TE: Dmel−L−B489−Map1−CLASSI/LTR/TRIM
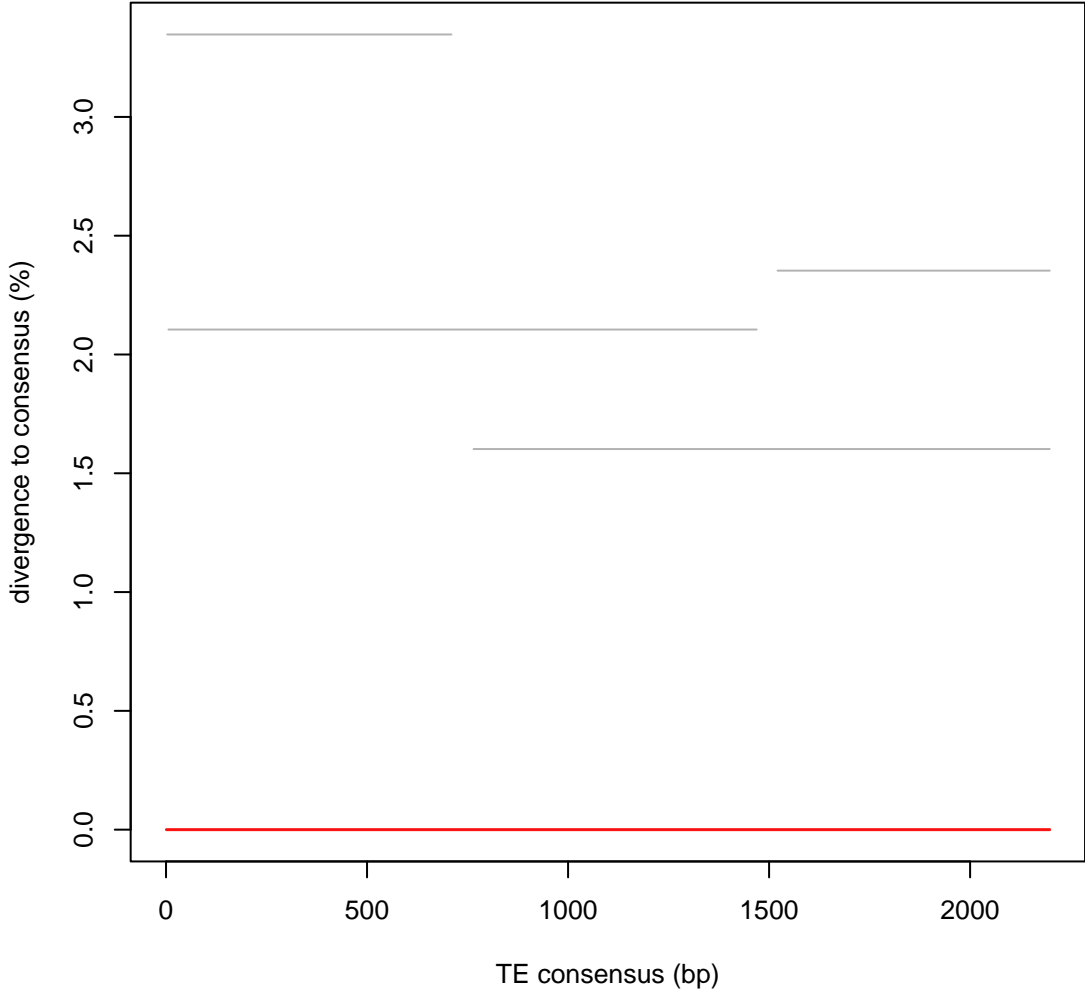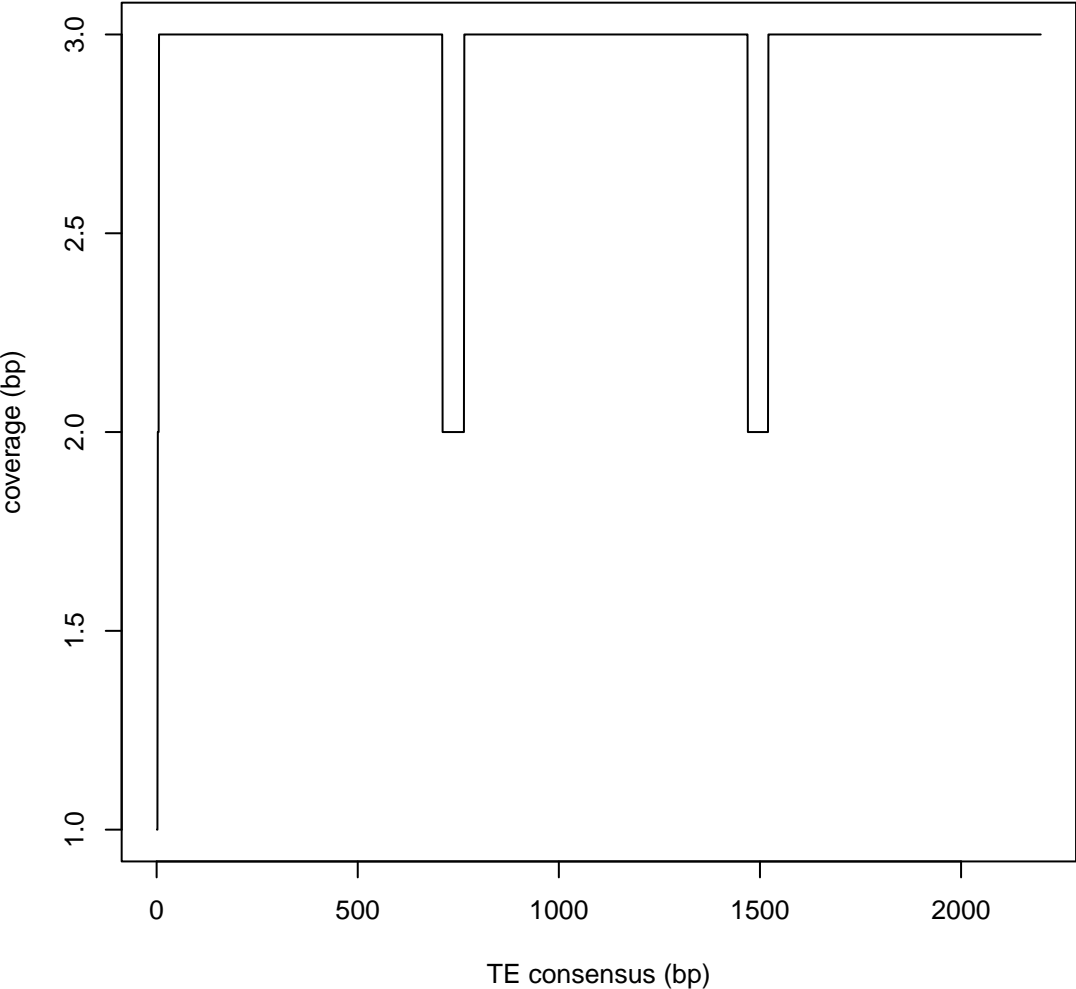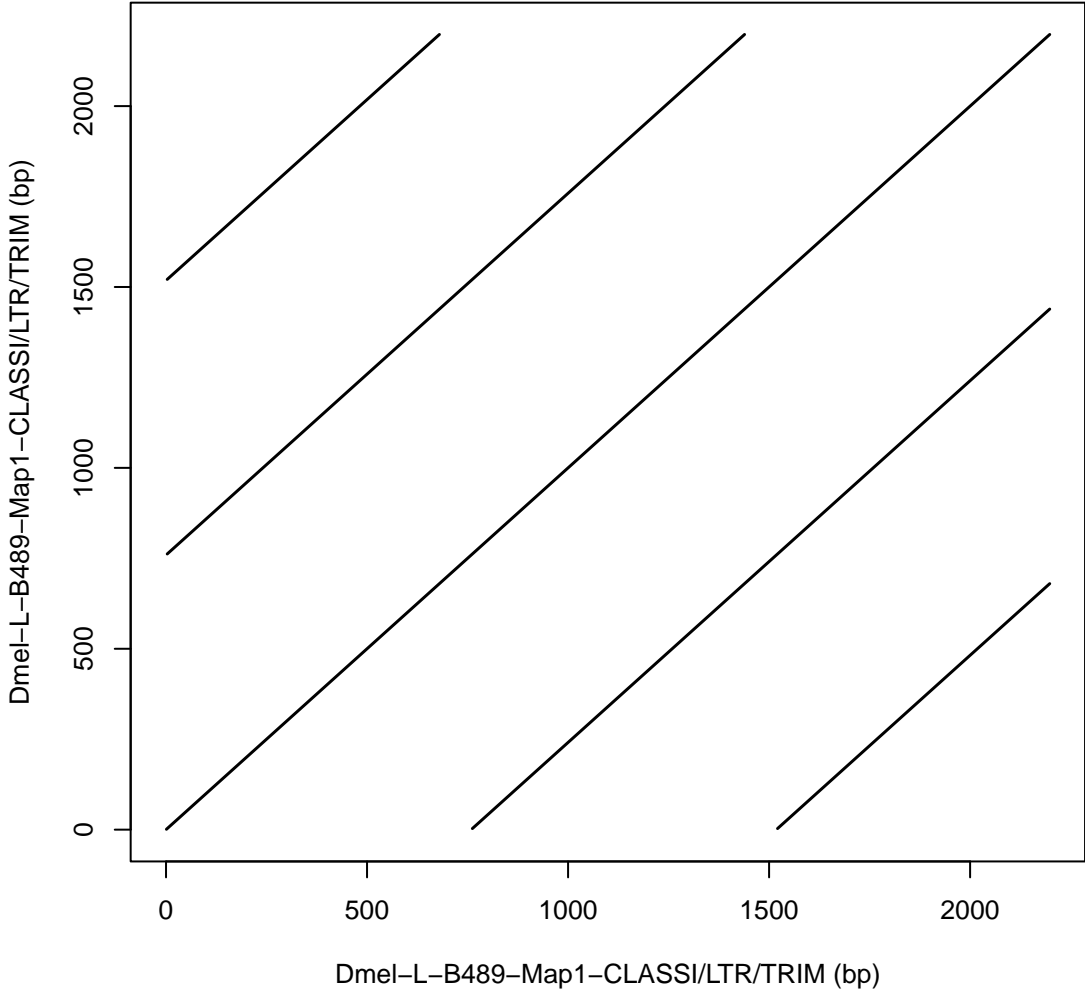consensus size: 2198bp; fragments: 5; full length: 1 (>=1978.2bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected