Sequence: SEQ10013680_B2_G10_0

Samples:	13899
Bases:	447
Average spacing:	32.0
Average quality >=	10: 8, 20: 29, 30: 374

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

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

G T A G A A A 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 C C T A A G T A C C A T G C A 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 G G A A C A A C C A G C T G C T G C T G A A A T A A C A G A A G C C 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 A 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 A 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 G T A A T C T T T A C C A T A G T A A G T G 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 R   A C M   C C R   C C C A T A

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11.10.2022

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11.10.2022