan bexarotene, bortezomib. Bosentan, calcitriol, exisulind, finasteride,fotemustine, ibandronic acid, miltefosine, mitoxantrone, l-asparaginase, procarbazine, dacarbazine, hydroxycarbamide, pegaspargase, pentostatin, tazarotne, Telcyta (TLK-286, Telik Inc.), Velcade (bortemazib, Millenium), tretinoin, and combinations thereof.

Other anti-angiogenic compounds include acitretin, fenretinide, thalidomide, zoledronic acid, angiostatin, aplidine, cilengtide, combretastatin A-4, endostatin, halofuginone, rebimastat, removab, Revlimid, squalamine, ukrain, Vitaxin and combinations thereof.

Platinum-coordinated compounds include but are not limited to, cisplatin, carboplatin, nedaplatin, oxaliplatin, and combinations thereof.

Camptothecin derivatives include but are not limited to camptothecin, 10-hydroxycamptothecin, 9-aminocamptothecin, irinotecan, SN-38, edotecarin, topotecan and combinations thereof.

Other antitumor agents include mitoxantrone, l-asparaginase, procarbazine, dacarbazine, hydroxycarbamide, pentostatin, tretinoin and combinations thereof.

Anti-tumor agents capable of enhancing antitumor immune responses, such as CTLA4 (cytotoxic lymphocyte antigen 4) antibodies, and other agents capable of blocking CTLA4 may also be utilized, such as MDX-010 (Medarex) and CTLA4 compounds disclosed in United States Patent No. 6,682,736; and anti-proliferative agents such as other farnesyl protein transferase inhibitors, for example the farnesyl protein transferase inhibitors. Additional, specific CTLA4 antibodies that can be used in the present invention include those described in United States Provisional Application 60/113,647 (filed December 23, 1998), United States Patent No. 6, 682,736 both of which are herein incorporated by reference in their entirety.

Specific IGF1R antibodies that can be used in the present invention include those described in International Patent Application No. WO 2002/053596, which is herein incorporated by reference in its entirety.

Specific CD40 antibodies that can be used in the present invention include those described in International Patent Application No. WO 2003/040170 which is herein incorporated by reference in its entirety.

Gene therapy agents may also be employed as anti-tumor agents such as TNFerade (GeneVec), which express TNFalpha in response to radiotherapy.

In one embodiment of the present invention statins may be used in conjunction with a P-cadherin antibody or antigen binding portion thereof, as described herein, and pharmaceutical compositions thereof. Statins (HMG-CoA reducatase inhibitors) may be selected from the group consisting of Atorvastatin (Lipitor, Pfizer Inc.), Provastatin (Pravachol, Bristol-Myers Squibb), Lovastatin (Mevacor, Merck Inc.), Simvastatin (Zocor, Merck Inc.), Fluvastatin (Lescol, Novartis), Cerivastatin (Baycol, Bayer), Rosuvastatin (Crestor, AstraZeneca), Lovostatin and Niacin (Advicor, Kos Pharmaceuticals), derivatives and combinations thereof.

In a preferred embodiment the statin is selected from the group consisting of Atovorstatin and Lovastatin, derivatives and combinations thereof.

Other agents useful as anti-tumor agents include Caduet.

For any of the methods of treating a hyperproliferative disorder or abnormal cell growth as described herein using a combination of a P-cadherin antibody or antigen binding portion with at least one additional therapeutic agent, the P-cadherin antibody can be conjugated, or derivatized, with the additional therapeutic agent. The at least one additional therapeutic agent can also be administered separately, or in a non-derivatized or non-conjugated manner. When the at least one additional therapeutic agent is not derivatized or conjugated to the antibody, it can be administered within the same pharmaceutical formulation as the antibody, or it can be administered in a separate formulation.

Gene Therapy

The nucleic acid molecules that encode the antibodies and antibody portions of the present invention can be administered to a patient in need thereof via gene therapy. The therapy may be either *in vivo* or *ex vivo*. In a preferred embodiment, nucleic acid molecules encoding both a heavy chain and a light chain are administered to a patient. In a more preferred embodiment, the nucleic acid molecules are administered such that they are stably integrated into chromosomes of B cells because these cells are specialized for producing antibodies. In a preferred embodiment, precursor B cells are transfected or infected *ex vivo* and re-transplanted into a patient in need thereof. In another embodiment, precursor B cells or other cells are infected *in vivo* using a virus known to infect the cell type of interest. Typical vectors used for gene therapy include liposomes, plasmids, and viral vectors. Exemplary viral vectors are retroviruses, adenoviruses and adeno-associated viruses. After infection either *in vivo* or *ex vivo*, levels of antibody expression can be monitored by taking a sample from the treated patient and using any immunoassay known in the art or discussed herein.

In a preferred embodiment, the gene therapy method comprises the steps of administering an isolated nucleic acid molecule encoding the heavy chain or an antigen-binding portion thereof of a P-cadherin antibody and expressing the nucleic acid molecule. In another embodiment, the gene

therapy method comprises the steps of administering an isolated nucleic acid molecule encoding the light chain or an antigen-binding portion thereof of a P-cadherin antibody and expressing the nucleic acid molecule. In a more preferred method, the gene therapy method comprises the steps of administering an isolated nucleic acid molecule encoding the heavy chain or an antigen-binding portion thereof and an isolated nucleic acid molecule encoding the light chain or the antigen-binding portion thereof of a P-cadherin antibody of the invention and expressing the nucleic acid molecules. The gene therapy method may also comprise the step of administering another therapeutic agent, such as any of the agents discussed previously in connection with combination therapy.

In order that this invention may be better understood, the following examples are set forth. These examples are for purposes of illustration only and are not to be construed as limiting the scope of the invention in any manner.

Examples

In the following examples and preparations, "BSA" means bovine serum albumin; "EDTA" means ethylenediaminetetraacetic acid; "DMSO" means dimethyl sulfoxide; "MOPS" means 3-(N-morpholino) propanesulfonic acid; "MES" means 2-(N-Morpholino)ethanesulfonic acid; "PBS" means phosphate buffered saline; "dPBS" means Dulbecco's phosphate buffered saline; "HEMA" means 2-hydroxy-ethyl methacrylate; "DMEM" means Dulbecco's modified eagle's medium; "FBS" means fetal bovine serum; "NEAA" means non-essential amino acids; "HEPES" means N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid; and "DMF" means dimethyl formamide.

Example 1: Screening a scFv Phage Display Library

Recombinant human P-cadherin (R&D Systems 861-PC-100) was used as the antigen to screen a scFv phage display library. Large scFv human *antibody libraries cloned into a phagemid vector were used for selections (Vaughan, T.J. et al., Nat.* Biotech. 14:309-314 (1996)). ScFv which recognized P-cadherin were isolated from phage display libraries in a series of repeated selection cycles on recombinant human P-cadherin and confluent monolayers of HCT116 cells expressing P-cadherin. In brief, following incubation with the library, bound phage were recovered from P-cadherin and unbound phage were washed away. Bound phage were then rescued as described in Vaughan, T.J. et al., *Nat. Biotech.* 14:309-314 (1996) and the selection process was repeated. A representative proportion of clones from the output of selection rounds were subjected to phage enzyme-linked immunosorbent assay (ELISA) to test for binding to P-cadherin, essentially as described in Vaughan, T.J. et al., *Nat. Biotech.* 14:309-314 (1996). Two different antigens were used in the ELISA: recombinant human P-cadherin (R&D Systems) and confluent monolayers of A431 cells expressing P-cadherin. ELISA-positive clones were subjected to DNA sequencing as decribed in Vaughan, T.J. et al., *Nat. Biotech.* 14:309-314 (1996) and in Osbourn, J.K. et al., *Immunotechnology* 3:293-302 (1998). Unique ELISA-positive clones were converted to whole IgG molecules and tested for their ability to neutralize P-cadherin in the P-cadherin dependent adhesion assay described in Example 4. Based on the results of this screening, the antibody 129-1c4 (IC50 of 1-3 μM in the A431 adhesion assay) was selected as the lead parent lineage for further optimization.

Example 2: Lead Optimization

Phage display libraries derived from 129-1c4 were created by oligonucleotide-directed mutagenesis of antibody variable heavy (V_H) and variable light (V_L) chain CDR3 regions. The libraries were constructed using standard molecular biology techniques as described in Clackson and Lowman, Phage Display – A Practical Approach (Oxford University Press 2004). Affinity-based selections were performed whereby; following incubation with the library, the recombinant human P-cadherin (R&D Systems) was captured by protein G-coated paramagnetic beads (Dynal 100.03) and bound phage were recovered by magnetic separation while unbound complexes were washed away. The selection process was repeated with decreasing concentrations of recombinant human P-cadherin (25 nM to 10 pM over 4 rounds) present during the selection. In addition, selection outputs from the V_H CDR3 and V_L CDR3 libraries were recombined into further phage display libraries and selected over two more rounds of affinity-based selection. Representative proportions of clones from the outputs of selection rounds were subjected to screening as scFv in the 129-1c4 epitope competition assay, as described in Example 8.

Example 3: P-cadherin-dependent adhesion assay

The following protocol was used to determine IC_{50} values in a P-cadherin-dependent adhesion assay using several optimized scFvs that were converted to IgG during the optimization phase of antibody discovery as described above in Example 2. The mean measured IC_{50} values for these antibodies are shown below in Table 4.

Recombinant human P-cadherin Fc (R&D Cat. 861-PC) was reconstituted to a concentration of 1 mg/mL with 2 mM $CaCl_2$ in MilliQ water 24 hours prior to use and stored at 4 °C. A431 cells were cultured and prepased as follows. A431 cells (ECACC No. 85090402) were routinely cultured in Nunc triple flasks (3 x 175cm^2 area) in minimal essential medium (MEM) (Invitrogen Cat 31095), containing 10% fetal bovine serum (Invitrogen Cat. 10100-147) and 1% non-essential amino acids (Invitrogen Cat. 11140-035). The cultured cells were approximately 80% confluent at the time of harvest for use in the assay. To safeguard against possible passage related effects, cells were routinely used between passage 4 and 8 and harvested after either 48 or 72 hours culture. A431 cells were harvested with 0.25% trypsin / 1 mM EDTA (Gibco Cat 25200-056) for just enough time for the cells to dissociate (7-10 min) and then immediately diluted into tissue culture media to a density of approximately 2 x 10^6 cells/mL. The A431 cells were then centrifuged (1200 rpm), re-suspended into assay buffer (Hanks balanced salts solution (without Mg^{2+} and Ca^{2+}, without phenol red) (Invitrogen Cat. 14175-053) supplemented to a final $CaCl_2$ final concentration of 1 mM) re-centrifuged and re-suspended into assay buffer again at a final density of 2 x 10^6 cells/mL.

Preparation of IgG serial dilutions and pre-incubation with A431 cells was performed as follows. 180 µL of each test IgG, or portion thereof, was supplemented with 20 µL of assay buffer containing 10% BSA (Sigma Cat. A-9576) to standardize the BSA concentration to 1%. An anti-murine/human P-cadherin reference polyclonal antibody (R&D Cat. AF-761) was initially resuspended into MilliQ water to give a 1 mg/mL stock. 40 µL of this stock solution was then further diluted into 140 µL of assay buffer. 20 µL of assay buffer containing 10% BSA was then added to

give 200 µL at 0.2 mg/mL antibody and 0.1% BSA. Duplicate serial dilutions were prepared by first adding 2 x 90 µL of AF-761 polyclonal antibody or test IgG to column 1 of a Greiner 96 well polypropylene dilution plate (Greiner Cat. 780271). 60 µL of assay buffer was then added to columns 2-11. A 30 µL into 60 µL (1:3) dilution was then prepared across the plate from left to right from columns 1-11. 60 µL of assay buffer alone was then added to wells 12 A-D to define maximum adhesion. Minimum adhesion was defined by addition of 60 µL of 25 mM EDTA (in assay buffer) to wells 12 E-H. 60 µL of A431 cell suspension (2 x 10^6 cells/mL in assay buffer – prepared as outlined above) was then added to all wells and, after agitation, the plates were pre-incubated for 1 hr at 37 °C.

In parallel with the preparation of test IgG serial dilutions and pre-incubation with A431 cells, P-cadherin coated assay plates were prepared as follows. Recombinant human P-cadherin Fc was diluted in coating buffer (PBS without Mg^{2+} and Ca^{2+} - Invitrogen Cat. 14190-094) to a concentration of 10 µg/mL and dispensed onto Fluoronunc 96 (Nunc Cat. 437958) well assay plates (100 µL/well). Plates were then incubated for 1 hr 30 min. at room temperature. The plates were then washed 3 times with PBS using a Tecan 96 plate washer. 200 µL/well of assay buffer was then added for blocking and the plates were incubated for an additional 1 hr at room temperature. The plates were then washed 3 times with PBS as described previously.

100 µL/well of the preincubated IgG / A431 material was then transferred from the Greiner 96 well dilution plates to the P-cadherin coated assay plates. At the time of transfer the IgG / A431 material was mixed by pipetting to ensure the cells were homogeneous. The adhesion process was then allowed to occur by incubating the assay plates at 37 °C for 30-45 minutes. At the end of the incubation, non-adherent cells were removed by gently aspirating the media from the plates and refilling the wells with cell wash buffer (Hanks balanced salts solution (without Mg^{2+} and Ca^{2+} and without phenol red – Invitrogen Cat. 14175-053) supplemented to 1 mM $CaCl_2$ final concentration). The plates were then inverted on a bath of cell wash buffer for 15 min. to remove residual non-adherent cells. At the end of this incubation, the contents of the wells were gently aspirated.

The quantitation of adherent cells was performed as follows. Adherent cells were detected by addition of 100 µL/well of combined lysis / alkaline phosphatase detection reagent (diethanolamine substrate buffer 5X concentrate (Pierce Cat. 34064) diluted 1:5 with water, after which one 15 mg PNPP tablet (Sigma Cat. N-2640) was dissolved per 25 mL of 1X solution) followed by incubation for 30-60 min. at 37 °C. The reaction was then stopped by addition of 1 M NaOH (50 µL/well), aiming to give a maximum OD value of about 0.8 in the absence of inhibition. Absorbance at 405 nm was then measured using a standard plate reader.

The results were then analyzed as follows. Taking column 12, wells A-D as 100% adhesion and column 12, wells E-H as 0% adhesion, the raw data was first converted to % adhesion values as follows:

% adhesion = {(value-min adhesion) / (max – min adhesion)}*100

The % adhesion versus concentration of IgG Inhibitor was then plotted and IC_{50} values determined using Prism software. Where partial inhibition was observed, the IC_{50} is quoted as the concentration

of IgG giving a true 50% inhibition as opposed to the curve mid-point. IC_{50} values are reported in Table 4.

Table 4

IgG	A431 Adhesion Assay Mean IC_{50} (nM) (n=3)
194-e06	0.162
194-a02	0.217
194-b09	0.229
195-e11	0.114
194-g09	0.158
196-h02	0.148
194-e01	0.147
196-d10	0.080
196-g03	0.149
196-e06	0.117
195-a09	0.114
198-a09	0.097
200-h06	0.168

Two separate A431 adhesion assays were performed to investigate several germlined optimized IgGs from the 129-1c4 lineage in comparison with their non-germlined equivalents. For these experiments, it was necessary to change to a new batch of P-cadherin (CFR-134041), which appears to be associated with slightly increased IC_{50} values compared with other data. Meaned data for several germlined IgGs from the two experiments are shown in Table 5. As noted previously, g-194-b09 refers to the germlined version of 194-b09, and so forth.

Table 5

IgG	A431 Adhesion Assay Mean IC_{50} (nM) (n=2)
g-194-b09	0.77
194-b09	2.10
g-194-g09	1.00
194-g09	0.73
g-196-g03	1.05
196-g03	0.39
g-194-e06	0.77
194-e06	0.46
g-195-e11	0.87
195-e11	0.97
g-200-h06	1.31
200-h06	0.63

Example 4: P-cadherin-dependent cell aggregation assay

The following protocol was used to determine IC_{50} values in a P-cadherin-dependent aggregation assay using several optimized scFvs that were converted to IgG during the optimization phase of antibody discovery as described above in Example 2. Because P-cadherin over-expressing cell lines form tight multicellular aggregates when placed in suspension growth, cell aggregation can be measured in the presence of P-cadherin antibodies that interfere with cellular aggregation. The mean measured IC_{50} values for several antibodies are shown below in Table 6.

Plate preparation was performed as follows. Each 96 well assay plate was coated with 50 µL polyHEMA (12 mg/mL in 90% ethanol, 10% methanol) and then evaporated for 6 hrs to overnight, followed by washing 3 x 100 µL sterile H_2O before use. Cells were then cultured as follows. Cells from the human cell line SW480 (stably expressing P-cadherin G418^r) were passaged in full growth medium (qs DMEM (inVitrogen 11995-065), 10% FBS (Omega Scientific FB-02), 1:100 NEAA (InVitrogen 11140-050), 1:100 sodium pyruvate (InVitrogen 11360-070), 1:100 glutamine (InVitrogen 25030-051), 1:100 penicillin/streptolysin (InVitrogen 15070-063), 1:100, geneticin 50 mg/mL (Invitrogen 10131-035)) + G418 (500 µg/mL), then split 1:3-4 twice per week. Cultures were then frozen in growth medium + 10% DMSO.

On Day 1 the SW480:pCAD cells and control SW480:pCLNX (control vector stable) were seeded at 5 x 10^6 cells/100 mm dish, at no more than 1:3 dilution. Cells were then grown for 48 hours in culture. Each 100 mm dish provided approximately 10 x 10^6 cells, or enough for 2 x 96 well plates.

On Day 3, the medium was removed, followed by washing with dPBS (Dulbecco's PBS (InVitrogen 14040-133)), after which the cells were trypsinized in 3 mL / 100 mm dish. Neutralization was then carried out after release with two volumes (6 mL) of full growth medium. The plate was then washed three times using a pipet with 10 mL to disrupt the clusters. Cells were then counted and a pellet was obtained using a Beckman centrifuge at 1000 rpm for 5 minutes. The media was then aspirated, the pellet resuspended first in < 1 mL full growth medium by finger vortex, then p1000 pipet, then the cell concentration was normalized to 1.3 M/mL. Single cell dispersion was assured by microscopy.

A reagent plate was then prepared by blocking a 96-well plate with dPBS and 5% FBS for 30 min. The plate was then washed using 1 x 100 µL dPBS, followed by aspiration and flick to dry. A dilution series of test IgG was prepared with dPBS, using 4x [IgG] concentration, enough for 3 wells of treatment plate in one well of the 96 well. 40 000 cells in 30 µL / well were aliquotted to a 96 well, washed, poly-HEMA coated Costar3590 non-tissue culture plate (Corning 3590). 10 µL of reagent was then transferred to each well of the 96 well plate. Triplicate samples per treatment were performed using an 8 channel pipet. Incubation then occurred at 250 rpm, shaken, in a humidified 37 °C, 5% CO_2 incubator overnight (16-18 hr).

On day 4, 40 µL of shaken cells were transferred to a poly-lysine-coated 96 well plate (BioCoat poly-lysine-coated 96 well plate: BD 356516). The wells were then rinsed with 60 µL full growth medium, the plate shaken by tapping, and transfered to the poly-lysine-coated plate. If necessary, an additional 50 µL wash was carried out. Incubation followed for 60 minutes in a humidified 37 °C, 5% CO_2 incubator. Care was taken at this step to quantitatively transfer all of the cells, as gently as possible, without excess pipeting. Cells were then fixed by adding 100 µL of fixing solution (7.4% formaldehyde (37% wt/vol. – Sigma F15587)) in a fume hood, followed by incubation for > 30 minutes at room temperature.

To wash the cells, liquid was then decanted into a collection beaker or tray and flicked to remove remaining liquid, and tapped gently on a paper towel. 100 µL per well dPBS was then

applied to wash, followed by incubation for 15 min. The cells were then stained by decanting as above and applying 100 µL Hoescht (1 µg/mL Hoescht in dPBS – Hoescht 10 mg/mL Molecular Probes 33342), followed by incubation for 30 min. The cells were then washed twice, leaving the remaining 100 µL dPBS in the well for microscopy.

The number of aggregated objects per well was then measured (Cellomics) and an average object count (with test IgG) was compared to that of IgG (e.g. Gt-anti-P-cadherin R&D Systems AF761) or media alone control. The object count versus concentration of IgG inhibitor was then plotted and IC_{50} values determined. The IC_{50} values for several antibodies of the present invention are reported in Table 6.

Table 6

IgG	SW480 Aggregation Assay Mean IC_{50} (nM))
194-e06	0.7
194-a02	1.1
g-194-b09	0.9
g-195-e11	2.2
g-194-g09	2.6
196-h02	3.2
194-e01	1.3
196-d10	1.5
g-196-g03	0.9
196-e06	1.9
195-a09	1.7
198-a09	2.9
g-200-h06	4.7
129-1c4	35

Example 5: P-cadherin-dependent spheroid disruption assay

The following spheroid disruption assay is a variation of the aggregation assay (described in Example 4) in which the cell aggregates are formed overnight prior to addition of P-cadherin and control antibodies. Test reagents are then added for an additional 24 hours before analysis.

Plate preparation was performed in the following manner. Each 96 well assay plate was coated with 50 µL polyHEMA (12 mg/mL in 90% ethanol, 10% methanol) and then evaporated for 6 hrs to overnight, followed by washing 3 x 100 µL sterile H_2O before use. Cells were then cultured as follows. Cells from the human cell line SW480 (stably expressing P-cadherin G418^r) were passaged in full growth medium (qs DMEM (InVitrogen 11995-065), 10% FBS (Omega Scientific FB-02), 1:100 NEAA (InVitrogen 11140-050), 1:100 sodium pyruvate (InVitrogen 11360-070), 1:100 glutamine (InVitrogen 25030-051), 1:100 penicillin/streptolysin (InVitrogen 15070-063), 1:100, geneticin 50 mg/mL (Invitrogen 10131-035)) + G418 (500 µg/mL), then split 1:3-4 twice per week. Cultures were then frozen in growth medium + 10% DMSO.

On Day 1 the SW480:pCAD cells and control SW480:pCLNX (control vector stable) were seeded at 5 x 10^6 cells/100 mm dish, at no more than 1:3 dilution. Cells were then grown for 48 hours in culture. Each 100 mm dish provided approximately 10 x 10^6 cells, or enough for 2 x 96 well plates.

On Day 3, the medium was removed, followed by washing with dPBS (Dulbecco's PBS (InVitrogen 14040-133)), after which the cells were trypsinized in 3 mL/100 mm dish. Neutralization was then carried out after release with two volumes (6 mL) of full growth medium. The plate was then washed three times using a pipet with 10 mL to disrupt the clusters. Cells were then counted and obtained a pellet using a Beckman centrifuge at 1000 rpm for 5 minutes. The media was then aspirated, the pellet resuspended first in < 1 mL full growth medium by finger vortex, then p1000 pipet, then the cell concentration was normalized to $1.0x10^6$/mL. Single cell dispersion was assured by microscopy.

40 000 cells in 40 µL / well were aliquotted to a 96 well, washed, poly-HEMA coated Costar3590 non-tissue culture plate (Corning 3590). Incubation then occurred at 250 rpm, shaken, in a humidified 37 °C, 5% CO_2 incubator overnight (16-18 hr).

On day 4, a reagent plate was then prepared by blocking a 96-well plate with dPBS and 5% FBS for 30 min. The plate was then washed using 1 x 100 µL dPBS, followed by aspiration and flick to dry. A dilution series of test IgG was prepared with dPBS, using 5x [IgG] concentration, enough for 3 wells of treatment plate in one well of the 96 well plate. 10 µL of reagent was then transferred to each well of the 96 well plate. Triplicate samples per treatment were performed using an 8 channel pipet. Incubation then occurred at 250 rpm, shaken, in a humidified 37 °C, 5% CO_2 incubator (20-24 h).

On day 5, 50 µL of shaken cells were transferred to a poly-lysine-coated 96 well plate (BioCoat poly-lysine-coated 96 well plate: BD 356516). The wells were then rinsed with 50 µL full growth medium, the plate shaken by tapping, and transfered to the poly-lysine-coated plate. If necessary, an additional 50 µL wash was carried out. Incubation followed for 60 minutes in a humidified 37 °C, 5% CO_2 incubator. Care was taken at this step to quantitatively transfer all of the cells, as gently as possible, without excess pipeting. Cells were then fixed by adding 100 µL of fixing solution (7.4% formaldehyde (37% wt/vol. – Sigma F15587)) in a fume hood, followed by incubation for > 30 minutes at room temperature.

To wash the cells, liquid was then decanted into a collection beaker or tray and flicked to remove remaining liquid, and tapped gently on a paper towel. 100 µL per well dPBS was then applied to wash, followed by incubation for 15 min. The cells were then stained by decanting as above and applying 100 µL Hoescht (1 µg/mL Hoescht in dPBS – Hoescht 10 mg/mL Molecular Probes 33342), followed by incubation for 30 min. The cells were then washed twice, leaving the remaining 100 µL dPBS in the well for microscopy. The number of aggregated objects per well was then measured (Cellomics) and an average object count (with test IgG) was compared to that of IgG (e.g. Gt-anti-P-cadherin R&D Systems AF761) or media alone control. The object count versus concentration of IgG inhibitor can be plotted to determine IC_{50} values. Alternatively, the object count was expressed as fold disruption vs. control at one defined concentration as shown in Table 7.

Table 7

IgG	SW480 Spheroid Disruption Assay Fold Increased Disruption vs Control at 5nM
194-e06	10
194-a02	10
g-194-b09	10
g-195-e11	7
g-194-g09	14
196-h02	10
194-e01	16
196-d10	10
g-196-g03	13
196-e06	10
195-a09	10
198-a09	13
g-200-h06	7
129-1c4	4

Example 6: Measurement of K_D and k_{off} of P-cadherin antibodies

Affinity measures (K_D and k_{off}) of P-cadherin scFv single chain antibodies by surface plasmon resonance using the BIACORE™ 3000 instrument were performed as follows using the manufacturer's protocols.

To perform kinetic analyses, recombinant human P-cadherin/Fc fusion protein (hCad/Fc) and mouse P-cadherin/Fc fusion protein (mCad/Fc) were immobilized on separate flow cells of a CM5 BIACORE™ sensor chip using routine amine coupling. Surfaces were prepared using 10 mM acetate buffer pH 4.5 with 2.0 mM $CaCl_2$ as the immobilization buffer and protein densities of 5800 and 1600 RU were achieved for the hCad/Fc and mCad/Fc fusion proteins, respectively. Deactivation of unreacted N-hydroxysuccinimide esters was performed using 1 M ethanolamine hydrochloride, pH 8.5. An activated/deactivated blank surface was used as a control surface. ScFv antibody samples in running buffer were prepared at concentrations ranging from 200 to 0.78 nM (a 0 nM solution comprising running buffer alone was included as a zero reference). Samples were randomized and injected in triplicate for 1 minute each across the flow cells using HBS-P (10 mM HEPES pH 7.4, 150 mM NaCl, 0.005% Surfactant P20) with 2.0 mM $CaCl_2$ as running buffer. A flow rate of 25 µL/min was used to determine affinity constants. The dissociation of the antibody was monitored for 5 minutes, the surface regenerated by a 12 second injection of 10 mM glycine-HCl pH 1.5 (25 µL/min). The raw data were processed using the Scrubber (©BioLogic Software) software package and analyzed using the CLAMP (©BioLogic Software) software package. The data were fit globally to a simple 1:1 Langmuir binding model.

Table 8 lists affinity constants for the single chain anti-P-cadherin antibodies of the present invention:

Table 8

scFv	hCad/Fc K_D (nM)	hCad/Fc k_{off} (1/s)	mCad/Fc K_D (nM)	mCad/Fc k_{off} (1/s)
194-b09	4.0	1.9×10^{-03}	11	3.6×10^{-3}
194-g09	2.6	1.6×10^{-03}	1.8	8.1×10^{-4}
196-g03	1.1	7.0×10^{-04}	0.76	4.7×10^{-4}

Example 7: Determining selectivity of P-cadherin antibodies

The following protocol was used to determine the selectivity of various antibodies to P-cadherin over E-cadherin.

Recombinant human P-cadherin (R&D Systems 861-PC-100) and recombinant human E-cadherin (R&D Systems 648-EC-100) were coated onto wells of Exiqon protein immobilizer plates (VWR International) at 1 µg/mL in PBS + 0.5 mM $CaCl_2$. Sample IgG's were blocked in 3% Marvel / PBS + 0.5 mM $CaCl_2$ for 1 hour before being titrated from 50 nM (7.5µg/mL) down to 0.64nM (0.096µg/mL) and added in duplicate to wells coated with the two different antigens. Following overnight equilibration at 4°C, the plates were washed three times with 1x PBS / 0.1% Tween + 0.5mM $CaCl_2$ then three times with 1 x PBS + 0.5 mM $CaCl_2$. 50 µL anti-human Fab peroxidase conjugate diluted 1:5000 in 3% Marvel / PBS + 0.5 mM $CaCl_2$ was then added to each well and left to equilibrate at room temperature for 1 hour. Plates were washed three times with 1 x PBS / 0.1% Tween + 0.5 mM $CaCl_2$ and three times with 1 x PBS + 0.5mM $CaCl_2$. 50µl 3,3', 5, 5'-tetramethylbenzidine (TMB; Sigma) was added to each well and reactions allowed to develop for 20 minutes before being stopped by the addition of 25 µL/well 0.5M H_2SO_4. Following reading of absorbance values at 450nm, the data was analyzed using Graphpad Prism software to calculate relative K_D values for each antibody for binding to P-cadherin and E-cadherin. The resulting data is summarized in Table 9.

Table 9

IgG	P-cadherin K_D (pM)	E-cadherin K_D (pM)	$K_D(E)/K_D(P)$
194-a02	116	682	6
g-194-b09	788	2156	3
g-194-g09	2602	No binding	> 100
194-e01	112	20511	183
g-194-e06	191	794	4
195-a09	120	11516	96
194-e06	56	264	5
196-d10	45	63	1.4
196-e06	62	187	3
g-196-g03	449	7513	17
196-h02	102	33212	326
198-a09	78	19396	249

Example 8: Epitope Competition Assay

The following protocol was used to measure IC_{50} values in an epitope competition assay to measure the ability of variant scFvs within a parental lineage to displace the parent IgG from native P-cadherin on the surface of A431 cells. The amount of bound biotinylated parent IgG in the

presence of inhibitory scFv is detected with europium streptavidin and DELFIA quantitation. The measured IC_{50} values for several scFvs are shown in Table 10.

A431 cells (ECACC No. 85090402), cultured according to standard methods, then harvested at approximately 80% confluence, were seeded at $2.5x10^4$ cells per well in 96 well Beckton Dickenson (BD Cat. 6407) collagen coated plates the day prior to use. The plates were then washed 3 times by submersing into a PBS buffer (Gibco Cat. 14190-094, without calcium, magnesium, and sodium bicarbonate) reservoir before addition of 200 µL per well of block buffer (PBS Gibco Cat. 14190-094, plus 3% Marvel (Premier International Foods Ltd.)) and incubated for 2 hours at room temperature. The plates were then washed 3 times with PBS as above.

For both high throughput screening and IC_{50} profiling scFv / IgG materials were first prepared in Greiner dilution plates (Greiner 96 well polypropylene plates (Greiner Cat. 780271)) in a total volume of 60 µL of assay buffer. 50 µL was then transferred from the slave Greiner plate directly onto the assay plate and the binding reaction was allowed to proceed for 2 hr 30 min. at room temperature. At the end of the binding reaction the plates were washed 3 times in PBS before the addition of europium labelled streptavidin (Perkin Elmer Cat. 1244-360 1:1000 dilution in DELFIA assay buffer (Perkin Elmer Cat. 4002-0010)) and incubated for a further 1hour at room temperature.

The plates were then washed 7 times with DELFIA wash buffer (Perkin Elmer Cat. 4010-0010) by repeatedly submerging plates into a buffer reservoir. Finally, after addition of DELFIA enhancer (Perkin Elmer Cat. 4001-0010 – 100 µL/well) plates were read using a standard DELFIA protocol on a compatible reader (e.g. Wallac or Envision).

Greiner dilution plate set up – HTS

High throughput screening (HTS) conditions were configured differently depending on the stage of the optimization. For HTS of outputs from the individual V_H and V_L chain optimization final [peri-prep] was fixed at 12.5% such that the parent scFv gave partial inhibition. For the later HTS of outputs from the V_H:V_L recombination libraries final peri-prep concentration was reduced to 1.7% and under these conditions the V_H optimized benchmark scFv (TOP-108-C01) gave partial inhibition. The optimization of peri-prep concentration such that the relevant benchmark scFv gave partial inhibition left a window in which to identify improved clones.

To achieve these final scFv peri-prep concentrations the following procedure was adopted:

i) Using a Cybiwell instrument, the required volume (see below) of peri-prep sample material was transferred from a deep well sample plate to a Greiner dilution plate in columns 1-11.

[Final peri-prep]		Transfer volume
[12.5%]	=	7.5 uL
[1.7%]	=	10.0 uL (of 1:10 diluted peri-prep)

(1:10 pre-diluted peri-prep was prepared by transferring 10 µL of neat peri-prep from the sample plate into a Greiner dilution plate containing 90 µL of assay buffer (using Cybiwell).

ii) The volume in columns 1-11 was then made up to 30 µL by addition of an appropriate volume of assay buffer.

iii) 30 µL of assay buffer was then added to column 12 (A-D) (total binding wells).

iv) 30 µL of excess unlabelled parent IgG (129-1c4) in assay buffer was added to column 12 (E-H) to define non-specific binding. This was added at 1000 nM concentration to give 500 nM final.

v) The 129-1c4 IgG was biotinylated using the water insoluble reagent EZ-link-NHS-LC-Biotin (Perbio/Pierce product no. 21336). The IgG solution was supplemented by adding a $1/10^{th}$ volume of 1 M $NaHCO_3$ and a $1/10^{th}$ volume of dimethylformamide. The EZ-link-NHS-LC-Biotin reagent was dissolved in DMF and then added at a five-fold molar excess over IgG and the reaction allowed to proceed at room temperature for 20 minutes. The biotinylated IgG was then stabilized by addition of BSA (0.1%). 30 µL of biotinylated parent 129-1c4 IgG in assay buffer was then added to all wells to give a final concentration of 1.5 nM in the final 60 µL volume (i.e. europium labelled IgG added at 2x final concentration).

vi) After mixing the Greiner plate contents by agitation, 50 µL was transferred directly to the assay plate and the binding reaction allowed to proceed for 2 hrs 30 min at room temperature (as described previously).

Greiner dilution plate set up – IC_{50} profiling

i) 30 µL/well of assay buffer was added to columns 2-11 and wells 12A to 12D in standard Greiner dilution plates.

ii) 30 µL of excess unlabelled 129-1c4 parent IgG in assay buffer was added to column 12 (E-H) to define non-specific binding. This should be added at 1000 nM concentration to give 500 nM final concentration

lii) To column 1 2x45µL of each undiluted scFv His-prep was addied such that 4 duplicate 11 point IC_{50} titrations could be set up per 96 well assay plate. 1:3 duplicate serial dilutions were then performed by taking 15 µL from column 1 and mixing into column 2 followed by taking 15 µL from column 2 and mixing into column 3, etc. After mixing into column 11, 15 µL was removed (i.e. to leave 30 µL final volume).

iv) 30 µL of biotinylated 129-1c4 parent IgG in assay buffer was then added to all wells to give a final concentration of 1.5nM in the final 60 µL volume (i.e. europium labelled IgG added at 2x final concentration).

v) After mixing the Greiner plate contents by agitation, 50 µL was then transferred directly to the assay plate and the binding reaction allowed to proceed for 2 hr 30 min at room temperature as discussed previously.

For HTS, scFv were expressed in 96 well format as crude extract from the bacterial periplasm (scFv peri-prep) in a base buffer containing MOPS / EDTA / Sorbitol pH 7.4 (MES pH 7.4). For IC_{50} profiling, scFv were expressed in a lower throughput purified form using the scFv His-tag for affinity purification (scFv His-prep) in a base buffer of PBS.

For both HTS and IC_{50} analysis, raw data was first converted to % binding data according to the following equation:

% binding ={(value-non-specific binding) / (total binding-non specific binding)}*100

HTS data was then further analyzed using standard Excel templates to identify hits giving greater inhibition (lower % binding) than the relevant benchmark scFv. For IC_{50} analysis, % binding data was analyzed using Prism version 4.0 curve fitting software. The scFvs are variants of the 129-1c4 parent IgG, wherein the V_H and V_L CDR3 regions were randomly mutated, as described in Example 2. V_H CDR3 variant sequences are shown in Figure 1 as SEQ ID NOs: 91 to 256. V_L CDR3 variant sequences are shown in Figure 1 as SEQ ID NOs: 257 to 319. The IC_{50} values for several scFvs that showed improved binding over the parent 129-1c4 IgG are shown in Table 10. Each of the scFv variants of the 129-1c4 parent are identified by two SEQ ID NOs: a V_H CDR3 and a V_L CDR3. For reference, the IC_{50} of the parent 129-1c4 IgG was 108.3 nM.

Table 10

SEQ ID NO: V_H CDR3	SEQ ID NO: V_L CDR3	IC_{50} (nM)
37	40	2.49
37	261	1.78
37	43	5.67
37	274	8.73
37	277	2.20
37	287	10.50
37	288	6.67
37	295	0.80
37	297	2.49
37	299	1.97
37	303	1.47
37	304	1.10
37	305	2.19
95	47	2.36
97	47	1.33
113	47	2.06
125	47	0.26
131	47	0.35
31	47	2.05
147	47	1.16
165	47	5.11
167	47	0.60
180	47	0.43
181	47	2.46
27	47	0.54
195	47	1.29
187	43	1.58
27	296	1.16
163	43	2.45
26	40	0.76
26	43	0.93
27	309	1.36
199	310	2.54
201	310	2.26
207	291	2.73
240	313	2.98

All publications, patents, and patent applications cited in this specification are incorporated herein by reference as if each individual publication or patent application were specifically and individually indicated to be incorporated by reference. Although the foregoing invention has been described in some detail by way of illustration and example for purposes of clarity of understanding, it will be readily apparent to those of ordinary skill in the art in light of the teachings of this invention that certain changes and modifications may be made thereto without departing from the spirit or scope of the appended claims.

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Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Gly Ser Arg Thr Val Gln Phe Asp Pro Trp Gly Gln Gly Thr Met
100 105 110

Val Thr Val Ser Ser
115

<210> 9
<211> 117
<212> PRT
<213> human

<400> 9

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

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Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Thr Asn Ser Pro Gly Thr Phe Asp Pro Trp Gly Gln Gly Thr Met
            100                 105                 110

Val Thr Val Ser Ser
        115
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<211> 117
<212> PRT
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<400> 10

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
            20                  25                  30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Thr Ile Ala Pro Gly Arg Phe Asp Pro Trp Gly Gln Gly Thr Met
            100                 105                 110

Val Thr Val Ser Ser
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115

<210> 11
<211> 117
<212> PRT
<213> human

<400> 11

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Gly Leu Asp Arg Val Trp Phe Asp Pro Trp Gly Gln Gly Thr Met
100 105 110

Val Thr Val Ser Ser
115

<210> 12
<211> 117
<212> PRT
<213> human

<400> 12

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

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Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Ser Trp Gly Gly Gly Trp Phe Asp Pro Trp Gly Gln Gly Thr Met
            100                 105                 110

Val Thr Val Ser Ser
        115
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<210> 13
<211> 117
<212> PRT
<213> human

<400> 13

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                   10                  15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
            20                  25                  30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Lys Trp Gly Gly Gly Trp Phe Asp Pro Trp Gly Gln Gly Thr Met
            100                 105                 110
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Val Thr Val Ser Ser
115

<210> 14
<211> 111
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<400> 14

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Phe Thr Ser Gly
85 90 95

Leu Pro Trp Val Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 15
<211> 111
<212> PRT
<213> human

<400> 15

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu

35 40 45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Phe Thr Ser Gly
85 90 95

Leu Pro Trp Val Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 16
<211> 111
<212> PRT
<213> human

<400> 16

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Phe Thr Ser Gly
85 90 95

Ile Val Phe Asn Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 17
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<213> human

<400> 17

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Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1               5                  10                  15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
            20                  25                  30

Asn Tyr Val Ser Trp Tyr Gln Arg His Pro Gly Lys Ala Pro Lys Leu
        35                  40                  45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
    50                  55                  60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65                  70                  75                  80

Gln Ala Glu Asp Glu Ala Glu Tyr Tyr Cys Ser Ser Tyr Thr Met Gly
                85                  90                  95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
            100                 105                 110
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<210> 18
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<400> 18

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1               5                  10                  15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
            20                  25                  30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
        35                  40                  45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
    50                  55                  60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65                  70                  75                  80
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Gln Ala Glu Asp Glu Ala Gly Tyr Tyr Cys Thr Ser Tyr Arg Ala Gly
                85                  90                  95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
            100                 105                 110


<210>  19
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<212>  PRT
<213>  human

<400>  19

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Pro Pro Gly Gln
1               5                  10                  15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
            20                  25                  30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
        35                  40                  45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
    50                  55                  60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65                  70                  75                  80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Thr Ser Tyr Thr Met Gly
                85                  90                  95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
            100                 105                 110


<210>  20
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<400>  20

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1               5                  10                  15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
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20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Thr Ser Tyr Arg Met Asp
85 90 95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 21
<211> 111
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<213> human

<400> 21

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Phe Thr Ser Gly
85 90 95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 22
<211> 111
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<400> 22

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Met Ile Tyr Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Phe Thr Ser Gly
85 90 95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 23
<211> 111
<212> PRT
<213> human

<400> 23

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Ile Leu Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Ser Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Thr Ser Tyr Thr Met Gly
85 90 95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

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<400> 24

Ser Tyr Ala Met Ser
1 5

<210> 25
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<400> 25

Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys
1 5 10 15

Gly

<210> 26
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<400> 26

Trp Gly Thr Gly Thr Leu Trp Pro
1 5

<210> 27
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<400> 27

Trp Gly Asp Gly Thr Leu Asn Pro
1 5

<210> 28
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<400> 28

Trp Gly Leu Gly Ser Asn Glu Asn
1 5

<210> 29
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<400> 29

Thr Asn Ser Ala Lys Phe Asp Pro
1 5

<210> 30
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<400> 30

Thr Gly Tyr Pro Ser Phe Asp Pro
1 5

<210> 31
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<400> 31

Thr Ala Lys Pro Ser Phe Asp Pro
1 5

<210> 32
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Asn Glu Arg Pro Ser Phe Asp Pro

1 5

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Ser Arg Thr Val Gln Phe Asp Pro
1 5

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Asn Ser Pro Gly Thr Phe Asp Pro
1 5

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<400> 35

Ile Ala Pro Gly Arg Phe Asp Pro
1 5

<210> 36
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Leu Asp Arg Val Trp Phe Asp Pro
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Trp Gly Gly Gly Trp Phe Asp Pro
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<400> 38

Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr Asn Tyr Val Ser
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Glu Val Asn Lys Arg Pro Ser
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Ser Ser Phe Thr Ser Gly Leu Pro Trp Val Leu
1 5 10

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Ser Ser Phe Thr Ser Gly Leu Pro Trp Val Val
1 5 10

<210> 42
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Ser Ser Phe Thr Ser Gly Ile Val Phe Asn Leu
1 5 10

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Thr Ser Tyr Thr Met Gly Ser Thr Phe Met Leu
1 5 10

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Ser Ser Tyr Thr Met Gly Ser Thr Phe Met Leu
1 5 10

<210> 45
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<400> 45

Thr Ser Tyr Arg Met Asp Ser Thr Phe Met Leu
1 5 10

<210> 46
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<400> 46

Thr Ser Tyr Arg Ala Gly Ser Thr Phe Met Leu
1 5 10

<210> 47
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<400> 47

Ser Ser Phe Thr Ser Gly Ser Thr Phe Met Leu
1 5 10

<210> 48
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<400> 48

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser
20 25 30

<210> 49
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<400> 49

Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ser
1 5 10

<210> 50
<211> 32
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<400> 50

Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
1 5 10 15

Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Ser
20 25 30

<210> 51
<211> 32
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<400> 51

Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
1 5 10 15

Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Asp
20 25 30

<210> 52
<211> 32
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<400> 52

Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
1 5 10 15

Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Gly
20 25 30

<210> 53
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<400> 53

Arg Phe Ala Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
1 5 10 15

Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Gly
20 25 30

<210> 54
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Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
1 5 10 15

Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Thr
20 25 30

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Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
1 5 10 15

Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Lys
20 25 30

<210> 56

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<400> 56

Trp Gly Gln Gly Lys Met Val Thr Val Ser Ser
1 5 10

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Trp Gly Gln Gly Thr Met Val Thr Val Ser Ser
1 5 10

<210> 58
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<400> 58

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Pro Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys
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<210> 59
<211> 22
<212> PRT
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<400> 59

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys
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<210> 60
<211> 15
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<400> 60

Trp Tyr Gln Arg His Pro Gly Lys Ala Pro Lys Leu Ile Leu Ser
1 5 10 15

<210> 61
<211> 15
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<400> 61

Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu Met Ile Tyr
1 5 10 15

<210> 62
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Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu Ile Leu Ser
1 5 10 15

<210> 63
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<400> 63

Gly Val Ser Asn Arg Phe Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser
1 5 10 15

Leu Ser Ile Ser Gly Leu Gln Ala Glu Asp Glu Ala Glu Tyr Tyr Cys
20 25 30

<210> 64
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<400> 64

Gly Val Ser Asn Arg Phe Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser
1 5 10 15

Leu Ser Ile Ser Gly Leu Gln Ala Glu Asp Glu Ala Gly Tyr Tyr Cys
20 25 30

<210> 65
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<400> 65

Gly Val Ser Asn Arg Phe Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser
1 5 10 15

Leu Thr Ile Ser Gly Leu Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys
20 25 30

<210> 66
<211> 32
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<400> 66

Gly Val Ser Asn Arg Phe Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser
1 5 10 15

Leu Ser Ile Ser Gly Leu Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys
20 25 30

<210> 67
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<213> human

<400> 67

Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
1 5 10

<210> 68
<211> 351
<212> DNA
<213> human

<400> 68
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gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gagttgggga 300

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<210> 69
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<212> DNA
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<222> (330)..(330)
<223> n is a, c, g, or t

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<222> (348)..(348)
<223> n is a, c, g, or t

<400> 69
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ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gagttgggga 300

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gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

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<400> 71
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gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc ggacacgaac 300

tccgccaagt tcgacccctg gggccaaggg acaatggtca ccgtctcgag t 351

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<400> 72
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ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

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<210> 73
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<400> 73
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gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gggcacggcc 300

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<210> 74
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<400> 74
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ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc ggggaacgag 300

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<213> human

<400> 75
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Ala Ser Tyr Gln Ala Ser Ser Thr Phe Met Leu
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        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Lys Trp Gly Asp Gly Thr Leu Asn Pro Trp Gly Gln Gly Thr Met
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Val Thr Val Ser Ser
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Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95
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Ala Lys Thr Asn Ser Ala Lys Phe Asp Pro Trp Gly Gln Gly Thr Met
            100                 105                 110

Val Thr Val Ser Ser
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Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Lys Asn Glu Arg Pro Ser Phe Asp Pro Trp Gly Gln Gly Thr Met
            100                 105                 110

Val Thr Val Ser Ser
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Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Lys Thr Asn Ser Ala Lys Phe Asp Pro Trp Gly Gln Gly Thr Met
            100                 105                 110

Val Thr Val Ser Ser
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            20                  25                  30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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Ala Lys Thr Gly Tyr Pro Ser Phe Asp Pro Trp Gly Gln Gly Thr Met
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Val Thr Val Ser Ser
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Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Lys Trp Gly Thr Gly Thr Leu Trp Pro Trp Gly Gln Gly Thr Met
100 105 110

Val Thr Val Ser Ser
115

<210> 326
<211> 111
<212> PRT
<213> human

<400> 326

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Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
            20                  25                  30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
        35                  40                  45

Met Ile Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
    50                  55                  60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65                  70                  75                  80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Phe Thr Ser Gly
                85                  90                  95

Leu Pro Trp Val Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
            100                 105                 110
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<210> 327
<211> 111
<212> PRT
<213> human

<400> 327

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Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1               5                  10                  15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
            20                  25                  30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
        35                  40                  45

Met Ile Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
    50                  55                  60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65                  70                  75                  80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Tyr Thr Met Gly
                85                  90                  95
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Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 328
<211> 111
<212> PRT
<213> human

<400> 328

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Met Ile Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Thr Ser Tyr Thr Met Gly
85 90 95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 329
<211> 111
<212> PRT
<213> human

<400> 329

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu

```
        35                  40                  45

Met Ile Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
    50                  55                  60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65                  70                  75                  80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Thr Ser Tyr Thr Met Gly
                85                  90                  95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
            100                 105                 110
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<210> 330
<211> 111
<212> PRT
<213> human

<400> 330

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Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1               5                   10                  15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
            20                  25                  30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
        35                  40                  45

Met Ile Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
    50                  55                  60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65                  70                  75                  80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Thr Ser Tyr Arg Ala Gly
                85                  90                  95

Ser Thr Phe Met Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
            100                 105                 110
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<210> 331
<211> 111
<212> PRT

<213> human

<400> 331

Gln Ser Ala Leu Thr Gln Pro Ala Ser Val Ser Gly Ser Pro Gly Gln
1 5 10 15

Ser Ile Thr Ile Ser Cys Thr Gly Thr Ser Asn Asp Val Gly Ala Tyr
20 25 30

Asn Tyr Val Ser Trp Tyr Gln Gln His Pro Gly Lys Ala Pro Lys Leu
35 40 45

Met Ile Ser Glu Val Asn Lys Arg Pro Ser Gly Val Ser Asn Arg Phe
50 55 60

Ser Gly Ser Lys Ser Gly Asn Thr Ala Ser Leu Thr Ile Ser Gly Leu
65 70 75 80

Gln Ala Glu Asp Glu Ala Asp Tyr Tyr Cys Ser Ser Phe Thr Ser Gly
85 90 95

Leu Pro Trp Val Leu Phe Gly Gly Gly Thr Lys Leu Thr Val Leu
100 105 110

<210> 332
<211> 351
<212> DNA
<213> human

<400> 332
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tcctgtgcag cctctggatt cacctttagc agctatgcca tgagctgggt ccgccaggct 120

ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gaaatgggga 300

gacgggacct tgaacccgtg gggccaaggg acaatggtca ccgtctcctc a 351

<210> 333
<211> 351
<212> DNA
<213> human

<400> 333

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tcctgtgcag cctctggatt cacctttagc agctatgcca tgagctgggt ccgccaggct 120

ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gaaaacgaac 300

tccgccaagt tcgacccctg gggccaaggg acaatggtca ccgtctcctc a 351

<210> 334
<211> 351
<212> DNA
<213> human

<400> 334
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ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gaaaaacgag 300

aggccgtcgt tcgacccctg gggccaaggg acaatggtca ccgtctcctc a 351

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<212> DNA
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<400> 335
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tcctgtgcag cctctggatt cacctttagc agctatgcca tgagctgggt ccgccaggct 120

ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gaaatgggga 300

gacgggacct tgaacccgtg gggccaaggg acaatggtca ccgtctcctc a 351

<210> 336
<211> 351
<212> DNA
<213> human

<400> 336

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tcctgtgcag cctctggatt cacctttagc agctatgcca tgagctgggt ccgccaggct 120

ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gaaaacgggg 300

taccccctcct tcgaccccctg gggccaaggg acaatggtca ccgtctcctc a 351

<210> 337
<211> 351
<212> DNA
<213> human

<400> 337
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tcctgtgcag cctctggatt cacctttagc agctatgcca tgagctgggt ccgccaggct 120

ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gaaatgggga 300

acggggacct tgtggccctg gggccaaggg acaatggtca ccgtctcctc a 351

<210> 338
<211> 333
<212> DNA
<213> human

<400> 338
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cacccaggca aagcccccaa actcatgatt tctgaggtca ataaacggcc ctcaggggtt 180

tctaatcgct tctctggctc caagtctggc aacacggcct ctctgaccat ctctgggctc 240

caggctgagg acgaggctga ttattactgc agctcattta caagcgggct cccctgggtc 300

gtcttcggcg gagggaccaa gctgaccgtc cta 333

<210> 339
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<212> DNA
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<400> 339

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tcctgcactg gaaccagtaa tgacgttggt gcttataatt atgtctcctg gtaccaacaa 120

cacccaggca aagcccccaa actcatgatt tctgaggtca ataaacggcc ctcaggggtt 180

tctaatcgct tctctggctc caagtctggc aacacggcct ctctgaccat ctctgggctc 240

caggctgagg acgaggctga ttattactgc agcagctaca cgatggggag cacttttatg 300

ctattcggcg gagggaccaa gctgaccgtc cta 333

<210> 340
<211> 333
<212> DNA
<213> human

<400> 340
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tcctgcactg gaaccagtaa tgacgttggt gcttataatt atgtctcctg gtaccaacaa 120

cacccaggca aagcccccaa actcatgatt tctgaggtca ataaacggcc ctcaggggtt 180

tctaatcgct tctctggctc caagtctggc aacacggcct ctctgaccat ctctgggctc 240

caggctgagg acgaggctga ttattactgc acctcgtaca ccatgggcag cacttttatg 300

ctattcggcg gagggaccaa gctgaccgtc cta 333

<210> 341
<211> 333
<212> DNA
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<400> 341
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tcctgcactg gaaccagtaa tgacgttggt gcttataatt atgtctcctg gtaccaacaa 120

cacccaggca aagcccccaa actcatgatt tctgaggtca ataaacggcc ctcaggggtt 180

tctaatcgct tctctggctc caagtctggc aacacggcct ctctgaccat ctctgggctc 240

caggctgagg acgaggctga ttattactgc acctcgtaca ccatgggcag cacttttatg 300

ctattcggcg gagggaccaa gctgaccgtc cta 333

<210> 342
<211> 333
<212> DNA
<213> human

<400> 342

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tcctgcactg gaaccagtaa tgacgttggt gcttataatt atgtctcctg gtaccaacaa 120

cacccaggca aagccccdaa actcatgatt tctgaggtca ataaacggcc ctcaggggtt 180

tctaatcgct tctctggctc caagtctggc aacacggcct ctctgaccat ctctgggctc 240

caggctgagg acgaggctga ttattactgc acctcgtaca ccatgggcag cacttttatg 300

ctattcggcg gagggaccaa gctgaccgtc cta 333

<210> 343
<211> 333
<212> DNA
<213> human

<400> 343
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tcctgcactg gaaccagtaa tgacgttggt gcttataatt atgtctcctg gtaccaacaa 120

cacccaggca aagccccaa actcatgatt tctgaggtca ataaacggcc ctcaggggtt 180

tctaatcgct tctctggctc caagtctggc aacacggcct ctctgaccat ctctgggctc 240

caggctgagg acgaggctga ttattactgc agctcattta caagcgggtt gccgtgggtg 300

ctcttcggcg gagggaccaa gctgaccgtc cta 333

<210> 344
<211> 330
<212> PRT
<213> human

<400> 344

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys
1 5 10 15

Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
20 25 30

Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
35 40 45

Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser
50 55 60

Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser Leu Gly Thr Gln Thr
65 70 75 80

```
Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
                85                  90                  95

Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys
            100                 105                 110

Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro
        115                 120                 125

Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys
    130                 135                 140

Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp
145                 150                 155                 160

Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu
                165                 170                 175

Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu
            180                 185                 190

His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn
        195                 200                 205

Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly
    210                 215                 220

Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu Glu
225                 230                 235                 240

Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr
                245                 250                 255

Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn
            260                 265                 270

Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe
        275                 280                 285

Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn
    290                 295                 300
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Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn His Tyr Thr
305                 310                 315                 320

Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
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<210> 345
<211> 106
<212> PRT
<213> human

<400> 345

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Gly Gln Pro Lys Ala Ala Pro Ser Val Thr Leu Phe Pro Pro Ser Ser
1               5                   10                  15

Glu Glu Leu Gln Ala Asn Lys Ala Thr Leu Val Cys Leu Ile Ser Asp
            20                  25                  30

Phe Tyr Pro Gly Ala Val Thr Val Ala Trp Lys Ala Asp Ser Ser Pro
        35                  40                  45

Val Lys Ala Gly Val Glu Thr Thr Thr Pro Ser Lys Gln Ser Asn Asn
    50                  55                  60

Lys Tyr Ala Ala Ser Ser Tyr Leu Ser Leu Thr Pro Glu Gln Trp Lys
65                  70                  75                  80

Ser His Arg Ser Tyr Ser Cys Gln Val Thr His Glu Gly Ser Thr Val
                85                  90                  95

Glu Lys Thr Val Ala Pro Thr Glu Cys Ser
            100                 105
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<210> 346
<211> 19
<212> PRT
<213> human

<400> 346

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Met Glu Trp Ser Trp Val Phe Leu Phe Phe Leu Ser Val Thr Thr Gly
1               5                   10                  15

Val His Ser
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<210> 347
<211> 20
<212> PRT
<213> human

<400> 347

Met Ser Val Pro Thr Gln Val Leu Gly Leu Leu Leu Leu Trp Leu Thr
1 5 10 15

Asp Ala Arg Cys
20

We Claim:

1. An antibody or antigen-binding portion thereof that has at least one property selected from the group consisting of:

a) has a $K_D(E)/K_D(P)$ that is greater than or equal to 1.5;

b) binds to P-cadherin with a K_D of 50 nM or less as measured by surface plasmon resonance;

c) a k_{off} for P-cadherin of less than or equal to 0.01 $^{s-1}$ as measured by surface plasmon resonance;

d) an IC_{50} of 50 nM or less as measured by a P-cadherin dependent cell adhesion assay;

e) an IC_{50} of 50 nM or less as measured by a P-cadherin dependent cell aggregation assay ;

f) increases spheroid disruption in a P-cadherin-dependent spheroid disruption assay by a factor of 2 or more as compared to a control sample with no IgG;

g) competes for binding to P-cadherin with an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4;

h) cross-competes for binding to P-cadherin with an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06129-1c4; and g-129-1c4;

i) binds to the same epitope of P-cadherin as an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4;

j) binds to P-cadherin with substantially the same K_D as an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4; and

k) binds to P-cadherin with substantially the same k_{off} as an antibody selected from the group consisting of 194-e06; 194-a02; 194-b09; 195-e11; 194-g09; 196-h02; 194-e01; 196-d10; 196-g03; 196-e06; 195-a09; 198-a09; 200-h06; g-194-b09; g-194-g09; g-196-g03; g-196-h02; g-194-e01; g-194-e06; 129-1c4; and g-129-1c4.

2. The antibody or antigen-binding portion thereof according to claim 1, comprising a V_H domain that is at least 90% identical in amino acid sequence to any one of SEQ ID NOs: 13, 320, 321, and 322.

3. The antibody or antigen-binding portion thereof according to claim 1, comprising a V_L domain that is at least 90% identical in amino acid sequence to any one of SEQ ID NOs: 22, 326, 327, and 328.

4. The antibody or antigen-binding portion thereof according to claim 1 selected from the group consisting of:

a) an antibody or antigen-binding portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 13, or differs from SEQ ID NO: 13 by 1 to 10 conservative amino acid substitutions, and a V_L domain as set forth in SEQ ID NO: 22, or differs from SEQ ID NO: 22 by 1 to 10 conservative amino acid substitutions;

b) an antibody or antigen-binding portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 320, or differs from SEQ ID NO: 320 by 1 to 10 conservative amino acid substitutions, and a V_L domain as set forth in SEQ ID NO: 326, or differs from SEQ ID NO: 326 by 1 to 10 conservative amino acid substitutions;

c) an antibody or antigen-binding portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 321, or differs from SEQ ID NO: 321 by 1 to 10 conservative amino acid substitutions, and a V_L domain as set forth in SEQ ID NO: 327, or differs from SEQ ID NO: 327 by 1 to 10 conservative amino acid substitutions; and

d) an antibody or antigen-binding portion thereof that comprises a V_H domain as set forth in SEQ ID NO: 322, or differs from SEQ ID NO: 322 by 1 to 10 conservative amino acid substitutions, and a V_L domain as set forth in SEQ ID NO: 328, or differs from SEQ ID NO: 328 by 1 to 10 conservative amino acid substitutions.

5. The antibody or antigen-binding portion thereof according to claim 1, comprising a V_H domain that is independently selected from any one of SEQ ID NOs: 1 to 13 and 320 to 325, or a sequence that differs from any one of SEQ ID NOs: 1 to 13 and 320 to 325 by 1 to 10 conservative amino acid substitutions; and further comprising a V_L domain that is independently selected from any one of SEQ ID NOs: 14 to 23 and 326 to 331, or a sequence that differs from any one of SEQ ID NOs: 14 to 23 and 326 to 331 by 1 to 10 conservative amino acid substitutions.

6. The antibody or antigen-binding portion thereof according to claim 1, wherein said antibody or antigen-binding portion comprises a V_H CDR3 selected from any one of SEQ ID

NOs: 26 to 37 and 91 to 256, or a sequence that differs from any one of SEQ ID NOs: 26 to 37 and 91 to 256 by 1 or 2 conservative amino acid substitutions.

7. The antibody or antigen-binding portion thereof according to claim 1, wherein said antibody or antigen-binding portion comprises a V_L CDR3 selected from any one of SEQ ID NOs: 40 to 47 and 257 to 319, or a sequence that differs from any one of SEQ ID NOs: 40 to 47 and 257 to 319 by 1 or 2 conservative amino acid substitutions.

8. The antibody or antigen-binding portion thereof according to claim 1, wherein said antibody or antigen-binding portion comprises:

a) a V_H CDR1 as set forth in SEQ ID NO: 24, or a sequence that differs from SEQ ID NO: 24 by 1 or 2 conservative amino acid substitutions;

b) a V_H CDR2 as set forth in SEQ ID NO: 25, or a sequence that differs from SEQ ID NO: 25 by 1 or 2 conservative amino acid substitutions; and

c) a V_H CDR3 that is selected from any one of SEQ ID NOs: 26 to 37 and 91 to 256, or a sequence that differs from any one of SEQ ID NOs: 26 to 37 and 91 to 256 by 1 or 2 conservative amino acid substitutions.

9. The antibody or antigen-binding portion thereof according to claim 1, wherein said antibody or antigen-binding portion comprises:

a) a V_L CDR1 as set forth in SEQ ID NO: 38, or a sequence that differs from SEQ ID NO: 38 by 1 or 2 conservative amino acid substitutions;

b) a V_L CDR2 as set forth in SEQ ID NO: 39, or a sequence that differs from SEQ ID NO: 39 by 1 or 2 conservative amino acid substitutions; and

c) a V_L CDR3 that is selected from any one of SEQ ID NOs: 40 to 47 and 257 to 319, or a sequence that differs from any one of SEQ ID NOs: 40 to 47 and 257 to 319 by 1 or 2 conservative amino acid substitutions.

10. An antibody or antigen-binding portion thereof, wherein the V_H domain comprises any one of SEQ ID NOs: 13, 320, 321, and 322, or a sequence that differs from any one of SEQ ID NOs: 13, 320, 321, and 322 by 1 to 10 conservative amino acid substitutions; and the V_L domain comprises any one of SEQ ID NOs: 22, 326, 327, and 328, or differs from any one of SEQ ID NOs: 22, 326, 327, and 328 by 1 to 10 conservative amino acid substitutions.

11. The antibody according to any of claims 1 to 10 that is an IgG, an IgM, an IgE, an IgA, or an IgD molecule, or is derived therefrom.

12. The antibody according to claim 11 wherein the IgG is an IgG_1 wherein the heavy chain constant region comprises SEQ ID NO: 344 and wherein the light chain constant region comprises SEQ ID NO: 345, with the proviso that the C-terminal lysine residue of SEQ ID NO: 344 is optionally cleaved.

13. A pharmaceutical composition comprising the antibody or antigen-binding portion according to any of claims 1 to 12 and a pharmaceutically acceptable carrier.

14. An isolated nucleic acid molecule comprising a nucleotide sequence as set forth in any one of SEQ ID NOs: 80, 89, 332, 333, 334, 338, 339, and 340.

15. An antibody or antigen-binding portion thereof according to claim 1, comprising a heavy chain variable region amino acid sequence that utilizes a human V_H-3 family gene.