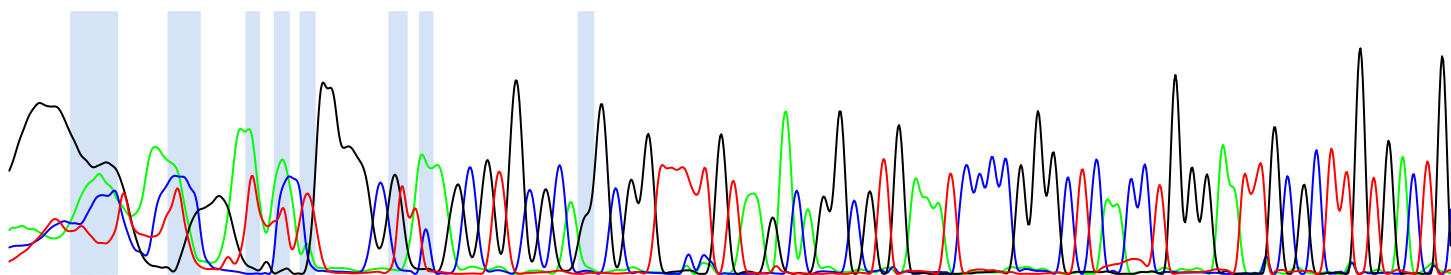


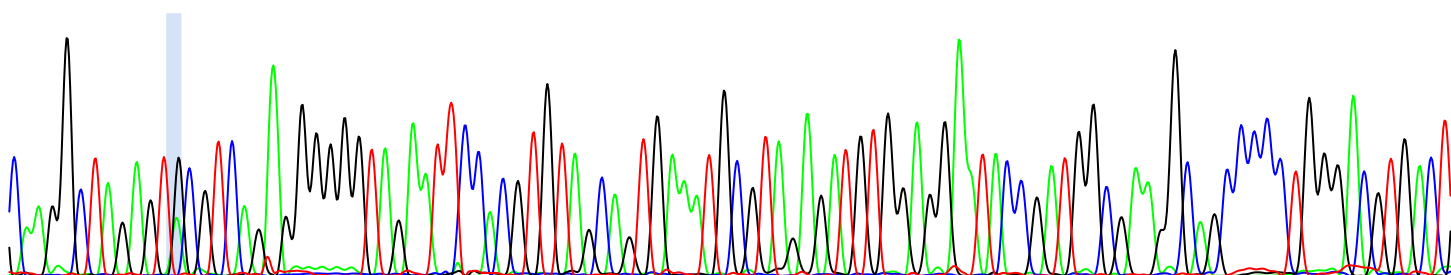
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1



CAAGGCTAGAGTGC G TCAGAGG GGGG TAGAATT CCA CGTGTAGCAG TG AAATGC GTAGA GATGT GGAGGAGTACC GATGGC GAA GGCAG CCCCCTGGG ACGTGACT  
 CAAGGCTAGAGTAC G TCAGAGG GGGG TAGAATT CCA CGTGTAGCAG TG AAATGC GTAGA GATGT GGAGGAGTACC GATGGC GAA GGCAG CCCCCTGGG ACGTGACT

93



GAC GCTCATG CAC GAGA GC GTGGGG A CACACA GATTAGACCCG T G TAGTC AAA  
 GAC GCTCATG CAC GACA GC GTGGGG A CACACA GATTAGACCCG A G TAGTC CAA

199

