Samples: 15733
Bases: 1314
Average spacing: 12

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

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

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

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

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

A T G A A T T T C C C A A C C C A T A T C G G T A A C C G G T T C T T C C A T G T G A A C C T G T T C A A C T T G C A G C A G G C T T G C A T C A T T G G T G C T G C
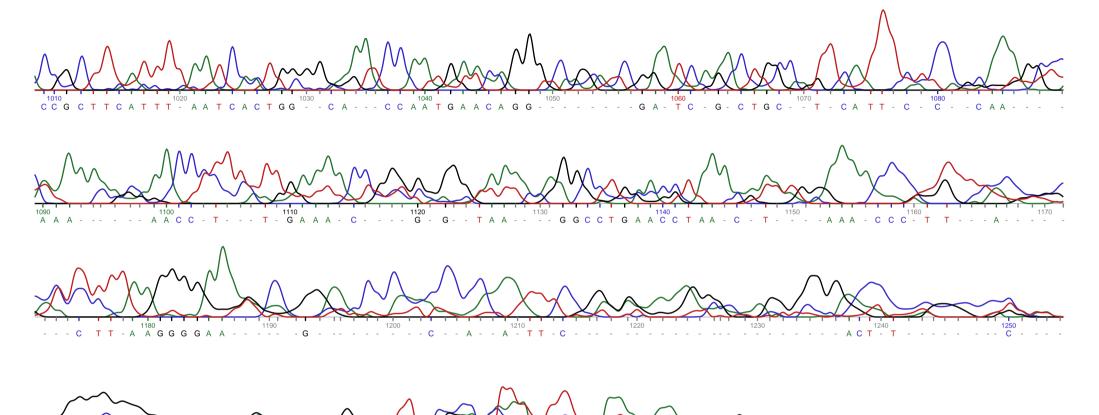
Samples: 15733
Bases: 1314
Average spacing: 12

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

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

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

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

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

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

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1010 CCG CTTCATTT - AATCACTGG - - CA - - - CCAATGAACAGG - - - - - - GA - TC - G - CTGC - - T - CATT - C - C - - CAA - - -

1090 AAA - - - - - - AACC - T - - T - GAAA - C - - - G - G - TAA - - - GGCCTGAACCTAA - C - T - - - AAA - CCC - TT - A - A - - -

- - - CTT - A AGGGGAA - - - - - G - - - - - - - C - A - A - TT - C - - - - - ACT - T - - - - - - - C - - - -

C - GC - - G - - - - A - - GA - - G - TT - - GT - - - - - - ATAATAACC - - - - - -