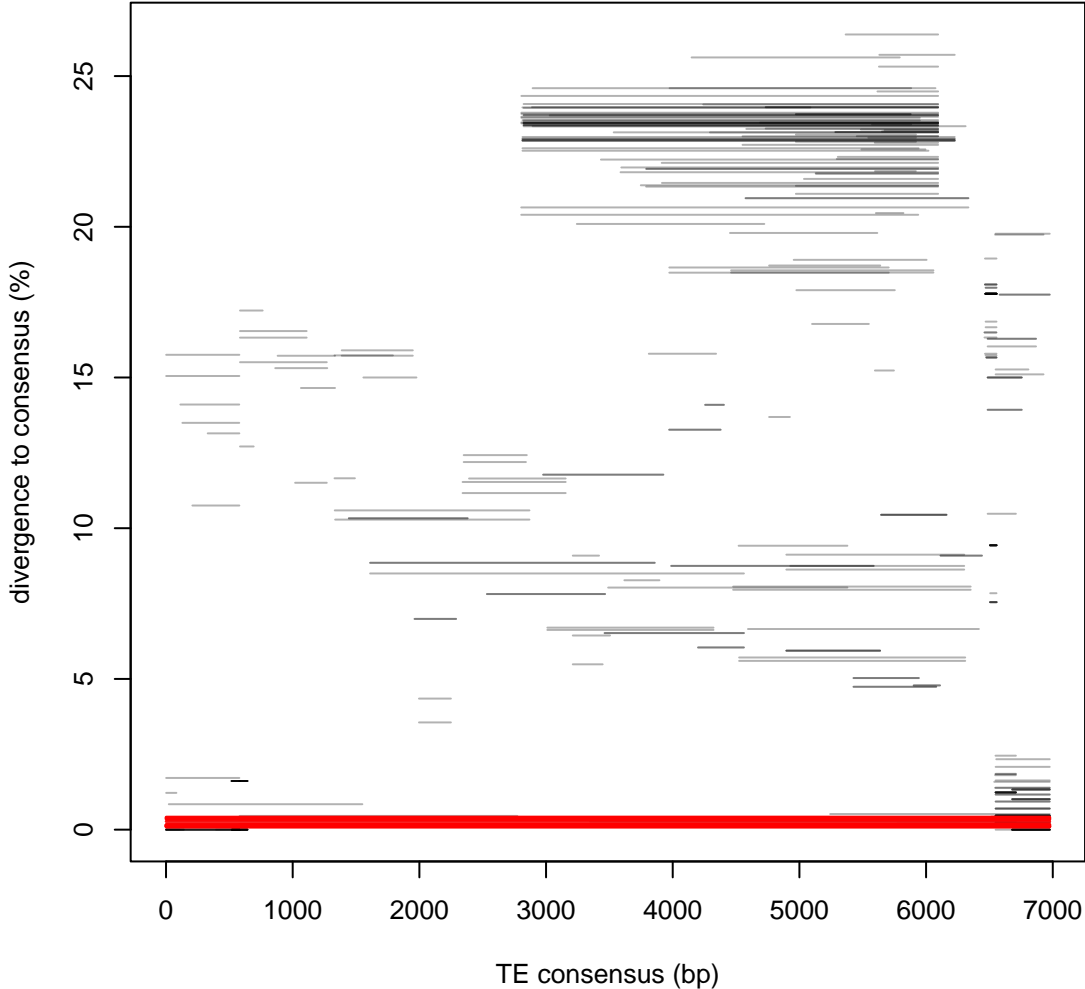
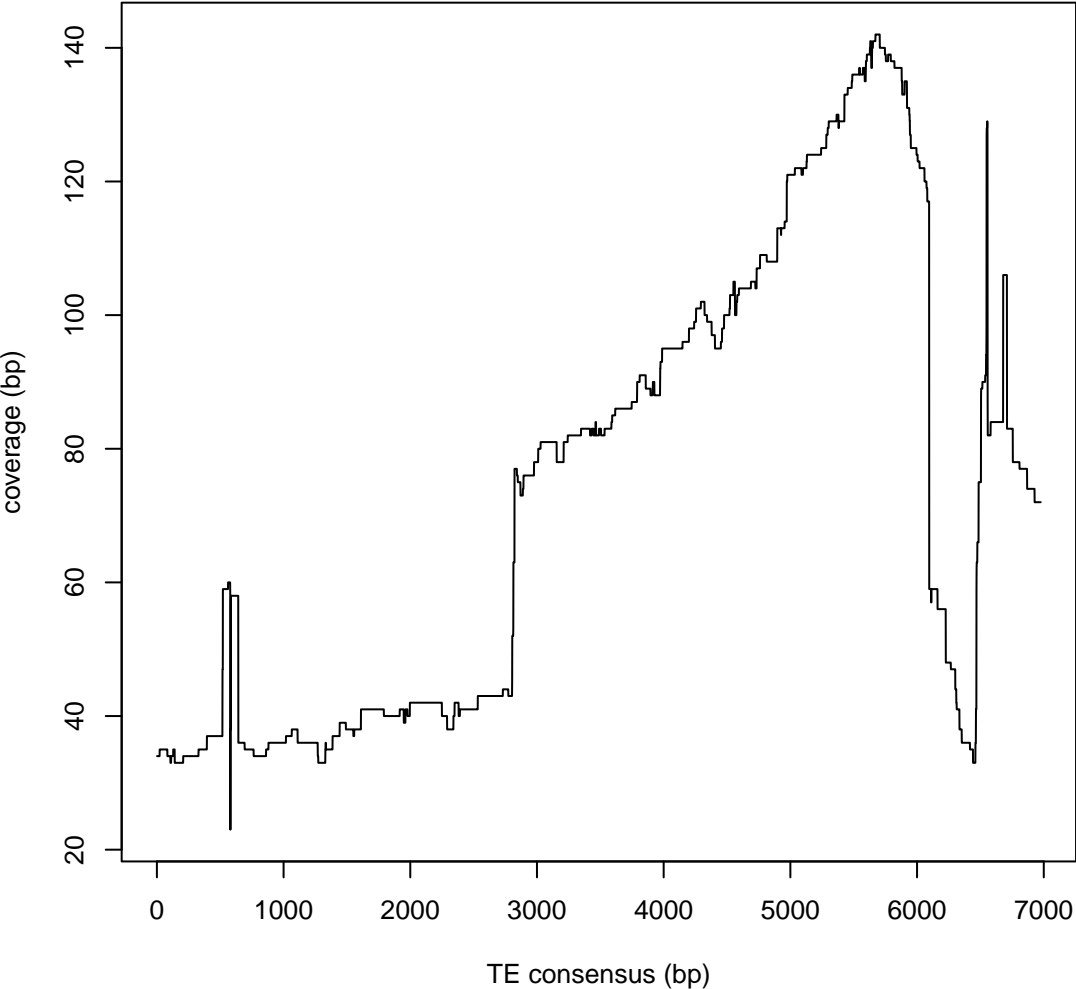


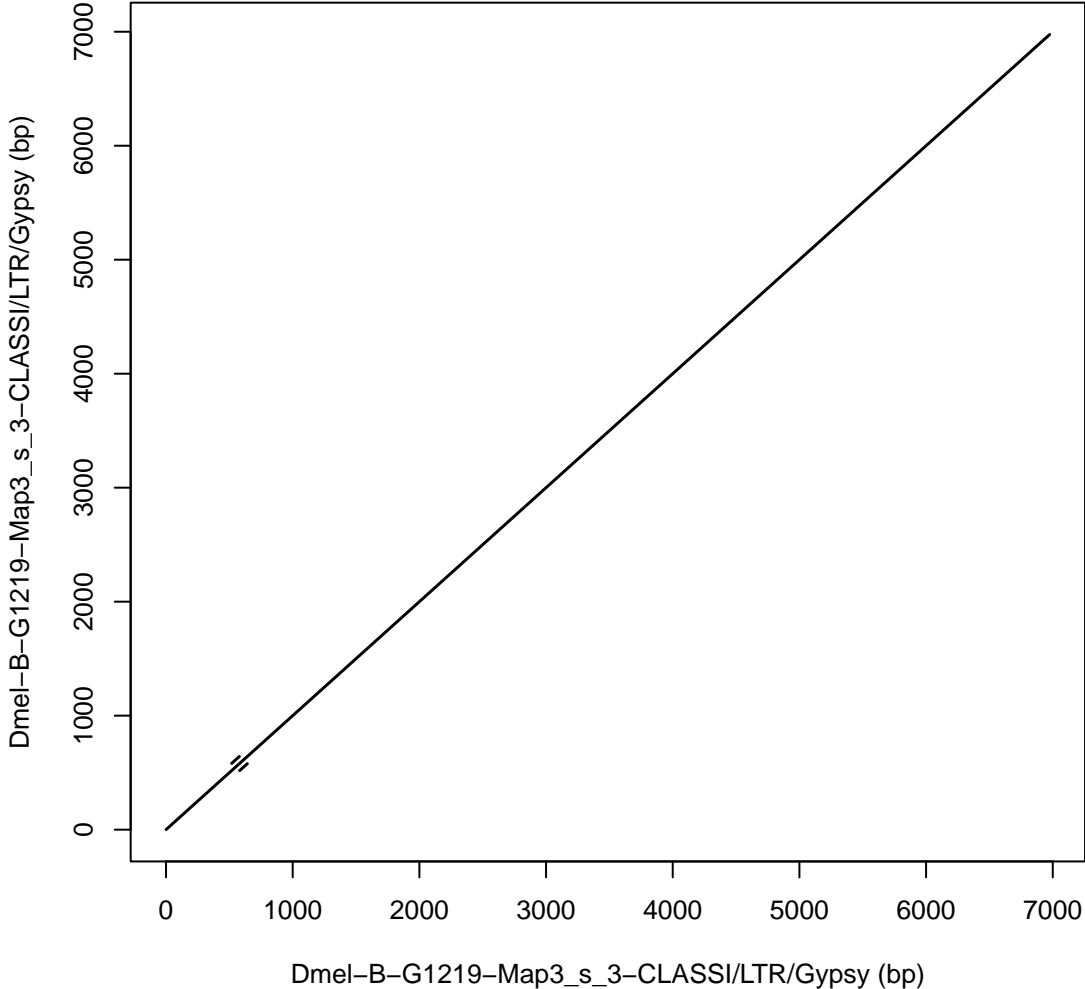
TE: Dmel-B-G1219-Map3_s_3-CLASSI/LTR/Gypsy
consensus size: 6976bp; fragments: 395; full length: 27 (>=6278.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

