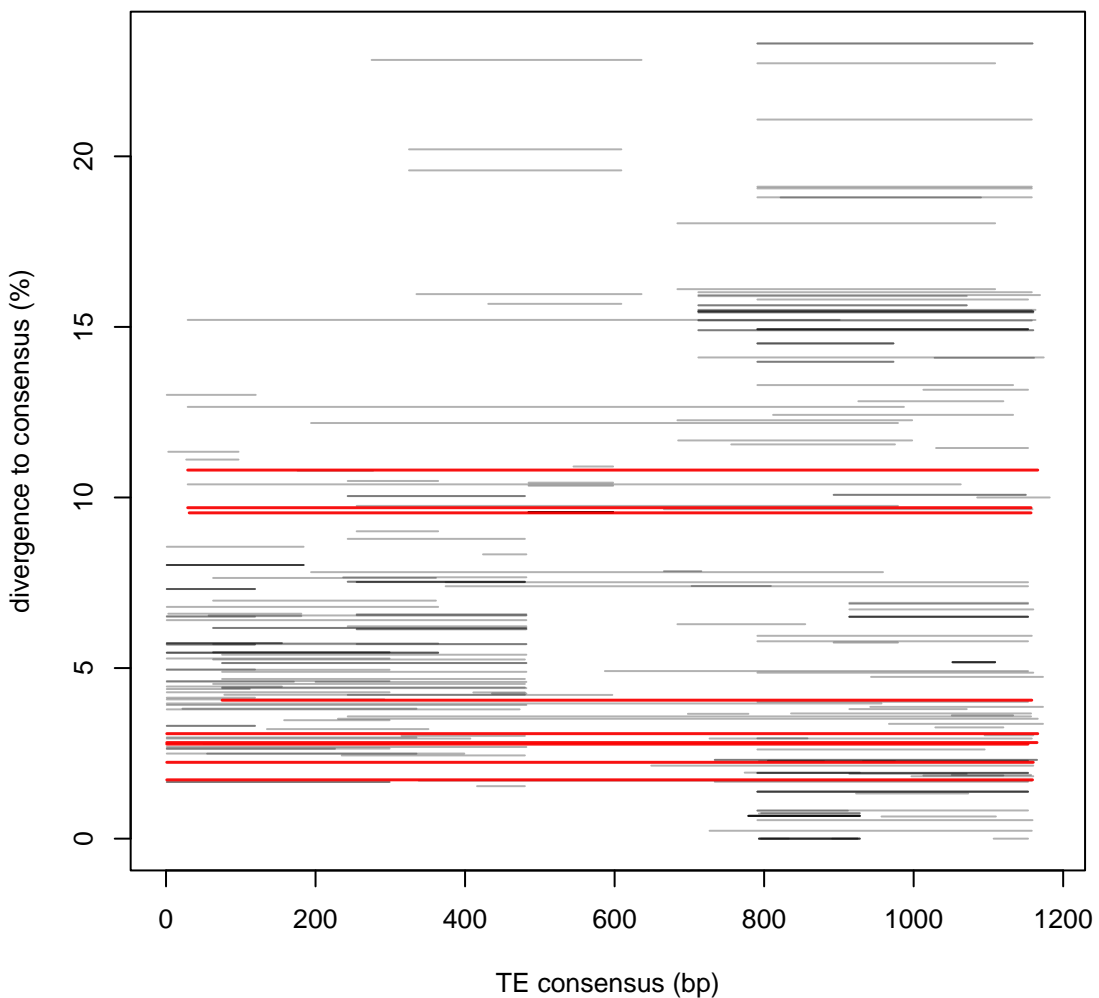
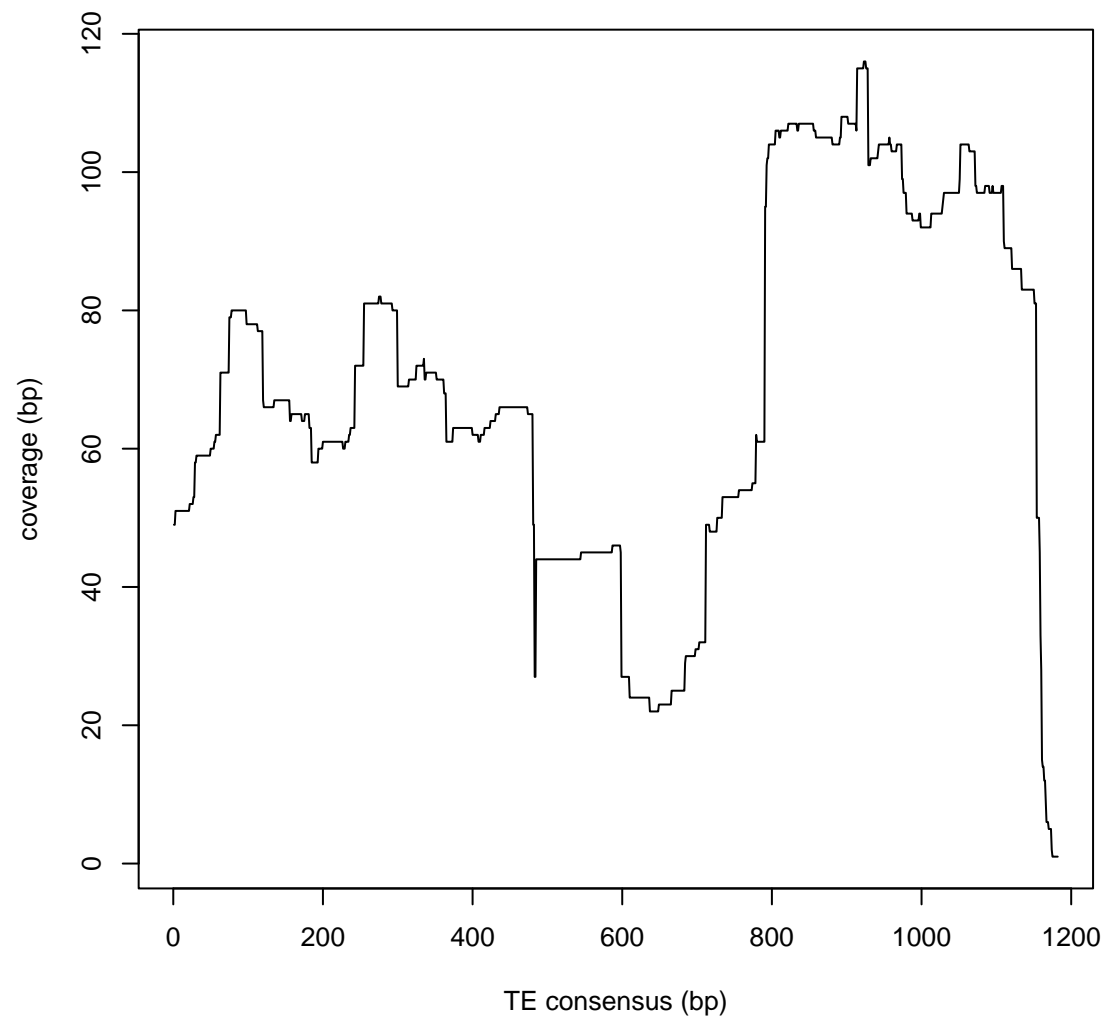


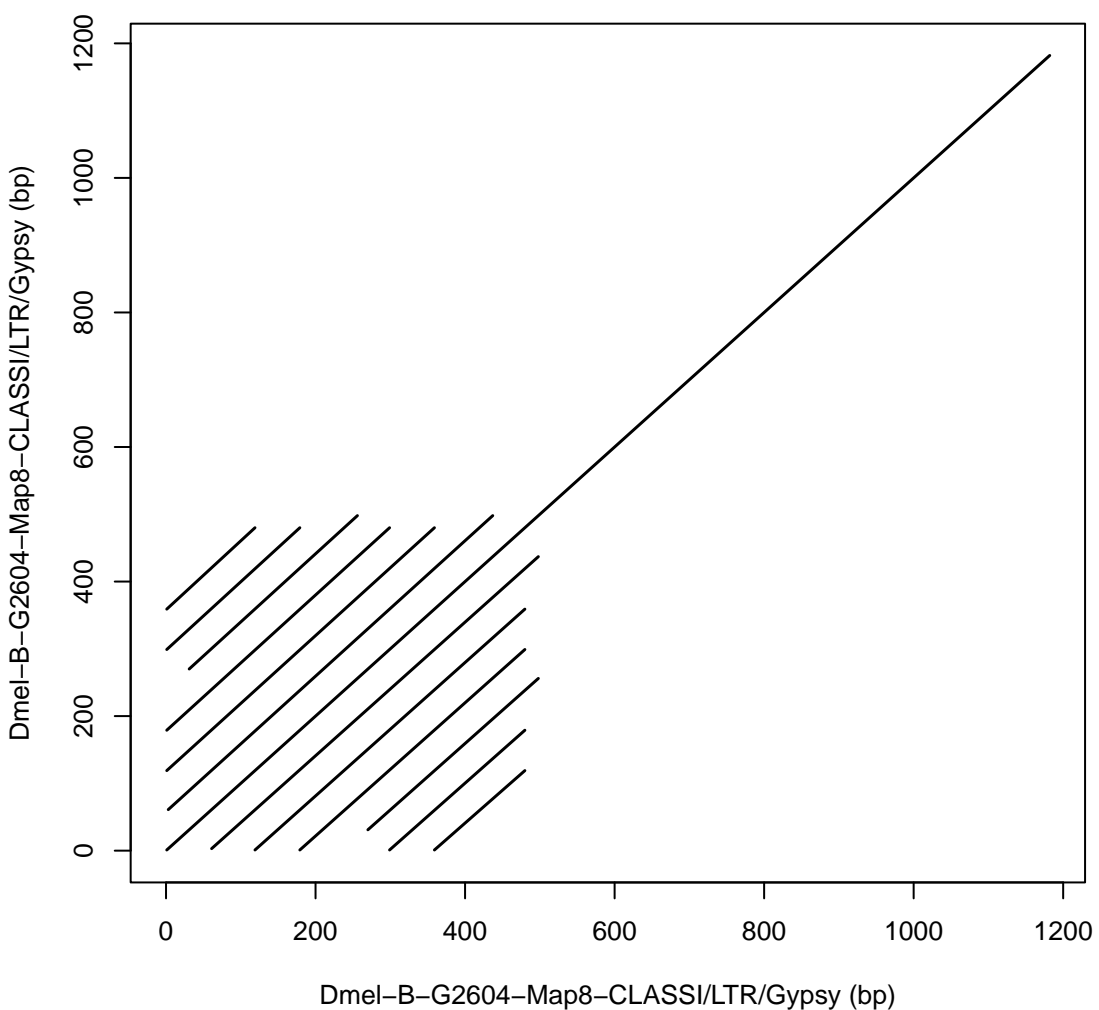
TE: Dmel-B-G2604-Map8-CLASSI/LTR/Gypsy
consensus size: 1182bp; fragments: 264; full length: 9 (≥ 1063.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

