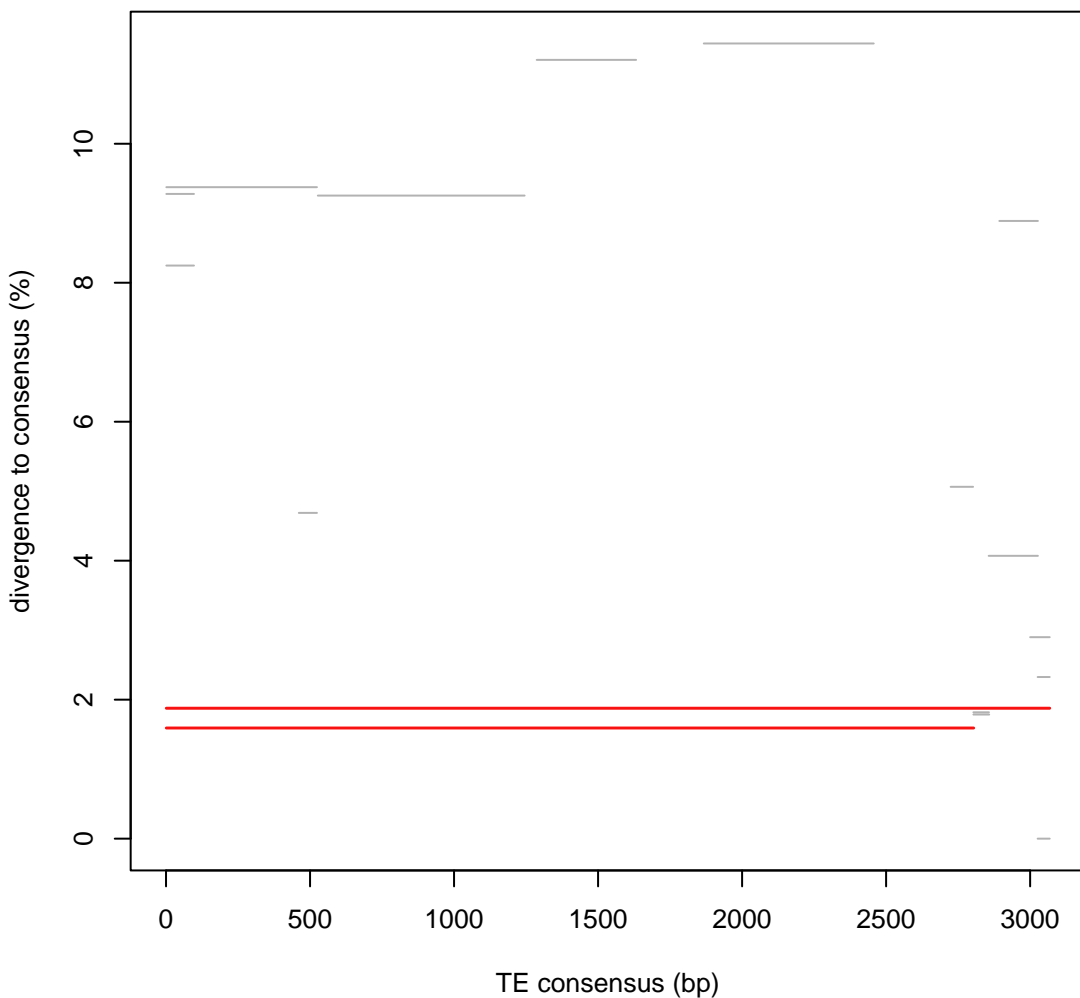
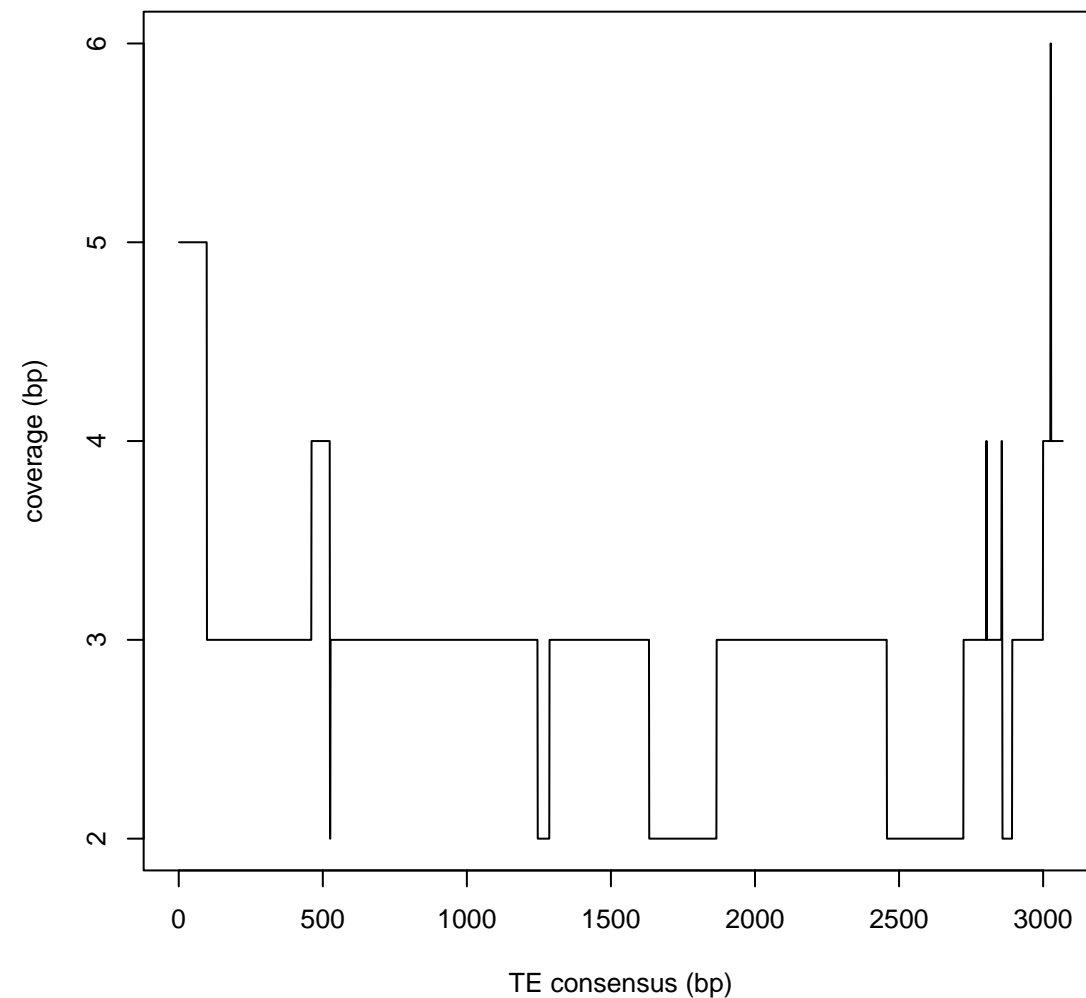


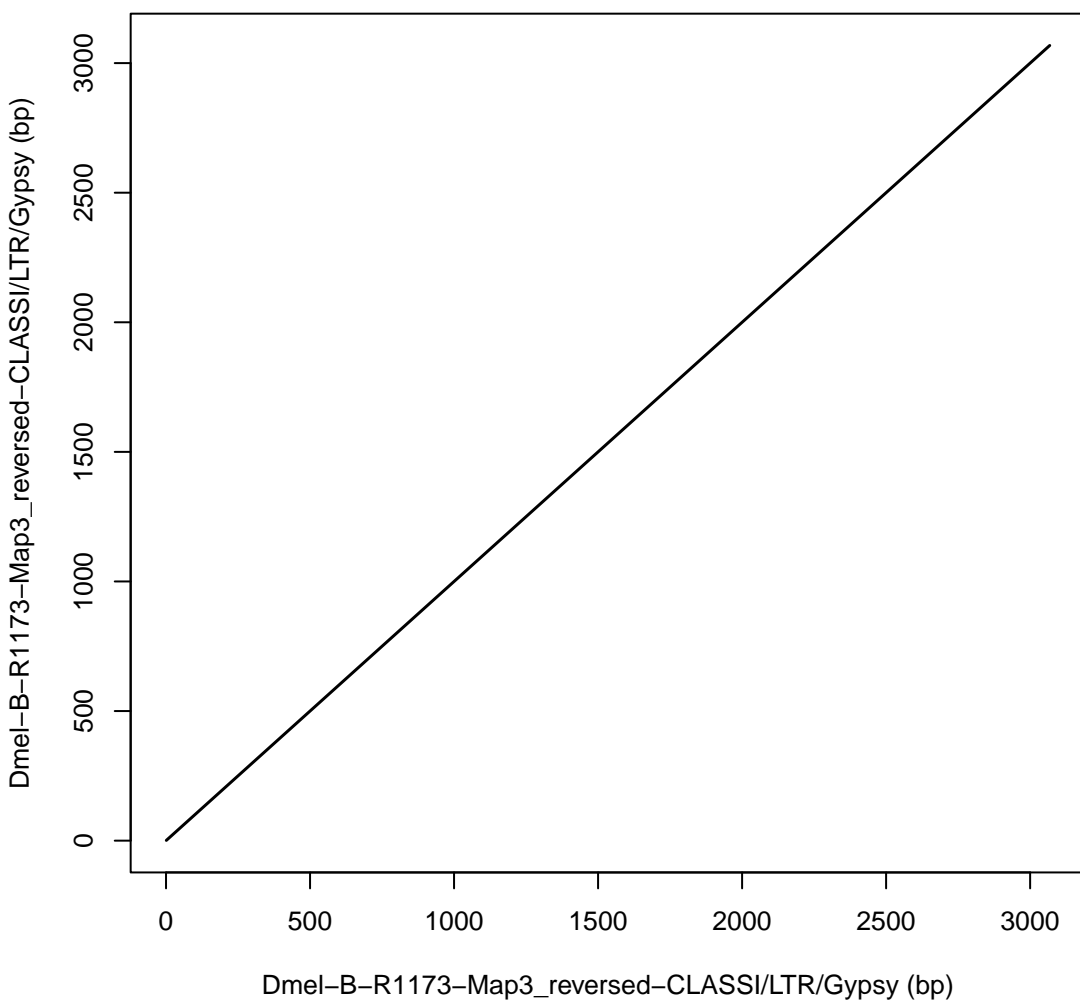
TE: Dmel-B-R1173-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 3068bp; fragments: 17; full length: 2 (≥ 2761.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

