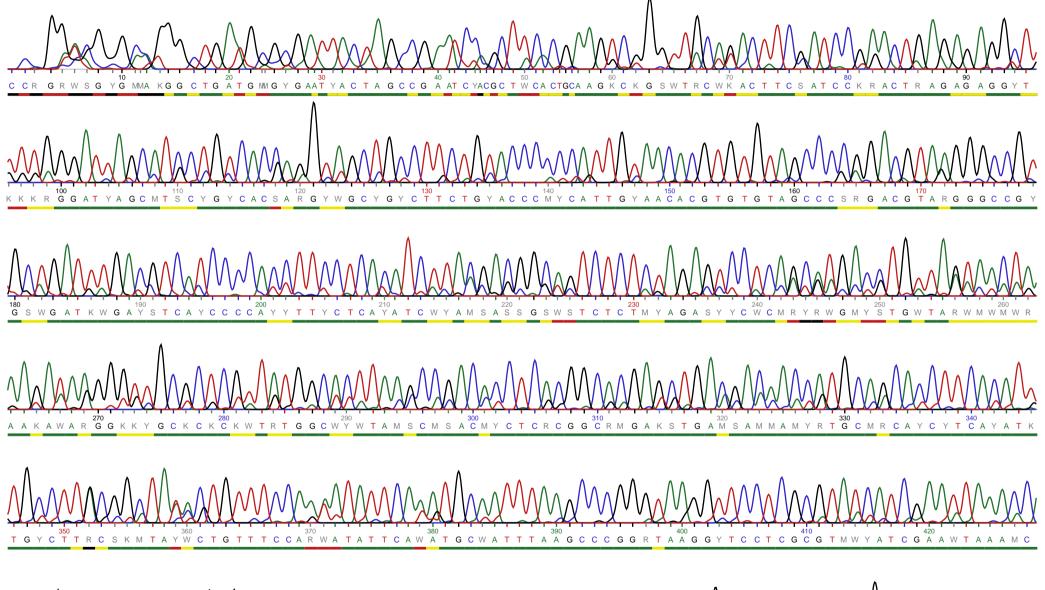
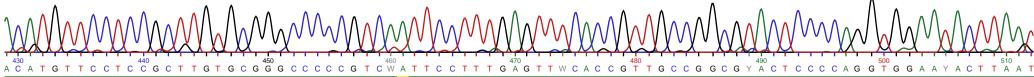
Samples: 15213 Bases: 1266

Average spacing: 13.0 Average quality >= 10: 90, 20: 156, 30: 988 Quality: 0 - 9 10 - 19 20 - 29









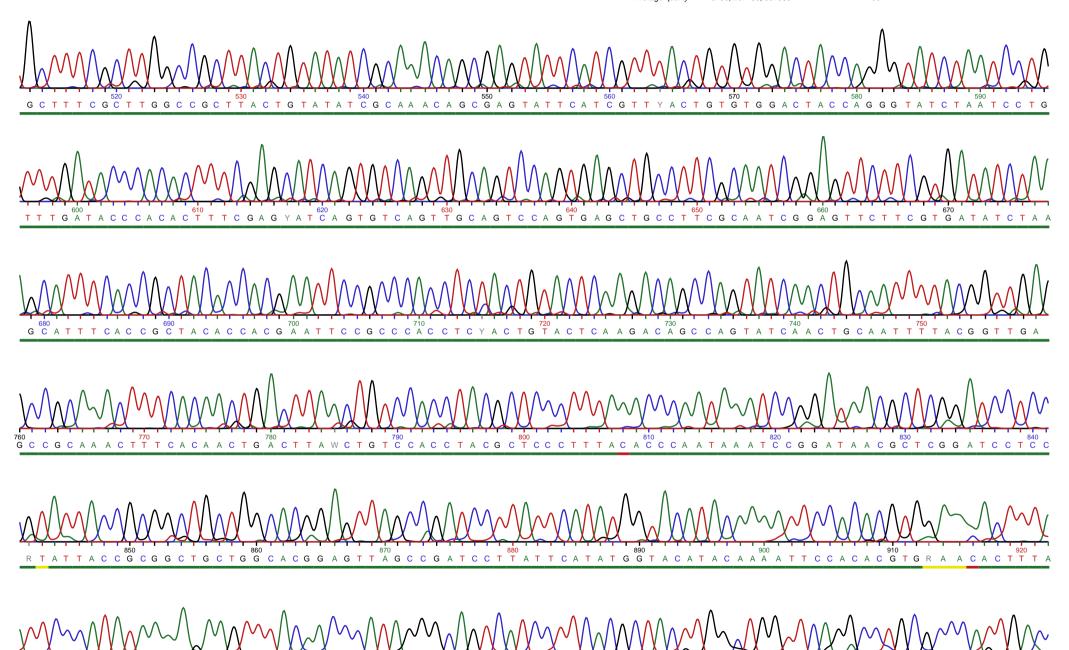
13.0 Average spacing: Average quality >= 10: 90, 20: 156, 30: 988 Quality: 0 - 9 10 - 19 20 - 29 >= 30





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TTCCCATATAAAAGAAGTTTACAACCCATAGGGCAGTCATCCTTCACGCTACKTGGCTTCAGRCCTGCGCCCAKTGAC

Samples: 15213 1266

Bases:

Average spacing: 13.0

10 - 19 20 - 29 Average quality >= 10: 90, 20: 156, 30: 988 >= 30

Quality: 0 - 9





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