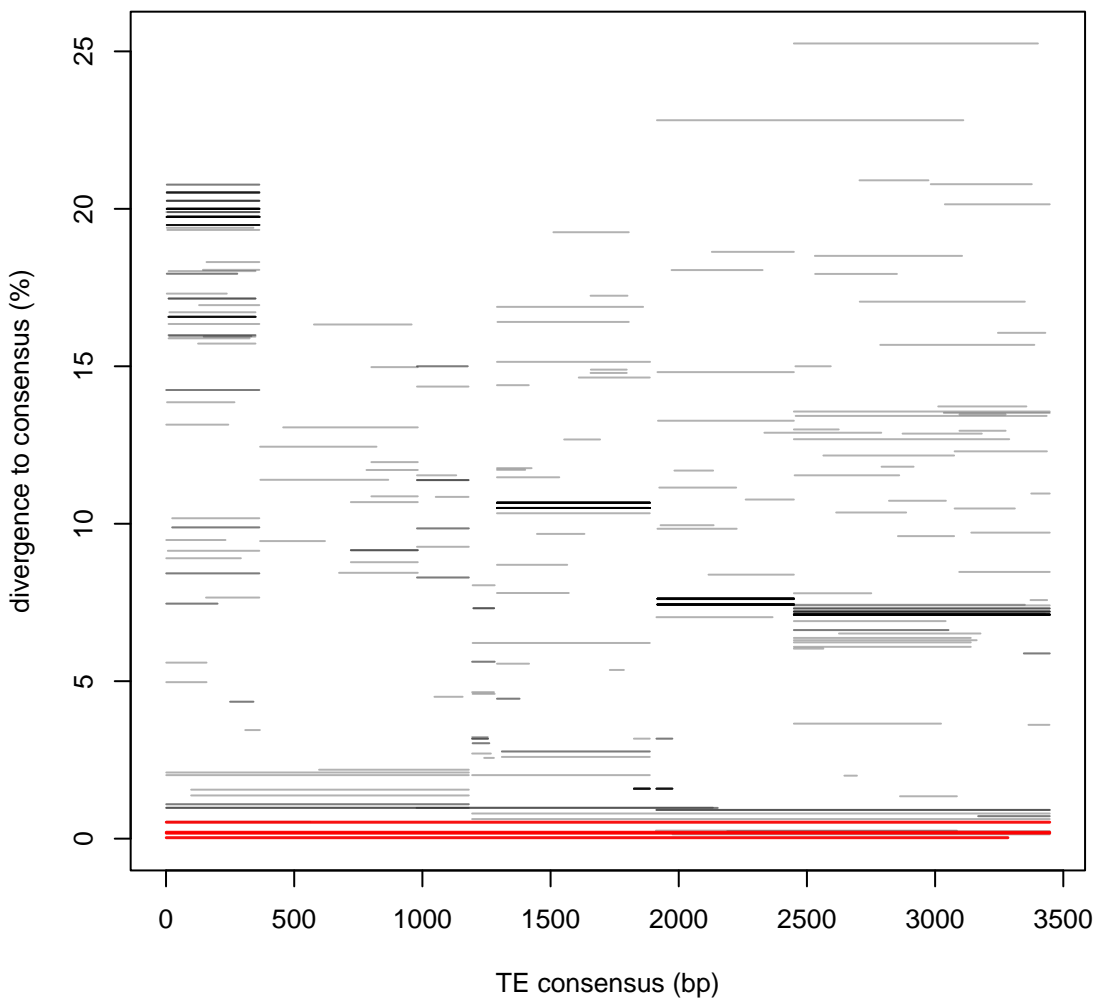
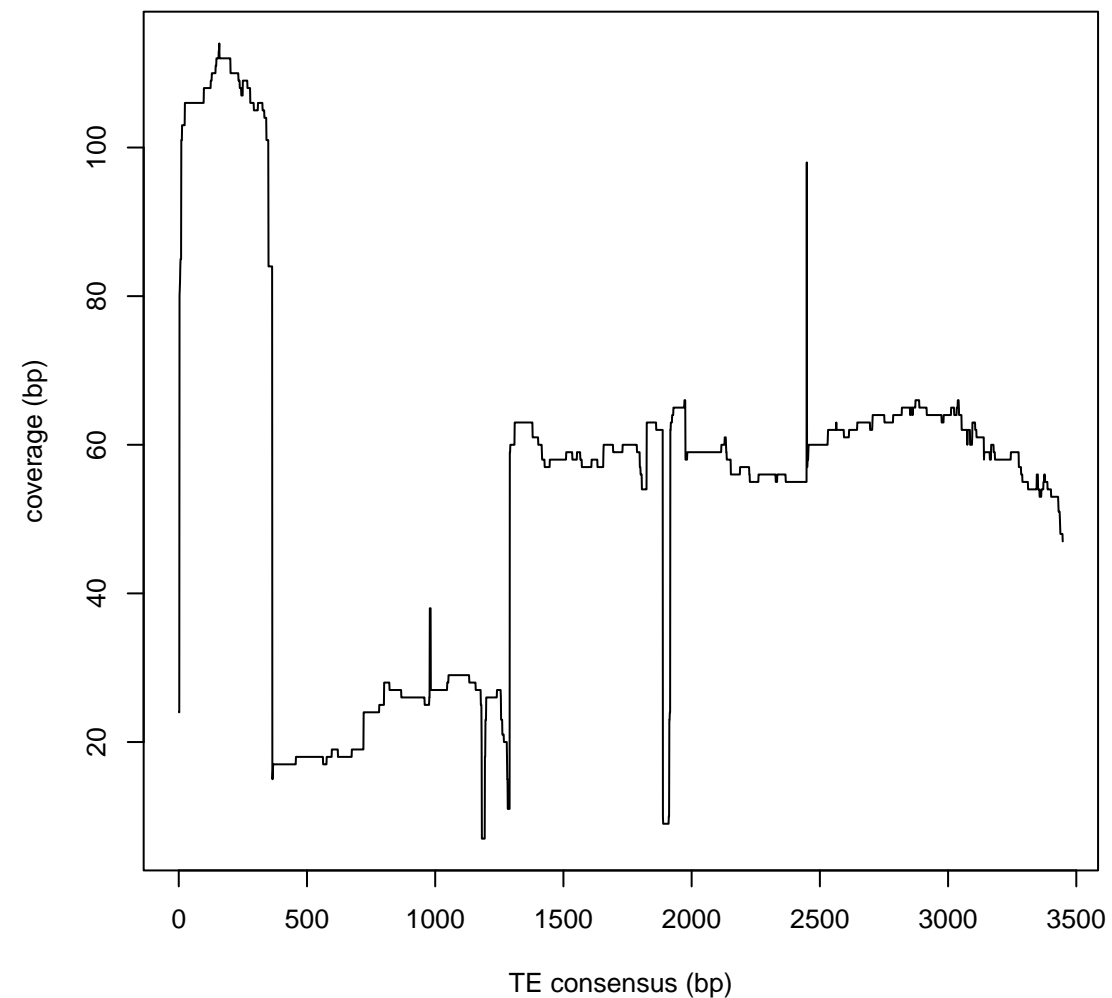


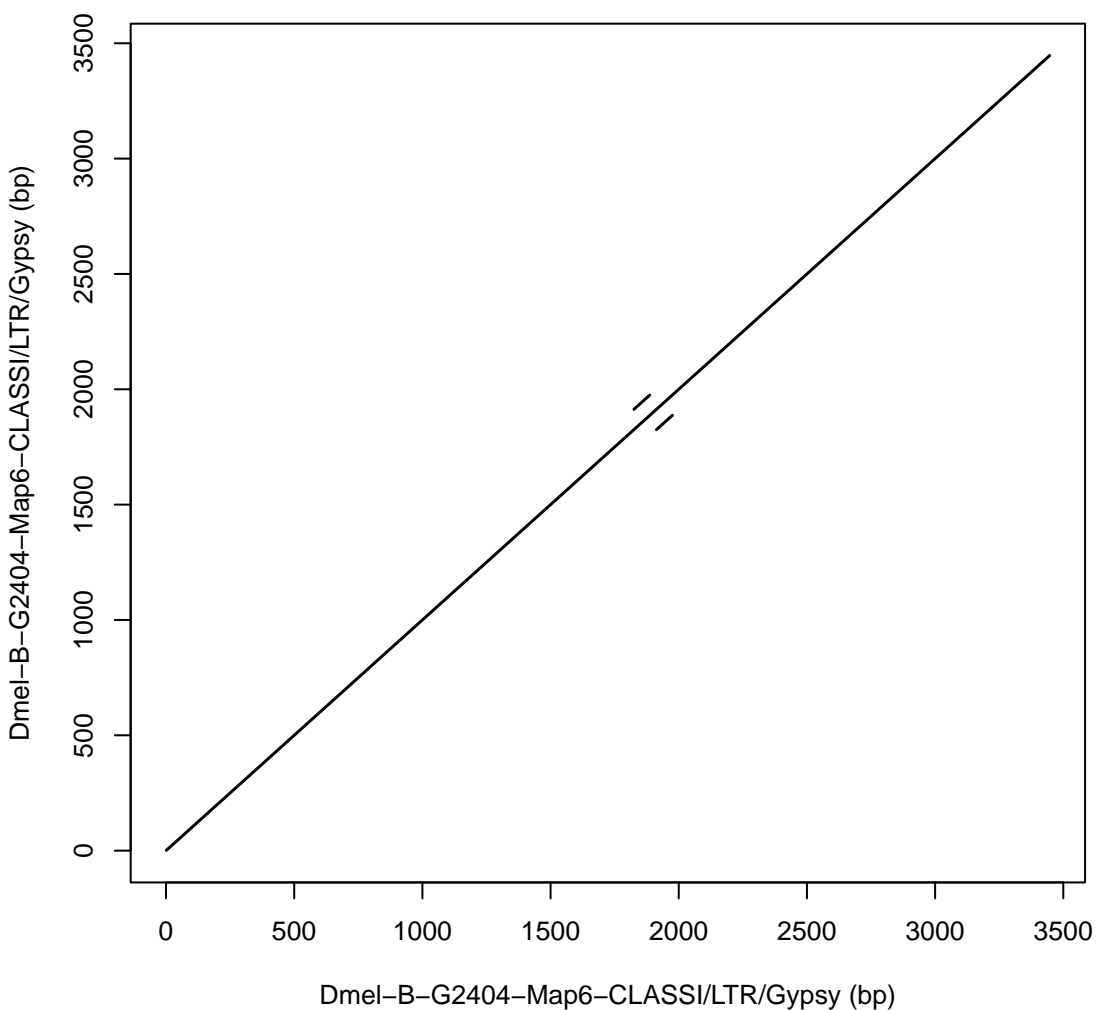
TE: Dmel-B-G2404-Map6-CLASSI/LTR/Gypsy
consensus size: 3447bp; fragments: 379; full length: 4 (≥ 3102.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

