Samples: 14822
Bases: 712
Average spacing: 21

- - - - - - - - - C - CA - T - - - TGCTG - TTT - C - - G ATCG - G - TCCT GATGATGG CT ATCAC - TGATGAATGGA -

A - - - AA - GAA TTT TT - ATG - TGG - GGATGAAAGGGG AGAGG CCGACTGCAG - TTCCATC TTGTCGAGG - ATGTCG - ATCTAT

GTG - A - - TGTAA - - - AAAACACA - CC - AACAAC - GGTA AG - TTGACCA - TTC - TATGA - G - TCT - AAA - - AAACA - CATTTT - A

C - CATCAAAAAAAACAGT - - ATAATCTCGAATCGG - TT - TTTTT - CTA TCCTAGGAA - C - GGATATCAACCTAT GACAC - CCC

TCTGGGCTTAAAAATATCCCGAATGTTGTGGCTTGTGACTTTC - - - GCTTATTTAAAAATATTACTT GGG - TACCGACTGAG

ATGGG AATACAGAGTTTGCCTGCTGATTCTGTA - ATG - - TAGA - AG - GCT - AC - - CTAC - TT - TG CGTTTTGCG - TACAACA

Samples: 14822
Bases: 712
Average spacing: 21

CTGTTTCCAATGGCAAG-G-TAA GTATACGG TCTCATG TATGTTCCCCCTCAAAAATAGTC-CTCTTGTTTCTTTTGACAT

A-ACAAAAGATCATCT--ACT-ACTGTCATAA-ATTTA-TAT----TGAATATTT-TTCTA-TCC-CA-CCTCT-G-A

C-TGA----A--A--ATAATAAA-ATTTGT-TTGCGA-A-A-G---A-G

Samples: 14822
Bases: 712
Average spacing: 21