Sequence: SEQ10010900_A2_B10_0

		Quality:	0 - 9	
Samples:	16300		10 - 19	
Bases:	417		20 - 29	
Average spacing:	40.0		>= 30	
Average quality >=	10: 264, 20: 3, 30: 0			

25.06.2022

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

A C G G T T G G T A G A A C T A G C T C A M A G C K G A G G T C C C G T A G A C C C C G G C G G G A A C A S A R C A A T T T A T C C C C A A T T T A C A G C

G A C C C T G C C G A G G C G A G G T G K A A A G T T A A G A A A T T T G C A C A A A C C C A G A G G T G G T G G C S A T M T C T A G A A C T C T G

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

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

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

Sequence: SEQ10010900_A2_B10_0

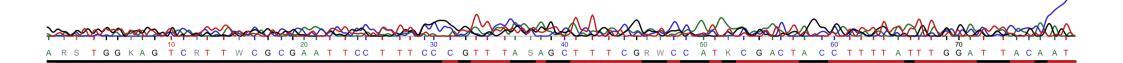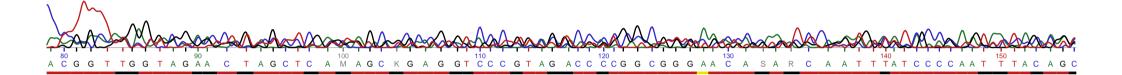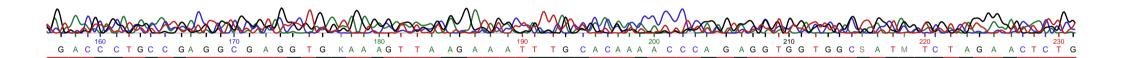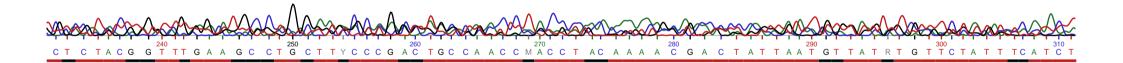|		Quality:	0 - 9	
Samples:	16300		10 - 19	
Bases:	417		20 - 29	
Average spacing:	40.0		>= 30	
Average quality >=	10: 264, 20: 3, 30: 0			

Sequence: SEQ10010900_A2_B10_0

Samples: 16300
Bases: 417
Average spacing: 40.0
Average quality >= 10: 264, 20: 3, 30: 0
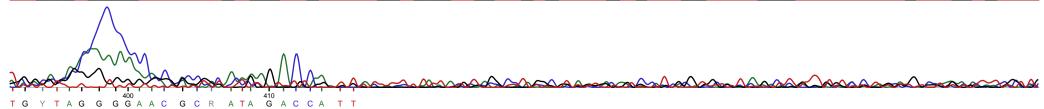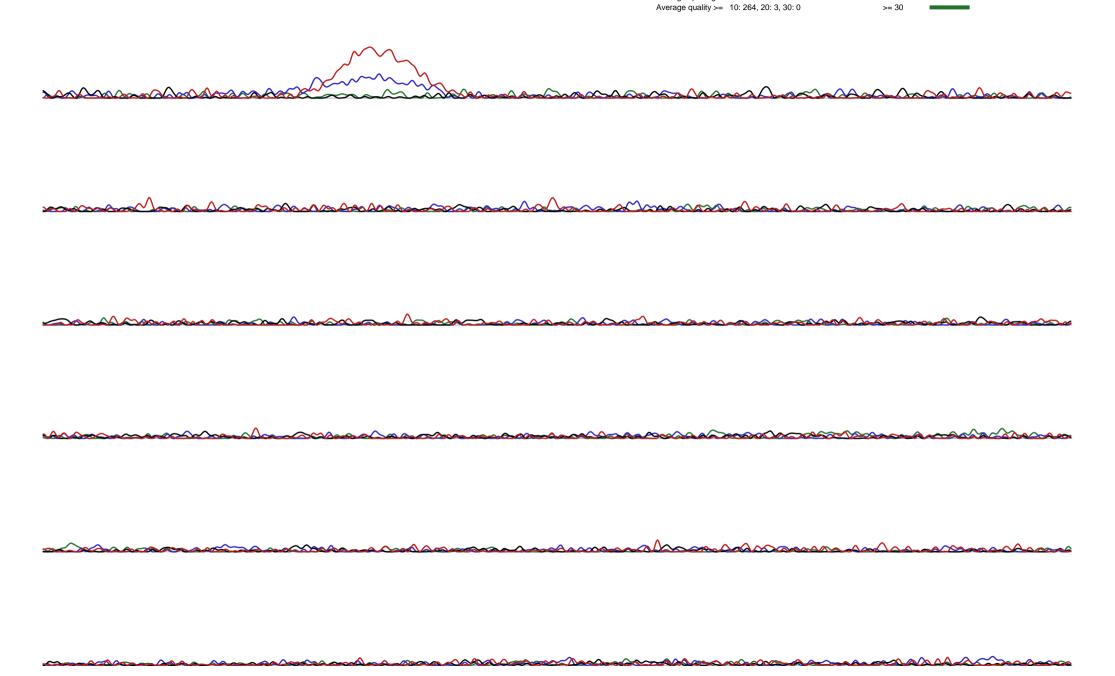
Quality: 0 - 9
10 - 19
20 - 29
>= 30

25.06.2022