



V1662 Sequence

1 TAGTTATTAA TAGTAATCAA TTACGGGGTC ATTAGTTCAT AGCCCATATA
 51 TGGAGTTC CGTTACATAA CTTACGGTAA ATGGCCCGCC TGGCTGACCG
 101 CCCAACGACC CCCGCCATT GACGTCAATA ATGACGTATG TTCCCATAGT
 151 AACGCCAATA GGGACTTTCC ATTGACGTCA ATGGGTGGAG TATTTACGGT
 201 AAAC TGCCCA CTTGGCAGTA CATCAAGTGT ATCATATGCC AAGTACGCC
 251 CCTATTGACG TCAATGACGG TAAATGGCCC GCCTGGCATT ATGCCCAGTA
 301 CATGACCTTA TGGGACTTTC CTACTTGGCA GTACATCTAC GTATTAGTCA
 351 TCGTATTAC CATGGTGATG CGGTTTTGGC AGTACATCAA TGGGCGTGGA
 401 TAGCGGTTTG ACTCACGGGG ATTTCCAAGT CTCCACCCCA TTGACGTCAA
 451 TGGGAGTTTG TTTTGGCACC AAAATCAACG GGACTTTCCA AAATGTCGTA
 501 ACAACTCCGC CCCATTGACG CAAATGGGCG GTAGGCGTGT ACGGTGGGAG
 551 GTCTATATAA GCAGAGCTGG TTTAGTGAAC CGTCAGATCC GCTAGCCACC
 601 ATGGCCTCCT CCGAGGACGT CATCAAGGAG TTCATGCGCT TCAAGGTGAA
 651 GATGGAGGGC TCCGTGAACG GCCACGAGTT CGAGATCGAG GGCGAGGGCG
 701 AGGGCCGCC CTACGAGGGC ACCCAGACCG CCAAGCTGAA GGTGACCAAG
 751 GGCGGCCCCC TGCCCTTCGC CTGGGACATC CTGTCCCCTC AGTTCCAGTA
 801 CGGATCCAAG GCCTACGTGA AGCACCCCGC CGACATCCCC GACTACTTGA
 851 AGCTGTCCTT CCCCAGGGC TTCAAGTGGG AGCGCATCAT GAACTTCGAG
 901 GACGGCGGCG TGGTGACCGT GACCCAGGAC TCCACCCTGC AGGACGGCGA
 951 GTTCATCTAC AAGGTGAAGC TGCGCGGCAC CAACTTCCCC TCCGACGGCC
 1001 CCGTGATGCA GAAGAAGACC ATGGGCTGGG AGGCCTCCAC CGAGCGGATG
 1051 TACCCCGAGG ACGGCGCCCT GAAGGGCGAG ATCAAGATGA GGCTGAAGCT
 1101 GAAGGACGGC GGCCACTACG ACGCCGAGGT CAAGACCACC TACAAGGCCA
 1151 AGAAGCAGGT CCAGCTGCCT GGCGCCTACA AGACCGACAT CAAGCTGGAC
 1201 ATCATCTCCC ACAACGAGGA CTACACCATC GTGGAACAGT ACGAGCGCGC
 1251 CGAGGCGCGC CACTCCACCG GCGCCCAAGC TTCGATAACT TCGTATAGCA
 1301 TACATTATAC GAAGTTATAG ATCCAATATT ATTGAAGCAT TTATCAGGGT
 1351 TATTGTCTCA TGAGCGGATA CATATTTGAA TGTATTTAGA AAAATAAACA
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 1451 CACCGGATCT AGATAACTGA TCATAATCAG CCATACCACA TTTGTAGAGG
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 1801 AAATCAAAAG AATAGACCGA GATAGGGTTG AGTGTGTTC CAGTTTGAA
 1851 CAAGAGTCCA CTATTAAGA ACGTGGACTC CAACGTCAA GGGCGAAAAA
 1901 CCGTCTATCA GGGCGATGGC CCACTACGTG AACCATCACC CTAATCAAGT
 1951 TTTTTGGGGT CGAGGTGCCG TAAAGCACTA AATCGGAACC CTAAAGGGAG
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 2151 GGGCGCGTCA GGTGGCACTT TTCGGGAAA TGTGCGCGGA ACCCTATTT
 2201 GTTTATTTTT CTAATACAT TCAAATATGT ATCCGCTCAT GAGACAATAA
 2251 CCCTGATAAA TGCTTCAATA ATATTGAAA AGGAAGAGTC CTGAGGCGGA
 2301 AAGAACCAGC TGTGGAATGT GTGTGAGTTA GGGTGTGGAA AGTCCCAGG
 2351 CTCCCAGCA GGCAGAAGTA TGCAAAGCAT GCATCTCAAT TAGTCAGCAA
 2401 CCAGGTGTGG AAAGTCCCCA GGCTCCCCAG CAGGCAGAAG TATGCAAAGC
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 2551 TAATTTTTTT TATTTATGCA GAGGCCGAGG CCGCCTCGGC CTCTGAGCTA
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4201 GCTTCAGCAG AGCGCAGATA CCAAATACTG TCCTTCTAGT GTAGCCGTAG
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4301 GCTAATCCTG TTACCAGTGG CTGCTGCCAG TGGCGATAAG TCGTGTCTTA
4351 CCGGGTTGGA CTCAAGACGA TAGTTACCGG ATAAGGCGCA GCGGTCGGGC
4401 TGAACGGGGG GTTCGTGCAC ACAGCCCAGC TTGGAGCGAA CGACCTACAC
4451 CGAACTGAGA TACCTACAGC GTGAGCTATG AGAAAGCGCC ACGCTTCCCG
4501 AAGGGAGAAA GCGGACAGG TATCCGGTAA GCGGCAGGGT CGGAACAGGA
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4701 TTCCTGGCCT TTTGCTGGCC TTTTGCTCAC ATGTTCTTTC CTGCGTTATC
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