

Séquence 1

atgtgatgttcactcaggggaccgctcctactggagcacgggTATCCCCAGCCATAAGGTCCGTTC 66
ACCCATTACATACAAAGTGCCTGCGGCTACAATAGGAGAGCATGAGTAAGGTGTTACGTGGACG 132
M S K V F T W T 8
GGAGCATACTGGAGAACCTTCCGCCCTAATTCGCGCAAACCTCGCATGTGACTGGTACGACAGTAGT 198
G A Y W R T F R P N S R K L A C D W Y D S S 30
AATAAGACCCACCAgttgtgatacgttactagcgaatagctctggggattaaggcggttgaggcaaaag 264
N K T H Q 35
atgagAGGCGGAACAGCTGGTGGCAAGTATCGGGCCTCCAATTACAGTAAGGAGATGATCGAGACA 330
G G T A G G K Y R A S N Y S K E M I E T 55
CGGAATATCAGCGCAAGTCTCATTCTCTTTACCGGGCGGCGCAAGGTATTAGATACCGACGTTAC 396
R N I S A S L I P L Y R A A Q G I R Y R R Y 77
AAACCCTGGGTGAGTACAGAAGTCCAGATAGATATAAGCAGCCATCGTTCTAATATGCAGGAATCT 462
K P W V S T E V Q I D I S S H R S N M Q E S 99
ACCCgtttcggttaaattttcaataactttacttttagcgtggccatacagTGGGCGTGGCTCCGACA 528
T L G V A P T 106
ATTATCCAAGAGTGTATATATAGACATTTCGAGCCCAGGTGTGTGCGTTACCTTATGTGGGTGTGAAT 594
I I Q E C I Y R H S S P G V S L P Y V G V N 128
ACTCCCCCGCATAACAACCTCCATGTTCGACTCCAGATTTGAGGTTCCCTGACGCTCGACTTAAATGGC 660
T P P A Y N S M S T P D L R F L T L D L N G 150
CCCCACCGGTCAGCTACCGGCTTGCTATACTCTTACCTAGTTTTCGATCGGCGATTTGTTCAACTGT 726
P D R S A T G L L Y S Y L V S I G D L F N C 172
GGACGAAGGAAGATGATGTTGTGTGGGATTTATAGAGTCCCCAAGTTAGCTGAAACGCGCTGGCGA 792
G R R K M M L C G I Y R V P K L A E T R W R 194
ACGGGACCGTCTGCTCAGTACgtttaggttctccccacaatatcaactgcggtattataagcggg 858
T G P S A Q Y 201
agctcctagtagCATGGTTCGGGATATTGCTACAAACCGACAACCTTACCCTTTTTTCGTGTGAGCTA 924
H G R G Y C Y K P T T Y P F S C E L 219
AGGCGGGGATATTGTTGGTACAGGTACTCGACGGATACCACGAAACTTCATGCGGTTCAACCGGGC 990
R R G Y C W Y R Y S T D T T K L H A V Q P G 241
TCCGCCAGGGATCTCCTgtacctttgagtcgaatatatgacagatctatcgtctatatttgaagtg 1056
S A R D L L 247
cttccactcgtcaccacaacatgcccgggttacaatggcagtcatacacccggacgcaaaagtaatt 1122

gtgggacccggtgtaaaagcttccagGCAAAGCCTAGCGTACCTCGCCAAGCATAGCACCTTGACC 1188
Q S L A Y L A K H S T L T 260
GAGCCATGCGTTAGGTACATGTGGAGACCGTGTATGAGTGATCGGCGGGTACCGGGTGAACGACTG 1254
E P C V R Y M W R P C M S D R R V P G E R L 282
CAAGAGTTCTAGACCTTCCGCCCTAATTCGCGCAAAC
Q E F *

Séquence 2

TATCCCCAGCCATAAGGTCCGTTACCCATTACATACAAAGTGCGCTGCGGCTACAATAGGAGAGC 66
ATGAGTAAGGTGTTACGTGGACGGGAGCATACTGGAGAACCTTCCGCCCTAATTCGCGCAAACCTC 132
M S K V F T W T G A Y W R T F R P N S R K L 22
GCATGTGACTGGTACGACAGTAGTAATAAGACCCACCAAGGCGGAACAGCTGGTGGCAAGTATCGG 198
A C D W Y D S S N K T H Q G G T A G G K Y R 44
GCCTCCAATTACAGTAAGGAGATGATCGAGACACGGAATATCAGCGCAAGTCTCATTCCCTCTTTAC 264
A S N Y S K E M I E T R N I S A S L I P L Y 66
CGGGCGGCGCAAGGTATTAGATACCGACGTTACAAACCCTGGGTGAGTACAGAAGTCCAGATAGAT 330
R A A Q G I R Y R R Y K P W V S T E V Q I D 88
ATAAGCAGCCATCGTTCCTAATATGCAGGAATCTACCCTGGGCGTGGCTCCGACAATTATCCAAGAG 396
I S S H R S N M Q E S T L G V A P T I I Q E 110
TGTATATATAGACATTCGAGCCCAGGTGTGTCGTTACCTTATGTGGGTGTGAATACTCCCCCGCA 462
C I Y R H S S P G V S L P Y V G V N T P P A 132
TACAACTCCATGTCGACTCCAGATTTGAGGTTCTGACGCTCGACTTAAATGGCCCCGACCGGTCA 528
Y N S M S T P D L R F L T L D L N G P D R S 154
GCTACCGGCTTGCTATACTCTTACCTAGTTTCGATCGGGCATTGTTCAACTGTGGACGAAGGAAG 594
A T G L L Y S Y L V S I G D L F N C G R R K 176
ATGATGTTGTGTGGGATTTATAGAGTCCCCAAGTTAGCTGAAACGCGCTGGCGAACGGGACCGTCT 660
M M L C G I Y R V P K L A E T R W R T G P S 198
GCTCAGTACCATGGTCGGGGATATTGCTACAAACCGACAACCTTACCCTTTTTCGTGTGAGCTAAGG 726
A Q Y H G R G Y C Y K P T T Y P F S C E L R 220
CGGGGATATTGTTGGTACAGGTA CTGACGGATACCACGAAACTTCATGCGGTTCAACCGGGCTCC 792
R G Y C W Y R Y S T D T T K L H A V Q P G S 242
GCCAGGGATCTCCTGCAAAGCCTAGCGTACCTCGCCAAGCATAGCACCTTGACCGAGCCATGCGTT 858
A R D L L Q S L A Y L A K H S T L T E P C V 264
AGGTACATGTGGAGACCGTGTATGAGTGATCGGCGGGTACCGGGTGAACGACTGCAAGAGTTCTAG 924
R Y M W R P C M S D R R V P G E R L Q E F *
ACCTTCCGCCCTAATTCGCGCAAAC

