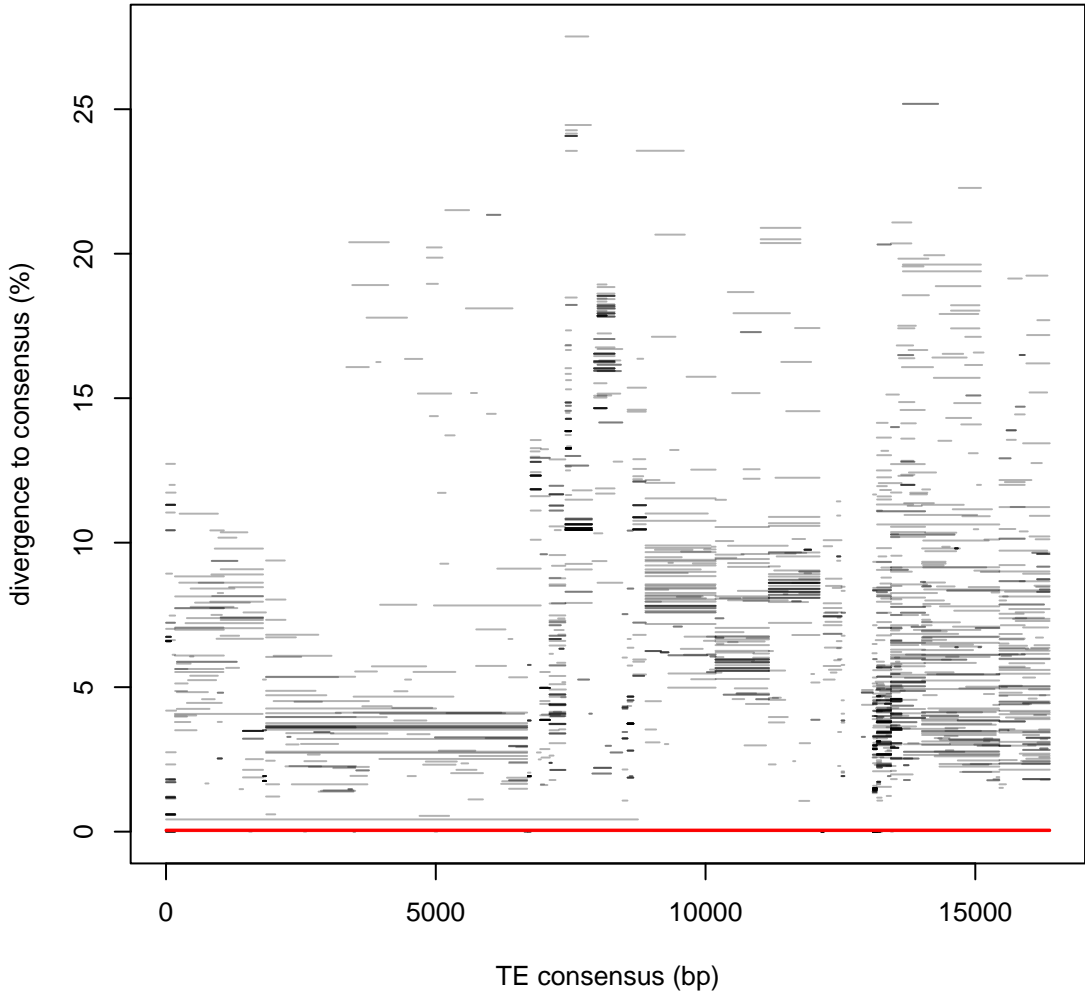
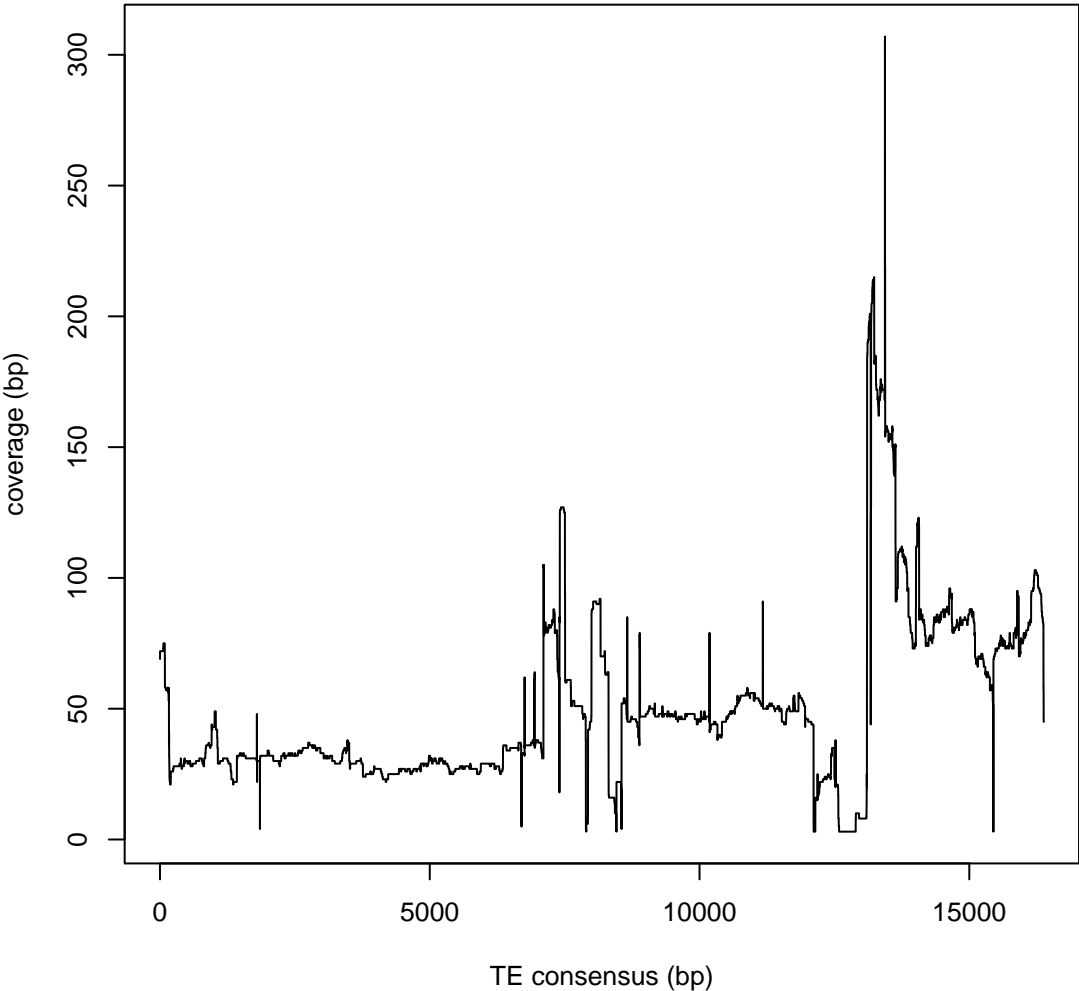


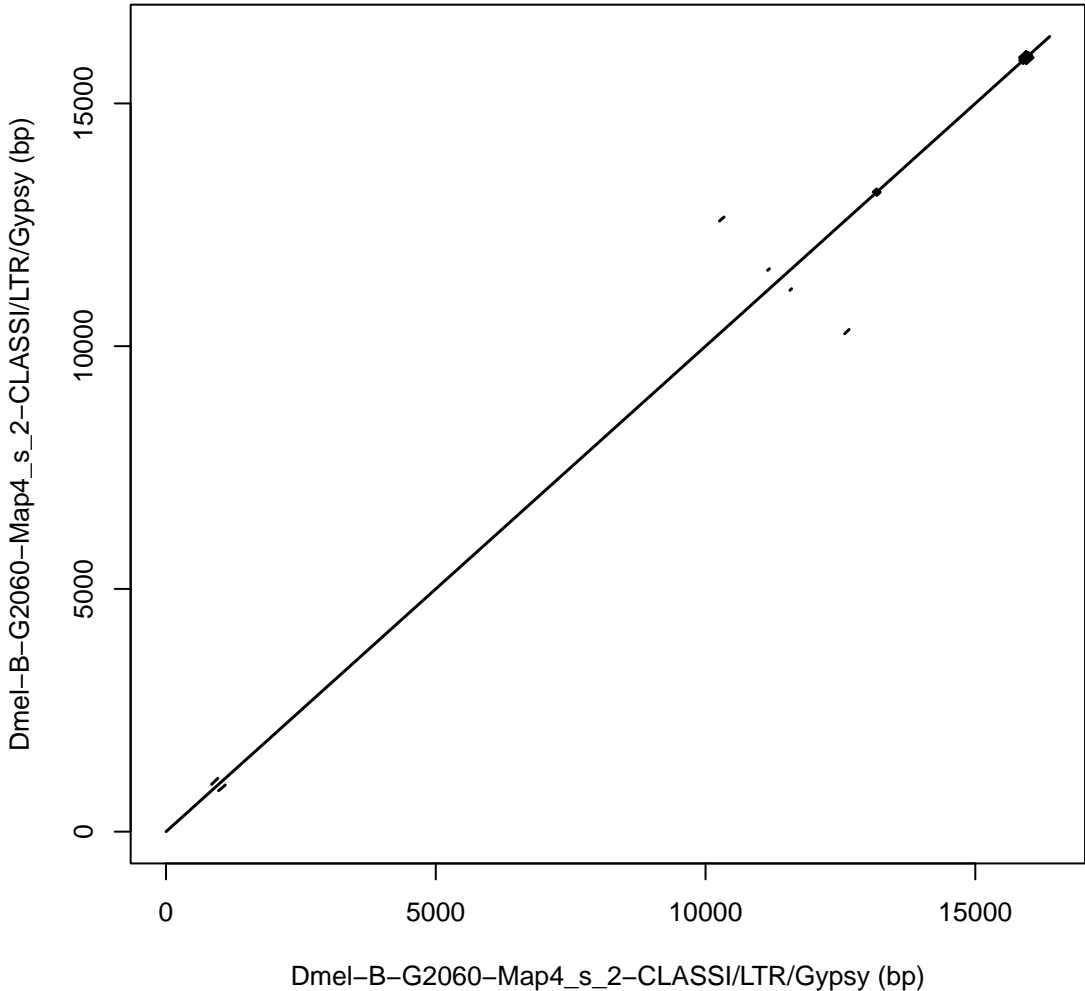
TE: Dmel-B-G2060-Map4_s_2-CLASSI/LTR/Gypsy
consensus size: 16379bp; fragments: 2056; full length: 2 (>=14741.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

