14236 1170

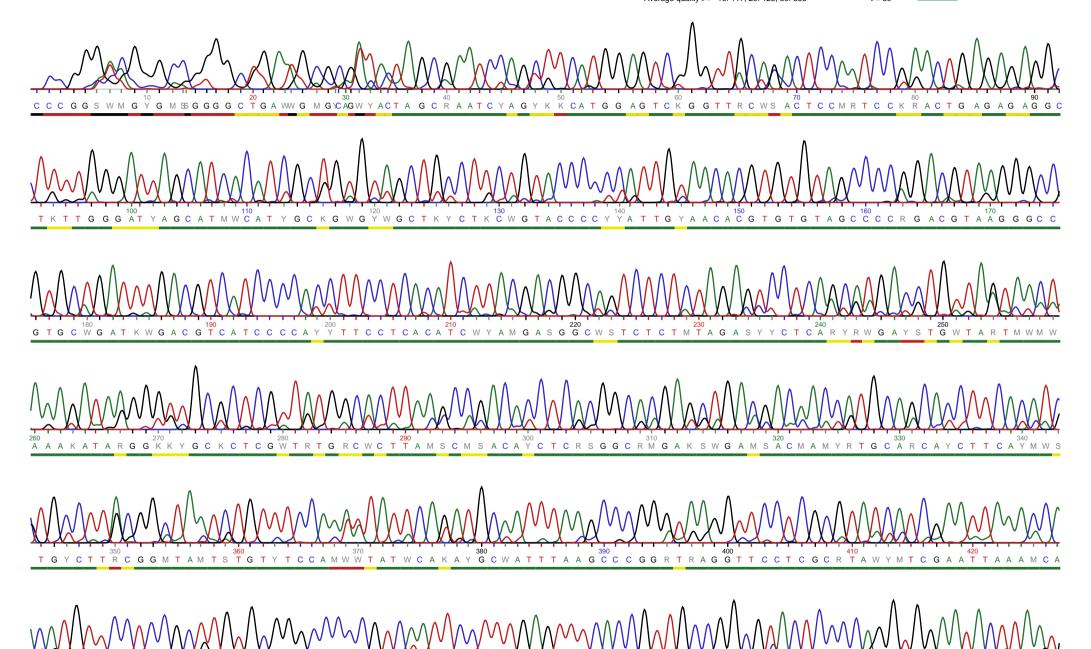
Average spacing: 13.0 Average quality >= 10: 117, 20: 125, 30: 866

Samples:

Bases:

Quality: 0 - 9 10 - 19 20 - 29





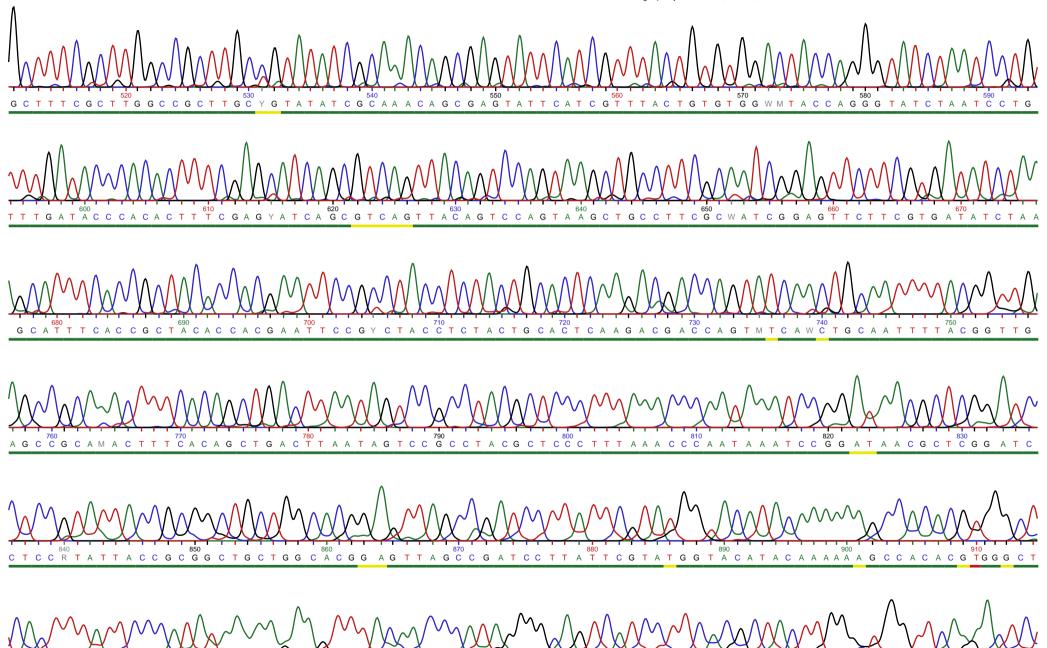
430 TGTTCCTCGGCKTGTGCGGGCCCCCGWCWATTCCTTTGAGTTTCACCGTTGCCGGCRYACTCCCAGGTGGAATACTTAAT

Samples: 14236 Bases: 1170

Average spacing: 13.0 Average quality >= 10: 117, 20: 125, 30: 866 Quality: 0 - 9 10 - 19 20 - 29 >= 30



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



CACT TTATT C C CATATAA A A GAAGT TTACAA C C CATR G G G G C A G T C A T C C T T C A C G C T A C T T G G G

14236 Quality: 0 - 9 1170 10 - 19

Average spacing: 13.0 20 - 29 Average quality >= 10: 117, 20: 125, 30: 866 >= 30

Samples:

Bases:



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