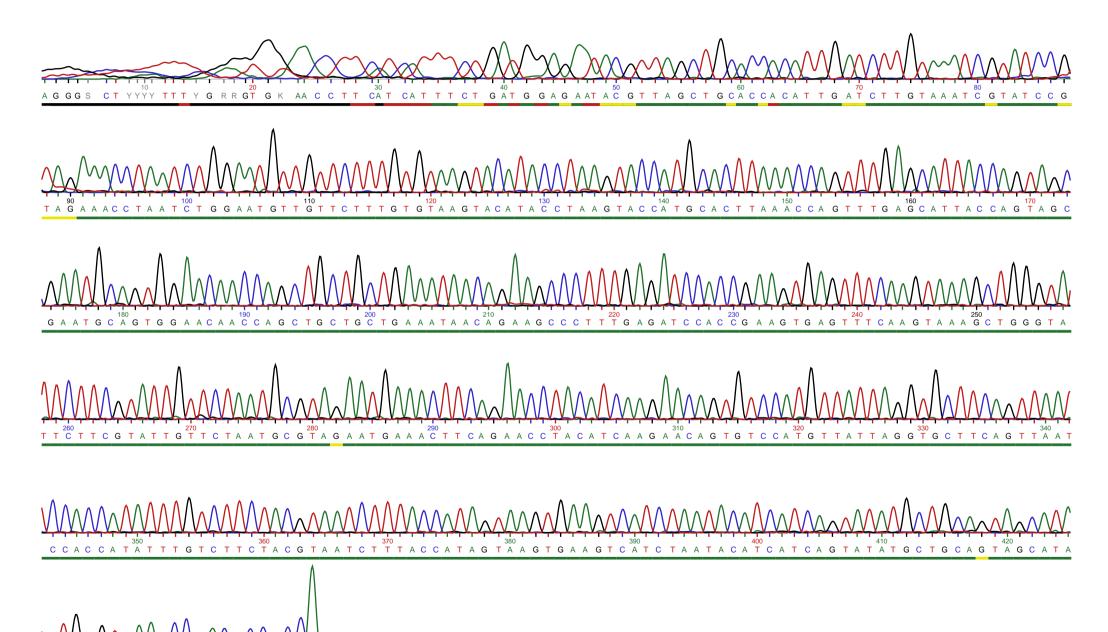
AGTRAAMCAA

ACMCCRCCAAC

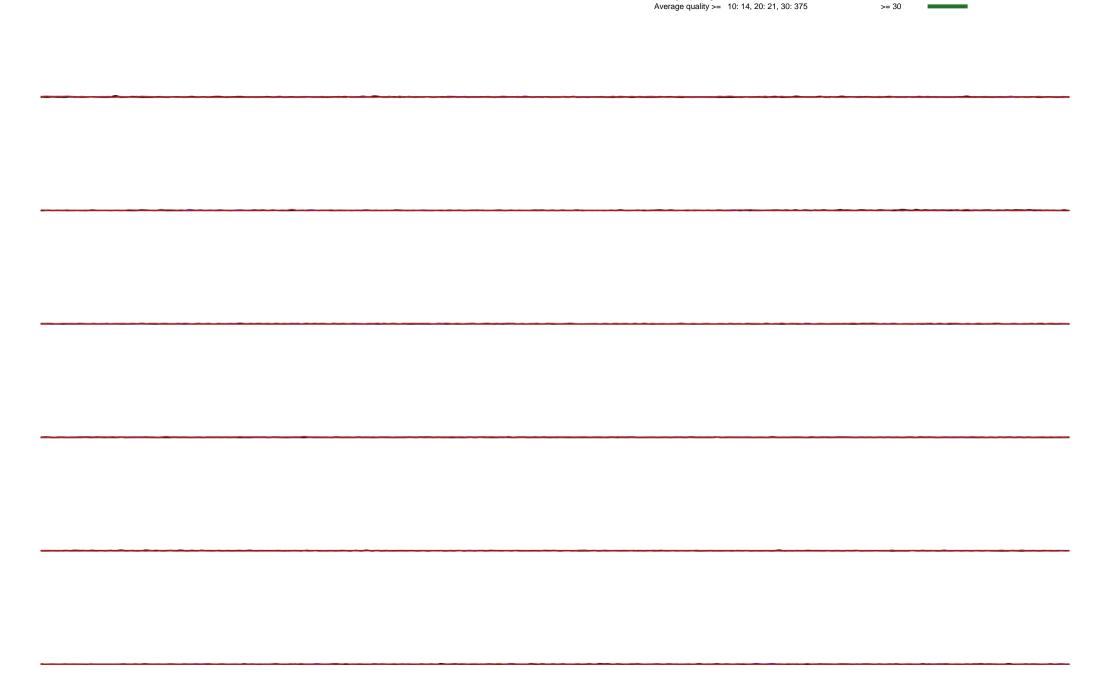
20 - 29





Sequence: SEQ10013680\_B2\_C05\_0

Samples: 14703 Bases: 450 Average spacing: 33.0 Quality: 0 - 9 10 - 19 20 - 29 >= 30 Page: 2 / 3 11.10.2022



Sequence: SEQ10013680\_B2\_C05\_0

Samples: 14703 Bases: 450 Average spacing: 33.0

Average quality >= 10: 14, 20: 21, 30: 375

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

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