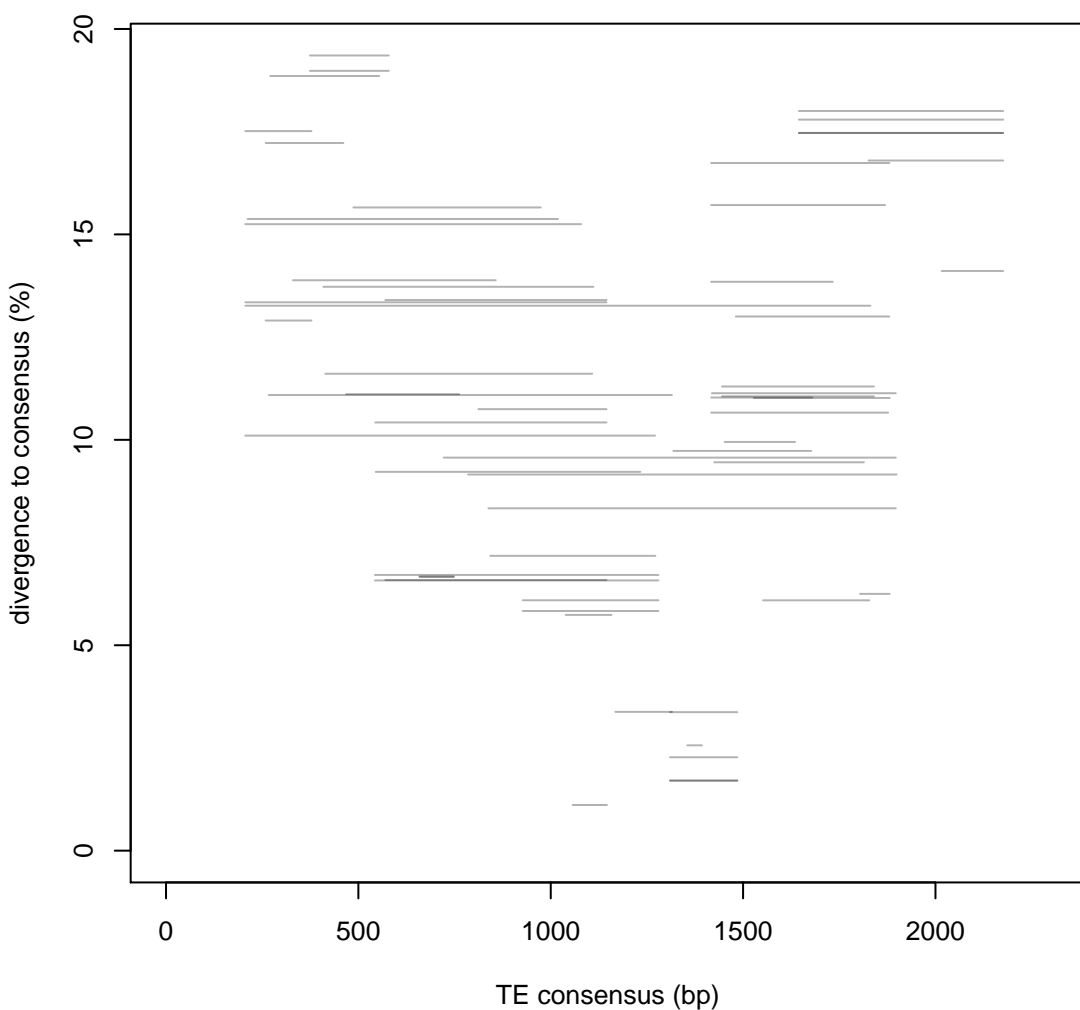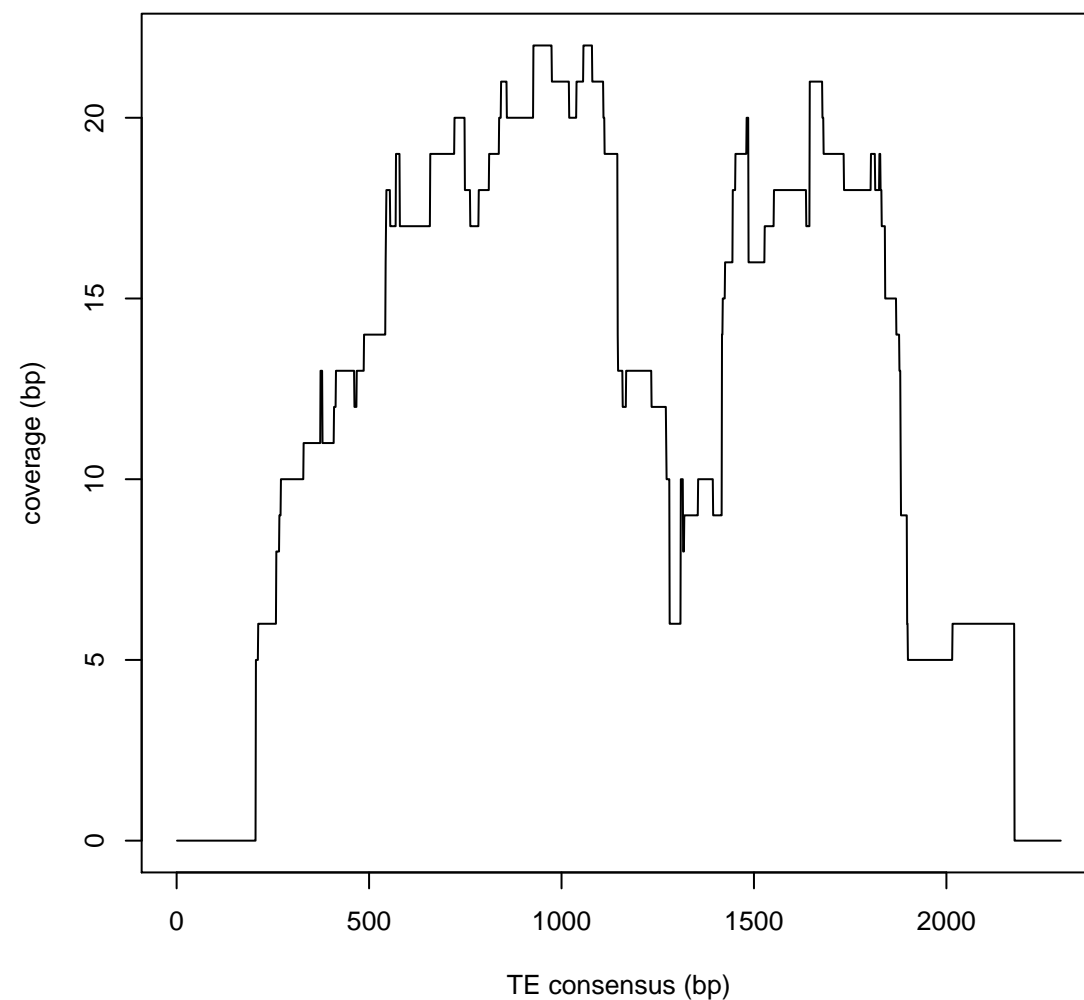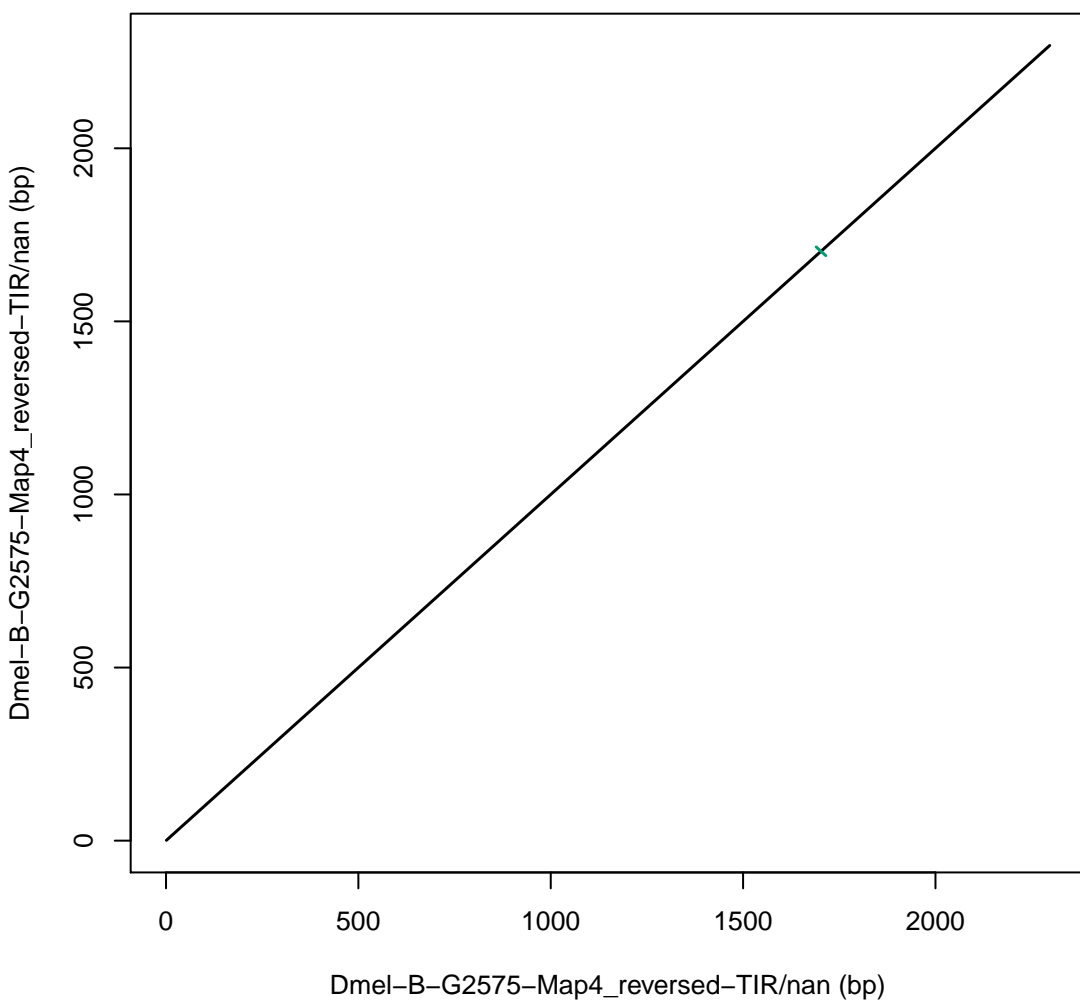**TE: Dmel−B−G2575−Map4_reversed−TIR/nan**
consensus size: 2297bp; fragments: 61; full length: 0 (>=2067.3bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

JASIMODO_pol LTR/Gypsy