Samples: 14825
Bases: 687
Average spacing: 22

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

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

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

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

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

A C G C C A T A T C C G G T G C C C C A C T A T A T A A T G G T A C T A G - C A G T T T C C - A T C C C C C A A T C A T A A C C G G A A T T A C T A A A A A A - A
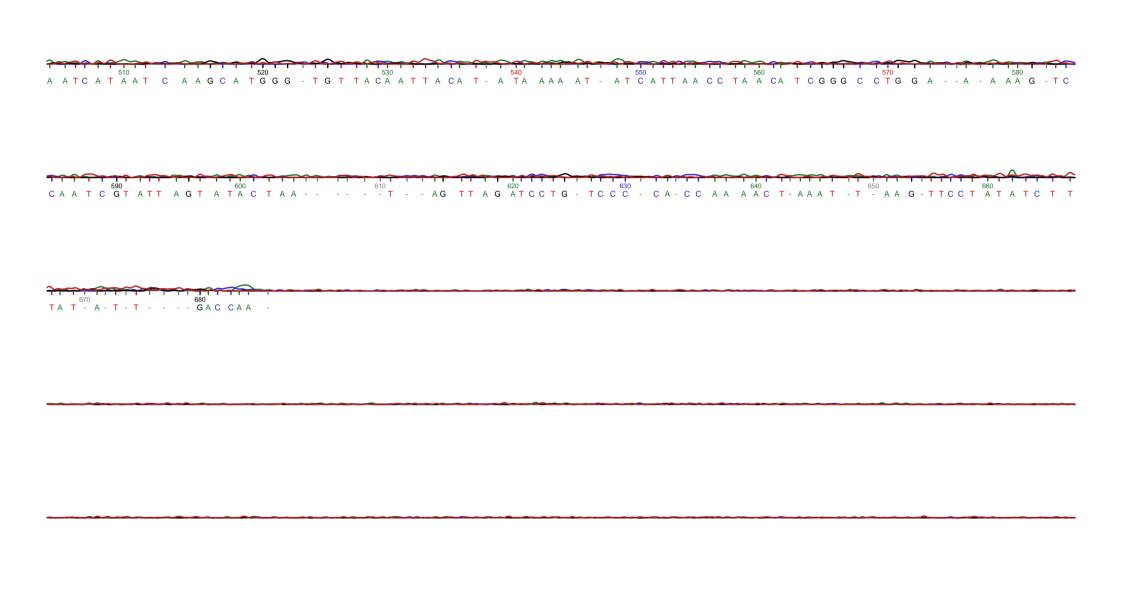
Samples: 14825
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A ATCATAAT C AAGCA TGGG - TGT TACAATTACA T - ATA AAA AT - AT CATTAA CC TAA CA TCGGG C CTGGA - - A - A AAG - TC

C AA TCG TATT AGT ATAC TAA- - - - -T - -AG TT AG ATCCTG- TCCC - CA-CC AA AAC T-AAAT -T - AAG -TTCCT ATAT CT T

TA T - A- T- T - - - GAC CAA -

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