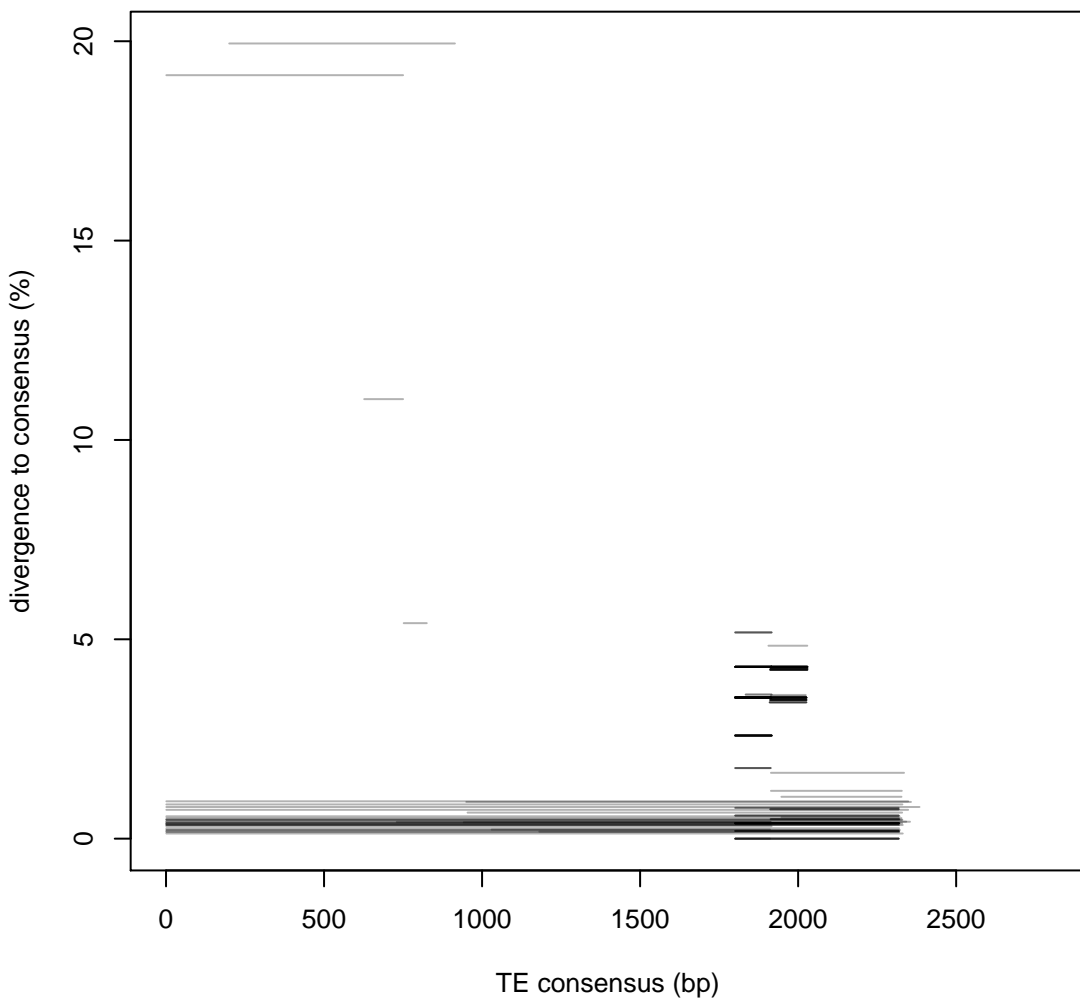
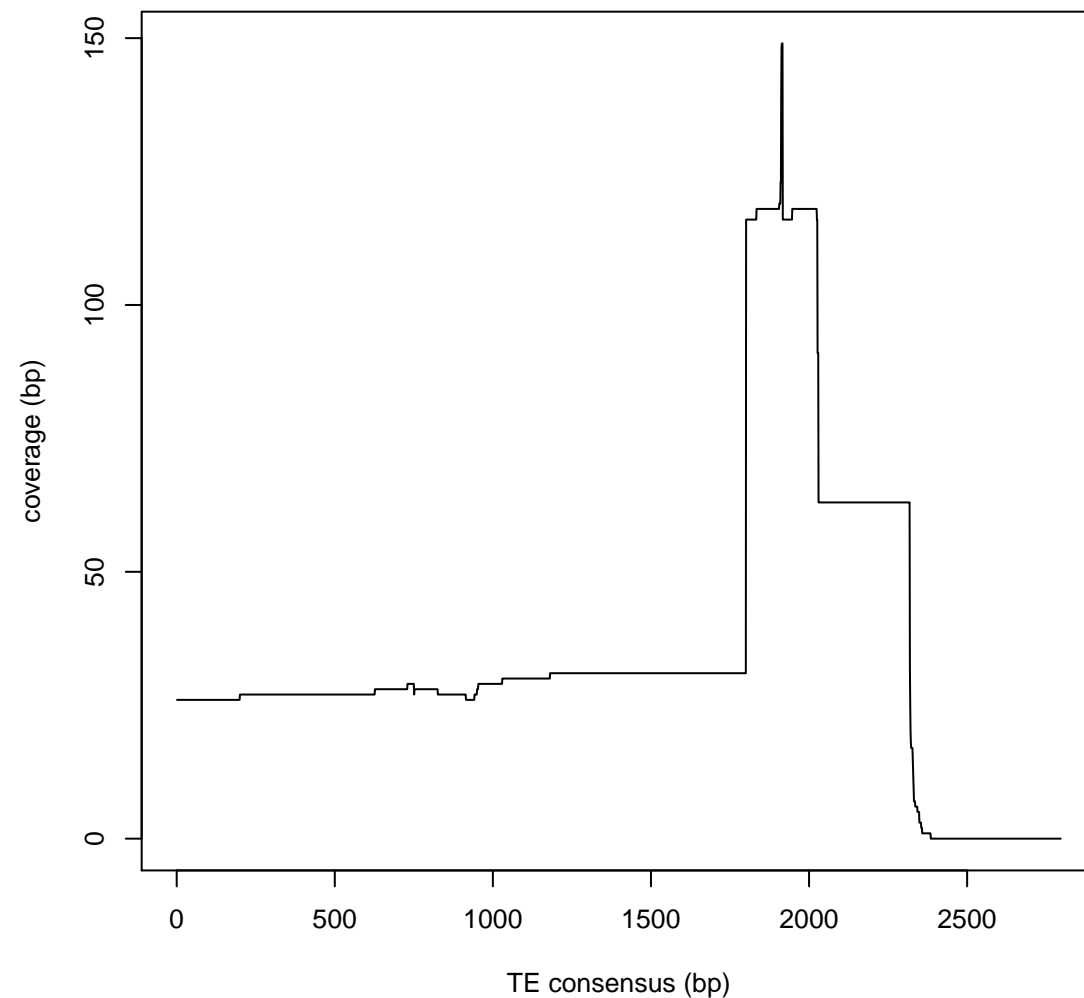


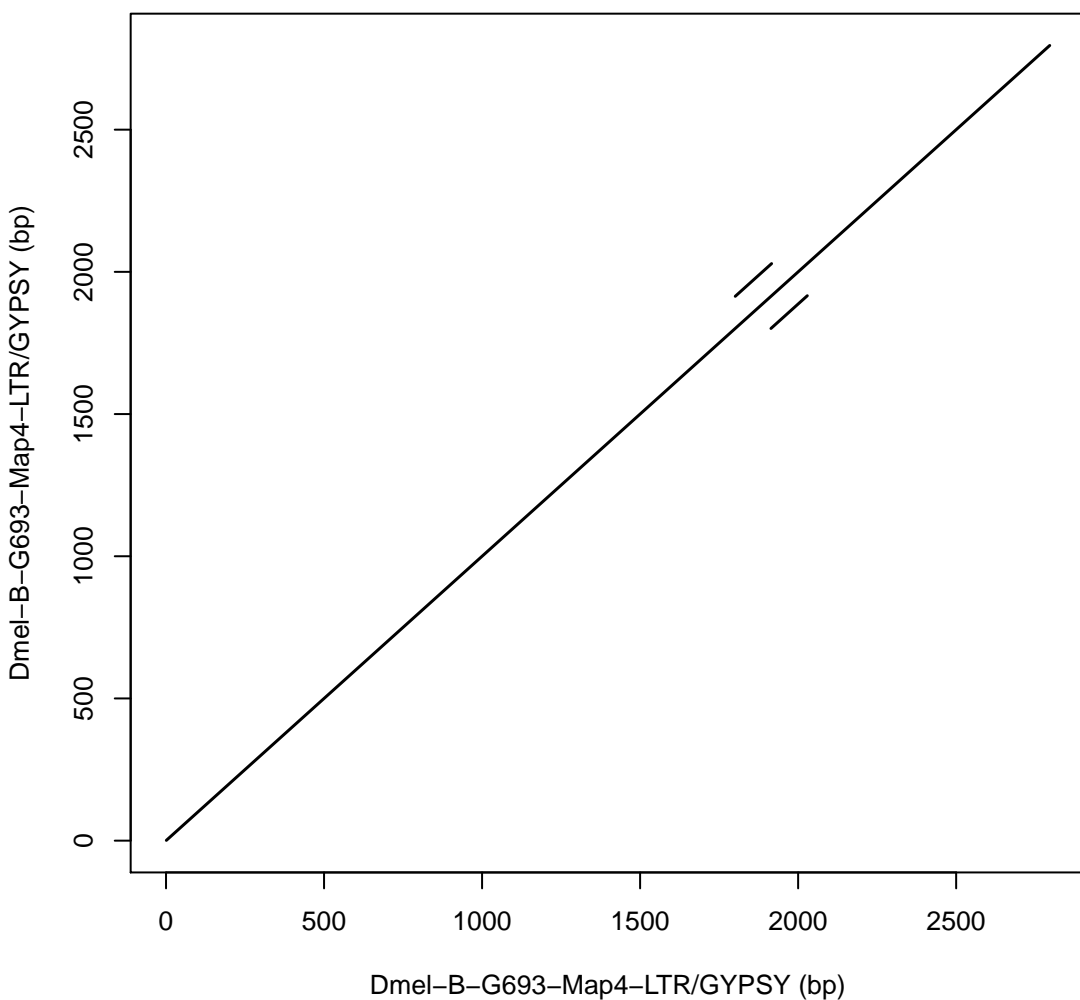
TE: Dmel-B-G693-Map4-LTR/GYPSY
consensus size: 2796bp; fragments: 187; full length: 0 (>=2516.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

