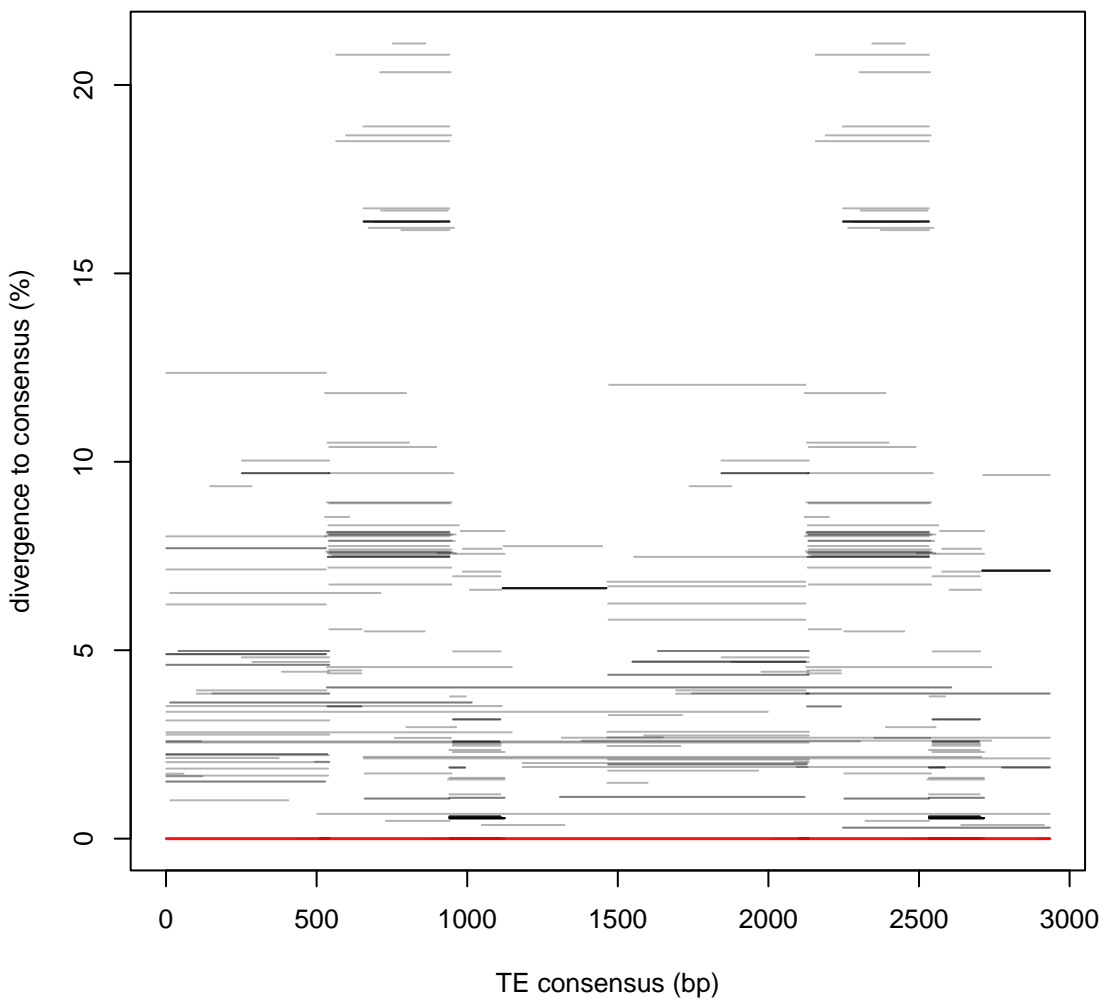
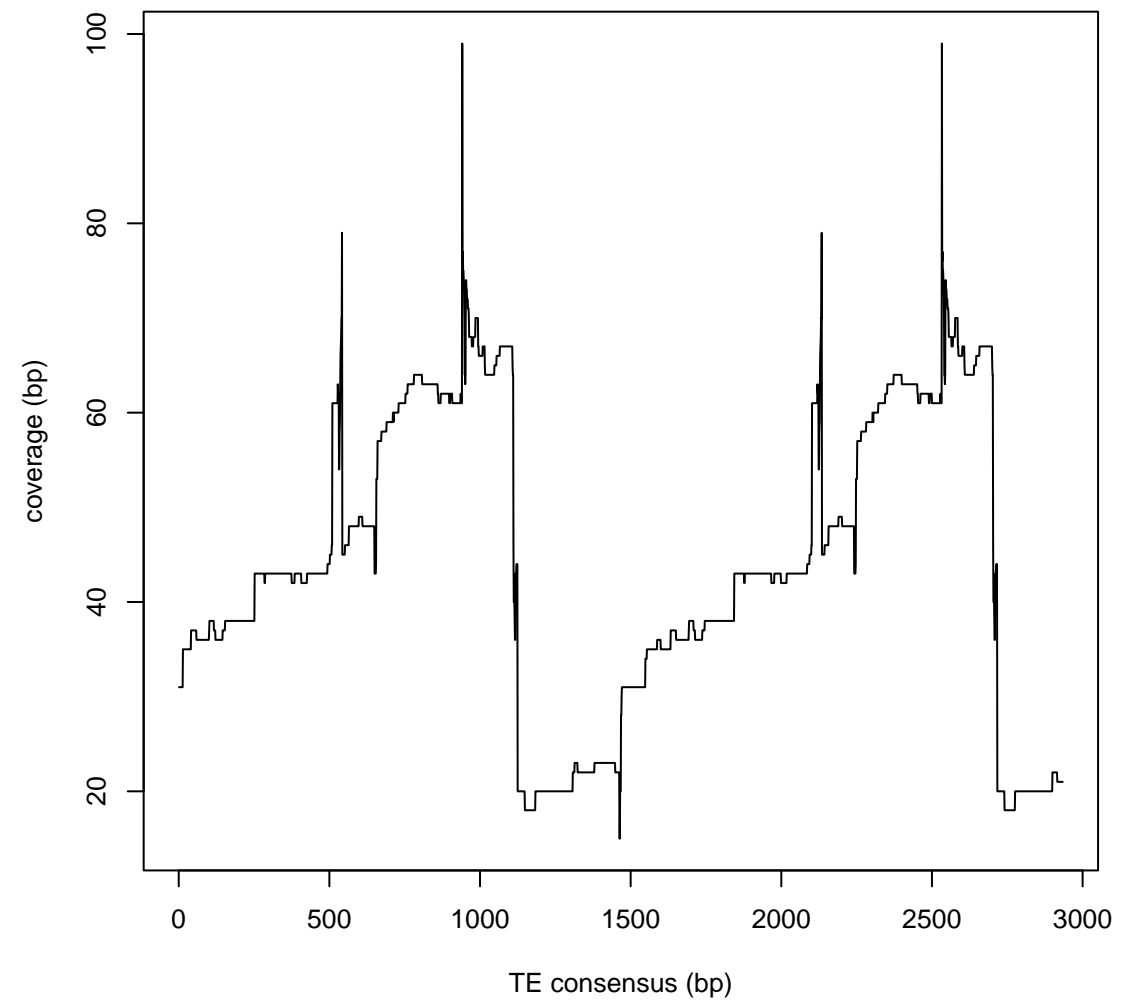


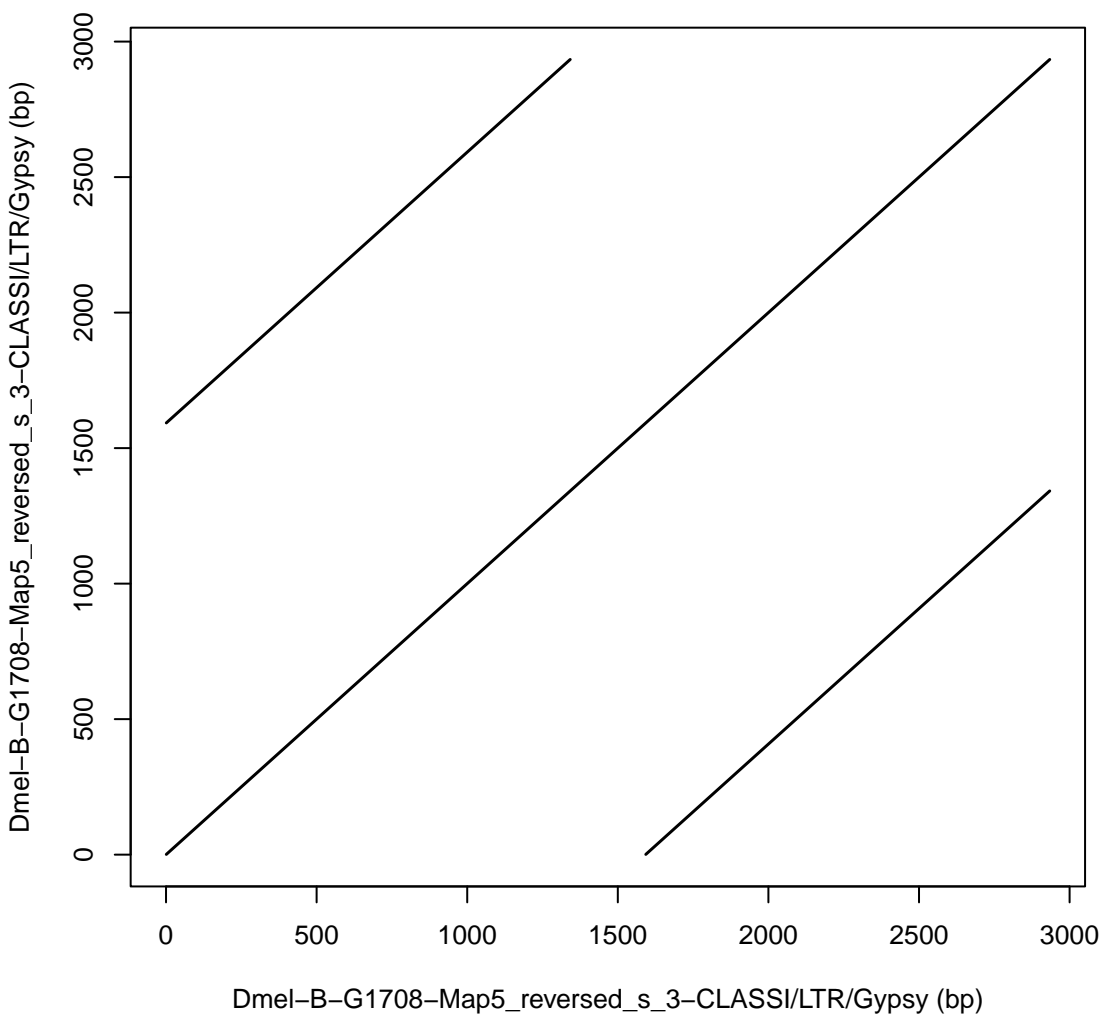
TE: Dmel-B-G1708-Map5_reversed_s_3-CLASSI/LTR/Gypsy
consensus size: 2934bp; fragments: 394; full length: 1 (≥ 2640.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

