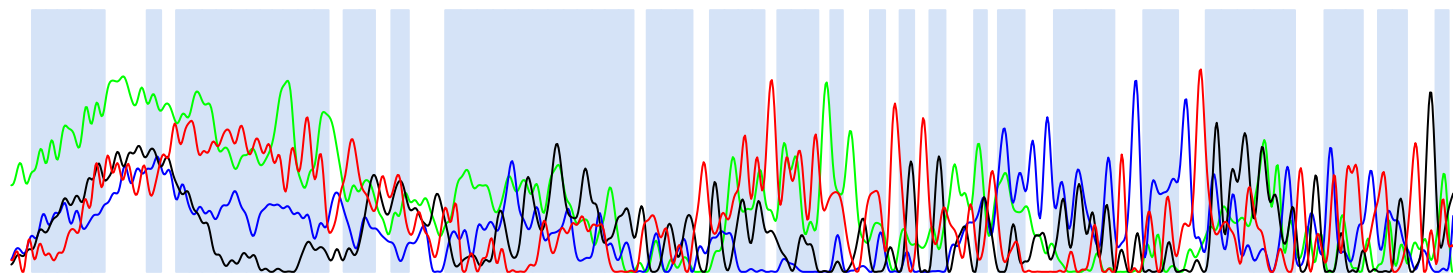


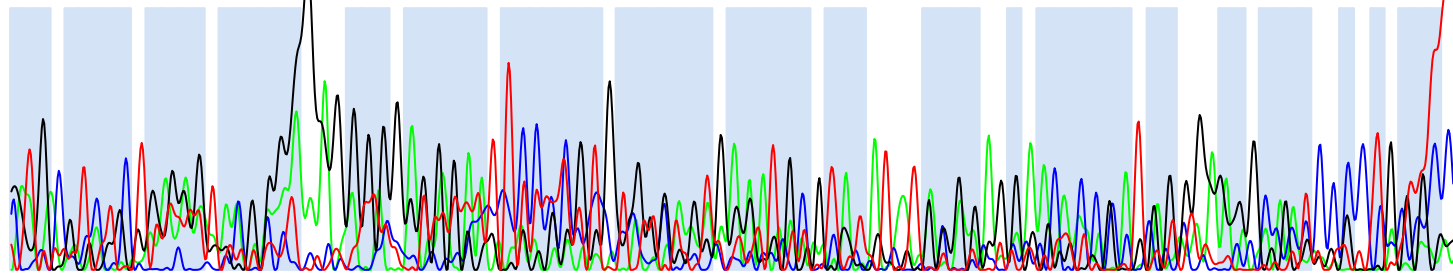
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1



GT GCGTCTCTG GAGTACGGGAGAGGGGGAGGGAA TTCCTCGTGTG TGAGTG AAATGCGTATATATACGGAGGAA C CCGATAGNGAGGGCAATCCCC TGTGTC  
CA ACWCA GGTW WKWTKAAMATGAGATMGWWTYTCTTSTT CGCMGMTAAGGGTAWWGC GCATATGKAAAGAAG W WACCTSTNGAAMGASCSCCG CCGSCCC

100



TG TANGACNCTCATGTAGG AANGC TG CGGAGCA ANAAGATTA TAATCCACCCCA CATCACT A A A  
TM MSNMARNRRGKCY SAYG CTNGR AGGGT AGCG ANKG WTMGA CYYCWTYWTTCYKCA WAGM A T

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