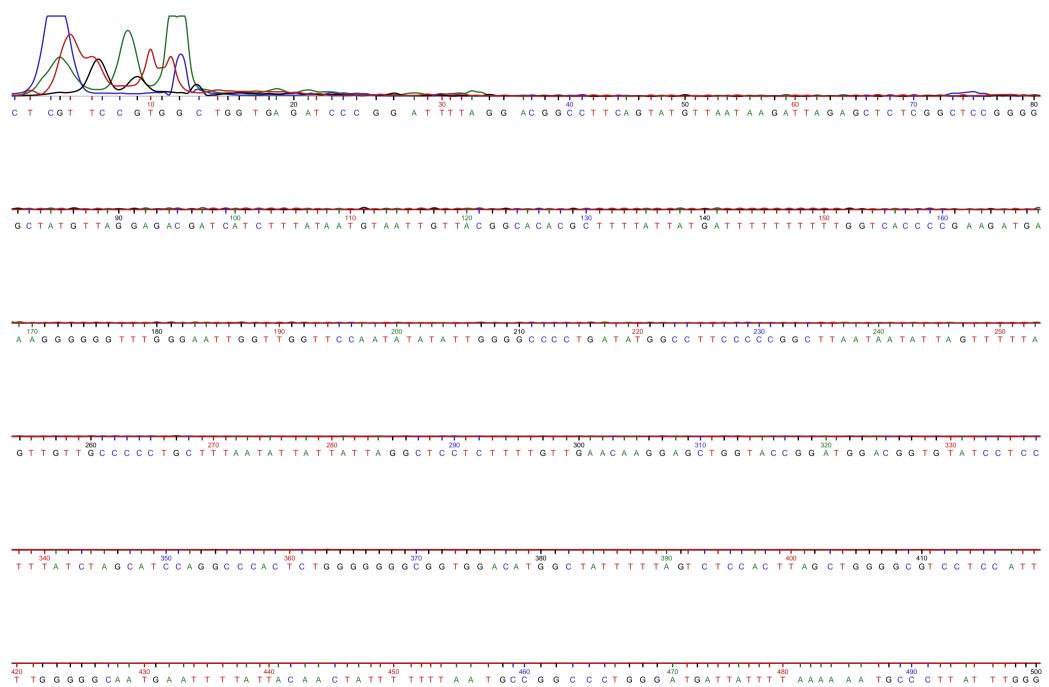
Samples: 14827 Bases: 560 Average spacing: 27



Samples: 14827 Bases: 560 Average spacing: 27

GGGGGG CCAACC TGAAATAC GGCCTTTTT AT TAA TAC T ACCCTTTTCCCCC GGAAA AGG	

Samples: 14827 Bases: 560 Average spacing: 27

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