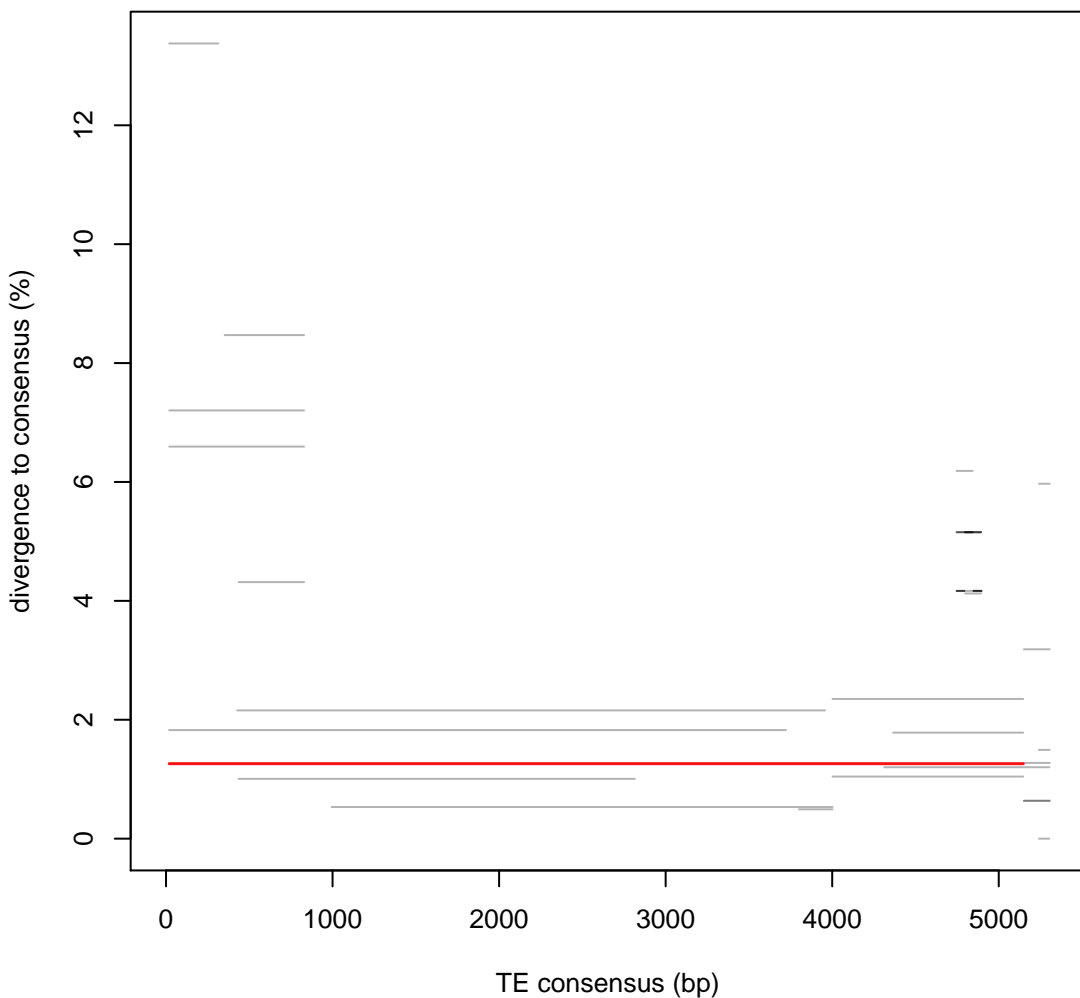
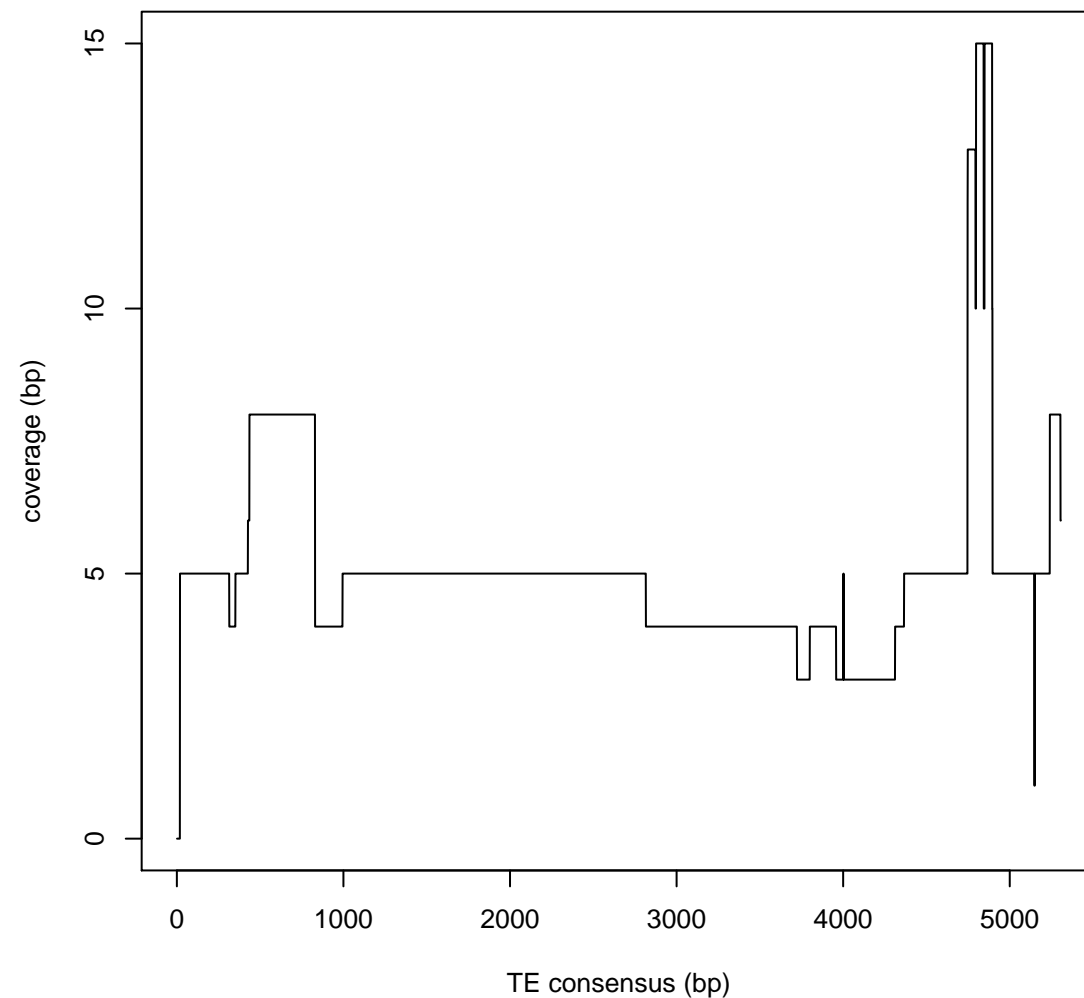


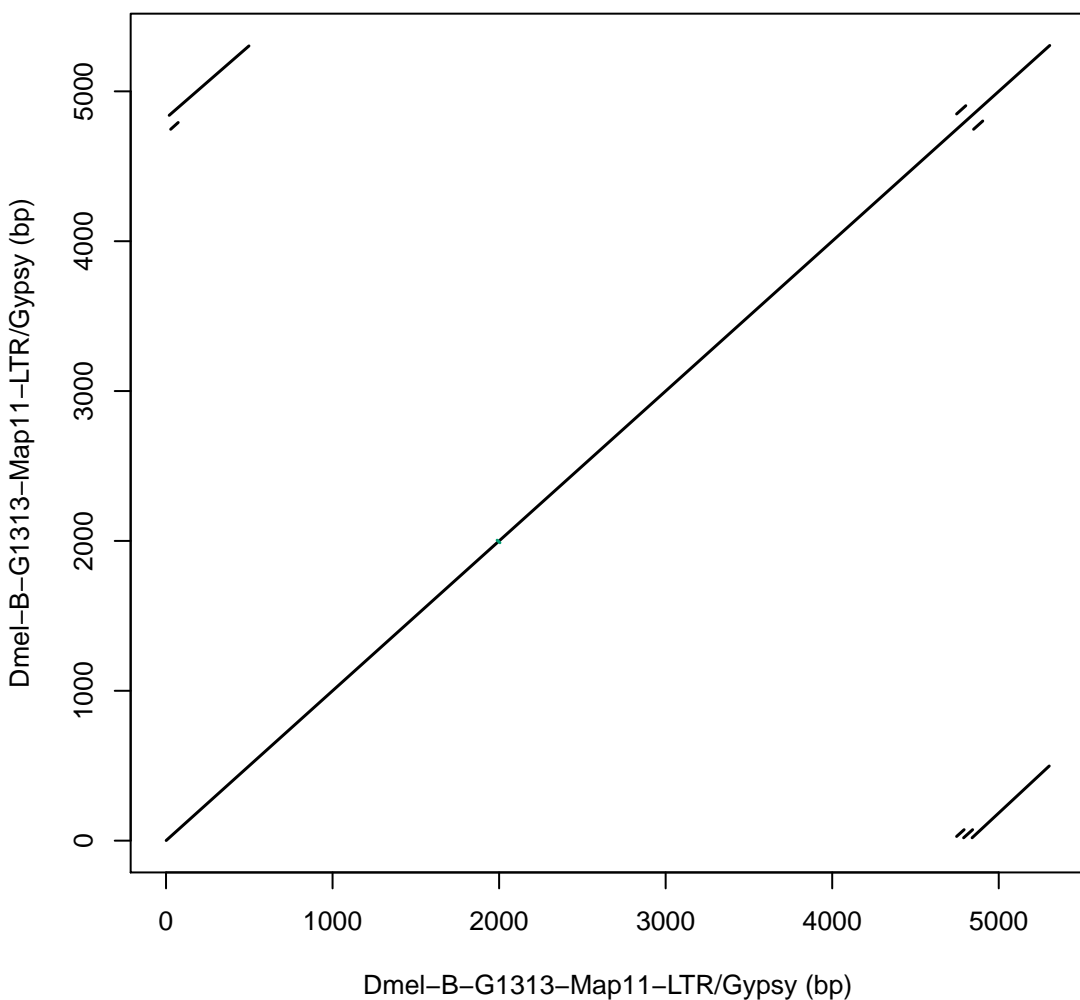
TE: Dmel-B-G1313-Map11-LTR/Gypsy
consensus size: 5306bp; fragments: 40; full length: 1 (≥ 4775.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

