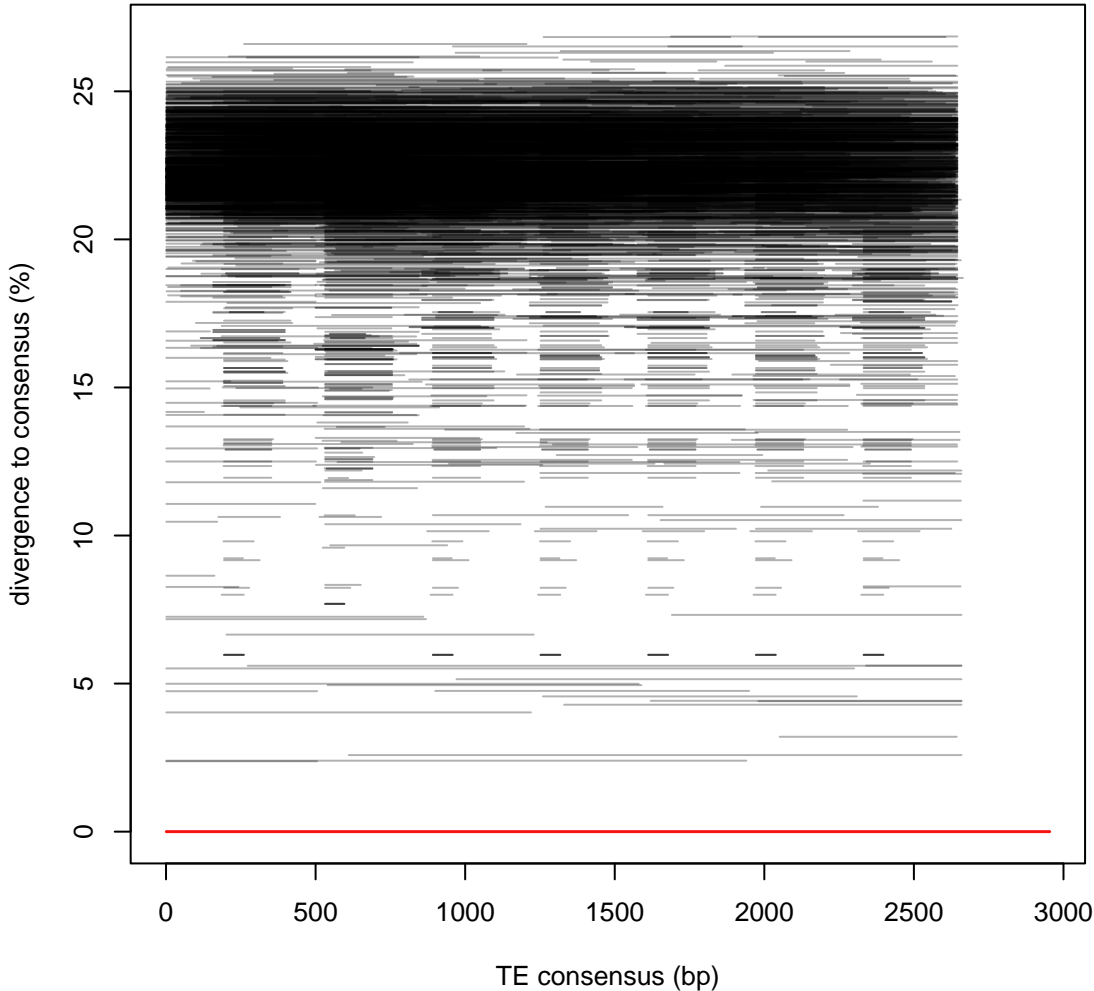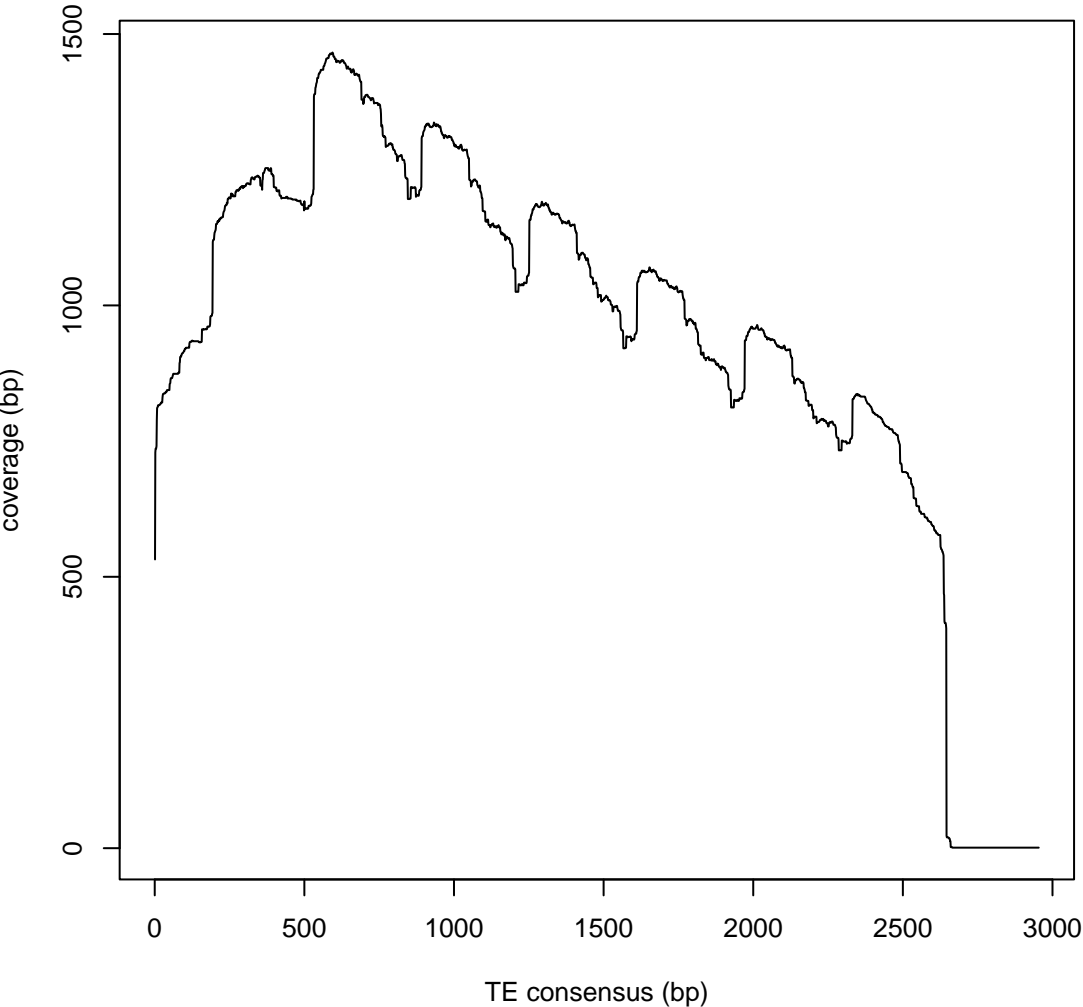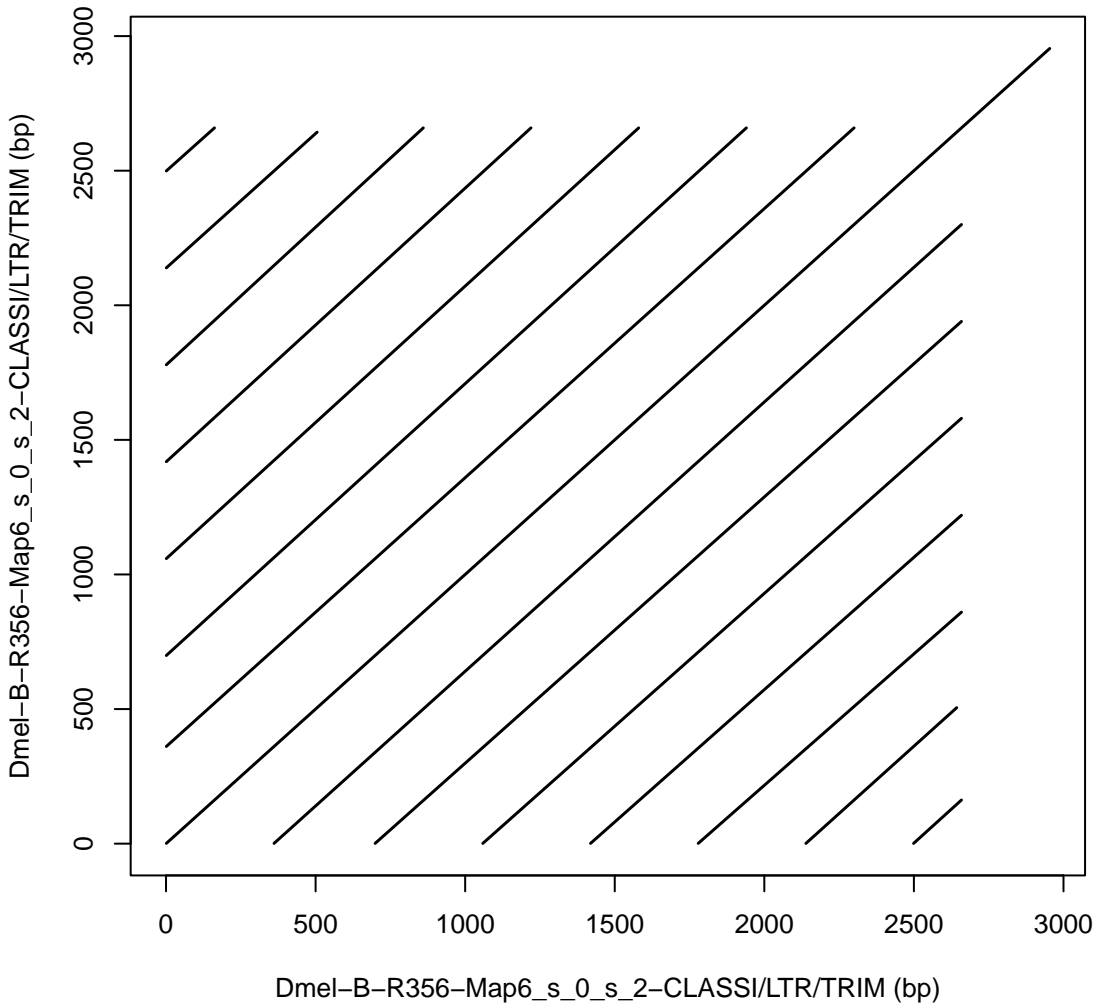**TE: Dmel−B−R356−Map6_s_0_s_2−CLASSI/LTR/TRIM**
consensus size: 2954bp; fragments: 2926; full length: 1 (>=2658.6bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected