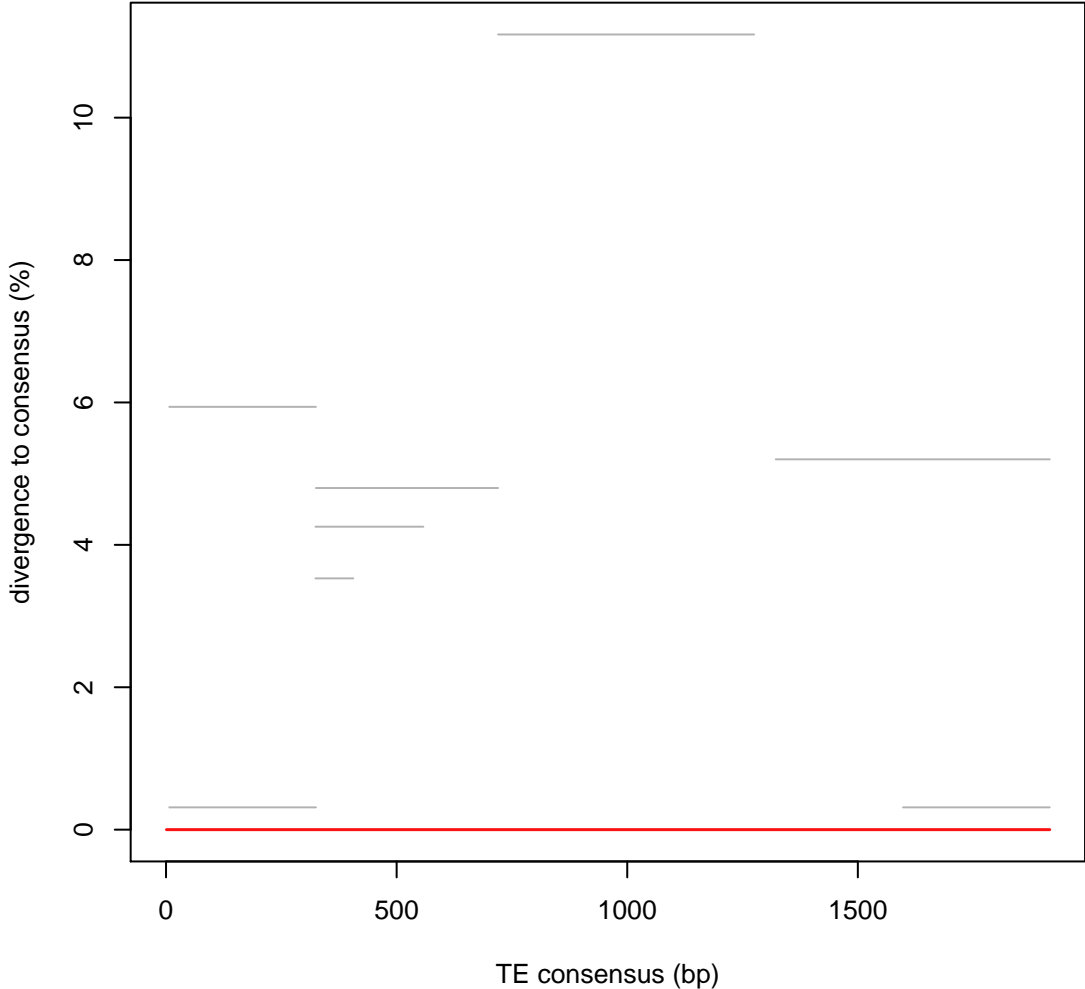
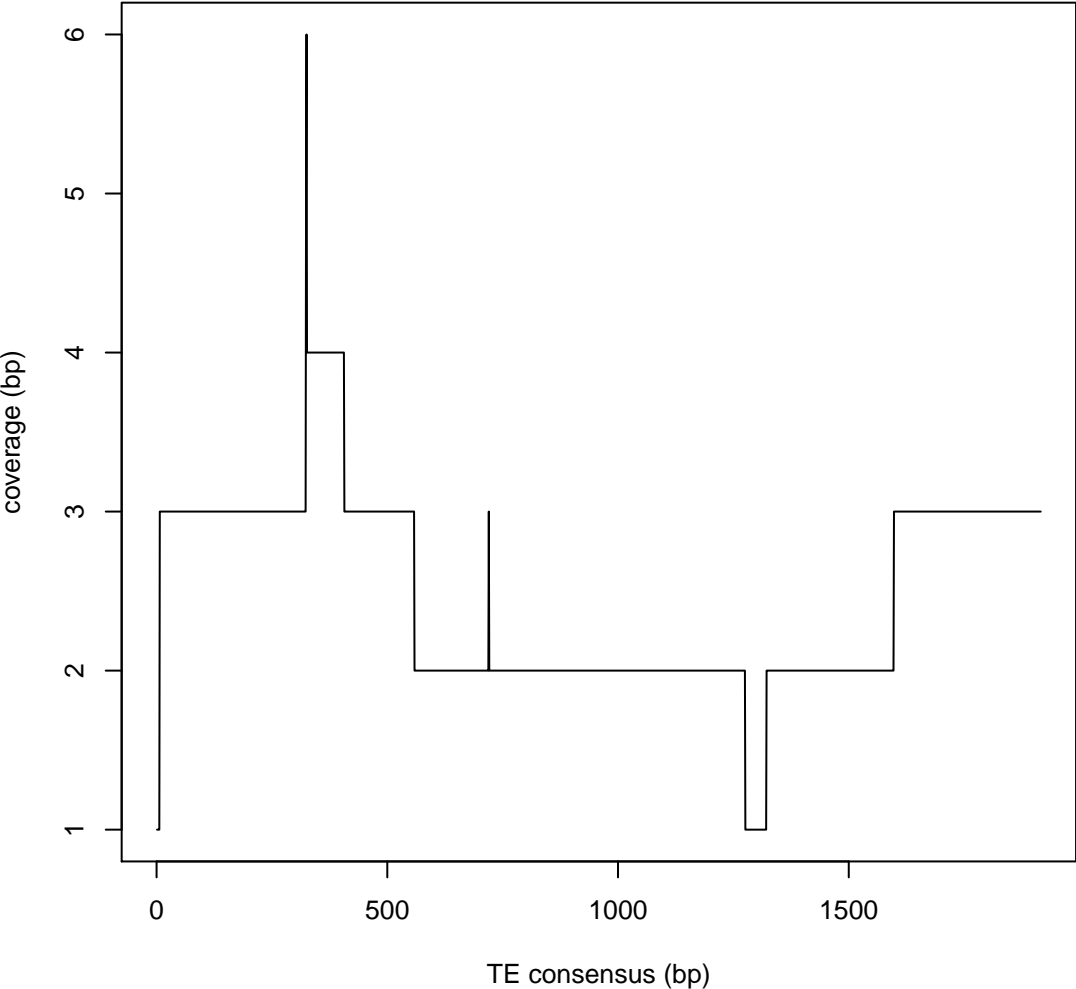


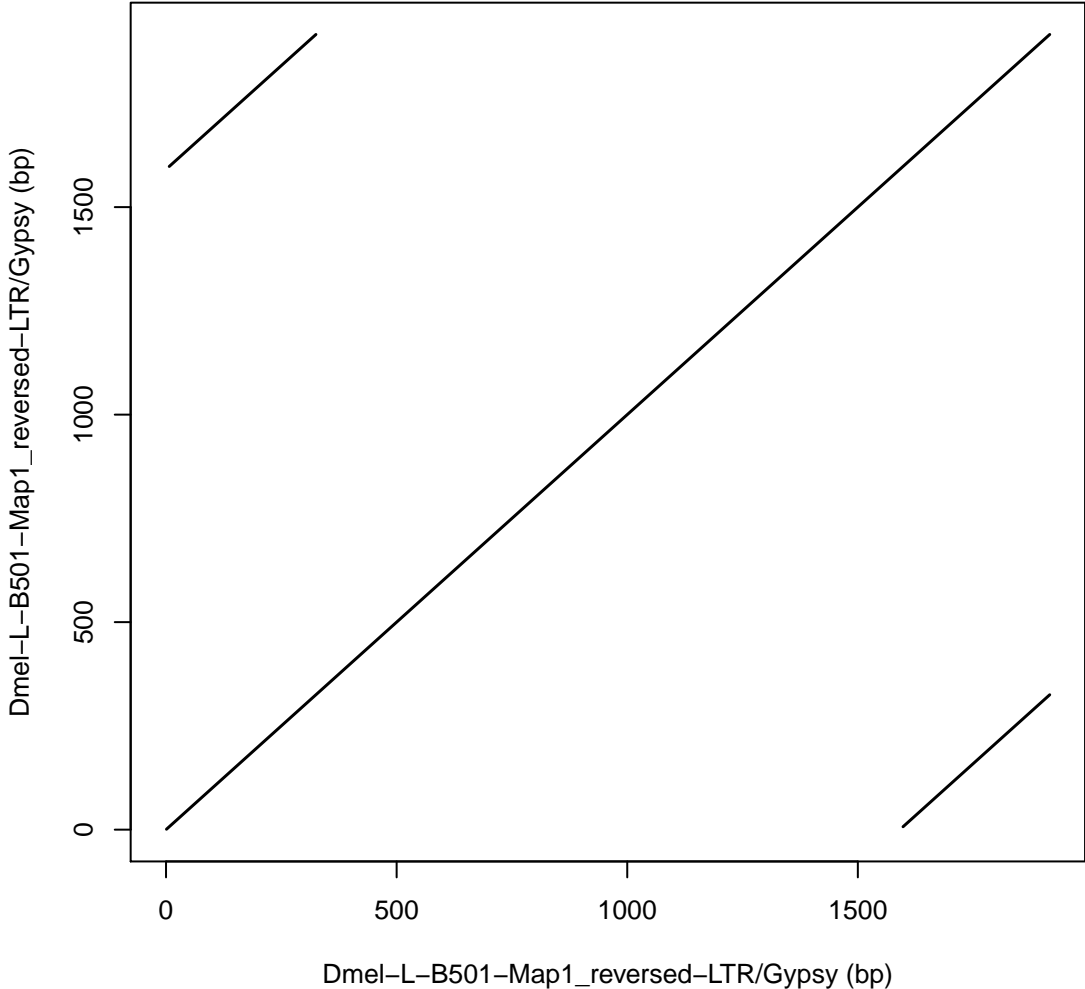
TE: Dmel-L-B501-Map1_reversed-LTR/Gypsy
consensus size: 1916bp; fragments: 9; full length: 1 (>=1724.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

