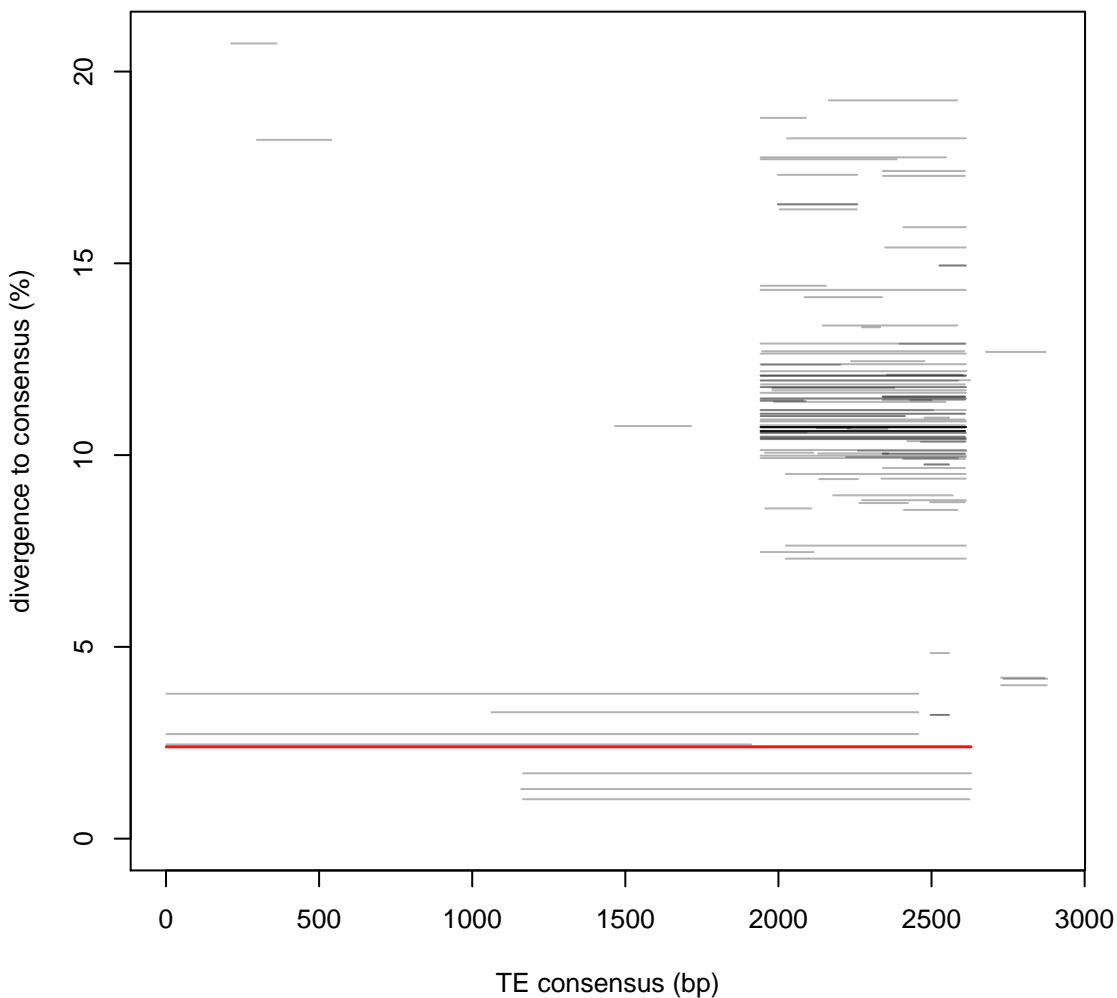
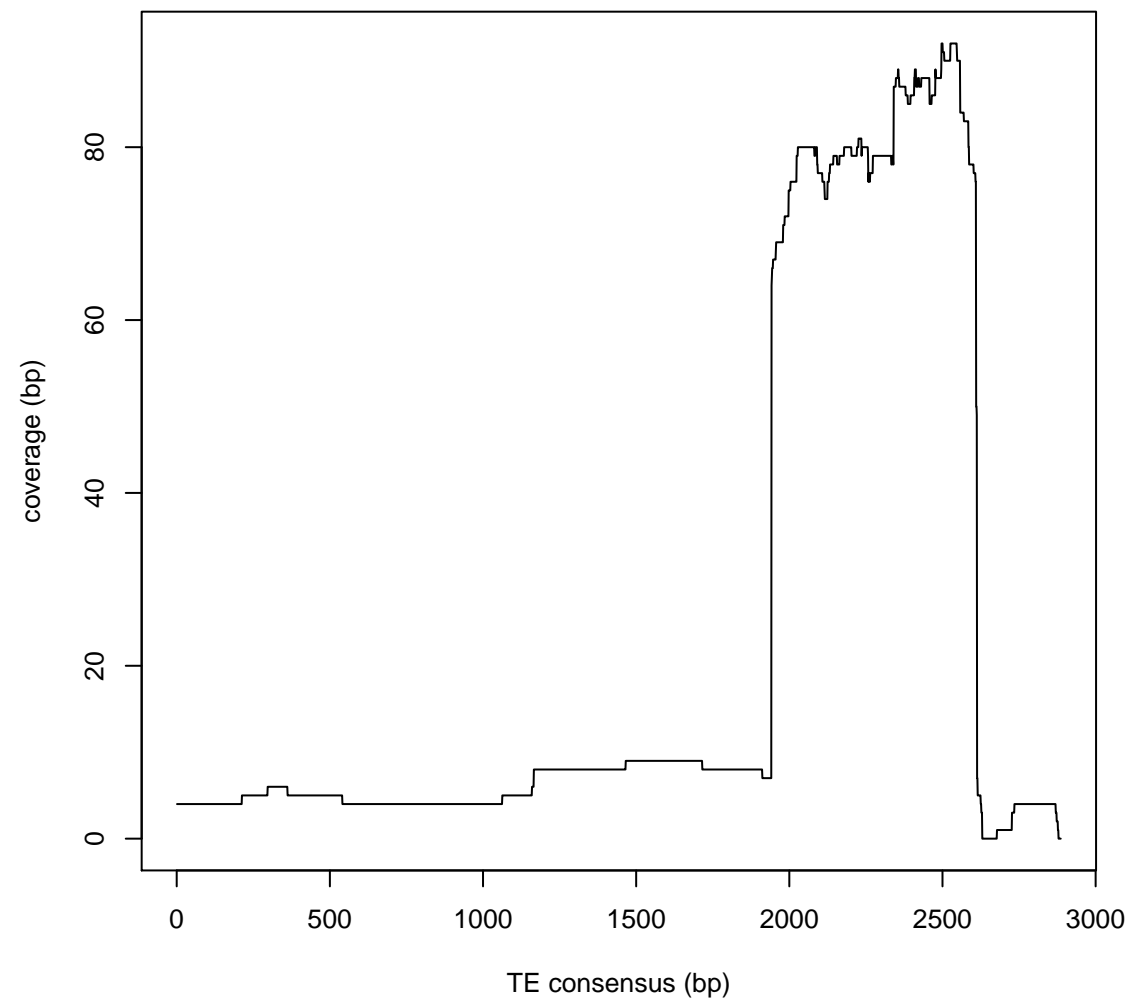


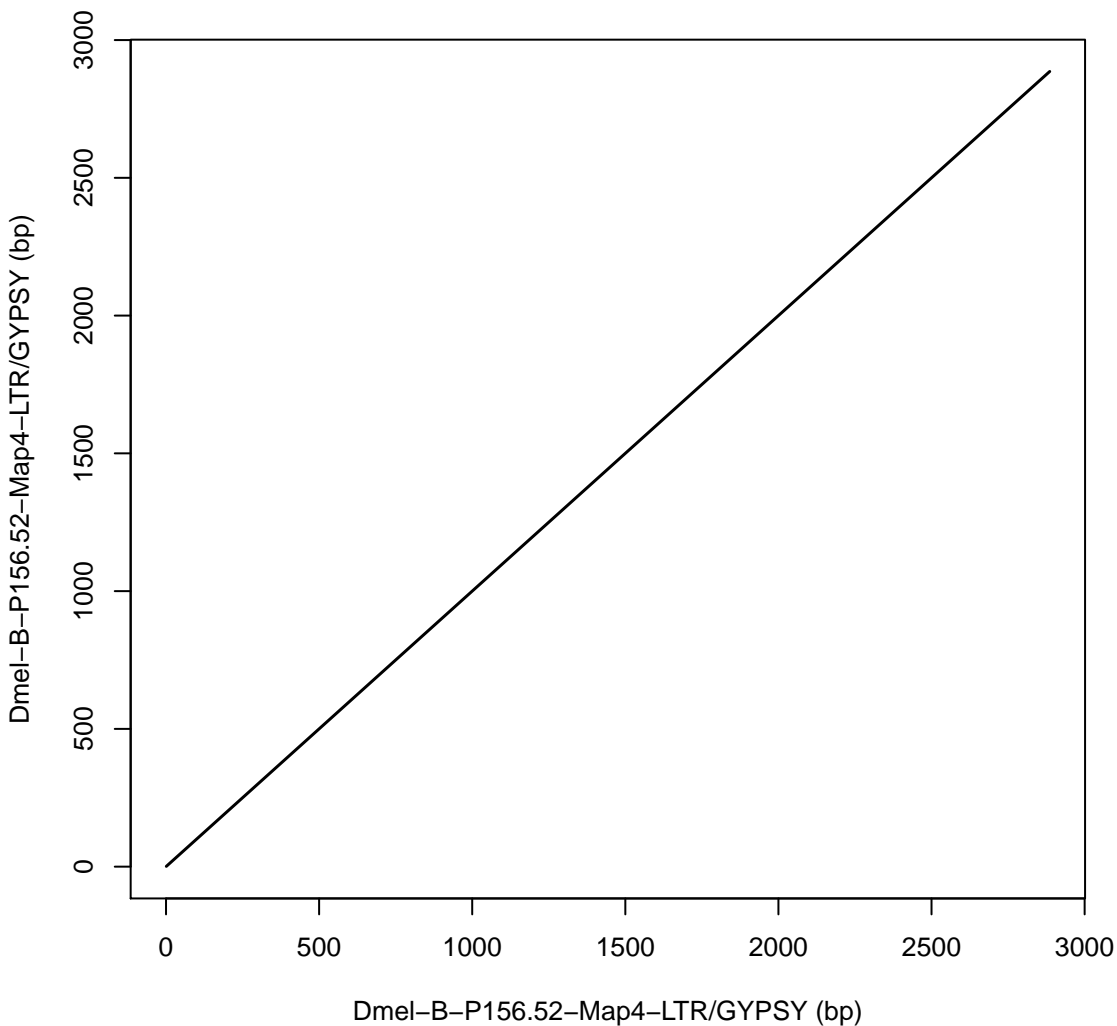
TE: Dmel-B-P156.52-Map4-LTR/GYPSY
consensus size: 2886bp; fragments: 131; full length: 1 (>=2597.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

