Sequence: SEQ10012873_A1_H12_0

		Quality:	0 - 9	
Samples:	13721		10 - 19	
Bases:	446		20 - 29	
Average spacing:	31.0		>= 30	
Average quality >=	10: 23, 20: 31, 30: 385			

09.09.2022

TTTKCYYTTYCGRTGGTWAACCTTMMTCATTTCTGATGGAGATACGTTAGCTGCACCACATTGATCTTGTAAATCGTATCCGTAGAAAC

CTAATCTGGAATGTTGTTCTTTGTGTAAGTACATACCTAAGTACCATGCACTTAAACCAGTTTGAGCATTACCAGTAGCGAAT

GCAGTGGAACAACCAGCTGCTGCTGAAATAACAGAAGCCCTTTGAGATCCACCGAAGTGAGTTTCAAGTAAAGCTGGGTATTCT

TCGTATTGTTCTAATGCGTAGAATGAAACTTCAGAACCTACATCAAGAACAGTGTCCATGTTGTTAGGTGCTTCAGTTAATCC

ACCATATTTGTCTTCTACGTAATCTTTACCATAGTAAGTGAAGTCATCTAATACATCATCAGTATATGCTGCAGTAGCATAYT

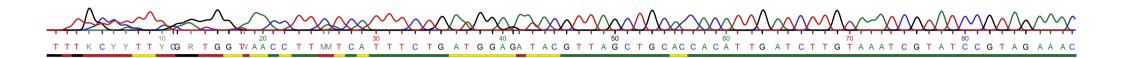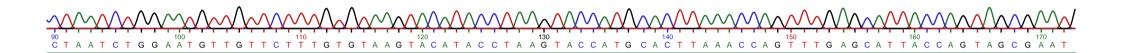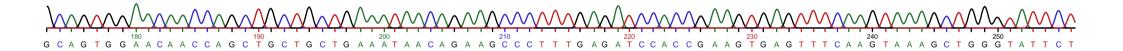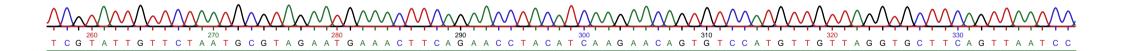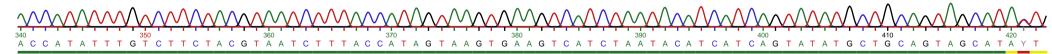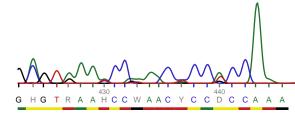GHGTRAAHCCWAACYCCDCCAAA

Sequence: SEQ10012873_A1_H12_0

Samples: 13721
Bases: 446
Average spacing: 31.0
Average quality >= 10: 23, 20: 31, 30: 385

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

Page: 2 / 3
09.09.2022

Samples: 13721
Bases: 446
Average spacing: 31.0
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Quality: 0 - 9
10 - 19
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>= 30