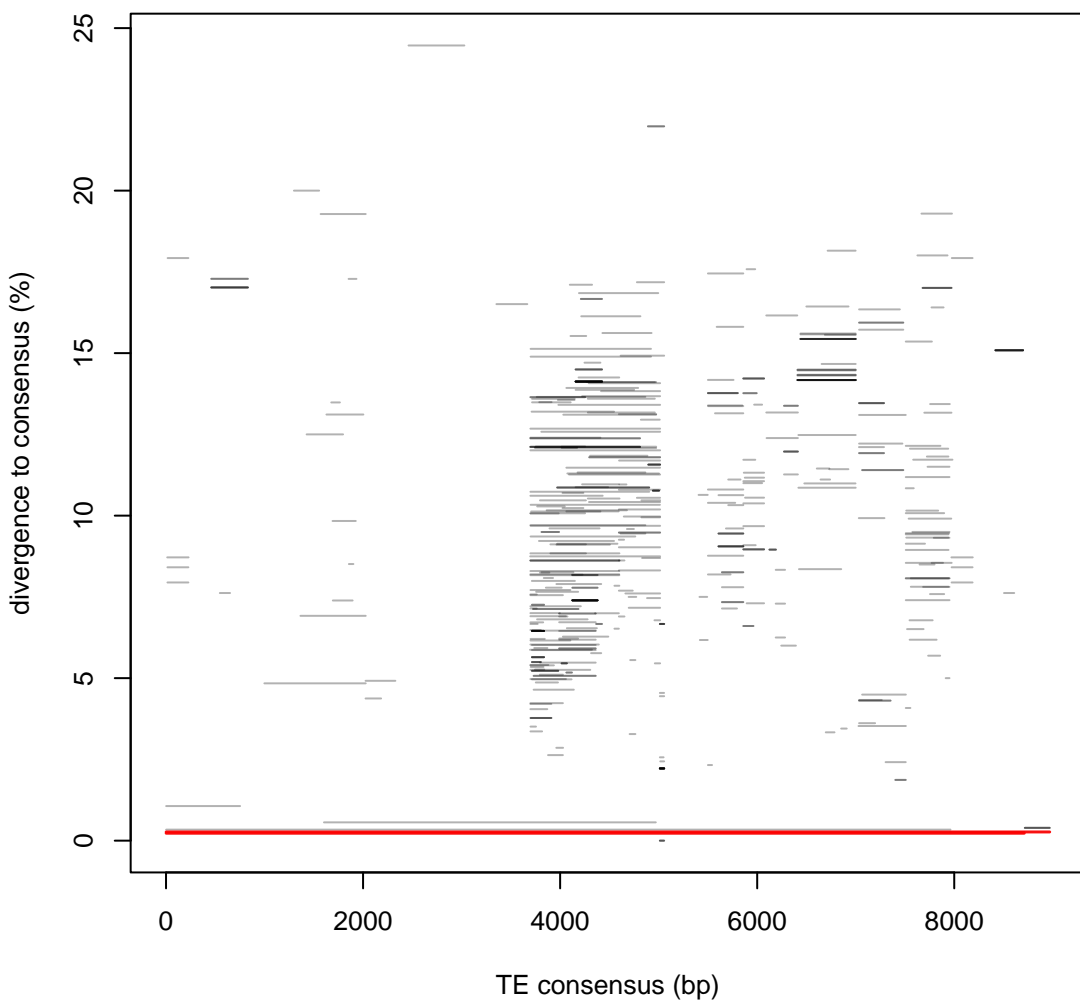
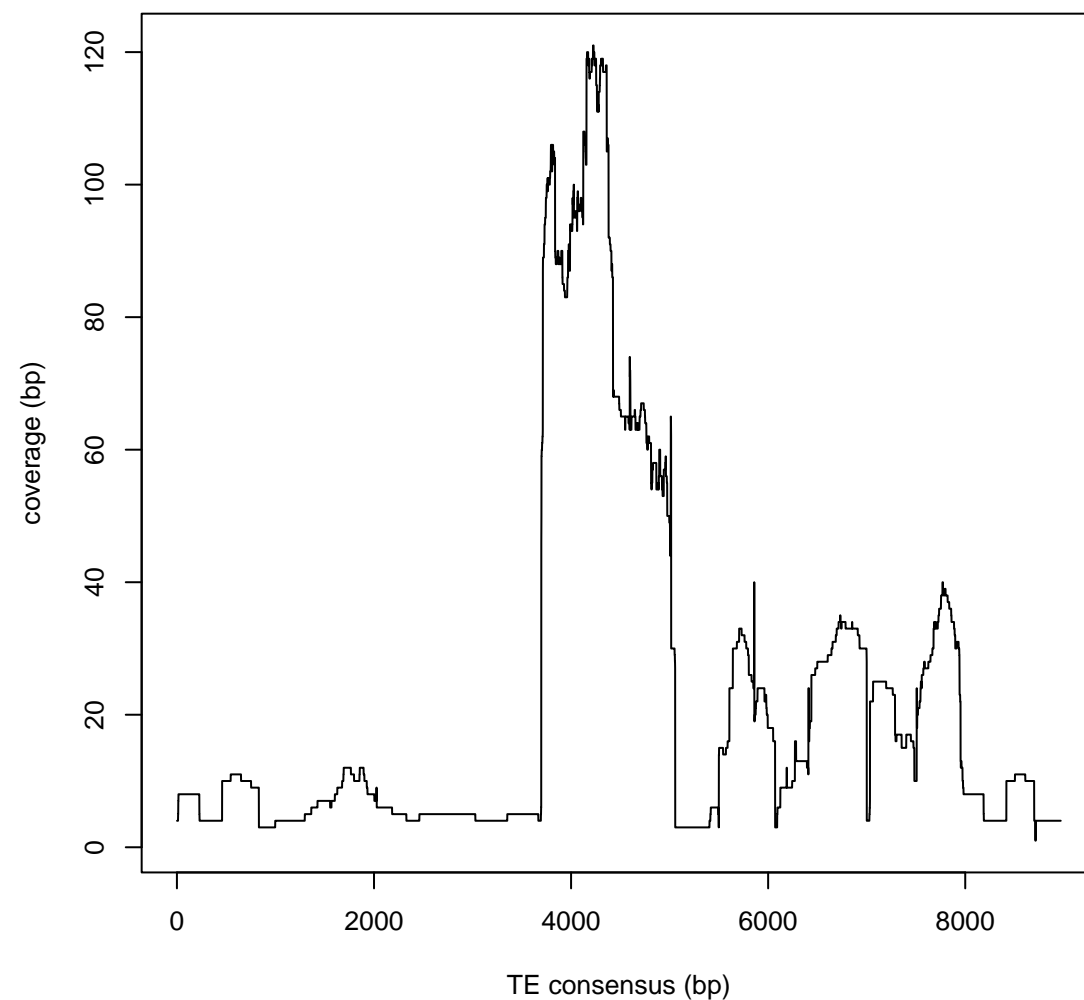


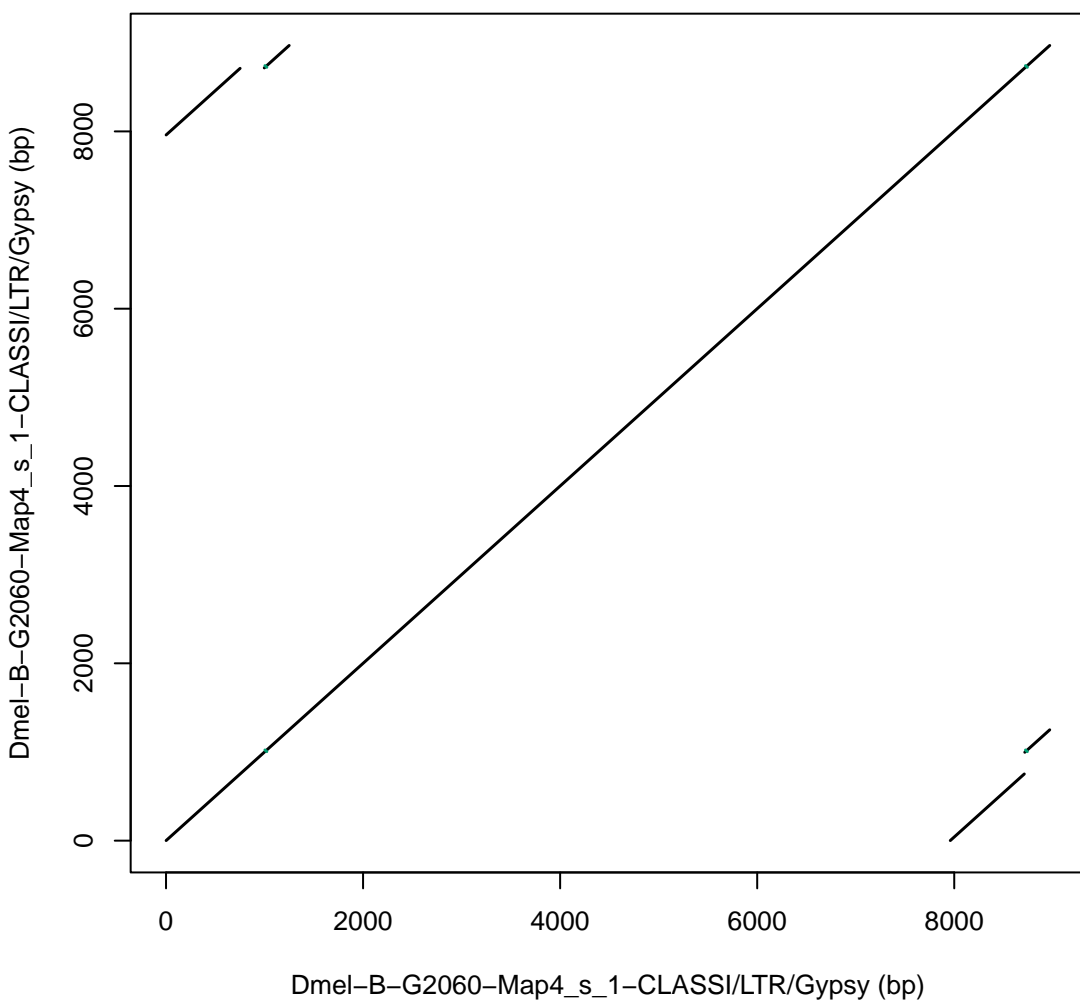
TE: Dmel-B-G2060-Map4_s_1-CLASSI/LTR/Gypsy
consensus size: 8969bp; fragments: 494; full length: 2 (>=8072.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

