Sequence: SEQ10012052_B1_E10_0
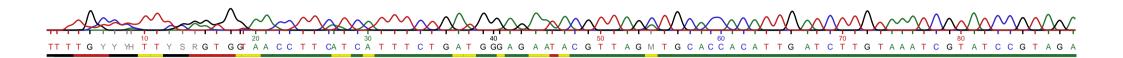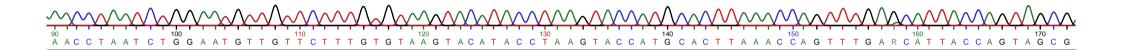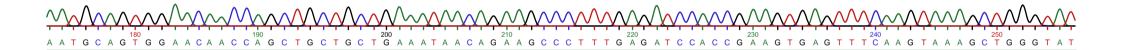
Samples: 16461
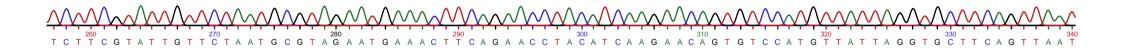Bases: 447
Average spacing: 37.0
Average quality >= 10: 17, 20: 27, 30: 393

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

TTTTGYYYHTTYSRGTGGTAACCTTCATCATTTCTGATGGGAGAATACGTTAGMTGCACCACATTGATCTTGTAAATCGTATCCGTAGA

AACCTAATCTGGAATGTTGTTCTTTGTGTAAGTACATACCTAAGTACCATGCACTTAAACCAGTTTGARCATTACCAGTAGCG

AATGCAGTGGAACAACCAGCTGCTGCTGAAATAACAGAAGCCCTTTGAGATCCACCGAAGTGAGTTTCAAGTAAAGCTGGGTAT

TCTTCGTATTGTTCTAATGCGTAGAATGAAACTTCAGAACCTACATCAAGAACAGTGTCCATGTTATTAGGTGCTTCAGTTAAT

CCACCATATTTGTCTTCTACGTAATCTTTACCATAGTAAGTGAAGTCATCTAATACATCATCAGTATATGCTGCAGTAGCATA

YTGHGTRAAHCCDACHCCDCCAA

Sequence: SEQ10012052_B1_E10_0

Samples: 16461 Quality: 0 - 9
Bases: 447 10 - 19
Average spacing: 37.0 20 - 29
Average quality >= 10: 17, 20: 27, 30: 393 >= 30

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Sequence: SEQ10012052_B1_E10_0

Samples: 16461
Bases: 447
Average spacing: 37.0
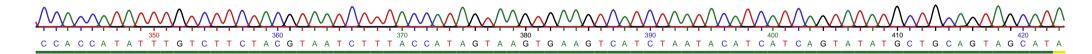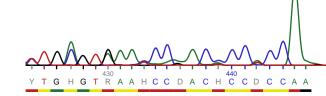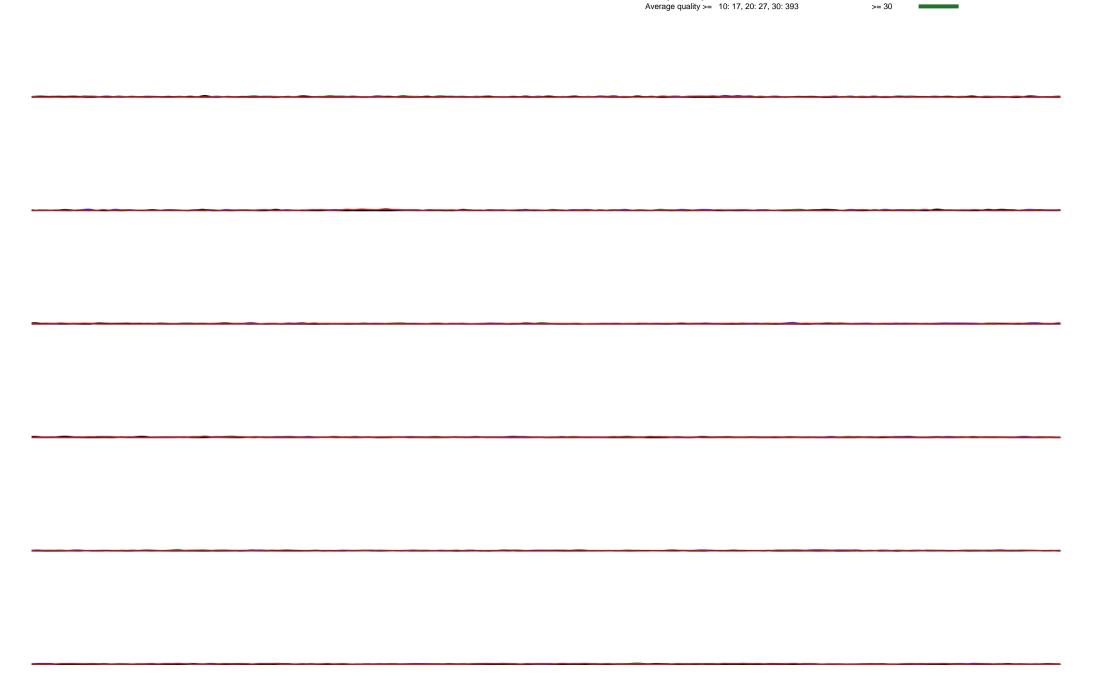Average quality >= 10: 17, 20: 27, 30: 393

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

09.08.2022