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G ACA AATTAAATT G C GGCCCC TA G AGGT CCCC T T TTTTAT TT TAAAAT TT TTT CACA AAAC GG TTT ACAAG C AT AAAAT CT CT G AAG AT GT GT AT AAG AG ACA GC CTT G AT GCT GCT GG CCCG



130 140 150 160 170 180 190 200 210 220 230 240 250

CT CCT GCCCC AAGG T ACG C AGC AAGG C GG T ACT CT C GGC CCA CG ACT TG AT CAC G GCC TC GCG C AGC TT GGC GGC GCG C TCGT CACG GCC CAACT GCT CCC ATT CGG CAT GC GGCT GC TG TTT CA



260 270 280 290 300 310 320 330 340 350 360 370

GGC GGG T ACG GT A AAT GCC TTC GAG C AT C GC GGG CTT GT C GCT GT C GT C CAC C AC CAG C AT C G AGG C ATT GGC C GGC CAC C GGG T CACCC C GT CGC TG C AGG AT C TTGG ACC GGG T GG CCT GGT AC



380 390 400 410 420 430 440 450 460 470 480 490 500

T CAC GCT C AAGGG GC TGT T GGGC AAT CA GC GGG C CGT C GCT GGC CT GGGC GCT GGT GCC TT CG ATTT CCA GGC GC AGT TC C GGG C GC TCTTT CAG C GC GG ATG C CAG C TT GT C C AAGG ACG ACT G



510 520 530 540 550 560 570 580 590 600 610 620

GGC CTC GCC GC TG A GT T CG C CT GG AG CC GGGG G CG AA GGC CAC GT T GC C AAGG T CTT CC G AGC T ACC GCC GCT GA T CAGCC C GCC AAT GA ATTT G AAT GGC GCC TG CG C CGC GC GC A GC ACC AG

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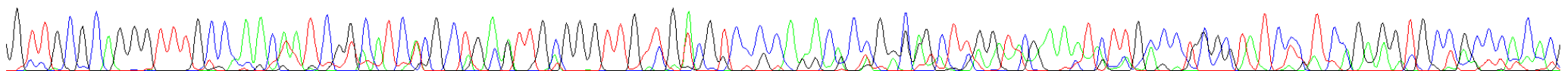
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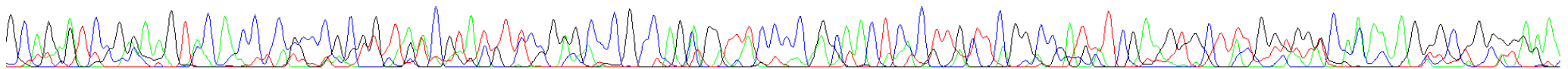
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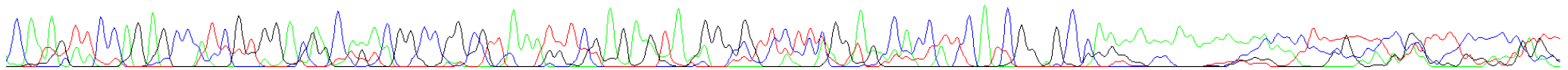
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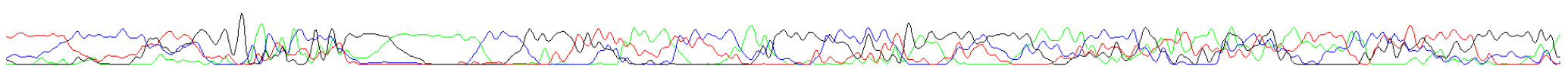
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