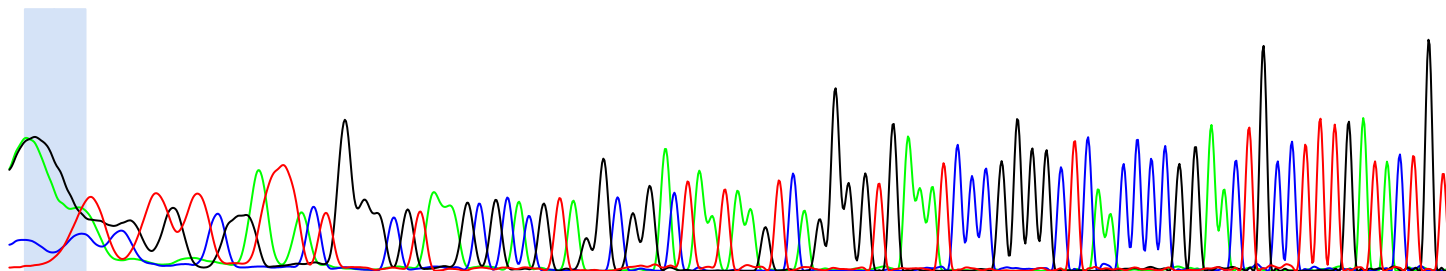


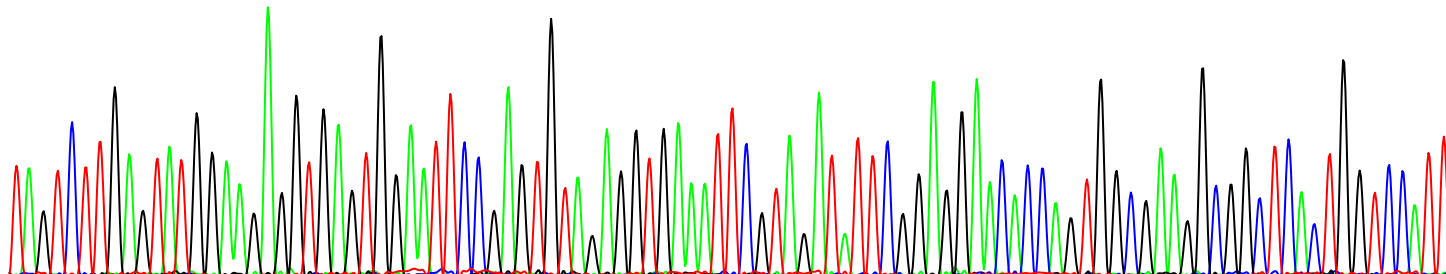
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1



TAGTCTTG AGTA TG GAA GA CG TGAGT CG AA TT CC GAG T GTAGA GG TGAAATT CG TAGA TATT CG GAGG AACACCAG TG GCG AAGG CCG CTCA C TGG TCCATT
TAGTCTTG AGTA TG GAA GA CG TGAGT CG AA TT CC GAG T GTAGA GG TGAAATT CG TAGA TATT CG GAGG AACACCAG TG GCG AAGG CCG CTCA C TGG TCCATT

98



ACTGACGCG TGAAG TGC GA AAGCG TGG GGAGCAAACAG GA TTAGAAACC C GG TAGTCCAG
ACTGACGCG TGAAG TGC GA AAGCG TGG GGAGCAAACAG GA TTAGATACC K WG TAGTCCAC

200

