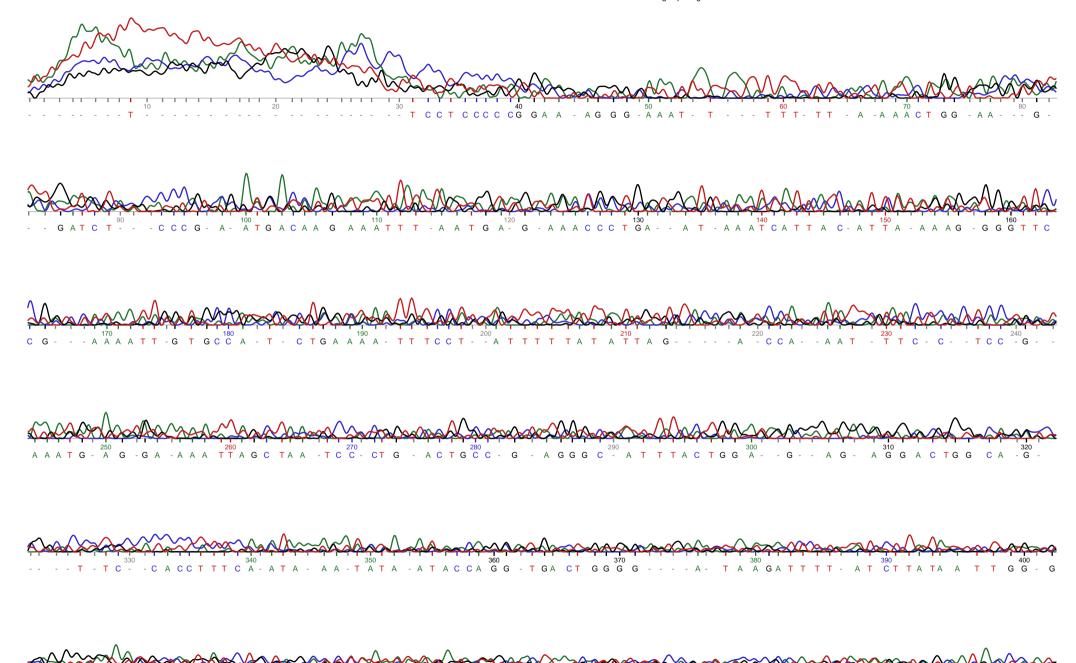
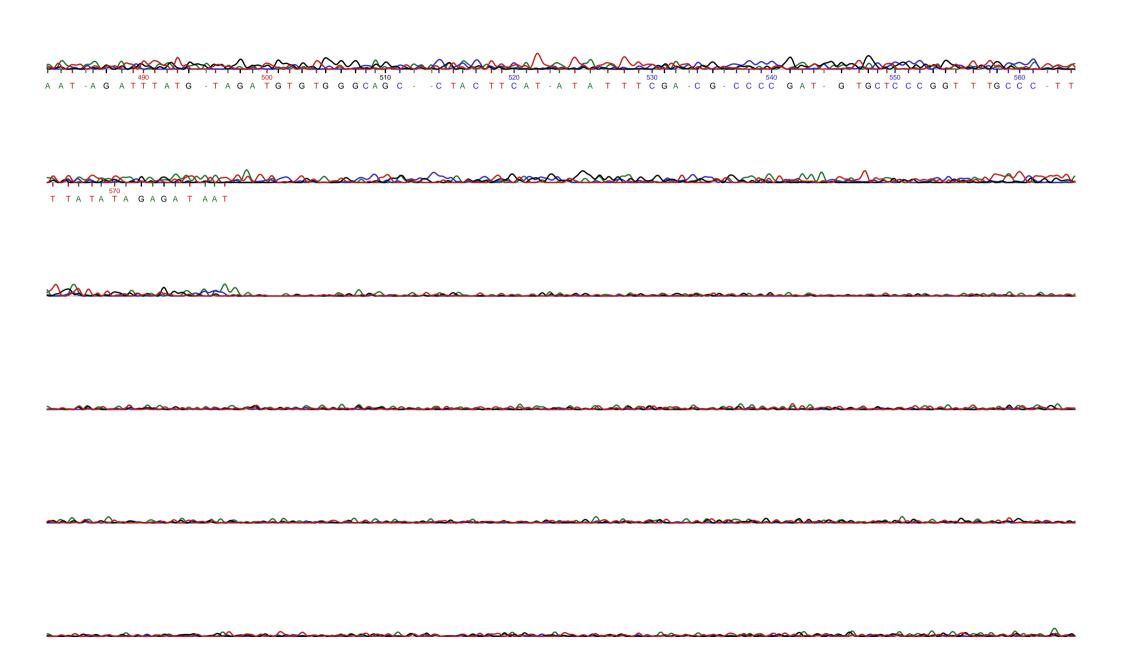
Samples: 14824 Bases: 580 Average spacing: 26



GGGGAAATGT - TATCTCTAGCCCTAATT - TATCT - ATA - C-GTTTCCG-T-C-CCCATTGAATAA TGGAATAACCAT

Samples: 14824 Bases: 580 Average spacing: 26



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