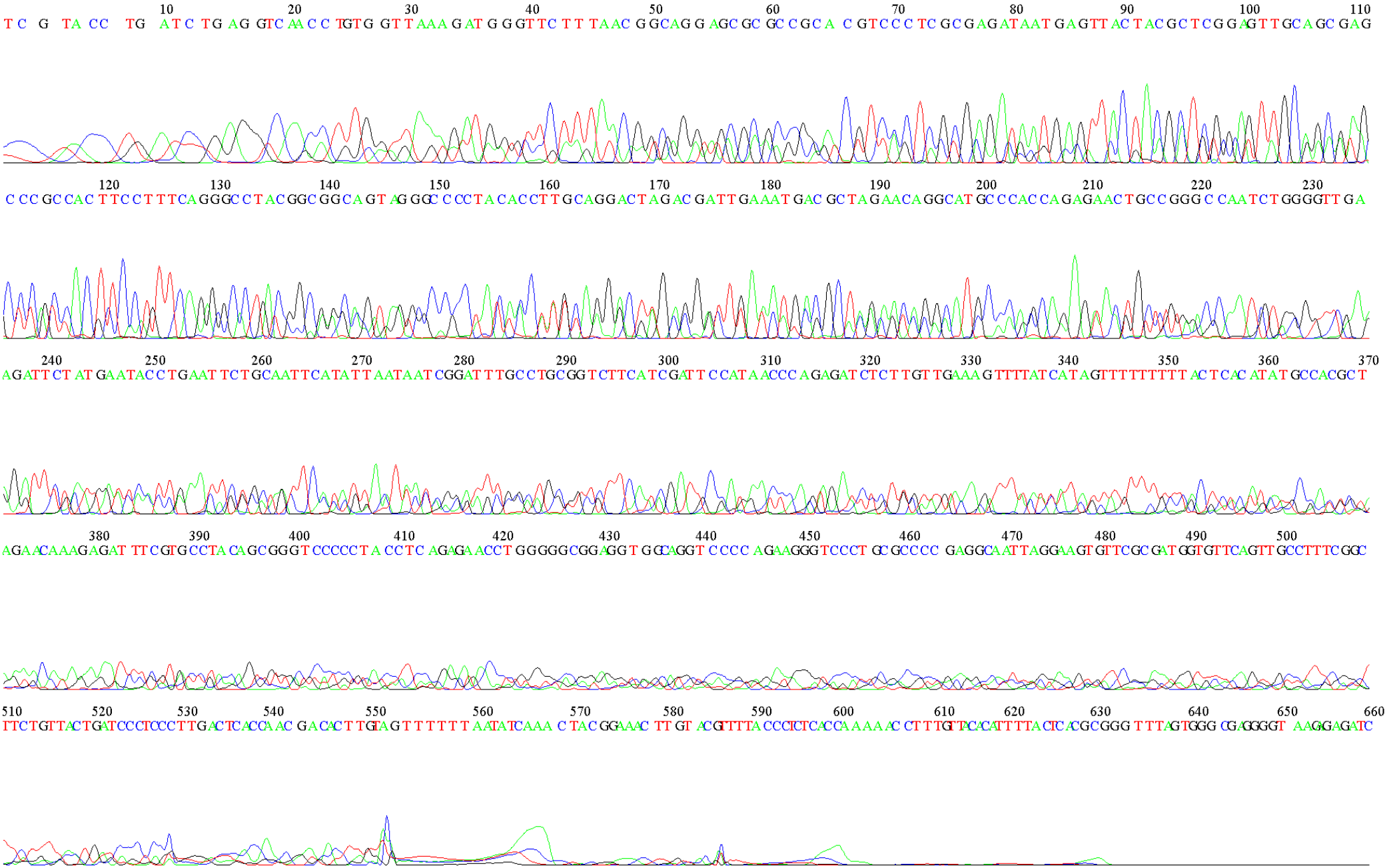
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120 130 140 150 160 170 180 190 200 210 220 230
C CC GC CAC T TC C T TTCA GGGC C T AC GGC GGCA GT AG GGC C CC T A CAC C TT GCAGG AC T AG AC GATT GAAAT GAC GC T AG AACA GGCAT GC C CAC C A GAG AAC T GC C GGG C CAAT C T GGGGTT G A
240 250 260 270 280 290 300 310 320 330 340 350 360 370
AG ATT C T AT GAATAC C T GAAT T C T GCAATT C ATATT AATAAT C GGAT T T GC C T GC GG T C TTCAT C GATT C C ATA AC C C A GAGAT C T C T T GTT GAAA G TTTTAT C ATA GTTTTTTTTTT AC T C AC ATAT GC C AC GC T
380 390 400 410 420 430 440 450 460 470 480 490 500
AGAACAAAGAGAT TTC GTGC C TA CAGC GGGT C C C C C T A C C T C AGAGAA C C T GGGGGC GGAGGT GC AGGT C C C C AGAAGGGTC C C C T GC GC C C C GAGGC AAT T AGGAAGT GTTC GC GAT GGTGTTC AGTT GC C TTTC GGC
510 520 530 540 550 560 570 580 590 600 610 620 630 640 650 660
TTC T GTTAC T GAT C C C T C C C TTGAC T CAC CAAC GAC AC T T GTAGTTTTTT AATAT CAAA C TAC GGAAAC TT GT AC GTTTTAC C C C T C TAC CAAAAAC C TTTGTTAC AC ATTTAC T C AC GC GGGG TTTAGTGGG CGAGGGGT AAGGAGATC

AC GAT G