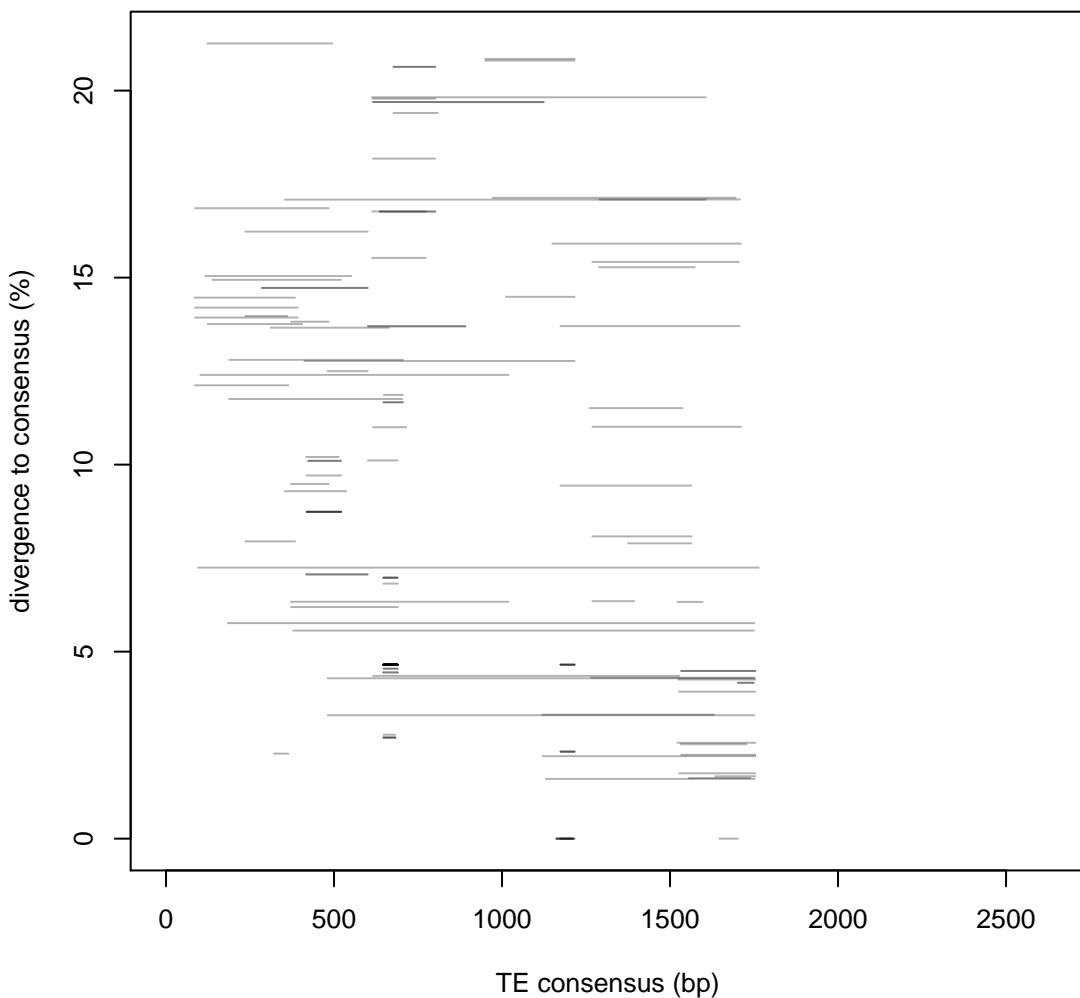
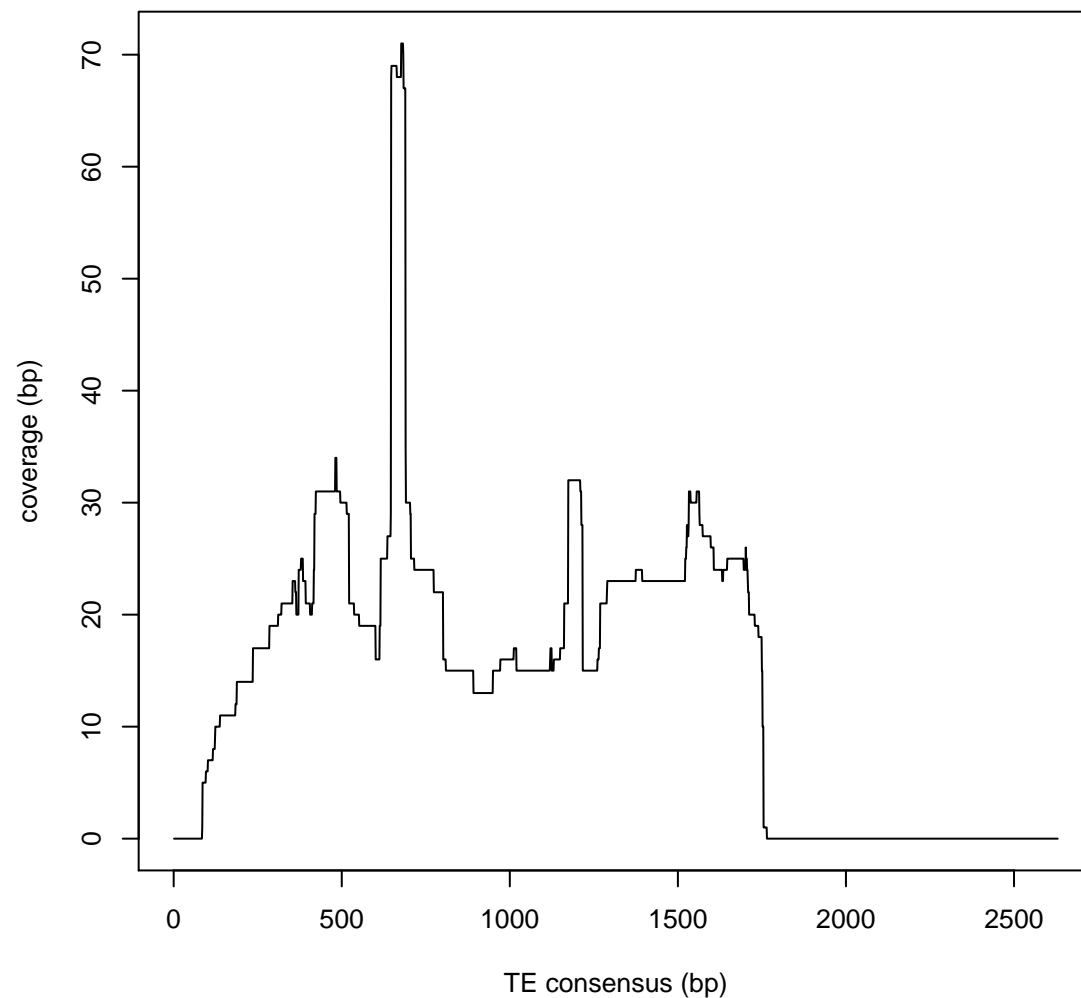


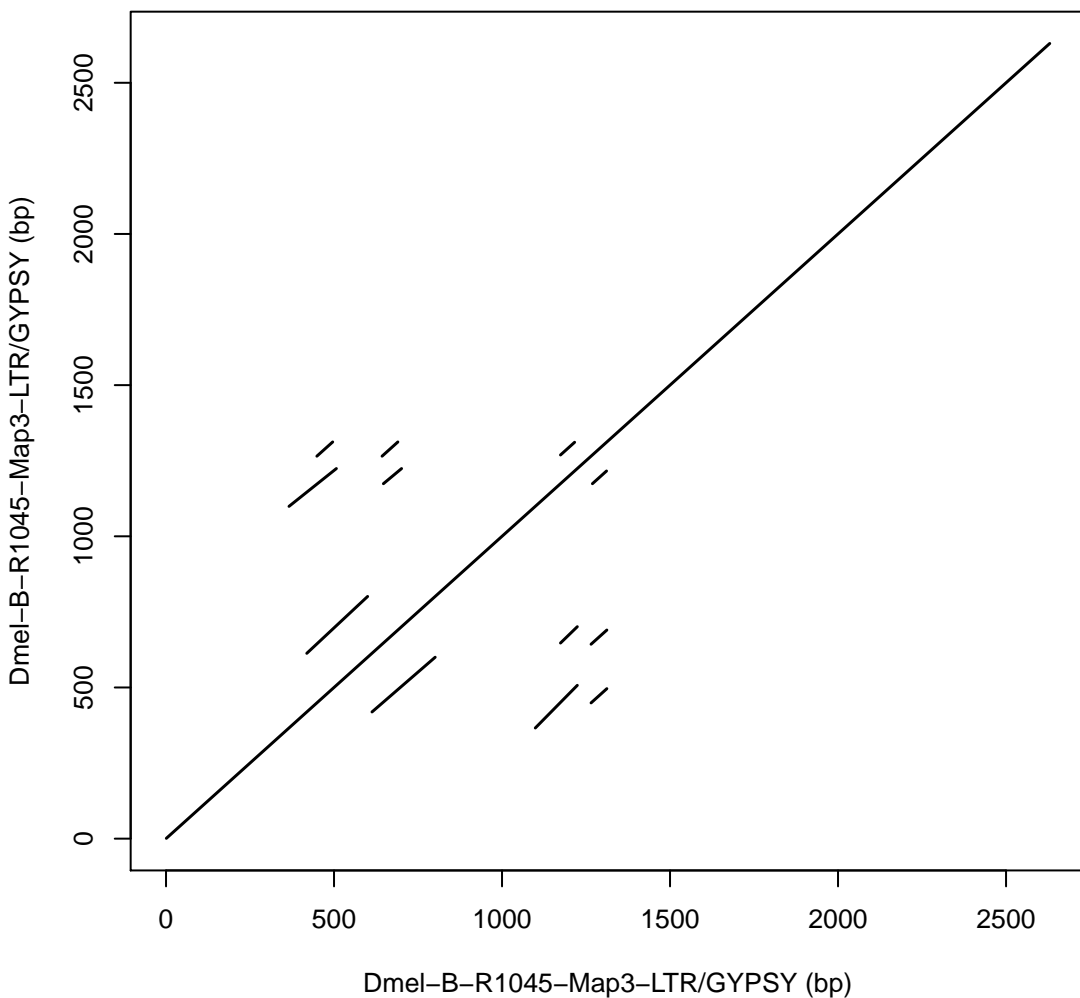
TE: Dmel-B-R1045-Map3-LTR/Gypsy  
consensus size: 2630bp; fragments: 147; full length: 0 (>=2367bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

