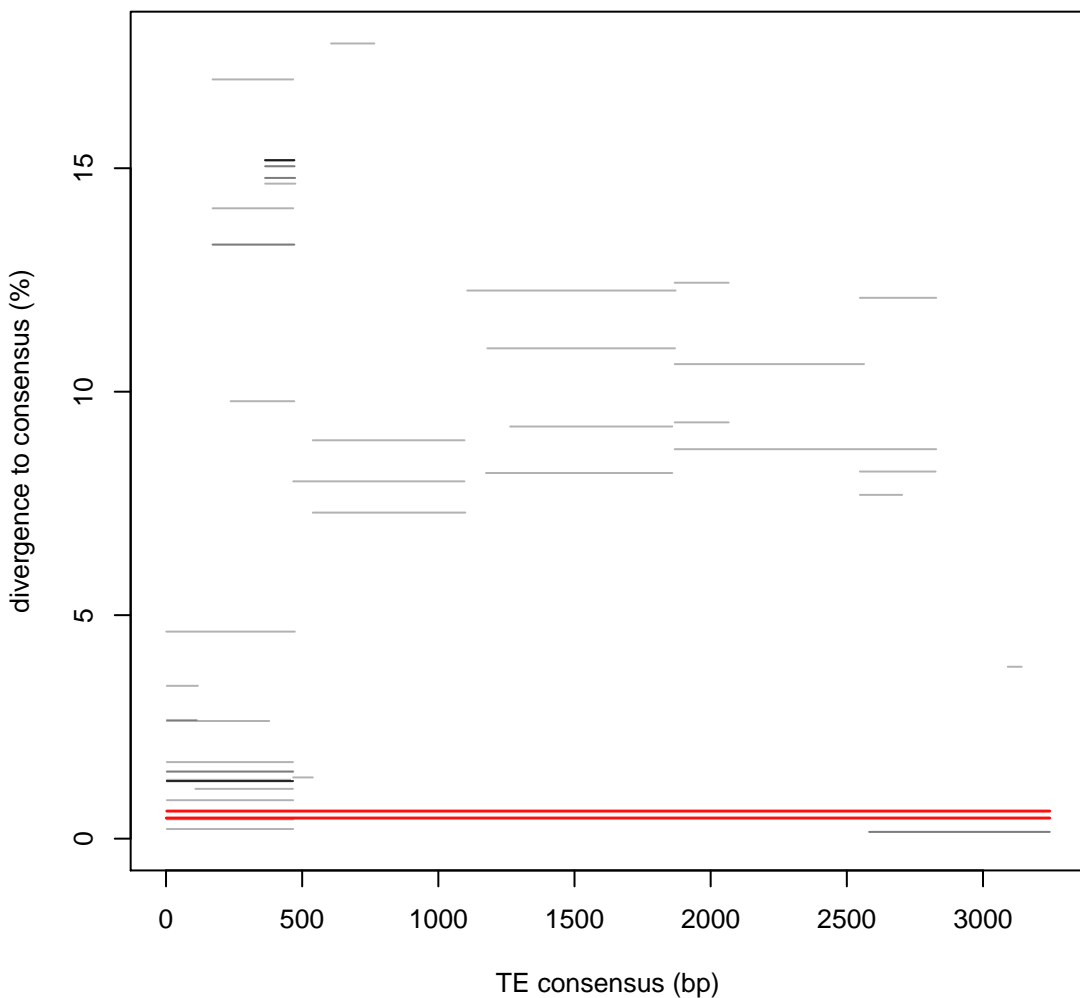
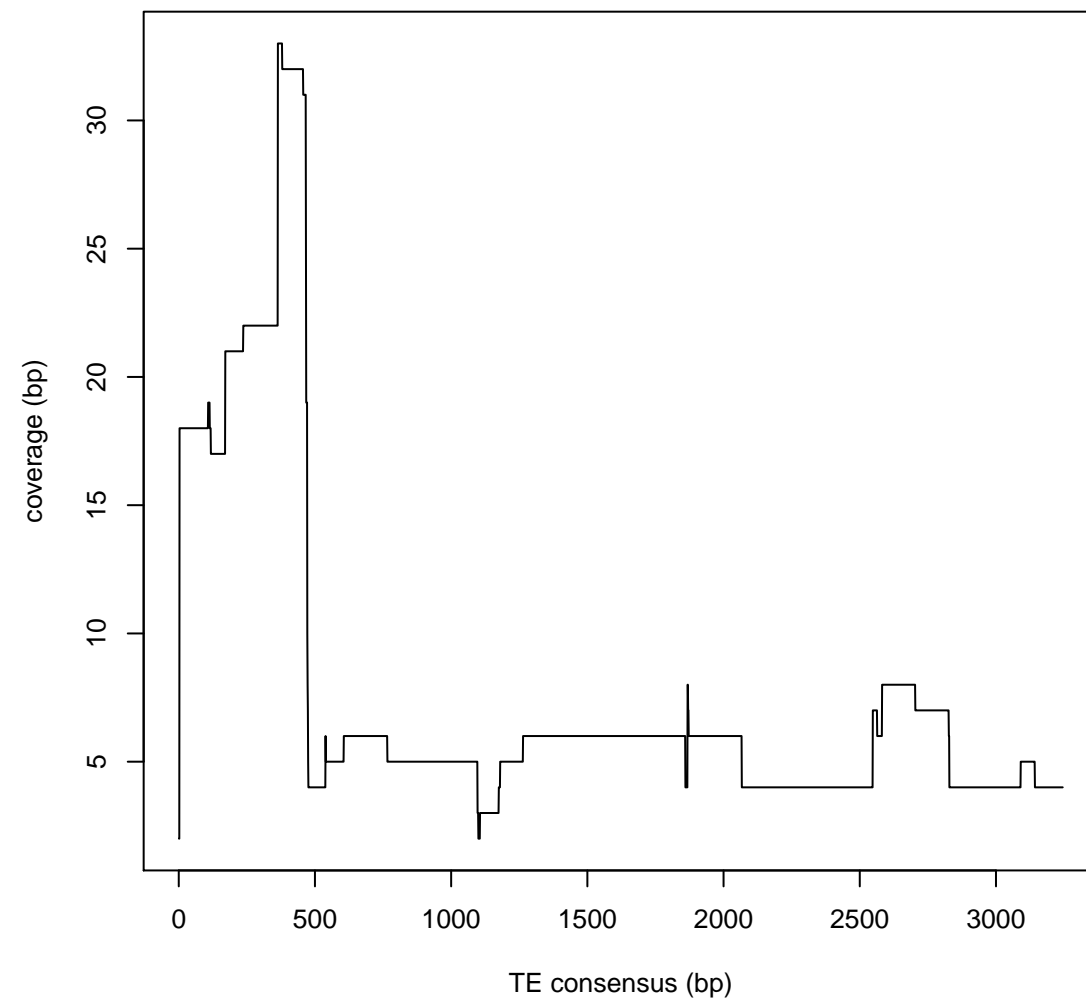


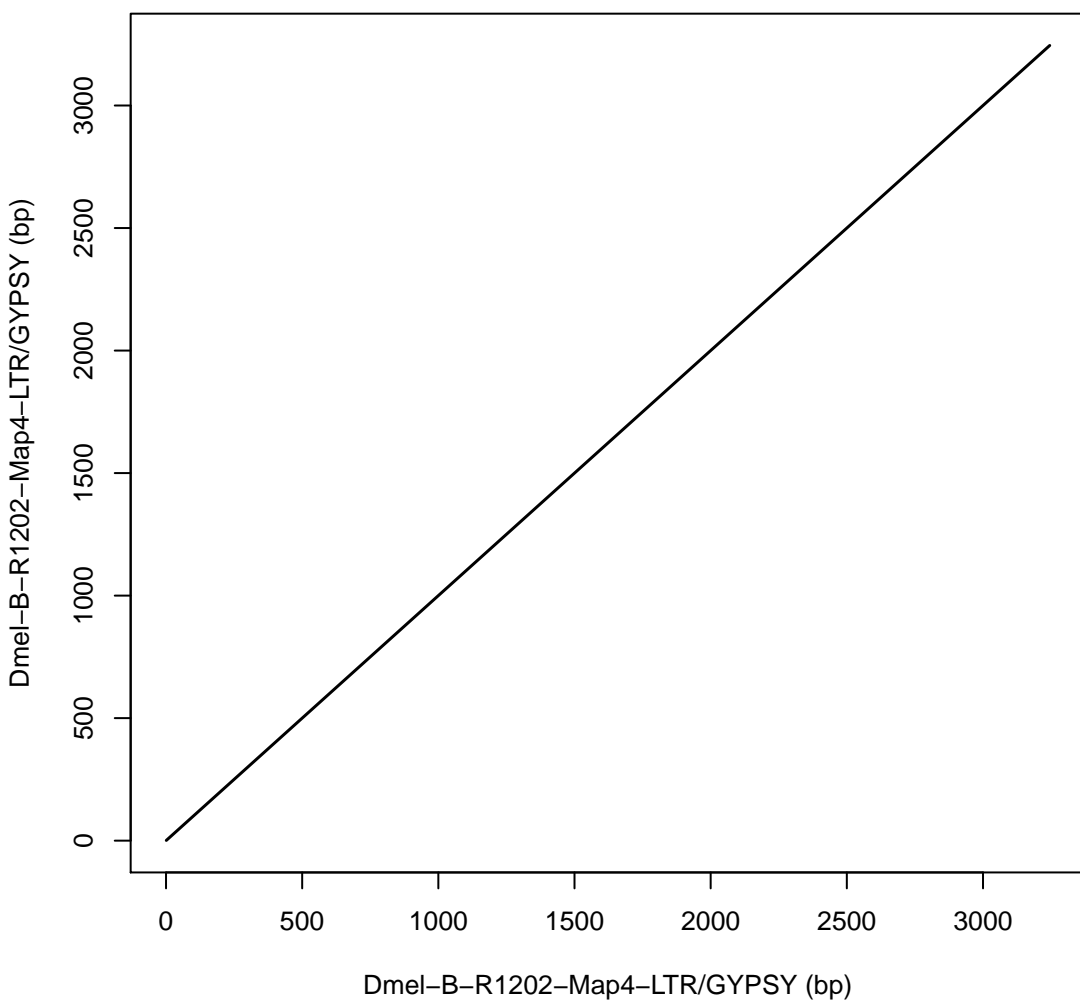
TE: Dmel-B-R1202-Map4-LTR/GYPSY
consensus size: 3245bp; fragments: 54; full length: 2 (>=2920.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

