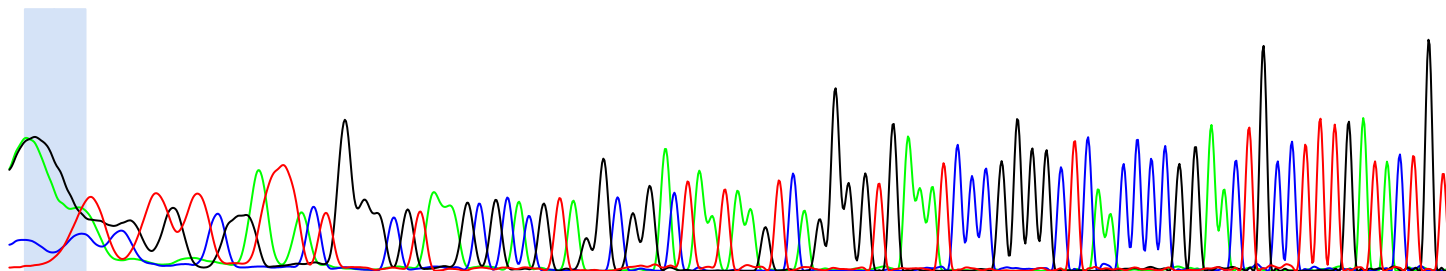


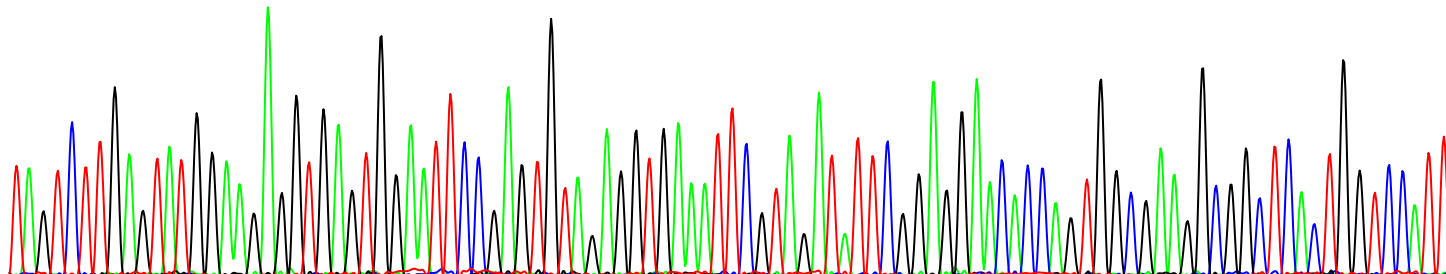
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1



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



ACTGACGCGTGAAGG TGC GA AAGCGTGG GGAGCAAACAGGA TTAGAAACC C GG TAGTCCAG  
ACTGACGCGTGAAGG TGC GA AAGCGTGG GGAGCAAACAGGA TTAGATACC K WG TAGTCCAC

200

