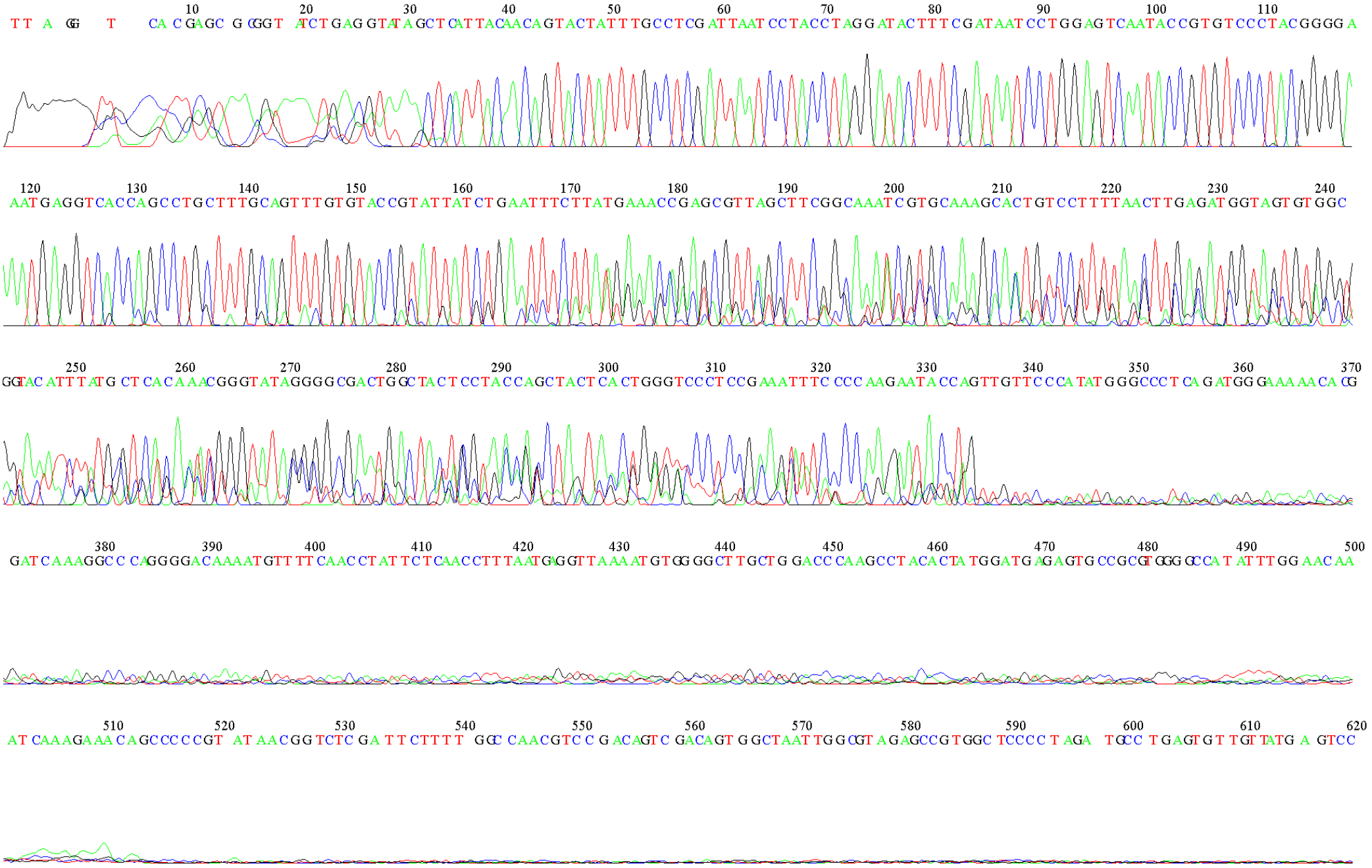
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250 260 270 280 290 300 310 320 330 340 350 360 370
GGTAC ATTTATG CT CACACAAACGGGTATAGGGGCGACTGGCTACT CCTACCAGCTACTCA CTGGGTCCCTCCGAAATTTC CCC AAGAATACCAGTTGTTCCCA TATGGGCCCT CA GATGGGAAAAACA CG
380 390 400 410 420 430 440 450 460 470 480 490 500
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510 520 530 540 550 560 570 580 590 600 610 620
AT CAAAGAAACAGCCCCCGT AT AACGGTCTC GA TTCTTTTT GGC CAACGTCCG ACAGTC GACAGT GGCTAATTGGCGT AGAGCCGTGGC TCCCCT AGA TGCC TGAGTGTTGT TATG A GTCC

630 640 650 660 670 680 690 700 710 720 730 740
GTGTCTAGGT TGAGGGCGTAGGGCTTATCGTGACTCGT AAGCAAAG AGCTGCTGAT CTTACAAAAACA CTTTCGGTAGTG GACCTCTGAACGAGACGGAGGGTGTGCGAAGAGTAGGCTC GTA
750 760 770 780 790 800 810 820 830 840 850 860 870 880
GTGG GAGCAATAACAGGGGG ATTCTTCGAG AA G ATAAACGATGACTCTGT AAGGGTCTTAAGTCTTGGTGTGCAAGACATGCCGGTGTTC TT TTG CCTCA C ACTACAAGGAACACACCGCGATACCGAAGGGTGAATTAAG
890 900 910 920 930 940 950 960 970 980 990 1000 1010 1020 1030
AATA TCC GGC CCCT ATTTTCTT G AATC GGGAA AAAAT ATCAAAT ATTT T GGG GCGCGT GGT GTT GGGCTTTGTGAGAGCGCGTTAGCTCCTT T GGACTGCTGCGAATG AAGACCTGAT C ACTGTATTCGAG TCATCATATGTTTTGTTC
1040 1050 1060 1070 1080 1090 1100 1110 1120 1130 1140 1150 1160 1170 1180
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1190 1200 1210 1220 1230 1240 1250 1260 1270 1280 1290 1300 1310 1320 1330 134(
AATGCGGGATGCTA G CGGAGTGGTGAT GGTC GGCGC GCATGGCAG AGAGGTGGGAC GATGA GGA TGC CATG CGTCACTTC TGTATGCGGACG CACA GGTGTTCATT CGTGT TCCGGTCA C T C AGAT CCGTG CACGTCCGC CT TTCA GCGA AACA