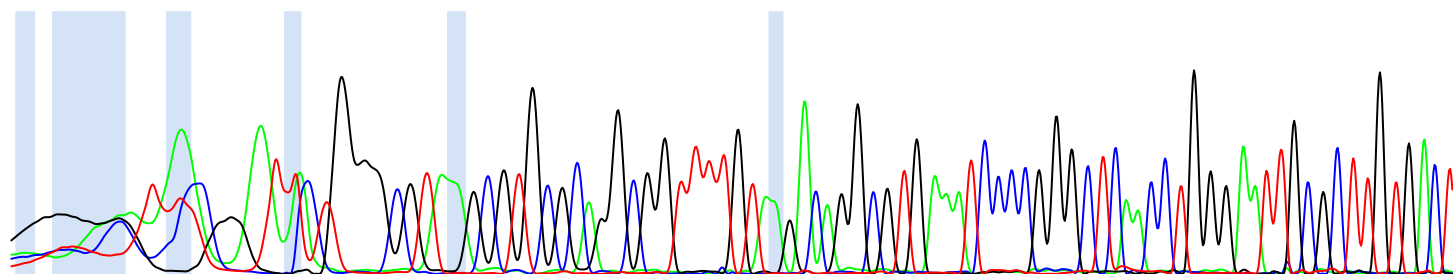


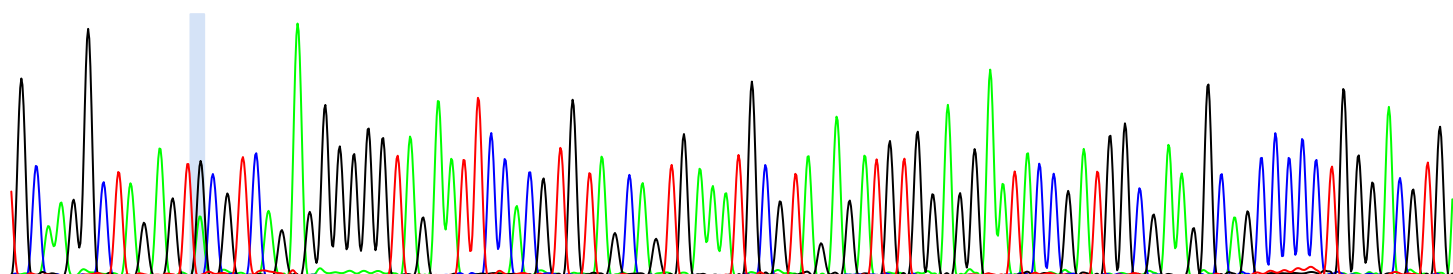
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1



GCAAGG CTAGAGTGC GTCA GAGG GGG GGG TAGAATTCC ACGTGTAGC AG TG A AATGCG TAGAGATGTGG AGG AATACCG ATGG C G AAGG CAGCC CCC TGGG ACGTG
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93



ACTGACGCTCATGCACGAAAGCGTGG G GAGCAAACAGG ATTAGAAACC
 ACTGACGCTCATGCACGAAAGCGTGG G GAGCAAACAGG ATTAGATACCA

197

