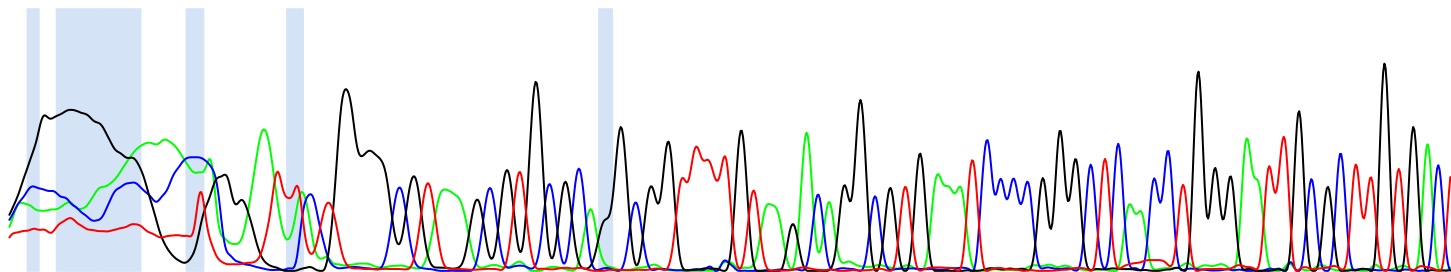


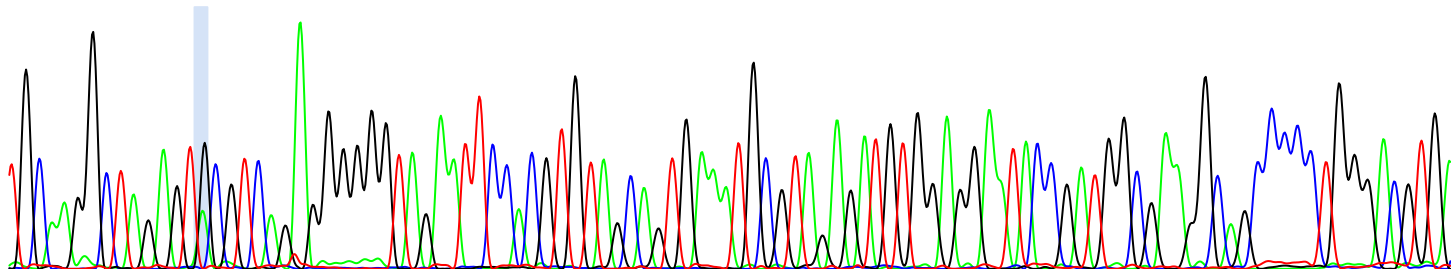
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1



TGC AA GGCTAGAGT CGGTCAGAGG GG GG TAGAATTCCAC GTG TAGAGTAGTGA AATGCG TAGA GATG TGGAGGAGTACC GA TGGCGAA GCGAGCCCCCTGG GAC GTG A
 TGC AA GGCTAGAGT ACGTCAGAGG GG GG TAGAATTCCAC GTG TAGAGTAGTGA AATGCG TAGA GATG TGGAGGAGTACC GA TGGCGAA GCGAGCCCCCTGG GAC GTG A

92



CTGACGCTCA TGCACGACGCGTGT GGAGCACACA GATTACAA GCG AATG TAGT CAAA
 CTGACGCTCA TGCACGACGCGTGT GGAGCACACA GATTACAA WCG AATG TAGT CAAA

198

