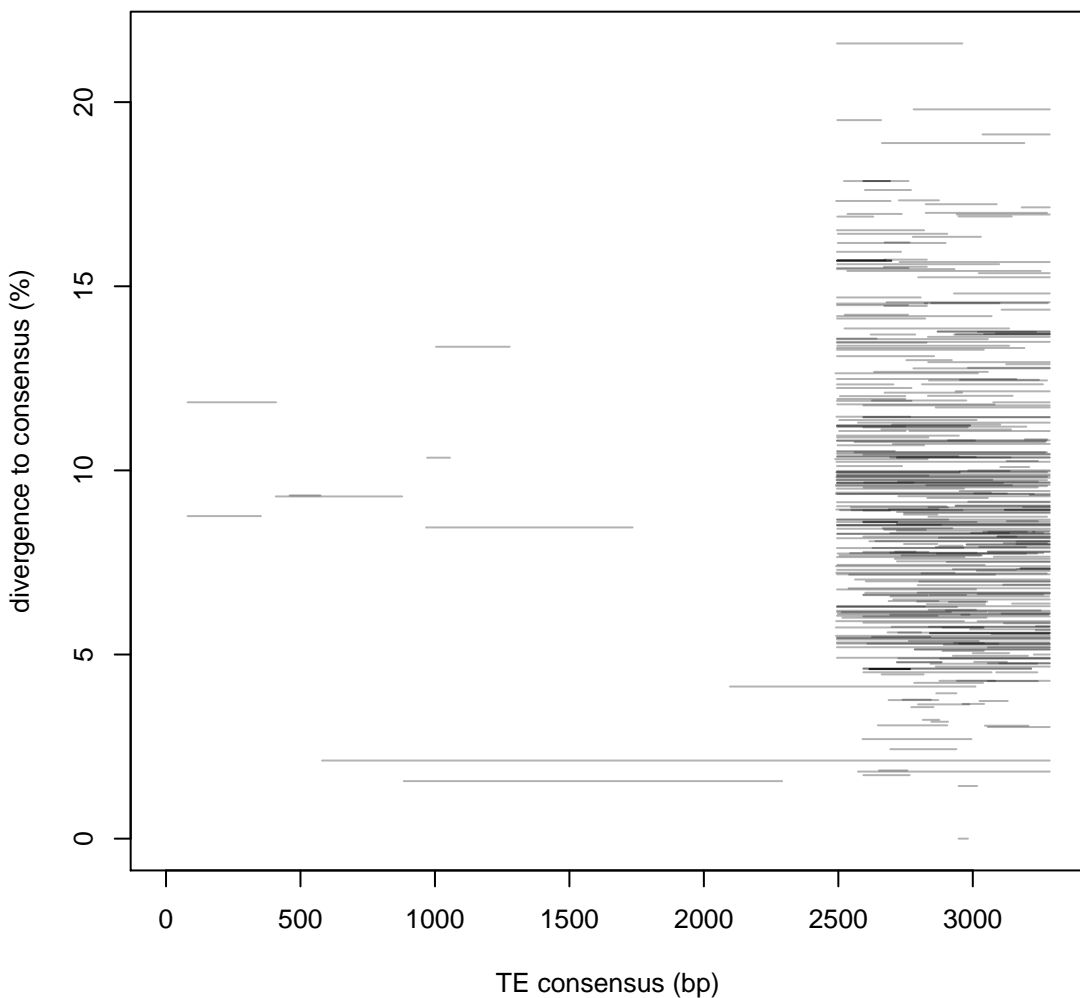
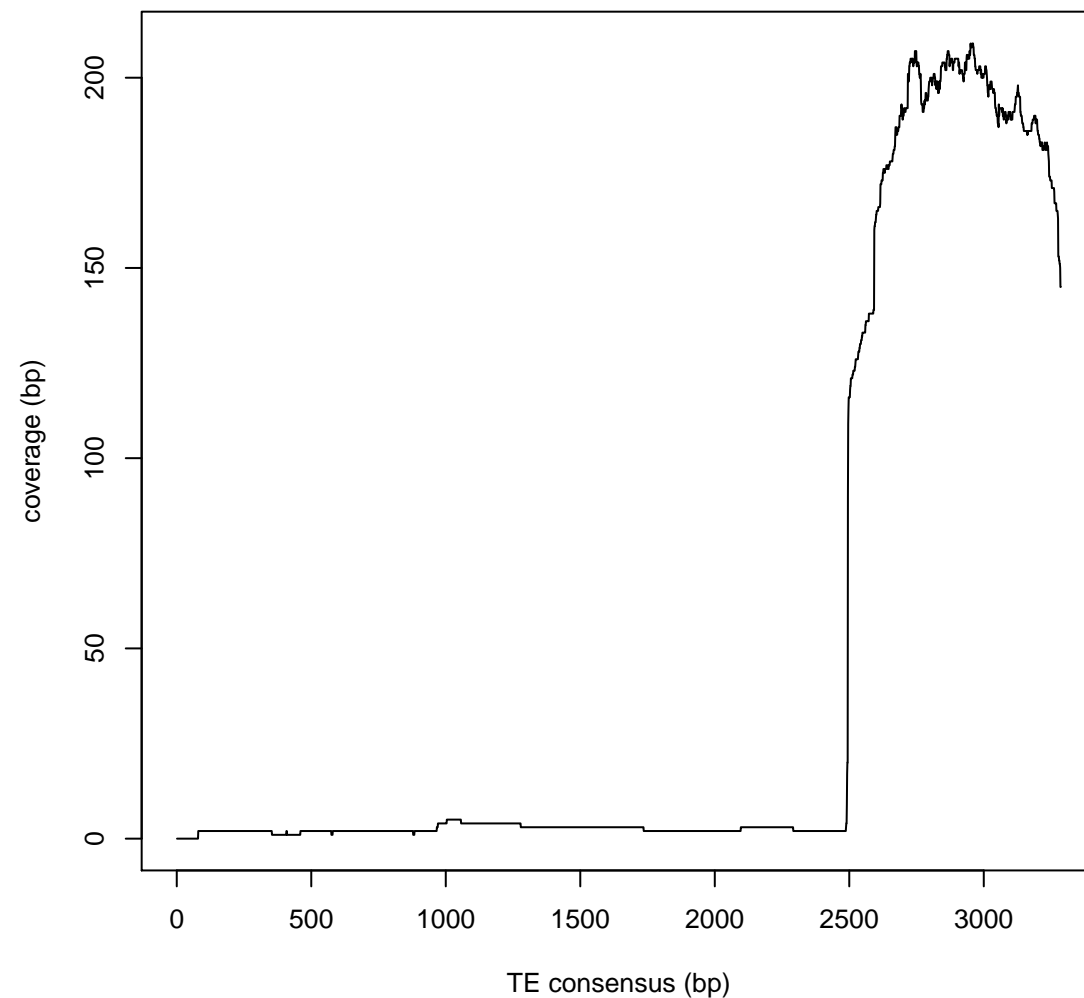


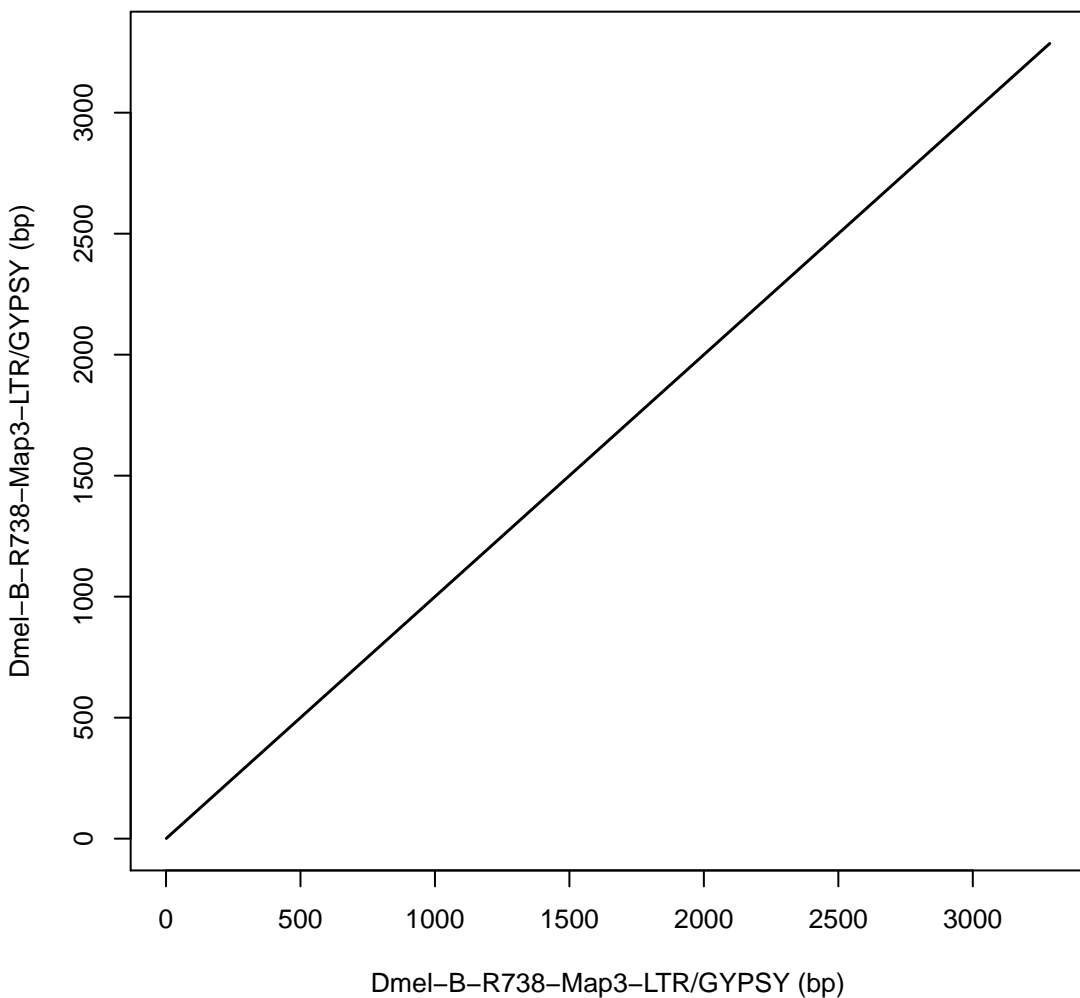
TE: Dmel-B-R738-Map3-LTR/GYPSY
consensus size: 3286bp; fragments: 375; full length: 0 (>=2957.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

