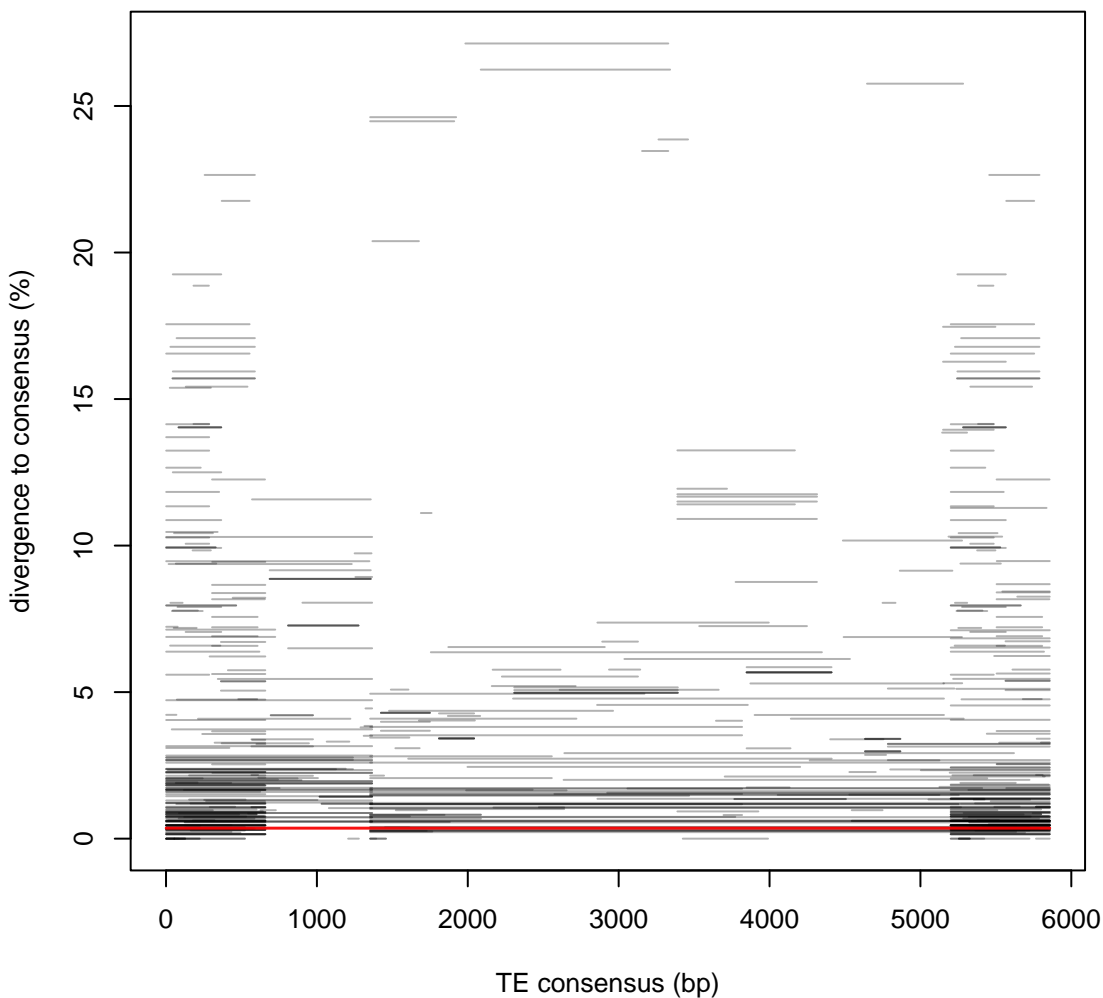
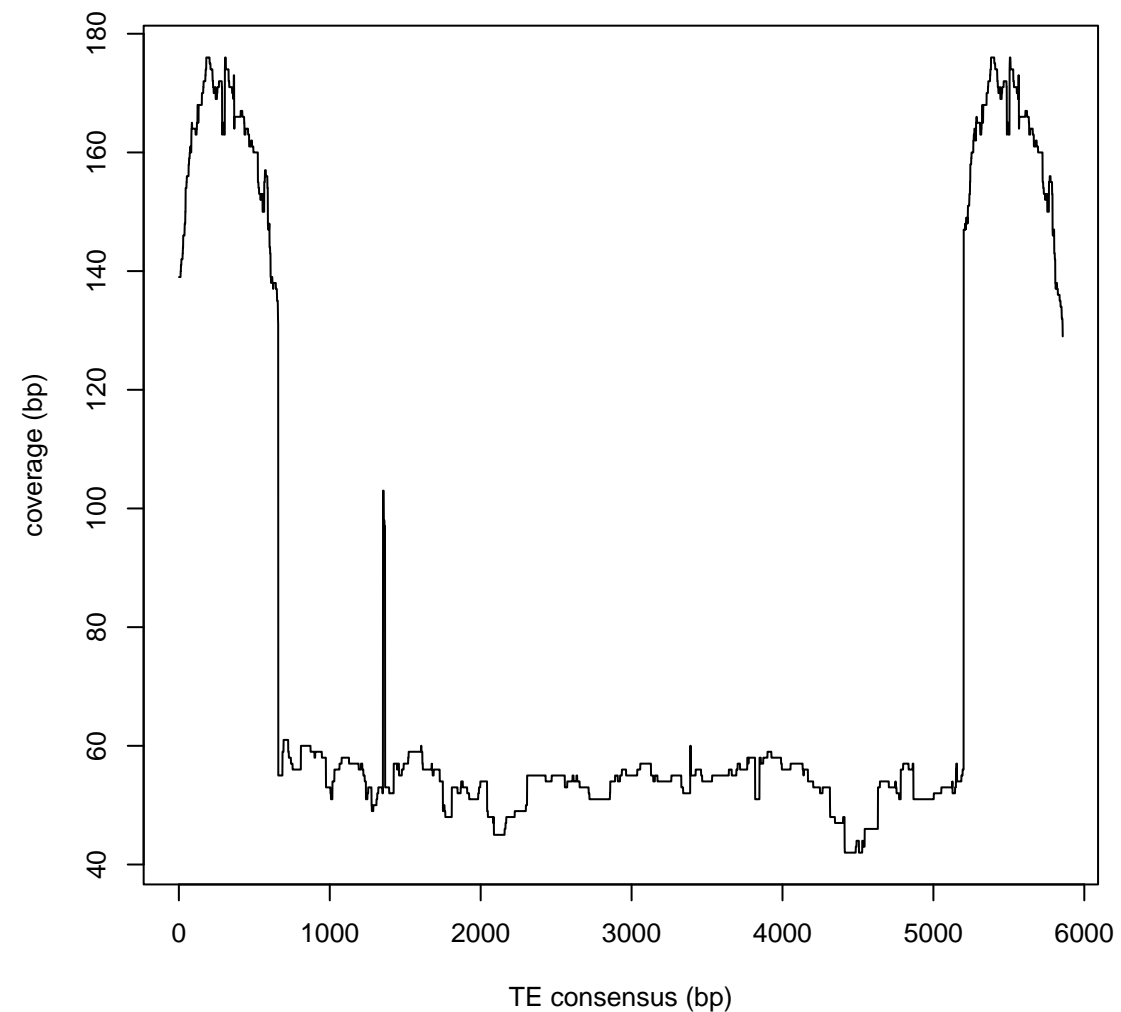


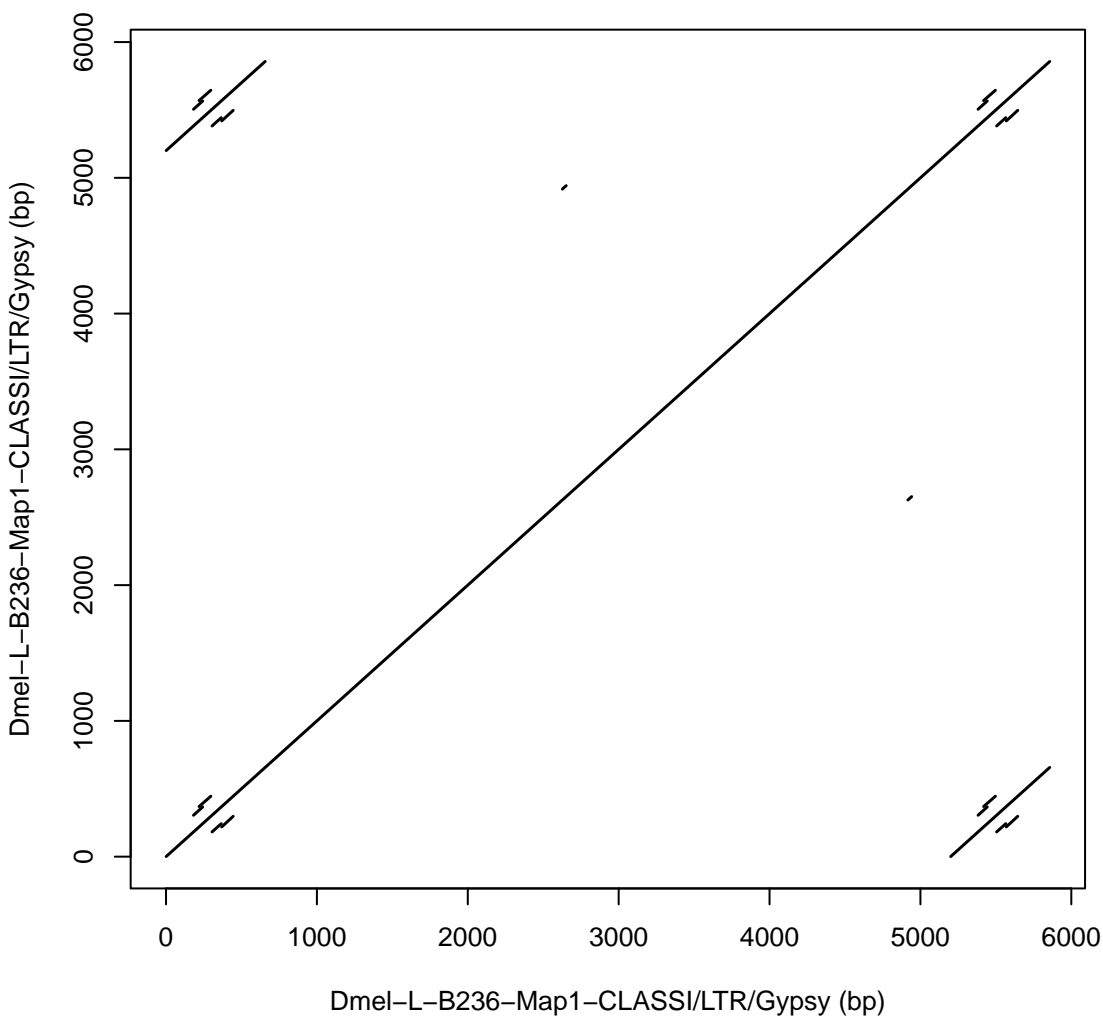
TE: Dmel-L-B236-Map1-CLASSI/LTR/Gypsy
consensus size: 5857bp; fragments: 593; full length: 1 (≥ 5271.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

