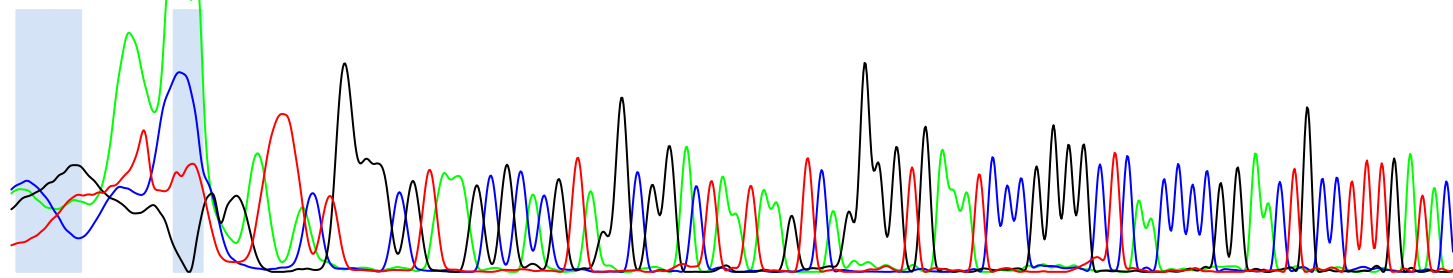


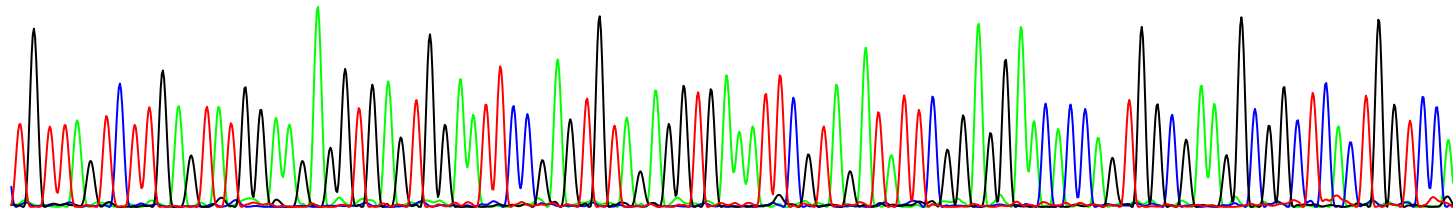
C GTT CATGACAGGAATCTGGGCGTAAAGCGCACGTA GGCG AC TAA TAA G TCA G GGG TG AAA T CC C GGGG CTCAACC CCGG AACTGCCTTTGATAC
 A WTT CATGAT T G GA T AC T GGG CG TAA G CG CAC G TA GGC G AC TAA TAA G TCA G GGG TG AAA T CC C GGGG CTCAACC CCGG AACTGCCTTTGATAC

1



TGTTAGTCTTGAGTATG AAGAGGTGAGTGG AATTCCGAG TGTAGA GG TGAAA TTCG TAGA TATT CG GAGG AACACCA G TGG CG AAGG CCGCTCACTGG TCCA
 TGTTAGTCTTGAGTATG AAGAGGTGAGTGG AATTCCGAG TGTAGA GG TGAAA TTCG TAGA TATT CG GAGG AACACCA G TGG CG AAGG CCGCTCACTGG TCCA

96



TTACTGACGC TGAGGTGC GAAAGCGTGGGG AGCAAACAGGA TTAGAAA CC C GG TAGTCC AA
 TTACTGACGC TGAGGTGC GAAAGCGTGGGG AGCAAACAGGA TTAGATA CC K WG TAGTCC AA

199

