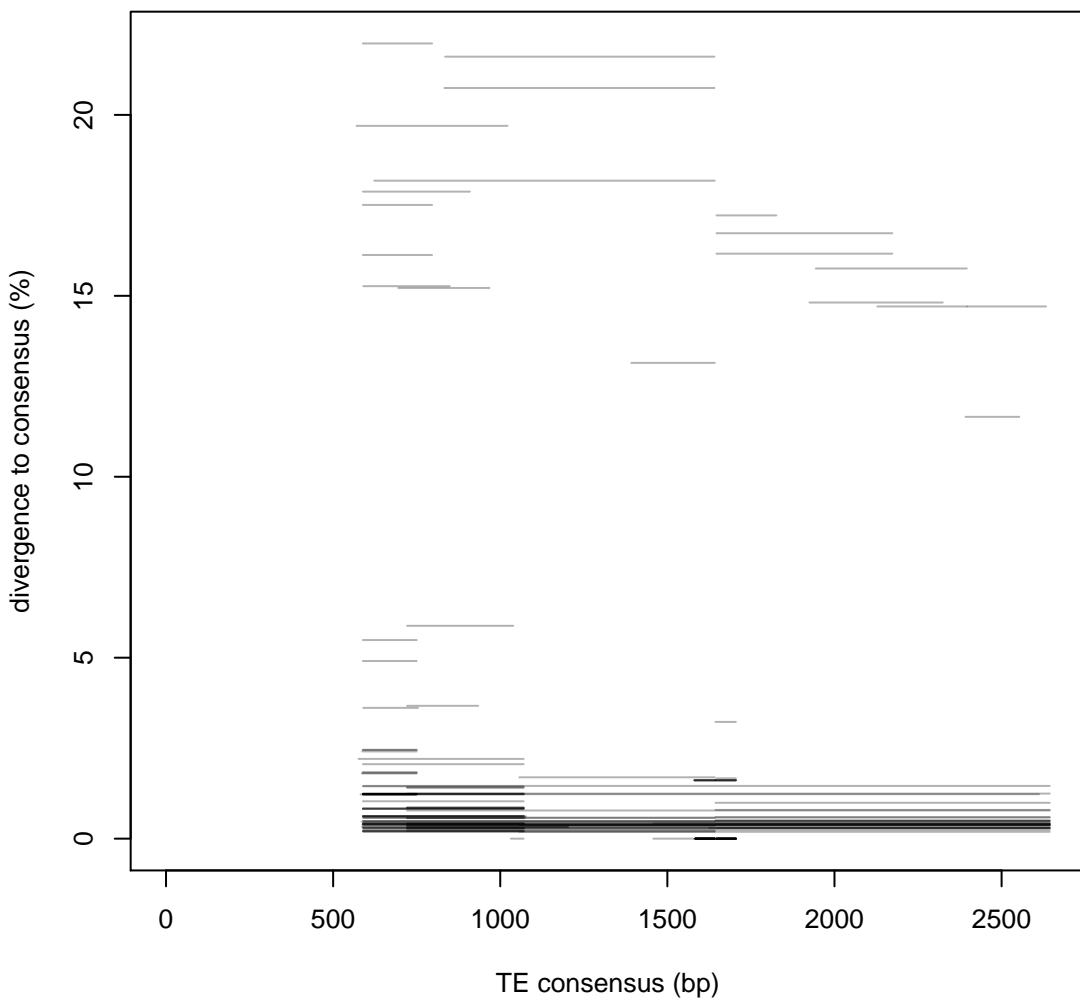
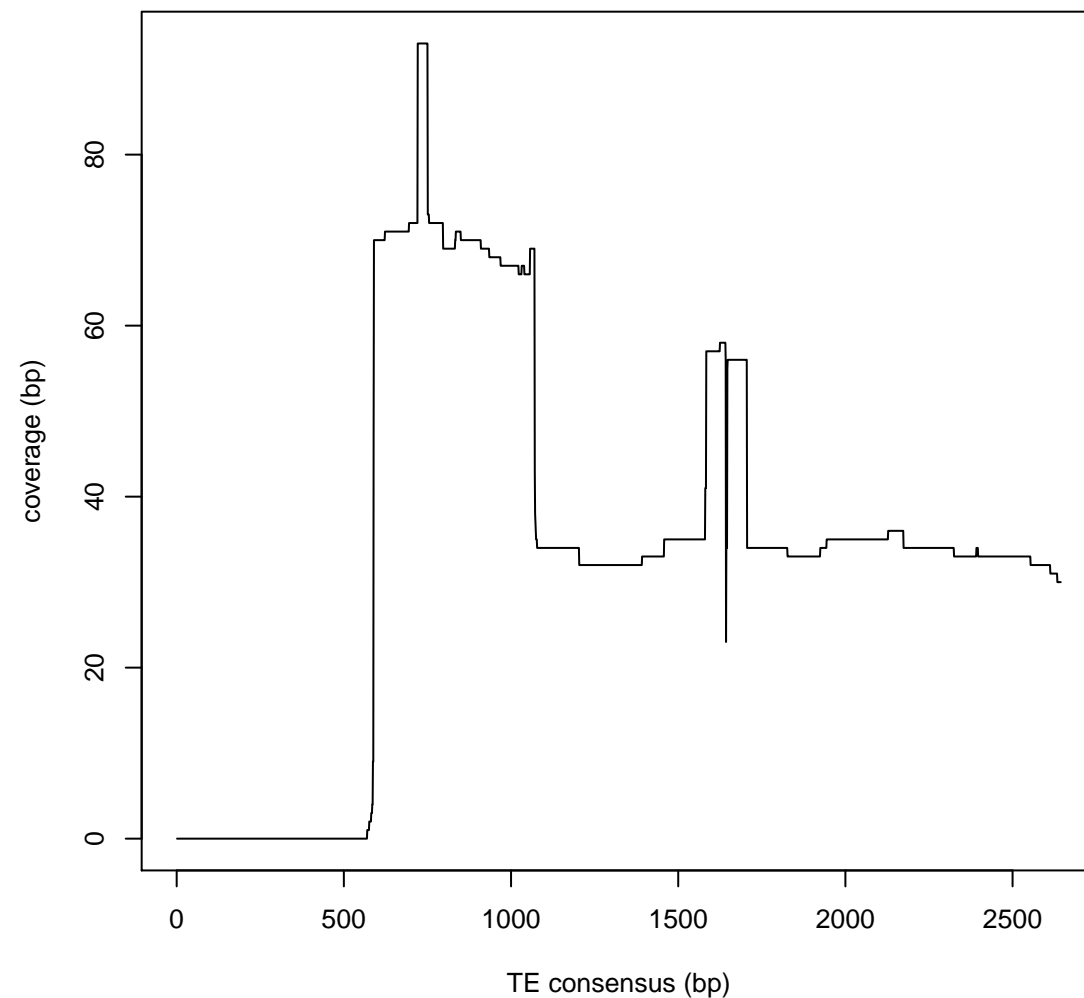


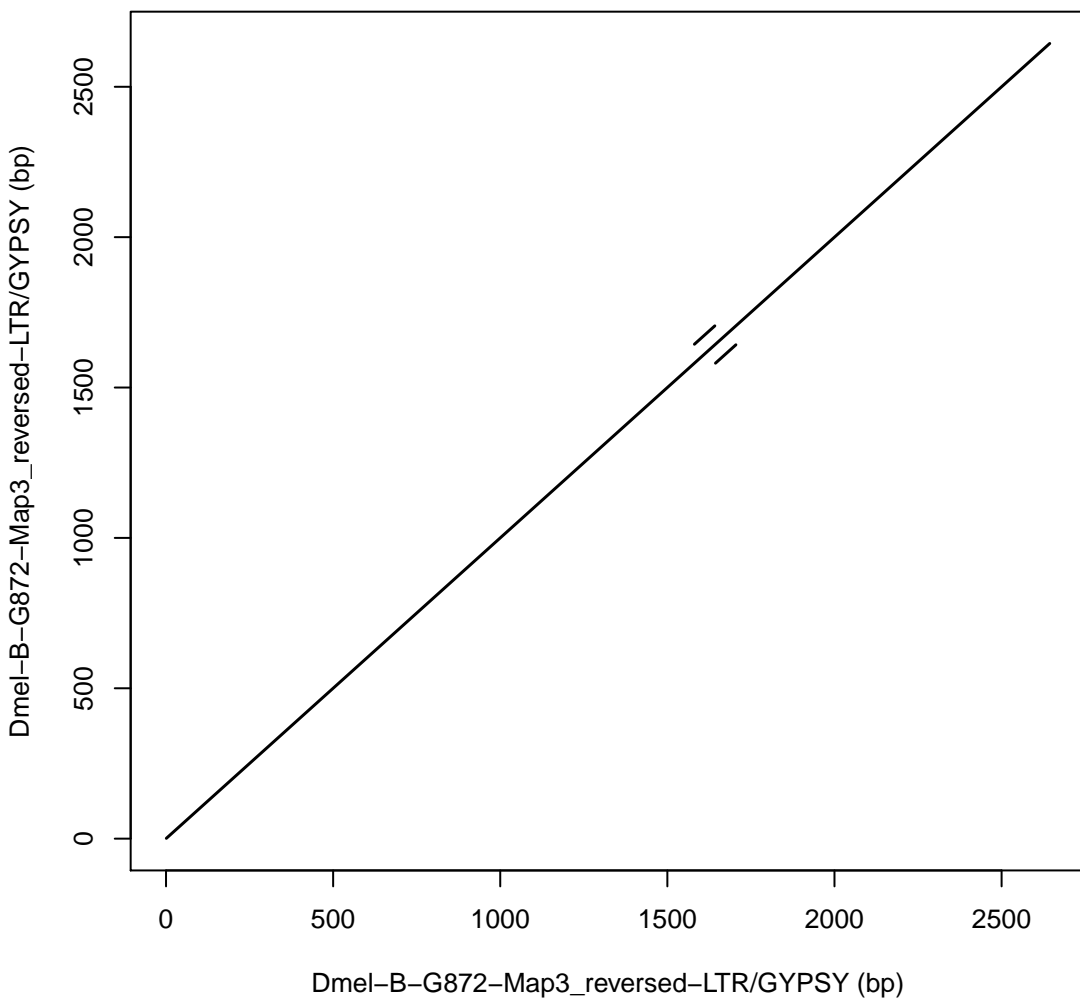
TE: Dmel-B-G872-Map3_reversed-LTR/Gypsy
consensus size: 2644bp; fragments: 163; full length: 0 (>=2379.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

