

10 20 30 40 50 60 70 80 90
C T T T C G 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 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

100 110 120 130 140 150 160 170
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 T T T C A T A G T T G A T G T G G A C A A T A C C A A A A C G T T T

180 190 200 210 220 230 240 250
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

260 270 280 290 300 310 320 330
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

340 350 360 370 380 390 400 410 420
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 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 T C C G

430 440 450 460 470 480 490 500
G A T G A A T T T C C C A A C C A T A T C G G T A A C C G G T T C T T T 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 T G G T G C T G C

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

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

Samples: 17469
Bases: 1472
Average spacing: 12

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