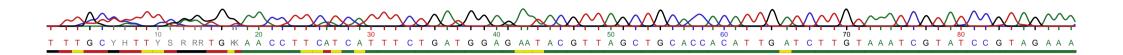
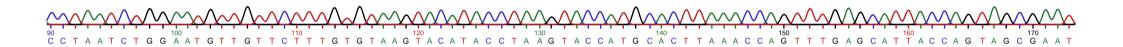
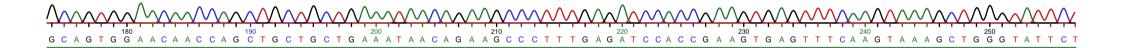
Samples: 12679
Bases: 444
Average spacing: 29.0
Average quality >= 10: 22, 20: 23. 30: 394

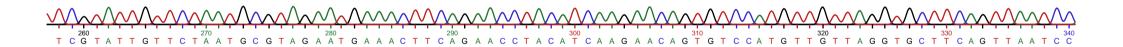
Quality: 0 - 9 10 - 19 20 - 29 >= 30

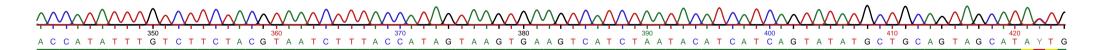


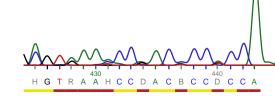












Sequence: SEQ10013680_B2_A06_0

Samples: 12679 Bases: 444 Average spacing: 29.0

Average quality >= 10: 22, 20: 23, 30: 394

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

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Sequence: SEQ10013680_B2_A06_0

 Samples:
 12679

 Bases:
 444

 Average spacing:
 29.0

 Average quality >=
 10: 22, 20: 23, 30: 394

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

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