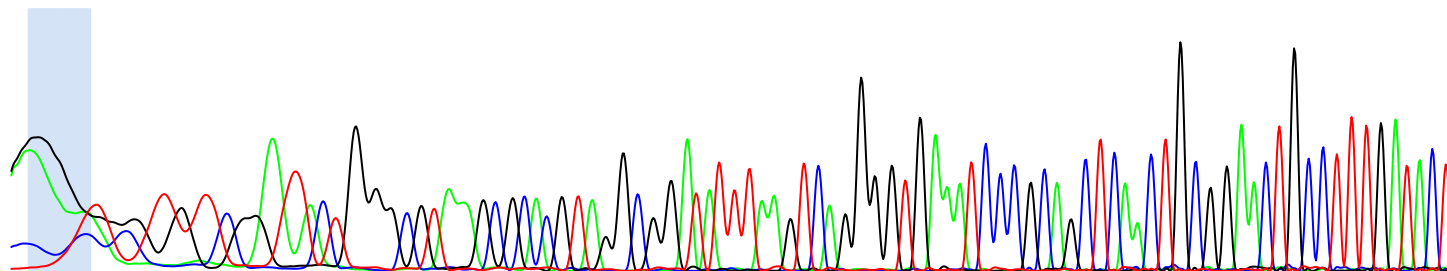


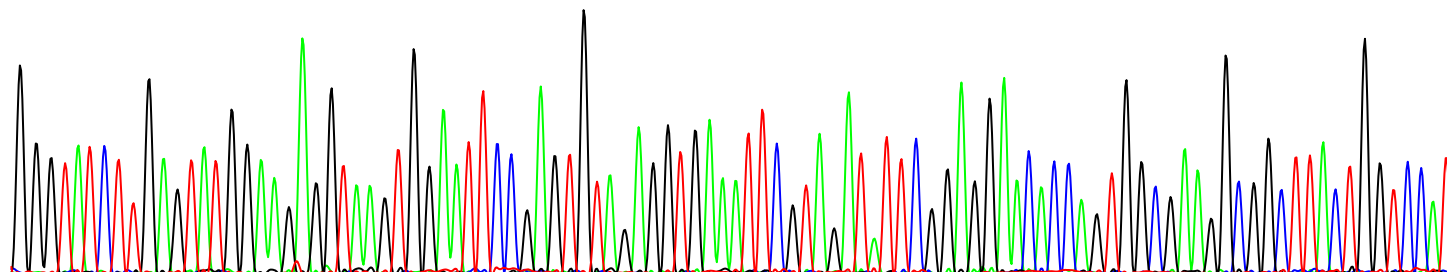
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1



G GGTATCTT GAGTATGGAAGA GTAA GTGGAATTCCGAGT GTAGA GGTGAAATT CGTATATTTCCG AGGAACACCAG TGGCGAAGGCCG CTACTG GTCCAT  
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96



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

199

