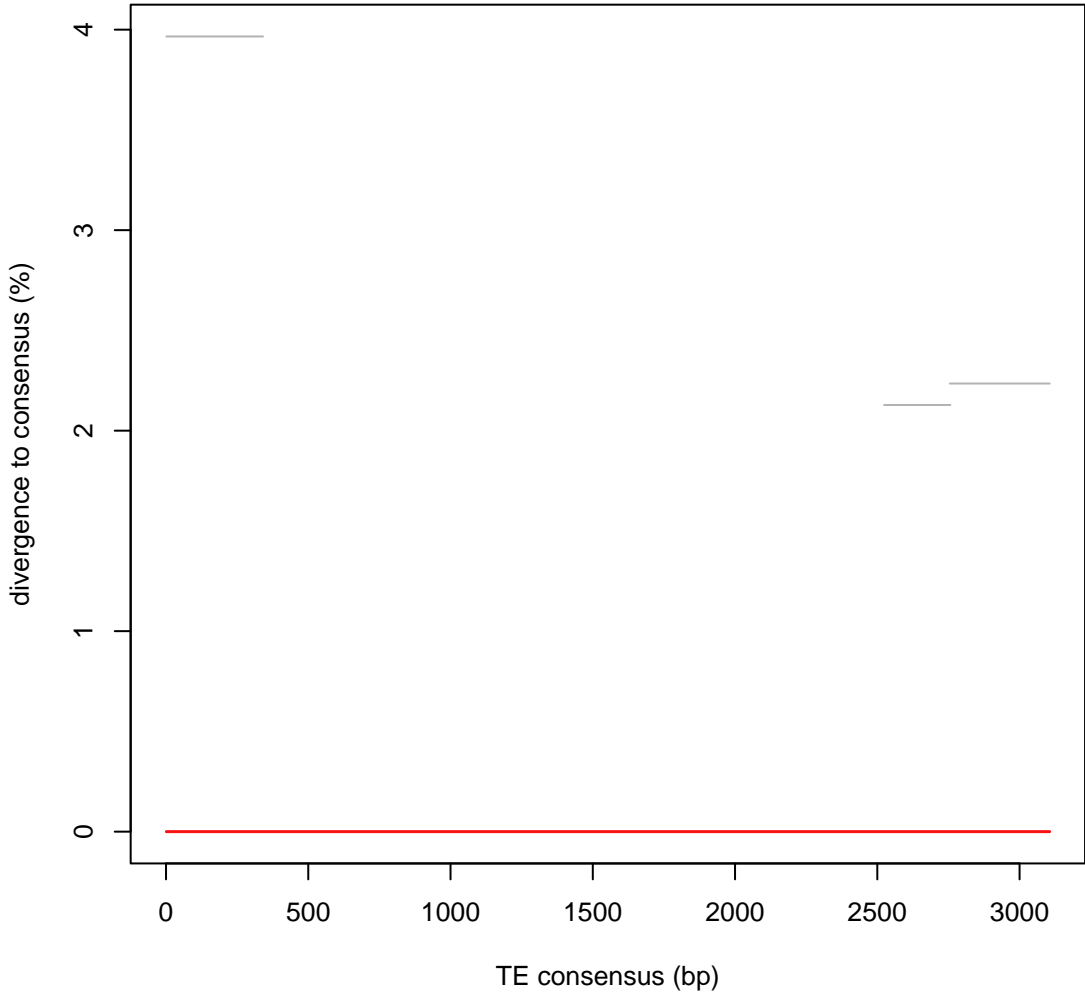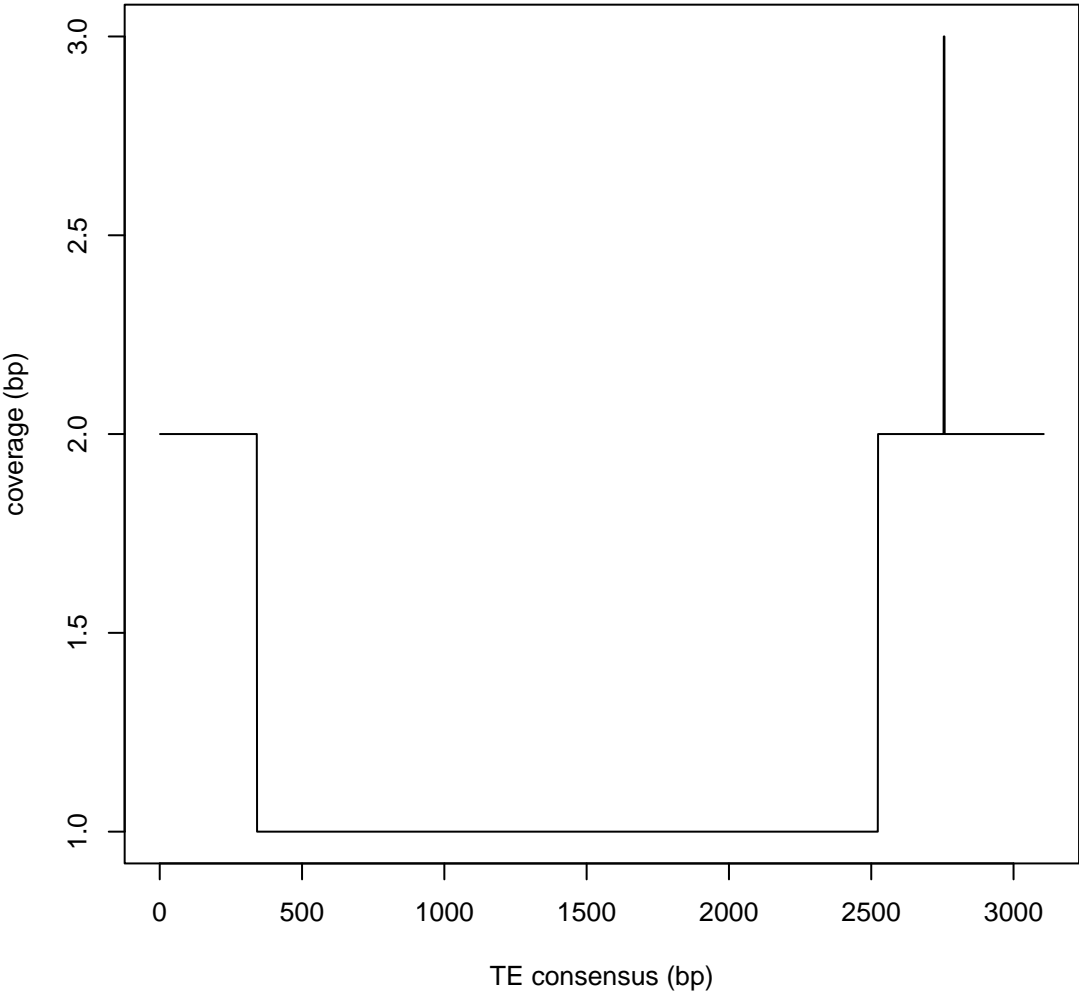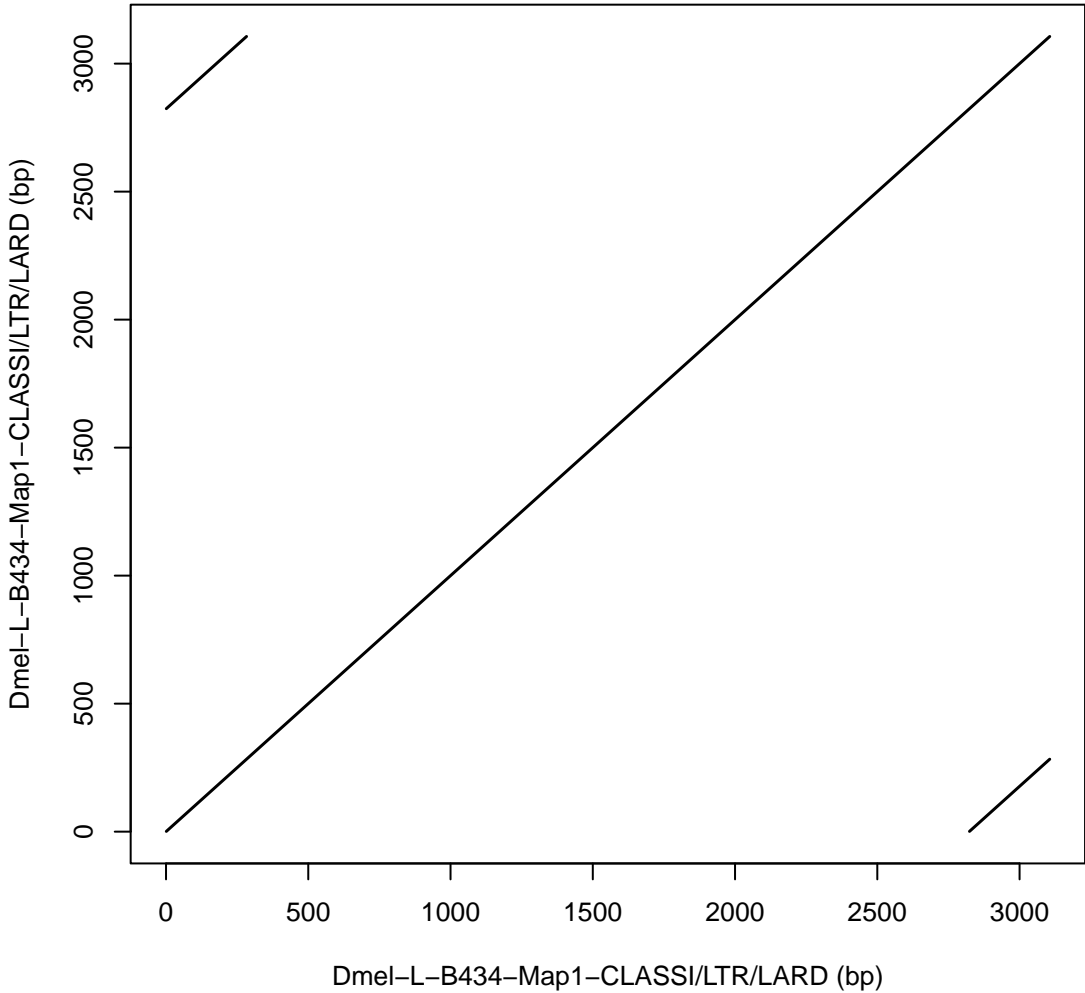**TE: Dmel−L−B434−Map1−CLASSI/LTR/LARD**
consensus size: 3106bp; fragments: 4; full length: 1 (>=2795.4bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected