Sequence: SEQ10012873_A1_C04_0

| | |
|---|---|
| Samples: | 14566 |
| Bases: | 1226 |
| Average spacing: | 12.0 |
| Average quality >= | 10: 160, 20: 339, 30: 695 |

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



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

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

C R T G M T G A T K W G A C G T C A T C C C C A Y Y T T Y C T C M S W W Y T G W M K A S G G C W S Y C Y C T M T A G A G Y S C H C M W Y W W D R Y G B K G K M M W M

W M D R R R S R K G R G K Y G Y K C K C K Y K G Y R D G D Y W Y W H M M C M M B A Y C Y C W M G A C R M G M K S W S W S S A C A M M H R Y R Y R C C A Y S T S T Y T
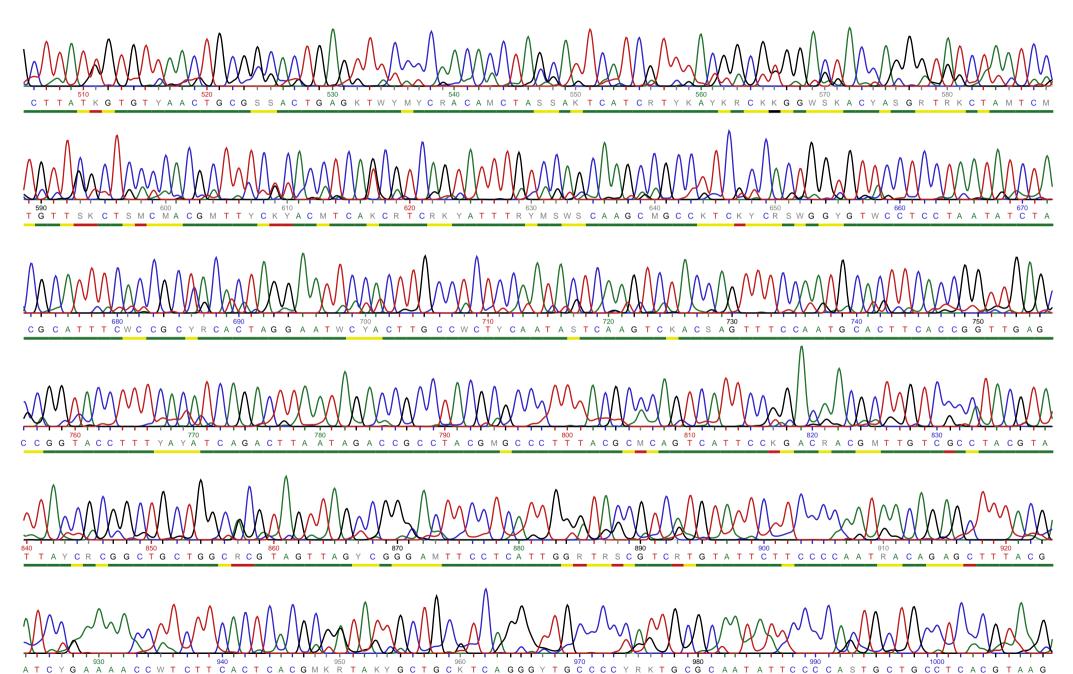
M T S T G Y K C C H D R A R G A W A M T Y Y S T A T C T M T A K K M T Y S W C A K A S K A T G T M R A S A C S T G G W R A G G T T C T Y C S C K W T G M T Y C R A A T T

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

Sequence: SEQ10012873_A1_C04_0

Samples: 14566
Bases: 1226
Average spacing: 12.0
Average quality >= 10: 160, 20: 339, 30: 695

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

CTTATKGTGTYAACTGCGSSACTGAGKTWYMYCRACAMCTASSAKTCATCRTYKAYKRCKKGGWSKACYASGRTRKCTAMTCM

TGTTSKCTSMCMACGMTTYCKYACMTCAKCRTCRKYATTTRYMSWSCAAGCMGCCKTCKYCRSWGGYGTWCCTCCTAATATCTA

CGCATTTCWCCGCYRCACTAGGAATWCYACTTGCCWCTYCAATASTCAAGTCKACSAGTTTCCAATGCACTTCACCGGTTGAG

CCGGTACCTTTYAYATCAGACTTAATAGACCGCCTACGMGCCCTTTACGCMCAGTCATTCCKGACRACGMTTGTCGCCTACGTA

TTAYCRCGGCTGCTGGCRCGTAGTTAGYCGGGAMTTCCTCATTGGRTRSCGTCRTGTATTCTTCCCCAATRACAGAGCTTTACG

ATCYGAAAACCWTCTTCACTCACGMKRTAKYGCTGCKTCAGGGYTGCCCCYRKTGCGCAATATTCCCCASTGCTGCCTCACGTAAG

Sequence: SEQ10012873_A1_C04_0

| Samples: | 14566 |
| Bases: | 1226 |
| Average spacing: | 12.0 |
| Average quality >= | 10: 160, 20: 339, 30: 695 |

Quality: 0 - 9 (black), 10 - 19 (red), 20 - 29 (yellow), >= 30 (green)

09.09.2022



AGTCTGGACCGYGTCTCAGTTACWGTGWGACCGTTCGCCCTCTCAGAGCGGRTACCCATCGTCGCCKWGGTGGGCTGWTATCTCAC

AWSTAGCTAATGWGACSCGGSWYCATCCTATGGCACCGGAYTTTTATGATCTTGGCATGCKACAAAACATAAAWWWRGSYTTTM

YYCMAGTTTCCGRRRTTWWTYCTTTYCWRRGGSWGTTMCCCABGC