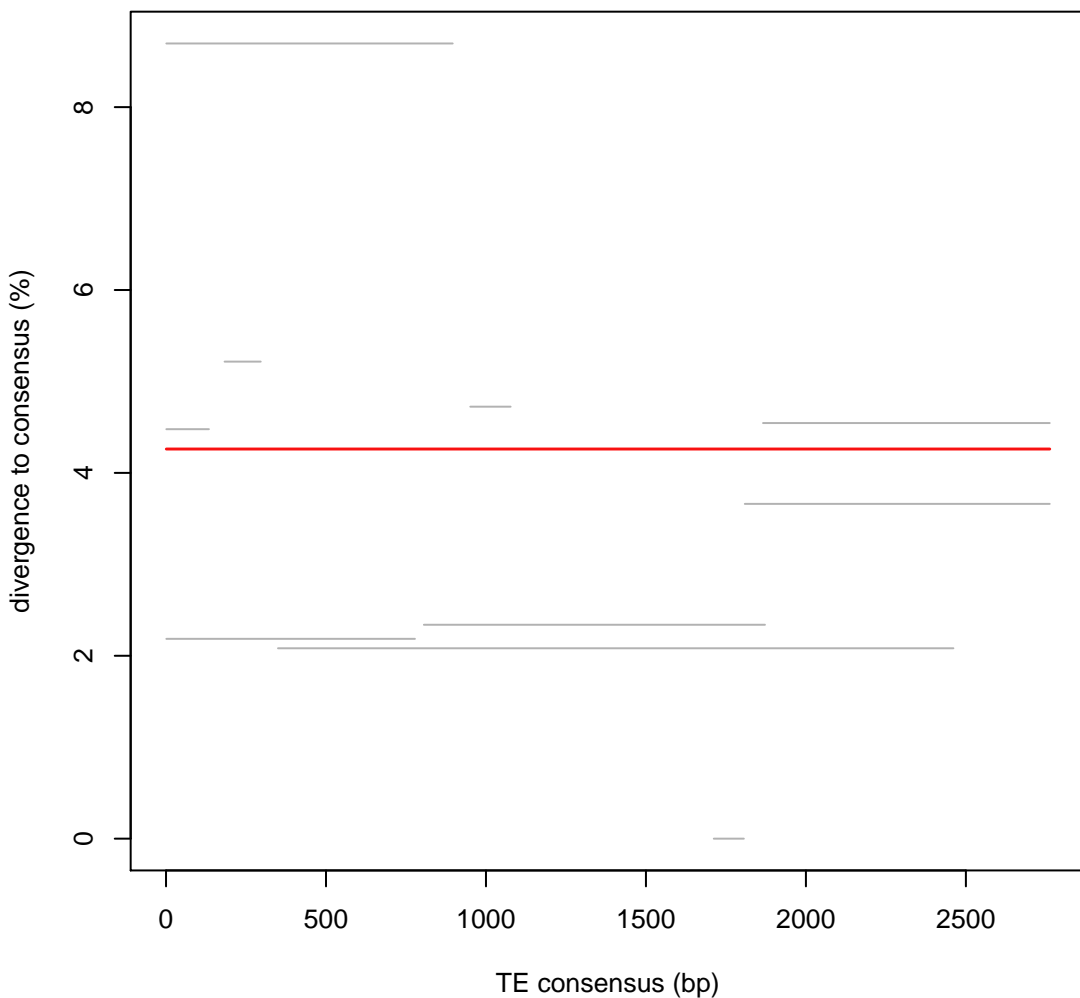
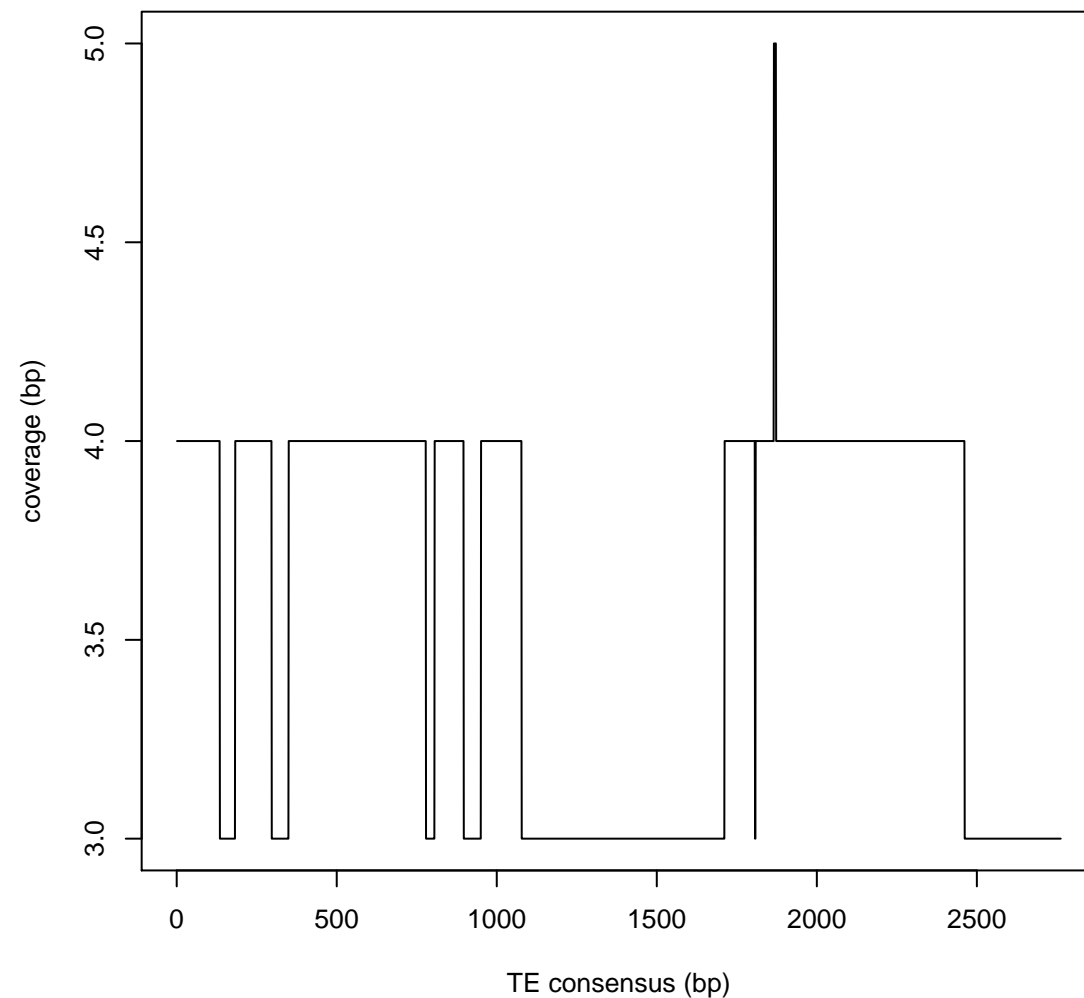


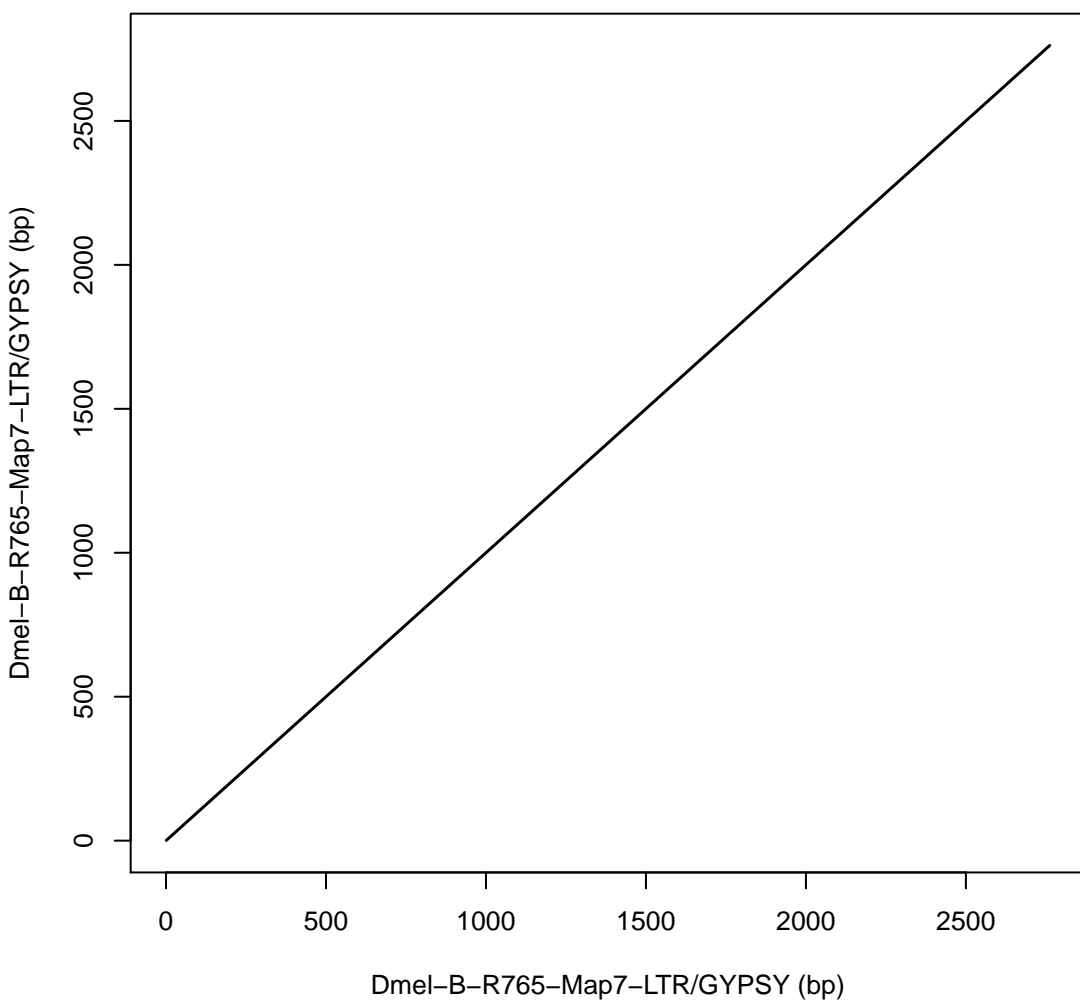
TE: Dmel-B-R765-Map7-LTR/GYPSY  
consensus size: 2762bp; fragments: 11; full length: 1 (>=2485.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

