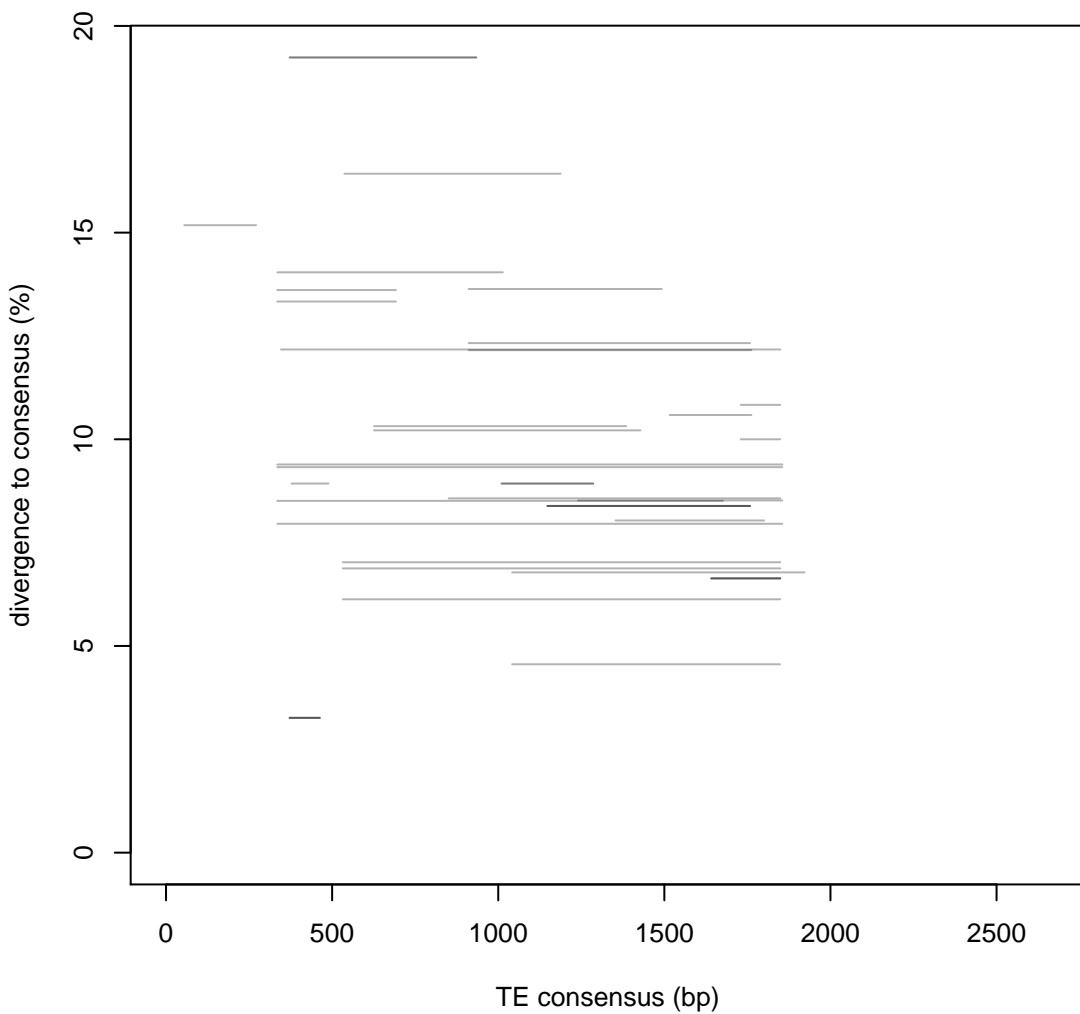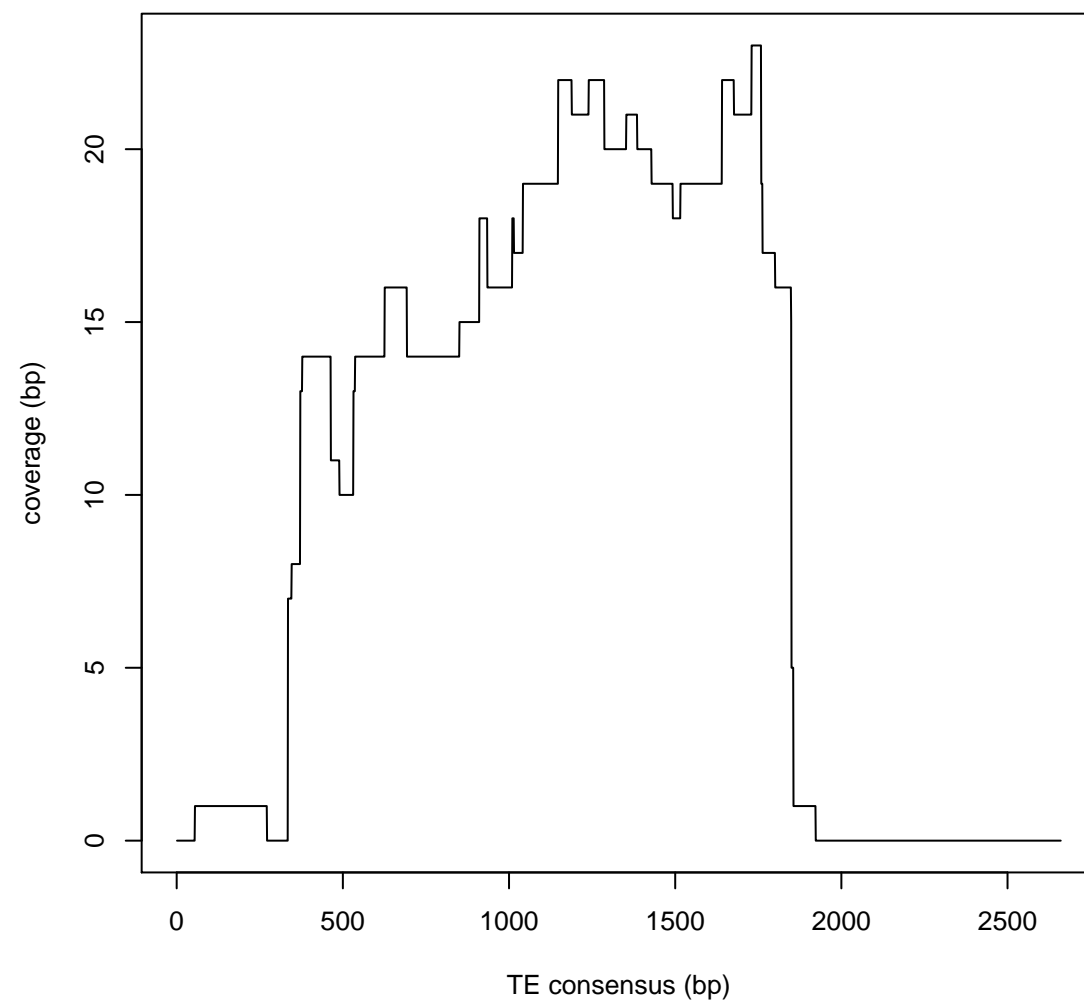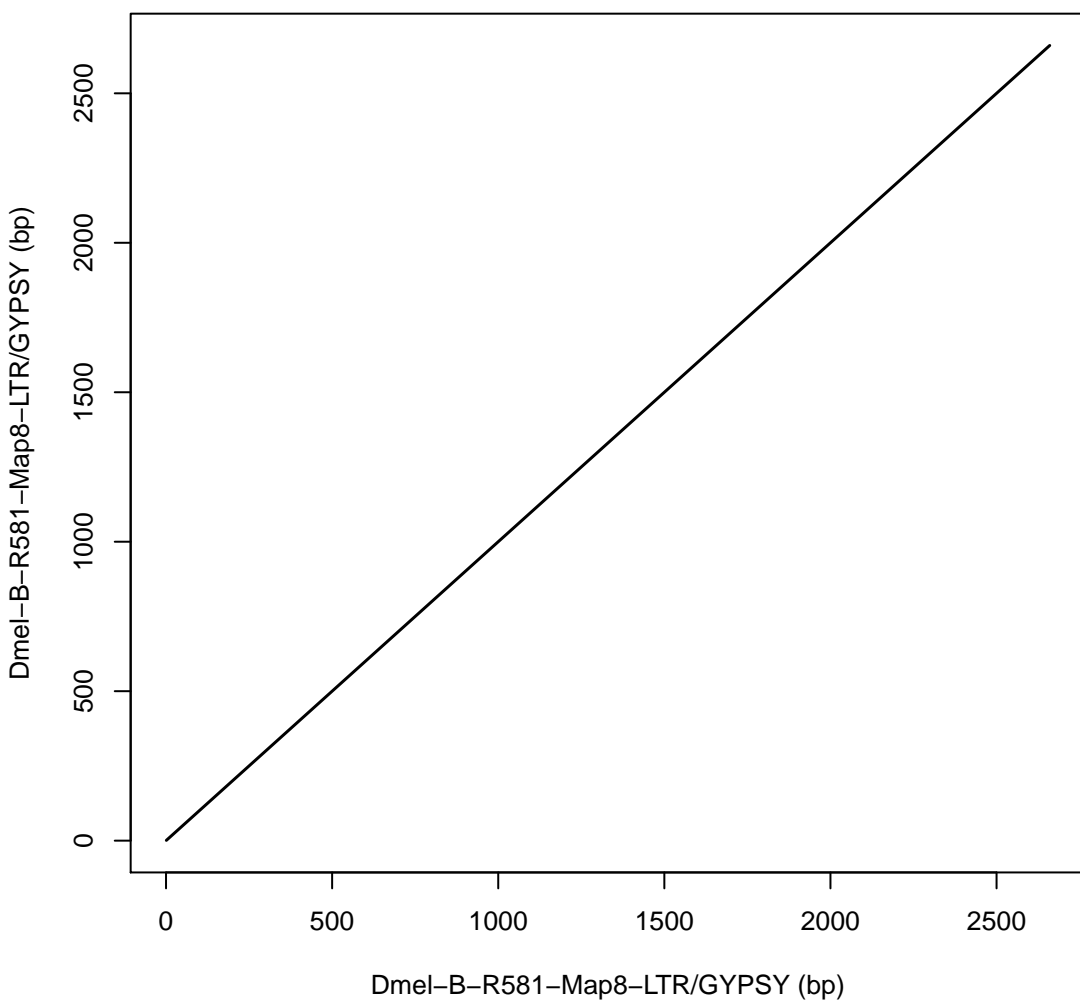**TE: Dmel−B−R581−Map8−LTR/GYPSY**
consensus size: 2660bp; fragments: 40; full length: 0 (>=2394bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

TOM_pol LTR/Gypsy