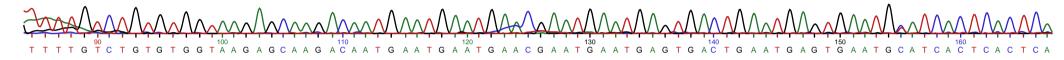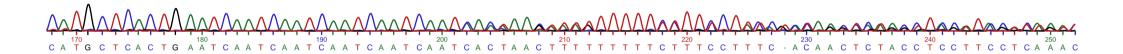Samples: 14827
Bases: 959
Average spacing: 16

- TTTTCTTCATATTAGTCAAACAACAATAATTTGTACTGCAAAGTCTCACAAATTT - - G - TG

TTTTGTCTGTGTGGTAAGAGCAAGACAATGAATGAATGAACGAATGAATGAGTGACTGAATGAGTGAATGCATCACTCA

CATGCTCACTGAATCAATCAATCAATCAATCAATCACTAACTTTTTTTTCTTTCCTTTC-ACAACTCTACCTCCTTCCTCAAAC

AATTCTATGCATCTTACTTACATACCATTCTTACTTGCACG-ACTTATTC--GCCCGTTG-T-GTTGAA-GATACA--ACT--

- TATATATA---AAGAACG-A-ACTTCC---TAA-TCT-CCATC-AT-A--TAT-AA-AAA-A-AC-AAC-A-TTG---C

- G-CG--T-ACTGG--G-CC-TC-T-A----G--G--CTCC-G--GGTGG--GC-G--TGAA-CCCTGCC-GCTG--

Samples: 14827
Bases: 959
Average spacing: 16

930   940   950

A