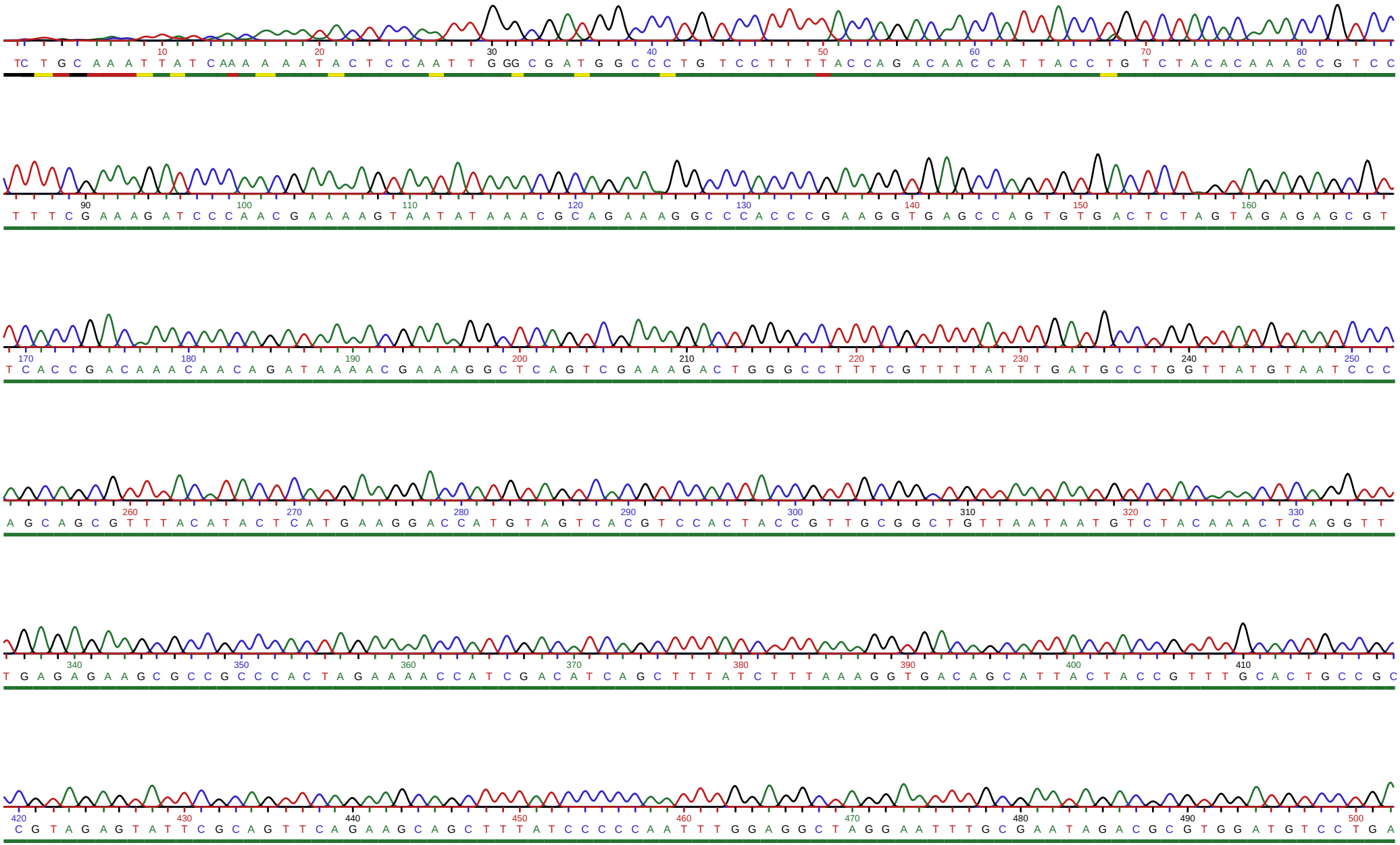
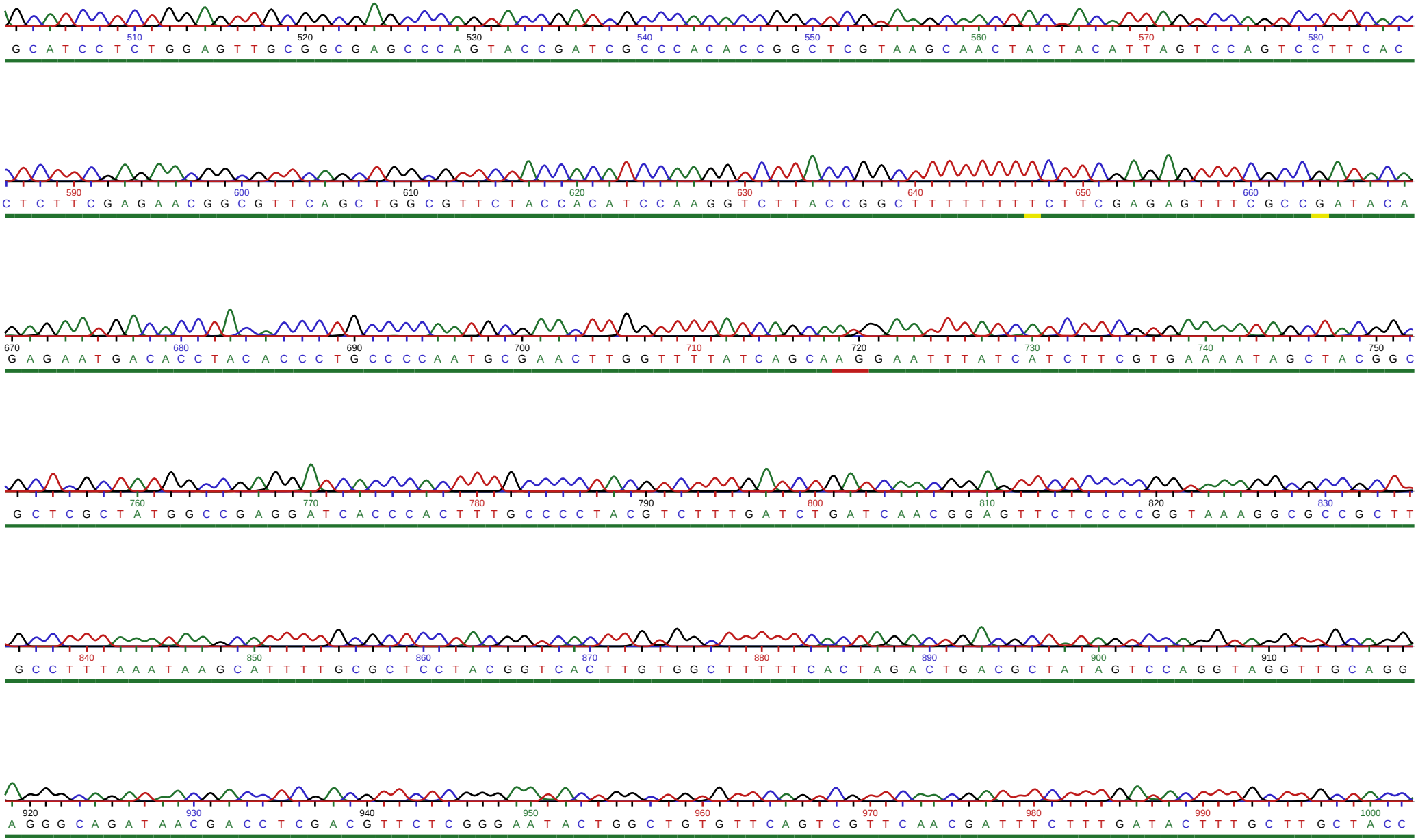
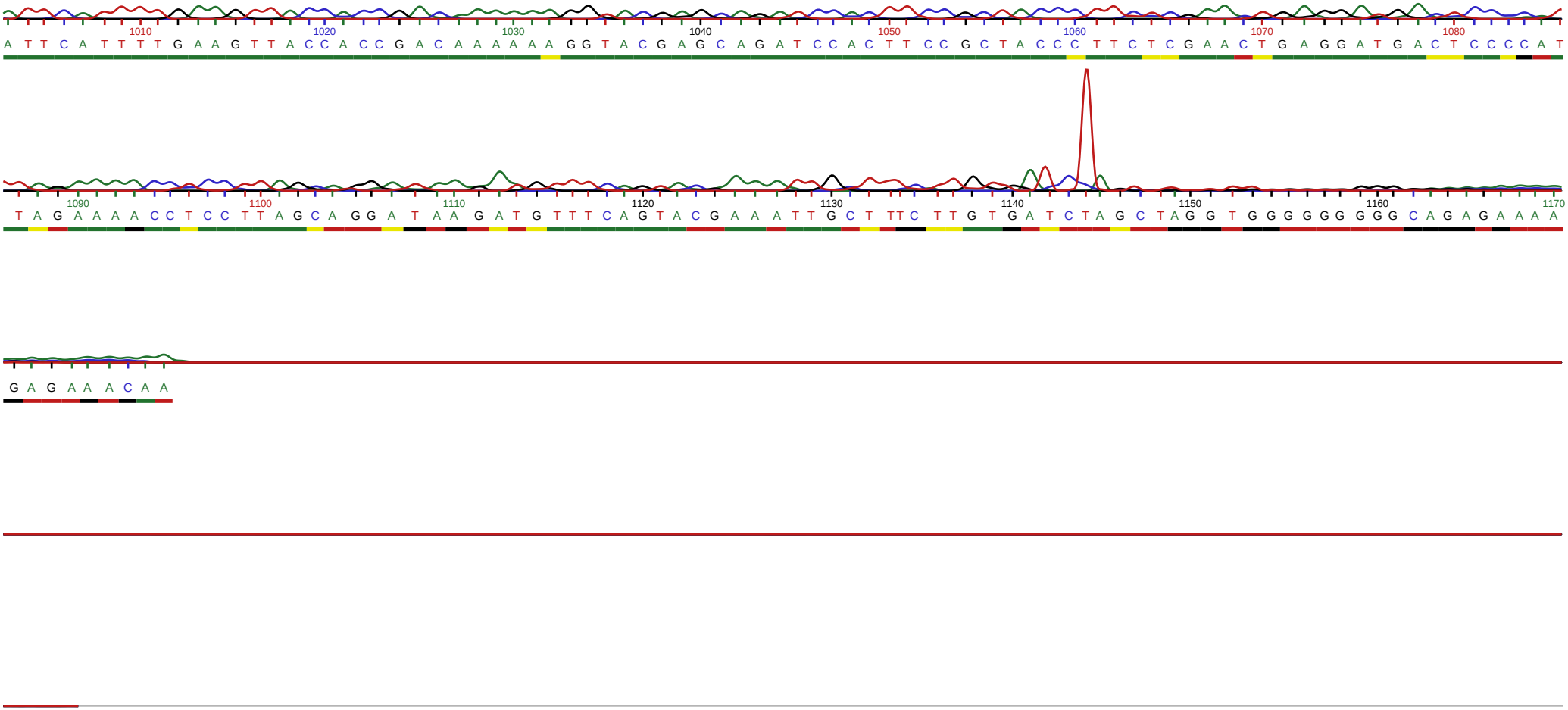
Sequence: 59746875

Samples:	16049	Quality:	0 - 9
Bases:	1180		10 - 19
Average spacing:	14.0		20 - 29
Average quality >=	10: 46, 20: 31, 30: 1080		>= 30

08.05.2024




510 520 530 540 550 560 570 580
G C A T C C T C T G G A G T T G C G G C G A G C C C A G T A C C G A T C G C C C A C A C C G G C T C G T A A G C A A C T A C T A C A T T A G T C C A G T C C T T C A C
590 600 610 620 630 640 650 660
C T C T T C G A G A A C G G C G T T C A G C T G G C G T T C T A C C A C A T C C C A A G G T C T T A C C G G C T T T T T T T T C T T C G A G A G T T T C G C C G A T A C A
670 680 690 700 710 720 730 740 750
G A G A A T G A C A C C T A C A C C C C T G C C C C A A T G C G A A C T T G G T T T T A T C A G C A A G G A A T T T A T C A T C T T C G T G A A A A T A G C T A C G G C
760 770 780 790 800 810 820 830
G C T C G C T A T G G C C C G A G G A T C A C C C A C T T T G C C C C T A C G T C T T T G A T C T G A T C A A C G G A G T T C T C C C C G G T A A A G G C G C C G C T T
840 850 860 870 880 890 900 910
G C C T T T A A A T A A G C A T T T T G C G C T C C T A C G G T C A C T T G T G G C T T T T T C A C T A G A C T G A C G C T A T A G T C C A G G T A G G T T G C A G G
920 930 940 950 960 970 980 990 1000
A G G G C A G A T A A C G A C C T C G A C G T T C T C G G G A A T A C T G G C T G T G T T C A G T C G T T C A A C G A T T T C T T T G A T A C T T T G C T T G C T A C C


1010
1020
1030
1040
1050
1060
1070
1080
A T T C A T T T T G A A G T T A C C A C C G A C A A A A A A G G T A C G A G C A G A T C C A C T T C C G C T A C C C T T C T C G A A C T G A G G A T G A C T C C C C A T
1090
1100
1110
1120
1130
1140
1150
1160
1170
T A G A A A A C C T C C T T A G C A G G A T A A G A T G T T T C A G T A C G A A A T T G C T TT C T T G T G A T C T A G C T A G G T G G G G G G G G G G C A G A G A A A A
G A G A A A C A A