



V308 Sequence

1 ATAAC TTCGT ATAGCATACA TTATACGAAG TTATGGCGCG CCGGCGCCCA
 51 ATACGCAAAC CGCCTCTCCC CGCGCGTTGG CCGATT CATT AATGCAGCTG
 101 GCACGACAGG TTTCCCGACT GGAAAGCGGG CAGTGAGCGC AACGCAATTA
 151 ATGTGAGTTA GCTCACTCAT TAGGCACCCC AGGCTTTACA CTTTATGCTT
 201 CCGGCTCGTA TGTTGTGTGA AATTGTGAGC GGATAACAAT TTCACACAGG
 251 AAACAGCTAT GACCATGATT ACGGATTCAC TGGCCGTCGT TTTACAACGT
 301 CGTGACTGGG AAAACCCTGG CGTTACCCAA CTTAATCGCC TTGCAGCACA
 351 TCCCCTTTT GCCAGCTGGC GTAATAGCGA AGAGGCCCGC ACCGATCGCC
 401 CTTCCCAACA GTTGCGCAGC CTGAATGGCG AATGGCGTTA ATTAACCAGG
 451 TAAGTTCCTA GGAGATCCTG GTCATGACTA GTGCTTGGAT TCTCACCAAT
 501 AAAAAACGCC CGGCGGCAAC CGAGCGTTCT GAACAAATCC AGATGGAGTT
 551 CTGAGGTCAT TACTGGATCT ATCAACAGGA GTCCAAGCGA GCTCGATATC
 601 AAATTACGCC CCGCCCTGCC ACTCATCGCA GTACTGTTGT AATTCATTAA
 651 GCATTCTGCC GACATGGAAG CCATCACAAA CGGCATGATG AACCTGAATC
 701 GCCAGCGGCA TCAGCACCTT GTCGCCTTGC GTATAATATT TGCCCATGGT
 751 GAAAACGGGG GCGAAGAAGT TGTCCATATT GGCCACGTTT AAATCAAAAC
 801 TGGTGAAACT CACCCAGGGA TTGGCTGAGA CGAAAAACAT ATTCTCAATA
 851 AACCCTTTAG GGAAATAGGC CAGGTTTTCA CCGTAACACG CCACATCTTG
 901 CGAATATATG TGTAGAAACT GCCGAAATC GTCGTGGTAT TCACTCCAGA
 951 GCGATGAAAA CGTTTCAGTT TGCTCATGGA AAACGGTGTA ACAAGGGTGA
 1001 ACACTATCCC ATATCACCAG CTCACCGTCT TTCATTGCCA TACGAAATTC
 1051 CCGATGAGCA TTCATCAGGC GGGCAAGAAT GTGAATAAAG GCCGGATAAA
 1101 ACTTGTGCTT ATTTTTCTTT ACGGTCTTTA AAAAGGCCGT AATATCCAGC
 1151 TGAACGGTCT GGTTATAGGT ACATTGAGCA ACTGACTGAA ATGCCTCAA
 1201 ATGTTCTTTA CGATGCCATT GGGATATATC AACGGTGGTA TATCCAGTGA
 1251 TTTTTTCTC CTTTTAGCT TCCTTAGCTC CTGAAAGATC CATAACTTCG
 1301 TATAGCATA ATTATACGAA GTTATGCGGC CGCGACGTCC ACATATACCT
 1351 GCCGTTCACT ATTATTTAGT GAAATGAGAT ATTATGATAT TTTCTGAATT
 1401 GTGATTAATA AGGCAACTTT ATGCCCATGC AACAGAAACT ATAAAAAATA
 1451 CAGAGAATGA AAAGAAACAG ATAGATTTTT TAGTTCTTTA GGCCCGTAGT
 1501 CTGCAAATCC TTTTATGATT TTCTATCAA CAAAAGAGGA AAATAGACCA
 1551 GTTGCAATCC AAACGAGAGT CTAATAGAAT GAGGTGCGAA AGTAAATCGC
 1601 GCGGGTTTGT TACTGATAAA GCAGGCAAGA CCTAAAATGT GTAAAGGGCA
 1651 AAGTGTATAC TTTGGCGTCA CCCCTTACAT ATTTTAGGTC TTTTTTTATT
 1701 GTGCGTAACT AACTTGCCAT CTTCAAACAG GAGGGCTGGA AGAAGCAGAC
 1751 CGCTAACACA GTACATAAAA AAGGAGACAT GAACGATGAA CATCAAAAAG
 1801 TTTGCAAAAC AAGCAACAGT ATTAACCTTT ACTACCGCAC TGCTGGCAGG
 1851 AGGCGCAACT CAAGCGTTTG CGAAAGAAAC GAACCAAAAG CCATATAAGG
 1901 AAACATACGG CATTTCCTAT ATTACACGCC ATGATATGCT GCAAATCCCT
 1951 GAACAGCAAA AAAATGAAAA ATATCAAGTT CCTGAGTTTCG ATTCGTCCAC
 2001 AATTAATAAT ATCTCTTCTG CAAAAGGCCCT GGACGTTTGG GACAGCTGGC
 2051 CATTACAAAA CGCTGACGGC ACTGTCGCAA ACTATCACGG CTACCACATC
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 2151 CATGTTCTAT CAAAAGTTCG GCGAACTTC TATTGACAGC TGGAAAAACG
 2201 CTGGCCGCGT CTTTAAAGAC AGCGACAAAT TCGATGCAA TGATTCTATC
 2251 CTAAAAGACC AAACACAAGA ATGGTCAGGT TCAGCCACAT TTACATCTGA
 2301 CGGAAAAATC CGTTTATTCT ACACTGATTT CTCCGGTAAA CATTACGGCA
 2351 AACAAACACT GACAACTGCA CAAGTTAACG TATCAGCATC AGACAGCTCT
 2401 TTGAACATCA ACGGTGTAGA GGATTATAAA TCAATCTTTG ACGGTGACGG
 2451 AAAAAAGTAT CAAAATGTAC AGCAGTTCAT CGATGAAGGC AACTACAGCT
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 2551 CACAAATACT TAGTATTTGA AGCAAACACT GGAAGTGAAG ATGGCTACCA
 2601 AGGCGAAGAA TCTTTATTTA ACAAAGCATA CTATGGCAA AGCACATCAT
 2651 TCTTCCGTCA AGAAAGTCAA AAACCTTCTG AAAGCGATAA AAAACGCACG
 2701 GCTGAGTTAG CAAACGGCGC TCTCGGTATG ATTGAGCTAA ACGATGATTA

2751 CACACTGAAA AAAGTGATGA AACCGCTGAT TGCATCTAAC ACAGTAACAG
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3051 CAATGTCGTG ATTACAAGCT ATATGACAAA CAGAGGATTC TACGCAGACA
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3151 ACATCTGTTG TCAAAGACAG CATCCTTGAA CAAGGACAAT TAACAGTTAA
3201 CAAATAAAAA CGCAAAAGAA AATGCCGATA TCCTATTGGC ATTGACGTA
3251 GGTGGCATT TCGGGGAAA TGTGCGCGGA ACCCTATTTT GTTTATTTTT
3301 CTAATAACAT TCAAATATGT ATCCGCTCAT GAGACAATAA CCCTGATAAA
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3501 GAGTGGGTTA CATCGAAGT GATCTCAACA GCGGTAAGAT CCTTGAGAGT
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4351 TTAACGTGAG TTTTCGTTCC ACTGAGCGTC AGACCCCGTA GAAAAGATCA
4401 AAGGATCTTC TTGAGATCCT TTTTTTCTGC GCGTAATCTG CTGCTTGCAA
4451 ACAAAAAAAC CACCGCTACC AGCGGTGGTT TGTGTTGCCG ATCAAGAGCT
4501 ACCAACTCTT TTTCCGAAGG TAACTGGCTT CAGCAGAGCG CAGATACCAA
4551 ATACTGTTCT TCTAGTGTAG CCGTAGTTAG GCCACCACTT CAAGAACTCT
4601 GTAGCACCGC CTACATACCT CGCTCTGCTA ATCCTGTTAC CAGTGGCTGC
4651 TGCCAGTGGC GATAAGTCGT GTCTTACCGG GTTGGACTCA AGACGATAGT
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4751 CCCAGCTTGG AGCGAACGAC CTACACCGAA CTGAGATACC TACAGCGTGA
4801 GCTATGAGAA AGCGCCACGC TTCCC GAAGG GAGAAAGGCG GACAGGTATC
4851 CGGTAAGCGG CAGGGTCGGA ACAGGAGAGC GCACGAGGGA GCTTCCAGGG
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5001 ACGCCAGCAA CGCGGCCTTT TTACGGTTCC TGGCCTTTTG CTGGCCTTTT
5051 GCTCACATGT TCTTTCCTGC GTTATCCCCT GATTCTGTGG ATAACCGTAT
5101 TACCGCCTTA CGCGTGATAA ACGACGGCCA GTAGATCT