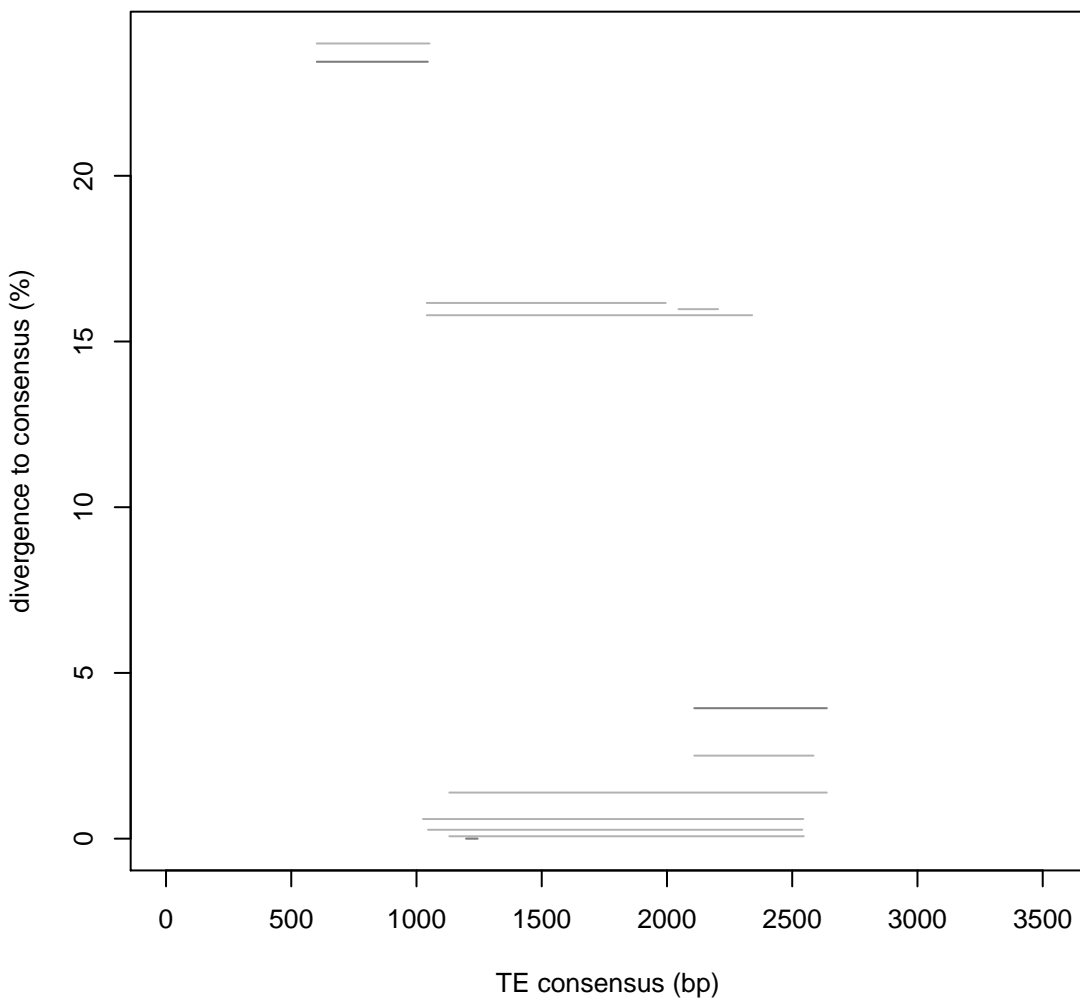
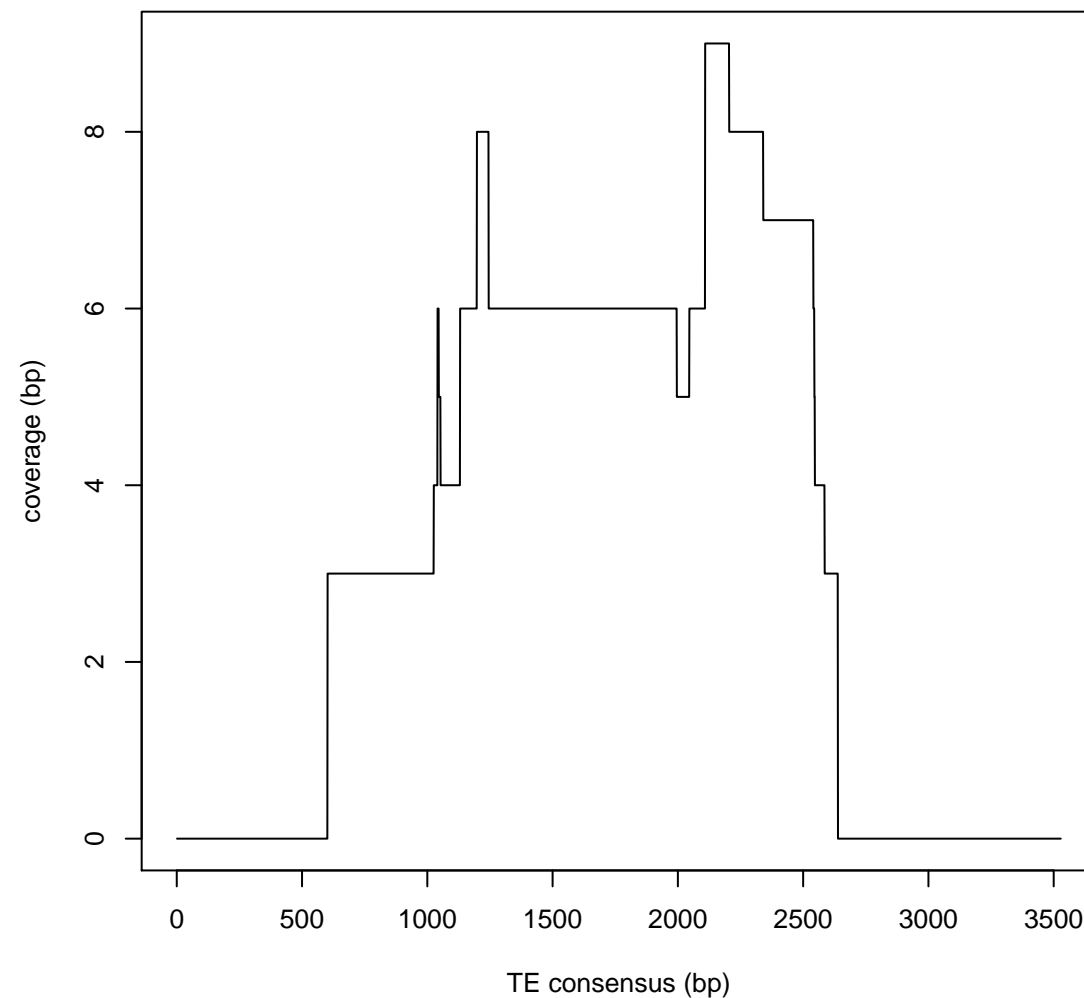


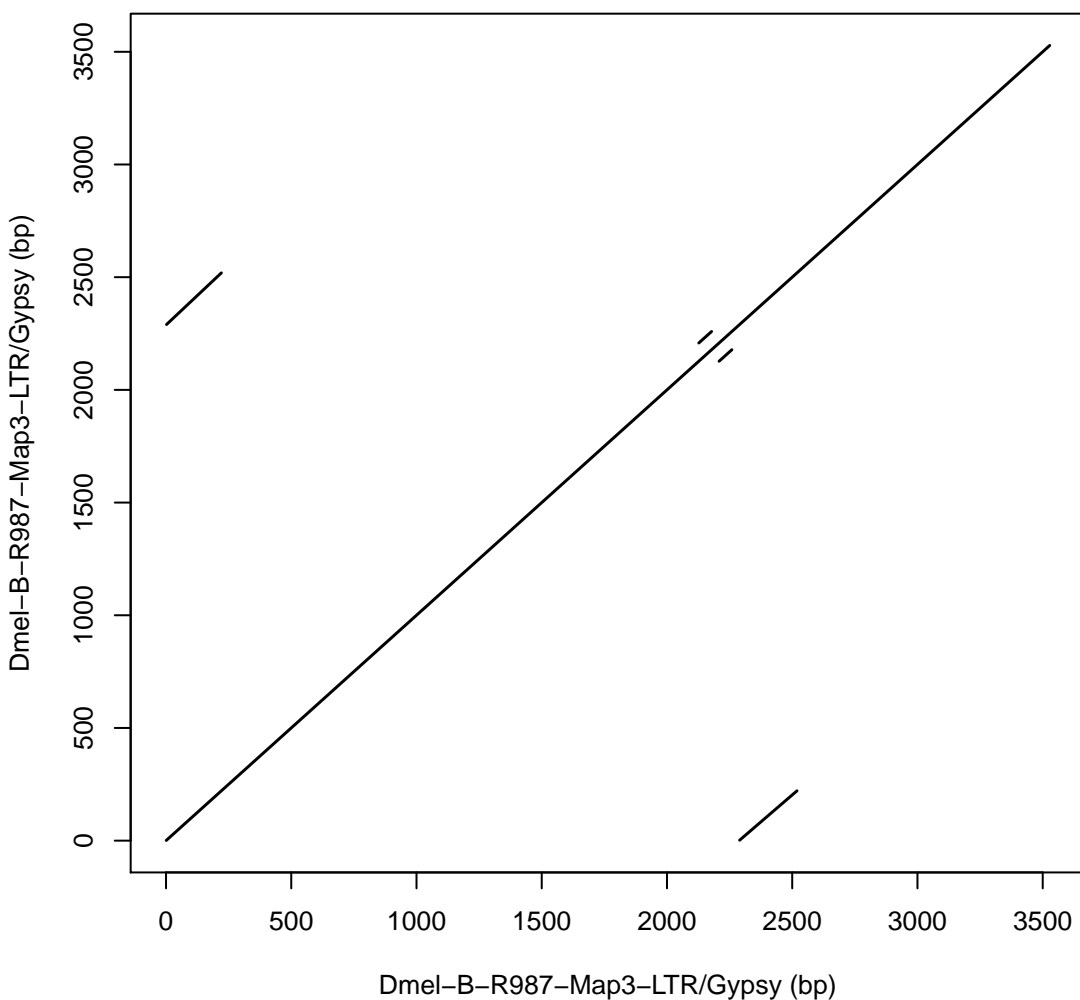
TE: Dmel-B-R987-Map3-LTR/Gypsy
consensus size: 3528bp; fragments: 15; full length: 0 (≥ 3175.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

