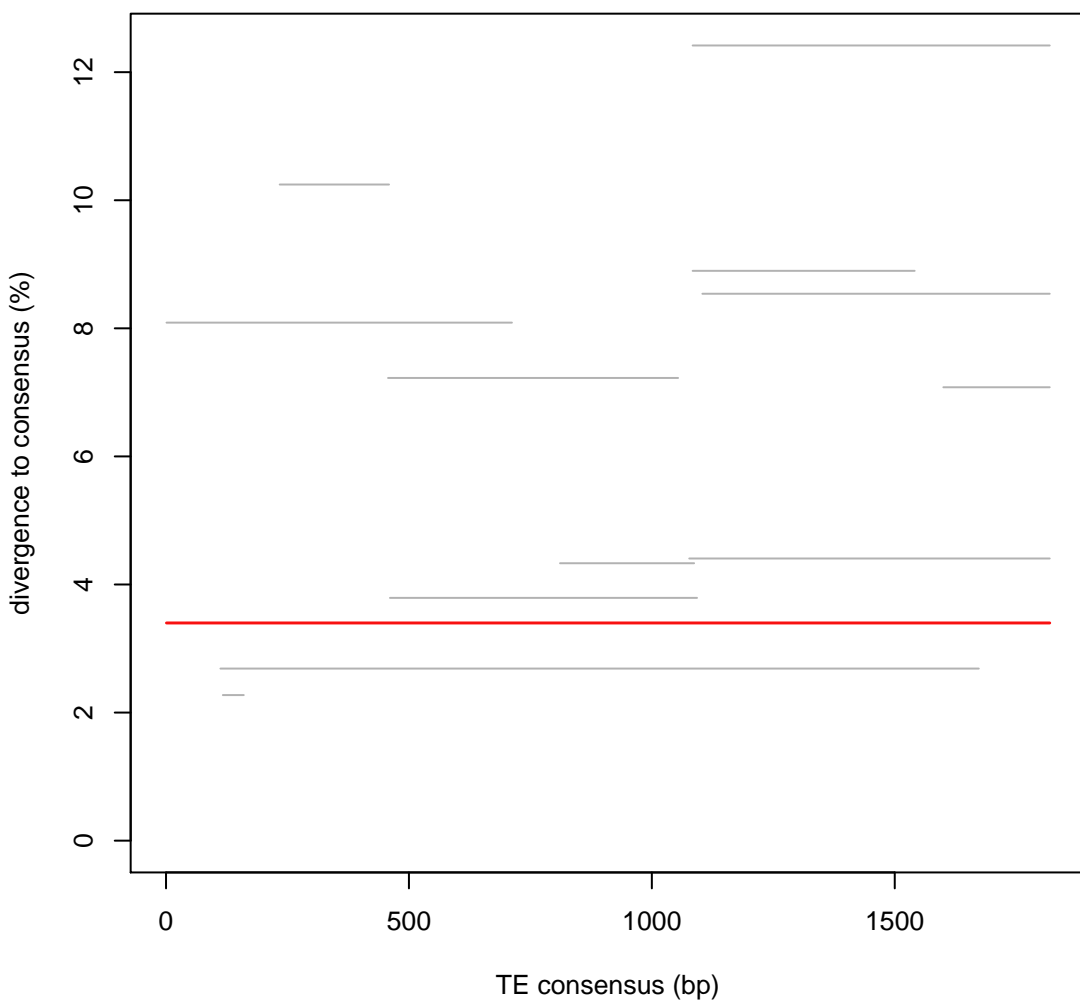
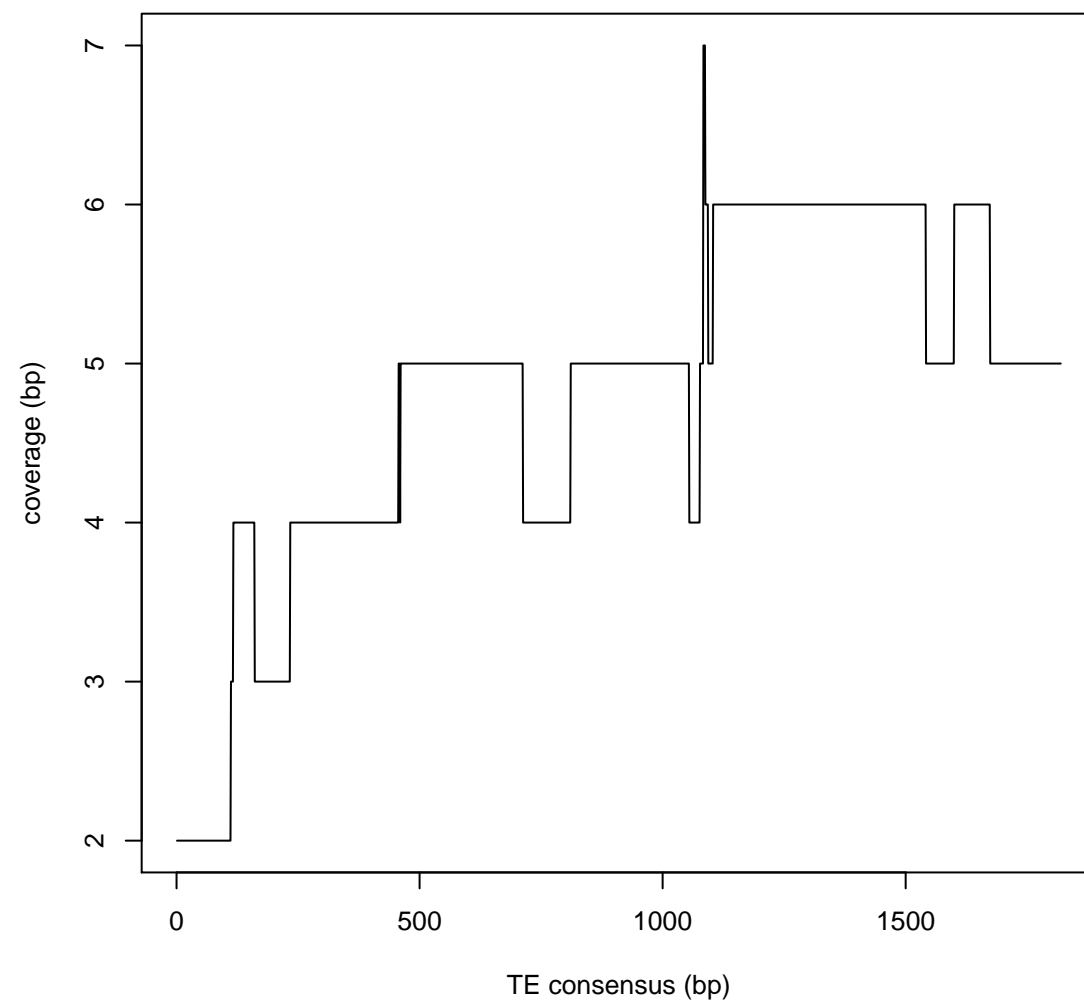


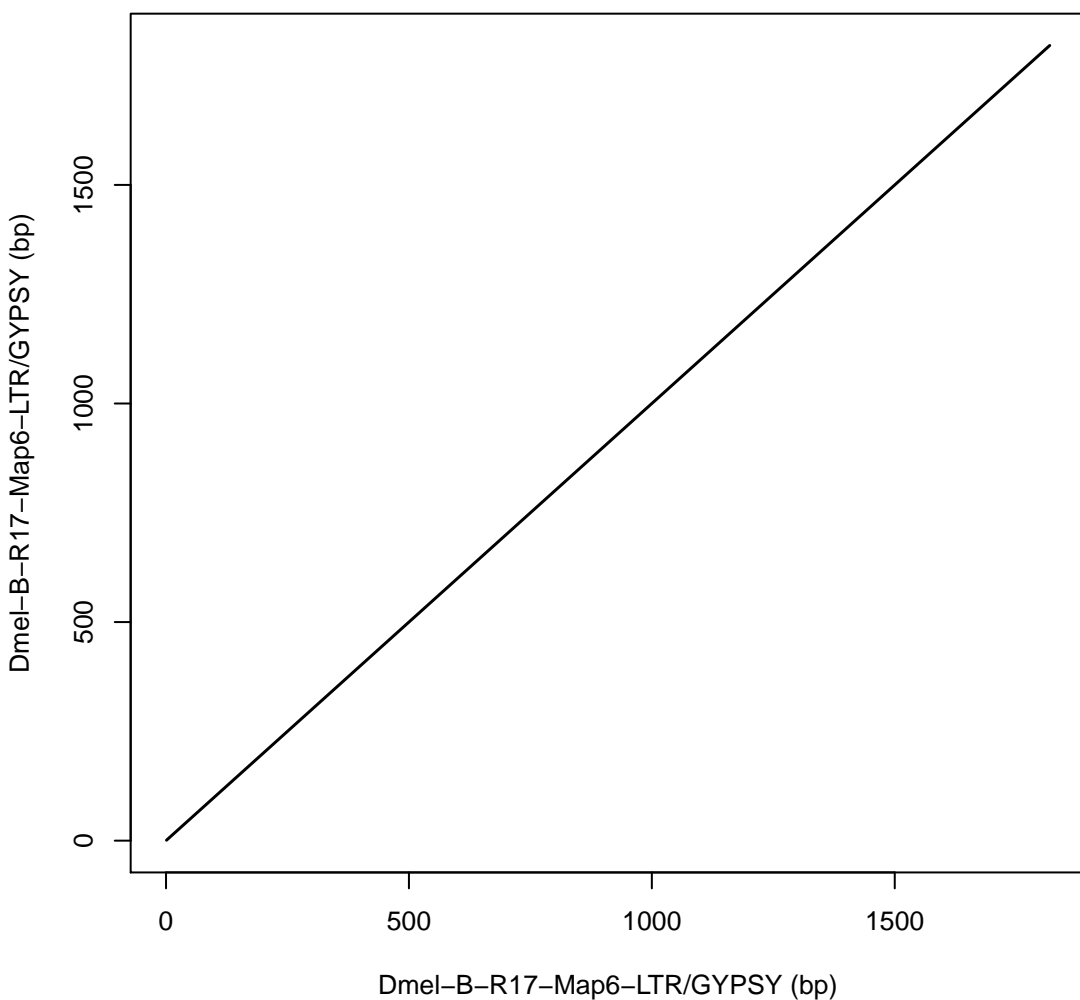
TE: Dmel-B-R17-Map6-LTR/GYPSY
consensus size: 1819bp; fragments: 13; full length: 1 (>=1637.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

