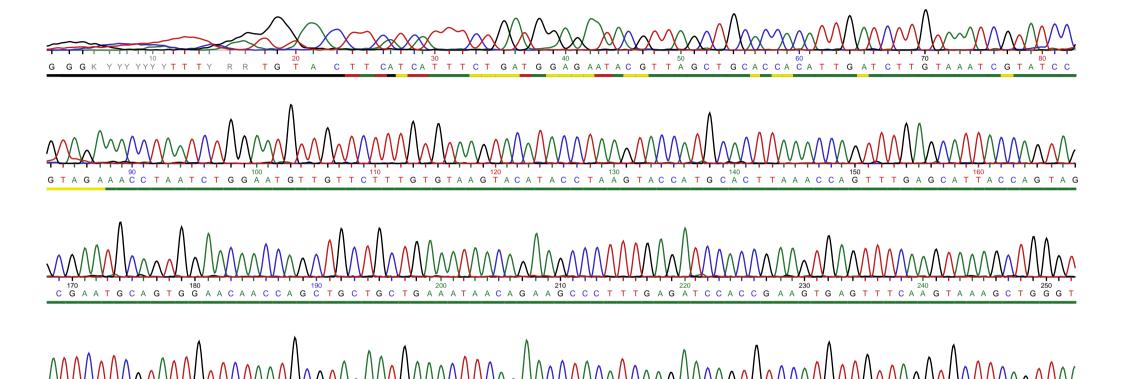
20 - 29





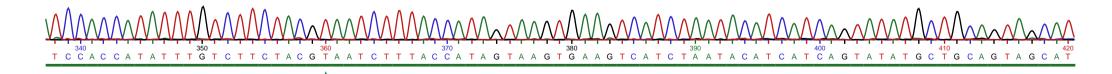
Samples:

Average spacing:

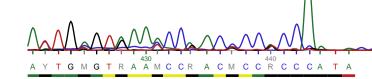
447

32.0

Average quality >= 10: 8, 20: 29, 30: 374



ATTCTTCGTATTGTTCTAATGCGTAGAATGAAACTTCAGAACCTACATCAAGAACAGTGTCATGTTGTTAGGTGCTTCAG



Sequence: SEQ10013680\_B2\_G10\_0

Samples: 13899 Bases: 447 Average spacing: 32.0

Average quality >= 10: 8, 20: 29, 30: 374

Quality: 0 - 9 10 - 19 20 - 29 >= 30 Page: 2 / 3 11.10.2022

Sequence: SEQ10013680\_B2\_G10\_0

Samples: 13899 Bases: 447 Average spacing: 32.0

Average quality >= 10: 8, 20: 29, 30: 374

Quality: 0 - 9 10 - 19 20 - 29 >= 30 Page: 3 / 3 11.10.2022