Sequence Listing Information:
DTD Version: V1_3
File Name: 732134 SA9-314PC.xml
Software Name: WIPO Sequence
Software Version: 2.2.0
Production Date: 2022-11-09
General Information:
Current application / IP Office: WO
Current application / Application number: PCT/US2022/078341
Current application / Filing date: 2022-10-19
Current application / Applicant file reference: 732134: SA9-314PC
Earliest priority application / IP Office: US
Earliest priority application / Application number: 63/257,876
Earliest priority application / Filing date: 2021-10-20
Applicant name: SANOFI BIOTECHNOLOGY
Applicant name / Language: en
Inventor name: LAWS, Elizabeth
Inventor name / Language: en
Invention title: METHODS FOR TREATING PRURIGO NODULARIS BY ADMINISTERING AN IL-4R ANTAGONIST (en)
Sequence Total Quantity: 295
Sequences:
Sequence Number (ID): 1
Length: 124
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..124
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LEQPGGSLRL SCAGSGFTFR DYAMTWVRQA PGKGLEWVSS ISGSGGNTYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCAKDR LSITIRPRYY GLDVWGQGTT
120
VTVS
124

Sequence Number (ID): 2
Length: 112
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..112
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSPLS LPVTPGEPAS ISCRSSQSLL YSIGYNYLDW YLQKSGQSPQ LLIYLGSNRA
60
SGVPDRFSGS GSGTDFTLKI SRVEAEDVGF YYCMQALQTP YTFGQGTKLE IK
112

Sequence Number (ID): 3
Length: 8
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..8

```
                    > mol_type, protein
                    > organism, synthetic construct
        Residues:
        GFTFRDYA
8

        Sequence Number (ID): 4
        Length: 8
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..8
                    > mol_type, protein
                    > organism, synthetic construct
        Residues:
        ISGSGGNT
8

        Sequence Number (ID): 5
        Length: 16
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..16
                    > mol_type, protein
                    > organism, synthetic construct
        Residues:
        AKDRLSITIR  PRYYGL
16

        Sequence Number (ID): 6
        Length: 11
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..11
                    > mol_type, protein
                    > organism, synthetic construct
        Residues:
        QSLLYSIGYN  Y
11

        Sequence Number (ID): 7
        Residues:
        000


        Sequence Number (ID): 8
        Length: 9
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..9
                    > mol_type, protein
                    > organism, synthetic construct
        Residues:
        MQALQTPYT
9
```

Sequence Number (ID): 9
Length: 451
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..451
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LEQPGGSLRL SCAGSGFTFR DYAMTWVRQA PGKGLEWVSS ISGSGGNTYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCAKDR LSITIRPRYY GLDVWGQGTT
120
VTVSSASTKG PSVFPLAPCS RSTSESTAAL GCLVKDYFPE PVTVSWNSGA LTSGVHTFPA
180
VLQSSGLYSL SSVVTVPSSS LGTKTYTCNV DHKPSNTKVD KRVESKYGPP CPPCPAPEFL
240
GGPSVFLFPP KPKDTLMISR TPEVTCVVVD VSQEDPEVQF NWYVDGVEVH NAKTKPREEQ
300
FNSTYRVVSV LTVLHQDWLN GKEYKCKVSN KGLPSSIEKT ISKAKGQPRE PQVYTLPPSQ
360
EEMTKNQVSL TCLVKGFYPS DIAVEWESNG QPENNYKTTP PVLDSDGSFF LYSRLTVDKS
420
RWQEGNVFSC SVMHEALHNH YTQKSLSLSL G
451

Sequence Number (ID): 10
Length: 219
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..219
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSPLS LPVTPGEPAS ISCRSSQSLL YSIGYNYLDW YLQKSGQSPQ LLIYLGSNRA
60
SGVPDRFSGS GSGTDFTLKI SRVEAEDVGF YYCMQALQTP YTFGQGTKLE IKRTVAAPSV
120
FIFPPSDEQL KSGTASVVCL LNNFYPREAK VQWKVDNALQ SGNSQESVTE QDSKDSTYSL
180
SSTLTLSKAD YEKHKVYACE VTHQGLSSPV TKSFNRGEC
219

Sequence Number (ID): 11
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIF GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSPPWTF GQGTKVEIK
109

Sequence Number (ID): 12
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIY GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSPPWTF GQGTKVEIK
109

Sequence Number (ID): 13
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIF GASSRAPGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSPPWTF GQGTKVEIK
109

Sequence Number (ID): 14
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIY GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSPPWTF GQGTKVEIK
109

Sequence Number (ID): 15
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIF GASSRAPGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSPPWTF GQGTKVEIK
109

Sequence Number (ID): 16
Length: 109
Molecule Type: AA

Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIY GASSRAPGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSPPWTF GQGTKVEIK
109

Sequence Number (ID): 17
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIF GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSPPWTF GQGTKVEIK
109

Sequence Number (ID): 18
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIF GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSAGWTF GQGTKVEIK
109

Sequence Number (ID): 19
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIF GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 20
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein

> organism, synthetic construct

Residues:

EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIF GASSRATGIP
60

DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSPPWTF GQGTKVEIK
109

Sequence Number (ID): 21

Length: 109

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..109
  > mol_type, protein
  > organism, synthetic construct

Residues:

EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIY GASSRATGIP
60

DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSPPWTF GQGTKVEIK
109

Sequence Number (ID): 22

Length: 109

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..109
  > mol_type, protein
  > organism, synthetic construct

Residues:

EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIF GASSRAPGIP
60

DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSPPWTF GQGTKVEIK
109

Sequence Number (ID): 23

Length: 109

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..109
  > mol_type, protein
  > organism, synthetic construct

Residues:

EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIY GASSRAPGIP
60

DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 24

Length: 109

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..109
  > mol_type, protein
  > organism, synthetic construct

Residues:

EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIF GASSRAPGIP

60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 25
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIY GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 26
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIF GASSRAPGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 27
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIY GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 28
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIF GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 29
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SSYLAWYQQK PGQAPRLLIF GASSRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYGSSPPWTF GQGTKVEIK
109

Sequence Number (ID): 30
Length: 109
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..109
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS NSYLAWYQQK PGQAPRLLIY GASSRAPGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ QYDHSAGWTF GQGTKVEIK
109

Sequence Number (ID): 31
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 32
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVQPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 33
Length: 115

Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 34
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 35
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 36
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 37
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115

> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVESGGG LVHPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 38
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVQSGGG LVQPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 39
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 40
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 41
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:

EVQLVESGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 42
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVQPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 43
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 44
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 45
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS

115

Sequence Number (ID): 46
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGRSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 47
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL TCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 48
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMHWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 49
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGEGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 50

Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDEAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 51
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAGDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 52
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFD DYAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 53
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVQPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 54
Length: 115
Molecule Type: AA
Features Location/Qualifiers:

- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVESGGG LVHPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 55
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVESGGG LVQPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 56
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 57
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 58
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLVQSGGG LVHPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 59
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVQPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 60
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 61
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 62
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60

DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 63
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 64
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVQSGGG LVHPGGSLRL SCAGSGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATNYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDM AVYYCARGRY YFDYWGQGTL VTVSS
115

Sequence Number (ID): 65
Length: 115
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..115
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS RNAMFWVRQA PGKGLEWVSG IGTGGATSYA
60
DSVKGRFTIS RDNAKNSLYL QMNSLRAEDT AVYYCARGRY YFPWWGQGTL VTVSS
115

Sequence Number (ID): 66
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
RNAMF
5

Sequence Number (ID): 67
Length: 16

Molecule Type: AA
Features Location/Qualifiers:
- source, 1..16
> mol_type, protein
> organism, synthetic construct
Residues:
GIGTGGATSY ADSVKG
16

Sequence Number (ID): 68
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
GRYYFDY
7

Sequence Number (ID): 69
Length: 12
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..12
> mol_type, protein
> organism, synthetic construct
Residues:
RASQSVSSSY LA
12

Sequence Number (ID): 70
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
GASSRAT
7

Sequence Number (ID): 71
Length: 10
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..10
> mol_type, protein
> organism, synthetic construct
Residues:
QQYDHSAGWT
10

Sequence Number (ID): 72
Length: 7

Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
GASSRAP
7

Sequence Number (ID): 73
Length: 10
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..10
> mol_type, protein
> organism, synthetic construct
Residues:
QQYGSSPPWT
10

Sequence Number (ID): 74
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLDYWGKGT LVTVSS
116

Sequence Number (ID): 75
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSLSANY VFGTGTKLTV L
111

Sequence Number (ID): 76
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY

60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYNWGKGT LVTVSS
116

Sequence Number (ID): 77
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSQPPNP LFGTGTKLTV L
111

Sequence Number (ID): 78
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK LLKNPWGKGT LVTVSS
116

Sequence Number (ID): 79
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWFGTPASNY VFGTGTKLTV L
111

Sequence Number (ID): 80
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYNWGKGT LVTVSS
116

Sequence Number (ID): 81
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSSPPQP IFGTGTKLTV L
111

Sequence Number (ID): 82
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYDWGKGT LVTVSS
116

Sequence Number (ID): 83
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSSPPQP IFGTGTKLTV L
111

Sequence Number (ID): 84
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 85
Length: 111

Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTTYHP IFGTGTKLTV L
111

Sequence Number (ID): 86
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWWQYWGKGT LVTVSS
116

Sequence Number (ID): 87
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSSPPQP IFGTGTKLTV L
111

Sequence Number (ID): 88
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWWQYWGKGT LVTVSS
116

Sequence Number (ID): 89
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111

> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTTYHP IFGTGTKLTV L
111

Sequence Number (ID): 90
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYNWGKGT LVTVSS
116

Sequence Number (ID): 91
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTTMYP LFGTGTKLTV L
111

Sequence Number (ID): 92
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYDWGKGT LVTVSS
116

Sequence Number (ID): 93
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct

Residues:

QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTVLTP IFGTGTKLTV L
111

Sequence Number (ID): 94
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWFYDWGKGT LVTVSS
116

Sequence Number (ID): 95
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSPSMIP LFGTGTKLTV L
111

Sequence Number (ID): 96
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWFYDWGKGT LVTVSS
116

Sequence Number (ID): 97
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTTMYP LFGTGTKLTV L

111

Sequence Number (ID): 98
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYDWGKGT LVTVSS
116

Sequence Number (ID): 99
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTTLQP LFGTGTKLTV L
111

Sequence Number (ID): 100
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYNWGKGT LVTVSS
116

Sequence Number (ID): 101
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSPPTKP LFGTGTKLTV L
111

Sequence Number (ID): 102

Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYNWGKGT LVTVSS
116

Sequence Number (ID): 103
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTHRHP LFGTGTKLTV L
111

Sequence Number (ID): 104
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWLYNWGKGT LVTVSS
116

Sequence Number (ID): 105
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTTYHP IFGTGTKLTV L
111

Sequence Number (ID): 106
Length: 116
Molecule Type: AA
Features Location/Qualifiers:

```
                - source, 1..116
                        > mol_type, protein
                        > organism, synthetic construct
        Residues:
        QVQLVQSGAE  VKKPGASVKV  SCKASGYAFT  SYYMHWARQA  PGQGLEWMGI  INPSGGSTSY
60
        AQKFQGRVTM  TRDTSTSTVY  MELSSLRSED  TAVYYCARGK  WWWQHWGKGT  LVTVSS
116

        Sequence Number (ID): 107
        Length: 111
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..111
                        > mol_type, protein
                        > organism, synthetic construct
        Residues:
        QSVLTQPPSV  SAAPGQKVTI  SCSGGSSNIG  NSYVSWYQQL  PGTAPKLLIY  DNNKRPSGIP
60
        DRFSGSKSGT  SATLAITGLQ  TGDEADYYCG  TWDTSPVDRP  IFGTGTKLTV  L
111

        Sequence Number (ID): 108
        Length: 116
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..116
                        > mol_type, protein
                        > organism, synthetic construct
        Residues:
        QVQLVQSGAE  VKKPGASVKV  SCKASGYAFT  SYYMHWARQA  PGQGLEWMGI  INPSGGSTSY
60
        AQKFQGRVTM  TRDTSTSTVY  MELSSLRSED  TAVYYCARGK  WWWQHWGKGT  LVTVSS
116

        Sequence Number (ID): 109
        Length: 111
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..111
                        > mol_type, protein
                        > organism, synthetic construct
        Residues:
        QSVLTQPPSV  SAAPGQKVTI  SCSGGSSNIG  NSYVSWYQQL  PGTAPKLLIY  DNNKRPSGIP
60
        DRFSGSKSGT  SATLAITGLQ  TGDEADYYCG  TWDTSTTPMP  VFGTGTKLTV  L
111

        Sequence Number (ID): 110
        Length: 116
        Molecule Type: AA
        Features Location/Qualifiers:
                - source, 1..116
                        > mol_type, protein
                        > organism, synthetic construct
```

Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK WWWQHWGKGT LVTVSS
116

Sequence Number (ID): 111
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTTYHP IFGTGTKLTV L
111

Sequence Number (ID): 112
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 113
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 114
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSASY
60

AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 115
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEAVYFCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 116
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 117
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QPVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYFCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 118
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VRKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 119
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NNYVSWYQQL PGTAPKLLIY DNNKRPPGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 120
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPRGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 121
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYFCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 122
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPRGGSASY
60
AQKFQGRVSM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 123
Length: 111
Molecule Type: AA

Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT TATLAITGLQ TGDEADYYCG TWVTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 124
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 125
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYFCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 126
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VRKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRPED TAVYYCARGK YWMYDWGKGT QVTVSS
116

Sequence Number (ID): 127
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein

> organism, synthetic construct

Residues:

QSVLTQPPLV SAAPGQKVTI SCSGGSSNIG NSYVSWYQRL PGTAPKLLIY DNNKRPSGIP
60

DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 128

Length: 116

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..116
  > mol_type, protein
  > organism, synthetic construct

Residues:

QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60

AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGNGT LVTVSS
116

Sequence Number (ID): 129

Length: 111

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..111
  > mol_type, protein
  > organism, synthetic construct

Residues:

LPVLTQPPSV SAAPGQKVTI SCSGGSSSIG NSYVSWYQQL PGAAPKLLIY DNNKRPSGIP
60

DRFSGFRSGT SATLAITGLQ TGDEADYYCG TWDTSPVWEW PFGTGTKLTV L
111

Sequence Number (ID): 130

Length: 116

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..116
  > mol_type, protein
  > organism, synthetic construct

Residues:

QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60

AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT RVTVSS
116

Sequence Number (ID): 131

Length: 111

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..111
  > mol_type, protein
  > organism, synthetic construct

Residues:

QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP

60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSPVWEW PFGTGTKLTV L
111

Sequence Number (ID): 132
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 133
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQRL PGAAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 134
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 135
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSSIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWATSPVWEW PFGTGTKLTV L
111

Sequence Number (ID): 136
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 137
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYFCG TWDTSTAWEW PFGTGTKLTV L
111

Sequence Number (ID): 138
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE EKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 139
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSALTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYFCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 140
Length: 116

Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVSM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 141
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYFCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 142
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 143
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSPVWEW PFGTGTKLTV L
111

Sequence Number (ID): 144
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116

> mol_type, protein
> organism, synthetic construct

Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSASY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 145
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDSSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 146
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPRGGSTSY
60
AQKFQGRVAM TRDTSTSTVY MELSSLRPED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 147
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGGSSIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGVP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSPVWEW PFGTGTKLTV L
111

Sequence Number (ID): 148
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:

QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSASY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 149
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYFCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 150
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPRGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 151
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTAWEW PFGTGTKLTV L
111

Sequence Number (ID): 152
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS

116

Sequence Number (ID): 153
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDSSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 154
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VRKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWMGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRPED TAVYYCARGK YWMYDWGKGT LVTVSG
116

Sequence Number (ID): 155
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGSTNIG NSYVSWYQRL PGTAPKLLIY DNNKRPPGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTVWEW PFGTGTKLTV L
111

Sequence Number (ID): 156
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWARQA PGQGLEWVGI INPSGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSGD TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 157

Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QAVLTQPPSV SAAPGQKVTI SCSGGSSNIG NSYVSWYQRL PGAAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLAITGLQ TGDEADYYCG TWDTSTGWEW PFGTGTKLTV L
111

Sequence Number (ID): 158
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYAFT SYYMHWVRQA PGQGLEWMGI INPRGGSTSY
60
AQKFQGRVTM TRDTSTSTVY MELSSLRSED TAVYYCARGK YWMYDWGKGT LVTVSS
116

Sequence Number (ID): 159
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSV SAAPGQKVTI SCSGGGSSIG NSYVSWYQQL PGTAPKLLIY DNNKRPSGIP
60
DRFSGSKSGT SATLGITGLQ TGDEADYYCG TWDTSPVWEW PFGTGTKLTV L
111

Sequence Number (ID): 160
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
SYYMH
5

Sequence Number (ID): 161
Length: 17
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..17
> mol_type, protein

> organism, synthetic construct
Residues:
IINPRGGSTS YAQKFQG
17

Sequence Number (ID): 162
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
GKYWMYD
7

Sequence Number (ID): 163
Length: 13
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..13
> mol_type, protein
> organism, synthetic construct
Residues:
SGGGSSIGNS YVS
13

Sequence Number (ID): 164
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
DNNKRPS
7

Sequence Number (ID): 165
Length: 12
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..12
> mol_type, protein
> organism, synthetic construct
Residues:
GTWDTSPVWE WP
12

Sequence Number (ID): 166
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein

> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAVSGFTFS NYAMSWVRQA PGKGLEWVSA ISSGGGNIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCAKLR RYFDYWGQGT LVTVSS
116

Sequence Number (ID): 167
Length: 121
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..121
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS DYAMSWVRQA PGKGLEWVSA ISSGGSSIYY
60
ADSVKGRFTI SRDNSKNTLH LQMNSLRAED TAVYYCARGP QRSATAVFDY WGQGTLVTVS
120
S
121

Sequence Number (ID): 168
Length: 127
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..127
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSW ISPNSGNIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARRP LSAAWSHSSY YNAMDVWGQG
120
TLVTVSS
127

Sequence Number (ID): 169
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS GYAMSWVRQA PGKGLEWVSL ISHSGSNTYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARPH RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 170
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116

> mol_type, protein
> organism, synthetic construct

Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSG ISHGSGSIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARPH RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 171
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSG ISHGNGSIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCAKTG RHFDYWGQGT LVTVSS
116

Sequence Number (ID): 172
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSS ISPSGSSIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARSY RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 173
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSA ISPSGGSIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARAK RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 174
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct

Residues:

EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSA ISPGSGSTYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCAKFR RHFDYWGQGT LVTVSS
116

Sequence Number (ID): 175
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSA ISSGGGNIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARVH RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 176
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSA ITSSGRSIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARVH RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 177
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSA ITSSGANIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARVH RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 178
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSA ITSSGGNIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARVH RAFDYWGQGT LVTVSS

116

Sequence Number (ID): 179
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS NYAMSWVRQA PGKGLEWVSA ITAGGGSIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARVH RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 180
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTFS RHAMAWVRQA PGKGLEWVSA ITSSGRSIYY
60
ADSVKGRFTI SRDNSKNTLY LQMNSLRAED TAVYYCARVH RAFDYWGQGT LVTVSS
116

Sequence Number (ID): 181
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG NNYVNWYQQL PGTAPKLLIY DNSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDASLSAYV FGGGTKLTVL
110

Sequence Number (ID): 182
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG NNNVSWYQQL PGTAPKLLIY ANSKRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG SWDDSLSAYV FGGGTKLTVL
110

Sequence Number (ID): 183

Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA PGTPGQRVTI SCTGSSSNIG SNSVNWYQQL PGTAPKLLIY DDSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCD AWDSSLSAYV FGGGTKLTVL
110

Sequence Number (ID): 184
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTL SCTGSSSNIG SNYVSWYQQL PGTAPKLLIY ADSQRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDDSLSGYV FGGGTKLTVL
110

Sequence Number (ID): 185
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSSSSSNIG SNYVSWYQQL PGTAPKLLIY SDSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG SWDYSLSAYV FGGGTKLTVL
110

Sequence Number (ID): 186
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCTGSSSNIG NNTVSWYQQL PGTAPKLLIY DNSHRPSGVP
60
DRFSGSKSGT SASLAISGLQ SEDEADYYCG SWDYSLSAYV FGGGTKLTVL
110

Sequence Number (ID): 187
Length: 110
Molecule Type: AA
Features Location/Qualifiers:

- source, 1..110
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSA SGTPGQRVTI SCTGSSSNIG NNDVNWYQQL PGTAPKLLIY YDSQRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCA TWDASLSAYV FGGGTKLTVL
110

Sequence Number (ID): 188
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG SNAVNWYQQL PGTAPKLLIY YDNQRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDDSLNGYV FGGGTKLTVL
110

Sequence Number (ID): 189
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG NNAVTWYQQL PGTAPKLLIY DDSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG SWDYSLSAYV FGGGTKLTVL
110

Sequence Number (ID): 190
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG SNTFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL
110

Sequence Number (ID): 191
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct

Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG SNTFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL
110

Sequence Number (ID): 192
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG SNTFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLRGYV LGGGTKLTVL
110

Sequence Number (ID): 193
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG SNTFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG YWDYSLSGYV LGGGTKLTVL
110

Sequence Number (ID): 194
Length: 110
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..110
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSSNIG SNTFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL
110

Sequence Number (ID): 195
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSSANSR TDGFNWYQQL PGTAPKLLIY ADSHRPSGVP
60

DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL G
111

Sequence Number (ID): 196
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSAQFGS RDNFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL G
111

Sequence Number (ID): 197
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSTKQMH NYQFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL G
111

Sequence Number (ID): 198
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSLLRGE NLQFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL G
111

Sequence Number (ID): 199
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSPLFPD SGSFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL G
111

Sequence Number (ID): 200
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
QSVLTQPPSA SGTPGQRVTI SCSGSAALDL SPSFNWYQQL PGTAPKLLIY ADSHRPSGVP
60
DRFSGSKSGT SASLAISGLR SEDEADYYCG TWDYSLSGYV LGGGTKLTVL G
111

Sequence Number (ID): 201
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
RHAMA
5

Sequence Number (ID): 202
Length: 17
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..17
> mol_type, protein
> organism, synthetic construct
Residues:
AITSSGRSIY YADSVKG
17

Sequence Number (ID): 203
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
VHRAFDY
7

Sequence Number (ID): 204
Length: 13
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..13
> mol_type, protein
> organism, synthetic construct
Residues:
SGSPLFPDSG SFN

13

Sequence Number (ID): 205
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
ADSHRPS
7

Sequence Number (ID): 206
Length: 11
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..11
> mol_type, protein
> organism, synthetic construct
Residues:
GTWDYSLSGY V
11

Sequence Number (ID): 207
Length: 117
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..117
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYTFT NYGISWVRQA PGQGLEWMGW ISVYNGKTNY
60
AQKLQGRVTM TTDTSTTTAY MEMRSLRSDD TAVYYCARGS GYDLDYWGQG TLVSVSS
117

Sequence Number (ID): 208
Length: 128
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..128
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS SFWMTWVRQA PGKGLEWVAN IKQDGSEKYY
60
VDSVKGRFTI SRDNAKNSLY LQMNSLRAED TAVYYCARDP GRTMVRGGIR YYYGMDVWGQ
120
GTTVTVSS
128

Sequence Number (ID): 209
Length: 122
Molecule Type: AA

Features Location/Qualifiers:
- source, 1..122
> mol_type, protein
> organism, synthetic construct
Residues:
EVKLAESGGG LVQPGGSLRL SCAASGFTFS SHWMNWVRQA PGKGLEWVAN IKQDGSDKYY
60
VDSVKGRFTI SRDNAKNSLY LQLNSLIAED TAVYYCARDR GVRPPRGAFD IWGQGTMVTV
120
SS
122

Sequence Number (ID): 210
Length: 128
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..128
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYTFN SYGISWVRQA PGQGLEWMGW IRTYNGNTNY
60
AQKLQGRVTM TTDTSTSTAY MELRSLRSDD TAVYYCARDE ARIVVAGTTP YYYGMDVWGQ
120
GTTVTVSS
128

Sequence Number (ID): 211
Length: 117
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..117
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVESGGG LVQPGGSLRL SCAVSGFTIS DHYMSWIRQA PGKGLEWISY ISSSGSKIYY
60
ADSVKGRFTI SRDNAKNSLF LQMNSLRAED TAVYYCARTR QLVGDYWGQG TLVTVSS
117

Sequence Number (ID): 212
Length: 124
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..124
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGRSLRL SCAASGFTFD NYAMHWVRQA PGKGLEWVSG IRWNSGSIGY
60
ADSVKGRFTI SRDNAKNSLY LQMNSLRAED TALYYCAKEG GYSGYRPGPF FDYWGQGTLV
120
TVSS
124

Sequence Number (ID): 213
Length: 117
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..117
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYTFT NYGISWVRQA PGQGLEWMGW ISVYNGHTNY
60
AQKLQGRVTM TTDTSTSTAY MELRSLRSDD TAVYYCARGS GYDFDSWGQG TLVTVSS
117

Sequence Number (ID): 214
Length: 116
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..116
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASRYTFT SYDINWVRQA TGQGLEWMGW MNPNSGNTGY
60
AQKFQGRVTM TRNTSTSTAY MELSSLRSED TAVYYCARVR RFFDYWGQGT LVTVSS
116

Sequence Number (ID): 215
Length: 117
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..117
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGPE VKKPGASVKV SCKASGYTFT NYGISWVRQA PGQGLEWMGW ISVYNGNINY
60
AQKLQGRVTM TTDTSTSTAY MDLRSLRSDD TAVYYCARGS GYDFDYWGQG TLVTVSS
117

Sequence Number (ID): 216
Length: 122
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..122
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKDSAYTFN RYGISWVRQA PGQGLEWMGW ISAYTGNTVY
60
AQKLQGRVTM TTDNSTSTAY MELRSLRSDD TAVYYCARDK SIFGVVRGFD YWGQGTLVTV
120
SS
122

Sequence Number (ID): 217

Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
AIQMTQSPSS LSASVGDRVT ITCRASQGIR NALGWYQQKP GKAPKLLIYA ASSLQSGVPS
60
RFSGSGSGTD FTLTFSSLQP EDFATYYCLQ DFNYPYTFGQ GTKLEIK
107

Sequence Number (ID): 218
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS VSASVGDRVT ISCRASQGVS SWLAWYQQKP GNAPKLLISA ASSIQSGVPS
60
RFSGSGSGTD FTLTISSLQP EDFATYYCQQ ANSFPLTFGG GTKVEIK
107

Sequence Number (ID): 219
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS VSASVGDRVT ITCRASQGIS SWLAWYQQKP GKAPKLLIYA ASSFQSGVPS
60
RFSGSGSGTD FTLTISSLQP EDFATYFCQQ ANSFPLTFGG GTTVEIK
107

Sequence Number (ID): 220
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS VSASVGDRVT ITCRASQDIS IWLAWYQQSP GKAPKLLINV ASRLQSGVPS
60
RFSGSGSGTD FTLTINSLQP EDFVTYYCQQ ANSFPITFGQ GTRLATK
107

Sequence Number (ID): 221
Length: 107
Molecule Type: AA
Features Location/Qualifiers:

- source, 1..107
> mol_type, protein
> organism, synthetic construct

Residues:
DIQLTQSPSF LSASVGDRVT ITCWASQGIS SYLAWYQQKP GKAPKLLIFA ASTLQSGVPS
60
RFSGSGSGTE FTLTISSLQP EDFATYYCQQ LNSYPLTFGG GTKVEIR
107

Sequence Number (ID): 222
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct

Residues:
EIVMTQSPAT LSVSPGERAT LSCRASQSVN YNLAWYQHKP GQAPRLLIYG ASTRATGIPA
60
RFSGSGSGTE FTLTISSLQS EDFAVYYCQQ YNNWPLTFGG GTKVEIK
107

Sequence Number (ID): 223
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct

Residues:
AIQMTQSSSS LSASVGDRVT ITCRASQAIR NALGWYQQKP GKAPKVLIYA ASSLQSGIPS
60
RFSGSGSGTD FTLTISSLQP EDFATYYCLQ DYDYPYTFGQ GTKLEIK
107

Sequence Number (ID): 224
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct

Residues:
DIQLTQSPSF LSASVGDRVT ITCWASQGII SYLAWYQQKP GKAPKLLIYA ASTLHSGVPS
60
RFSGSGSGTE FTLTISSLQP EDFATYYCHQ LKSYPITFGQ GTRLEIK
107

Sequence Number (ID): 225
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct

Residues:
AIQMTQSPSS LSASVGDRVT ITCRASQDIR NALGWYQQKP GKAPKLLIYA ASSLQSGVPS
60
RFSGSASGTD FTLTISSLQP EDFAAYYCLQ DYNYPYTFGQ GTKLEIK
107

Sequence Number (ID): 226
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
EIVMTQSPVT LSLSPGERAT LPCRASQSVS SSLAWYQQKA GQSPRLLIYG ASTRATGIPA
60
RFSGSGSGTE FTLTISNLQS EDFAVYYCQQ YNNWPLTFGG GTKVEIK
107

Sequence Number (ID): 227
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISSNGGSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VGYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 228
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPSGSSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRSK VRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 229
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPSGVSTYY
60

ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VKYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 230
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPTSGSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 231
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPTGTSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK GAYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 232
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISSSGSSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VAYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 233
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPSSTSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VLYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 234
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPSSASTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK SKYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 235
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISGNSASTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK LKYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 236
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISHSGTSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVR VLYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 237
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPSGVSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VKYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 238
Length: 120
Molecule Type: AA

Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISSNGGSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVF VRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 239
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPTSASTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK GRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 240
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPTGGSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK GRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 241
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISHSGNSTYY
60
ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK RRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 242
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein

> organism, synthetic construct

Residues:

EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISPSSNSTYY
60

ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 243

Length: 120

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..120

> mol_type, protein

> organism, synthetic construct

Residues:

EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISSSGSSTYY
60

ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK PAYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 244

Length: 120

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..120

> mol_type, protein

> organism, synthetic construct

Residues:

EVQLLESGGG LVQPGGSLRL SCAASGFTLS SYAMHWVRQA PGKGLEYVSG ISYSSASTYY
60

ANSVKGRFTI SRDNPKNTLF LQMSSLRAED TAVYYCVRVK VRYRGGMDVW GQGTTVTVSS
120

Sequence Number (ID): 245

Length: 108

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..108

> mol_type, protein

> organism, synthetic construct

Residues:

ETTLTQSPDT LPLSPGDRAS LSCRASQSVS SAYLAWYQQK PGQAPRLLIY GTSRRATGVP
60

GRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGSSSVTFG QGTKLEIK
108

Sequence Number (ID): 246

Length: 108

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..108

> mol_type, protein

> organism, synthetic construct

Residues:

EIVLTQSPGT LSLSPGERAT LSCRASQGIS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP

60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 247
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQGIS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 248
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGASSVTFG QGTKLEIK
108

Sequence Number (ID): 249
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGASSVTFG QGTKLEIK
108

Sequence Number (ID): 250
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQGIS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 251
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGASSVTFG QGTKLEIK
108

Sequence Number (ID): 252
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSIS TAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGASSVTFG QGTKLEIK
108

Sequence Number (ID): 253
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQDIS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 254
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQDVS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 255
Length: 108

Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQNIS TAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 256
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQDAS NAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGSSSVTFG QGTKLEIK
108

Sequence Number (ID): 257
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQGVS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGRSSVTFG QGTKLEIK
108

Sequence Number (ID): 258
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQNIS TAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGTSSVTFG QGTKLEIK
108

Sequence Number (ID): 259
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108

> mol_type, protein
> organism, synthetic construct

Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQSVS TAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 260
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct

Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQDIS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 261
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct

Residues:
EIVLTQSPGT LSLSPGERAT LSCRASQGVS TAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGATSVTFG QGTKLEIK
108

Sequence Number (ID): 262
Length: 108
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..108
> mol_type, protein
> organism, synthetic construct

Residues:
EIVLPQSPGT LSLSPGERAT LSCRASQGVS SAYLAWYQQK PGQAPRLLIY GTSRRATGIP
60
DRFSGSGSGT DFTLTISRLE PEDFAVYYCQ LYGSTSVTFG QGTKLEIK
108

Sequence Number (ID): 263
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct

Residues:

EVQLVESGGG LVQPKGSLKL SCAASGFTFN TYGMHWVRQA PGKGLEWVAH IRSKSSNYAT
60
YYADSVKDRF TISRDDSQSM LYLQMNNLKT EDTAMYYCVR WFRAMDYWGQ GTSVTVSS
118

Sequence Number (ID): 264
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLIESGGG LVQPKGSLKL SCAASGFTFN MYAMDWVRQA PGKGLEWVAR IRSKGSNFET
60
NYADSVKDRF TISRDDSQSM VYLQMINLKT EDTAMYYCVR HRGGAWFAYW GQGTLVSVSA
120

Sequence Number (ID): 265
Length: 121
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..121
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVETGGG LVRPGNSLKL SCVTSGFTFS NYRMHWLRQP PGKRLEWIAV ITVKSNNYGA
60
NYAESVKGRF AISRDDSKSS VYLEMNRLRE EDTATYFCSR ERAYGNPFDY WGQGTTLTVS
120
S
121

Sequence Number (ID): 266
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPKGSLKL SCAASGFTFN MYAMNWVRQA PGQGLEWVAR IRSKSNNYAT
60
YYADSVKDRF IISRDDSESM VYLQMSNLRA ADTAMYYCVR HLRAMDYWGQ GTSVTVSS
118

Sequence Number (ID): 267
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPKGSLKL SCAASGFSFN MYAMNWVRQA PGKGLEWVAR IRTKSNHYST

60
YYADSVKDRF TISRDDSASM FYLQMNNLKT EDTAMYFCVR HLRAMDYWGQ GTSVTVSS
118

Sequence Number (ID): 268
Length: 120
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..120
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLIESGGG LVQPKGSLKL SCAASGFTFN MYAMDWVRQA PGKGLEWVAR IRSKGSNFET
60
NYADSVKDRF TISRDDSQSM VYLQMNNLKT EDTAMYYCVR HRGGAWFAYW GQGTLVTVSA
120

Sequence Number (ID): 269
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVRPKGSLKL SCAASGFSFN TYAMNWVRQA PGKGLEWIVW IRSKSHNYAT
60
YYADSVKDRF TISRDDSESM LYLQMNNLKT EDTAMYYCVR HLRAMDYWGQ GTSVTVSS
118

Sequence Number (ID): 270
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVRLVESGGG LVQPKGSLKL SCEASGFSFN MYAMNWVRQA PGKGLEWITH IRSKSNNYAT
60
YYADSVKDRF IISRDDSESM VYLQMNNLKT EDTAMYYCVR LLRALDYWGQ GTSVTVSS
118

Sequence Number (ID): 271
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPKGSLKL SCAASGFTFN MYGMHWVRQA PGKGLEWVAH IRSKSSNYAT
60
YYADSVKDRL TISRDDSQSM LYLQMNNLKT EDTAMYYCVR WFRAMDYWGQ GTSVTVSS
118

Sequence Number (ID): 272
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVKPGGSLRL SCAASGFTFS MYGMHWVRQA PGKGLEWVGH IRSKSSNYAT
60
YYADSVKDRF TISRDDSKNT LYLQMNSLKT EDTAVYYCTT WFRAMDYWGQ GTLVTVSS
118

Sequence Number (ID): 273
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVKPGGSLRL SCAASGFTFS MYGMHWVRQA PGKGLEWVGH IRSKSSNYAT
60
YYADSVKDRF TISRDDSKNT LYLQMNSLKT EDTAVYYCTT WFRAMDYWGQ GTLVTVSS
118

Sequence Number (ID): 274
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS MYGMHWVRQA PGKGLEWVSH IRSKSSNYAT
60
YYADSVKDRF TISRDNAKNS LYLQMNSLRA EDTAVYYCAR WFRAMDYWGQ GTLVTVSS
118

Sequence Number (ID): 275
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLKL SCAASGFTFS MYGMHWVRQA SGKGLEWVGH IRSKSSNYAT
60
YYADSVKDRF TISRDDSKNT AYLQMNSLKT EDTAVYYCTR WFRAMDYWGQ GTLVTVSS
118

Sequence Number (ID): 276
Length: 118

Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS MYGMHWVRQA PGKGLEWVGH IRSKSSNYAT
60
YYADSVKDRF TISRDDSKNS LYLQMNSLKT EDTAVYYCAR WFRAMDYWGQ GTLVTVSS
118

Sequence Number (ID): 277
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLRL SCAASGFTFS MYGMHWVRQA PGKGLEWVSH IRSKSSNYAT
60
YYADSVKDRF TISRDNAKNS LYLQMNSLRA EDTAVYYCAR WFRAMDYWGQ GTLVTVSS
118

Sequence Number (ID): 278
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LEQPGGSLRL SCAGSGFTFR MYGMHWVRQA PGKGLEWVSH IRSKSSNYAT
60
YYADSVKDRF TISRDNSKNT LYLQMNSLRA EDTAVYYCAK WFRAMDYWGQ GTTVTVSS
118

Sequence Number (ID): 279
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLKL SCAASGFTFS MYGMHWVRQA SGKGLEWVGH IRSKSSNYAT
60
YYADSVKDRF TISRDDSKNT AYLQMNSLKT EDTAVYYCTR WFRAMDYWGQ GTTVTVSS
118

Sequence Number (ID): 280
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107

> mol_type, protein
> organism, synthetic construct

Residues:
DIVMTQSHKF MSTSVGDRVS ITCKASQDVS TAVAWYQEKP GQSPKLLIYW ASTRHTGVPD
60
RFTGSGSGTD YTLTISSVQA EDLALYYCQQ HYSTPLTFGA GTKLELK
107

Sequence Number (ID): 281
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct

Residues:
DIVVTQSPAS LAVSLGQRAT ISCRASKSVS TSGYSYMHWY QQKPGQPPKL LIYLASNLES
60
GVPARFSGSG SGTDFTLNIH PVEEEDVAIY YCQHSRELPL TFGAGTKLEL K
111

Sequence Number (ID): 282
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct

Residues:
DIQMTQSPSS LSASLGERVS LTCRASQEIS GYLSWLQQKP DGTIKRLIYA ASTLDSGVPK
60
RFSGSRSGSD YSLTISSLES EDFADYYCLQ YGSYPYTFGG GTKLEIK
107

Sequence Number (ID): 283
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct

Residues:
DIVLTQSPAS LTVSLGQRAT ISCRASKSVS TSGYSYMHWY QQKPGQPPKL LIYLASNLES
60
GVPARFSGSG SGTDFTLNIH PVEEEDAATY YCQHSRELPI TFGSGTKLEI K
111

Sequence Number (ID): 284
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct

Residues:

DIVLTQSPAS LVVSLGQRAT ISCRASQSVS TSGYSYMHWY QQKPGQPPKL LIYLASNVQS
60
GVPARFSGSG SGTDFTLNIH PVEEEDVATY YCHHNRDLPF TFGSGTKLEI K
111

Sequence Number (ID): 285
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
DIVVTQSPAS LAVSLGQRAT ISCRASKSVS TSGYSYMHWY QQKPGQPPKL LIYLASNLES
60
GVPARFSGSG SGTDFTLNIH PVEEEDVAIY YCQHSRELPL TFGAGTKLEL K
111

Sequence Number (ID): 286
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
DIVLTQSPAS LAVSLGQRAT ISCRASKSVS ASGYSYMHWY QQKPGQPPKL LIYLASNLQS
60
GVPARFSGSG SGTDFTLNIH PVEEEDAATY YCQHSRELPP TFGGGTKLEI K
111

Sequence Number (ID): 287
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
DIVLTQSPAS LAVFLGQRAT ISCRASKSVS TSGYSYMHWY QQKAGQPPKL LIYLASNLES
60
GVPARFSGSG SGTDFTLNIH PVEEEDAATY YCHHSRELPI TFGSGTKLEM K
111

Sequence Number (ID): 288
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSHKF MSTSVGDRVS ITCKASQDVS TAVAWYQEKP GQSPKLLIYW ASTRHTGVPD
60
RFTGSGSGTD YTLTISSVQA EDLALYYCQQ HYSTPLTFGA GTKLELK

107

Sequence Number (ID): 289
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
EIVLTQSPGT LSLSPGERAT LSCKASQDVS TAVAWYQQKP GQAPRLLIYW ASTRHTGIPD
60
RFSGSGSGTD FTLTISRLEP EDFAVYYCQQ HYSTPLTFGQ GTKVEIK
107

Sequence Number (ID): 290
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSPLS LPVTPGEPAS ISCKASQDVS TAVAWYLQKS GQSPQLLIYW ASTRHTGVPD
60
RFSGSGSGTD FTLKISRVEA EDVGFYYCQQ HYSTPLTFGQ GTKLEIK
107

Sequence Number (ID): 291
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSPDS LAVSLGERAT INCKASQDVS TAVAWYQQKP GQPPKLLIYW ASTRHTGVPD
60
RFSGSGSGTD FTLTISSLQA EDVAVYYCQQ HYSTPLTFGG GTKVEIK
107

Sequence Number (ID): 292
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS LSASVGDRVT ITCKASQDVS TAVAWYQQKP GKVPKLLIYW ASTRHTGVPS
60
RFSGSGSGTD FTLTISSLQP EDVATYYCQQ HYSTPLTFGG GTKVEIK
107

Sequence Number (ID): 293

Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQLTQSPSF LSASVGDRVT ITCKASQDVS TAVAWYQQKP GKAPKLLIYW ASTRHTGVPS
60
RFSGSGSGTE FTLTISSLQP EDFATYYCQQ HYSTPLTFGG GTKVEIK
107

Sequence Number (ID): 294
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS LSASVGDRVT ITCKASQDVS TAVAWYQQKP GKAPKLLLYW ASTRHTGVPS
60
RFSGSGSGTD YTLTISSLQP EDFATYYCQQ HYSTPLTFGG GTKVEIK
107

Sequence Number (ID): 295
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
AIRMTQSPFS LSASVGDRVT ITCKASQDVS TAVAWYQQKP AKAPKLFIYW ASTRHTGVPS
60
RFSGSGSGTD YTLTISSLQP EDFATYYCQQ HYSTPLTFGG GTKVEIK
107
END