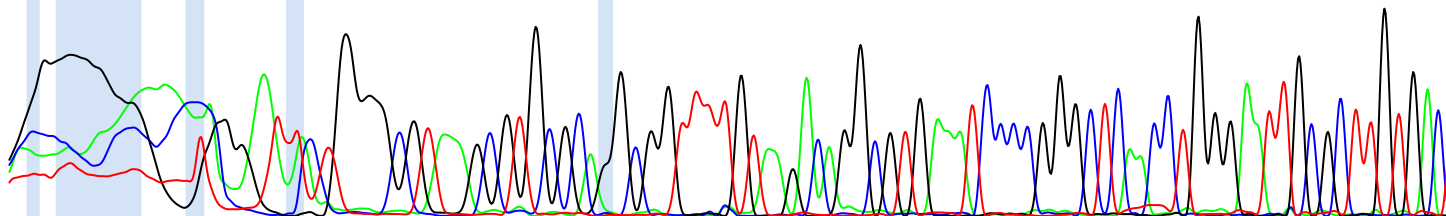


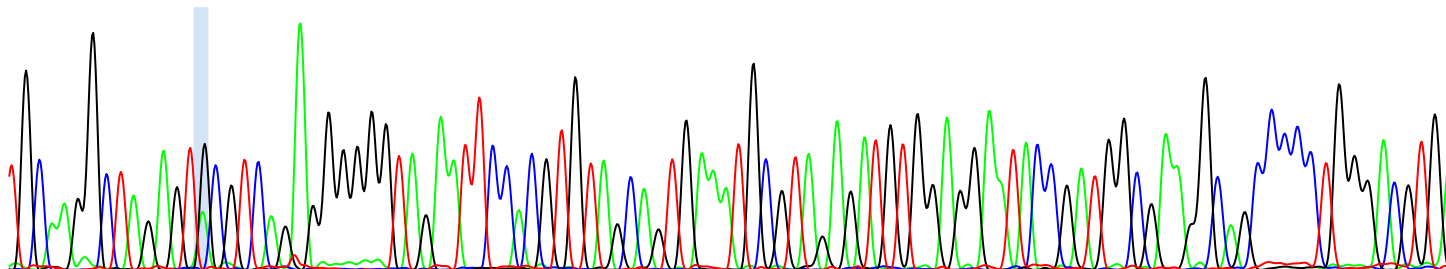
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1



TGC AA GGCTAGAGT CGGTCAGAGG GG GG TAGAATTCCAC GTG TAGAGTAGTGA AATGCG TAGAGATG TGGAGGAGTACC GA TGGCGAA GCAGCCCCCTGG GACGTGA
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92



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

198

