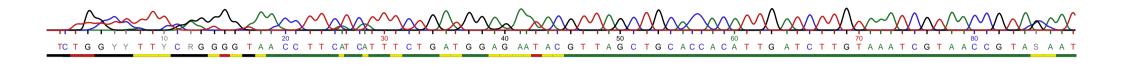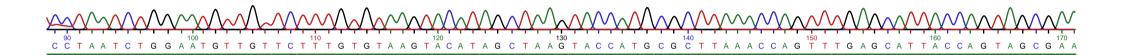Sequence: SEQ10013054_A1_F03_0
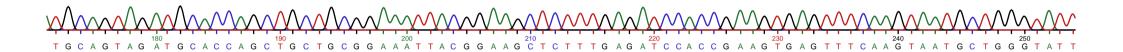
Samples: 15282
Bases: 444
Average spacing: 35.0
Average quality >= 10: 10, 20: 34, 30: 389

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

16.09.2022

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CCTAATCTGGAATGTTGTTCTTTGTGTAAGTACATAGCTAAGTACCATGCGCTTAAACCAGTTTGAGCATTACCAGTAGCGAA

TGCAGTAGATGCACCAGCTGCTGCGGAAATTACGGAAGCTCTTTGAGATCCACCGAAGTGAGTTTCAAGTAATGCTGGGTATT

CTTCGTATTGTTCTAATGCGTAGAATGCAACTTCAGAACCTACATCAAGAACAGTGTCCATGTTGTTAGGTGCTTCAGTTAAT

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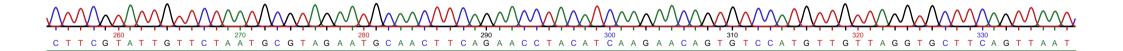
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Sequence: SEQ10013054_A1_F03_0

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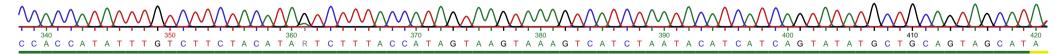Average quality >= 10: 10, 20: 34, 30: 389
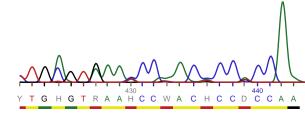
Quality: 0 - 9
10 - 19
20 - 29
>= 30

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Samples: 15282
Bases: 444
Average spacing: 35.0
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Quality: 0 - 9
10 - 19
20 - 29
>= 30