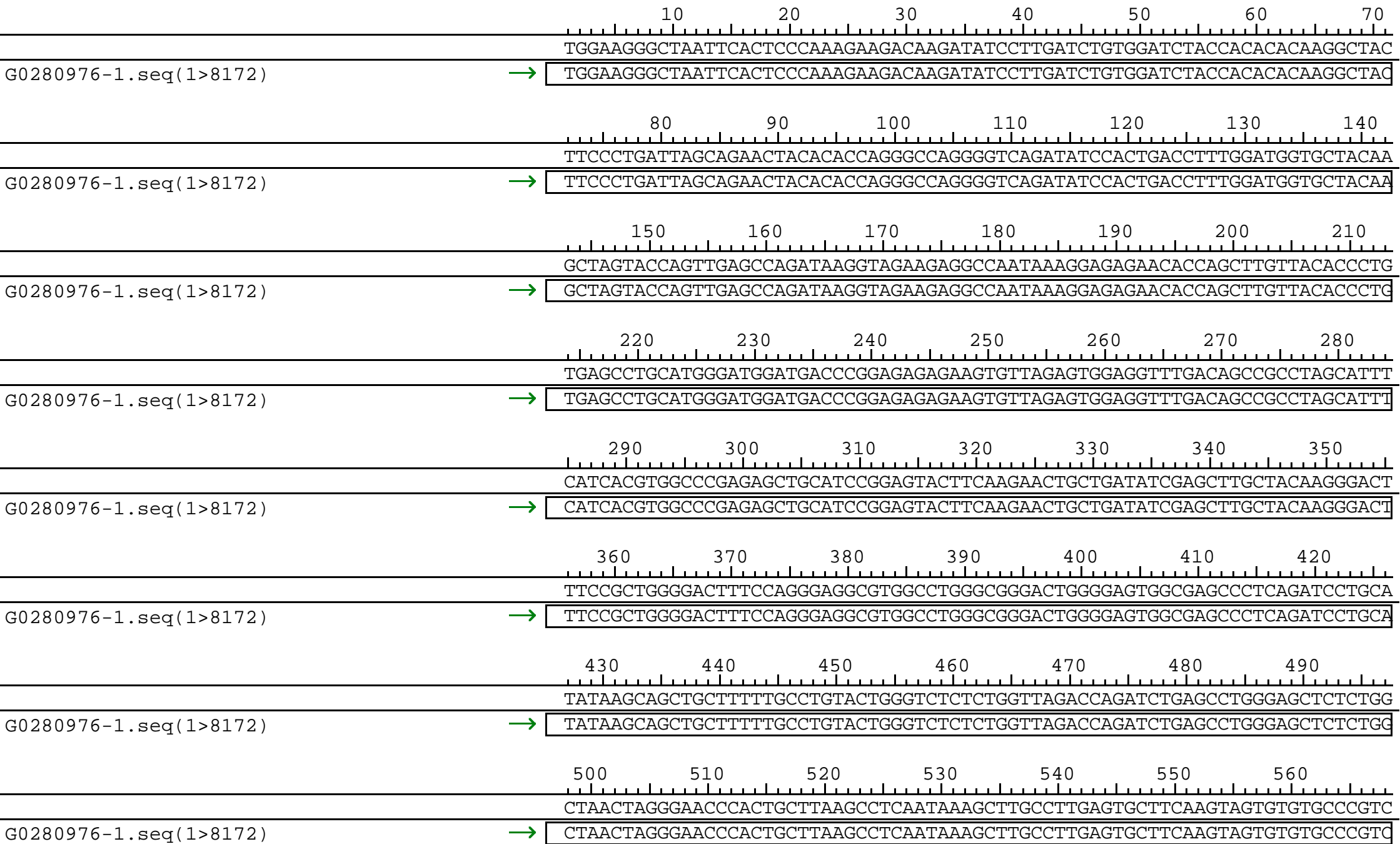
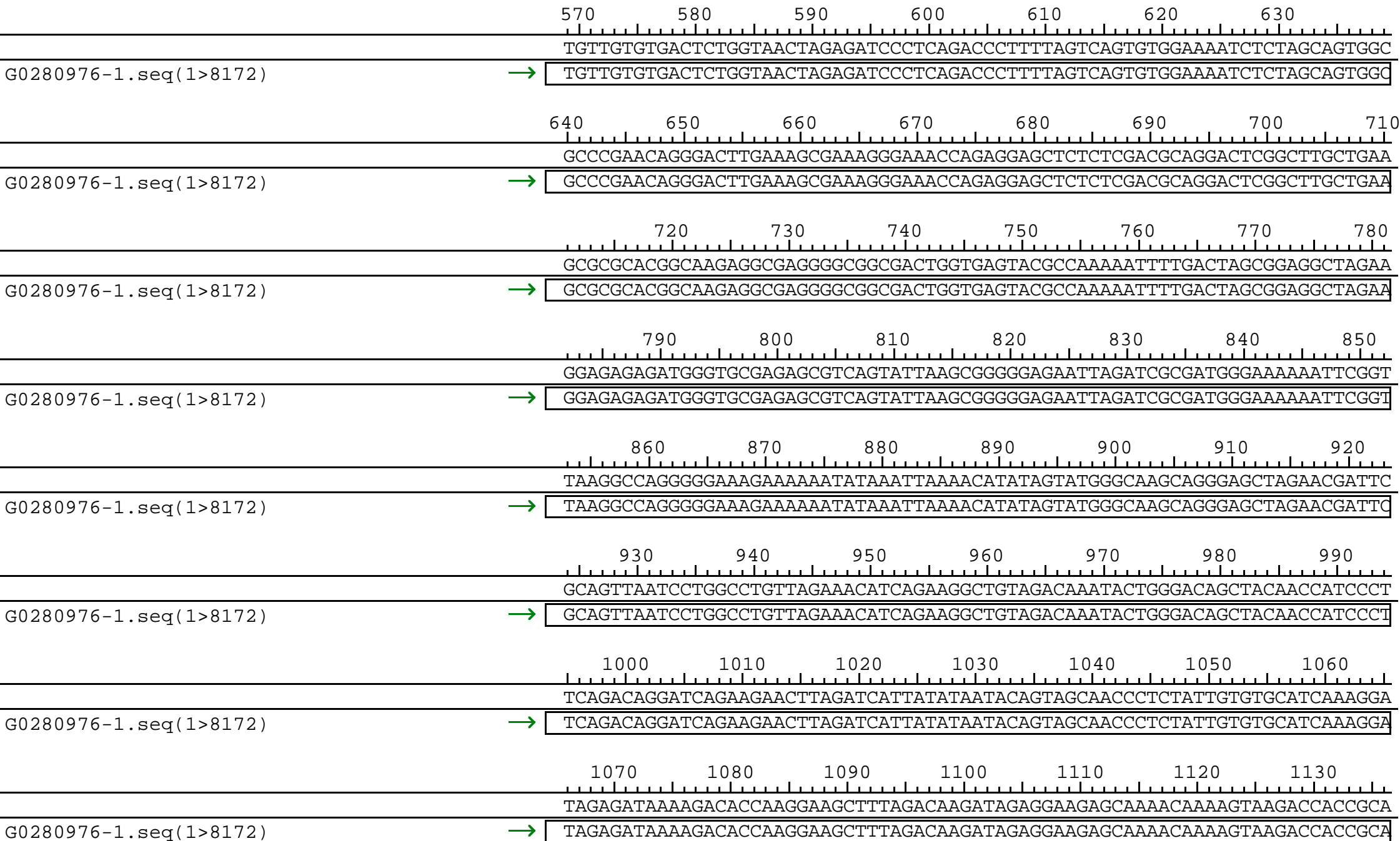


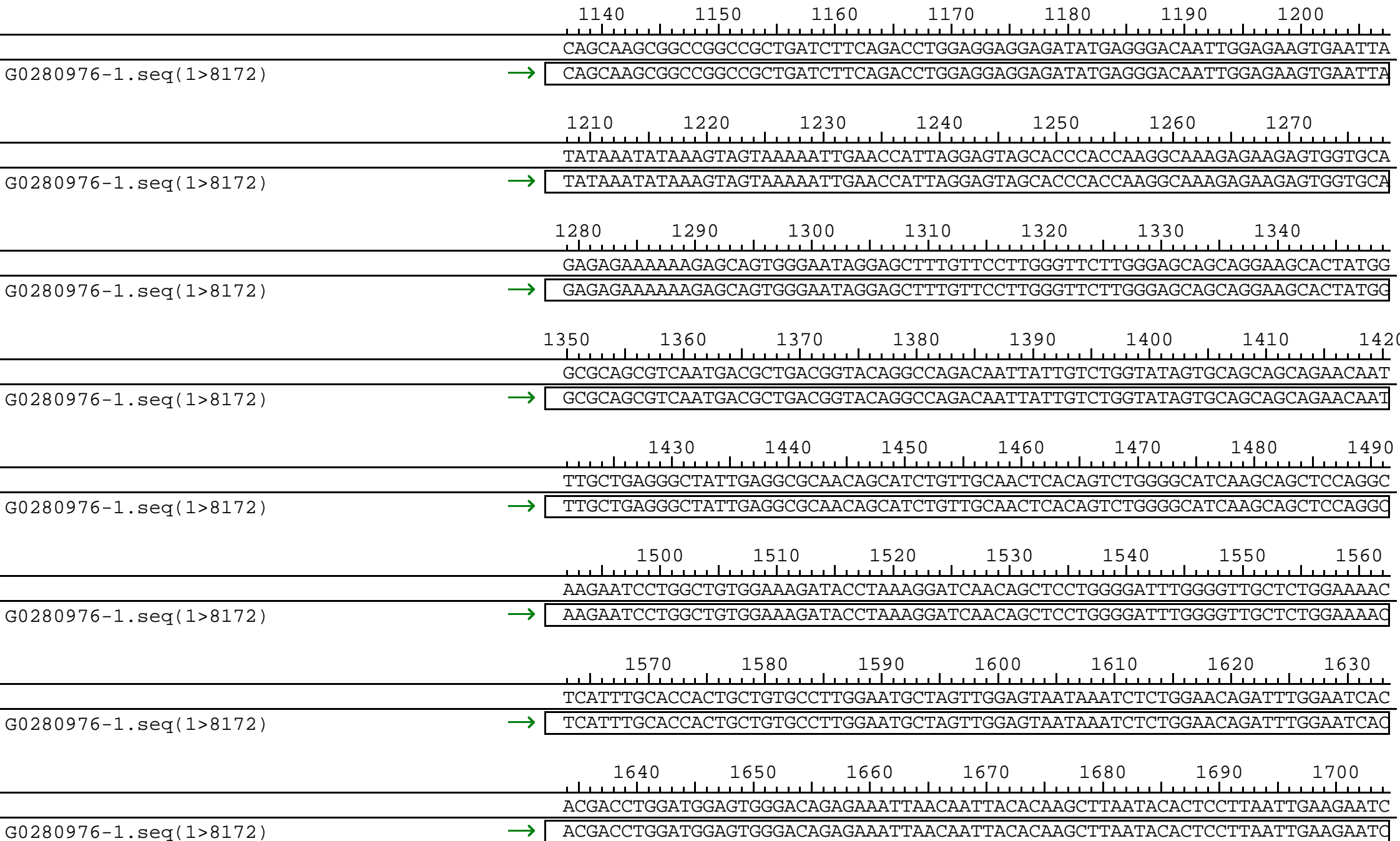
Project: G0280976-1.SQD Contig 1



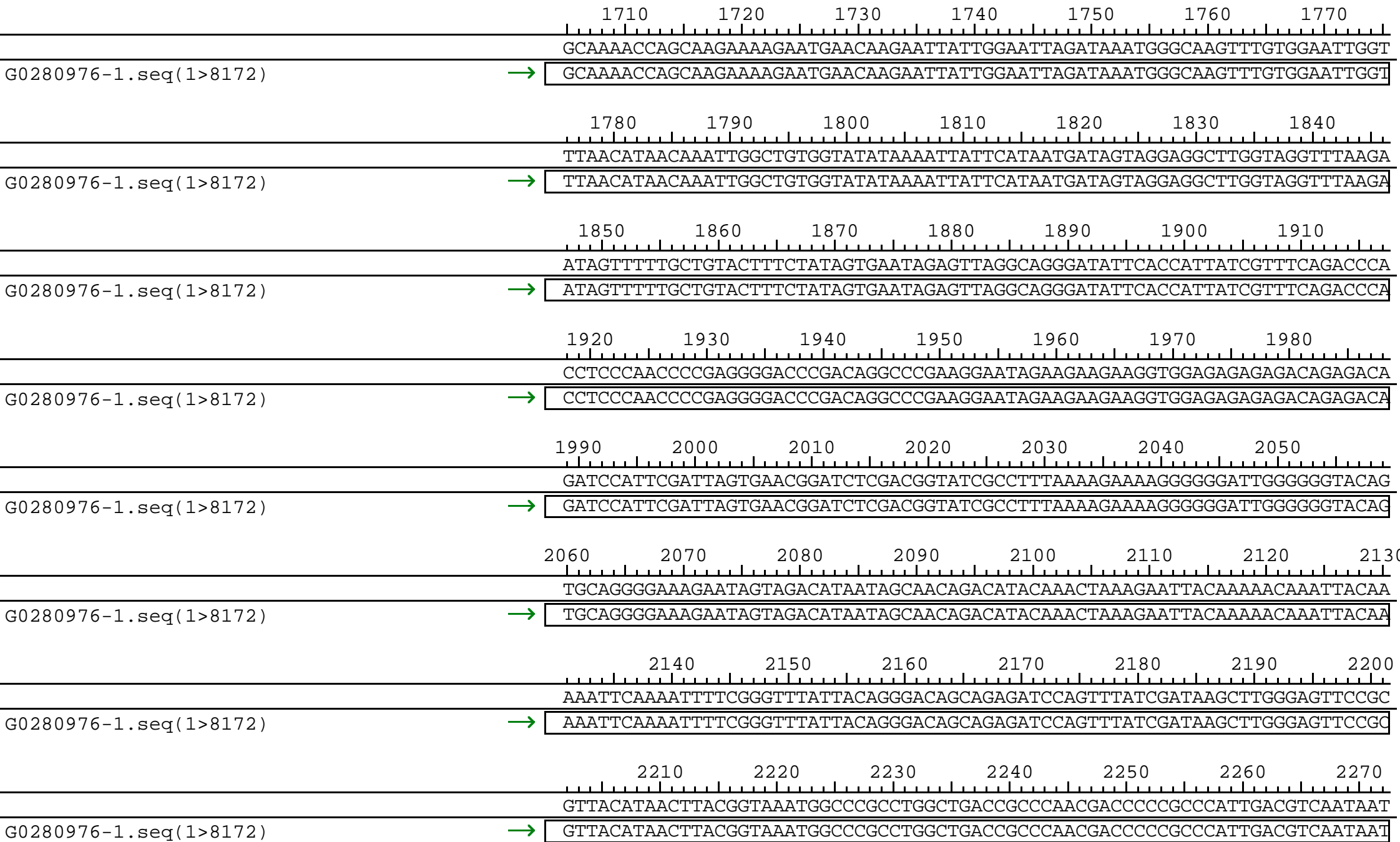
Project: G0280976-1.SQD Contig 1

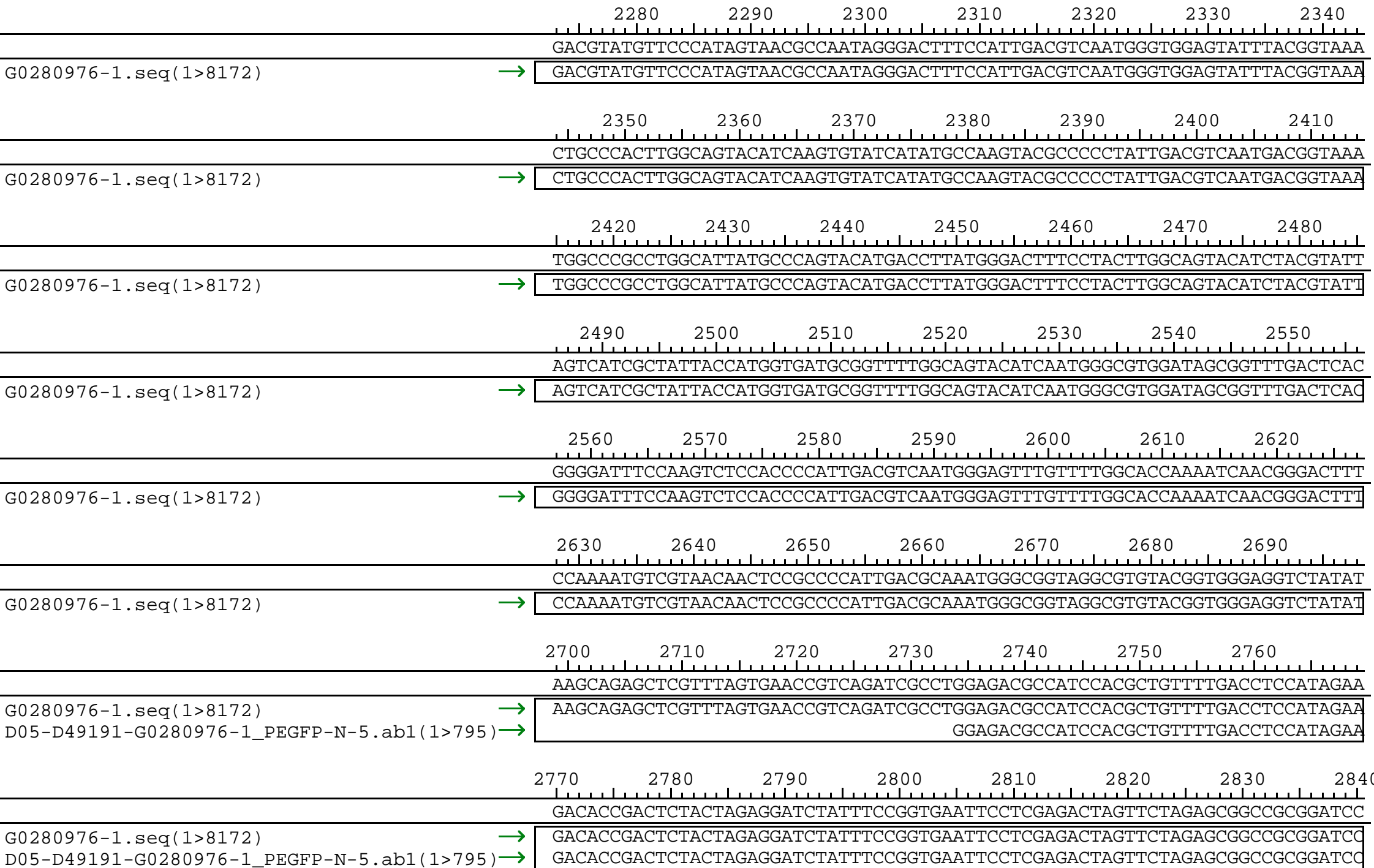


Project: G0280976-1.SQD Contig 1



Project: G0280976-1.SQD Contig 1





Project: G0280976-1.SQD Contig 1

2850 2860 2870 2880 2890 2900 2910
CGCCCCCTCTCCCTCCCCCCCCCTAACGTTACTGGCCGAAGCCGCTTGAATAAGGCCGGTGTGCGTTTGT
G0280976-1.seq(1>8172) → CGCCCCCTCTCCCTCCCCCCCCCTAACGTTACTGGCCGAAGCCGCTTGAATAAGGCCGGTGTGCGTTTGT
D05-D49191-G0280976-1_PEGFP-N-5.ab1(1>795) → CGCCCCCTCTCCCTCCCCCCCCCTAACGTTACTGGCCGAAGCCGCTTGAATAAGGCCGGTGTGCGTTTGT

2920 2930 2940 2950 2960 2970 2980
CTATATGTTATTTTCCACCATATTGCCGTCTTTTGGCAATGTGAGGGCCCGAAACCTGGCCCTGTCTTCT
G0280976-1.seq(1>8172) → CTATATGTTATTTTCCACCATATTGCCGTCTTTTGGCAATGTGAGGGCCCGAAACCTGGCCCTGTCTTCT
D05-D49191-G0280976-1_PEGFP-N-5.ab1(1>795) → CTATATGTTATTTTCCACCATATTGCCGTCTTTTGGCAATGTGAGGGCCCGAAACCTGGCCCTGTCTTCT

2990 3000 3010 3020 3030 3040 3050
TGACGAGCATTCCCTAGGGGTCTTTCCCCTCTCGCCAAAGGAATGCAAGGTCTGTTGAATGTCGTGAAGGAA
G0280976-1.seq(1>8172) → TGACGAGCATTCCCTAGGGGTCTTTCCCCTCTCGCCAAAGGAATGCAAGGTCTGTTGAATGTCGTGAAGGAA
D05-D49191-G0280976-1_PEGFP-N-5.ab1(1>795) → TGACGAGCATTCCCTAGGGGTCTTTCCCCTCTCGCCAAAGGAATGCAAGGTCTGTTGAATGTCGTGAAGGAA

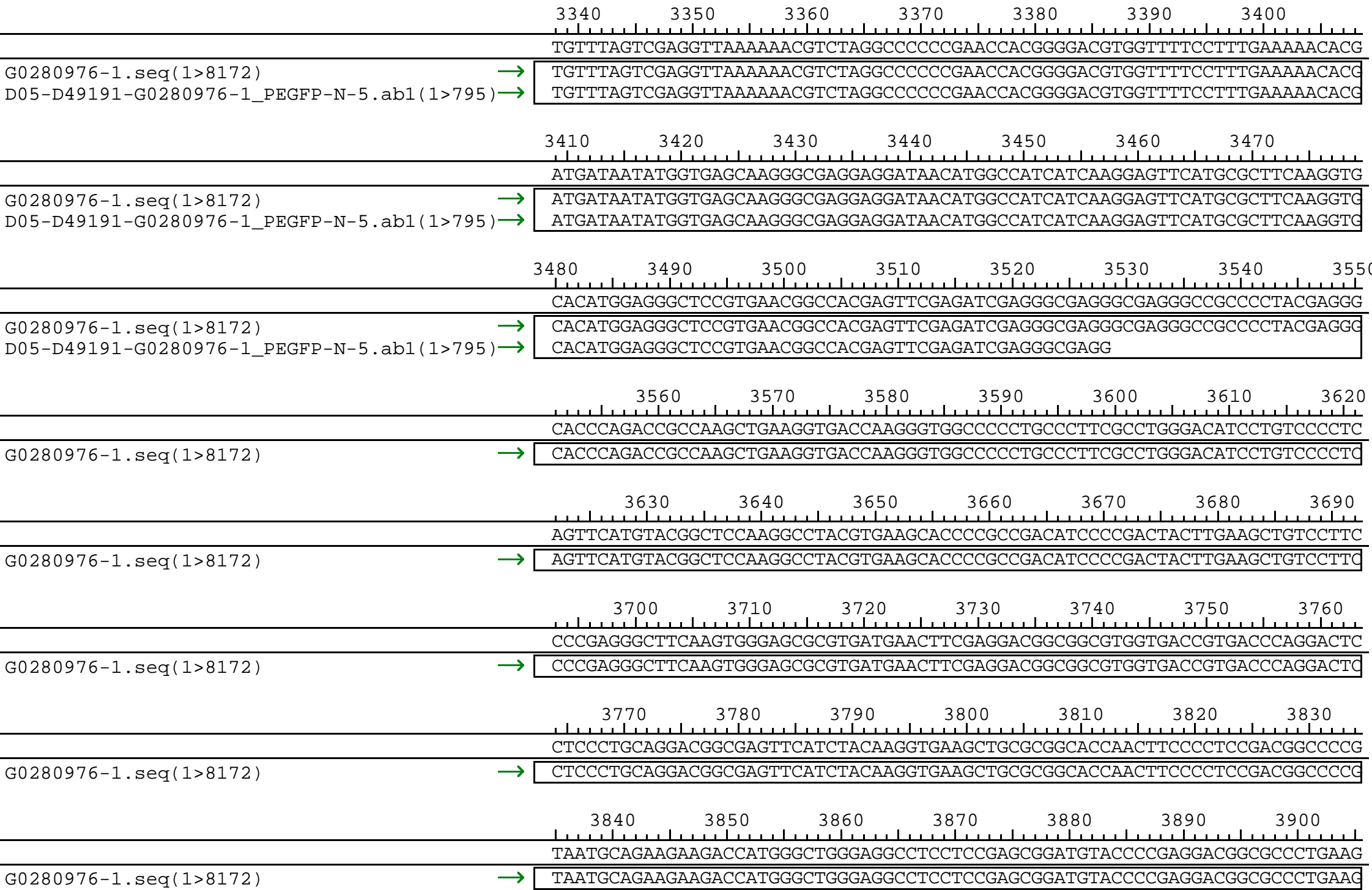
3060 3070 3080 3090 3100 3110 3120
GCAGTTCCTCTGGAAGCTTCTTGAAGACAAACAACGTCTGTAGCGACCCTTTCAGGCAGCGGAACCCCC
G0280976-1.seq(1>8172) → GCAGTTCCTCTGGAAGCTTCTTGAAGACAAACAACGTCTGTAGCGACCCTTTCAGGCAGCGGAACCCCC
D05-D49191-G0280976-1_PEGFP-N-5.ab1(1>795) → GCAGTTCCTCTGGAAGCTTCTTGAAGACAAACAACGTCTGTAGCGACCCTTTCAGGCAGCGGAACCCCC

3130 3140 3150 3160 3170 3180 3190
ACCTGGCGACAGGTGCCTCTGCGGCCAAAAGCCACGTGTATAAGATACACCTGCAAAGGCGGCACAACCCC
G0280976-1.seq(1>8172) → ACCTGGCGACAGGTGCCTCTGCGGCCAAAAGCCACGTGTATAAGATACACCTGCAAAGGCGGCACAACCCC
D05-D49191-G0280976-1_PEGFP-N-5.ab1(1>795) → ACCTGGCGACAGGTGCCTCTGCGGCCAAAAGCCACGTGTATAAGATACACCTGCAAAGGCGGCACAACCCC

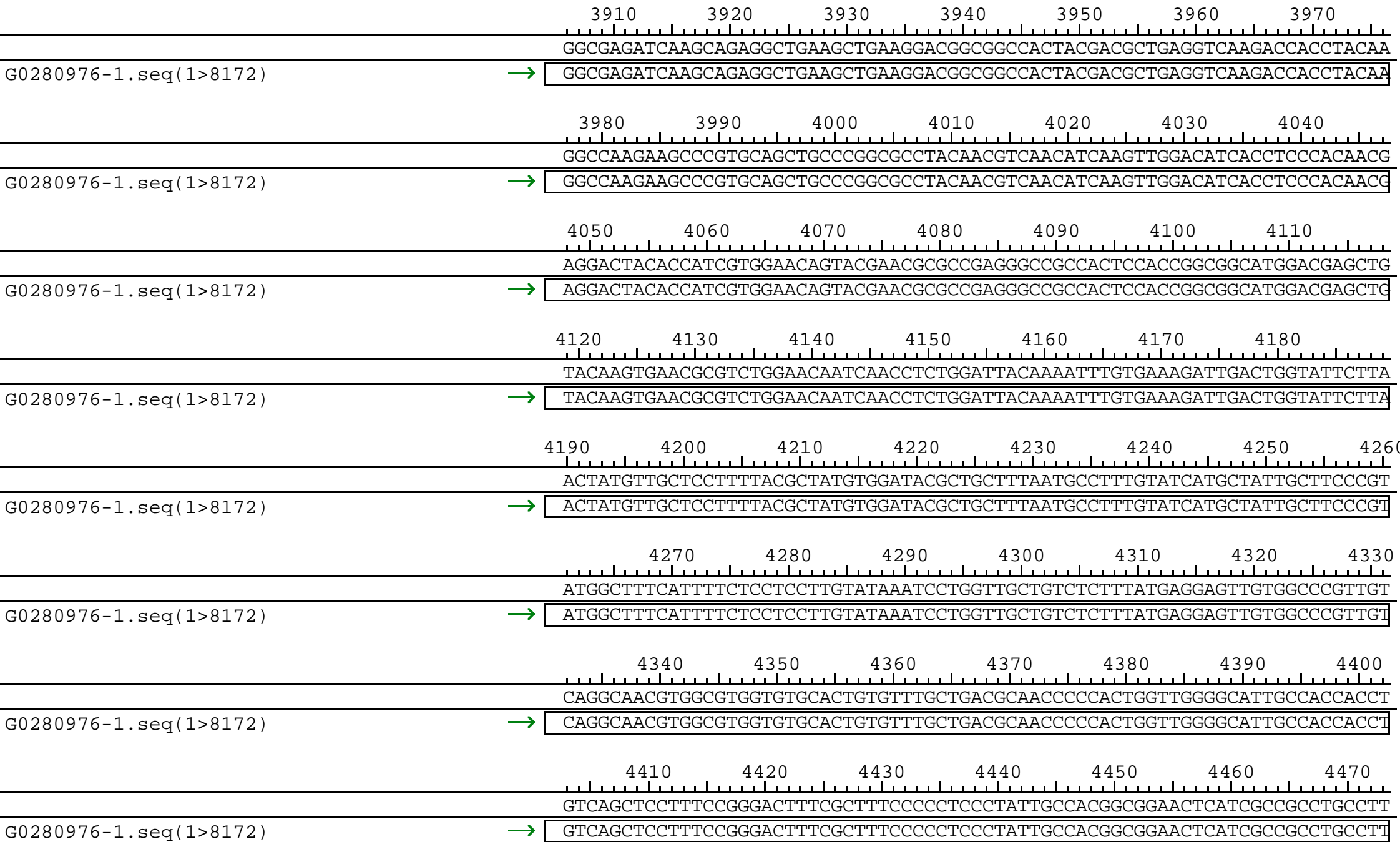
3200 3210 3220 3230 3240 3250 3260
AGTGCCACGTTGTGAGTTGGATAGTTGTGGAAAGAGTCAAATGGCTCACCTCAAGCGTATTCAACAAGGGG
G0280976-1.seq(1>8172) → AGTGCCACGTTGTGAGTTGGATAGTTGTGGAAAGAGTCAAATGGCTCACCTCAAGCGTATTCAACAAGGGG
D05-D49191-G0280976-1_PEGFP-N-5.ab1(1>795) → AGTGCCACGTTGTGAGTTGGATAGTTGTGGAAAGAGTCAAATGGCTCACCTCAAGCGTATTCAACAAGGGG

3270 3280 3290 3300 3310 3320 3330
CTGAAGGATGCCCAGAAGGTACCCCATTTGTATGGGATCTGATCTGGGGCCTCGGTGCACATGCTTTACATG
G0280976-1.seq(1>8172) → CTGAAGGATGCCCAGAAGGTACCCCATTTGTATGGGATCTGATCTGGGGCCTCGGTGCACATGCTTTACATG
D05-D49191-G0280976-1_PEGFP-N-5.ab1(1>795) → CTGAAGGATGCCCAGAAGGTACCCCATTTGTATGGGATCTGATCTGGGGCCTCGGTGCACATGCTTTACATG

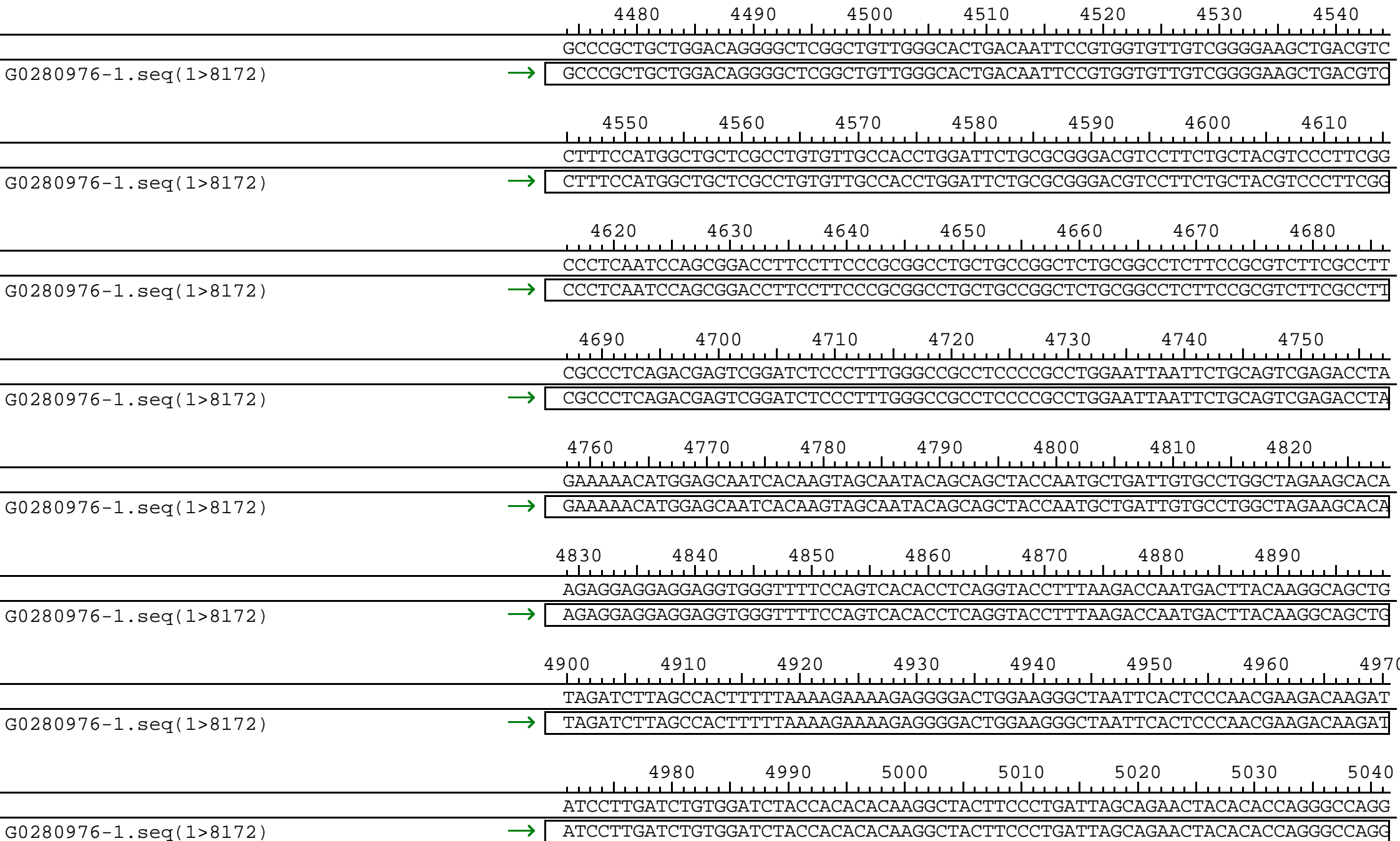
Project: G0280976-1.SQD Contig 1



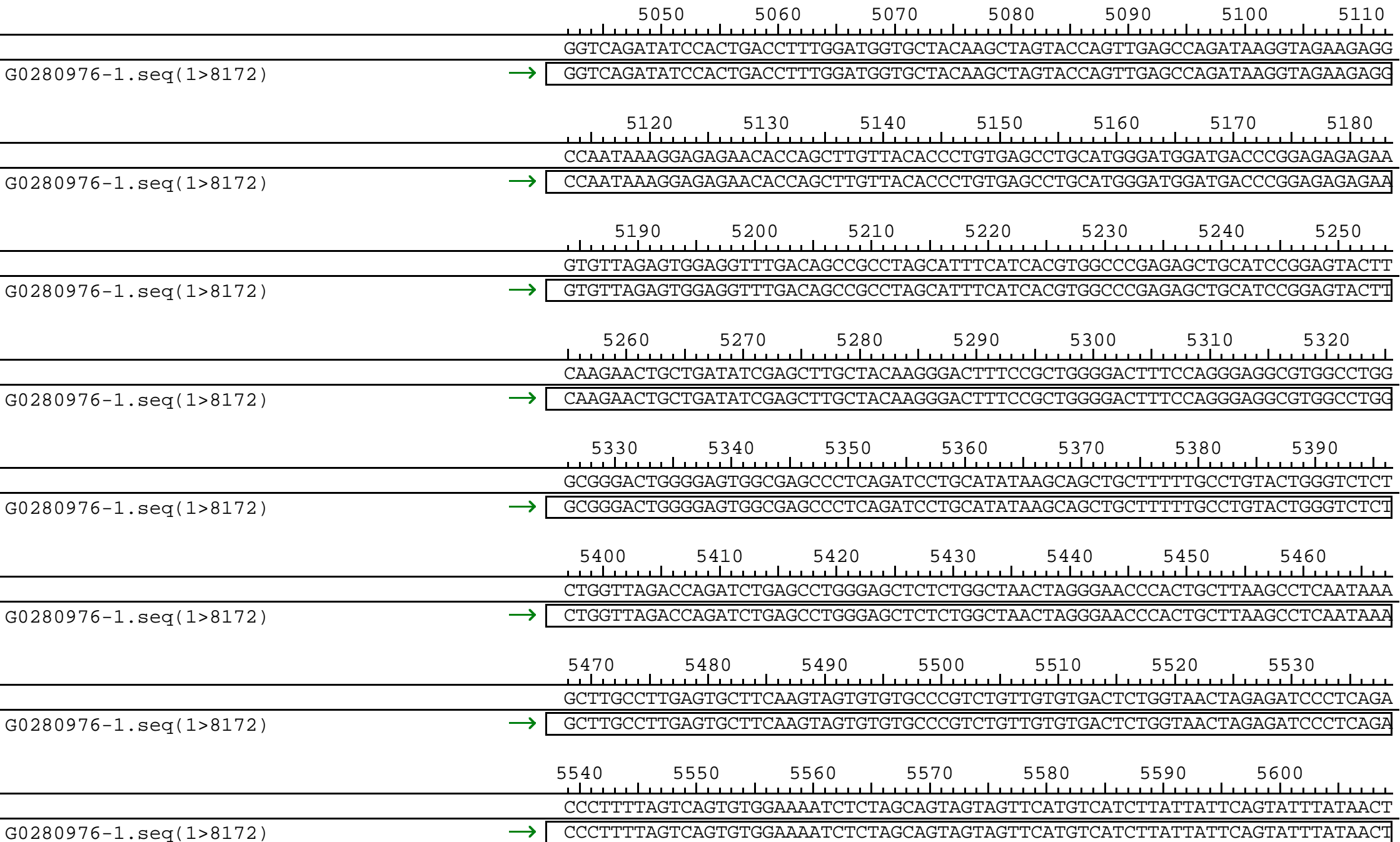
Project: G0280976-1.SQD Contig 1



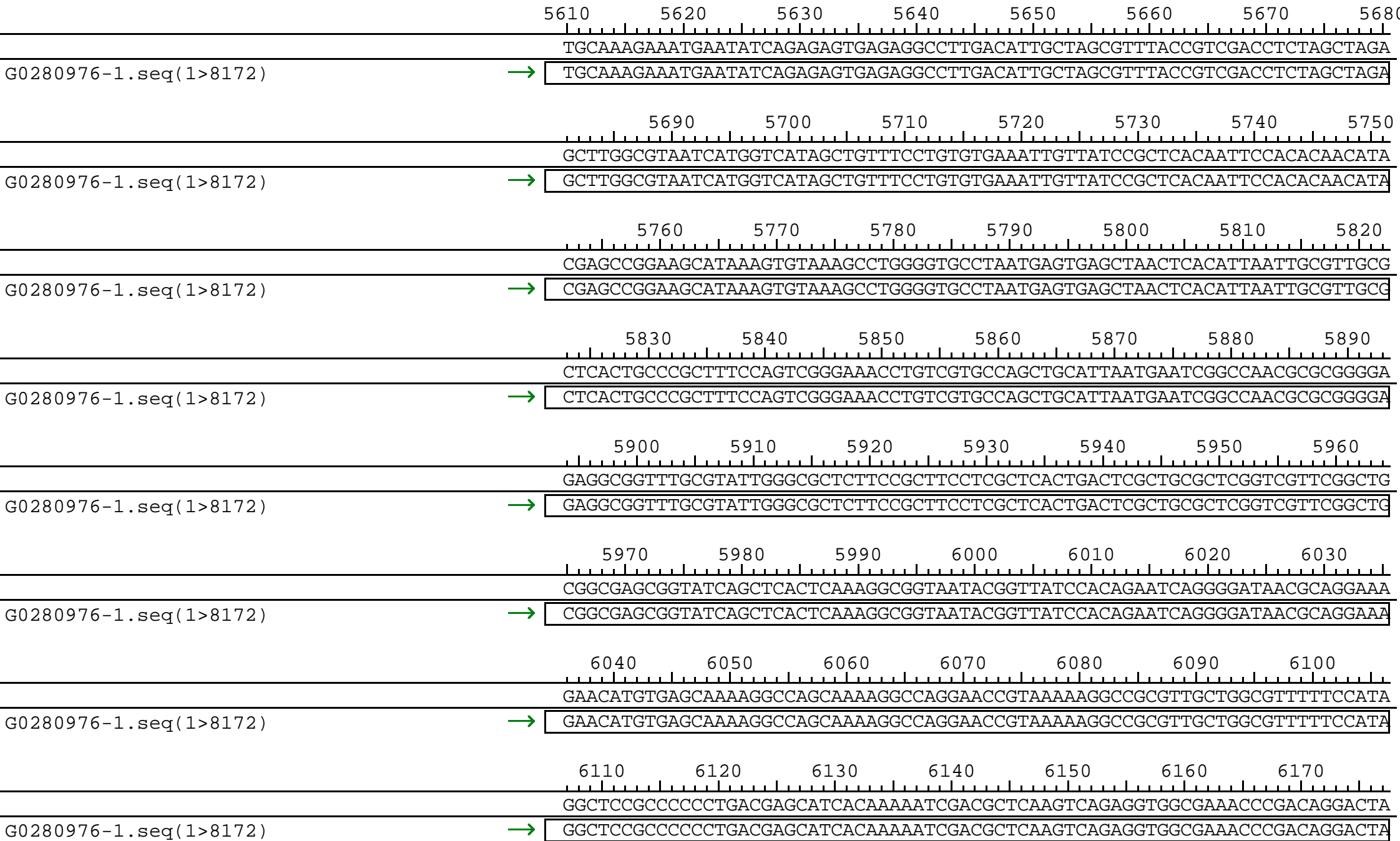
Project: G0280976-1.SQD Contig 1



Project: G0280976-1.SQD Contig 1



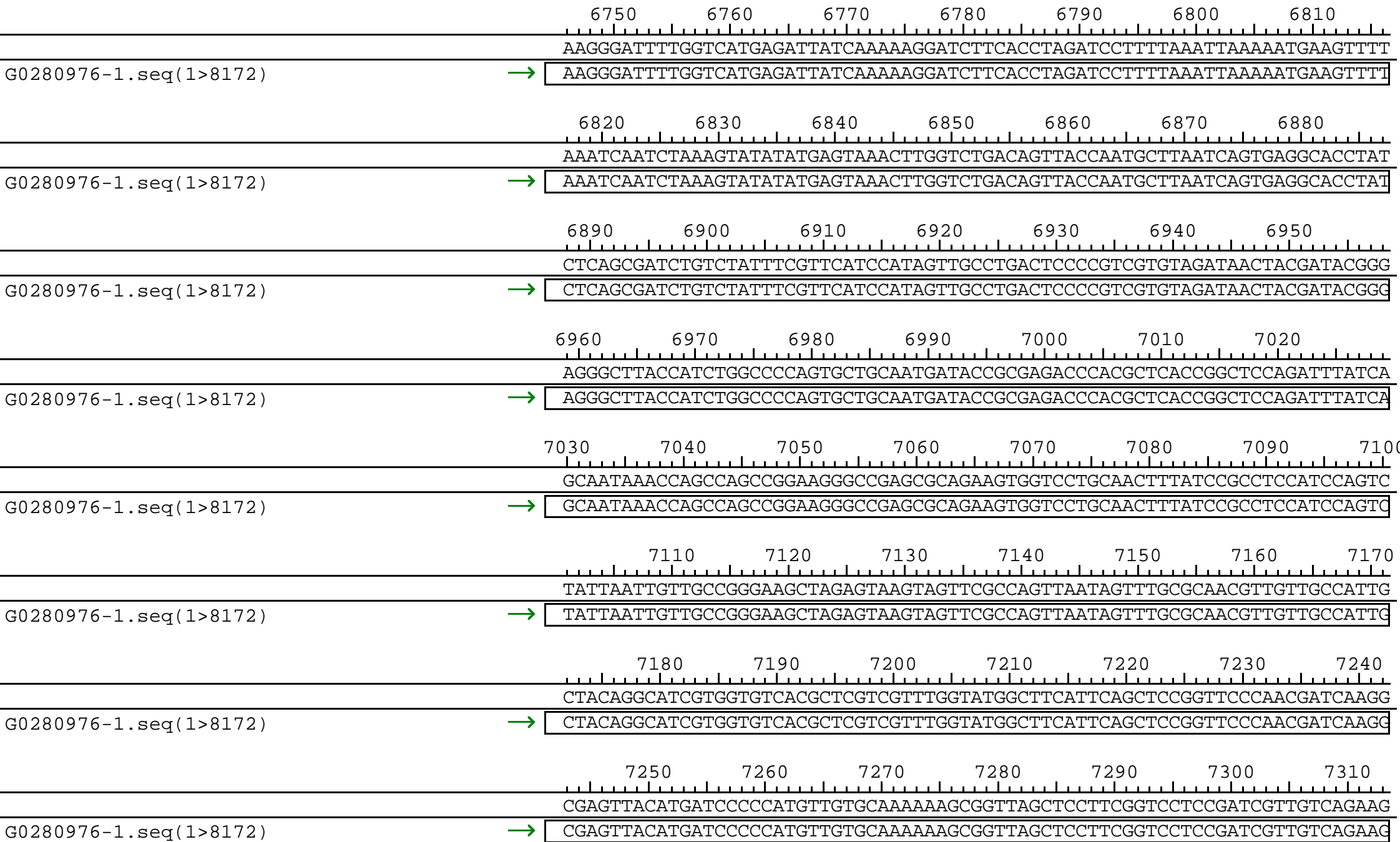
Project: G0280976-1.SQD Contig 1



Project: G0280976-1.SQD Contig 1



Project: G0280976-1.SQD Contig 1



Project: G0280976-1.SQD Contig 1



7890 7900 7910 7920 7930 7940 7950
TGCAGCTTATAATGGTTACAAATAAAGCAATAGCATCACAAATTCACAAATAAAGCATTTTTTTCTACTGC
G0280976-1.seq(1>8172) → TGCAGCTTATAATGGTTACAAATAAAGCAATAGCATCACAAATTCACAAATAAAGCATTTTTTTCTACTGC

7960 7970 7980 7990 8000 8010 8020
ATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGATCAACTGGATAACTCAAGCT
G0280976-1.seq(1>8172) → ATTCTAGTTGTGGTTTGTCCAAACTCATCAATGTATCTTATCATGTCTGGATCAACTGGATAACTCAAGCT

8030 8040 8050 8060 8070 8080 8090
AACCAAATCATCCCAAACCTCCCACCCCATACCCTATTACCACTGCCAATTACCTGTGGTTTCATTTACT
G0280976-1.seq(1>8172) → AACCAAATCATCCCAAACCTCCCACCCCATACCCTATTACCACTGCCAATTACCTGTGGTTTCATTTACT

8100 8110 8120 8130 8140 8150 8160
CTAAACCTGTGATTCCTCTGAATTATTTTCATTTTAAAGAAATTGTATTTGTTAAATATGTACTACAAACT
G0280976-1.seq(1>8172) → CTAAACCTGTGATTCCTCTGAATTATTTTCATTTTAAAGAAATTGTATTTGTTAAATATGTACTACAAACT

8170
TAGTAGT
G0280976-1.seq(1>8172) → TAGTAGT