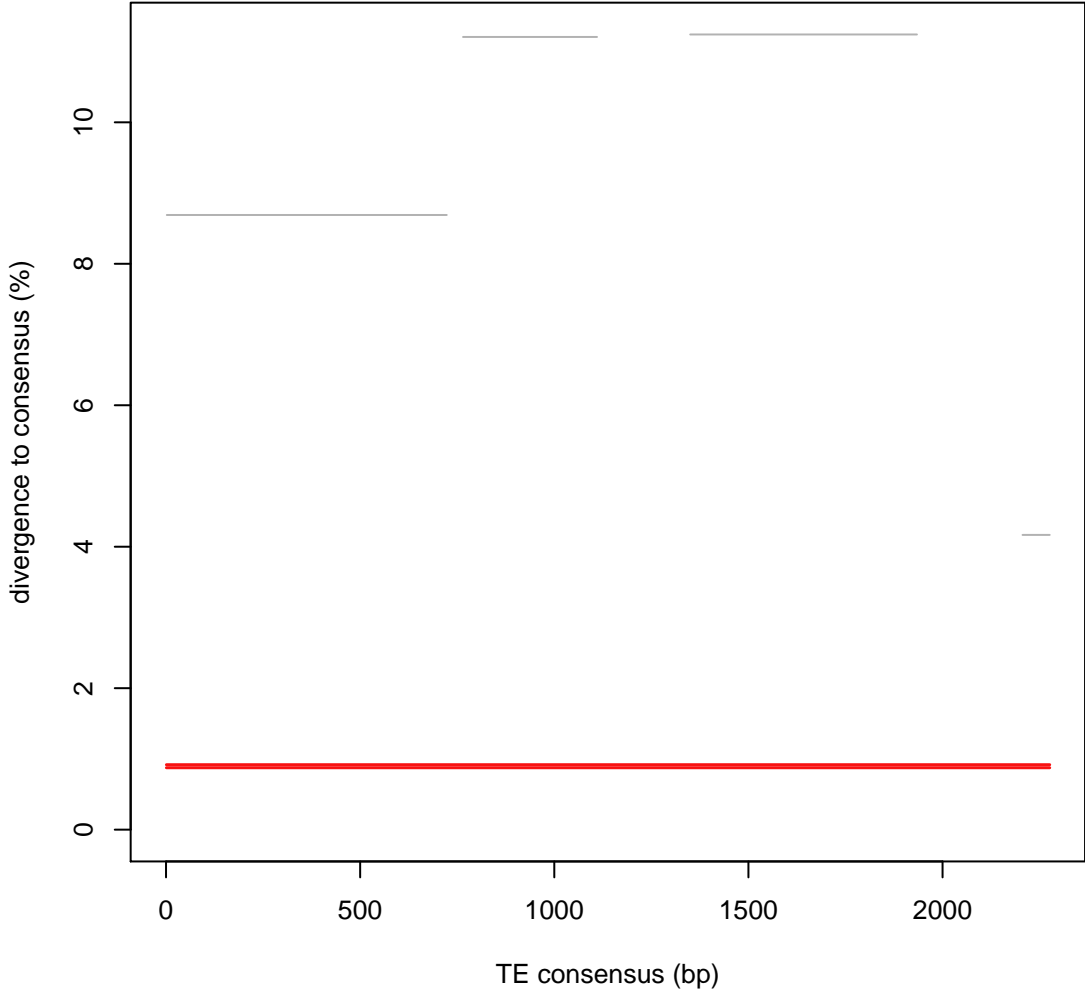
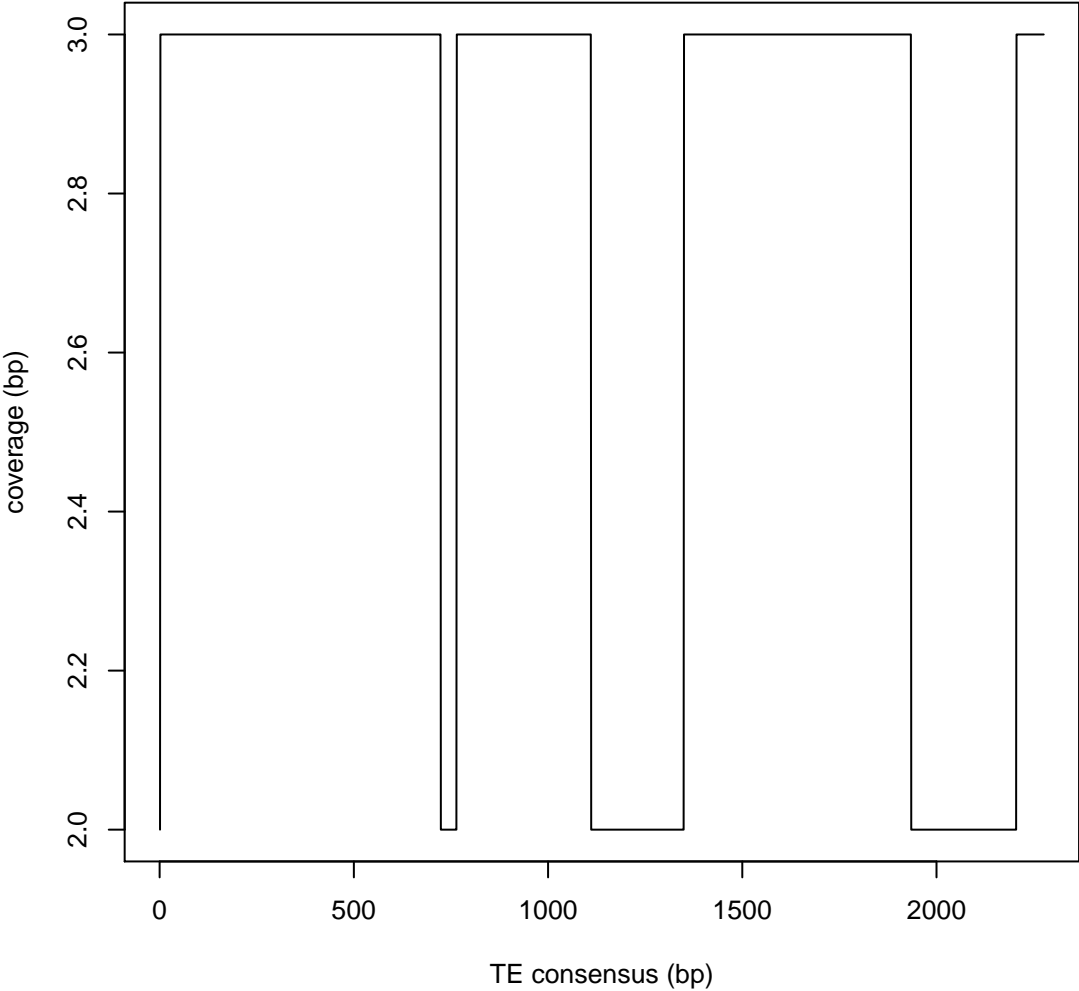


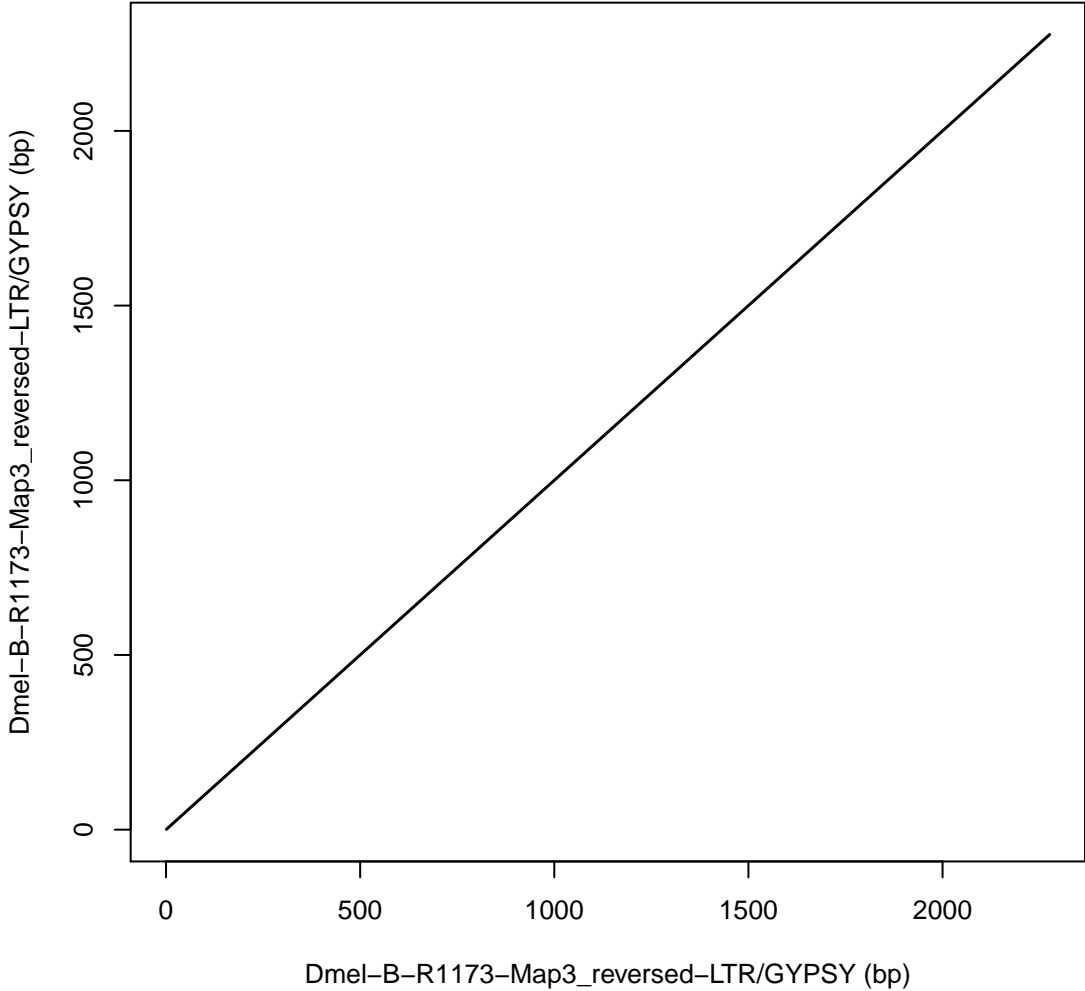
TE: Dmel-B-R1173-Map3_reversed-LTR/GYPSY
consensus size: 2276bp; fragments: 6; full length: 2 (>=2048.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

