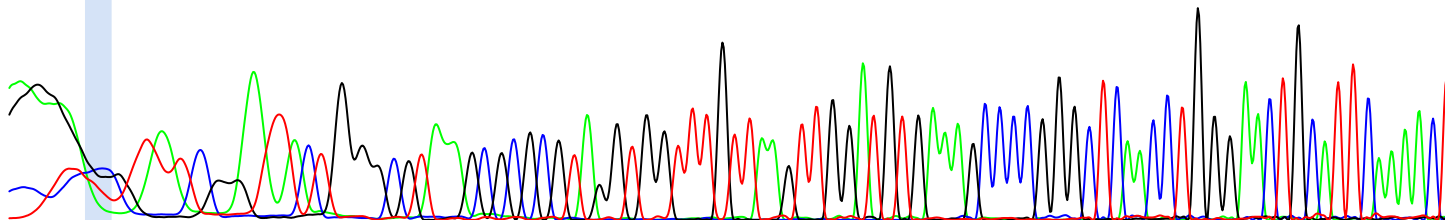


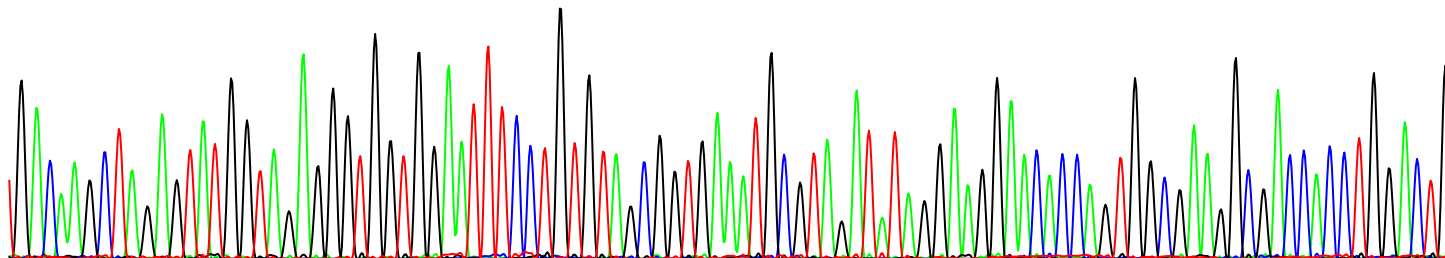
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1



G ACAAG CTAG AGTATGGTAG A GGGTGGTGG AATTTTCC TG TG TAG CG GT GAAATGCGTAGATATAG GAAGGAACACCAG TGG CGAAGGC ACCACC TG GACTG  
G ACAAG CTAG AGTATGGTAG A GGGTGGTGG AATTTTCC TG TG TAG CG GT GAAATGCGTAGATATAG GAAGGAACACCAG TGG CGAAGGC ACCACC TG GACTG

97



ATACTGACACTG AGGTGCGAAAGCGTGGGGAGCAACAG GATTAGAAACC C GGTAGTCCAG  
ATACTGACACTG AGGTGCGAAAGCGTGGGGAGCAACAG GATTAGATACC K WGTAGTCCAA

199

