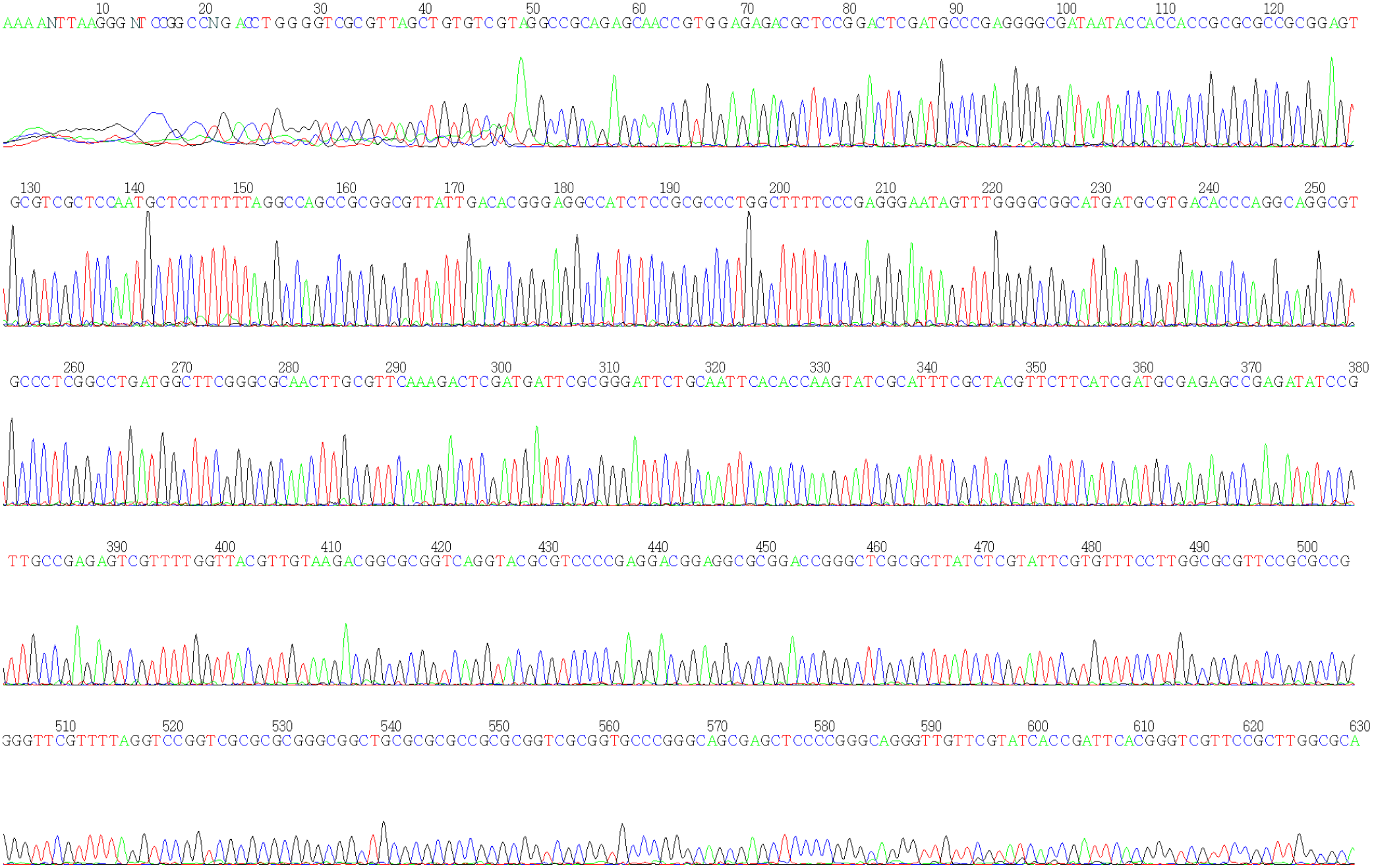
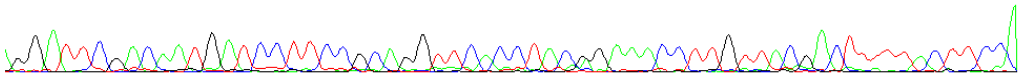

10 20 30 40 50 60 70 80 90 100 110 120
AAAANTTAAGGGNT CCGG CCNG ACCT GG GGTCGC GTTAGCT GTGTC GTAGGCCGCAGAGCAACCGTGGAGAGACGCTCCGGACTCGATGCCCGAGGGGCGATAATACCACCACCGCGCGCCGCGGAGT
130 140 150 160 170 180 190 200 210 220 230 240 250
GCGTCGCTCCAATGCTCCTTTTTAGGCCAGCCGCGGCGTTATTGACACGGGAGGCCATCTCCGCGCCCTGGCTTTTCCCGAGGGAATAGTTTGGGGCGGCATGATGCGTGACACCCAGGCAGGCGT
260 270 280 290 300 310 320 330 340 350 360 370 380
GCCCTCGGCCTGATGGCTTCGGGCGCAACTTGCGTTCAAAGACTCGATGATTCGCGGGATTCTGCAATTCACACCAAGTATCGCATTTCGCTACGTTCTTCATCGATGCGAGAGCCGAGATATCCG
390 400 410 420 430 440 450 460 470 480 490 500
TTGCCGAGAGTCGTTTTGGTTACGTTGTAAGACGGCGCGGTCAGGTACGCGTCCCCGAGGACGGAGGCGCGGACCGGGCTCGCGCTTATCTCGTATTCGTGTTTCCTTGGCGCGTTCCGCGCCG
510 520 530 540 550 560 570 580 590 600 610 620 630
GGGTTCGTTTTAGGTCCGGTCGCGCGCGGGCGGCTGCGCGCGCCGCGCGGTCGCGGTGCCCGGGCAGCGAGCTCCCCGGGCAGGGTTGTTCGTATCACCGATTCACGGGTCGTTCCGCTTGGCGCA

640 650 660 670 680 690
GGATTCGACAATGATCCTTCCGCAGGTTCACCTACGGAAACCTTGTTACGACTTTTTACTTC