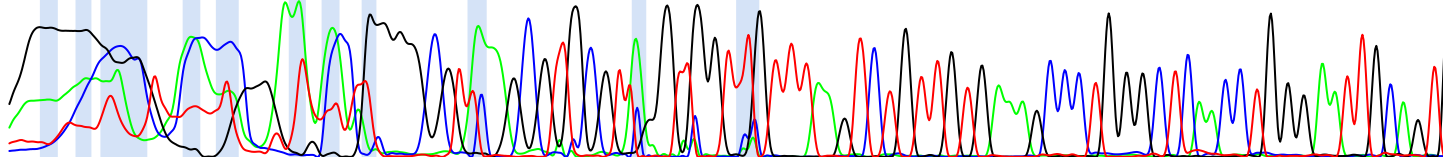


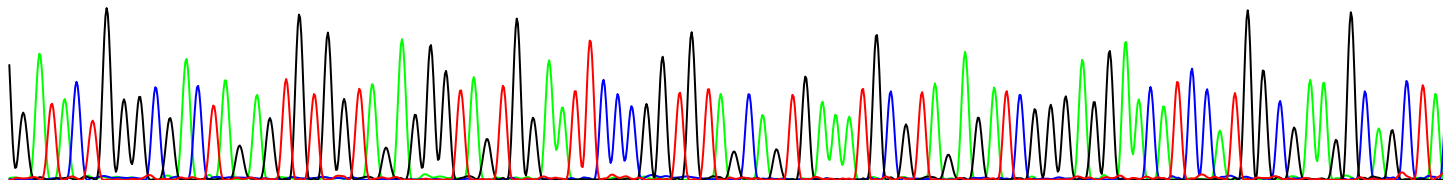
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1



G A TACTGGG CGACTAGAG TGTGGTAGAG GGTAGTGG AATTCCC GGTG TAGCA GTG AAATGCG TAG AGATCGG GAGG AACATCCATGG CG AAGGCAGC TAC  
G A TACTGGG CGACTAGAG TGTGGTAGAG GGTAGTGG AATTCCC GGTG TAGCA GTG AAATGCG TAG AGATCGG GAGG AACATCCATGG CG AAGGCAGC TAC

98



CTGGACCAACACTGACACTGAGG CACG AAAGCGTGGG GAGCAAACAGG ATTA GAAACCCACGGTAGTCCTCAA  
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198

