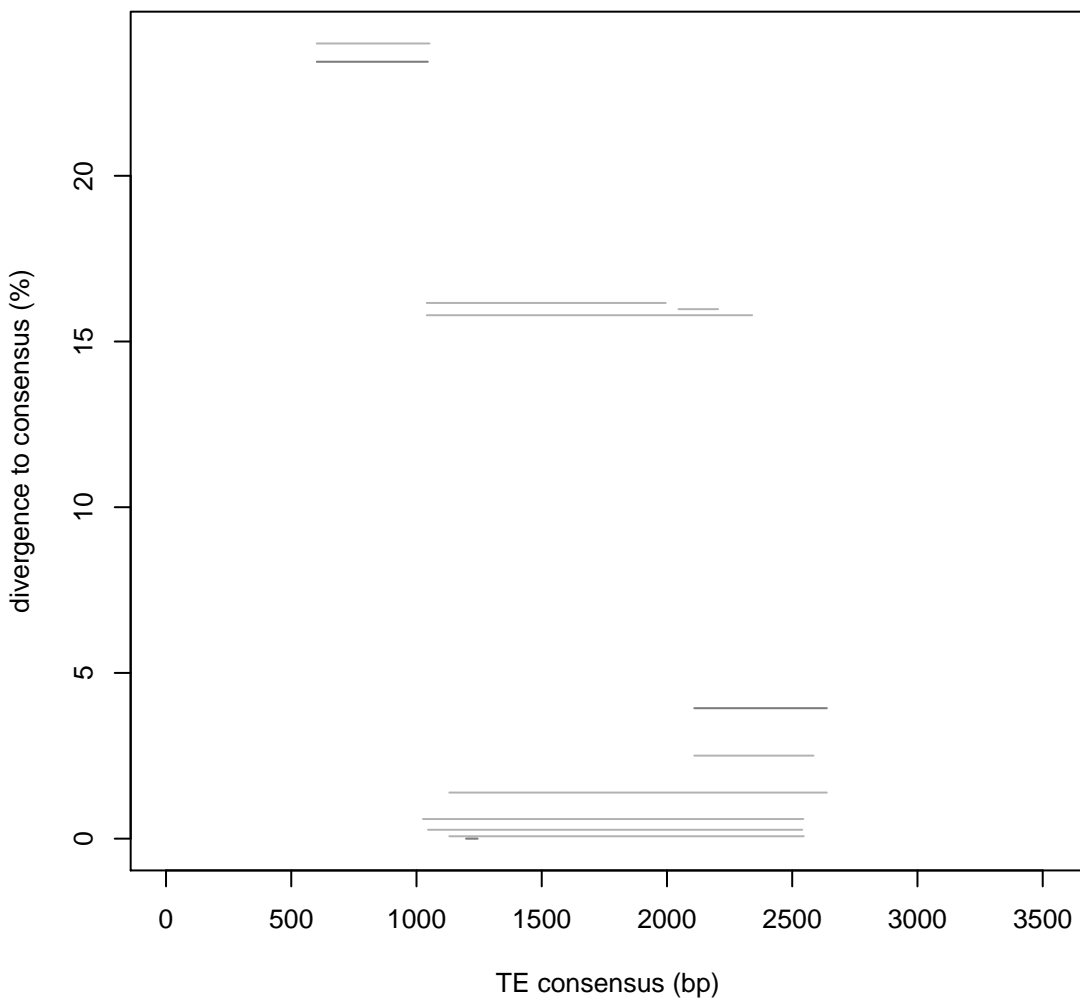
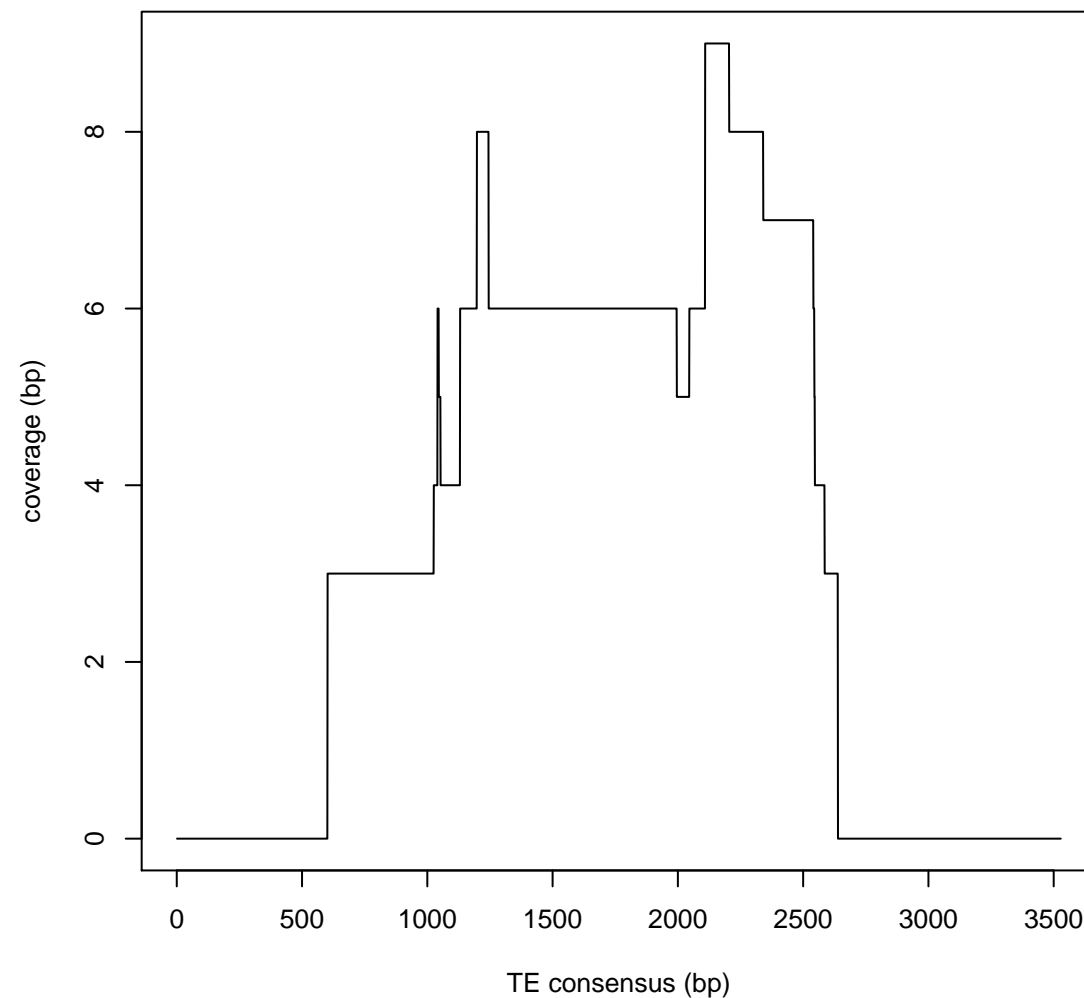


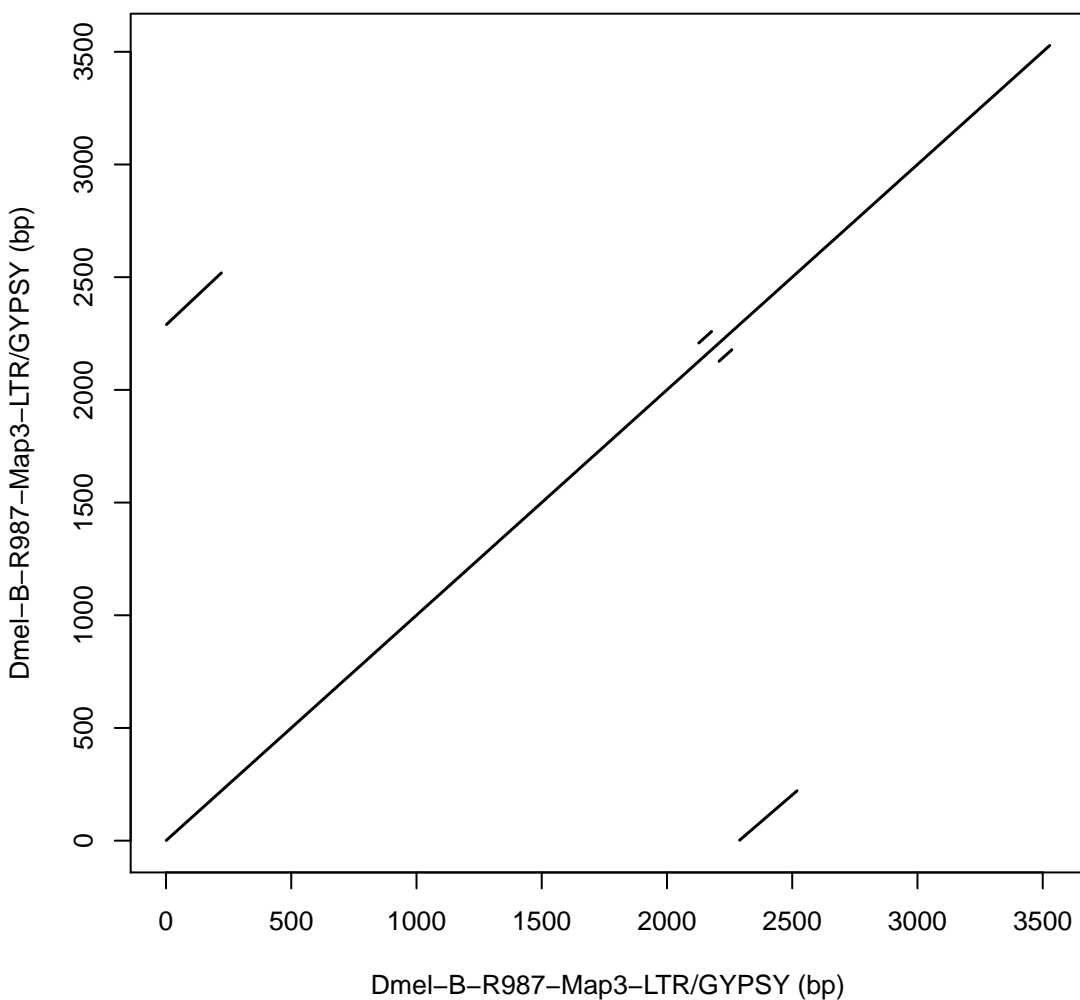
TE: Dmel-B-R987-Map3-LTR/GYPSY
consensus size: 3528bp; fragments: 15; full length: 0 (>=3175.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

