



B.

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1 - agccgactcagtcgccgacactcaataggcatatttccgcagcgaggaatatttgctgttttcttaccgacaaaATGGACTTTGCAGCCTCCA - 90
1 - M D F A A S K - 7
91 - AGATGCTGCTGCAACGCCACTGCAAGCGGCCGCTGGTCCCGCGGAGGGAACCAATGAAGCAAAAGACAAGCGAACGAGCATACGGCGCC - 180
8 - M S C N A T A S G R G P A E G T N E A K T S G T E Q I T A D - 37
181 - ATGTGGCGCCCCGTGCGGCGGCGGCGAGCGGTCTTCTTTAGCCGCGCCCCGCGGAAAGTCGCTGGCCAACCATCATCGATAGCGAGGAGA - 270
38 - V A P V R R R Q R S S F S R P R R K S L A N H I D S E E S - 67
271 - GCCTGCTGCTCAAGTTTGACTTGTCTTGACCGAACTGGAGCGGCGGTCTGAAAGCCTCGAGAGTATGGCGAGGTCTCCATCGACTCGA - 360
68 - L L L K F D L F L T E L E R R L E S L E S Y G E V S I D S S - 97
361 - GCATTGCCCGGCGCATATGCCACACTGCAGGTGTGCGTACGCGGTGCTCGCAAGCGTCGCAAGAGGTGATTGGTGCAGCGGCGGCGCGT - 450
98 - I A R A Y A T L Q V V R T R C S Q A S E E V I G A G R R R L - 127
451 - TGCAAAATCATGGTCGAGACCATCGAGACTCGGTACAGGAGGCCCTTGTGGCGGCGAGTCGCTCAACGAAAAGCGCGGGTAAGCATTG - 540
128 - Q I M V E T I E T R Y Q E A L V A A E S L N E K A R V S I E - 157
541 - AGTACTCGACGACATGCTGCGGACTTTGAGAACCACGCCCAAGTTCGCGGAGCGGAGCATCTCACAGCCACAGACGACGCGGGT - 630
158 - L L D D M L S D F E N H A H K F R E R S I S H A T D A A G L - 187
631 - TGTTTATGGGCGAGAGCGCGCGGTGATGGACGAAGGACATAGGCGGTGCACGTGAGGTGGTGAGGAGTCTTGAAGCGTGCTAAGAAGG - 720
188 - F M G E S R R V M D E G I G R A E V V D E S F E R A K K A - 217
721 - CGGCCGAGACACTCGAGGAGCACATCCAGCGTGCAGATTGCCAACTCACGGA AAAACGGCCTGCTGAGCTACGAGGAGCTGCCGGTGCCGT - 810
218 - A E T L L E E H I Q R A I A N S R K N G L S Y E E L P V P W - 247
811 - GGCGCATCAACCCGACATATCTCGCGGCTACCGCTTACGGAGACCAAGGCGGCGATCTCCGCTGTCATGTTCCGCTTCCCAACGAAC - 900
248 - R I N P H I I R G Y R F S E T K A A C T C F R S M F R L S N E L - 277
901 - TGGTCAACATCTGGTGCACCGGCTTGGGCATCGTGGTGGTGTCTGATTTGCCCTTGATCTTTACCCGACAGCGCTCAACTTCCACTCA - 990
278 - V N I W S H G L G I V V V L S I A L Y F Y P T S V N F H L S - 307
991 - GCACCAAAAACGACGCTTTCATTGCTGCCGTCTTCTTCTTCGCGGCGTGCCAATGCCTGGTCTGCAGCACCATCTGGCACACCATGAAC - 1080
308 - T K T D C V F I A A V F F F F A A C Q C L V C S T I W H T M N S - 337
1081 - CGATTGCGGACCGGACCTGATCTCATCGTGGCGTGTGTCGATTACCGGCATCTCGTCTGGTGGCGGACATCCATCATGACGACGCC - 1170
338 - I A D A D L I S S L A C V D Y T G I S L V A A S I M T T E - 367
1171 - AATACAGGCGCTTTTATCGCAACCGTTTAGCCGGTGGGTGTACATTTCCATGACAGCGGTGCTGGGCAATGGCGGCGTGATGCTGCCGT - 1260
368 - Y T A F Y C C E P F S R W V Y I S M T A V L G I G V M L P W - 397
1261 - GGCACCCGTTCTTCAACCGCGGCAGATGGCCTGGGCGCGCGTCCGCTTTATGTTGGTCTGGCCATGACGGGCTTTGTGCGCATGGTGC - 1350
398 - H P F F N R A D M A W A R A V A F Y V G L A M T G F V P M V Q - 427
1351 - AGATCATGACGACGCGCGGCGATGTCGTTTGTGTGGAGTTTACCTGACCATTTACCAAGTCGCTGTGGTGTACTGACCTGACGCGCCATGG - 1440
428 - I M T T R G M S F V W E F Y L P I T K S L V L V Y L T G A M V - 457
1441 - TGTATGCGAGCAAGGTGCGGCGAGCGGTGGCGGCGGCGCATGTTTGACTACATTTGCGGCGGAGCCAAACCTGTGGCACTTGGCCGTGCTGG - 1530
458 - Y A S K V F E R W R P G M F D Y I G G S H N L W H L A V L G - 487
1531 - GCGGCATCCTGTTCACATCTCGGCGGATGACAGGATCTTCTTCAACGCCCTTCCGCGAGGCGCAAGACGGCTGCCCTGCTACTaagctg - 1620
488 - G I L F H Y S A M Q E F F S N A F R Q A E D G C P A Y * - 514
1621 - gctgttggttaatggtattcctgaacgtccgagtcctttcttggcgctttttcttcttcttcttcttcttcttcttcttcttcttcttctt - 1710
1711 - cttgttcaactcagtcgcatgataaacaagaatttaccaccaagagacacaaaagacagtcgttgcatctttaaactattttttttttttctt - 1800
1801 - tttctttcttcttcaatttttggtttgtacacgattttacggctttgtttcttctgtctgttcttcttctgtcacacatttttggcggttttatgtatt - 1890
1891 - ttgcatattttgggaaccactgcacgttcttcttgacatatagggaacatcacatacatatacacataaaaaaaaaaaaaaaaaaaaaaaaaa - 1980
1981 - a - 1981

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