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

C G T A G A A T C C T A A T C T G G A A T G T T G T T C T T T G T G T A A G T A C A T A G C T A A G T A C C A T G C G C T T A A A C C A G T T T G A G C A T T A C C A G T

A G C G A A T G C A G T A G A T G C A C C A G C T G C T G C G G A A A T T A C G G A A G C T C T T T G A G A T C C A C C G A A G T G A G T T T C A A G T A A T G C T G G

G T A T T C T T C G T A T T G T T C T A A T G C G T A G A A T G C A A C T T C A G A A C C T A C A T C A A G A A C A G T G T C C A T G T T G T T A G G T G C T T C A G T

T A A T C C A C C A T A T T T G T C T T C T A C A T A G T C T T T A C C A T A G T A A G T A A A G T C A T C T A A T A C A T C A T C A G T A T A T G C T G C A G T A G C

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

Samples: 16303
Bases: 447
Average spacing: 37.0
Average quality >= 10: 14, 20: 33, 30: 358

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

Sequence: SEQ10013052_B2_G09_0

Samples:	16303
Bases:	447
Average spacing:	37.0
Average quality >=	10: 14, 20: 33, 30: 358

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

16.09.2022