Sequence Listing Information:
dtdVersion: V1_3
fileName: d242671bpwo-en.xml
softwareName: WIPO Sequence
softwareVersion: 3.1.2
productionDate: 2026-07-27
General Information:
Current application / Applicant file reference: D242671BPWO
Earliest priority application / IP Office: CN
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Applicant name: CSPC Megalith Biopharmaceutical Co., Ltd.
Applicant name / Language: en
Inventor name: Xiwu HUI
Inventor name / Language: en
Invention title: ANTI-DLL3 ANTIBODY AND USE (en)
SequenceTotalQuantity: 59
Sequences:
Sequence Number (ID): 1
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
DYNMY
5

Sequence Number (ID): 2
Length: 17
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..17
> mol_type, protein
> organism, synthetic construct
Residues:
YIDPYNGVTG FNQKFKG
17

Sequence Number (ID): 3
Length: 13
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..13
> mol_type, protein
> organism, synthetic construct
Residues:
GQGGYYIDYA MDY
13

Sequence Number (ID): 4
Length: 11
Molecule Type: AA

Features Location/Qualifiers:
- source, 1..11
> mol_type, protein
> organism, synthetic construct
Residues:
RASQDIGDTL A
11

Sequence Number (ID): 5
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
ATSNLDS
7

Sequence Number (ID): 6
Length: 9
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..9
> mol_type, protein
> organism, synthetic construct
Residues:
LQYAISPYT
9

Sequence Number (ID): 7
Length: 122
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..122
> mol_type, protein
> organism, synthetic construct
Residues:
EIQLQQSGPG LVNPGTSVKV SCKASGYSFT DYNMYWVKQS HGRSLEWIGY IDPYNGVTGF
60
NQKFKGKATL TVDKSSSTAF MHLNSLTSED SAVYYCSRGQ GGYYIDYAMD YWGQGTSVTV
120
SS
122

Sequence Number (ID): 8
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS LSASLGERVS LTCRASQDIG DTLAWLQQEP DGTIKRLIYA TSNLDSGVPK

60

RFSGSRSGSD FSLTISSLES EDFVHYYCLQ YAISPYTFGG GTKLEIK

107

Sequence Number (ID): 9
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
TYGMS

5

Sequence Number (ID): 10
Length: 17
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..17
> mol_type, protein
> organism, synthetic construct
Residues:
TIHSDGGDTY YPDSVKG

17

Sequence Number (ID): 11
Length: 9
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..9
> mol_type, protein
> organism, synthetic construct
Residues:
QTGPWLLTY

9

Sequence Number (ID): 12
Length: 11
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..11
> mol_type, protein
> organism, synthetic construct
Residues:
KASQAVNTAV A

11

Sequence Number (ID): 13
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein

> organism, synthetic construct
Residues:
WASTRHT
7

Sequence Number (ID): 14
Length: 9
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..9
> mol_type, protein
> organism, synthetic construct
Residues:
QQHYSTPYT
9

Sequence Number (ID): 15
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLVESGGG LVQPGGSLKL SCAASGFTFR TYGMSWVRQT PDKRLELVAT IHSDGGDTYY
60
PDSVKGRFTI SRDSAKNTLY LQMSSLKSED TAVYYCARQT GPWLLTYWGQ GTLVTVSA
118

Sequence Number (ID): 16
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSHRF MSTSVGDRVS ITCKASQAVN TAVAWYQQKP GQSPKLLIYW ASTRHTGVPD
60
RFTGSGAGTD YSLTISSVQA EDLALYYCQQ HYSTPYTFGG GTKLEIK
107

Sequence Number (ID): 17
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
NYGMK
5

Sequence Number (ID): 18

Length: 17
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..17
> mol_type, protein
> organism, synthetic construct
Residues:
WINTYTGEPT YADDFKG
17

Sequence Number (ID): 19
Length: 9
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..9
> mol_type, protein
> organism, synthetic construct
Residues:
FSDYGGFDY
9

Sequence Number (ID): 20
Length: 11
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..11
> mol_type, protein
> organism, synthetic construct
Residues:
KASQSVSNDV A
11

Sequence Number (ID): 21
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
YASNRYT
7

Sequence Number (ID): 22
Length: 9
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..9
> mol_type, protein
> organism, synthetic construct
Residues:
QQDYSSPFT
9

Sequence Number (ID): 23
Length: 118
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
QIQLVQSGPE LKKPGETVKI SCKASGYTFT NYGMKWVKQA PGKGLKWMGW INTYTGEPTY
60
ADDFKGRFAF SLETSASTAY LQINNLKNED TATYFCARFS DYGGFDYWGQ GTTLTVSS
118

Sequence Number (ID): 24
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
NIVMTQTPKF LLVSAGDRVT ITCKASQSVS NDVAWYQQKP GQSPKLLIYY ASNRYTGVPD
60
RFTGSGYGTD FTFTISTVQA EDLAVYFCQQ DYSSPFTFGS GTKLEIK
107

Sequence Number (ID): 25
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
DYYMK
5

Sequence Number (ID): 26
Length: 17
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..17
> mol_type, protein
> organism, synthetic construct
Residues:
AFNLNNGDTF YNQKFKG
17

Sequence Number (ID): 27
Length: 8
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..8
> mol_type, protein

> organism, synthetic construct
Residues:
DVYGYGDY
8

Sequence Number (ID): 28
Length: 15
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..15
> mol_type, protein
> organism, synthetic construct
Residues:
RASKSVSTSG YSYMH
15

Sequence Number (ID): 29
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
LASNLDS
7

Sequence Number (ID): 30
Length: 9
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..9
> mol_type, protein
> organism, synthetic construct
Residues:
QHSRELPYT
9

Sequence Number (ID): 31
Length: 117
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..117
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLQQSGPE LVKPGASVKM SCKASGYTFT DYYMKWVKQS HGKSLEWIGA FNLNNGDTFY
60
NQKFKGKATL TVDKSSSTAY MQLNSLTSED SAVYYCARDV YGYGDYWGQG TSVTVSS
117

Sequence Number (ID): 32
Length: 111
Molecule Type: AA

Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
DIVLTQSPAS LAVSLGQRAT ISCRASKSVS TSGYSYMHWY QQKPGQPPKL LIYLASNLDS
60
GVPARFSGSG SGTDFTLNIH PVEEEDAATY YCQHSRELPY TFGGGTKLEI K
111

Sequence Number (ID): 33
Length: 5
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..5
> mol_type, protein
> organism, synthetic construct
Residues:
NYWMH
5

Sequence Number (ID): 34
Length: 17
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..17
> mol_type, protein
> organism, synthetic construct
Residues:
AIYPGNSNTA YNQKFKG
17

Sequence Number (ID): 35
Length: 8
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..8
> mol_type, protein
> organism, synthetic construct
Residues:
DGYYGFAY
8

Sequence Number (ID): 36
Length: 11
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..11
> mol_type, protein
> organism, synthetic construct
Residues:
KASQNVGTNV A
11

Sequence Number (ID): 37
Length: 7
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..7
> mol_type, protein
> organism, synthetic construct
Residues:
SASYRYS
7

Sequence Number (ID): 38
Length: 9
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..9
> mol_type, protein
> organism, synthetic construct
Residues:
QQYKNYPYT
9

Sequence Number (ID): 39
Length: 117
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..117
> mol_type, protein
> organism, synthetic construct
Residues:
EVQLQQSGTV LARPGASVKM SCKASGYSFT NYWMHWMKQR PGQGLEWIGA IYPGNSNTAY
60
NQKFKGKAKL TAITSASSAY MELSSLTNED SAVYYCTRDG YYGFAYWGQG TLVTVSA
117

Sequence Number (ID): 40
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSQKF MSTSVGDRVS VTCKASQNVG TNVAWYQQKS GQFPKALIYS ASYRYSGVPD
60
RFTGSGSGTD FTLTISNVQS EDLAEYFCQQ YKNYPYTFGG GTKLEIK
107

Sequence Number (ID): 41
Length: 447
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..447
> mol_type, protein

> organism, synthetic construct

Residues:

EVQLQQSGPE LVKPGASVKM SCKASGYTFT DYYMKWVKQS HGKSLEWIGA FNLNNGDTFY
60

NQKFKGKATL TVDKSSSTAY MQLNSLTSED SAVYYCARDV YGYGDYWGQG TSVTVSSAST
120

KGPSVFPLAP SSKSTSGGTA ALGCLVEDYF PEPVTVSWNS GALTSGVHTF PAVLQSSGLY
180

SLSSVVTVPS SSLGTQTYIC NVNHKPSNTK VDEKVEPKSC DKTHTCPPCP APEAAGGPSV
240

FLFPPKPKDT LMISRTPEVT CVVVDVSHED PEVKFNWYVD GVEVHNAKTK PREEQYNSTY
300

RVVSVLTVLH QDWLNGKEYK CKVSNKALAA SIEKTISKAK GQPREPQVCT LPPSRDELTK
360

NQVSLSCAVK GFYPSDIAVE WESNGQPENN YKTTPPVLDS DGSFFLVSKL TVDKSRWQQG
420

NVFSCSVMHE ALHNHYTQKS LSLSPGK
447

Sequence Number (ID): 42

Length: 330

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..330

> mol_type, protein

> organism, synthetic construct

Residues:

ASTKGPSVFP LAPSSKSTSG GTAALGCLVE DYFPEPVTVS WNSGALTSGV HTFPAVLQSS
60

GLYSLSSVVT VPSSSLGTQT YICNVNHKPS NTKVDEKVEP KSCDKTHTCP PCPAPEAAGG
120

PSVFLFPPKP KDTLMISRTP EVTCVVVDVS HEDPEVKFNW YVDGVEVHNA KTKPREEQYN
180

STYRVVSVLT VLHQDWLNGK EYKCKVSNKA LAASIEKTIS KAKGQPREPQ VCTLPPSRDE
240

LTKNQVSLSC AVKGFYPSDI AVEWESNGQP ENNYKTTPPV LDSDGSFFLV SKLTVDKSRW
300

QQGNVFSCSV MHEALHNHYT QKSLSLSPGK
330

Sequence Number (ID): 43

Length: 452

Molecule Type: AA

Features Location/Qualifiers:

- source, 1..452

> mol_type, protein

> organism, synthetic construct

Residues:

QIQLVQSGPE LKKPGETVKI SCKASGYTFT NYGMKWVKQA PGKGLKWMGW INTYTGEPTY
60

ADDFKGRFAF SLETSASTAY LQINNLKNED TATYFCARFS DYGGFDYWGQ GTTLTVSSAS
120

VAAPSVFIFP PSDEQLKSGT ASVVCLLNNF YPREAKVQWK VDNALQSGNS QESVTEQDSK
180

DSTYSLSSTL TLSKADYEKH KVYACEVTHQ GLSSPVTKSF NRGECDKTHT CPPCPAPEAA
240
GGPSVFLFPP KPKDTLMISR TPEVTCVVVD VSHEDPEVKF NWYVDGVEVH NAKTKPREEQ
300
YNSTYRVVSV LTVLHQDWLN GKEYKCKVSN KALAASIEKT ISKAKGQPRE PQVYTLPPCR
360
DELTKNQVSL WCLVKGFYPS DIAVEWESNG QPENNYKTTP PVLDSDGSFF LYSKLTVDKS
420
RWQQGNVFSC SVMHEALHNH YTQKSLSLSP GK
452

Sequence Number (ID): 44
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
ASVAAPSVFI FPPSDEQLKS GTASVVCLLN NFYPREAKVQ WKVDNALQSG NSQESVTEQD
60
SKDSTYSLSS TLTLSKADYE KHKVYACEVT HQGLSSPVTK SFNRGEC
107

Sequence Number (ID): 45
Length: 227
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..227
> mol_type, protein
> organism, synthetic construct
Residues:
DKTHTCPPCP APEAAGGPSV FLFPPKPKDT LMISRTPEVT CVVVDVSHED PEVKFNWYVD
60
GVEVHNAKTK PREEQYNSTY RVVSVLTVLH QDWLNGKEYK CKVSNKALAA SIEKTISKAK
120
GQPREPQVYT LPPCRDELTK NQVSLWCLVK GFYPSDIAVE WESNGQPENN YKTTPPVLDS
180
DGSFFLYSKL TVDKSRWQQG NVFSCSVMHE ALHNHYTQKS LSLSPGK
227

Sequence Number (ID): 46
Length: 218
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..218
> mol_type, protein
> organism, synthetic construct
Residues:
DIVLTQSPAS LAVSLGQRAT ISCRASKSVS TSGYSYMHWY QQKPGQPPKL LIYLASNLDS
60
GVPARFSGSG SGTDFTLNIH PVEEEDAATY YCQHSRELPY TFGGGTKLEI KRTVAAPSVF
120
IFPPSDRKLK SGTASVVCLL NNFYPREAKV QWKVDNALQS GNSQESVTEQ DSKDSTYSLS

180

STLTLSKADY EKHKVYACEV THQGLSSPVT KSFNRGEC

218

Sequence Number (ID): 47
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
RTVAAPSVFI FPPSDRKLKS GTASVVCLLN NFYPREAKVQ WKVDNALQSG NSQESVTEQD

60

SKDSTYSLSS TLTLSKADYE KHKVYACEVT HQGLSSPVTK SFNRGEC

107

Sequence Number (ID): 48
Length: 212
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..212
> mol_type, protein
> organism, synthetic construct
Residues:
NIVMTQTPKF LLVSAGDRVT ITCKASQSVS NDVAWYQQKP GQSPKLLIYY ASNRYTGVPD

60

RFTGSGYGTD FTFTISTVQA EDLAVYFCQQ DYSSPFTFGS GTKLEIKSSA STKGPSVFPL

120

APSSKSTSGG TAALGCLVKD YFPEPVTVSW NSGALTSGVH TFPAVLQSSG LYSLSSVVTV

180

PSSSLGTQTY ICNVNHKPSN TKVDKKVEPK SC

212

Sequence Number (ID): 49
Length: 105
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..105
> mol_type, protein
> organism, synthetic construct
Residues:
SSASTKGPSV FPLAPSSKST SGGTAALGCL VKDYFPEPVT VSWNSGALTS GVHTFPAVLQ

60

SSGLYSLSSV VTVPSSSLGT QTYICNVNHK PSNTKVDKKV EPKSC

105

Sequence Number (ID): 50
Length: 456
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..456
> mol_type, protein
> organism, synthetic construct

Residues:
EIQLQQSGPG LVNPGTSVKV SCKASGYSFT DYNMYWVKQS HGRSLEWIGY IDPYNGVTGF
60
NQKFKGKATL TVDKSSSTAF MHLNSLTSED SAVYYCSRGQ GGYYIDYAMD YWGQGTSVTV
120
SSASVAAPSV FIFPPSDEQL KSGTASVVCL LNNFYPREAK VQWKVDNALQ SGNSQESVTE
180
QDSKDSTYSL SSTLTLSKAD YEKHKVYACE VTHQGLSSPV TKSFNRGECD KTHTCPPCPA
240
PEAAGGPSVF LFPPKPKDTL MISRTPEVTC VVVDVSHEDP EVKFNWYVDG VEVHNAKTKP
300
REEQYNSTYR VVSVLTVLHQ DWLNGKEYKC KVSNKALAAS IEKTISKAKG QPREPQVYTL
360
PPCRDELTKN QVSLWCLVKG FYPSDIAVEW ESNGQPENNY KTTPPVLDSD GSFFLYSKLT
420
VDKSRWQQGN VFSCSVMHEA LHNHYTQKSL SLSPGK
456

Sequence Number (ID): 51
Length: 212
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..212
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS LSASLGERVS LTCRASQDIG DTLAWLQQEP DGTIKRLIYA TSNLDSGVPK
60
RFSGSRSGSD FSLTISSLES EDFVHYYCLQ YAISPYTFGG GTKLEIKSSA STKGPSVFPL
120
APSSKSTSGG TAALGCLVKD YFPEPVTVSW NSGALTSGVH TFPAVLQSSG LYSLSSVVTV
180
PSSSLGTQTY ICNVNHKPSN TKVDKKVEPK SC
212

Sequence Number (ID): 52
Length: 448
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..448
> mol_type, protein
> organism, synthetic construct
Residues:
QIQLVQSGPE LKKPGETVKI SCKASGYTFT NYGMKWVKQA PGKGLKWMGW INTYTGEPTY
60
ADDFKGRFAF SLETSASTAY LQINNLKNED TATYFCARFS DYGGFDYWGQ GTTLTVSSAS
120
TKGPSVFPLA PSSKSTSGGT AALGCLVEDY FPEPVTVSWN SGALTSGVHT FPAVLQSSGL
180
YSLSSVVTVP SSSLGTQTYI CNVNHKPSNT KVDEKVEPKS CDKTHTCPPC PAPEAAGGPS
240
VFLFPPKPKD TLMISRTPEV TCVVVDVSHE DPEVKFNWYV DGVEVHNAKT KPREEQYNST
300
YRVVSVLTVL HQDWLNGKEY KCKVSNKALA ASIEKTISKA KGQPREPQVC TLPPSRDELT

360
KNQVSLSCAV KGFYPSDIAV EWESNGQPEN NYKTTPPVLD SDGSFFLVSK LTVDKSRWQQ
420
GNVFSCSVMH EALHNHYTQK SLSLSPGK
448

Sequence Number (ID): 53
Length: 214
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..214
> mol_type, protein
> organism, synthetic construct
Residues:
NIVMTQTPKF LLVSAGDRVT ITCKASQSVS NDVAWYQQKP GQSPKLLIYY ASNRYTGVPD
60
RFTGSGYGTD FTFTISTVQA EDLAVYFCQQ DYSSPFTFGS GTKLEIKRTV AAPSVFIFPP
120
SDRKLKSGTA SVVCLLNNFY PREAKVQWKV DNALQSGNSQ ESVTEQDSKD STYSLSSTLT
180
LSKADYEKHK VYACEVTHQG LSSPVTKSFN RGEC
214

Sequence Number (ID): 54
Length: 122
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..122
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYTFT DYNMYWVRQA PGQRLEWMGY IDPYNGVTGF
60
NQKFKGRVTI TRDTSASTAY MELSSLRSED TAVYYCARGQ GGYYIDYAMD YWGQGTSVTV
120
SS
122

Sequence Number (ID): 55
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIQMTQSPSS LSASVGDRVT ITCRASQDIG DTLAWFQQKP GKAPKRLIYA TSNLDSGVPS
60
RFSGSRSGTD FTLTISSLQP EDFATYYCLQ YAISPYTFGG GTKLEIK
107

Sequence Number (ID): 56
Length: 118
Molecule Type: AA

Features Location/Qualifiers:
- source, 1..118
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYTFT NYGMKWVRQA PGQGLEWMGW INTYTGEPTY
60
ADDFKGRVTM TRDTSISTAY MELSRLRSDD TAVYYCARFS DYGGFDYWGQ GTTLTVSS
118

Sequence Number (ID): 57
Length: 107
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..107
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSPDS LAVSLGERAT INCKASQSVS NDVAWYQQKP GQPPKLLIYY ASNRYTGVPD
60
RFSGSGSGTD FTLTISSLQA EDVAVYYCQQ DYSSPFTFGS GTKLEIK
107

Sequence Number (ID): 58
Length: 117
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..117
> mol_type, protein
> organism, synthetic construct
Residues:
QVQLVQSGAE VKKPGASVKV SCKASGYTFT DYYMKWVRQA PGQRLEWMGA FNLNNGDTFY
60
NQKFKGRVTI TRDTSASTAY MELSSLRSED TAVYYCARDV YGYGDYWGQG TSVTVSS
117

Sequence Number (ID): 59
Length: 111
Molecule Type: AA
Features Location/Qualifiers:
- source, 1..111
> mol_type, protein
> organism, synthetic construct
Residues:
DIVMTQSPDS LAVSLGERAT INCRASKSVS TSGYSYMHWY QQKPGQPPKL LIYLASNLDS
60
GVPDRFSGSG SGTDFTLTIS SLQAEDVAVY YCQHSRELPY TFGGGTKLEI K
111
END