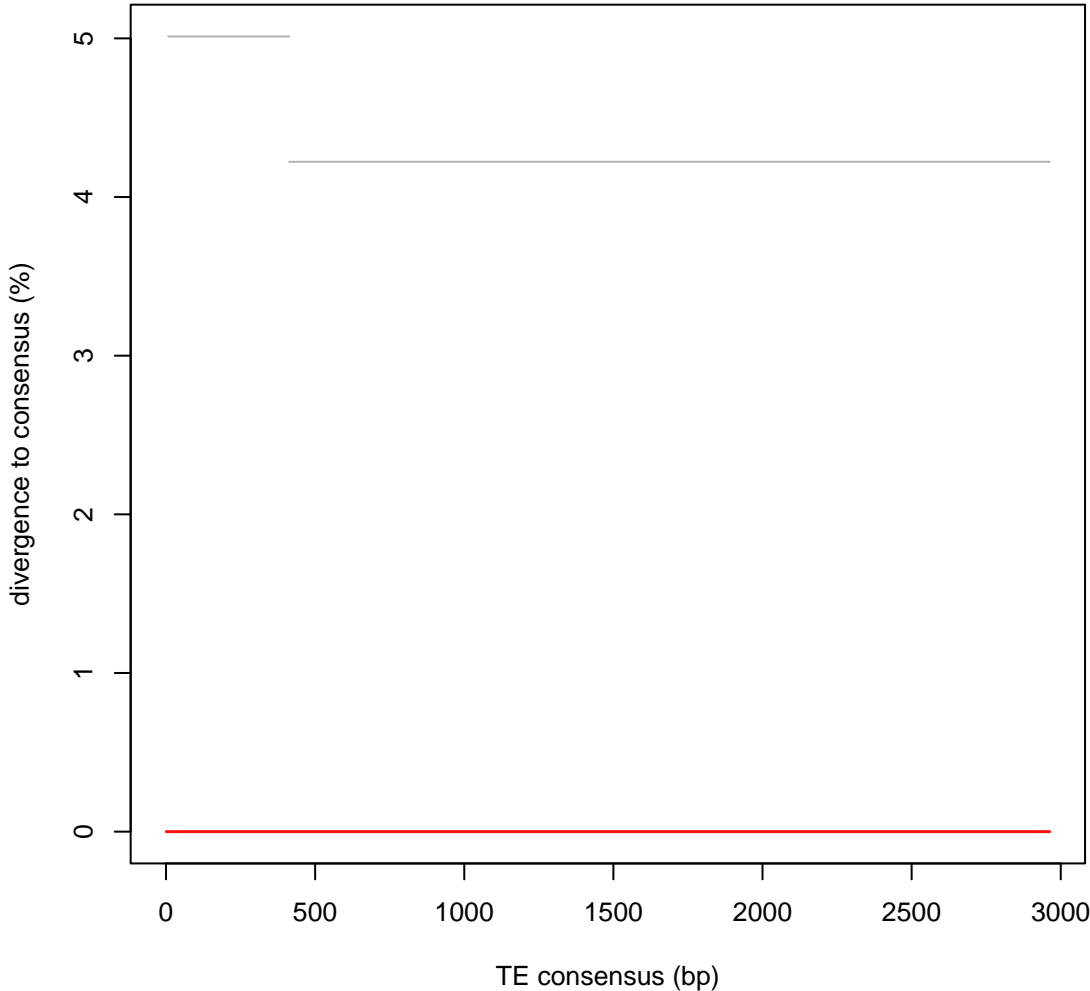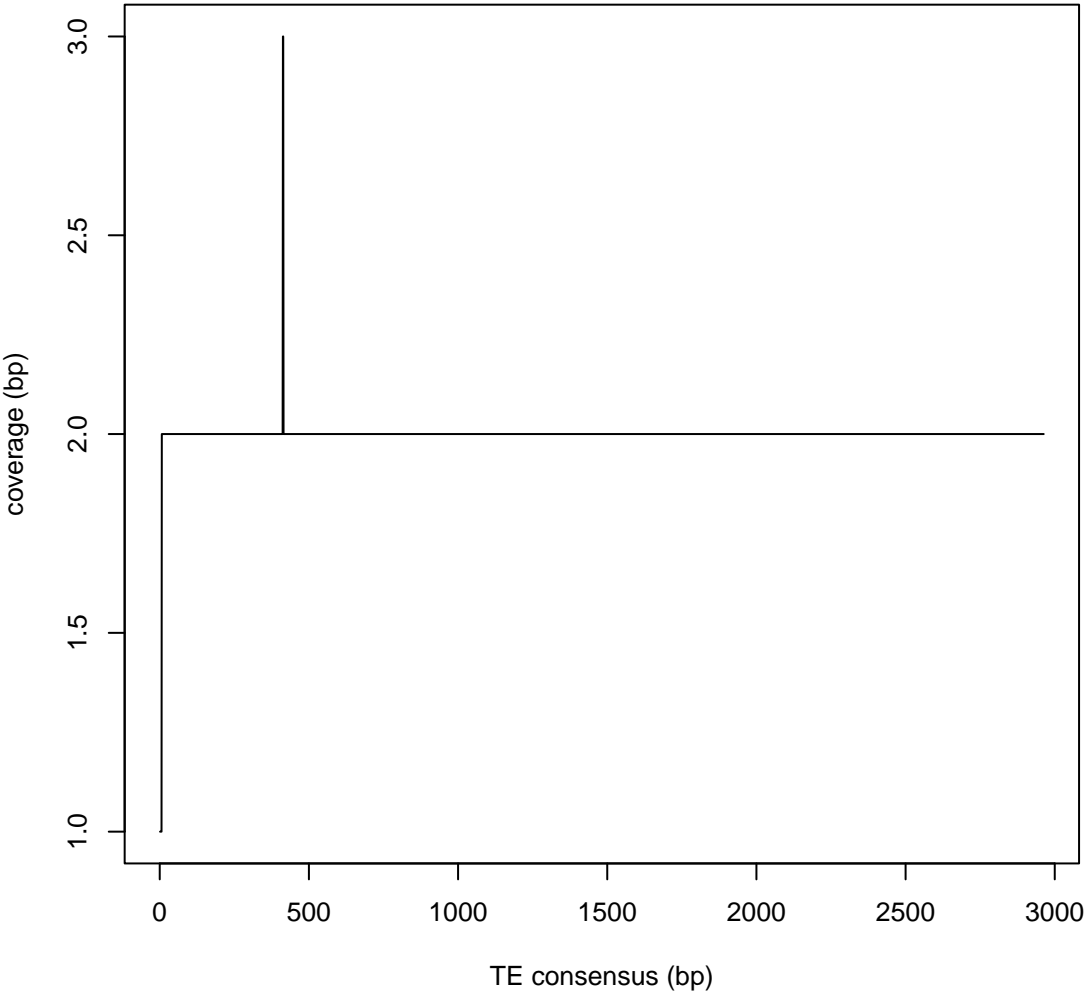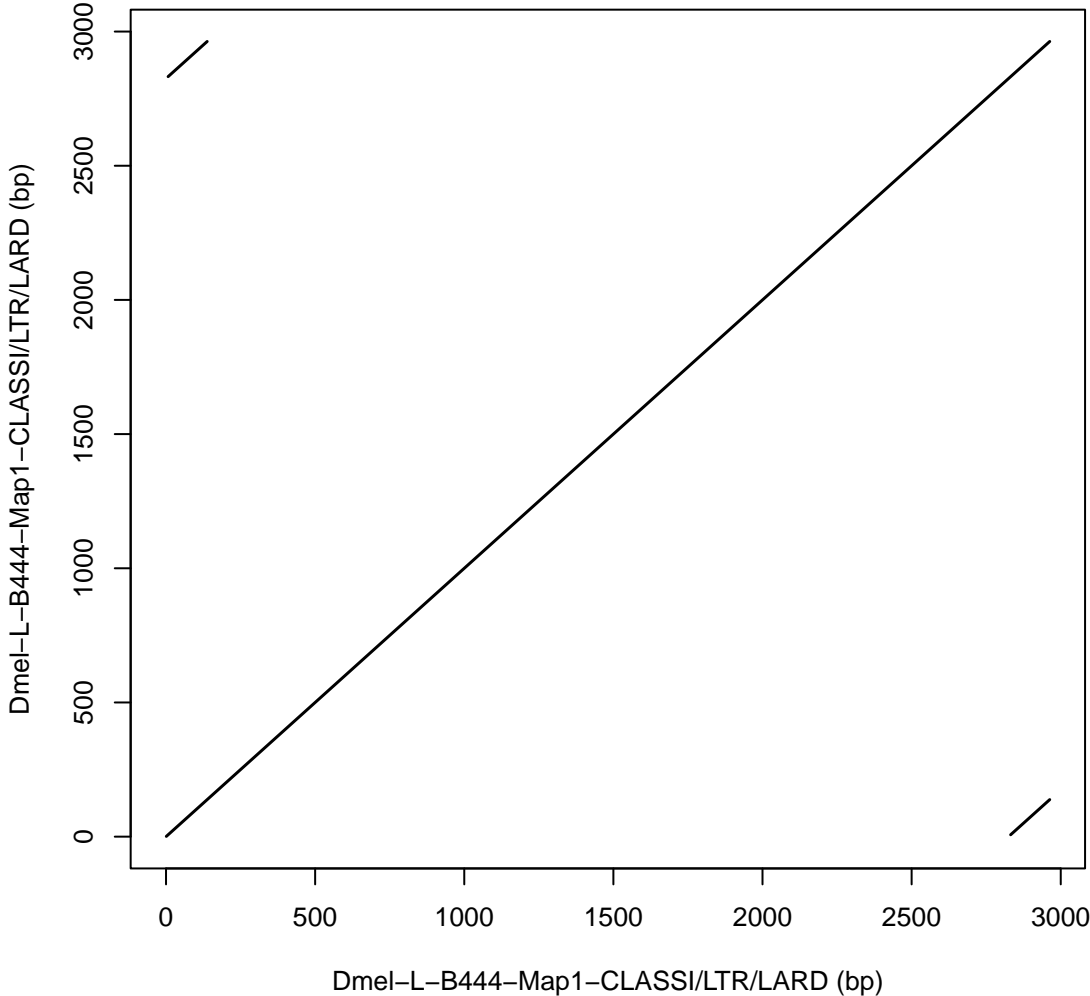# TE: Dmel−L−B444−Map1−CLASSI/LTR/LARD
consensus size: 2963bp; fragments: 3; full length: 1 (>=2666.7bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected