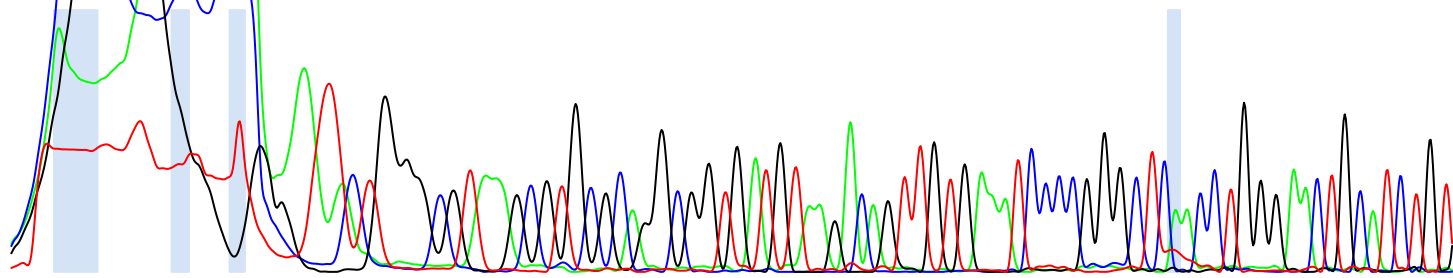


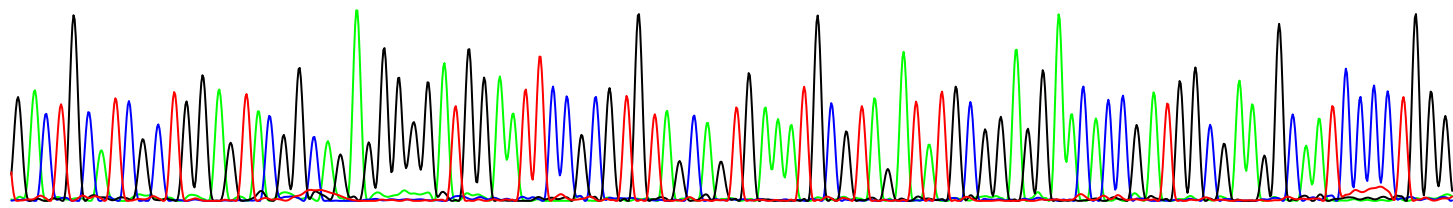
T TC G G TAT CA ACAG G A TAC TGG ACG TA AGCG TGC GCA G GCGG T GATGTAG ACAG TT GT GATATCC CC GG G CTCAA CC TGGG AACT G CATCTGT
T TA Y G TAT TA ATAG G A TAC TGG ACG TA AGCG TGC GCA G GCGG T GATGTAG ACAG TT GT GATATCC CC GG G CTCTA CC TGGG AACT G CATCTGT

1



GAC TGCATCGCT G3 AGTAC GG CAG A GGG G GATGG AATTCCGCG TGTAGCAGTGAAATGCGTAG ATATGCGG AGGAACACC G ATGGCG AAGGCAATCC CCTGGG
GAC TGCATCGCT G3 AGTAC GG CAG A GGG G GATGG AATTCCGCG TGTAGCAGTGAAATGCGTAG ATATGCGG AGGAACACC G ATGGCG AAGGCAATCC CCTGGG

95



CC TGTACTG ACG CT CATGC ACGAAAGC GTGGGG AGC AAACAGG ATTAGAAA CC C
CC TGTACTG ACG CT CATGC ACGAAAGC GTGGGG AGC AAACAGG ATTAGATA CA C

198

