Sequence: SEQ10013054_B1_B07_0

Samples:	21003
Bases:	453
Average spacing:	47.0
Average quality >=	10: 28, 20: 30, 30: 381

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

CCAKGSYTTTYGRRGGGTAACCTTTCATCATTTCTGATGGAGAATACGTTAGCTGCACCACATTGATCTTGTAAATCGTAACCGTAGA

ATCCTAATCTGGAATGTTGTTCTTTGTGTAAGTACATAGCTAAGTACCATGCGCTTAAACCAGTTTGAGCATTACCAGTAGCG

AATGCARTAGATGCACCAGCTGCTGCGGAAATTACGGAAGCTCTTTGAGATCCACCGAAGTGAGTTTCAAGTAATGCTGGGTA

TTCTTCGTATTGTTCTAATGCGTAGAATGCAACTTCAGAACCTACATCAAGAACAGTGTCCATGTTGTTAGGTGCTTCAGTTA

ATCCACCATATTTGTCTTCTACATARTCTTTACCATAGTAAGTAAAGTCATCTAATACATCATCAGTATATGCTGCAGTAGCAT

AYTGHGTRAAHCCRAAACDCCDCCAASAAC

Sequence: SEQ10013054_B1_B07_0

Samples: 21003
Bases: 453
Average spacing: 47.0
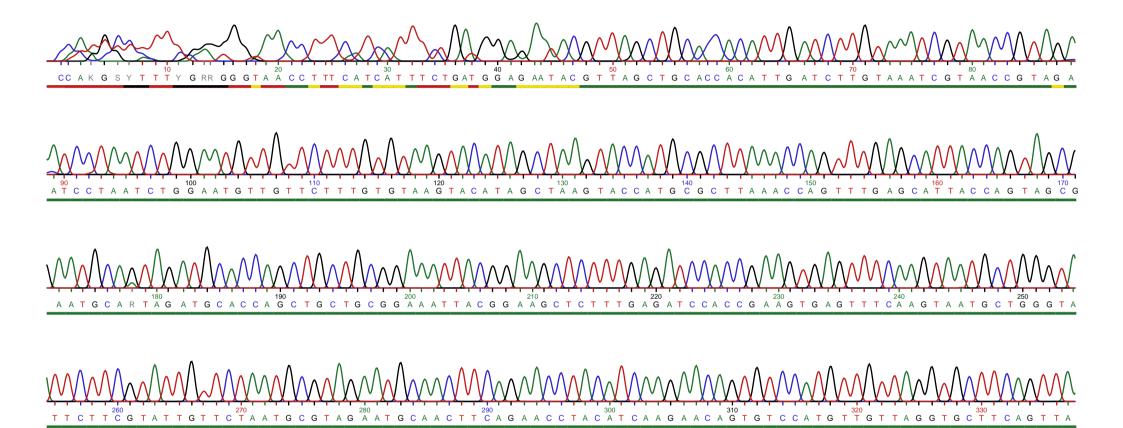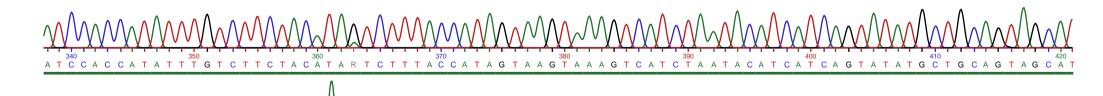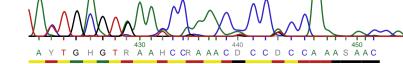Average quality >= 10: 28, 20: 30, 30: 381

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

16.09.2022

Samples: 21003
Bases: 453
Average spacing: 47.0
Average quality >= 10: 28, 20: 30, 30: 381

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

Sequence: SEQ10013054_B1_B07_0

Samples:	21003
Bases:	453
Average spacing:	47.0
Average quality >=	10: 28, 20: 30, 30: 381

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