```
        10        20        30        40        50        60        70        80        90        100       110       120
GG CC CTTA ATT T GTATTT TTA TTC C TGACAT G GC A ACC CAAA GGC GT AAT G AAA GT AAA GGTC GT C TT GCACG G CTTATAC G GG ATCC GT TCCTCT GAGG ACGG CG CACCGT AGCCCGTCCCAACT
```
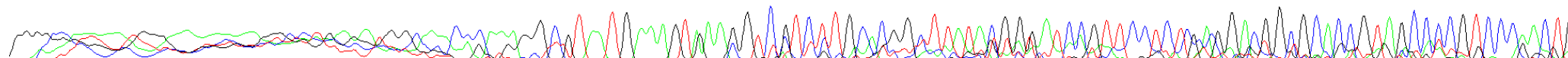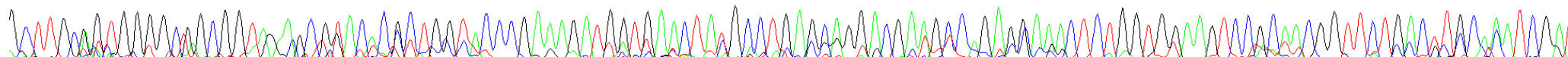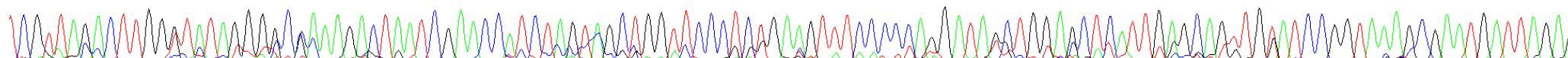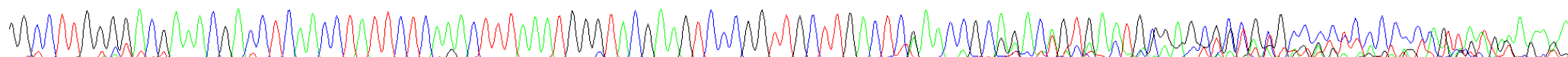


```
130       140       150       160       170       180       190       200       210       220       230       240       250
GC TT GCAGT G GGG C GGC GG TA AAGC GTACAC GCT GGT ACC CGAAAGAT GGT GAACT AT GCCT GAGCAGGACGAAGCCAGAGG AAAC T CT GGT GGAAGTCCGCAACGGTTCTGACGT GCAAAT CGA
```



```
          260       270       280       290       300       310       320       330       340       350       360       370
T CGTTAG ACTTGGGT ATAGG GGCGAAAG ACT AATC GAAC CAT CTAGTAGCTGGTTCCCTCTGAAGTTTCCCCCAGGATAGCTGGAGCTCATTGAGCAGTTGTATCCGGTAAAGCGAATGATTAGA
```
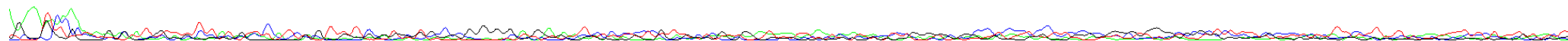


```
380       390       400       410       420       430       440       450       460       470       480       490       500
GG CCTTGGGGACGAAACGACCTCAACCTATTCTCAAACTTTAAATGGGTACGAAGTCCCGGTTGCTTGACTCGAACCGCAGACAGTGAATGCGAGCGCCGCGCCCCTCCCCACCGTATGTAAAA
```



```
         510       520       530       540       550       560       570       580       590       600       610       620
G AGC AAAG CAT GAGTT TGACAT GGTTTC AT CTGCCGGGGAGAAATCCCGTGTGCCGCGATCATATAAGCCTGGTCGCCTCCCCTCGGCGGGGCTTTTGCCGACTATTATAATTTTTAGTAT
```

```
       630        640        650        660        670        680        690        700        710        720        730        740
GTA AGACGTT GTCATCC CATCAAA CATCACGGA TTT TC GATTT  GCAGCAGCT CC GGCC G GCT GC CA  ACTGCC G CA ACCGCAGC GTT T T  CC GTT TTTCC GC CGC TC AAT AGGAGCTGG AAC
```

```
      750        760        770        780        790        800        810        820        830        840        850        860
T C AAGC CAGGAG CT GAA AG CTTTC TCTCC TT ATT AAT TGT GAT T T CATGT TTGGT  GC GTT GT ACGTAGC GCG T TT TG T T TTCAAGAT GCAATACCTGC GTCATGA C ACCATA ATTGT GCAT AG
```

```
       870        880        890        900        910        920        930        940        950        960        970        980
T  T CTAGAGTAT GTG ACTTTC A GG ATAC ACTCGCTT A AAGAT TT TCA TTTCTGCC GT CTGTGC A TGTT ATCATAGTA G GAGGCATCC GA GCT CAT GT CGT TGATAT AT GTCTATCCGC
```

```
      990       1000       1010       1020       1030       1040       1050       1060       1070       1080       1090       1100
GTTGTG TTTATGCT TGCC CACAG TTA CGGA GC ACGCG CTT GCAT TT GTC GA G T ATGAGT GTGGTTCTTATT TTC GATTAG TGAATAC TCAAAGCG TAG TATATT CGTTT CT G T CACTT AT
```

```
      1110       1120       1130       1140       1150       1160       1170       1180       1190       1200       1210       1220
GGCATG T AAG CTCT AG TTA TTG TA GTC AGAAT GCGT GTGATGTACT AAC GC GGATGC T AGCATCAGTCG C AT AGTCACTAGAG AAG AGTGACT GGCTGAT GCATGCA AGTGC TT GACTT CT CT AT C
```