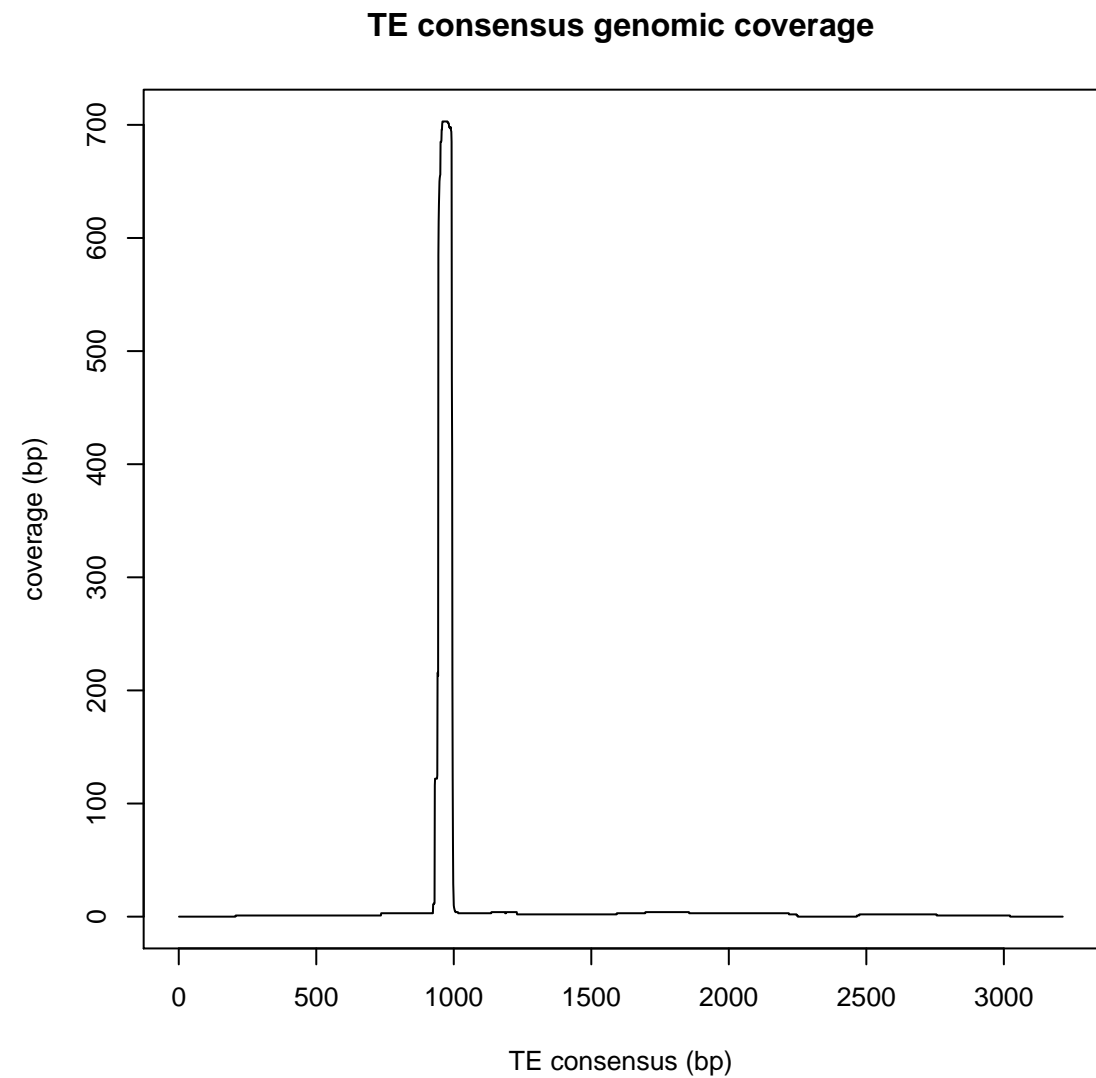
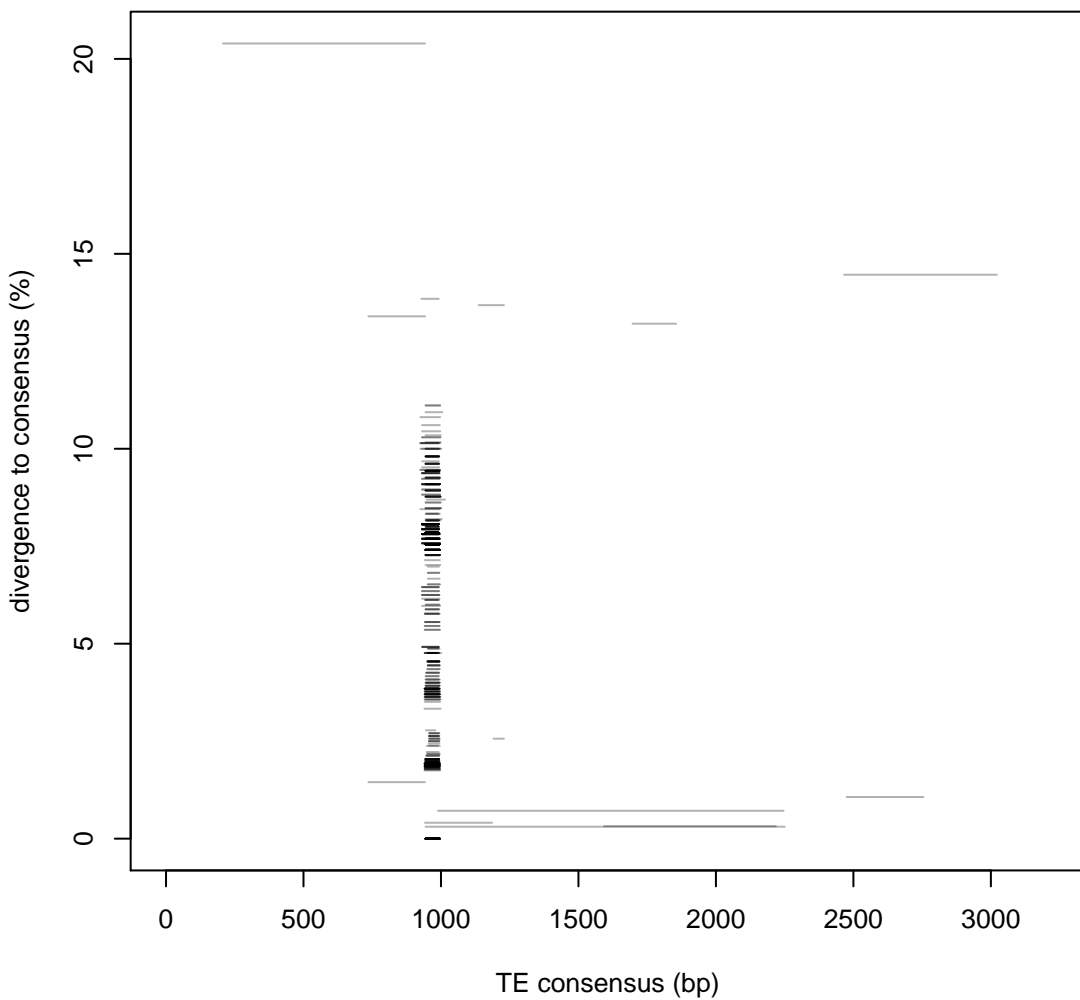
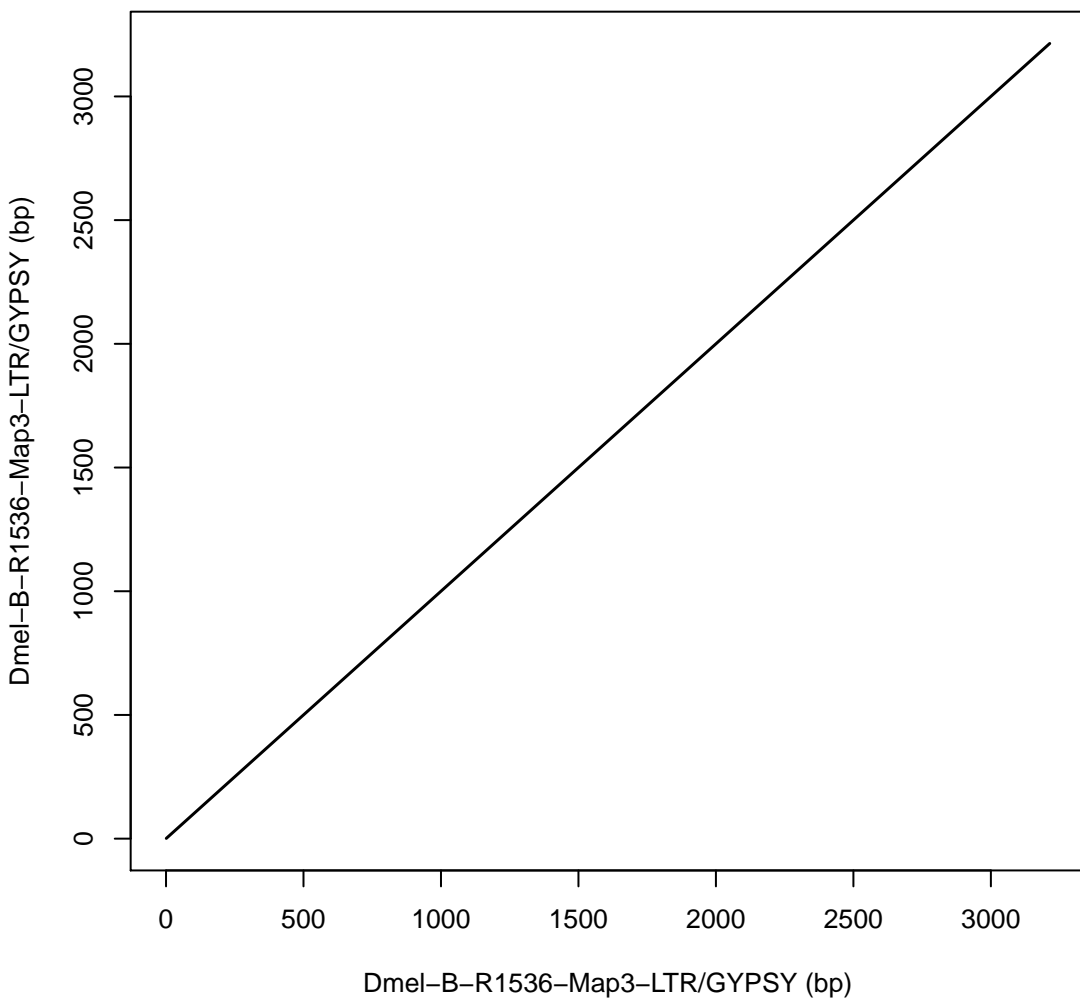


TE: Dmel-B-R1536-Map3-LTR/GYPSY  
consensus size: 3214bp; fragments: 713; full length: 0 (>=2892.6bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

