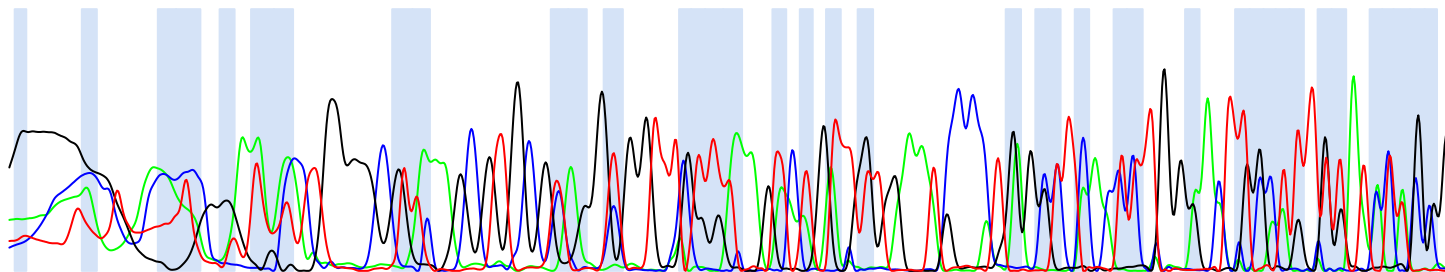


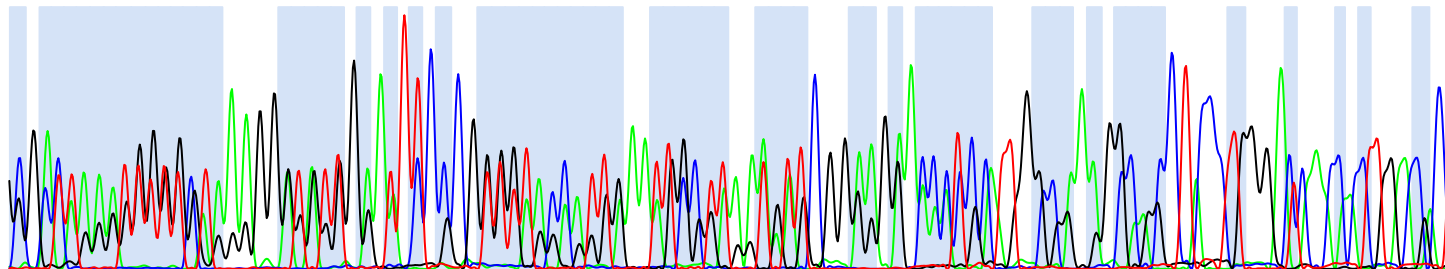
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1



C GACTAAAT GG TGCTAAAG GGTATTGAA TTT CC CGGGG TACCATTGAA TTG CTTAAAGTTCGG AAGAA C C C T C C GT GGCAAAGGCGCTACG TGGACCA CCA TTGA C CT
G GCTAG GGG TT GTGAGA AG GAGGGGGGAATCCG CGT TTT AGGAGGG AAAAGC GGAAA TAAGCGG GGGGAAAA CGAGT GC GGA GGCAAGCTACG GGGATCA CAA CTGA G CT

92



G AGGCACAAAA GC G AGGGGA GCAAA CA AGA TTAG A CCA TGT AT CAA
C TGCCACGAAA GC G AGGGGA GCAAA CA TGA TTAC T ACA AGT GT CTG

201

