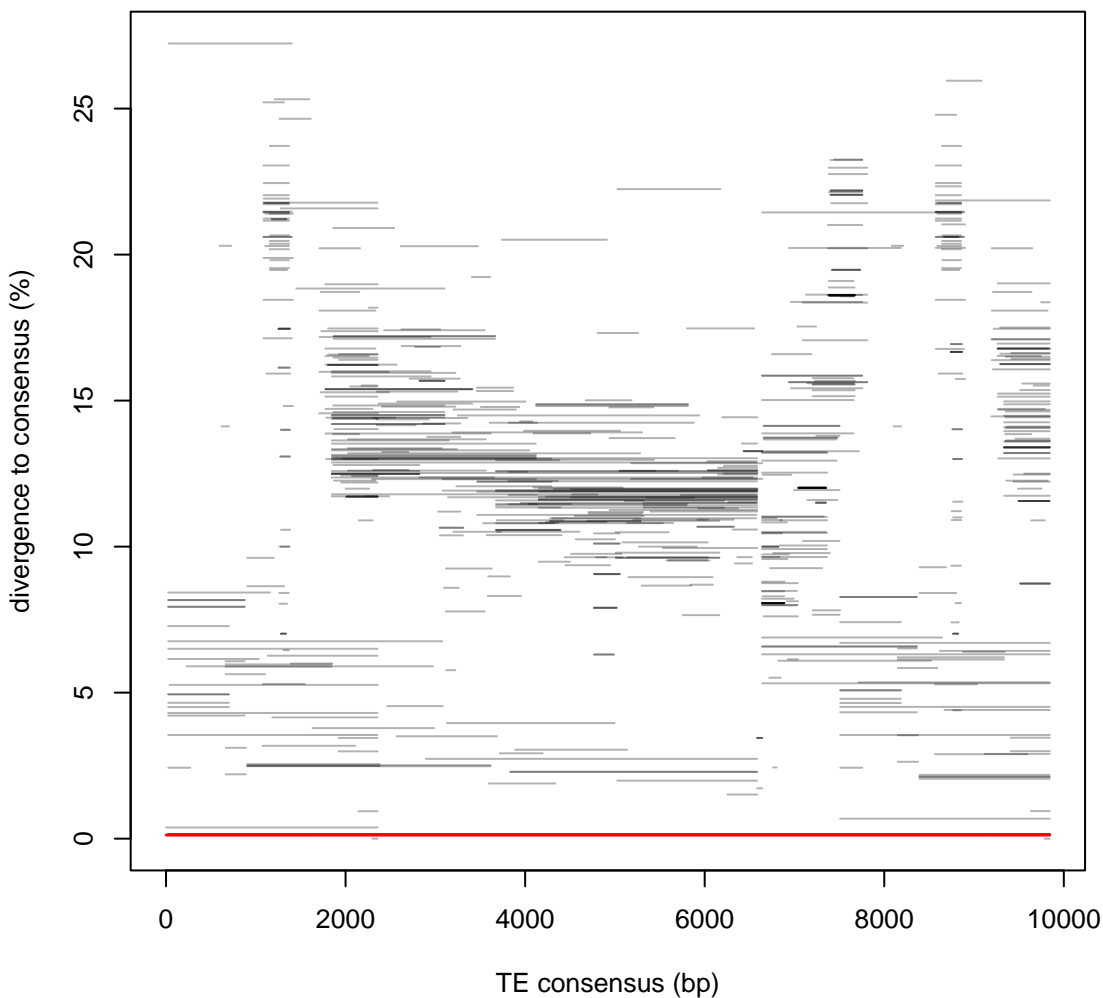
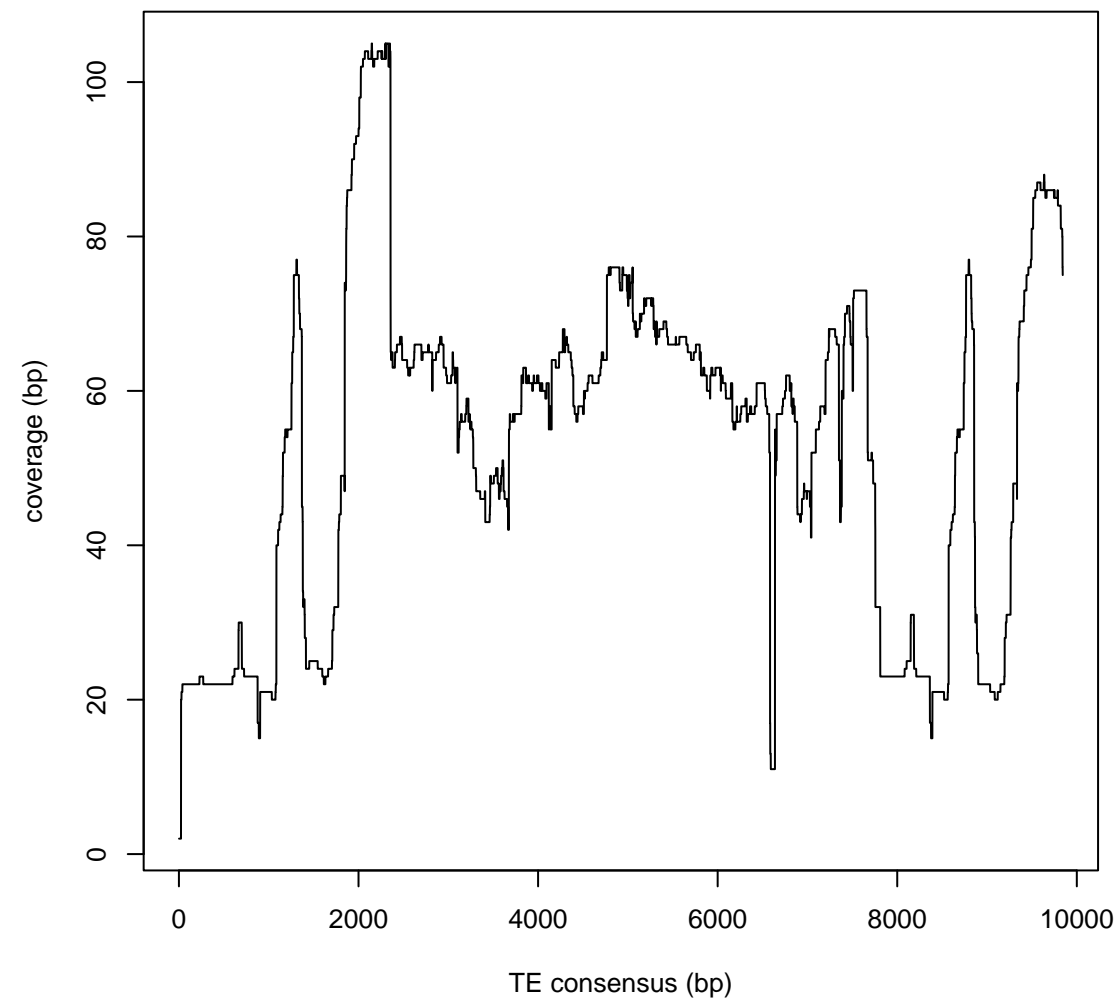


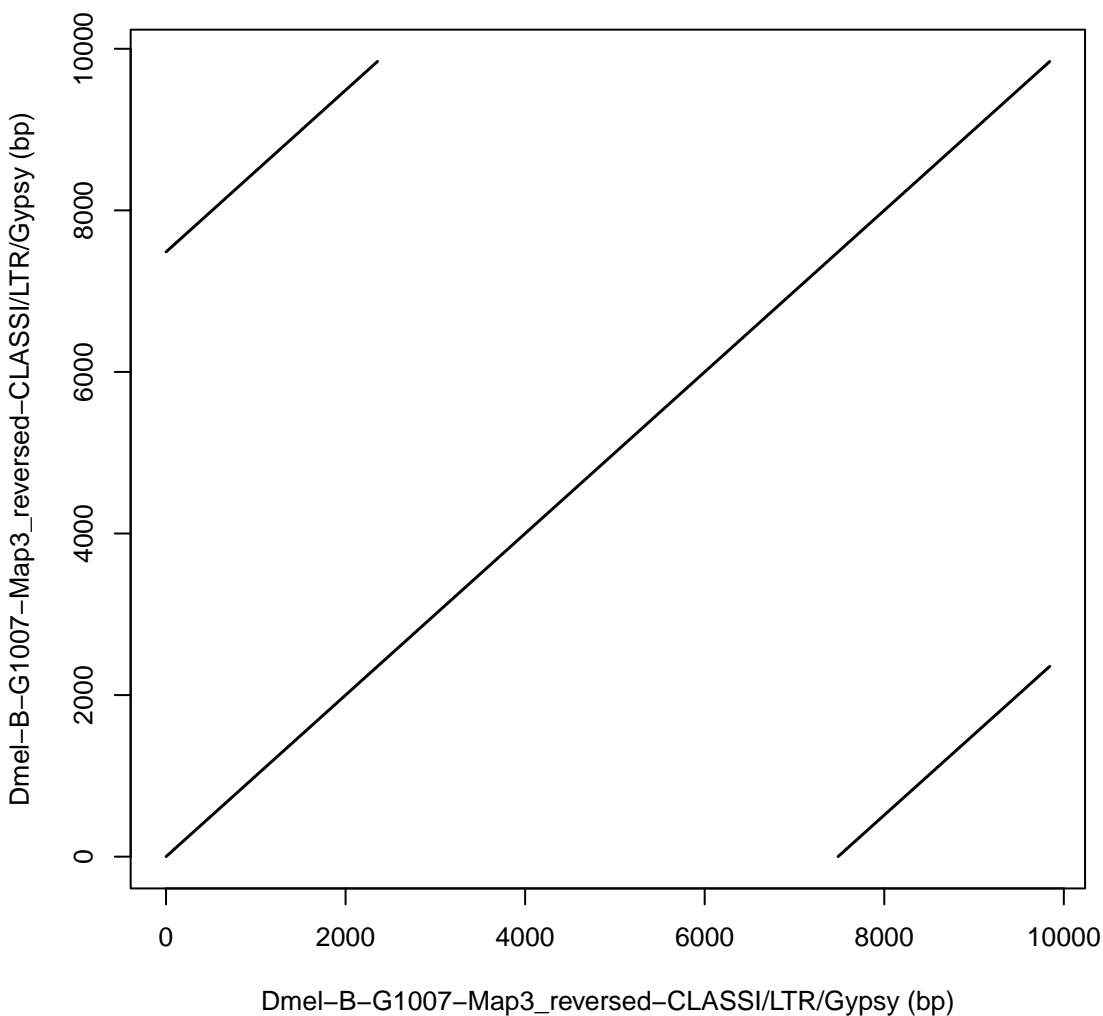
TE: Dmel-B-G1007-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 9843bp; fragments: 676; full length: 2 (>=8858.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

