10 20 30 40 50 60 70 80 90 100 110 120
C GGTA   TA TTC T G G A GTT GTA AT TTA GT GAT CA GC CT GT G G AAA AAG A CAAATACATAAAT GTGTATGTATTG ACT CAAAT T T T T T TCTTT AAAAC TAT T CAG GT TAAT T TATA GAAAAT GCC
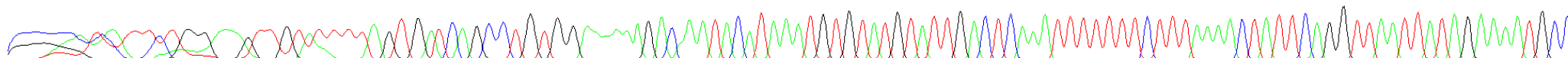

130 140 150 160 170 180 190 200 210 220 230 240
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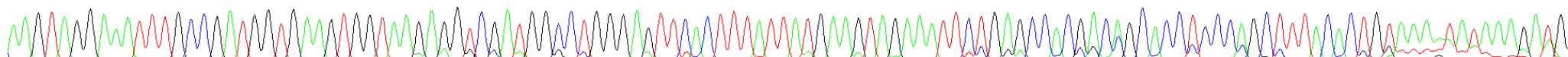
250 260 270 280 290 300 310 320 330 340 350 360 370
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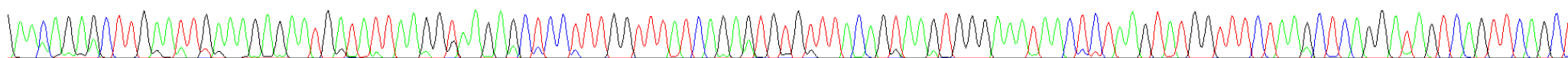
380 390 400 410 420 430 440 450 460 470 480 490
C GGTTT GAAGGCA GAGGATT GAGC TCTT GCTTT GGGT CACATACT GTTTTT GGCTCT GGGAAT GTAT GAAT GAGGCATAGT CCTT GTTCCCAA GAAGCT CACACT TACT GGG AAACT GACCC TTT
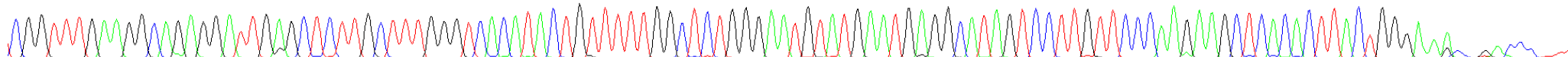
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TAAA AAA AAAA AA AAAAAAA CAGAAATTTA CCTT GCAAAAAAT AAT CACT GT AAC GGGAT AAAGGG TTTT AAT T GTT ACTTACA GCTTCAAA GCT GC GGGGT CCCCCC TCCCAAT GAAGA

```
620        630        640        650        660        670        680        690        700        710        720        730
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```

```
740        750        760        770        780        790        800        810        820        830        840        850
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860        870        880        890        900        910        920        930        940        950
TTGGACGCAACAATCCTCAAAATAACGAAAAATAAATTAAAAATTTATACAATCCCTGGGAAAATTATATGGAAGTGGAAAAAAGGGCAGAAGGGAACCCGCCGGG
```

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960        970        980        990        1000       1010       1020       1030       1040       1050       1060
AGCGCTTTTTAATTAGAAACGTTATCGATAGGCGCGGGGCCTAATCGGGTTTTTAAGGCGGGCGACGTCGGGAAGAGGGGGCATTAAGGAAGGGCGGGGCCCCTAC
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1070       1080       1090       1100       1110       1120       1130       1140       1150       1160       1170
GATTAATTATTATACTTTTTTTATTAAGTCCTTTGGGTATAGCGAGGTTAATTTGCTTGAGCTGAGCGGGGGGCGGTGAGGAAGAATATTTCGGATAAAACGGGTAC
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