Sequence: SEQ10013680_B2_D11_0

Samples:	13881
Bases:	450
Average spacing:	31.0
Average quality >=	10: 14, 20: 36, 30: 369

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

GG GG SSYYYYY TTTY GRRGTG TAACCTTCATCATTTCTGATGGAGAATACGTTAGCTGCACCATTGATCTTGTAAATCGTATCCG

TAGAAACCTAATCTGGAATGTTGTTCTTTGTGTAAGTACATACCTAAGTACCATGCACTTAAACCAGTTTGAGCATTACCAGTAG

CGAATGCAGTGGAACAACCAGCTGCTGCTGAAATAACAGAAGCCCTTTGAGATCCACCGAAGTGAGTTTCAAGTAAAGCTGGGTA

TTCTTCGTATTGTTCTAATGCGTAGAATGAAACTTCAGAACCTACATCAAGAACAGTGTCCATGTTATTAGGTGCTTCAGTTAAT

CCACCATATTTGTCTTCTACGTAATCTTTACCATAGTAAGTGAAGTCATCTAATACATCATCAGTATATGCTGCAGTAGCATA
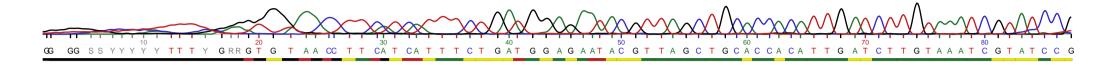
YTGMGTRAAMCCRACMCCWCCCAC

Samples: 13881
Bases: 450
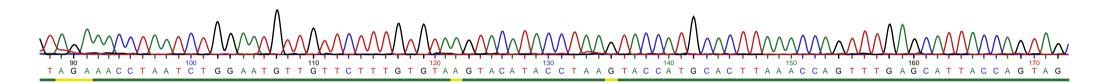Average spacing: 31.0
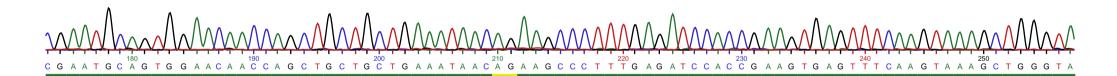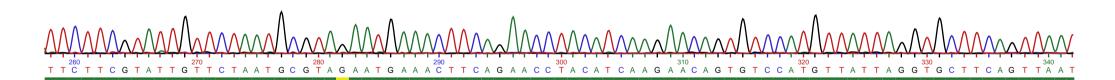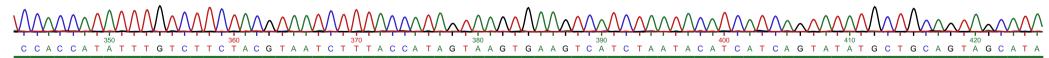Average quality >= 10: 14, 20: 36, 30: 369
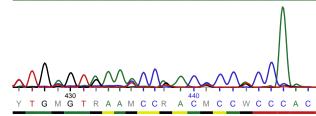
Quality: 0 - 9
10 - 19
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Samples: 13881
Bases: 450
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11.10.2022