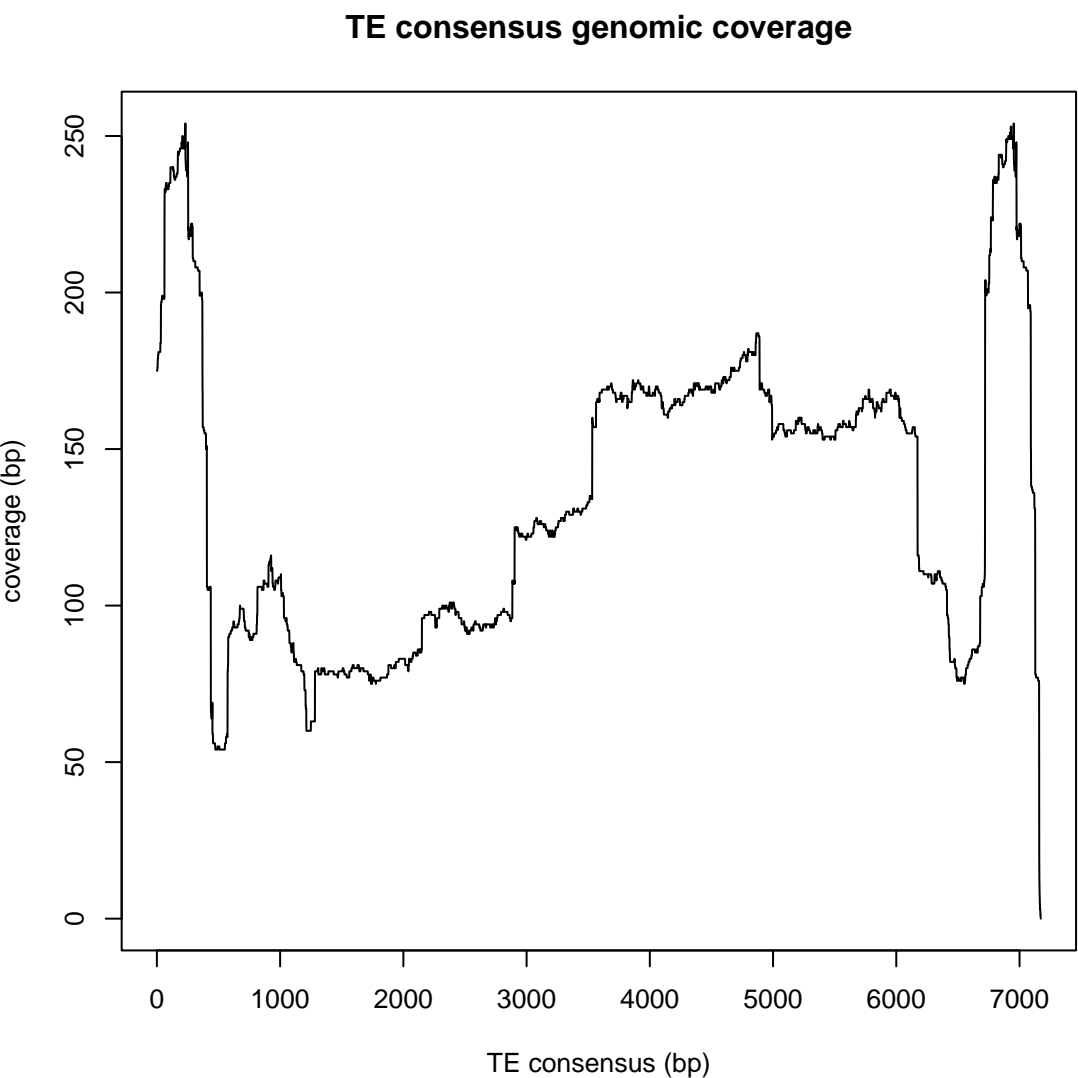
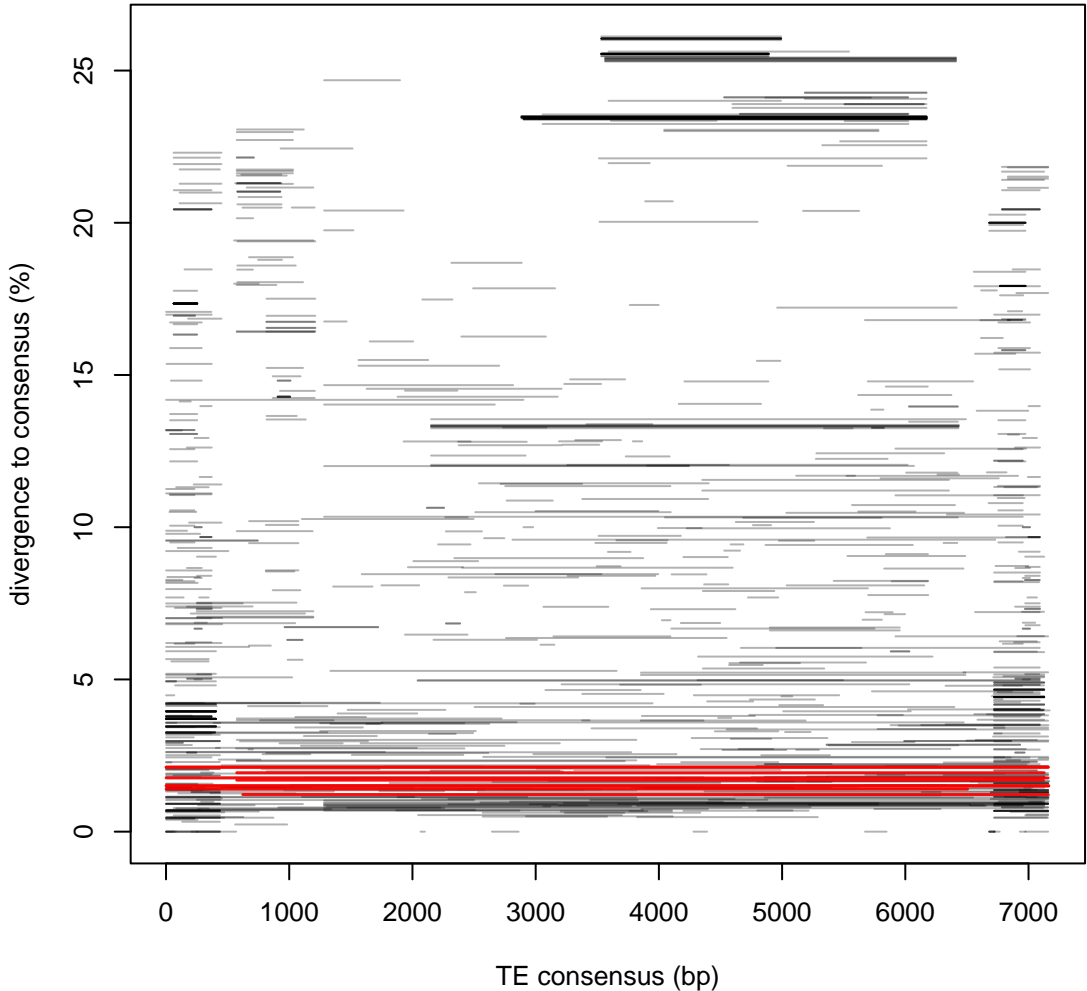
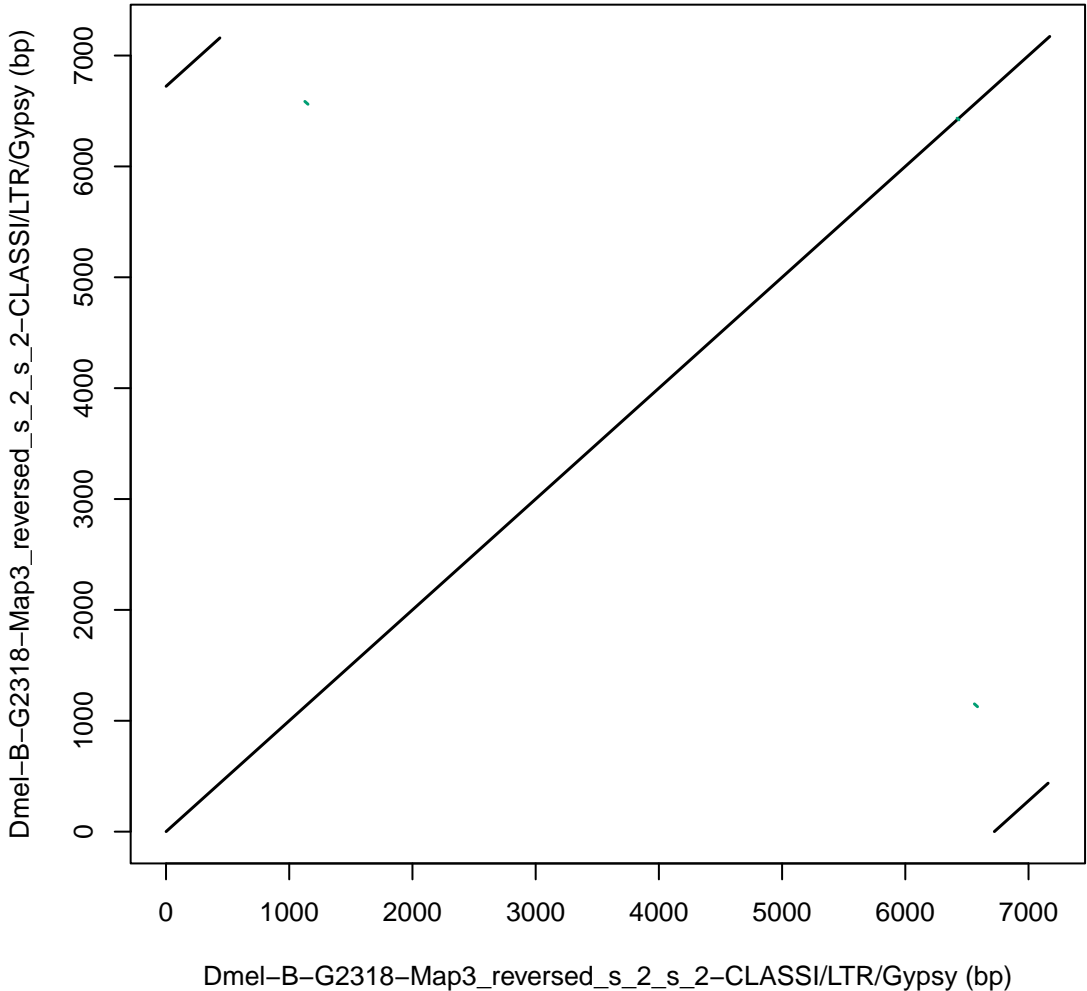


TE: Dmel-B-G2318-Map3_reversed_s_2_s_2-CLASSI/LTR/Gypsy
consensus size: 7172bp; fragments: 1051; full length: 9 (>=6454.8bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

