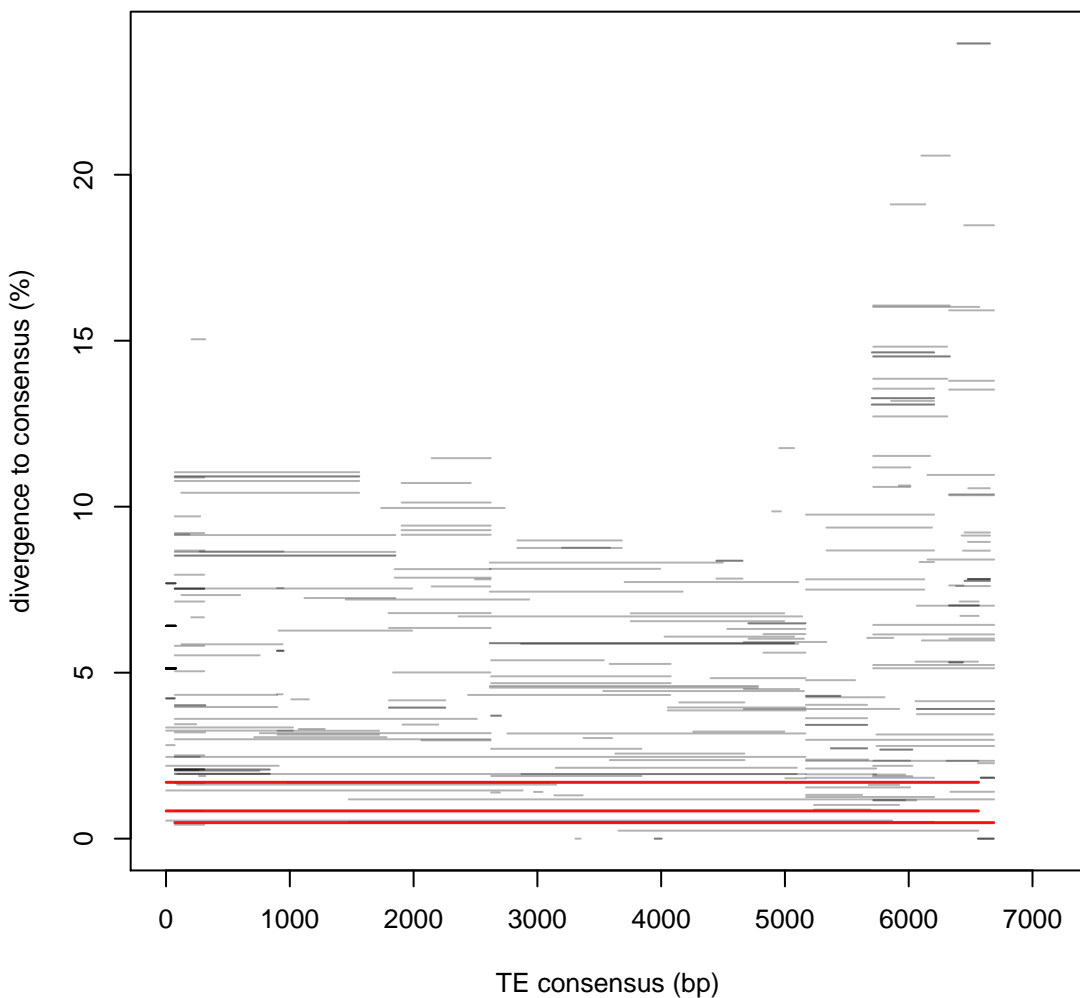
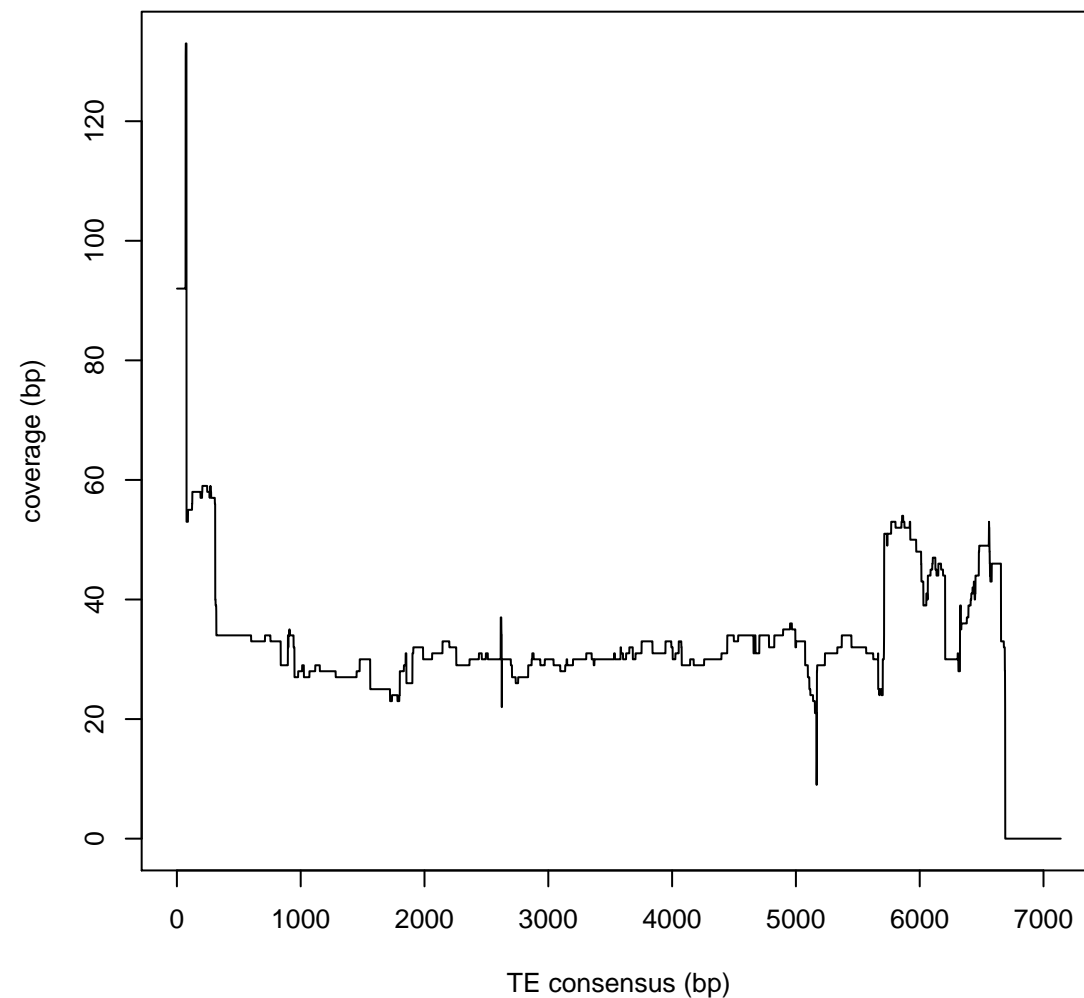


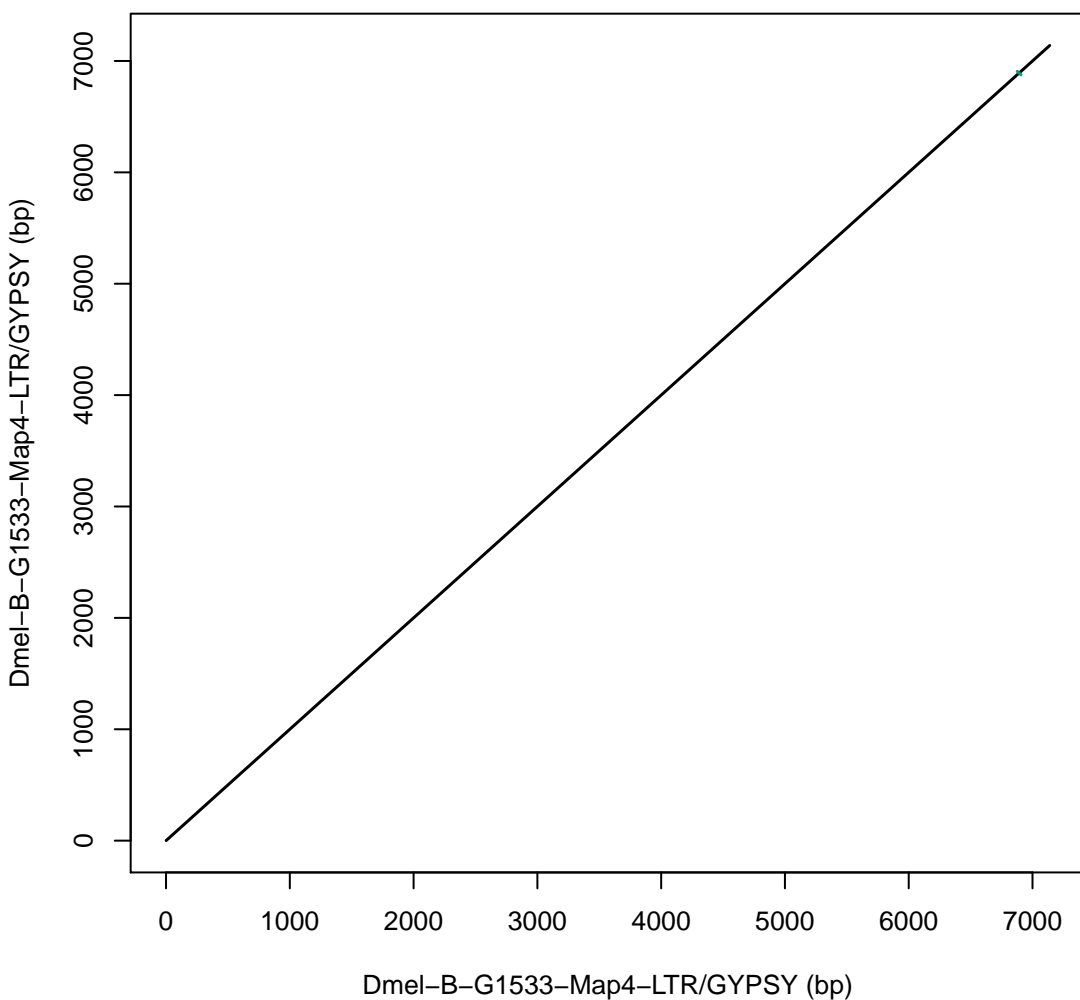
TE: Dmel-B-G1533-Map4-LTR/GYPSY  
consensus size: 7139bp; fragments: 350; full length: 3 (>=6425.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

