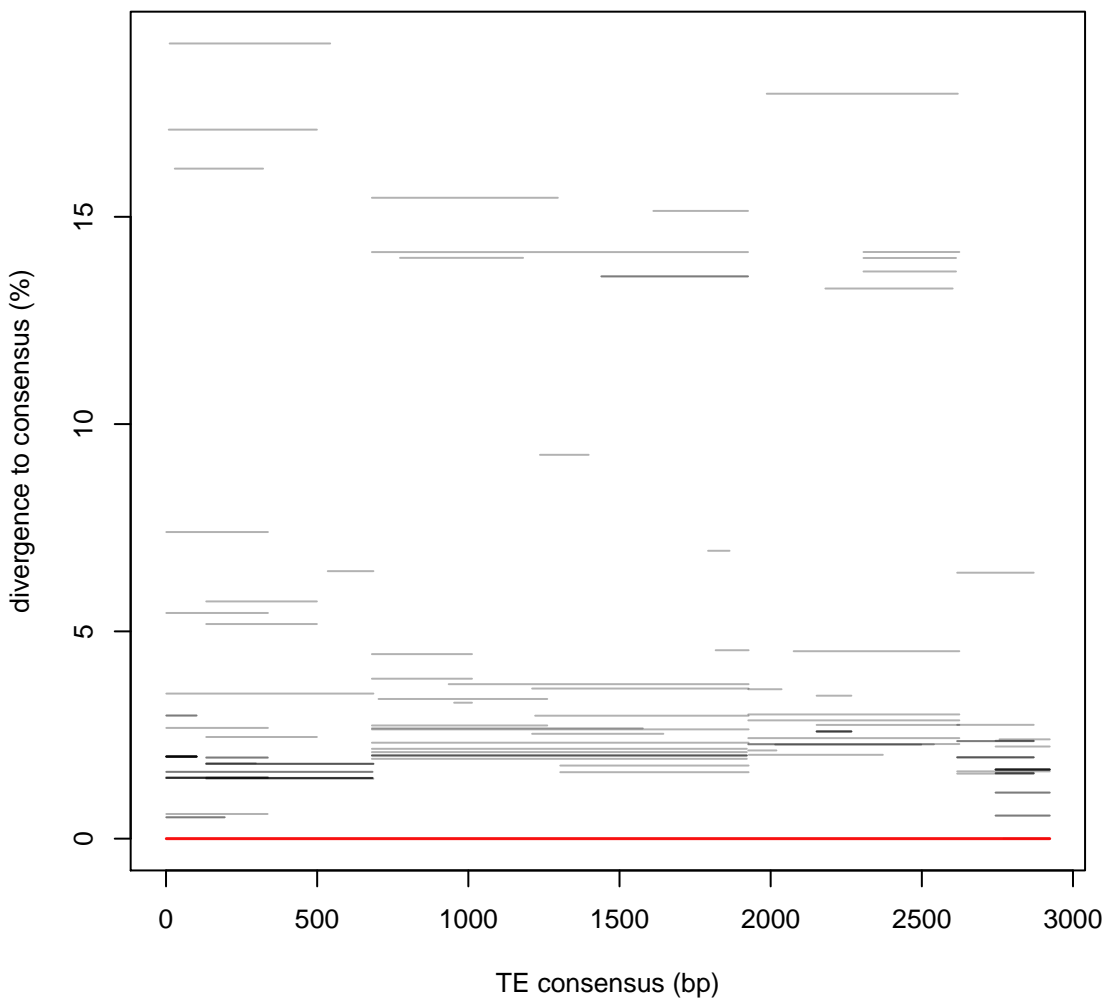
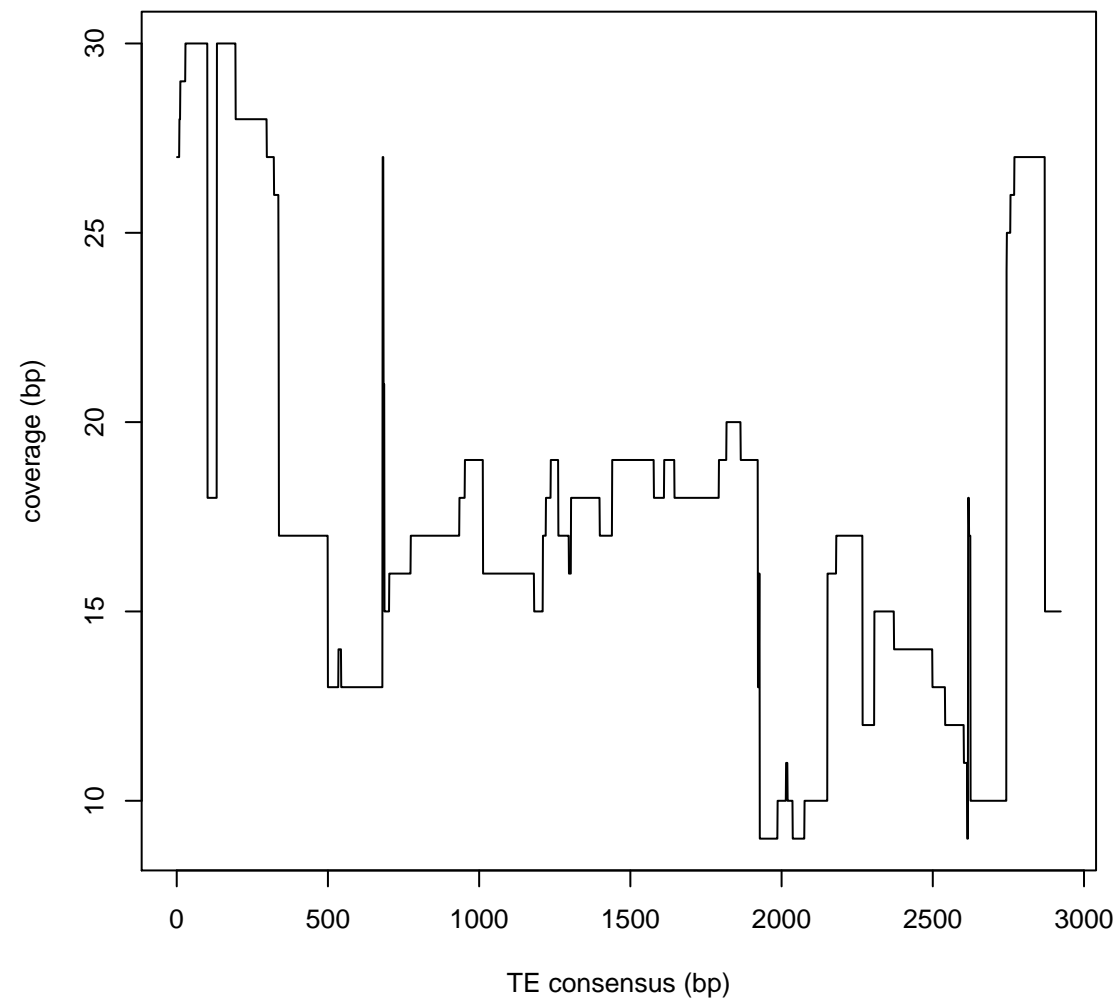


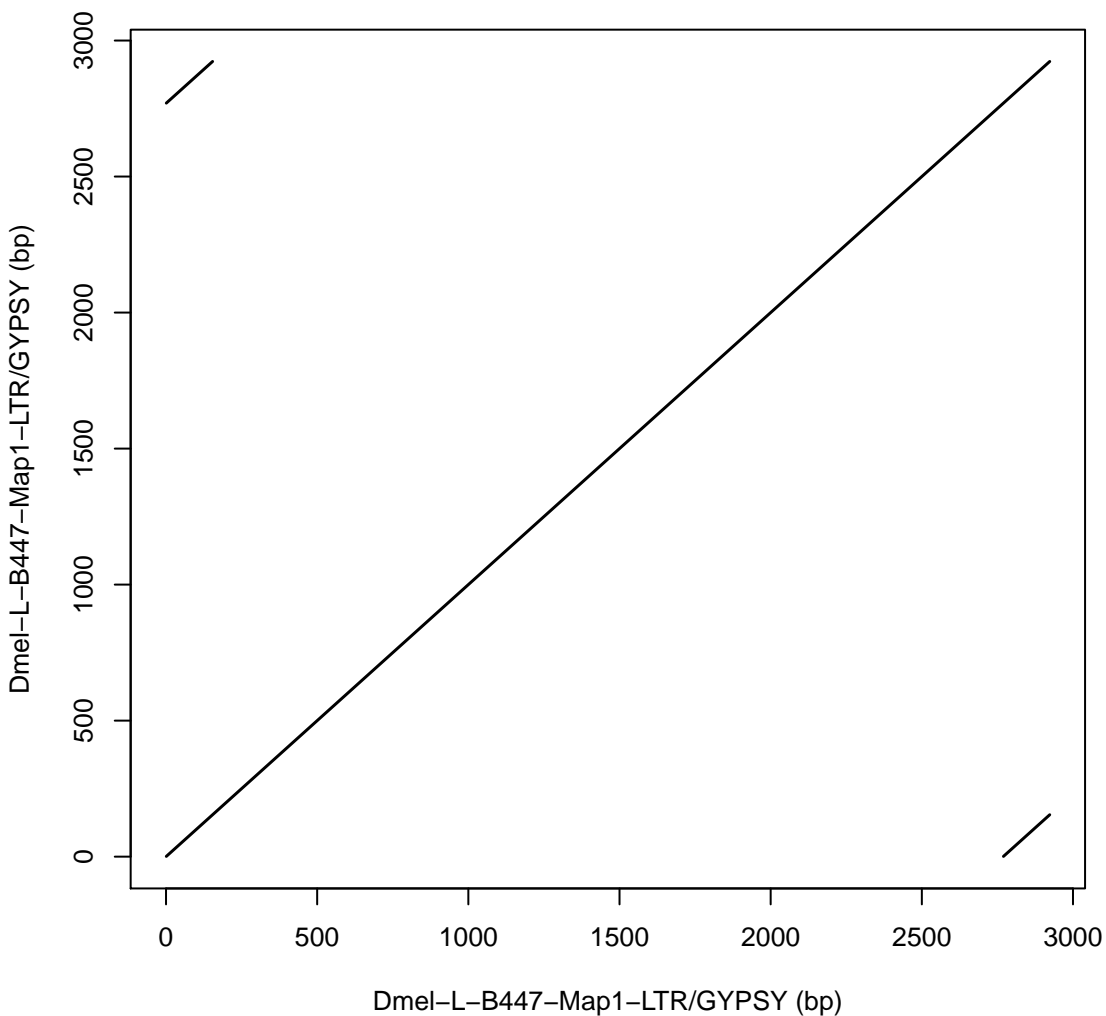
TE: Dmel-L-B447-Map1-LTR/GYPSY
consensus size: 2923bp; fragments: 119; full length: 1 (≥ 2630.7 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

