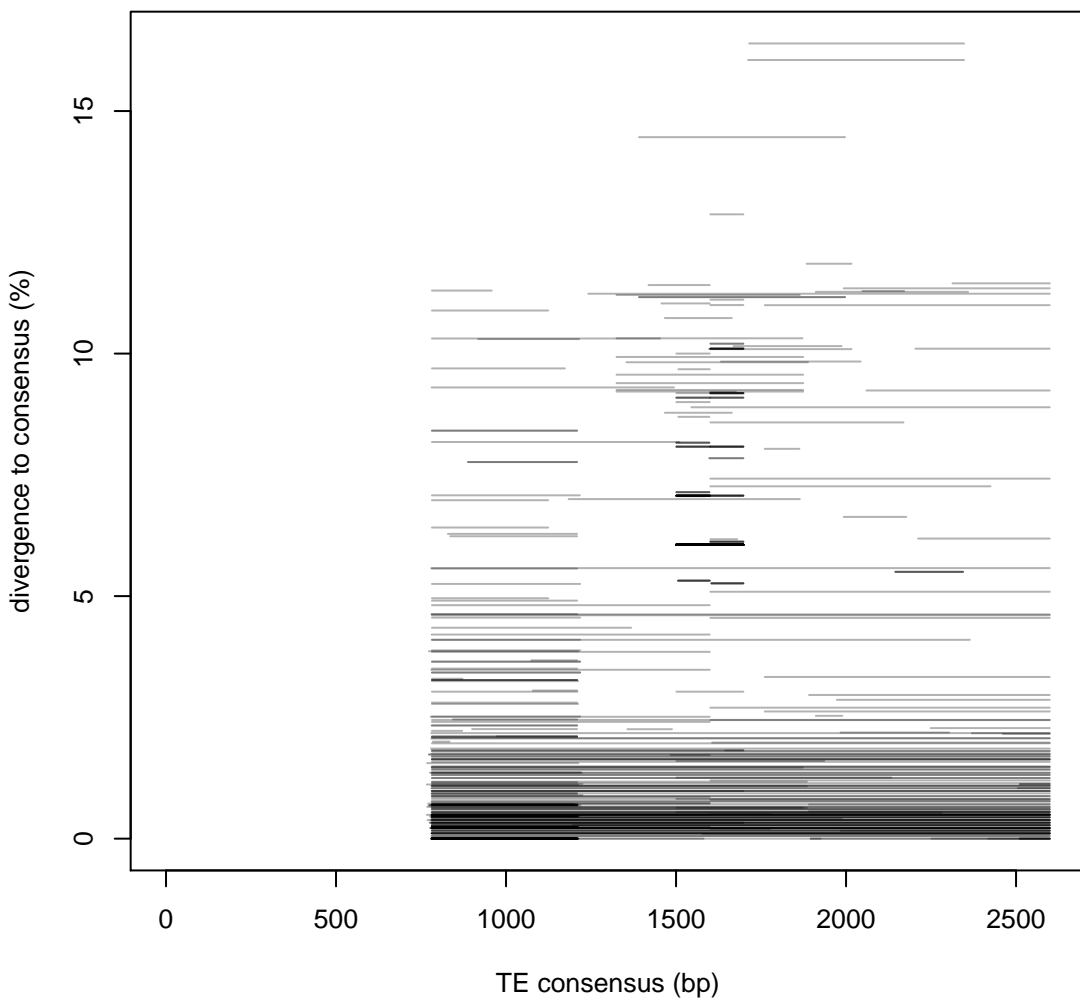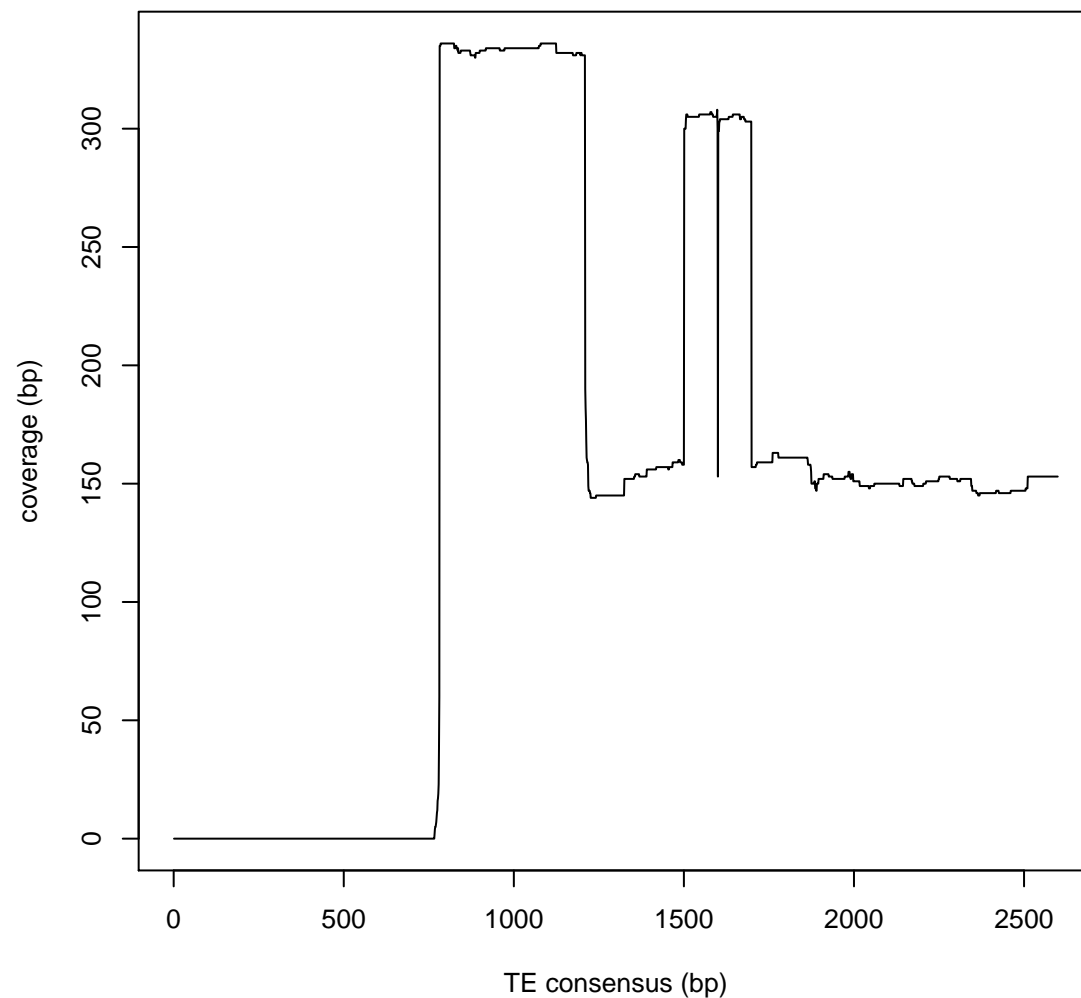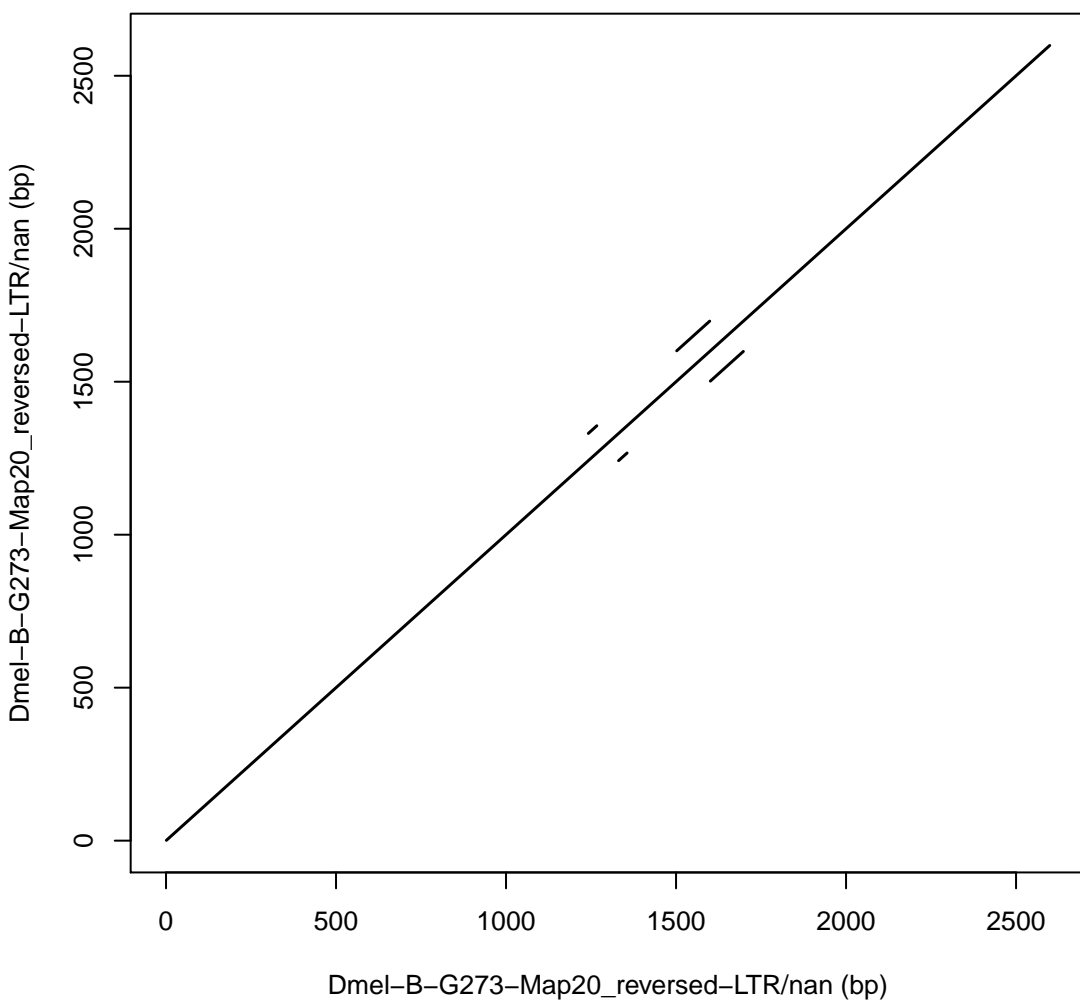**TE: Dmel−B−G273−Map20_reversed−LTR/nan**
consensus size: 2599bp; fragments: 711; full length: 0 (>=2339.1bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

ROO_pol LTR/Pao