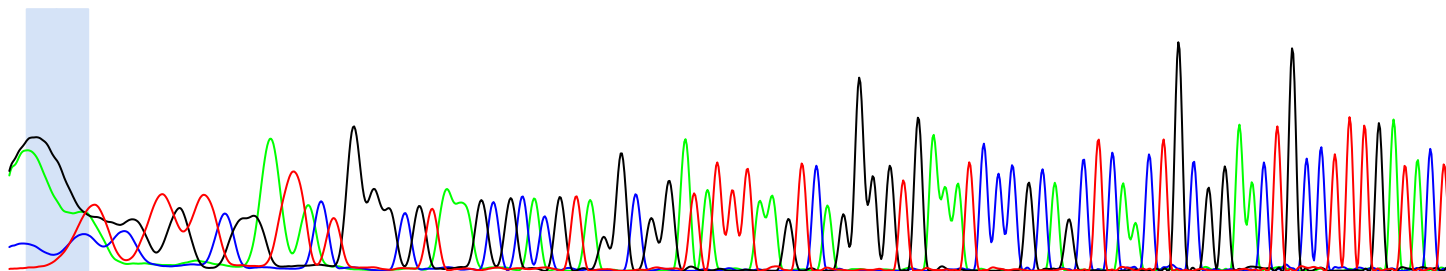


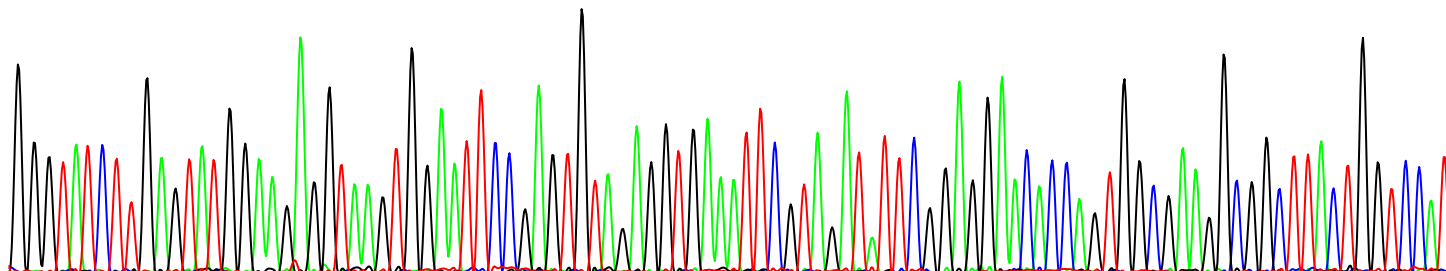
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1



G GGTATCTT T GAGTATG GAAGA GGTAA GTGGAATTCCGAGT GTAGA GGTGAAATT CGTATGATATT CGG AGGAACACCAG TGGCGAAGGCCG CTACTG GTCCAT
G GGTATCTT T GAGTATG GAAGA GGTAA GTGGAATTCCGAGT GTAGA GGTGAAATT CGTATGATATT CGG AGGAACACCAG TGGCGAAGGCCG CTACTG GTCCAT

96



TACTGACGCTGAGG TGC GAA AGCGTGGG GAGCAAACAG GATTAGAAACCC TA
TACTGACGCTGAGG TGC GAA AGCGTGGG GAGCAAACAG GATTAGATACCC TT

199

