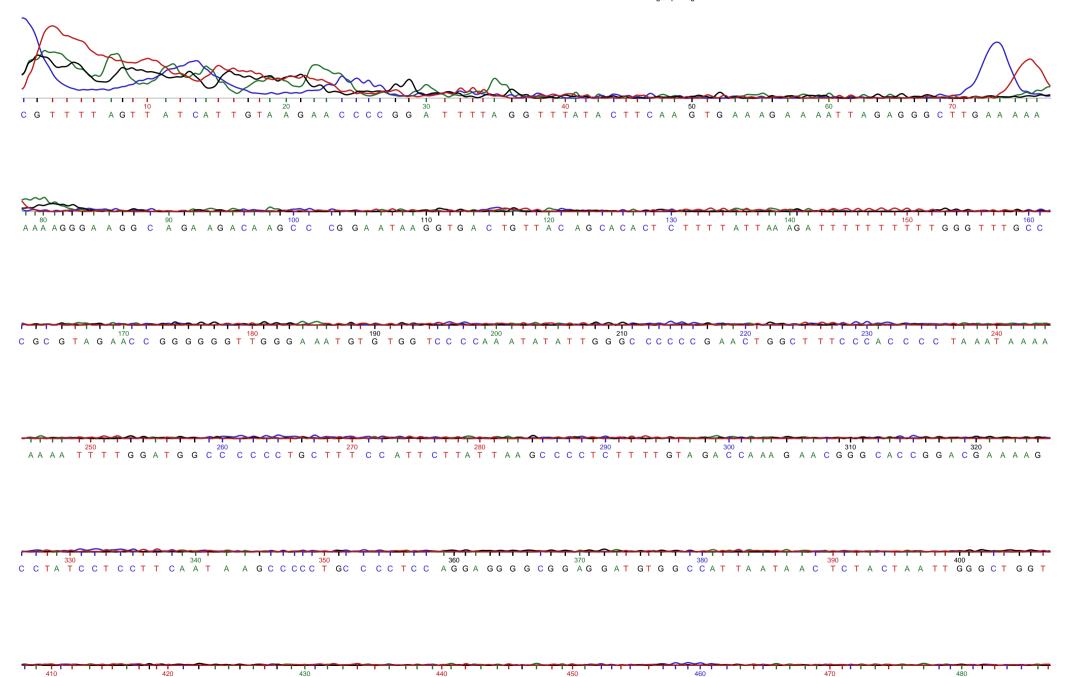
Samples: 14825 Bases: 545 Average spacing: 28



CATCT ATT ATGTT C G AAG AT G AACA AT T A AA ATAT A AT G A TATA CCTC T C C C C C A TC A A AATTCA A T A A C CACG T C C AT

Samples: 14825 Bases: 545 Average spacing: 28