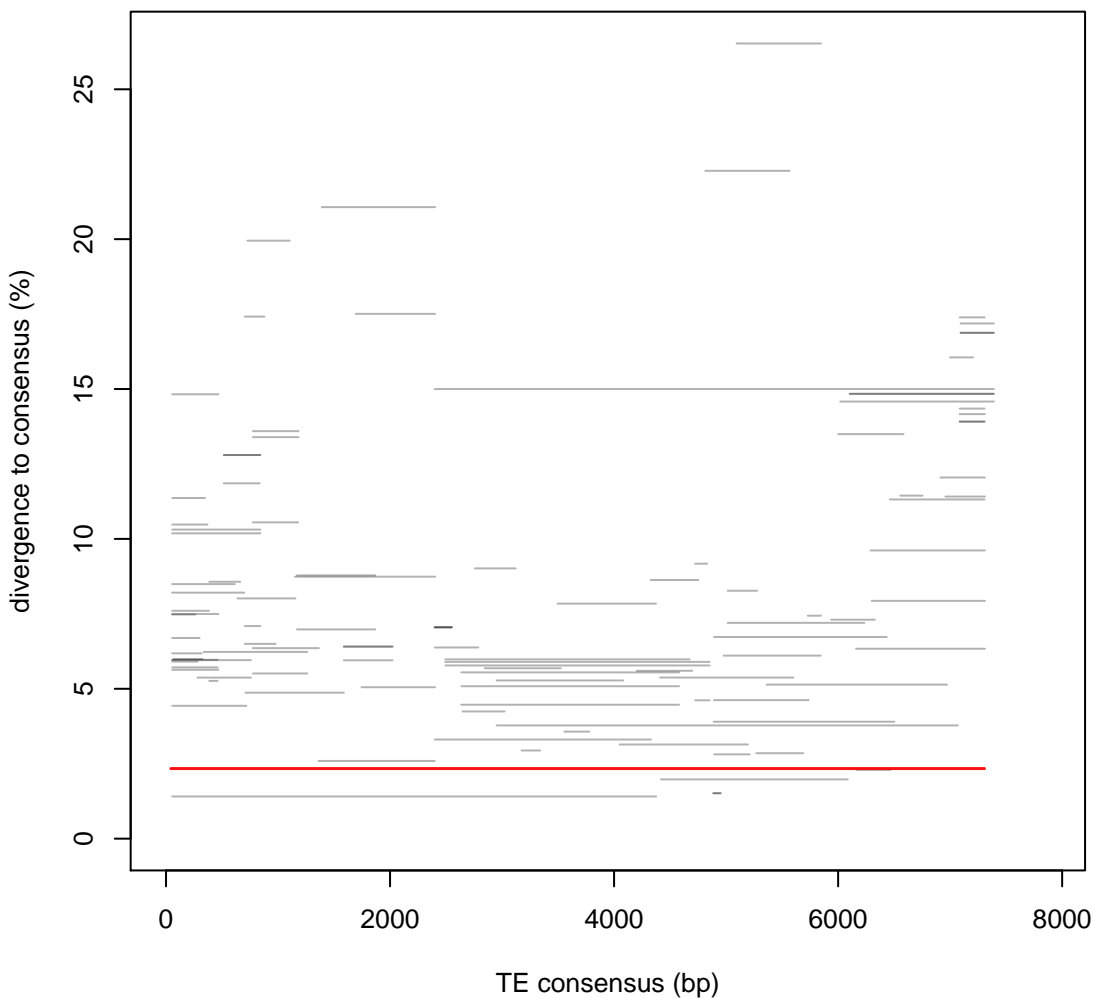
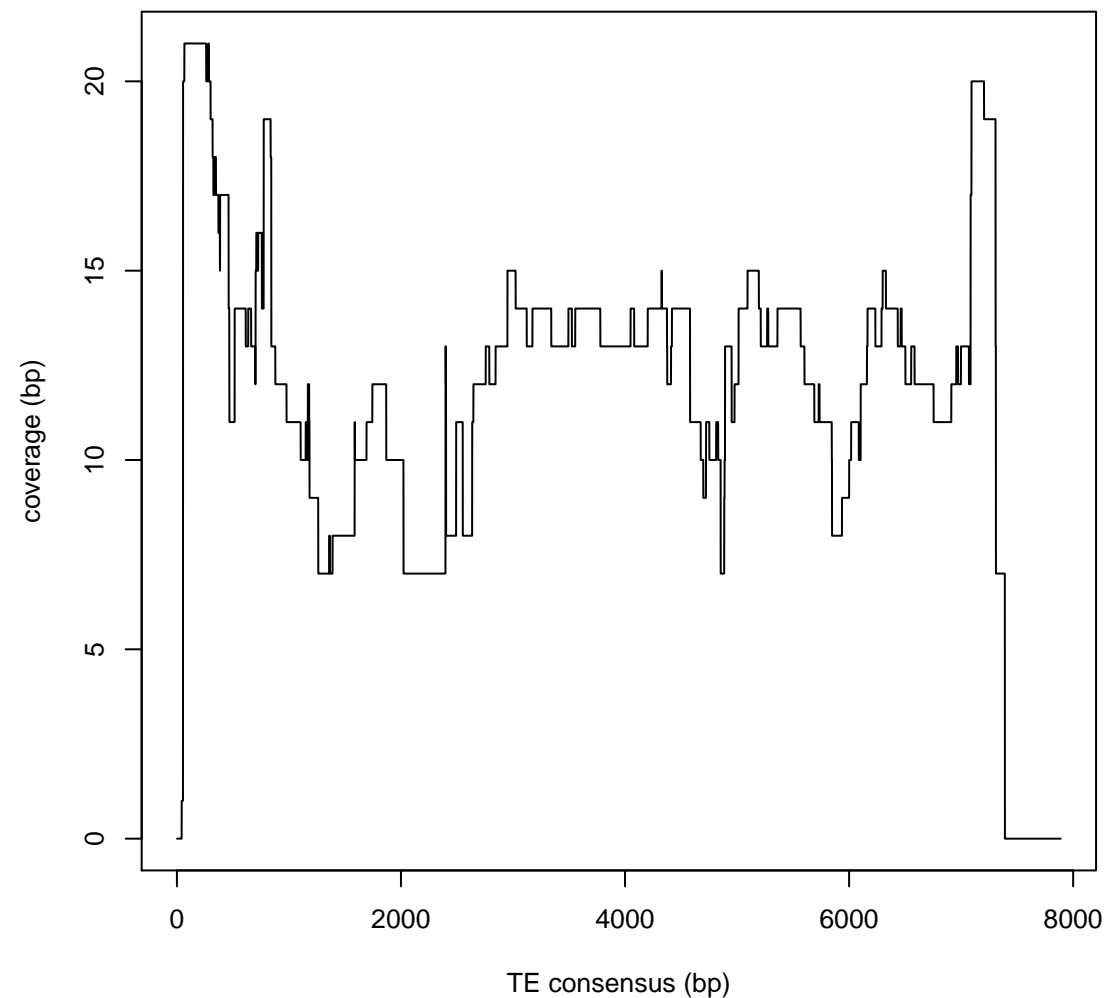


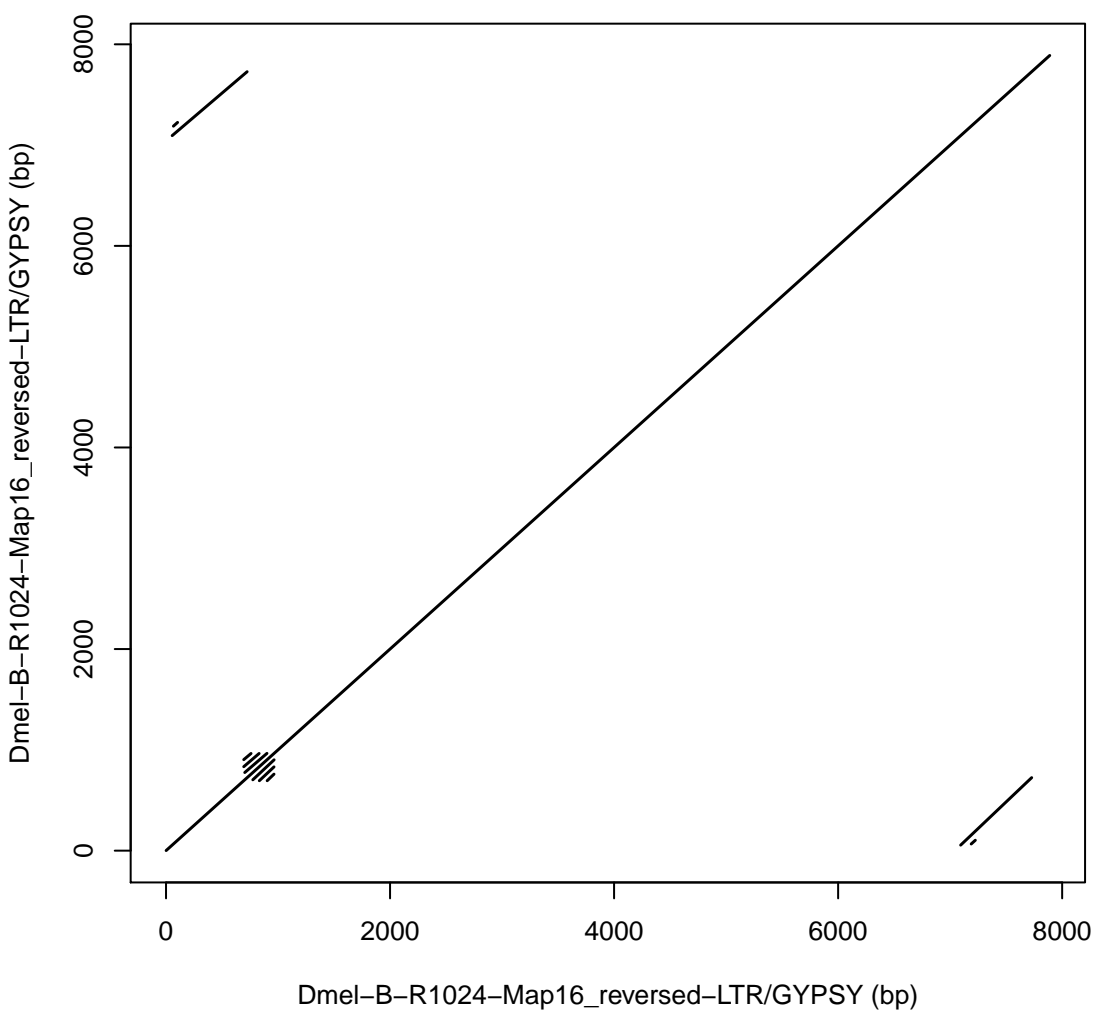
TE: Dmel-B-R1024-Map16\_reversed-LTR/GYPSY  
consensus size: 7889bp; fragments: 112; full length: 1 (>=7100.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

