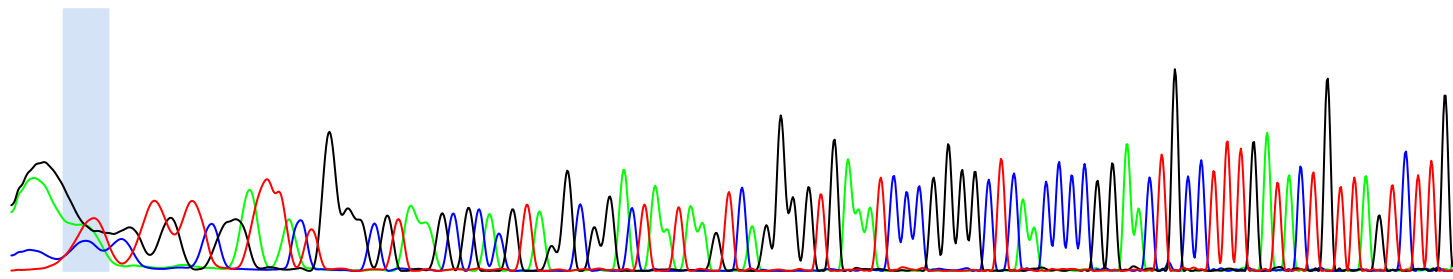


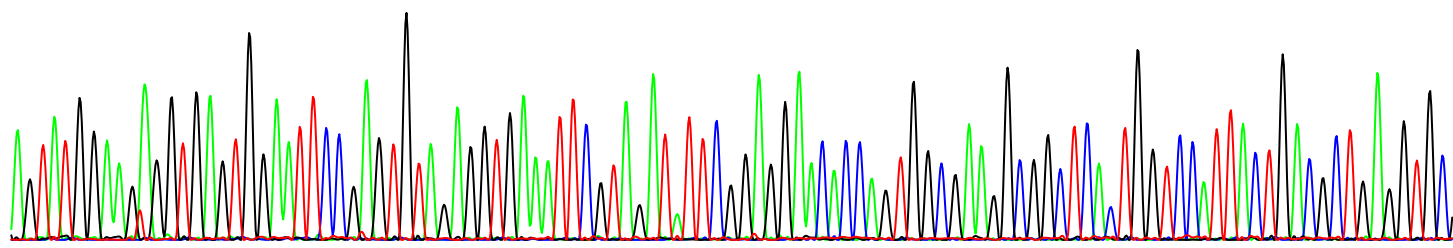
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1



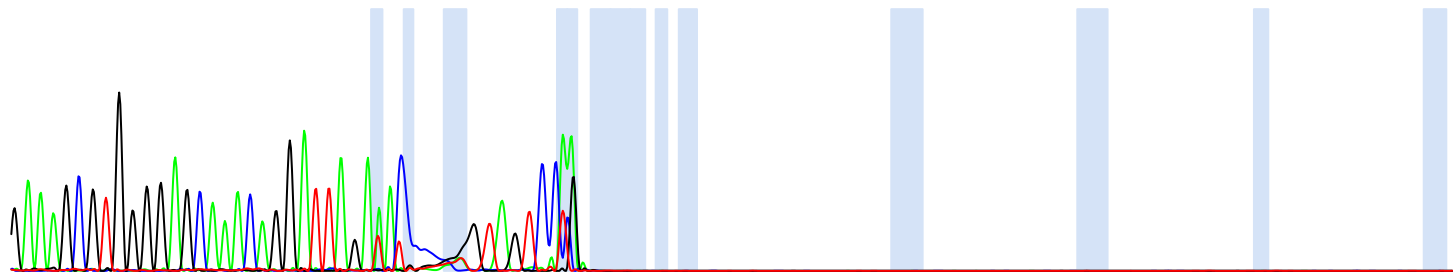
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GAAAGCG TGGGGAGCAA ACAGGATTAGAAACGG AGTAGTCCAAA C TACA TC CT T C GC ATT A C A ACC AC GCG T C C C T T GTATTACC CC CT C
GAAAGCG TGGGGAGCAA ACAGGATTAGATACGCG TGTAGTCCCTGA W A GCTT CT T C GC ATT C C A ACC AC GCT T C C C T T GTTTACC CC CT T

218



TGG CAA C T
TTG TAA C T

309

