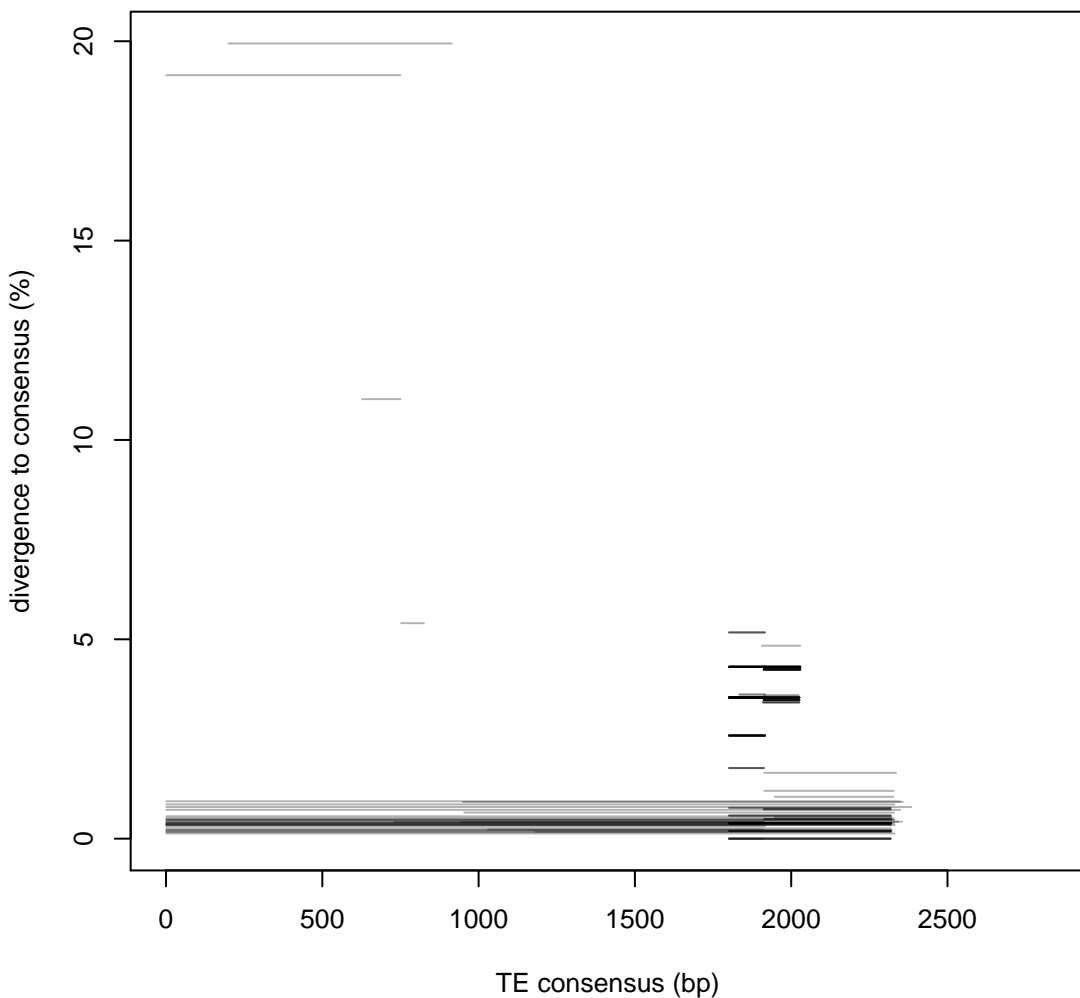
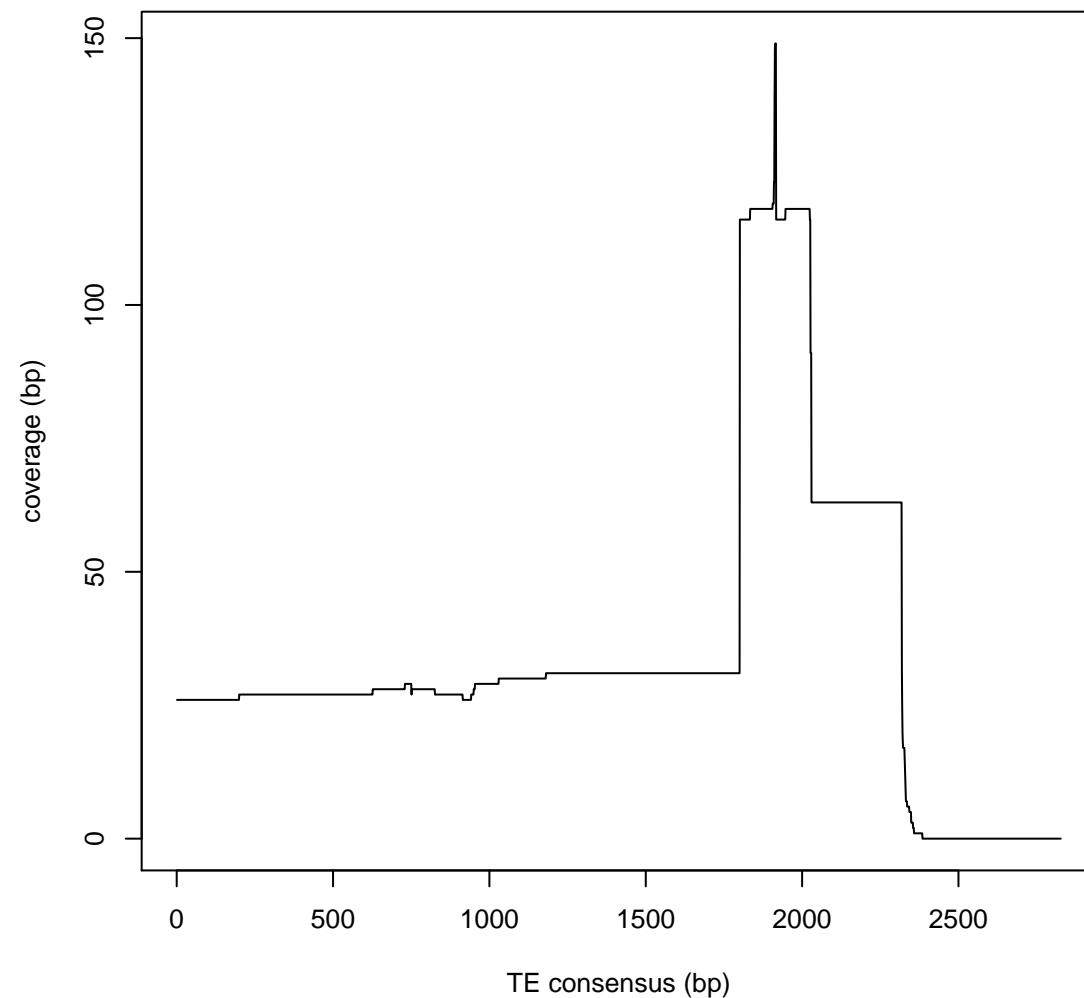


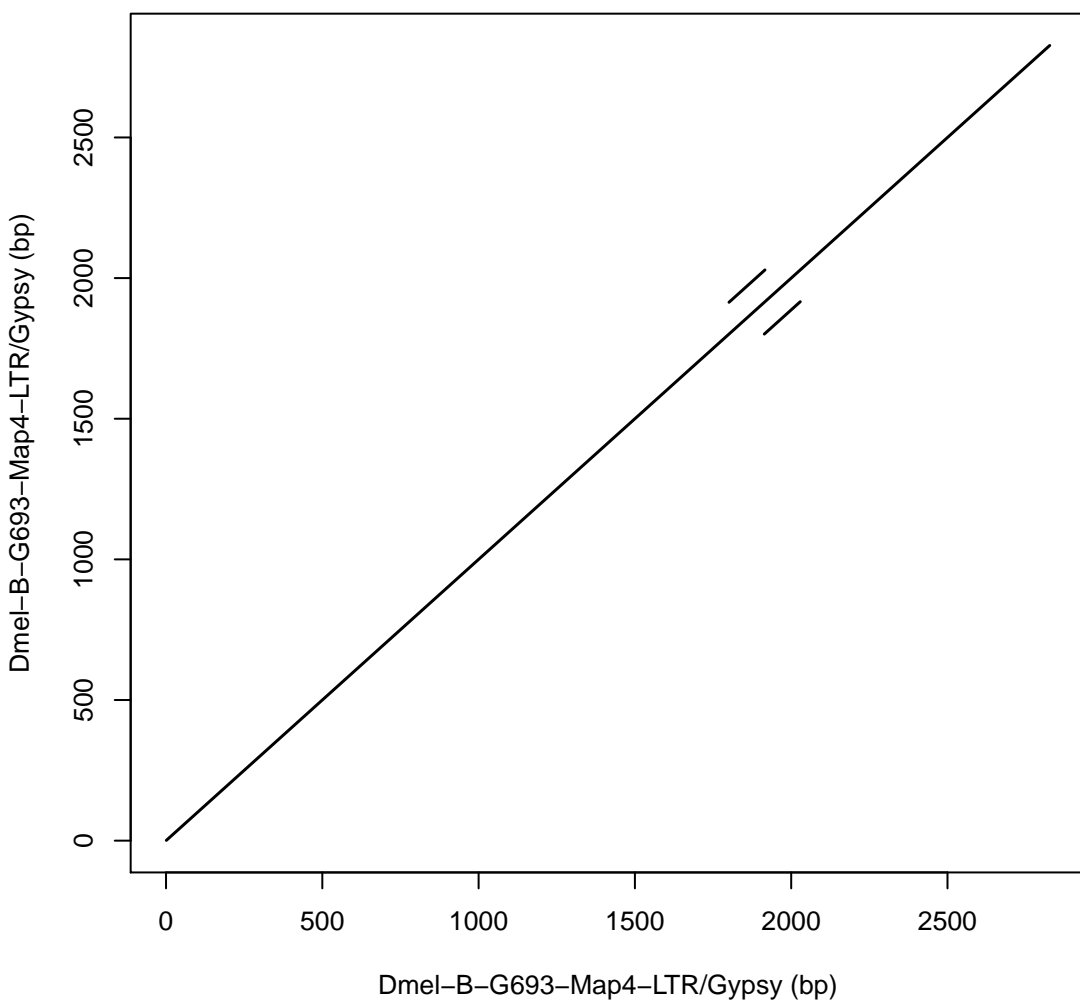
TE: Dmel-B-G693-Map4-LTR/Gypsy
consensus size: 2827bp; fragments: 187; full length: 0 (>=2544.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

