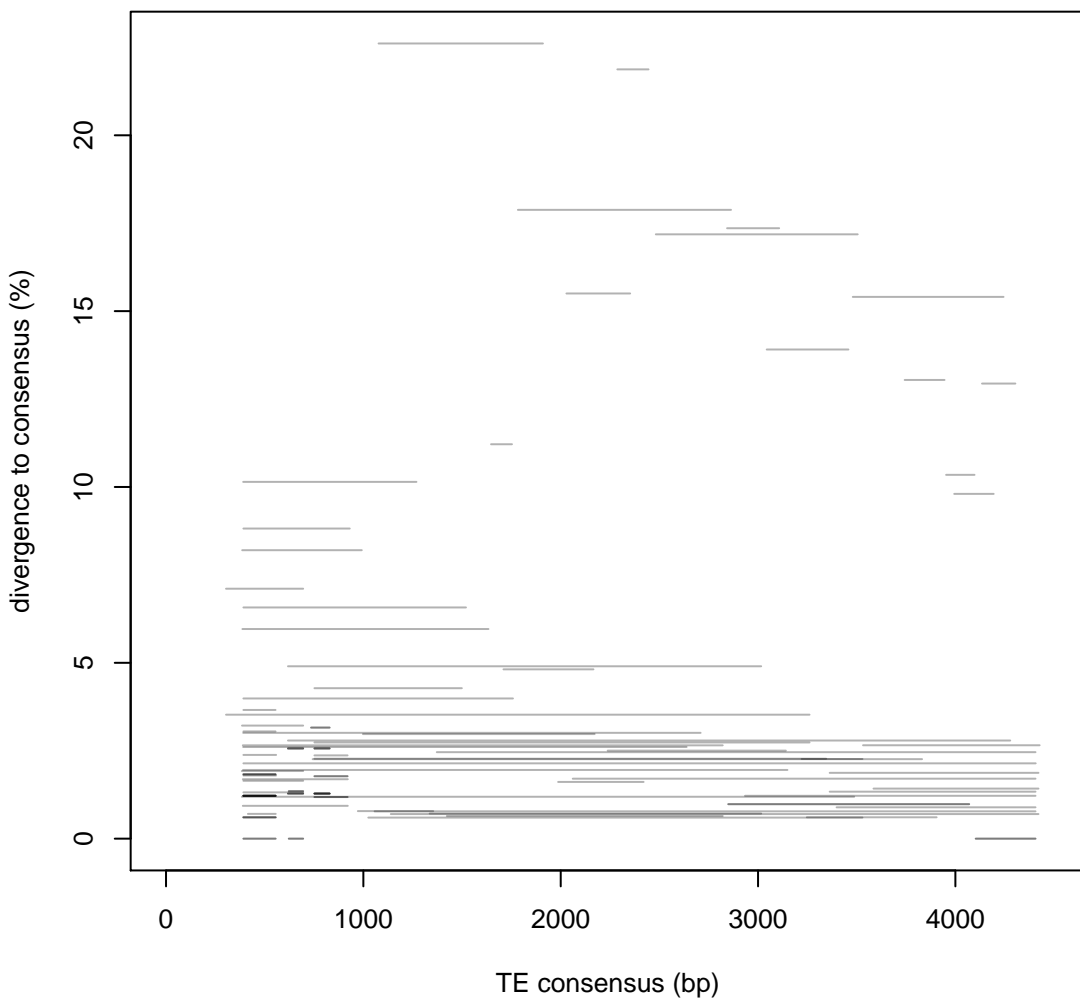
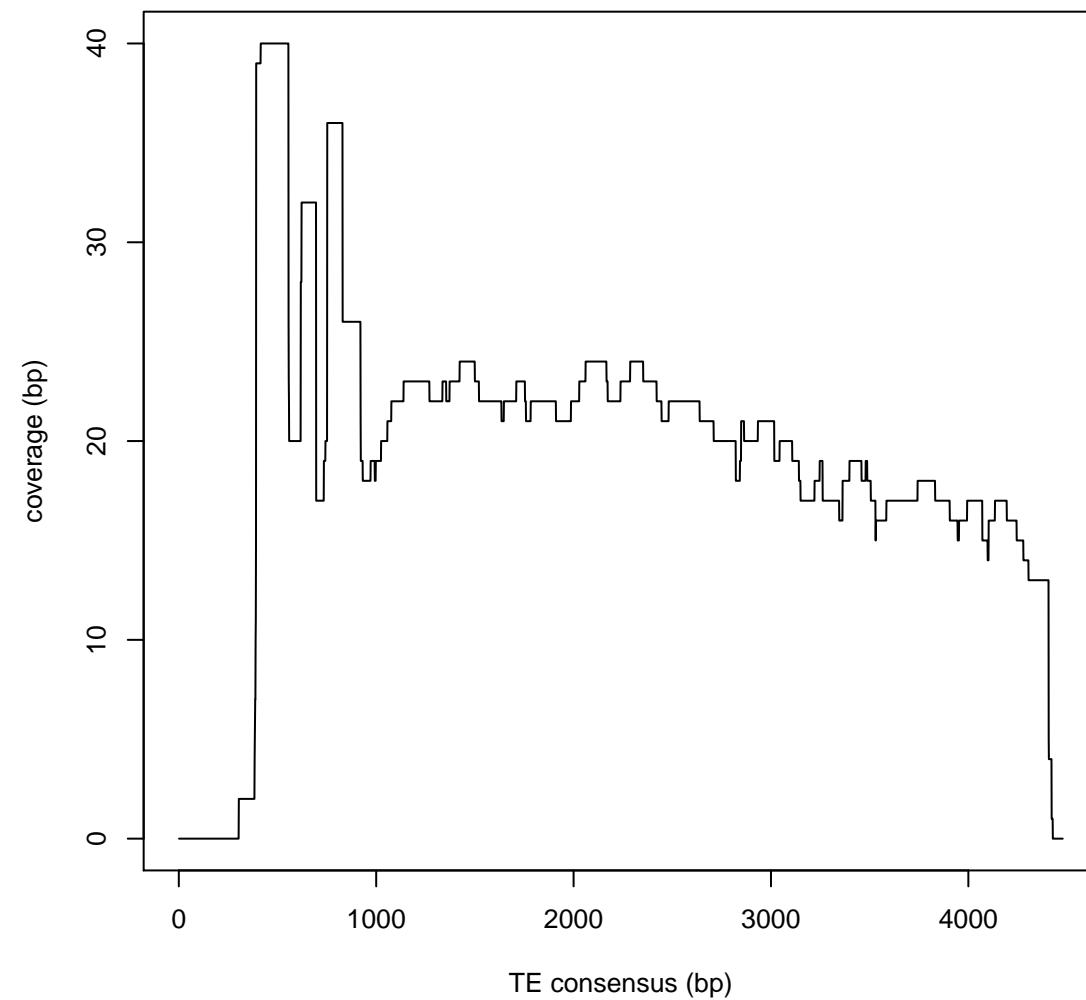


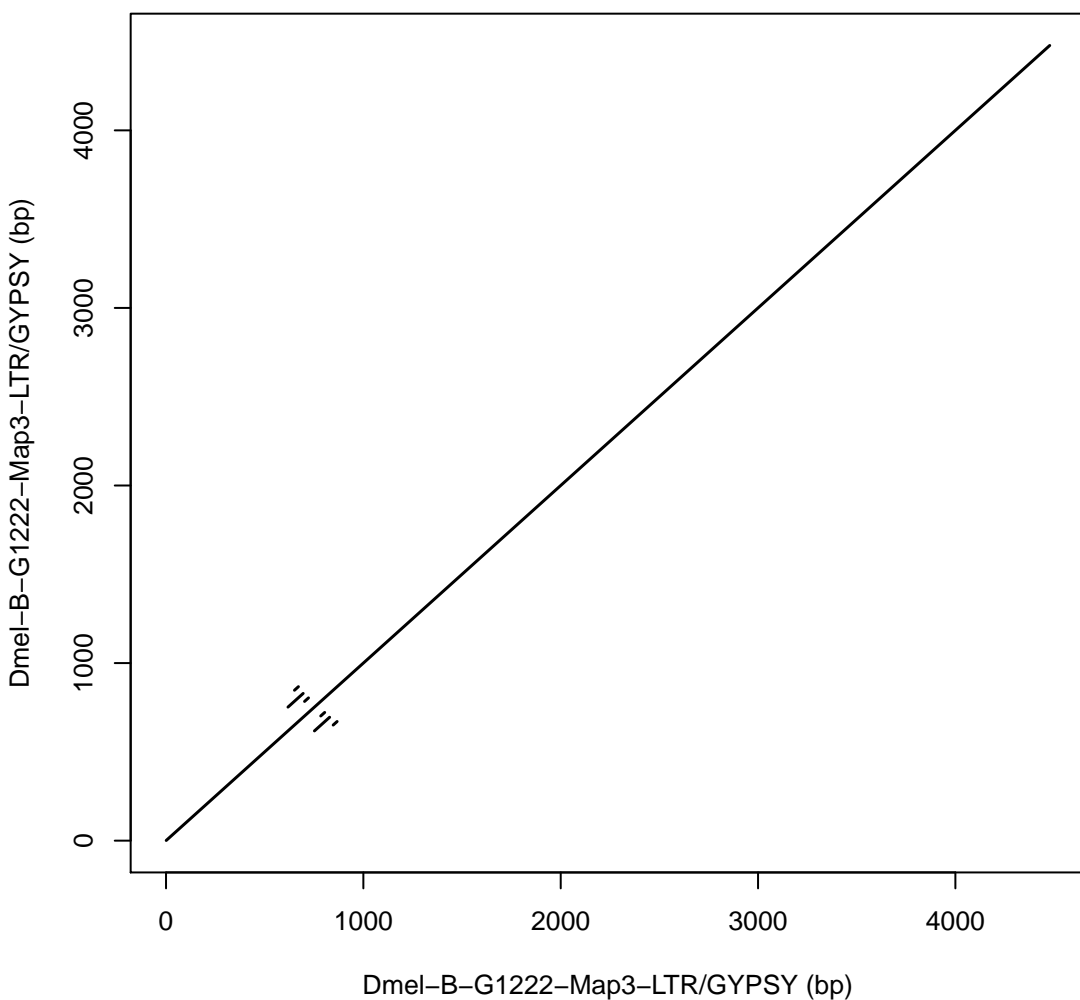
TE: Dmel-B-G1222-Map3-LTR/GYPSY
consensus size: 4478bp; fragments: 108; full length: 0 (>=4030.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

