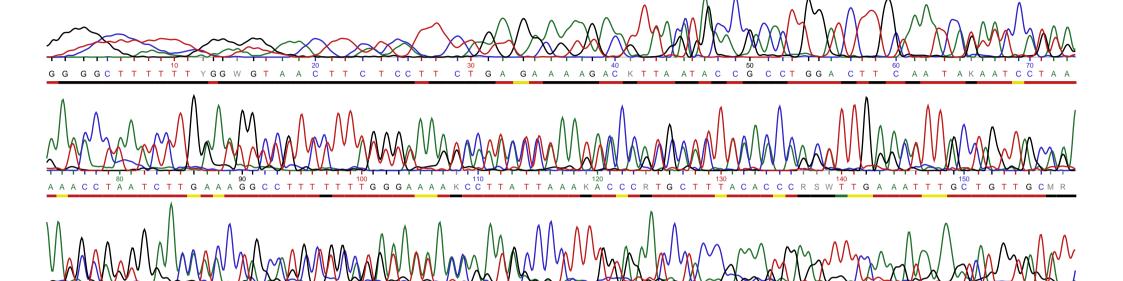
Samples: 16302 Bases: 428

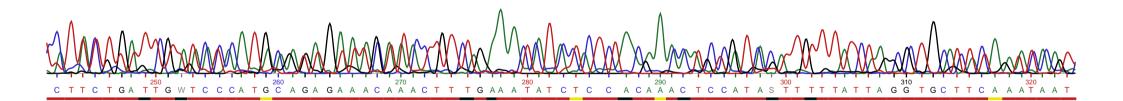
Average spacing: 39.0 Average quality >= 10: 289, 20: 26, 30: 1 Quality: 0 - 9 10 - 19 20 - 29 >= 30



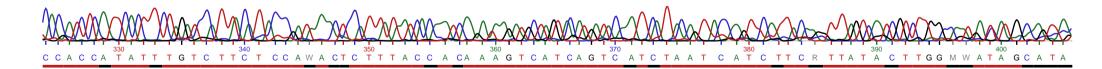
Page: 1/3

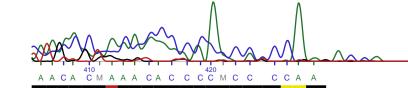
25.06.2022





TGTGGAACATCCTGCATGCTGAAATAAMCAAAGCCCTTTTTGAAACCTCCGAATTTTTCAAGTAARCTG





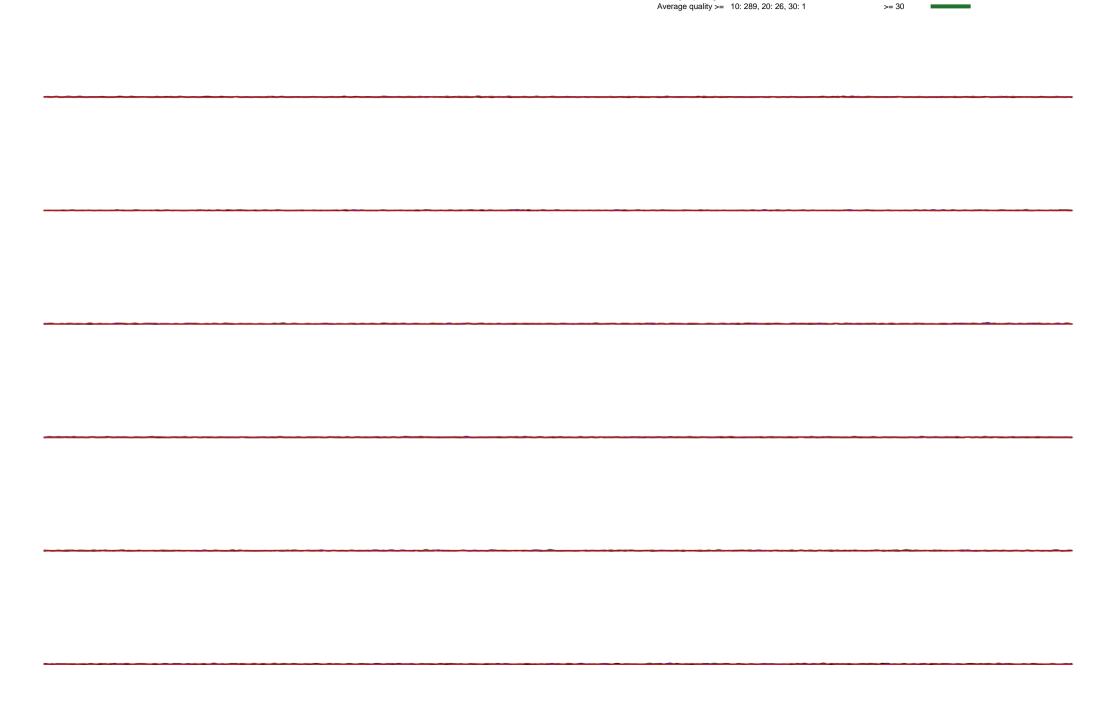
Sequence: SEQ10010919\_B2\_C06\_1

 Samples:
 16302

 Bases:
 428

 Average spacing:
 39.0

Quality: 0 - 9 10 - 19 20 - 29 >= 30 Page: 2 / 3 25.06.2022



Sequence: SEQ10010919\_B2\_C06\_1

Samples: 16302 Bases: 428 Average spacing: 39.0

Average quality >= 10: 289, 20: 26, 30: 1

Quality: 0 - 9 10 - 19 20 - 29 >= 30 Page: 3 / 3 25.06.2022

