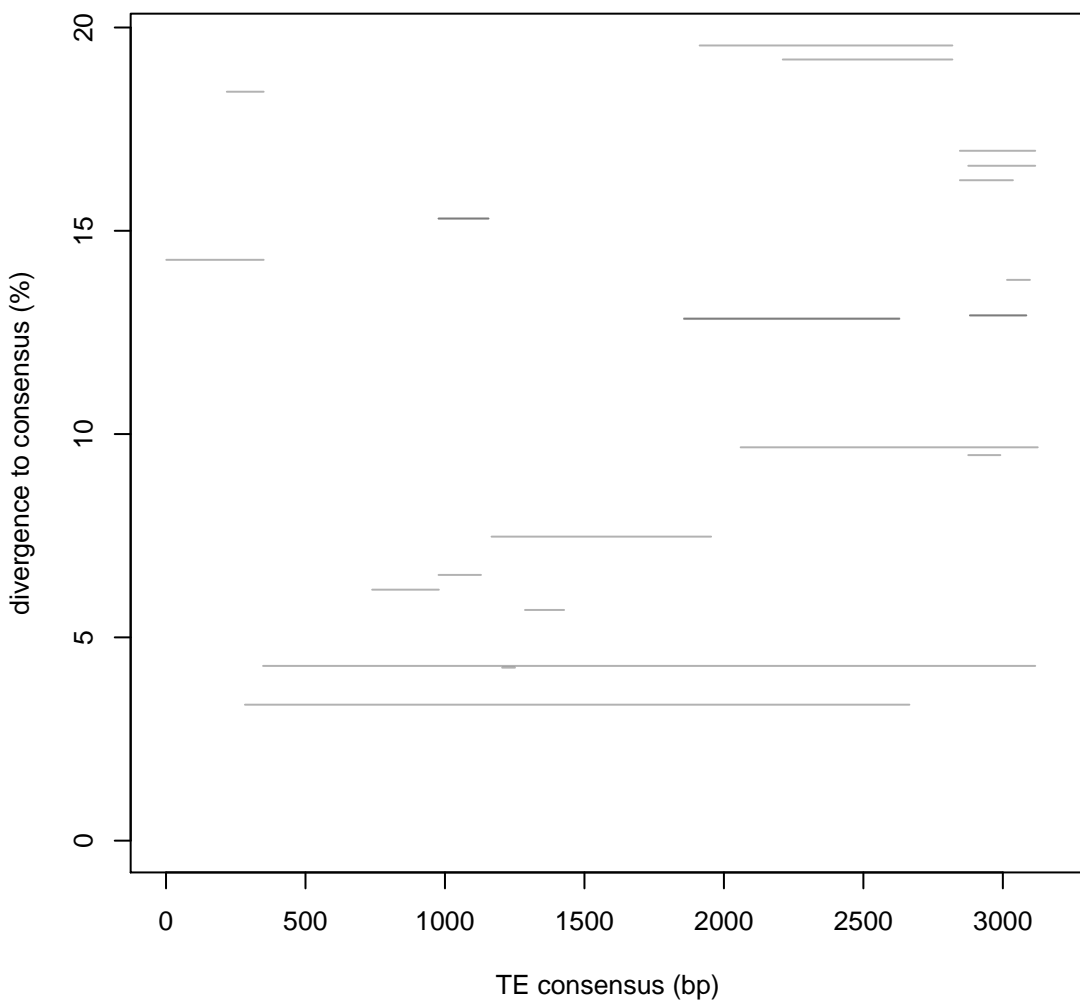
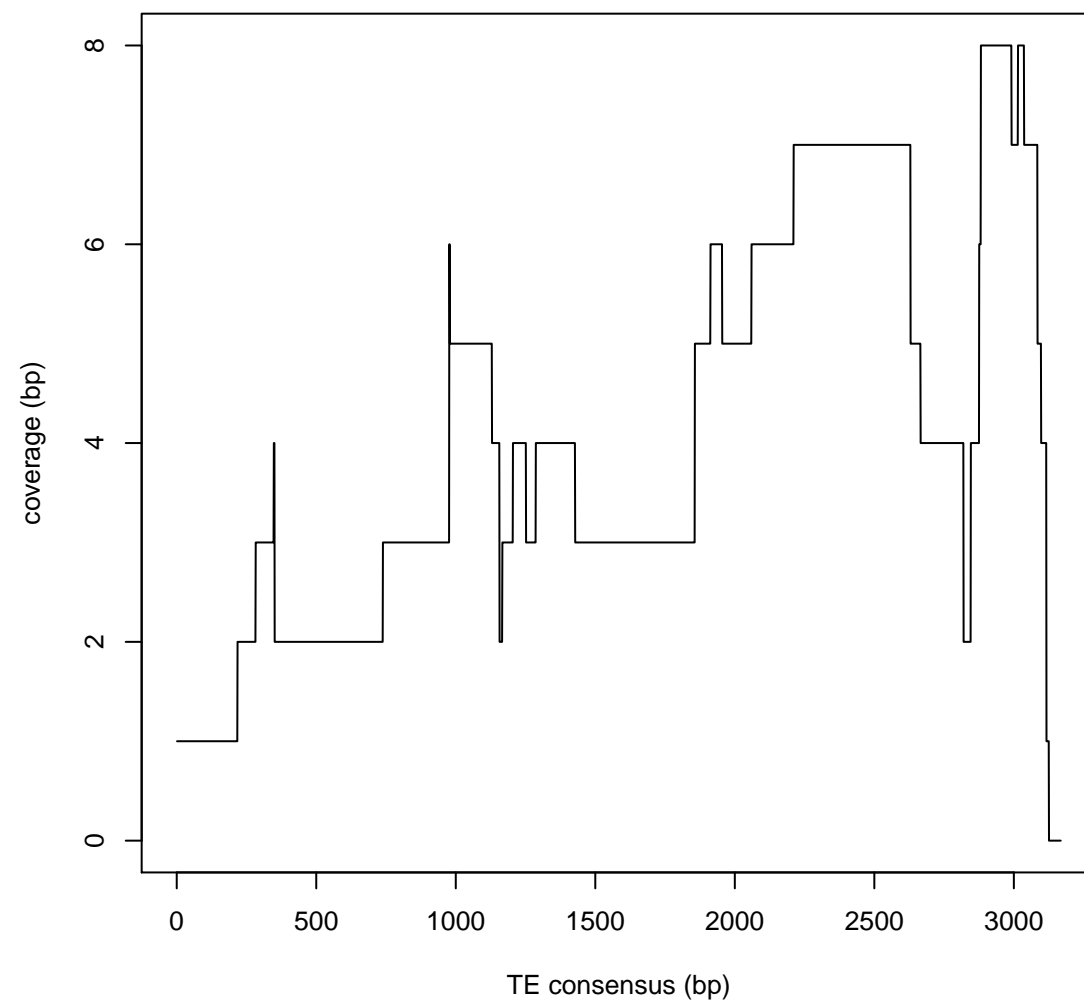


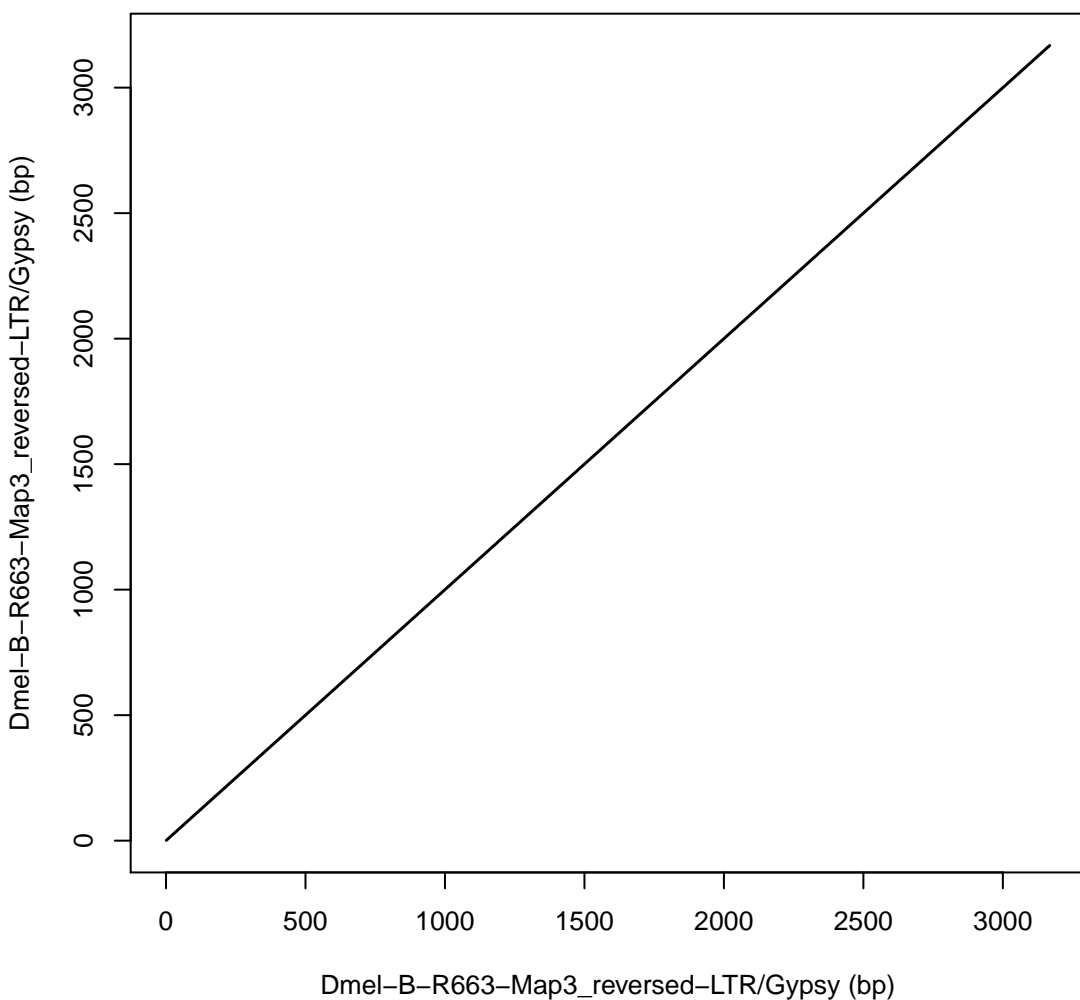
TE: Dmel-B-R663-Map3_reversed-LTR/Gypsy
consensus size: 3168bp; fragments: 23; full length: 0 (≥ 2851.2 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

