Sequence: SEQ10013054_A1_C08_0

Samples:	18652	
Bases:	446	
Average spacing:	42.0	
Average quality >=	10: 17, 20: 30, 30: 389	

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

16.09.2022

T C T K G C Y T T T Y C R G G G T T A A C C T T C W T C A T T T C T G A T G G A G A A T A C G T T A G C T G C A C C A C A T T G A T C T T G T A A A T C G T A A C C G T A S A A

T C C T A A T C T G G A A T G T T G T T C T T T G T G T A A G T A C A T A G C T A A G T A C C A T G C G C T T A A A C C A G T T T G A G C A T T A C C A G T A G C G A A

T G C A G T A G A T G C A C C A G C T G C T G C G G A A A T T A C G G A A G C T C T T T G A G A T C C A C C G A A G T G A G T T T C A A G T A A T G C T G G G T A T T

C T T C G T A T T G T T C T A A T G C G T A G A A T G C A A C T T C A G A A C C T A C A T C A A G A A C A G T G T C C A T G T T G T T A G G T G C T T C A G T T A A T C

C A C C A T A T T T G T C T T C T A C A T A R T C T T T A C C A T A G T A A G T A A A G T C A T C T A A T A C A T C A T C A G T A T A T G C T G C A G T A G C A T A Y

T G H G T R A A H C C R A A C H C C D C C A A

Sequence: SEQ10013054_A1_C08_0

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Bases: 446
Average spacing: 42.0
Average quality >= 10: 17, 20: 30, 30: 389

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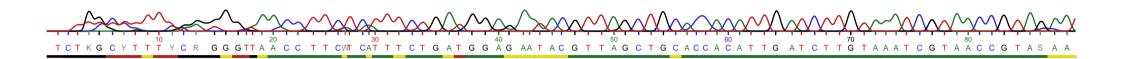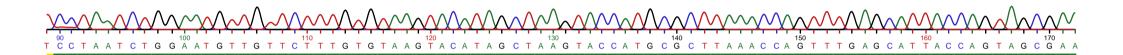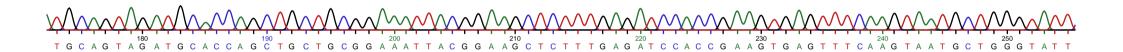10 - 19
20 - 29
>= 30

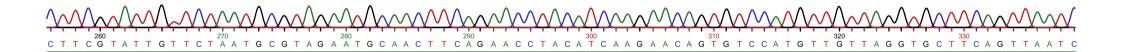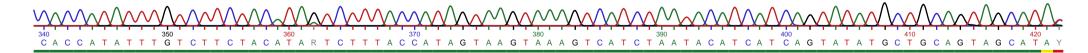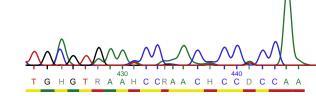Page: 2 / 4
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