```
           10        20        30        40        50        60        70        80        90       100       110       120
TAT GGG AG A CA GATAC CA CGT A GGA T T TAGC CAAG AAT GT CTT CATT AAT CAAG AACG AAAGT CAGAG GTT CGAAG GCG AT TAG AT ACC GC CATAGTT CT GA CC TTAAACAAT ACCT GCT GGC GAA
```
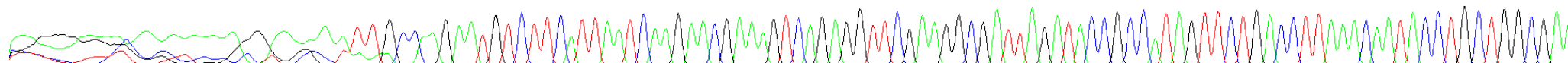


```
          130       140       150       160       170       180       190       200       210       220       230       240       250
CC GAT GGT GGCG AAAGCT GC CTT TT CGGGG AGC ACC CCGG AAAC GAAA GCATAT C GG AT CC GGGGG T AGT AT GGT T GCAAAGCT G AAA CTT AAAG GAATT GACG GAAG AGC ACC ACC AGG AGT GG
```
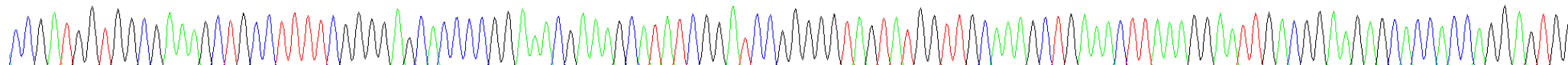


```
          260       270       280       290       300       310       320       330       340       350       360       370
AGC CTG CGG CTT AAT TT GACT CAAC AC GGG AAACCT CACC CGG CCC GG ACACCG T AAG GA TT GT CAGACT GATT GCT CTT ATAT GATT C GGT GGTT GGT GGT GCAT GGC CGT T CTT AGTT GGT GAA
```
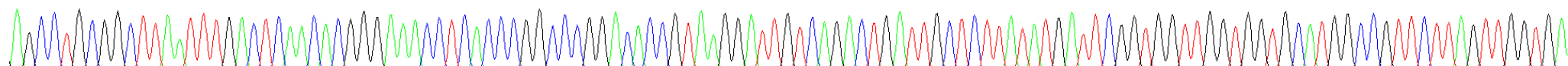


```
          380       390       400       410       420       430       440       450       460       470       480       490       500
GCG ATTT GT CAGG TTT ATT CC GGT AACGA GCG AGA CT CTA GGCT CT AATT GAATAC GGGGG AT CTTT GG ATTT CT GTA CTT TCTT AAAG GAATT GGT GGT AAAG AACG GCAT GC AAT AAA GCAATA
```
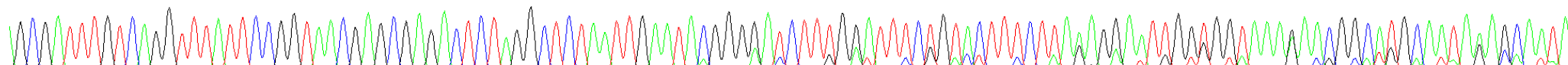


```
          510       520       530       540       550       560       570       580       590       600       610       620
ACAGG T CT GTGAT GCCCTT CGAT GTCC GGGGC TGC AC GCGT GCT ACACT GGGGT CCT CAGC AGTTT CCTAT CTT AAT CAGA CAC AGG T AAA CC GTT GCAAGG ACCCC GTGTT CGGGT TAT G AATT
```
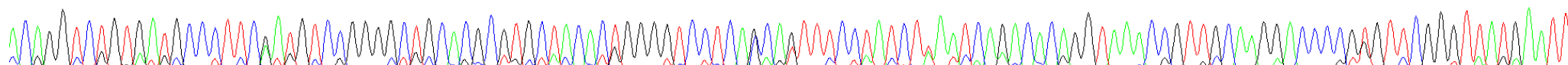
```
630          640          650          660          670          680          690          700          710          720          730          740          750
TGCTATTATTCTCCTCTAACGAGGAATTCCTATTAAGGACGAGTCATTATCTTGTTCTGATTATGTCCCTGCTCTTTGTACACACCGCCCGTCGCTACCCGGGACTGAACCTTTTCGAGAAATA
```
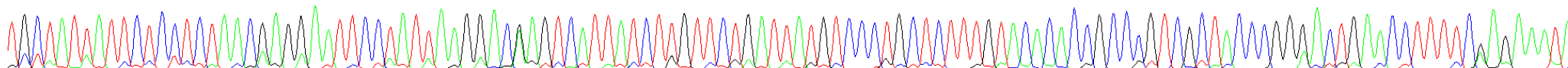


```
760          770          780          790          800          810          820          830          840          850
CAGGGAGTTGTTGCATAGGATTCGTCTTATGCGGCAGTGAACTATCTTGATCGCGAAGGTTTGAACCGGGTAAAATCACCCGGGGGGAAAAAAAAAAAAAAA
```