GG TCAA TTG T CTGTC A GCAG G TA GC CG T TACG AC A GAA T GT T TT CAT TAAT CAA GAAC GAAA GT TAT G GGT TC GAAGGC G AT CAAATA CC GC CCT AGT CAT AAC C GT AAAC G ATAT AGACTAGGC
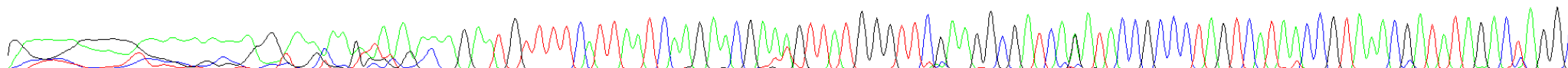
GT CTGC T C GGGT TTAT TT AT GAC C GA GT AGA AGC CT T CCC GG AAAC GAAA GT C TT CGT ATT C C GGGGG AAGT ATAG C TG CAAA GCC G AAAC TT AAAG GAA TT GAC GG AAGGG C ACCACCA GGGG TG
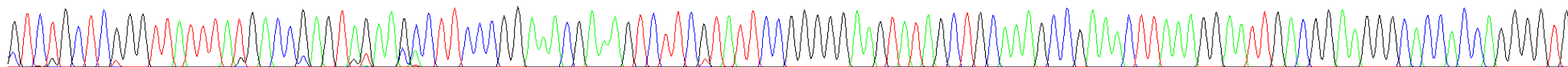
GAGC CTG C GGC TT AAT TT GACT C AAC AC GGGG AAAC T CACC C GGC CCG GA C ATT T GAAG GA TT GACA GT AT GAAGT GC TTT C GAG ATT AGAT GGGT GGT GGT G CAT GGC C GTT CTTA GTT GGT GGA
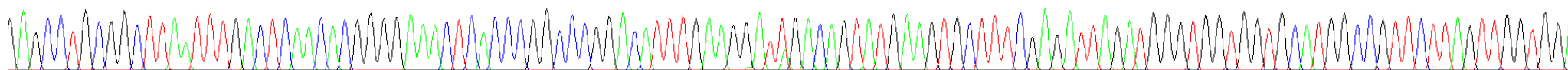
GT GAT TTG T C T GGT TTAA TT CC GAT AAC GAAC GAGA C TTT AT CCT GCT AAAT A GTAAT CAT CT AATT TCT GGGT G ATT TAC TT CTT AGA GGG ACAAA C GGT TTT CCA GCC GT AT GAAAGAAAG C AAT A
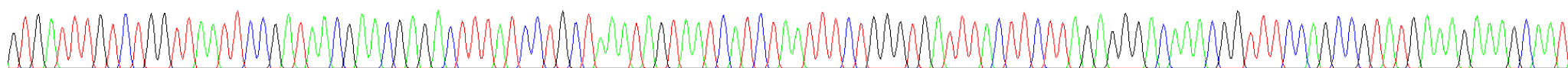
ACAGGT CT GTGA T G CCC TT AGA T GT CC GGGG CT GCAC GCG CGC TAC ACT G AA GGAA T CA GCG T GT GGGG T AACCT GA CCC G AAA GGG TT GGG T AAT CC GGT G AAAAT CCT T C GTGAT T GGGA TC
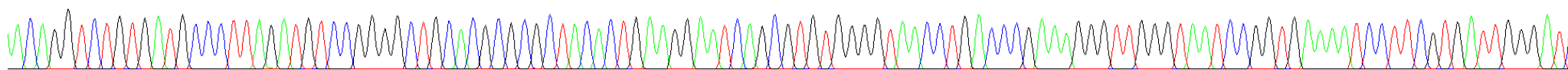
630   640   650   660   670   680   690   700   710   720   730   740   750
GGGACTTGCAATTATTTCCCTTGAACGAGGAATCCCTAGTAAGTGCAGGTCATCAGCTTGCGCTGATTACGTCCCTGCCCTTTGTACACACCGCCCGTCGCTGCCCGGAACTGAACCATTTCG
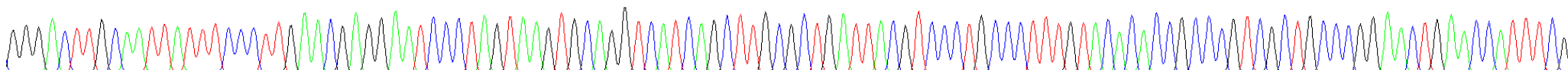
760   770   780   790   800   810   820   830   840   850   860   870
AGAAATGTGGGGACTGACGACTTGAGGCCTTCACGGGCTTCTTGTTGACGGAAACCGCCTTAATCGCAGTGGTTTGAACCGGGCAAAGTCGTAGCCAAGGGTAACCAGTGGCAAAGCCCCACGCC
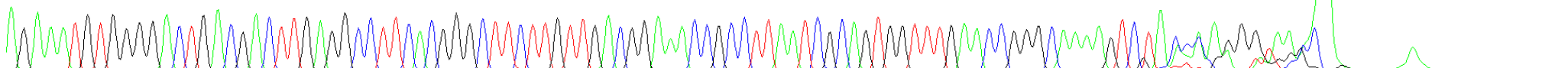
880   890   900   910   920   930   940   950   960   970   980   990
GGCCGCCACCCCAACCCCCCCCCCCAAACAGGGGAGGAGGGAACGGGAGGAGGAGCGGCGGAGGGGAGGAGAGAAAAAGAGAGGGGGGCAGGGAAGAGGAACAGGAGGGCAAGGC

1000   1010   1020   1030   1040   1050   1060   1070   1080   1090   1100
GGGACGGGAGGCAGGCAGCGAGGGGGGCGGGCCGGGCCCGGCGAGGCAGGGGAGGGGGCCGATCGCCAGGGCGCGCCAGGCAAGGGAGGCAGGGCGGAGGGAGCGGGAGGGGGGAC

1110   1120   1130   1140   1150   1160   1170   1180   1190   1200   1210   1220
CGGGGGGAGAAGGCGGGGGTGGGCGAAGGGGACGAACGCGCACAGTACTCGTGAGAGGGAACGTACGGAGGTGATGGCGGGGGAGGGCTGGAGGGAGAGTAGATGGGCGTCGTAGGGAT