Sequence: SEQ10013680_B2_H05_0

Samples: 20936
Bases: 1720
Average spacing: 13.0
Average quality >= 10: 686, 20: 429, 30: 175

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

Page: 1 / 4
11.10.2022

T GGGGARWTTCCGGCGRGG RWWCCGGCMGG MMGGG K GC TRCAY GCG YRTRTAY AYRTM GGC AGT HAY CABA GY TYM GTGG AS KAKKKWYRSWSAYYMCW

CYSCRRWMYR ACWRRGKTWWKARGRATMSBBMYSYMCTSYSGYGKYRSASSYCSTYKBYHCMCCCMWTKGHAMSACGCTG

TGT ASCCCCRMGRCATTCAGGSGCCGTGSWGWTTTGACSTCATCCACACYTTCCTCMYATYTTACAKATGGCWSTCKTGW

KAGASTKCYCRRYMTCMHSWRTTAGMMWCKSACRWMARKRDKHGCGCTYGTCRBGSYMBTTRMSHSDAYYHMYC

RSRSCRCDMRCTGAMSAMDMYBRYGSSGSMYSTKCACHYYCTCTHWTSYWRWCWASAMSCRSTTCTCWCSAGAWSATT

CTAGTGCACWTTKAGTCCCSGGTARDGWTRTGTCGCGTMTYWWMRTAYTAATACCAMGTSWTYCTCCACTTGTGG

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CGGGTCGCCCCGCTCCAKATCTCWCTTTTMKATRTATMTCATSMGAGTGCKYGGMGACACTCACCCASMGTAGAGAATSACGTGTACAKCRWCATTA

TGCGTCTTTGCTKCGCTWGTSGKATATCGWCAAACAGCACMGMYATTCATMRTKTYACRCRTGTGKWAYASCAGGGG

CWMTMYAWCKSSTRTYTGATACCSACACTKYCCSMKMSTYAAKYGCTCMTCTTRCMGCYTAACWGCGTCTSC

CTTTCARCAMKCSWGWSTCTCWCTACGGWGATACTCTMAMSACAKTWTTCACGCGACATATCACACAGCKGATATCTCCCKGCGGT

RGCTMTCCATACGTGACACTTCWMAGCACATCTCAGATATCCAAACTAGMTAATCTTWTACGCGWRTGTMGCCCMGMARWAWC

TATTCYRGCARKGTAGTWCTYTWACAACATCACATTCTATCRWCTCCGSSTTTWAMAARCACCAYATTAWAAATSCCGSGAA

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11.10.2022

990 1000 1010 1020 1030 1040 1050 1060
T A A A A C R T M T K Y G R G A A W C C C T C C Y G R T A A T T Y A C G C G G C G G G T S T T G M C T G C G C C A C S S G G A A G T T K K A G C C C M G A A K C

1070 1080 1090 1100 1110 1120 1130 1140
Y C T G T T R T Y T T C A A T A A R W A S G T T A C Y R T T G G C A A A A A C T S G G K G G T T A T T G G C C A T Y A A C C C C M G A A A C T T T T T T A Y Y T T C C

1150 1160 1170 1180 1190 1200 1210 1220 1230
C Y T T T T A A T A A A A A A A G R A A R G G T T T T Y Y A A C G R A C C C M Y Y A A T G A A G G G G G G C T A A S T T C C S A T T C C C T T T T T C C A Y C G G W C

1240 1250 1260 1270 1280 1290 1300 1310 1320
T T T A A T K K T T T G G G G S T T T G G G T T T T T C C A G G G G G C C C C A A T T C C C G C C C C C C C C A T T T T T R G G A G A A C C A C A A W T A T T T C T C C C C

1330 1340 1350 1360 1370 1380 1390 1400 1410
W T T C W C A T C T T T G G G G C K S T G G G C C C C T C C C C C C T S S W G T W A A A A G G G A A A A T T T T T T T G G G G C G A A A C C C S G T K G G G G T T C C M W T C C

1420 1430 1440 1450 1460 1470 1480 1490 1500
C A A T Y C T T T T C C C C S A A G A T G G T G G G T G G G G G G G G A G G G A A A A Y A C C T C T T T T C C C T C T T T Y T T T A A A A A A A A A C A C C T C C

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Average spacing:	13.0
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A H T T A A W T C C M M C A T C T C C G G W T T W R G G A A M A A T A R A A G G G G G G G G G G G G G C G C C C C R G K T T T T A C C C C C C C C G S C C

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

A A A A A A A A A W A T C C C C T T T T T T A M T T C T C C T T S G G G G G G G A A R A M M T G G G D G G A A A A C T T T C H T R G