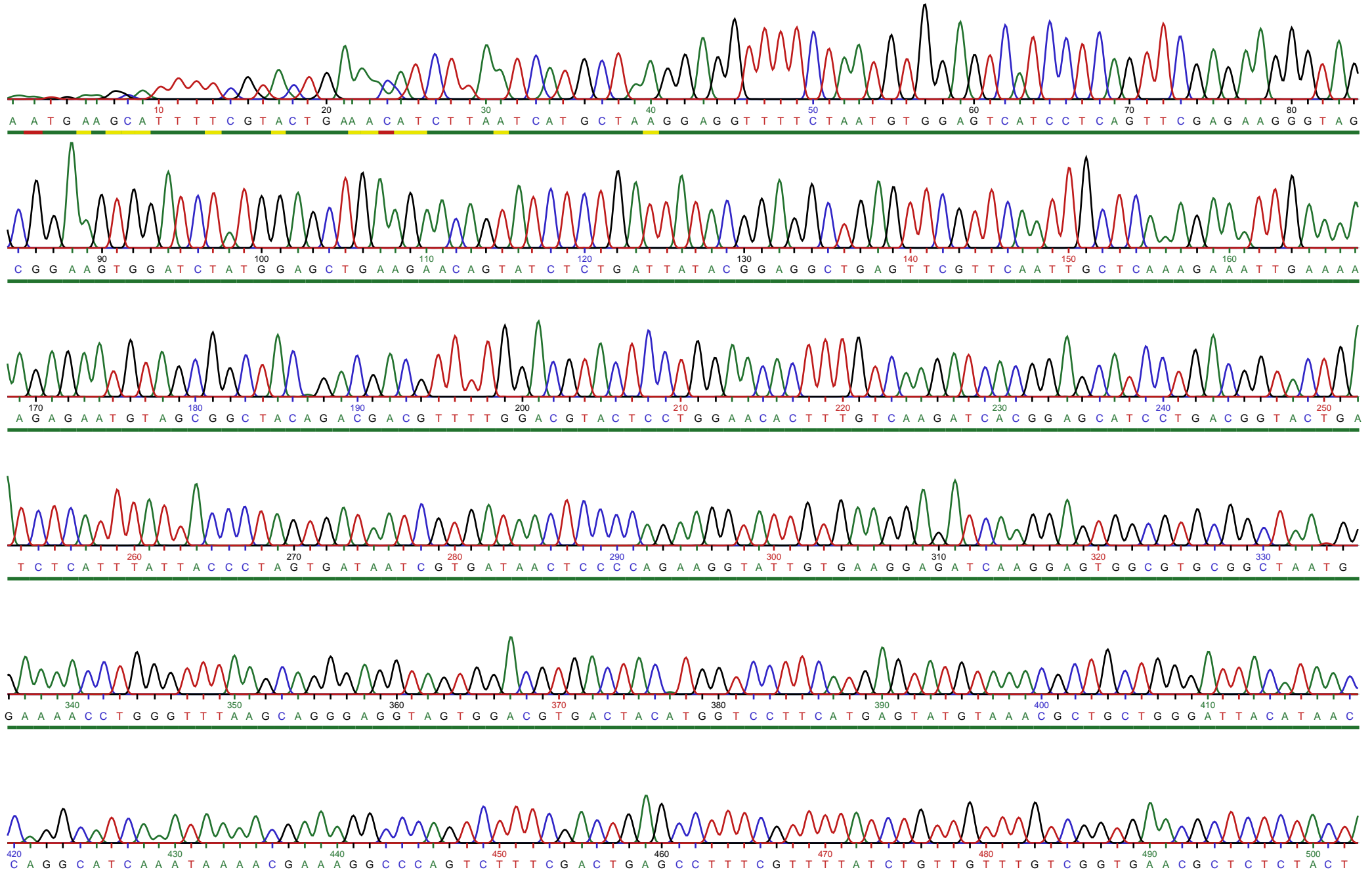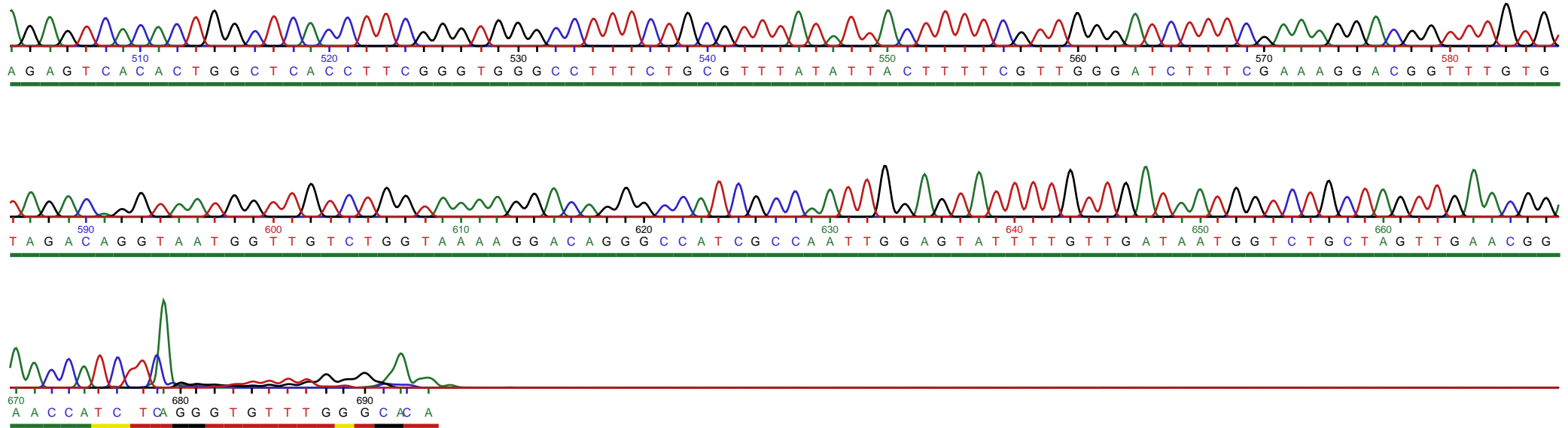Sequence: 59746950

Samples: 13836
Bases: 695
Average spacing: 20.0
Average quality >= 10: 16, 20: 15, 30: 660

Quality: 0 - 9
10 - 19
20 - 29
>= 30

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08.05.2024

A A T G A A G C A T T T T C G T A C T G A A A C A T C T T A A T C A T G C T A A G G A G G T T T T C T A A T G T G G A G T C A T C C T C A G T T C G A G A A G G G T A G

C G G A A G T G G A T C T A T G G A G C T G A A G A A C A G T A T C T C T G A T T A T A C G G A G G C T G A G T T C G T T C A A T T G C T C A A A G A A A T T G A A A A

A G A G A A T G T A G C G G C T A C A G A C G A C G T T T T G G A C G T A C T C C T G G A A C A C T T T G T C A A G A T C A C G G A G C A T C C T G A C G G T A C T G A

T C T C A T T T A T T A C C C T A G T G A T A A T C G T G A T A A C T C C C C A G A A G G T A T T G T G A A G G A G A T C A A G G A G T G G C G T G C G G C T A A T G

G A A A C C T G G G T T T A A G C A G G G A G G T A G T G G A C G T G A C T A C A T G G T C C T T C A T G A G T A T G T A A A C G C T G C T G G G A T T A C A T A A C

C A G G C A T C A A T A A A A C G A A A G G C C C A G T C T T T C G A C T G A G C C T T T C G T T T A T C T G T T G T T T G T C G G T G A A C G C T C T C T A C T

Sequence: 59746950

Samples:	13836
Bases:	695
Average spacing:	20.0
Average quality >=	10: 16, 20: 15, 30: 660

Quality: 0 - 9
10 - 19
20 - 29
>= 30

AGAGTCACACTGGCTCACCTTCGGGTGGGCCTTTCTGCGTTTATATTACTTTTCGTTGGGATCTTTCGAAAGGACGGTTGTG

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Sequence: 59746950

Samples: 13836   Quality: 0 - 9
Bases: 695   10 - 19
Average spacing: 20.0   20 - 29
Average quality >= 10: 16, 20: 15, 30: 660   >= 30

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