TGCG AC G ACCCA G CG CCG A GGGGGCTCTCTG GC TATTAACT GACGCTG AG GCGCGAAAGCGTGGGTAGCAACAGGAT

TAGATACCCTGGTAGTCCACGCCGTAAACGATGAGTACTAGATATTGGAGCCTTTGACCGCTTCAGTGTCGCAGCTAACGCAT

TAAGTACTCCGCCTGGGAAGTACGGTCGCAAGACTAAAACTCAAAGGAATTGACGGGGGCCCGCACAAGCGGTGGATTATGTGG

TTTAATTCGAAGCAACGCGCAGAACCTTACCTAGGCTTGAAATCCCGAGAAGATCGACCAAAATTGATTGTGCTCTTCGGAGAACT

CGGTGACAGGCGCTGCATGGCTGTCGTCAGCTCGTGTCGTGAGATGTTGGGTTAAGTCTCGCAACGAGCGCAACCCCTATCCTT

AGTTGCCAGCATTAAGTTGGGCACTCTAGGGAGACTGCCTGGGTTAACCAGGAGGAAGGTGGGGATGACGTCAAGTCCTCATG

Samples: 11205
Bases: 721
Average spacing: 16

500 510 520 530 540 550 560 570 580
G C C C T T A T G T C T A G G G C T A C A C A C G T A A T A C A A T G G T G C A T A C A A A G A G A C G C G A T C A G G T G A C T G T C A G C A A A T C T C A A A A G

590 600 610 620 630 640 650 660
T G C A T C T C A G T C C G G A T T G G A G T C T G C A A C T C G A C T C C A T G A A G T T G G A A T C G C T A G T A A T C G C A G A T C A G C A C G C T G C G G T

670 680 690 700 710 720
G A A T A C G T T C C C G G G C C T T G T A C A C A C C G C C C G T C A C A C C A T G G G A G T T G A T C T A C