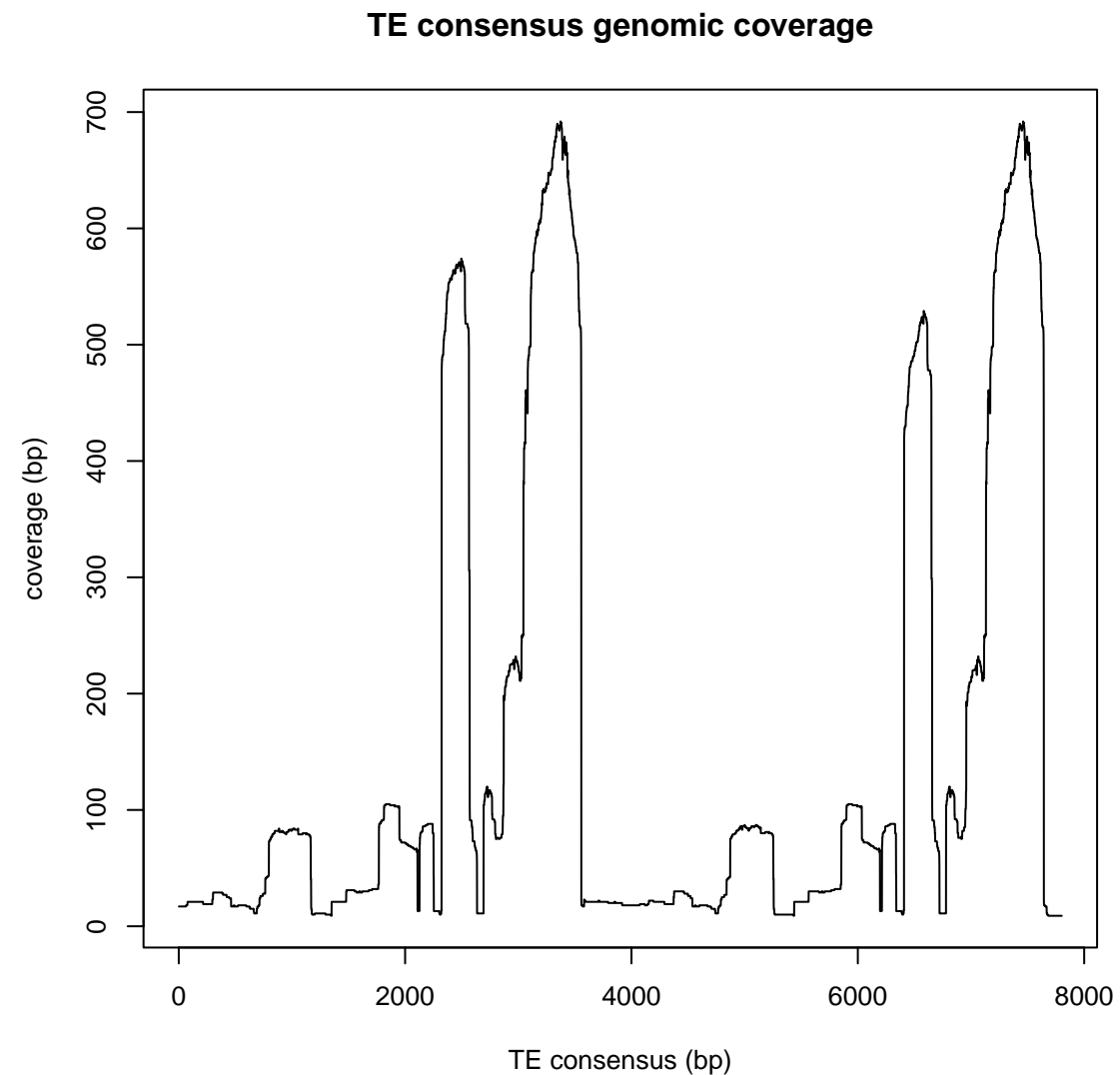
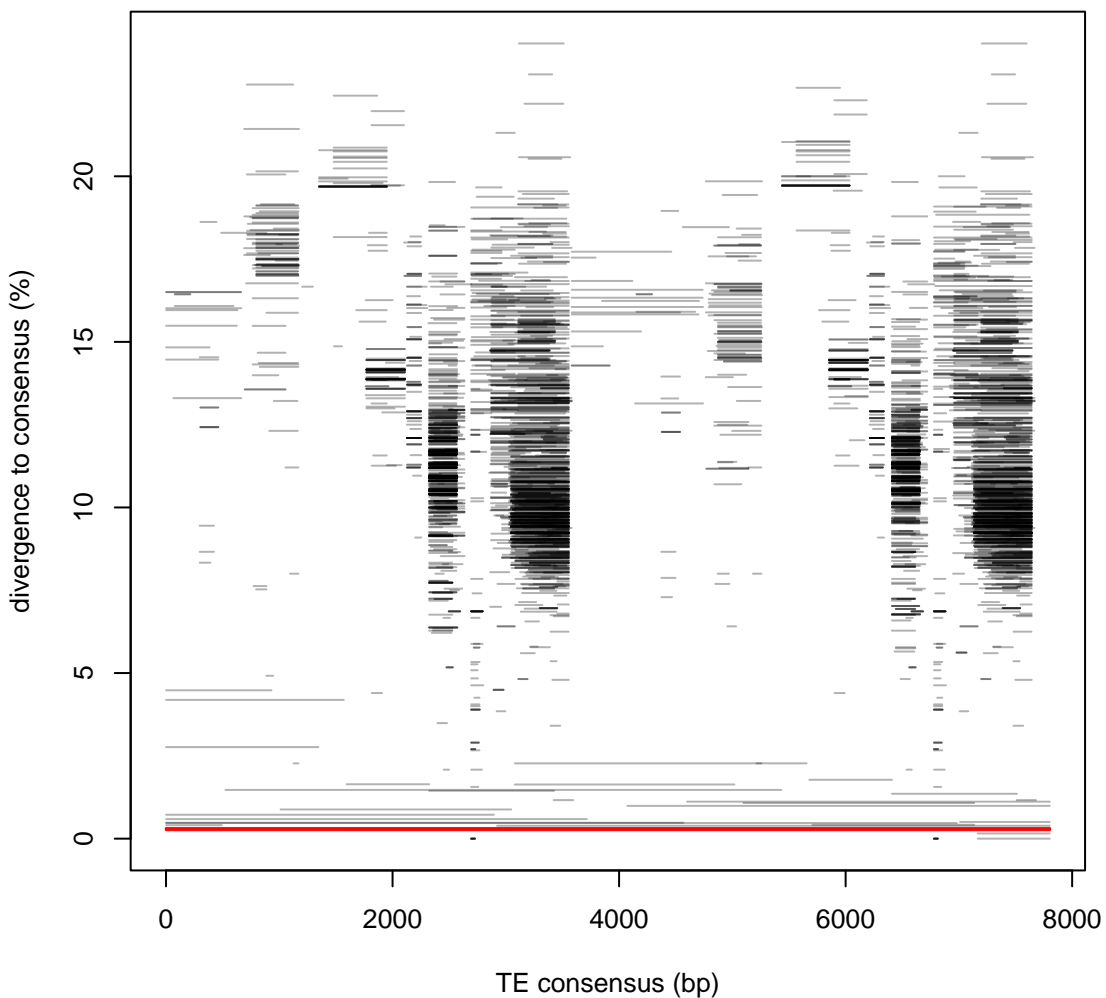
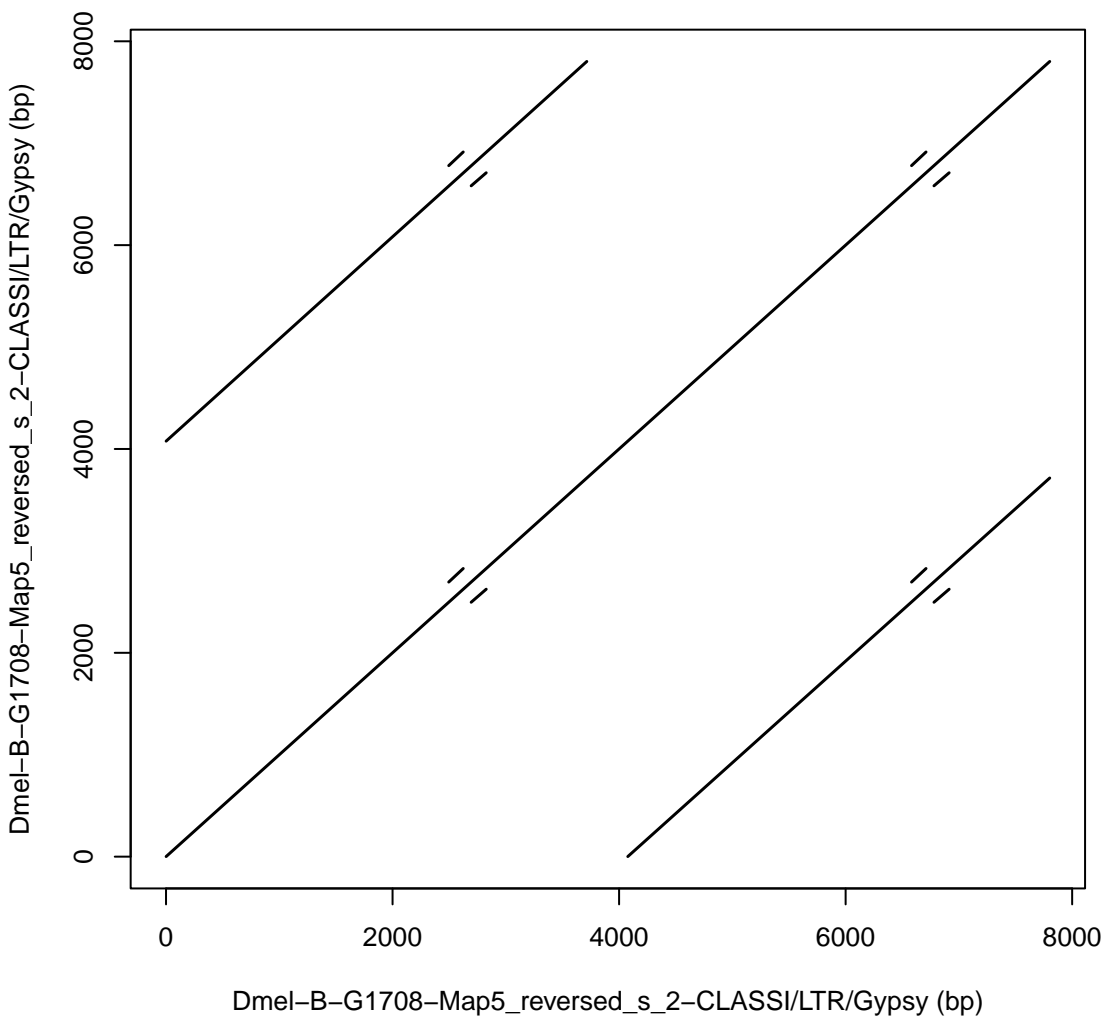


TE: Dmel-B-G1708-Map5_reversed_s_2-CLASSI/LTR/Gypsy
consensus size: 7802bp; fragments: 3558; full length: 2 (≥ 7021.8 bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

