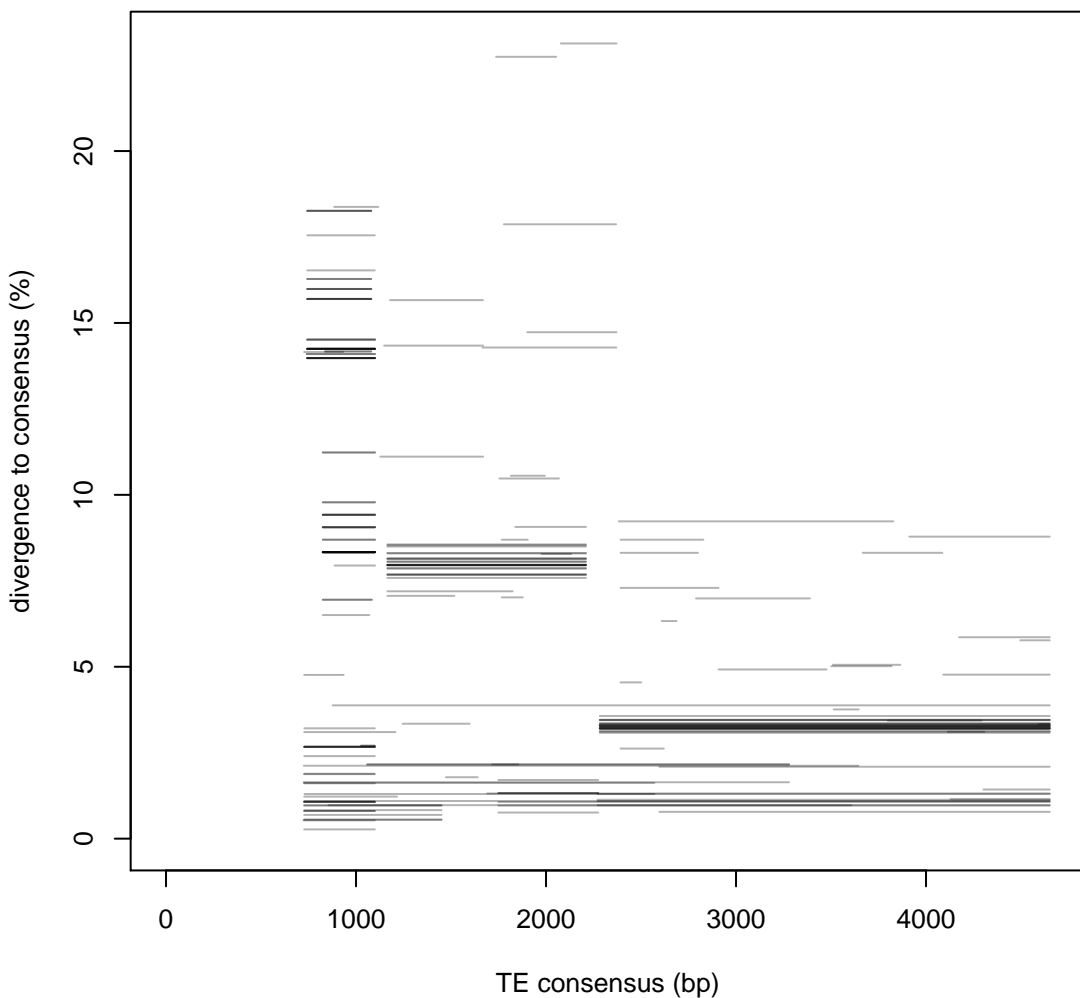
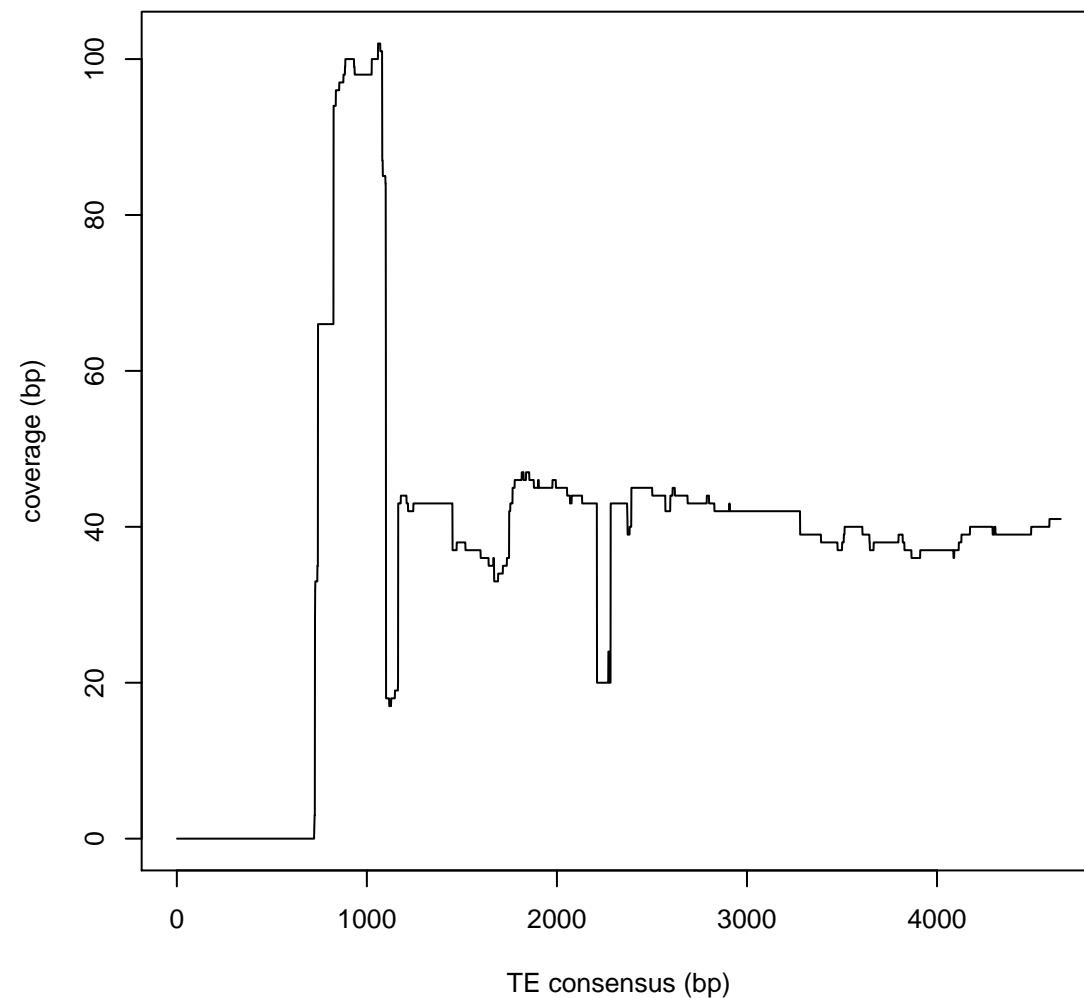


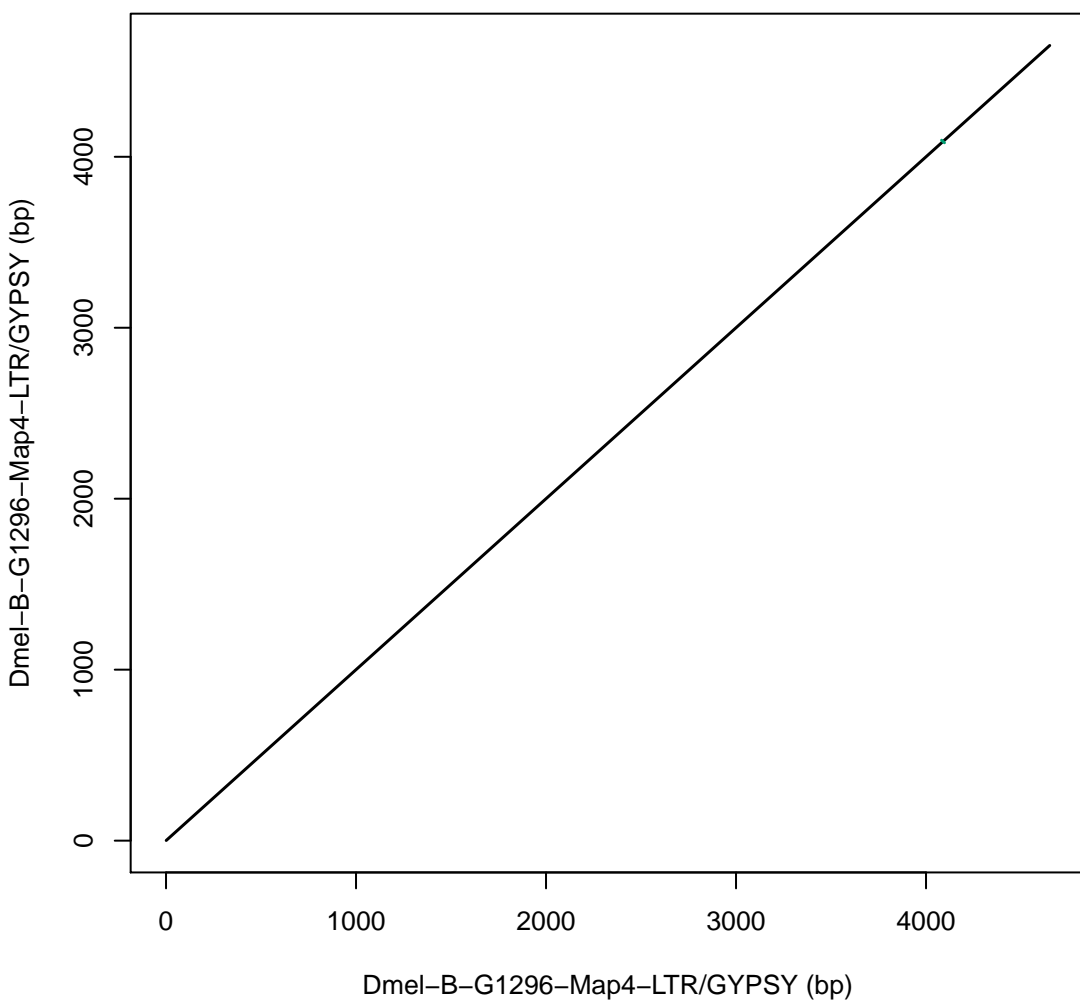
TE: Dmel-B-G1296-Map4-LTR/GYPSY
consensus size: 4651bp; fragments: 203; full length: 0 (>=4185.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

