Samples: 14821
Bases: 473
Average spacing: 32

CTAA TTT TCGG GC T TTTTA G ACA CC T GG TAT TTAA G GCCCG G CC TT CAG TATGT TAATAA GATTA GAG C TC TCG G C TCC

G G G G G C TATG T TAGGAG AC G A T CATC T TTTA TAATG TAATT G T TAC G G CACAC G C TTTTA TTAT GA TTTTT G TT C G CCC T GCC

TG TTAA TTAAG T GATC G G C G GGG A TATG CCA GC G TTAATT C C TG TTATGATTGACAA TTTT GG AT C G C GAC TG TC TGG AATA

AAA T ACAT GTCG T TCCC G A TTAT GACC C C TGG CCC T TG C TTTTT G AT GC CC TTC G G TTTTA TAC AAA TTC TCC TC TG TACCTG

G TT GAAA TG TA TAT C TCCATTA A T GGG G TAC AAAG T C TTG CACG G C G TTC TG T GAA TTTT G C TAAT TTTAG C TTACA TTTC

CC TGGA GC TTC C CCT TTAT TATC TGC TAG TAATTT TATTAG CAAC T GG TGAT AAAAA G GAGAC

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