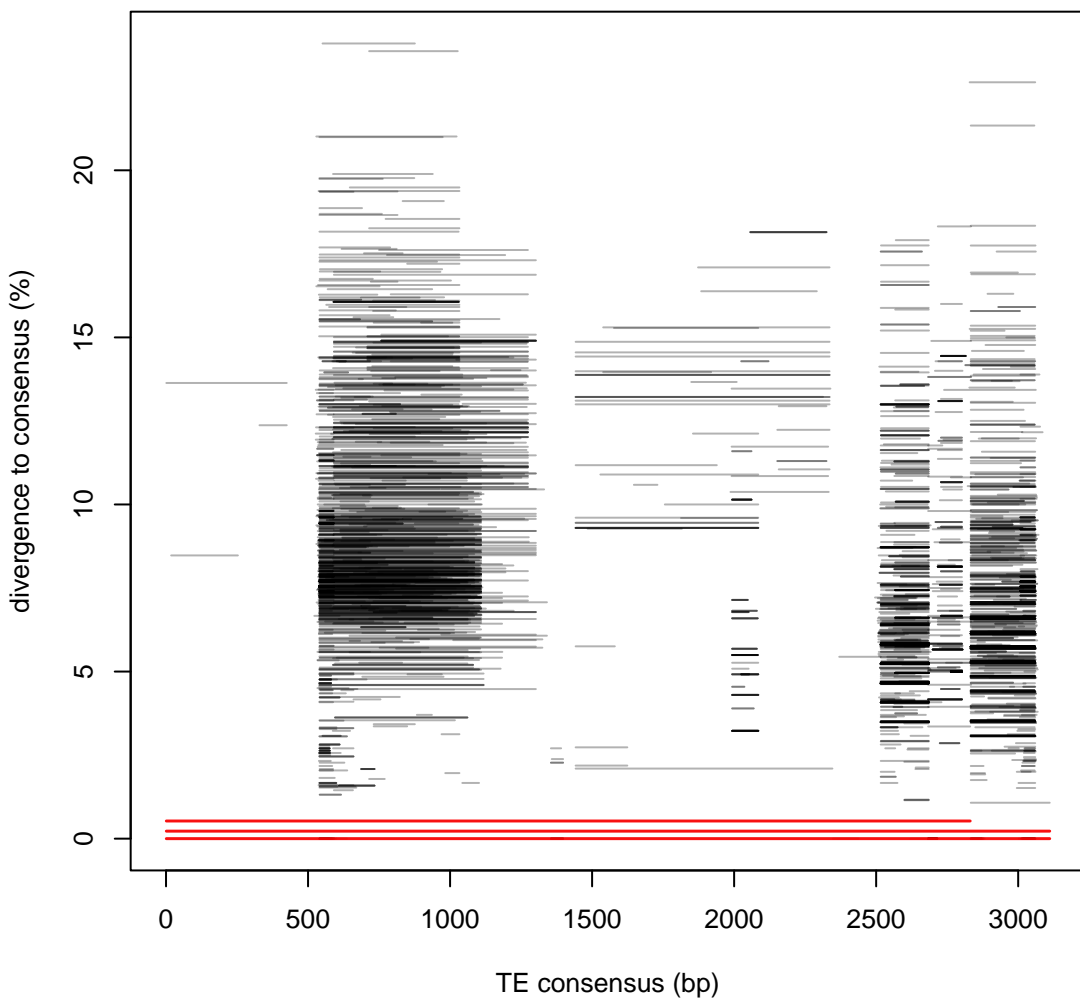
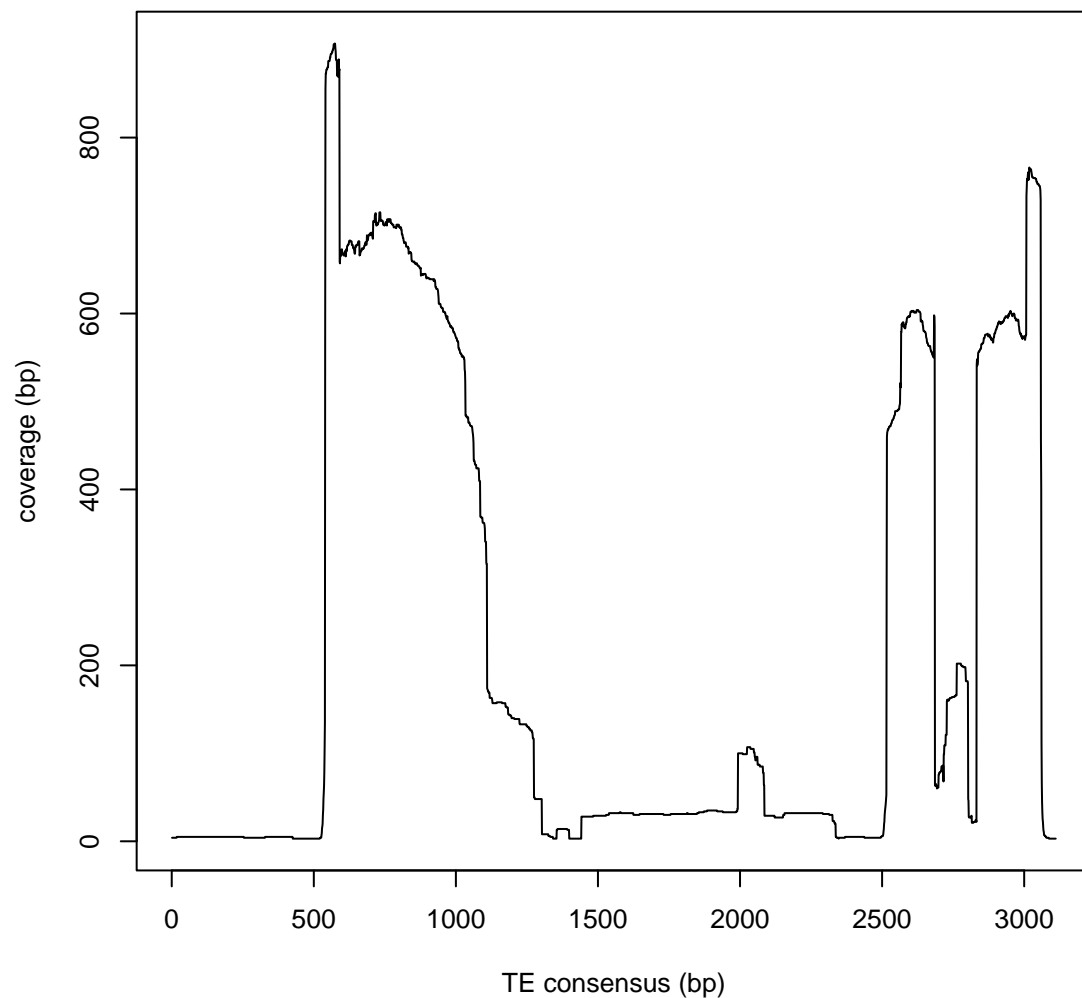


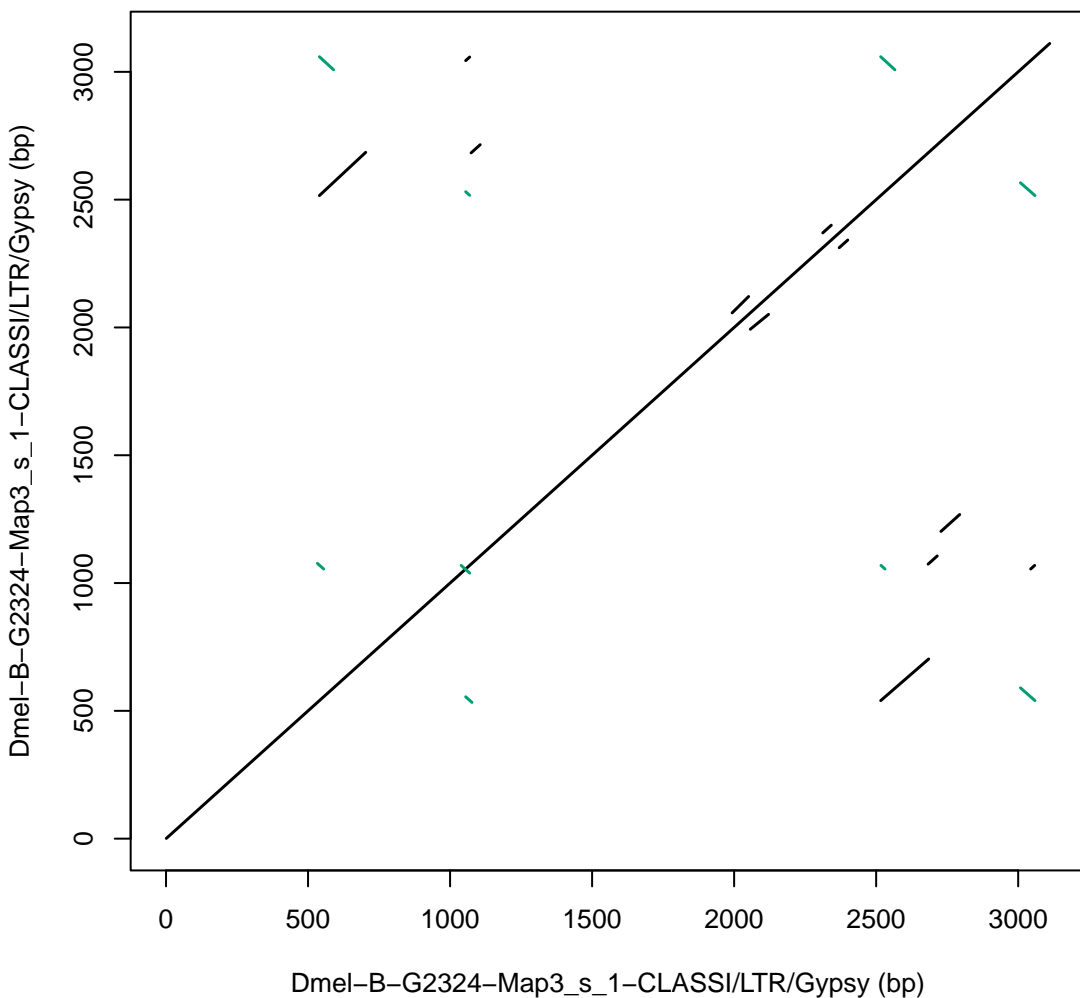
TE: Dmel-B-G2324-Map3_s_1-CLASSI/LTR/Gypsy
consensus size: 3111bp; fragments: 3136; full length: 3 (>=2799.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

