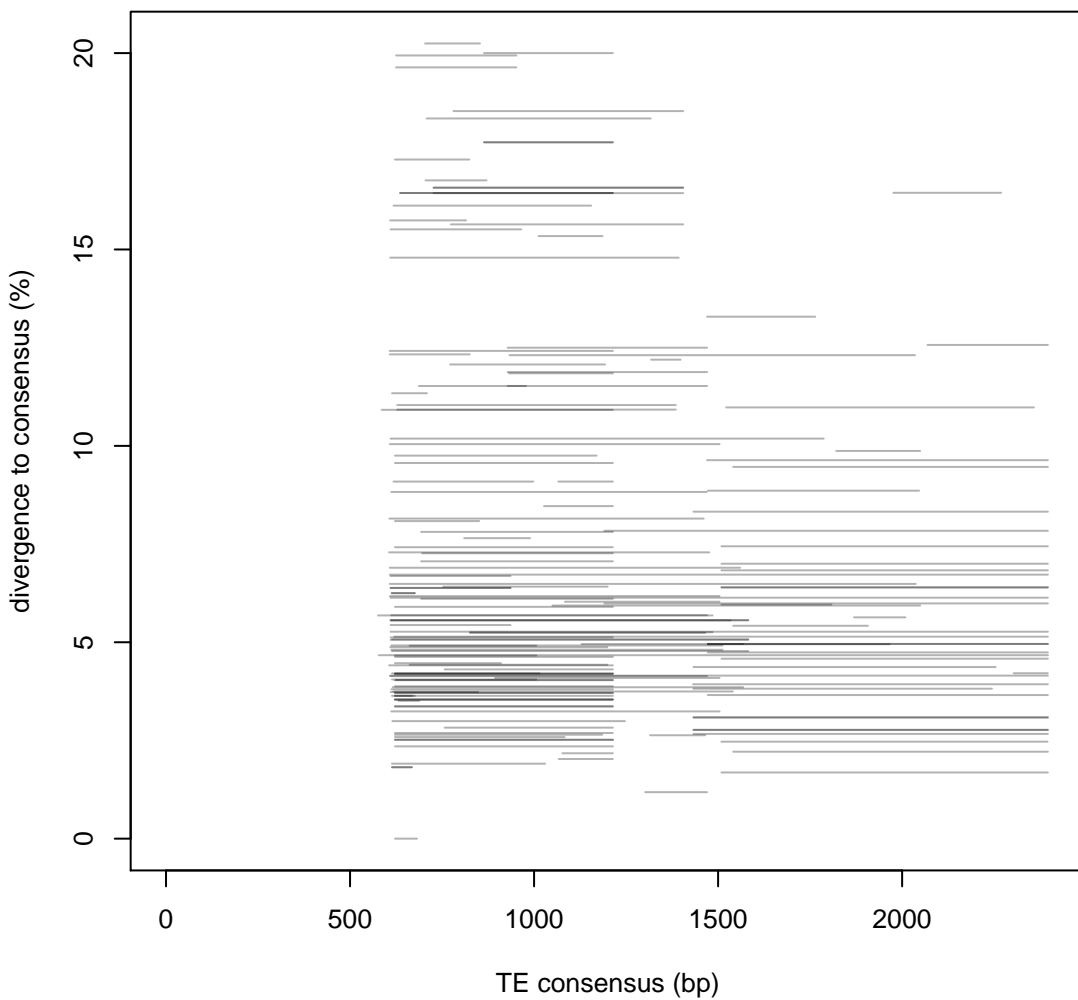
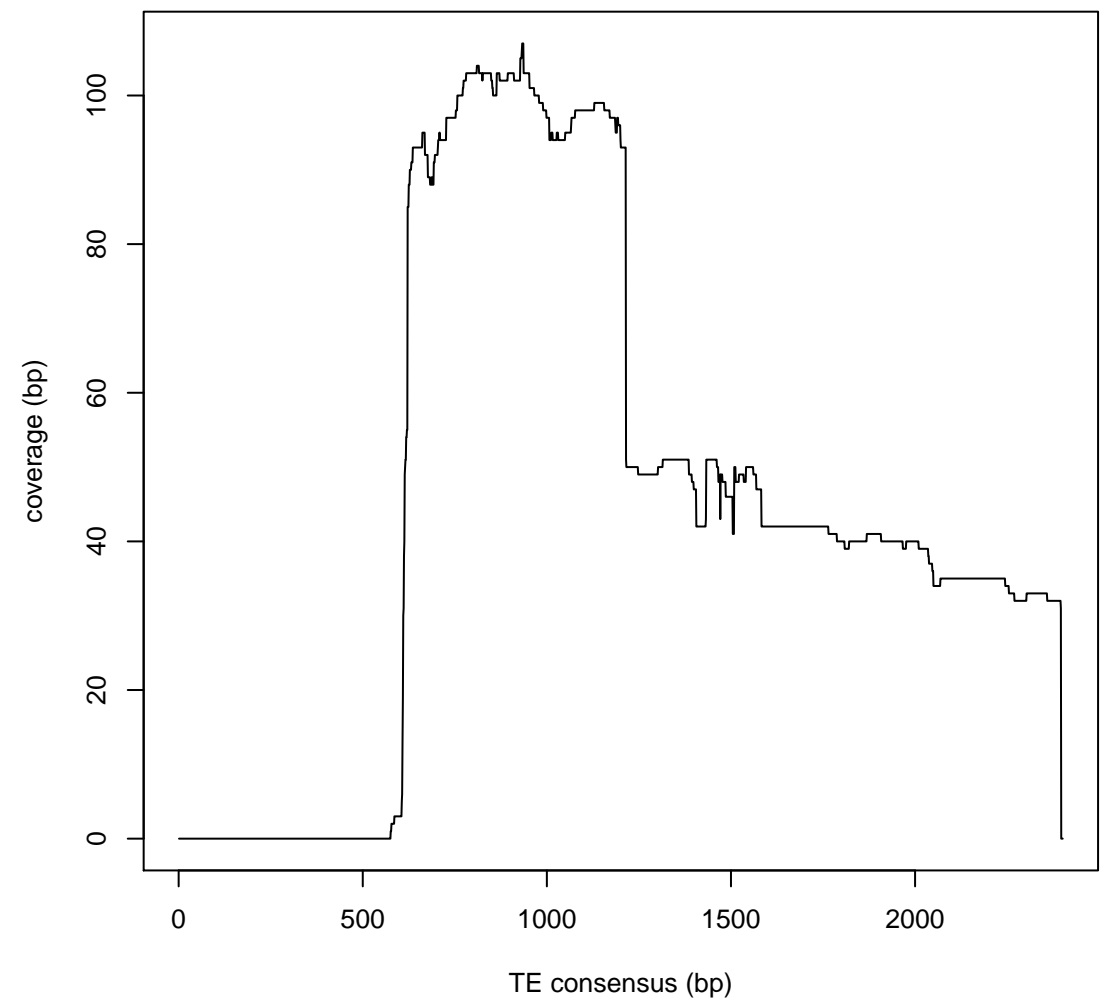


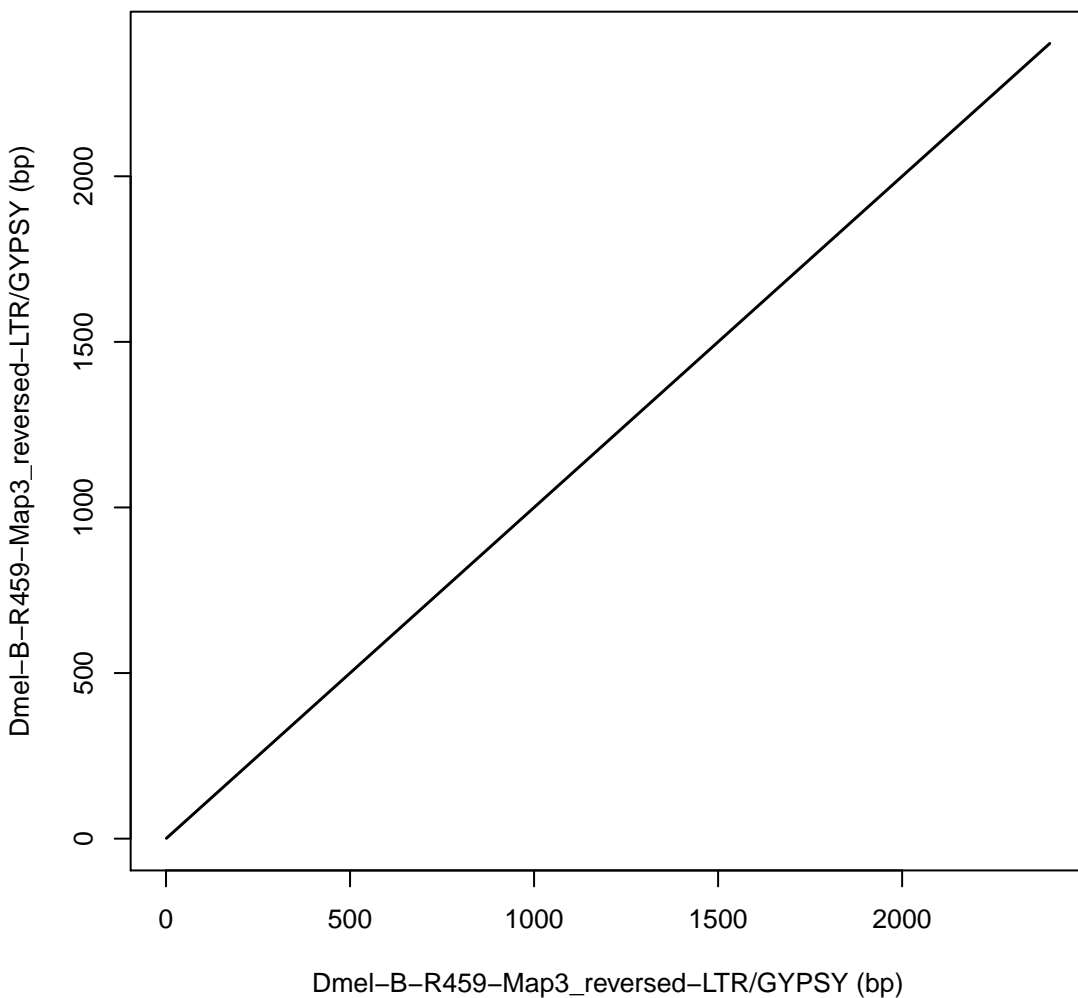
TE: Dmel-B-R459-Map3_reversed-LTR/Gypsy
consensus size: 2401bp; fragments: 172; full length: 0 (>=2160.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

