Sequence: SEQ10013054_A1_H08_0

CCTGSWMGCGMMKGGCTGAAGMGYGWTTACTAGCGAATCCARCTTCRTGRAGTCKGGTTRCWSACTYCRRTCCKAACTRCGASMK

GYTTAGRGATYMGCWBCBBGTCACSHSGTDGCTGYCYKCWGTACCSSYYRTTGYAACACGTGTGTMGCCCCGGRYGTAAGG

GCCGTGCTGATTTGAYGTCATCCCCRYYTTCCTCRCAYCWTASGRTGRCWSTCTCKMYAGAGYSCYSGKCTWCAMSHGDTRGM

AWSTGGCGATRGGGGKTGCKCTCGWTRTGRSWCTTMMCCMSACAYCTCRCGGCRMGAKSWGAMSACAAMYRTGCMRCAYCKTS

TATMKGYCSYRMRGSKAATWYSTGTYYMCWAKAMDYRCRMRMYAYWTWWRMMCSSGKRWARKGYYYCKCGYRTMWYMKMRWW

WWAMMMCRYRTKYYYCYSCKYKTGYGCGSSCCCSYSWMWWYYYYTKWGWKTYWCMKCKKYSSSSRYRYWCYCCMSRTGTGGRY

Sequence: SEQ10013054_A1_H08_0

Samples:	15627	
Bases:	1292	
Average spacing:	13.0	
Average quality >=	10: 439, 20: 379, 30: 248	

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

```
500        510       TAY TTTC KYKT GMR CCS TGG SWR TR TMK CRC MMA G MW CSA RTK MRY MK C TT WYW STG YGTG SWM YMM SRGG AT
W CWY WWR                                                                                                      580
```

```
M TMW WM TC STG T KMK MYM CACA CKC TCK MG CRY MWSH GYG WSW TAY MGW GCM TGG RRM KST GYY TK CRM T MKC GGKK YTY
590       600          610            620            630            640       650           660
```

```
C RW G AG ATM TMT RA AYA TTW CACC TCYAC TW CRM WTG TTCCGC MTCTCTM STS TM CTC WCG AGR GCG C SAR TT WCM DC
670       680          690          700            710          720           730
```

```
C RM GTG CWM SAK TK TAS TCTG CARY CTTTT MCCG YKRAA YTWG GYG TCCMG CYW ASS YGCC YTT WAAM CCM AWW WAAW YC
740        750       760         770           780          790         800           810
```

```
S GRR AAAS GYTSG MWY CCCCS GWW ATWM CSSS GGS YKSY GG MMS GRAW TTWAS CCRAAY GCTT WWTTC ATC RRGT WMS
820       830          840          850          860          870         880          890       900
```

```
T GSM AAAW YCS CMA CGCKK GG MMRAW C TATT YTYYCY TT RWM MAAAR RRKK TTYM CAS SSC YRTG GGAMM RWTYY TYCYY
910          920         930          940         950          960          970       980
```

Sequence: SEQ10013054_A1_H08_0

Samples:	15627
Bases:	1292
Average spacing:	13.0
Average quality >=	10: 439, 20: 379, 30: 248

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

16.09.2022

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

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

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

T T T T T T A A A A A A G A A A A A A A A G C G G G G G C C A C C C C C M C C M A C C A A Y Y Y G