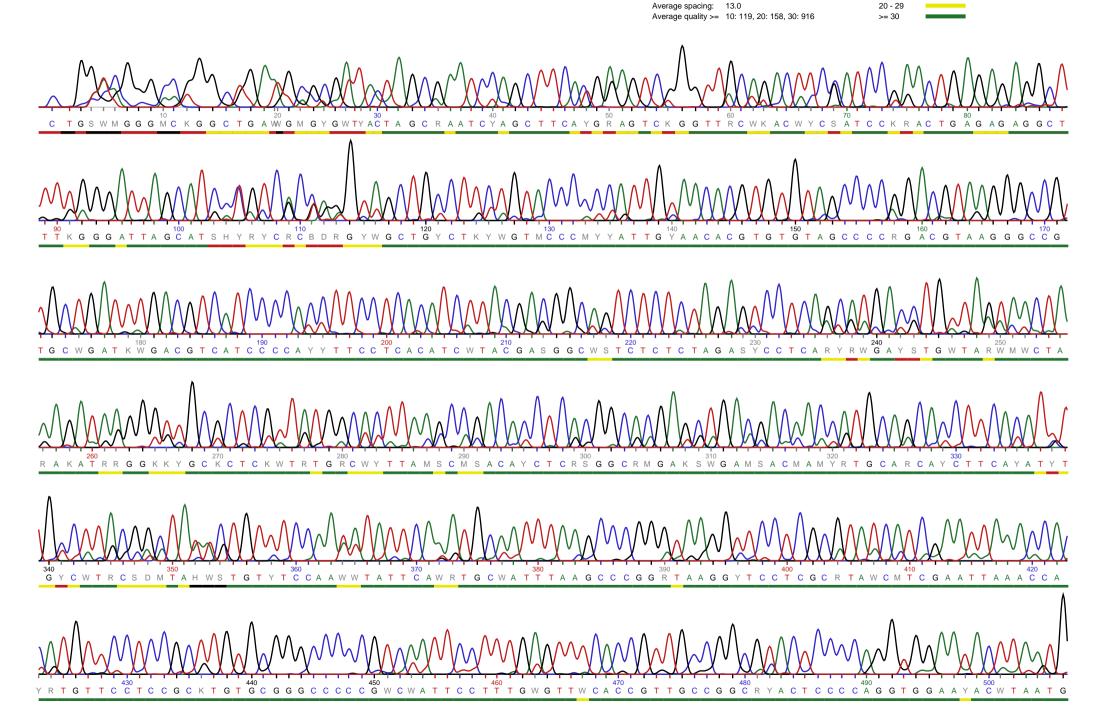
Quality: 0 - 9 10 - 19 20 - 29

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Samples:

Bases:

15190

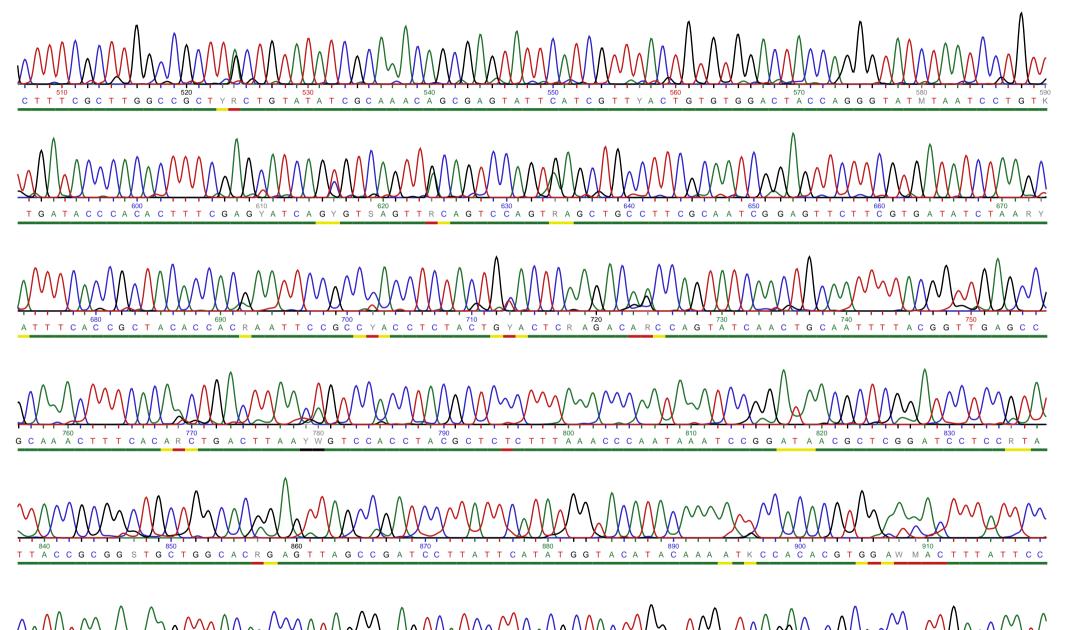
1265

Samples: 15190 Bases: 1265

Average spacing: 13.0 Average quality >= 10: 119, 20: 158, 30: 916 Quality: 0 - 9 10 - 19 20 - 29 >= 30



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920 CATATAAAA GAAGTTTACAACCCATAGGGCAGTCATCYTTCACGCTACTTGGCTGGTTCAGRCYYKCGCCCATTGACCAATA

Quality: 0 - 9

10 - 19 20 - 29 >= 30

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Samples:

