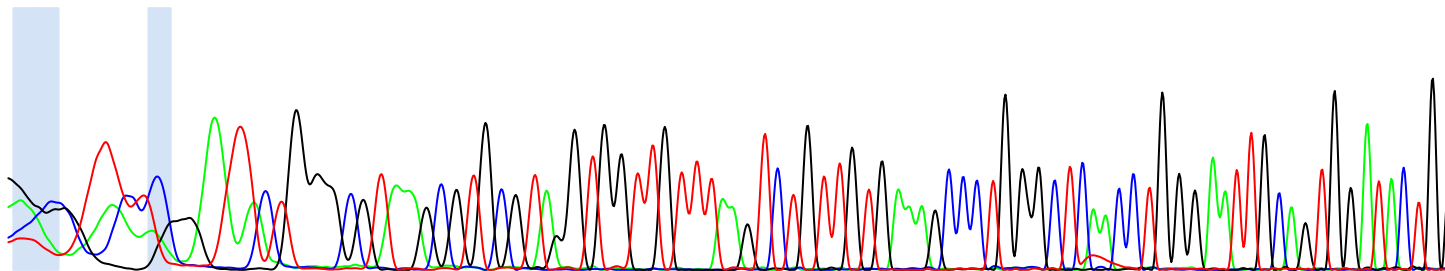


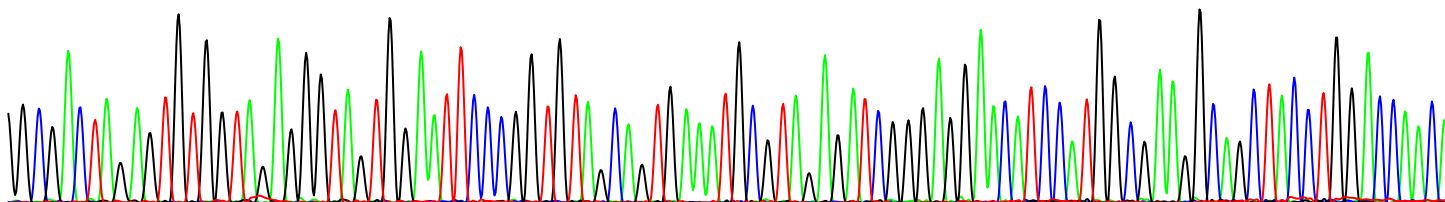
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1



G CG ACTAGAGTGTGG TAGAGGG TAGTGG AATTCCCGG TG TAGCAGTG AAATGCG TAG AGATCG GG AGGAACATCCATGG CGAAGGCA GC TACC TGG ACCAACA
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95



CTGACAC TGAG GCACG AAAAGC GTGGGGA GC AAACAGG ATTAGAAA CCC CGGTAGT CCAAA
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