Sequence: SEQ10010900_B1_D05_0

| | |
|---|---|
| Samples: | 16221 |
| Bases: | 420 |
| Average spacing: | 39.0 |
| Average quality >= | 10: 297, 20: 16, 30: 0 |

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

C A TTT G G G K G T M Y C T G G A T A C A G G C T C T G G C T C G A C C C C T C G A T C G T C C T T C T T C T T A C C T T C T T C C G G M T C T

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

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

G G T T T T T C T G G G T G A C T G G G C A G T A C C T T C G G G G G G C C C A K A T T T C G C G C C G W G A C G C G T C T A C T G C G C G G A C A T T A C

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

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

Sequence: SEQ10010900_B1_D05_0

| | | Quality: | 0 - 9 | |
|---|---|---|---|---|
| Samples: | 16221 | | 10 - 19 | |
| Bases: | 420 | | 20 - 29 | |
| Average spacing: | 39.0 | | >= 30 | |
| Average quality >= | 10: 297, 20: 16, 30: 0 | | | |

25.06.2022

Sequence: SEQ10010900_B1_D05_0

Samples: 16221
Bases: 420
Average spacing: 39.0
Average quality >= 10: 297, 20: 16, 30: 0

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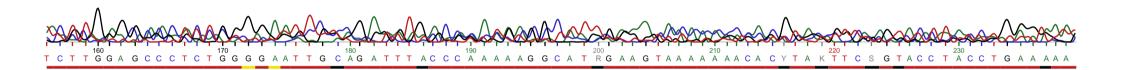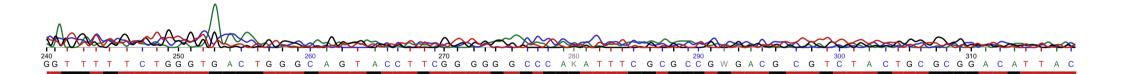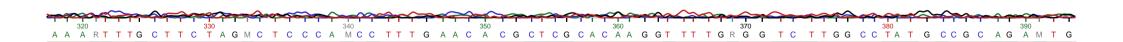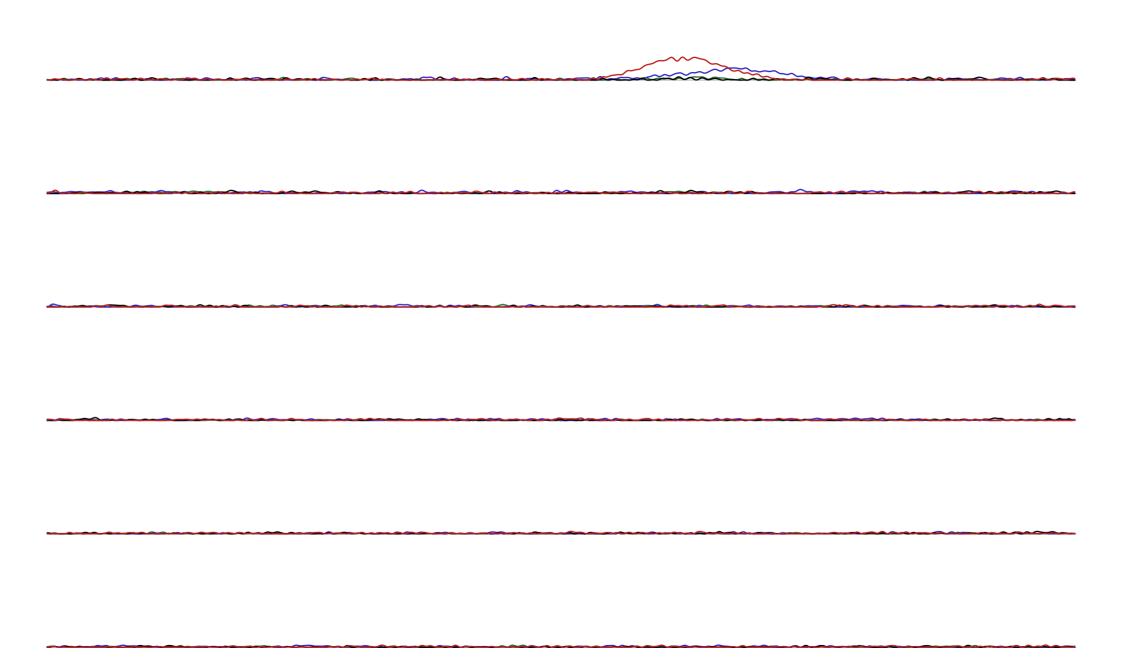10 - 19
20 - 29
>= 30

25.06.2022