Sequence: SEQ10010900_A2_H09_0

| | | Quality: | 0 - 9 | |
|---|---|---|---|---|
| Samples: | 16301 | | 10 - 19 | |
| Bases: | 451 | | 20 - 29 | |
| Average spacing: | 37.0 | | >= 30 | |
| Average quality >= | 10: 51, 20: 124, 30: 207 | | | |

25.06.2022



GGGKYSYYYYTTTTYRRGTGTTAACTTCATCATTTCTGATGGAGATACGTAGCTGCACCATTGATCTTGTAAATCGTATCCGT

AGAAACCTAATCTGGAATGTTGTTCTTTGTGTAAGTACATACCTAAGTACCATGCACTTAAACCAGTTTGAGCATTACCAGTAGC

GAATGCAGTGGAACAACCAGCTGCTGCTGAAATAACAGAAGCCCTTTGAGATCCACCGAAGTGAGTTTCAAGTAAAGCTGGGKA
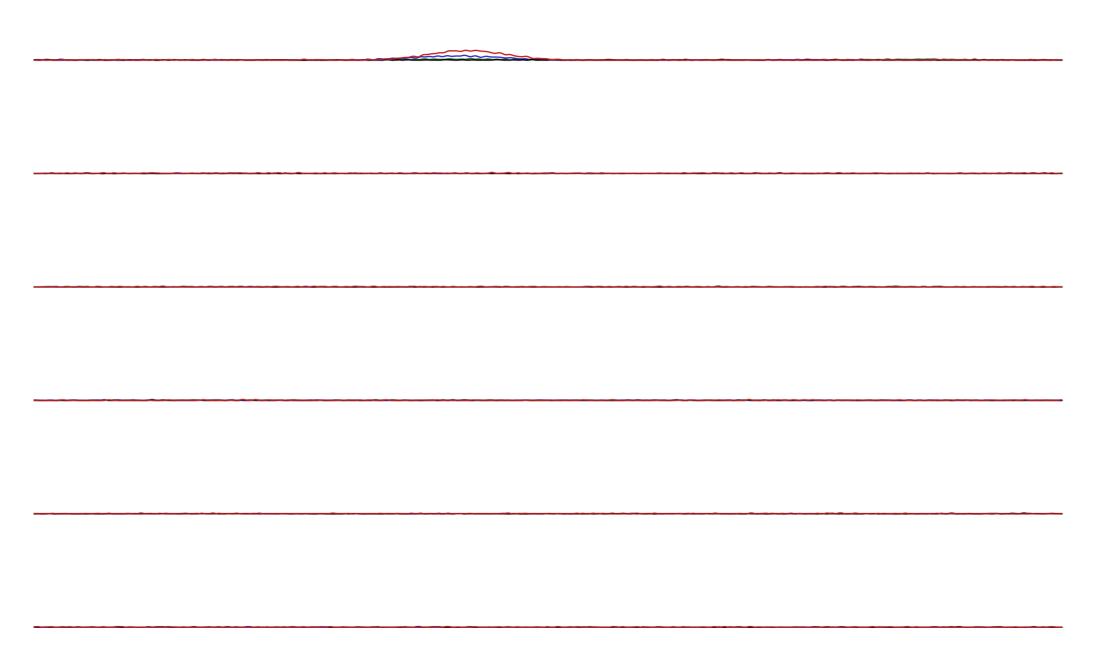
TTCTTCGTATTGTTCTAATGCGTAGAATGAAACTTCAGAACCTACATCAAGAACAGTGTTCCATGTTGTTAGGTGCTTCAGTTA

ATCCACCATTTGTCTTCTACGTAATCTTTACCATARTAAGTGAAGTCATCTAATACATCATCRGTATATGCTGCRGKARMA

TAYTGSSGYSCCACAAAACCCCSCCATA

Samples: 16301
Bases: 451
Average spacing: 37.0
Average quality >= 10: 51, 20: 124, 30: 207

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

Sequence: SEQ10010900_A2_H09_0

Samples: 16301
Bases: 451
Average spacing: 37.0
Average quality >= 10: 51, 20: 124, 30: 207

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

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