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FIG. 1 (SEQ ID NO:1)

1	A E T V E S C L A K P H T E N S F T N V W K	22
23	D D K T L D R Y A N Y E G C L W <u>N A T</u> G V V	44
45	V C T <u>G D E T Q C Y G T</u> W V P I G L A I P E	66
67	N <u>E G G G S E G G G S E G G G S E G G G</u> T K	88
89	P P E Y G D T P I P G Y T Y I N P L D G T Y	110
111	P P G T E Q N P A N P N P S L E E S Q P L N	132
133	T F <u>M F Q N N R F R N</u> R Q G A L T V Y T G T	154
155	F T Q G T D P V K T Y Y Q Y T P V S <u>S K A M</u>	176
177	<u>Y D A Y W</u> N G K F R D C A F H S G F N E D P	198
199	F V C E Y Q G Q S S D L P Q P P <u>V N A G G G</u>	220
221	<u>S G G</u> <u>G S G G G S E G G G S E G G G S E S G</u>	242
243	<u>G S E G G G S G G G S G S G</u> A <u>M V R S D K T</u>	264
265	<u>H T</u> C P P C P A P E L L G G P S V F L F P P	286
287	K P K D T L M I S R T P E V T C V V V D V S	308
309	H E D P E V K F N W Y V D G V E V H N A K T	330
331	K P R E E Q Y N S T Y R V V S V L T V L H Q	352
353	D W L N G K E Y K C K V S N K A L P A P I E	374
375	K T I S K A K G Q P R E P Q V Y T L P P S R	396
397	E E M T K N Q V S L T C L V K G F Y P S D I	418
419	A V E W E S N G Q P E N N Y K T T P P V L D	440
441	S D G S F F L Y S K L T V D K S R W Q Q G N	462
463	V F S C S V M H E A L H N H Y T Q K S L S L	484
485	S P G K	

FIG. 2 (SEQ ID NO:2)

1	A E T V E S C L A K P H T E N S F T N V W K	22
23	D D K T L D R Y A N Y E G C L W <u>N A T</u> G V V	44
45	V C T <u>G D E T Q C Y G T</u> W V P I G L A I P E	66
67	N E G G G S E G G G S E G G G S E G G G T K	88
89	P P E Y G D T P I P G Y T Y I N P L D G T Y	110
111	P P G T E Q N P A N P N P S L E E S Q P L N	132
133	T F <u>M F Q N N R F R N</u> R Q G A L T V Y T G T	154
155	F T Q G T D P V K T Y Y Q Y T P V S <u>S K A M</u>	176
177	<u>Y D A Y W</u> N G K F R D C A F H S G F N E D P	198
199	F V C E Y Q G Q S S D L P Q P P A N A G G E	220
221	S G G G S G G G S E G G G S E G G G S E G G	242
243	G S E G G G S G G G S G S G A R S D K T	262
263	H T C P P C P A P E L L G G P S V F L F P P	284
285	K P K D T L M I S R T P E V T C V V V D V S	306
307	H E D P E V K F N W Y V D G V E V H N A K T	328
329	K P R E E Q Y N S T Y R V V S V L T V L H Q	350
351	D W L N G K E Y K C K V S N K A L P A P I E	372
373	K T I S K A K G Q P R E P Q V Y T L P P S R	394
395	E E M T K N Q V S L T C L V K G F Y P S D I	416
417	A V E W E S N G Q P E N N Y K T T P P V L D	438
439	S D G S F F L Y S K L T V D K S R W Q Q G N	460
461	V F S C S V M H E A L H N H Y T Q K S L S L	482
483	S P G K	

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SEQ ID NO:3

ATGTACAGGATGCAACTCCTGTCTTGCATTGCACCTAAGTCTTGACCTTGTCACGAATTCGATGGCTGAAACTGTTGAAAG
TTGTTTAGCAAAACCCCATACAGAAATTCATTACTAACGCTCGAAAGACGACAAAACTTTAGATCGTTACGCTAACT
ATGAGGGCTGTCTGTGGAATGCTACAGCGCTTGTAGTTGTACTGGTGACGAAACTCAGTGTACGGTACATGGGTTCCCT
ATTGGGCTTGCTATCCCTGAATAATGAGGGTGGTGGCTCTGAGGGTGGCGGTTCTGAGGGTGGCGGTTCTGAGGGTGGCGG
TACTAAACCTCCTGAGTACGGTGATACACCTATTCGGGCTATACCTTATCAACCTCTCGACGGCACTTATCCGCTG
GTACTGAGCAAAACCCGCTAATCCTAATCCTTCTCTTGAGGAGTCTCAGCCTCTTAATACTTTCATGTTCAGATAAT
AGGTTCCGAAATAGCGAGGGGCATTAACTGTTTATACGGGCACTTTTACTCAAGGCACCTGACCCCGTTAAAACTTATTA
CCAGTACACTCCTGTATCATCAAAAGCCATGTATGACGCTTACTGGAACGGTAATTCAGAGACTGCGCTTCCATTCTG
GCTTTAATGAGGATCCATTCTGTTGTGAATATCAAGGCCAATCGTCTGACCTGCTCAACCTCTGTCAATGTGCGCGC
GGCTCTGGTGGTCTGTTGGCGGCTCTGAGGGTGGTGGCTCTGAGGGTGGCGGTTCTGAGGGTGGCGGCTCTGAGGG
AGGCGGTTCCGGTGGTGGCTCTGGTTCGGTGCCATGGTTAGATCTGACAAAACTCACACATGCCACCCGTCGCCAGCAC
CTGAACCTCCTGGGGACCGTCAGTCTTCCTCTTCCCCCAAAACCAAGGACACCCCTCATGATCTCCCGGACCCCTGAG
GTCACATGCGTGGTGGACGTGAGCCACGAACCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGAGGTGCA
TAATGCCAAGACAAAGCCGCGGAGGAGCAACAGCACGTACCGTGTGGTCAAGCTCCTCAGCTCCTGCACCCAGG
ACTGGCTGAATGGCAAGGAGTACAAGTGCAAGGTCTCCAACAAGCCCTCCAGCCCATCGAGAAACCATCTCCAAA
GCCAAAGGCAGCCCCGAGAACCCACAGGTGTACACCCCTGCCCCCATCCCGGAGGAGATGACCAAGAACCCAGTCAAGCCT
GACCTGCCCTGGTCAAAGGCTTCTATCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAAGCCGGAGAACACTACA
AGACCAAGCCTCCCGTGGTGGACTCCGACGGCTCCTTCTCTACAGCAAGCTCACCGTGGACAAAGAGCAGGTGGCAG
CAGGGAAACGCTCTTCTCATGTCTCCGTGATGCACGAGGCTCTGCAACAACCACTACACGCAAGAGCCTCTCCCTGTCTCC
GGTAAATGA

FIG. 3

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SEQ ID NO:4

ATGTACAGGATGCAACTCCTGTCTTGCAATTGCACCTTGTCACGAATTCGATGGCTGAAACTGTTGAAAG
TTGTTTAGCAAAACCCCATACAGAAATTCATTTACTAACGCTGGAAGACGACAAACTTTAGATCGTTACGCTAACT
ATGAGGGCTGTCTGTGGAATGCTACAGCGCTTGTAGTTGTACTGGTGACGAAACTCAGTGTACGGTACATGGGTTCCCT
ATTGGGCTTGCTATCCCTGAATAATGAGGGTGGTGGCTCTGAGGGTGGCGGTCTGAGGGTGGCGGTCTGAGGGTGGCGG
TACTAAACCTCCTGAGTACGGTGATACACCTATTCGGGGCTATACCTTATCAACCTCTCGACGGCACTTATCCGCCTG
GTACTGAGCAAAACCCCGCTAATCCTAATCCTTCTTGAGGAGTCTCAGCCTCTTAATACTTTTCATGTTCAGATAAT
AGGTTCCGAAATAGGCAGGGGCATTAACTGTTTATACGGGCCTTTTACTCAAGGCACCTGACCCCGTTAAACTTATTA
CCAGTACACTCCTGTATCATCAAAAGCCATGTATGACGCTTACTGGAACGGTAAATTCAGAGACTGCGCTTTCCATTCTG
GCTTTAATGAGGATCCATTCTGTTGTGAATATCAAGCCCAATCGTCTGACCTGCCCTCAACTCTGCCAATGCTGGCGGC
GagTCTGGTGGTCTCTGGTGGCGCTCTGAGGGTGGTGGCTCTGAGGGTGGCGGTCTGAGGGTGGCGGTCTGAGGG
AGGCGGTTCCGGTGGTGGCTCTGGTTCGGGTGCCAGATCTGACAAACTCAACATGCCCCACCGTGCCCCAGCACCTGAAC
TCCTGGGGGACCGTCAGTCTTCCTCTTCCCCCAAAACCAAGGACACCCCTCATGATCTCCCGGACCCCTGAGGTCACA
TGCGTGGTGGTGACGTGAGCCACGAAGACCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGAGGTGCATAATGC
CAAGACAAAGCCGGGAGGAGCAGTACAAACAGCACGTACCGTGTGGTCAGGTCCTCACCGTCTGCACCCAGGACTGGC
TGAATGGCAAGGAGTACAAGTCAAGTCTCCAACAAGCCCTCCAGCCCCCATCGAGAAACCATCTCCAAGCCAAA
GGCAGCCCCGAGAACCCACAGGTGTACACCCCTGCCCCCATCCCGGAGGAGATGACCAAGAACCCAGTCAAGCTGACCTG
CCTGGTCAAAGGCTTCTATCCAGCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAAGCCGGAGAACCACTACAAGACCA
CGCCTCCCGTGGACTCCGACGGCTCCTTCTTCCCTCTACAGCAAGCTCACCGTGGACAGAGCAGGTGGCAGCAGGGG
AACGTCTTCTCATGTCCGTGATGCACGAGGCTCTGACAAACCACTACACGCAGAGAGCTCTCCCTGTCTCCGGGTAA
ATGA

FIG. 4

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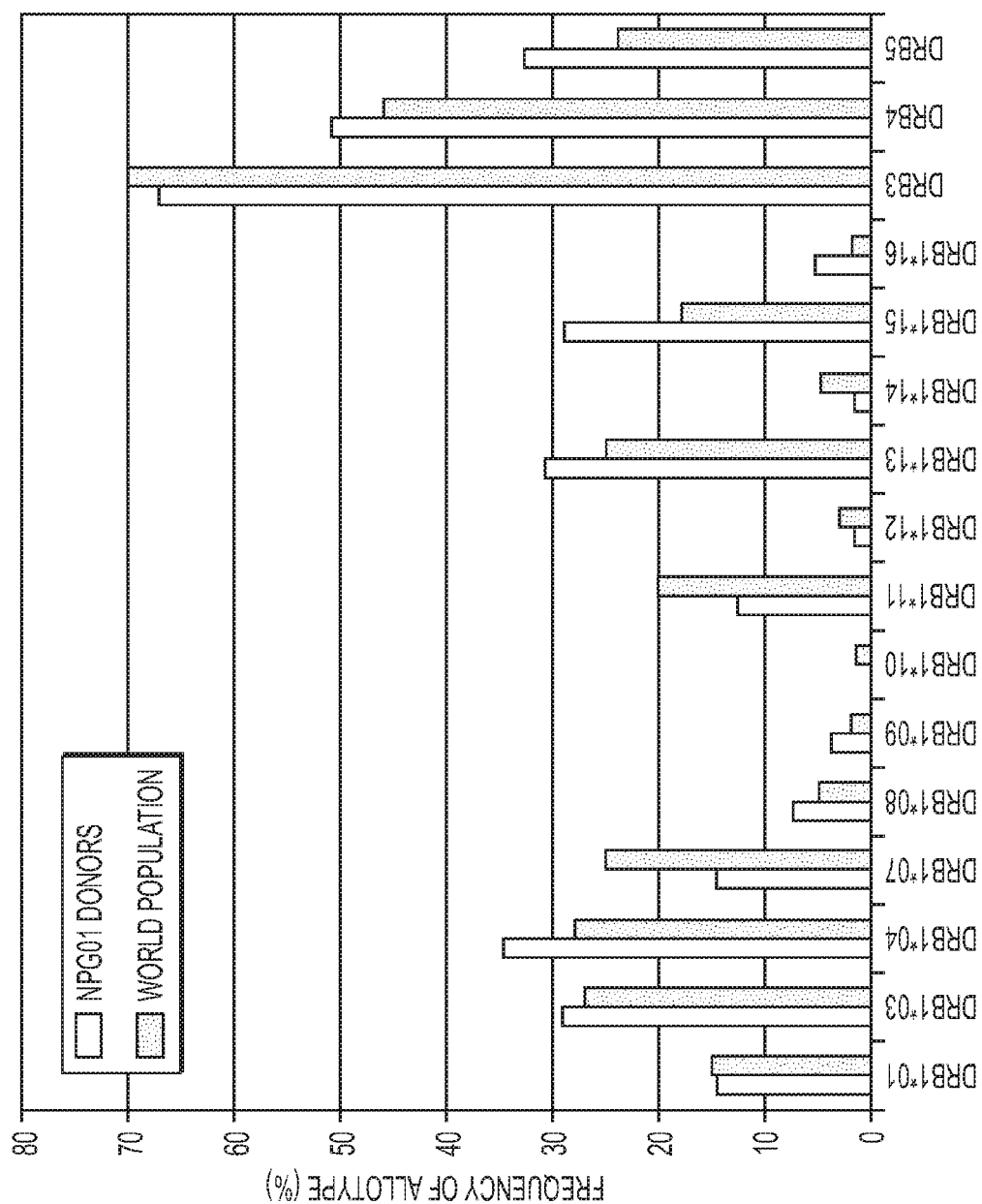


FIG. 5

FIG. 6 (SEQ ID NO:5)

[illegible]

SEQ ID NO: 6

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GGATCTGCGATCGCTCCGGTGGCCGTCAGTGGGCAGAGCGCACATCGCCACAGTCCCCGAGAAGTTGGGGGGAGGGGTTCGGCAATTGAACGGGT
 GCCTAGAGAAAGGTGGCGCGGGGTAAACTGGGAAAGTGATGTCGTGTACTGGCTCCGCTTTTCCCGAGGGTGGGGGAGAACCGTATATAAGTGC
 AGTAGTCGCGGTGAACGTTCTTTTCGCAACGGGTTTGGCGCCAGAACACAGCTGAAGCTTCGAGGGGCTCGCATCTCTCCTTCACGCGCCCGCC
 GCCCTACCTGAGGCGCCCATCCACGCGGTTGAGTCGCGTTCTGCGCGCTCCGCGCTGTGGTGCCCTCTGAAGTCCGTCGCGCGCTCTAGGTAAGT
 TTAAGCTCAGGTCGAGACCGGGCTTTGTCCGGCGCTCCCTGGAGCCTACCTAGACTCAGCGCGCTCTCCACGCTTTGGCTGACCTGCTTGC
 TCAACTCTACGTCCTTTGTTCTGTTCTGTTCTGCGCGCTTACAGATCCAAGCTGTGACCGCGCTACCTGAGATCACCGGGAAGGAGGGCCA
 CCATGTACAGGATGCAACTCCTGTCTTGCAATTGCACTAAGCTTGTCACTTGTACGAATTCGATGGCTGAAACTGTTGAAAGTTGTTTAGCAAAA
 CCCCATACAGAAAATTCATTTACTAACGTCGGAAGACGACAAAATTTAGATCGTTACGTAACCTATGAGGGCTGTCTGTGGAATGCTACAGG
 CGTTGTAGTTTGTACTGGTGACGAAACTCAGTGTACGGTCATTGGGTTCTTATTGGGCTTGCTATCCCTGAAAATGAGGGTGGTGGCTCTGAGG
 GTGGCGGTTCTGAGGGTGGCGGTTCTGAGGGTGGCGGTACTAAACCTCCTGAGTACGGTGATACACCTATTCCGGGTATACCTTATATCAACCCCT
 CTCGACGGCACTTATCCGCTGGTACTGAGCAAAACCCCGCTAATCCTAATCCTTCTCTTGAGGAGTCTCAGCCTCTTAATACTTTTCATGTTTCA
 GAATAATAGGTTCCGAAATAGGCGAGGGGCATTAACTGTTTATACGGGCACTTTTACTCAAGGCACTGACCCGTTAAAATTTATTACAGTACA
 CTCCTGTATCATCAAGAGCCATGTATGACGCTTACTGGAACGGTAAATTCACAGACTGCGCTTTCCATTCTGGCTTTAATGAGGATCCATTGCTT
 TGTGAATATCAAGGCCAATCGTCTGACCTGCTCAACCTCCTGCCAATGCTGGCGCGAGTCTGGTGGTGGTCTGGTGGCGCTCTGAGGGTGG
 TGGCTCTGAGGGTGGCGGTTCTGAGGGTGGCGGCTCTGAGGGAGGGGCTTCCGGTGGTCTGGTTCGGTTCGGTGGCAGATCTGACAAAATCACA
 CATGCCACCGTCCCGACACCTGAACTCCTGGGGGACCGTCAGTCTTCTCTTCCCCCAAAACCCAAGGACACCCCTCATGATCTCCCGGACC
 CCTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGTGCATAATGCCAA
 GACAAAGCCCGGGAGGAGCAGTACAACAGCAGTACCGTGTGGTCAGCGTCTCACCCTCTGCACCGAGTGGTGAATGGCAAGGAGTACA
 AGTGCAAGGTCTCCAACAAAGCCCTCCAGCCCCATCGAGAAAACCATCTCCAAGCCAAAGGGCAGCCCGAGAACCCACAGGTGTACACCGTG
 CCCCCATCCCGGGAGGAGATGACCAAGAACCAGGTGACCTGACCTGGCTGGTCAAAGGCTTCTATCCAGCGACATCGCGCTGGAGTGGGAGAG
 CAATGGGCAGCCGGAGAACAACATAAGACCACGCTCCCGTGTGGACTCCGACGGCTCCTTCTTCTCTACAGCAAGCTCACCCTGGACAAGA
 GCAGTGGCAGCAGGGGAACGTCTTCTCATGCTCCGTGATGCAGAGGCTCTGCACAACCACTACACGCAGAGAGCTCTCCCTGTCTCCGGGT
 AAATGAGTGTAGCTGGCCAGACATGATAACATACATTGATGAGTTTGGACAACCAACAACAGTGAATGCAGTGAATAAATGCTTTATTTGTGAA
 ATTTGTGATGCTATTTGCTTTATTTGTAACATTATAAGCTGCAATTAACAAGTTAACAACAACAAATGCAATTCATTTATGTTTCAGGTTACGGG
 GGAGGTGTGGGAGGTTTTTTAAAGCAAGTAAACCTCTACAAATGTGGTATGGAATTAATTTCAAAATACAGCATAGCAAACTTTAACCTCCAA
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 TCTTTTCATGGAGTTTAAGATATAGTGTATTTTCCCAAGGTTTGAAGTACTCTTCATTTCTTTATGTTTTAAATGCACTGAOCTCCACATTTCCC
 TTTTATAGTAAATATTCAGAAATAATTTAATAACATCATTGCAATGAAAATAAATGTTTTTATTAGGCAGAAATCCAGATGCTCAAGGCCCTTCA
 TAATATCCCCAGTTTAGTAGTTGGACTTAGGGAACAAAGGAACCTTTAATAGAAATTTGGACAGCAAGAAAGCGAGCTTCTAGCTTATCCTCAGT
 CCTGCTCCTCTGCCACAAAGTGACGCAAGTTGCGCGCGGGTTCGCGCAGGGCGAACTCCCGCCCCACGGCTGCTCGCCGATCTCGGTCATGGCC
 GGCCCGGAGGCGTCCCGGAAGTTCTGCGACACGACCTCCGACCACTCGGCGTACAGCTCGTCCAGGCGCGGACCCACACCCAGGCCAGGCTGTT
 GTCCGGCACCACTGCTCTGACCGCGCTGATGAACAGGTCACGTCGTCCCGACCAACCGGCGAAGTGTCTCCACGAAGTCCCGGAGAG
 ACCCGAGCCGCTCGTCCAGAACTCGACCGCTCCGGCGACGTCGCGCGCGGTGAGCACCGGAACGGCACTGGTCAACTTGGCCATGATGGCTCCT
 CctgtcaggagaggaagaagaaggttagtacaattgCTATAGTGAAGTGTATTATATACTATGCAGATATACTATGCCAATGATTAATTGTCAA
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 TCACAGGAGGGAGAGGAGGAGAGCTTGAGACAGACCCCGGGACCGCCAACTGCGAGGGGACGTGCTAGGGCGGCTTCTTTTATGGTGCGCCG
 GCCCTCGGAGGAGGGCGCTCGGGAGGCTTAGCGGCCAATCTGCGGTGGCAGGAGGCGGGCCGAGGCCCTGCTGACCAATCCGGAGCAGAT
 AGGAGTCTCAGCCCCCGCCCCAAAGCAAGGGGAAGTCACGCGCTGTAGCGCCAGCGTGTGTGAATGGGGCTTGGGGGGTGGGGCCCTG
 ACTAGTCAAAACAACTCCCATTTGACGTCAATGGGTGGAGACTTGAATCCCGTGAGTCAAAACCGCTATCCACGCCATTGATGTACTGCCA
 AAACCGCATCATCATGGTAATAGCGATGACTAATACGTAGATGTACTGCCAAGTAGGAAAGTCCCATAGGTCATGTACTGGGCATTAATGCCAGG
 CGGGCCATTACCGTCATTGACGTCAATAGGGGCGTACTTGGCATATGATACACTTGTATGTACTGCCAAGTGGGCAGTTTACCGTAAATACTCC
 ACCCATTTGACGTCAATGGAAGTCCCTATTGGCGTTACTATGGGAACATACGTCAATTATTGACGTCAATGGGCGGGGTGTTGGGCGGTACGCC
 AGGCGGGCCATTACCGTAAGTTATGTAACGCTGCAGGTTAATTAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGC
 CGCGTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTAT
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 GGAAGCGTGGCGTTTTCTCATAGCTCAGCTGTAGGTATCTCAGTTCCGTTAGGTGCTTCCGCTCCAAGCTGGGCTGTGTGCACGAACCCCGGT
 TCAGCCCGACCGCTGCGCTTATCCGGTAACCTATCGTCTTGTAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACA
 GGATTAGCAGAGCGAGGTATGTAGGCGGTGTACAGAGTCTTGAAGTGGTGGCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCTGC
 GCTCTGCTGAAGCCAGTTACCTTCGAAAAAGAGTTGGTAGCTCTTCATCCGGCAACAAACACCGCTGGTAGCGGTGGTTTTTTTCTTTGCAA
 GCAGCAGATTACCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTACGGGGTCTGACGCTCAGTGGAACGAAAACTCAGCTTAAG
 GGATTTTGGTCATGGCTAGTTAATTAACATTTAAATCAGCGCCGCAATAAAATATCTTTATTTTATTACATCTGTGTGTGGTTTTTTGTGTG
 AATCGTAACTAACATACGCTCTCCATCAAAACAAACGAAACAAACAACTAGCAAAATAGGCTGTCCCGAGTGAAGTGCAGGTGCCAGAAC
 TTCTCTATCGAA

FIG. 7

SEQ ID NO:7

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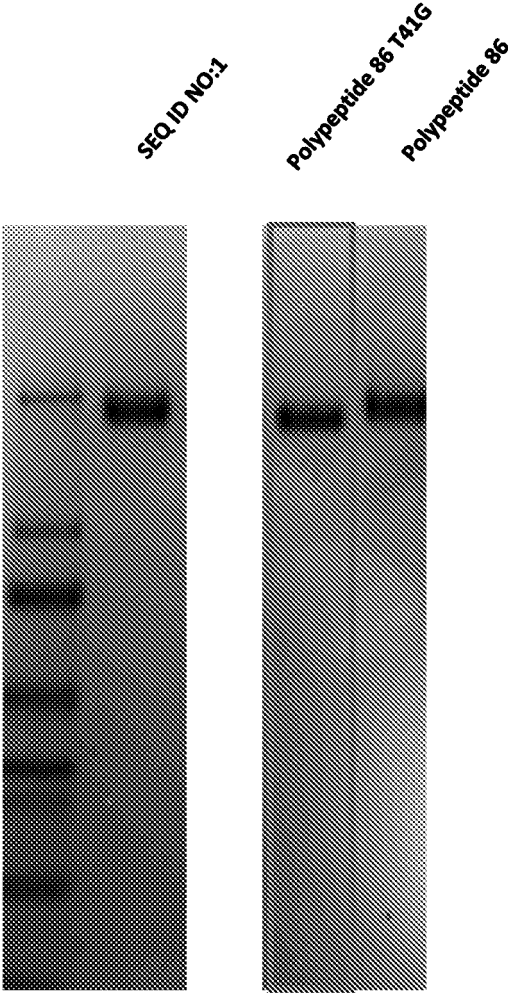
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AGTAGTCGCGGTGAACGTTCTTTTTCGCAACGGGTTTGGCGCCAGAACACAGCTGAAGCTTCGAGGGGCTCGCATCTCTCCTTCACGCGCCCGCC
GCCCTACCTGAGGCGCCCATCCACGCGGTTGAGTCGGCTTCTGCCGCTCCCGCTGTGGTGCTCTGAACTGCGTCCGCGCTCTAGTAAGT
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CCCCATACAGAAAATTCATTTACTAAGCTCTGGAAAGACGACAAAACCTTAGATCGTTACGCTAACTATGAGGGCTGTCTGTGGAATGCTGGAGG
CGTTGTAGTTTGTACTGTTGACGAAACTCAGTGTACGGTCAATGGGTTCTTATGGGCTTGCTATCCCTGAAAATGAGGGTGGTGGCTCTGAGG
GTGGCGGTTCTGAGGGTGGCGGTTCTGAGGGTGGCGGTACTAAACCTCCTGAGTACGGTGATACACCTATTCCGGGCTATACTTATATCAACCT
CTCGACGGCACTTATCCGCTGGTACTGAGCAAAACCCGCTAATCCTAATCCTTCTCTTGGAGTCTCAGCCTCTTAATACTTTCATGTTTCA
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CATGCCACCGTGGCCAGCACCTGAACCTCTGGGGGACCGCTCAGTCTTCTCTTCCCCCAAAACCCAAAGGACACCCCTCATGATCTCCCGGACC
CCTGAGGTACATGCGTGGTGGTGGACGTGAGCCACGAAGACCTGAGGTCAAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCATAATGCCAA
GACAAAGCCGCGGAGGAGCAGTACAACAGCACGTACCGTGTGGTCAAGCTCCTCACCCTCCTGCACCGAGGACTGGCTGAATGGCAAGGAGTACA
AGTGCAAGGTCTCCAACAAAGCCCTCCAGCCCCATCGAGAAAACATCTCCAAGCCAAAGGGCAGCCCCGAGAACCACAGGTGTACACCCCTG
CCCCATCCCGGGAGGAGATGACCAAGAACAGGTGAGCCTGACCTGCTGGTCAAAGGCTTCTATCCAGCGACATCGCGGTGAGTGGGAGAG
CAATGGGACGCGGAGAACAACTACAAGACCACGCTCCCGTGGTGGACTCCGACGGCTCCTTCTTCTCTACAGCAAGCTCACCGTGGACAAGA
GCAGGTGGCAGGAGGGAACGCTTCTCATGCTCCGTGATGCACGAGGCTCTGCACAACCACTACACGCAGAAGACCTCTCCCTGTCTCCGGGT
AAATGAGTCTAGCTGGCCAGACATGATAAGATACATTGATGAGTTTGGACAAACCAACTAGAATGCAGTGAACAAAATGCTTTATTTGTGAA
ATTTGTGATGCTATTGCTTTATTTGTAAACATATATAAGCTGCAATAAACAAGTTAAACAACAACATTCGATTCAATTTATGTTTCAGGTTTCAGGG
GGAGGTGTGGGAGGTTTTTAAAGCAAGTAAACCTCTACAAATGTGGTATGGAATTAATTTCTAAATACAGCATAGCAAACTTTAACTCCAA
ATCAAGCTCTACTTGAATCCTTTTCTGAGGGATGAATAAGGCATAGGCATCAGGGGCTGTGGCAATGTGCATTAGCTGTTTGCAGCCTCACT
TCTTTCATGGAGTTTAAGATATAGTGTATTTTCCCAAGGTTTGAAGTCTTCTTATTTCTTTATGTTTAAATGCAGTACCTCCACATTTCC
TTTTTAGTAAATATTTAGAAATAATTTAAATACATCATTGCAATGAAATAAATGTTTTTATTAGGCAGAAATCCAGATGCTCAAGGCCCTTCA
TAATATCCCCAGTTTAGTAGTTGGAAGTAAAGGAACCTTTAATAGAAATTTGACAGCAAGAAAGCGAGCTTCTAGCTTATCTCAGT
CCTGCTCCTCTGCCACAAAGTGACGCACTTGGCGGCGGGTGGCGCAGGGCGAACTCCCGCCCCACGGCTGCTCGCCGATCTCGGTGATGGCC
GGCCCGGAGGCGTCCCGGAAGTTCTGTGACACGACCTCCGACCACTCGGCGTACAGCTCGTCCAGGCGCGCACCCACACCCAGGCCAGGGTGT
GTCCGGCACCACTGGTCTGGACCGGCTGATGAACAGGTCACGTCTCCCGGACCAACCGCGGAAGTCTGCTCCACGAAGTCCCGGAGA
ACCCGAGCCGCTGGTCCAGAACTCGACCGCTCCGGGACGTGCGCGCGGTTGAGCACCGGAACGGCACTGGTCAACTTGGCCATGATGGCTCCT
CctgtcaggagaggaaagagaagaaggttagtacaattgCTATAGTGAGTTGTATTATACTATGCAGATATATATGCCAATGATTAATTGTCAA
ACTAGGGCTGCAgggttcagtagtgcacttttctgcactgccccatctctgcccaccccttccaggtatagacagtcagtgacttaeCAAAC
TCACAGGAGGAGAGAGGAGAGCTTGAGACAGACCCGCGGACCGCGAACTCGGAGGGGACGTGGCTAGGGCGGCTTCTTTTATGGTGGCGCG
GCCCTCGGAGGCGAGGCGCTCGGGGAGGCTAGCGGCAATCTGCGGTGGCAGGAGGCGGGCGAAGGCCGTGCTGAACCAATCCGAGCACAT
AGGAGTCTCAGCCCCCGCCCCAAAGCAAGGGGAAGTCAAGCGCTGTAGCGCAGCGTGTGTGAAATGGGGGCTTGGGGGGTGGGGCCCTG
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AAACCGCATCATCATGGTAAATAGCGATGACTAATACGTAGATGTACTGCCAAGTAGGAAAGTCCCATAGGTCATGTACTGGGCATAATGCCAGG
CGGGCCATTTACCGTCATTGACGTCAATAGGGGGGCTACTTGGCATATGATACACTTGATGTACTGCCAAGTGGGCAGTTTACCGTAAATACTCC
ACCCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGGAACATACGTCAATTATTGACGTCAATGGGCGGGGCTGTTGGCGGTGACGC
AGGCGGGCCATTTACCGTAAGTTATGTAACGCTGCAGGTAAATTAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGC
CGCGTGTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTAT
AAAGATACCAGGCGTTTCCCTGGAAGCTCCCTGCTGGCTCTCCTGTTCCGACCTGCCGCTTACCGGATACCTGTCCGCTTTCTCCCTTCG
GGAAGCGTGGCGCTTTCTCATAGCTCAGCTGTAGGTATCTCAGTTCCGGTGTAGGTGCTTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCGT
TCAGCCCGACCGCTGCGCTTATCCGGTAACTATCGTCTTGTAGTCCAACCGGTAAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAACA
GGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGACTTCTTGAAGTGGTGGCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCTGC
GCTCTGCTGAAGCCAGTTACCTTCGAAAAAGACTTGTAGCTCTTGATCCGGCAACAAACACCGCTGGTAGCGGTGGTTTTTTTTGTTTGCAG
GCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGCTGACGCTCAGTGAACGAAAACTCACGTTAAG
GGATTTTGGTCAATGGCTAGTTAATTAACATTTAAATCAGCGCGCGCAATAAAAATATCTTTATTTTTCATACATCTGTGTGTGGTTTTTGTGTG
AATCGTAACATAACGCTCTCCATCAAAACAAACGAAACAAACAACTAGCAAAATAGGCTGTCCCGAGTGAAGTGCAGGTGCCAGAAC
TTTCTCTATCGAA

FIG. 8

SUBSTITUTE SHEET (RULE 26)

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FIG. 9



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FIG. 10.

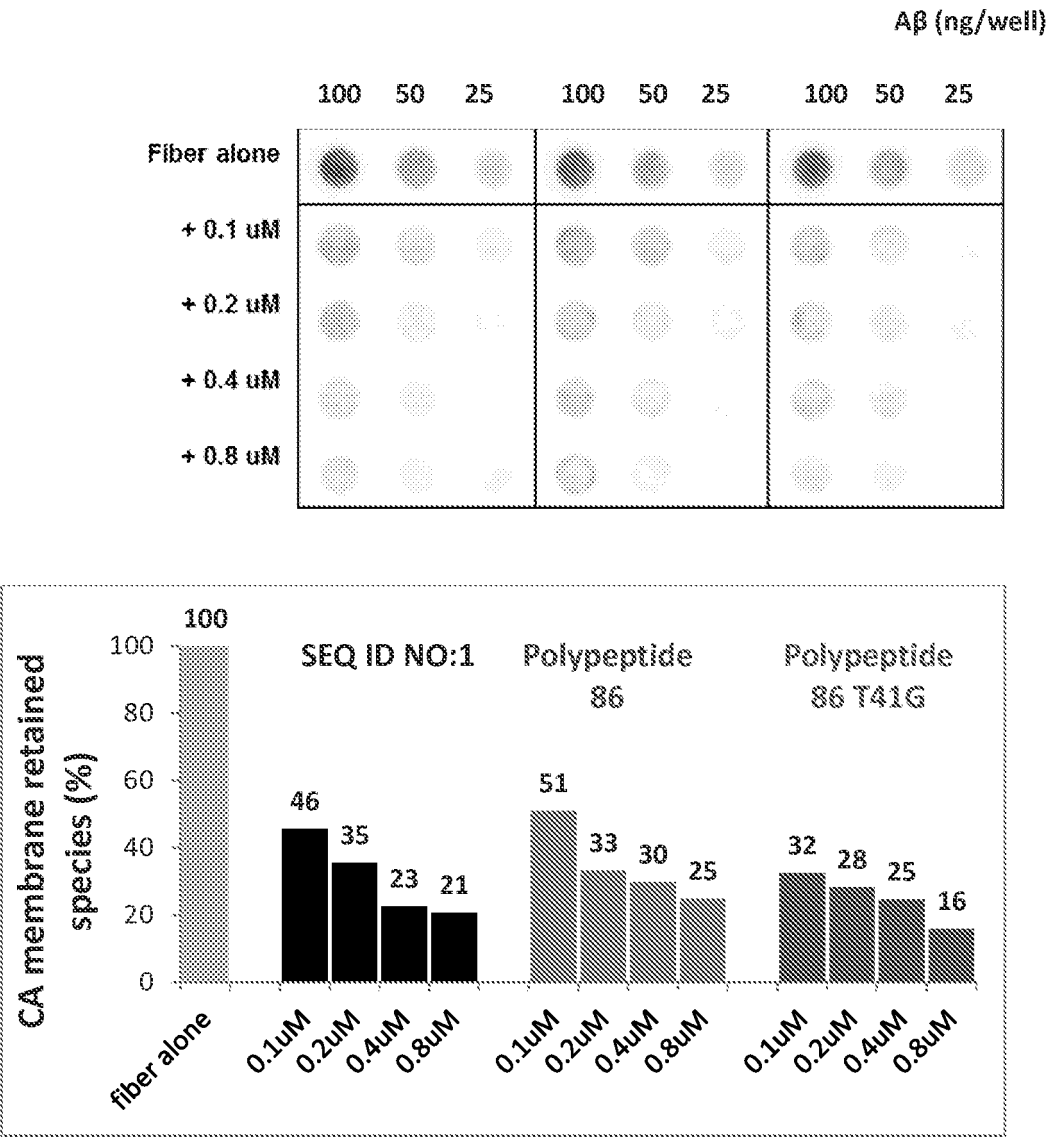


FIG. 11.

