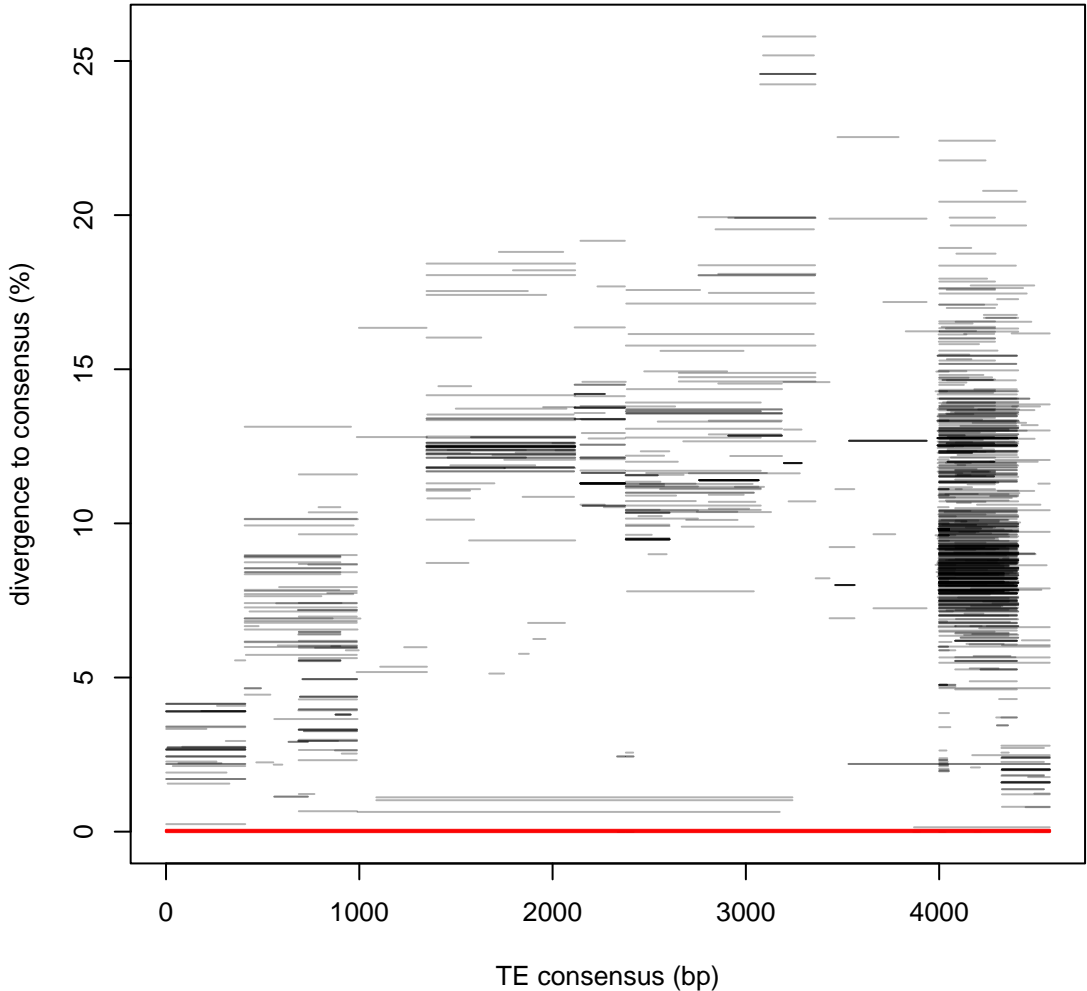
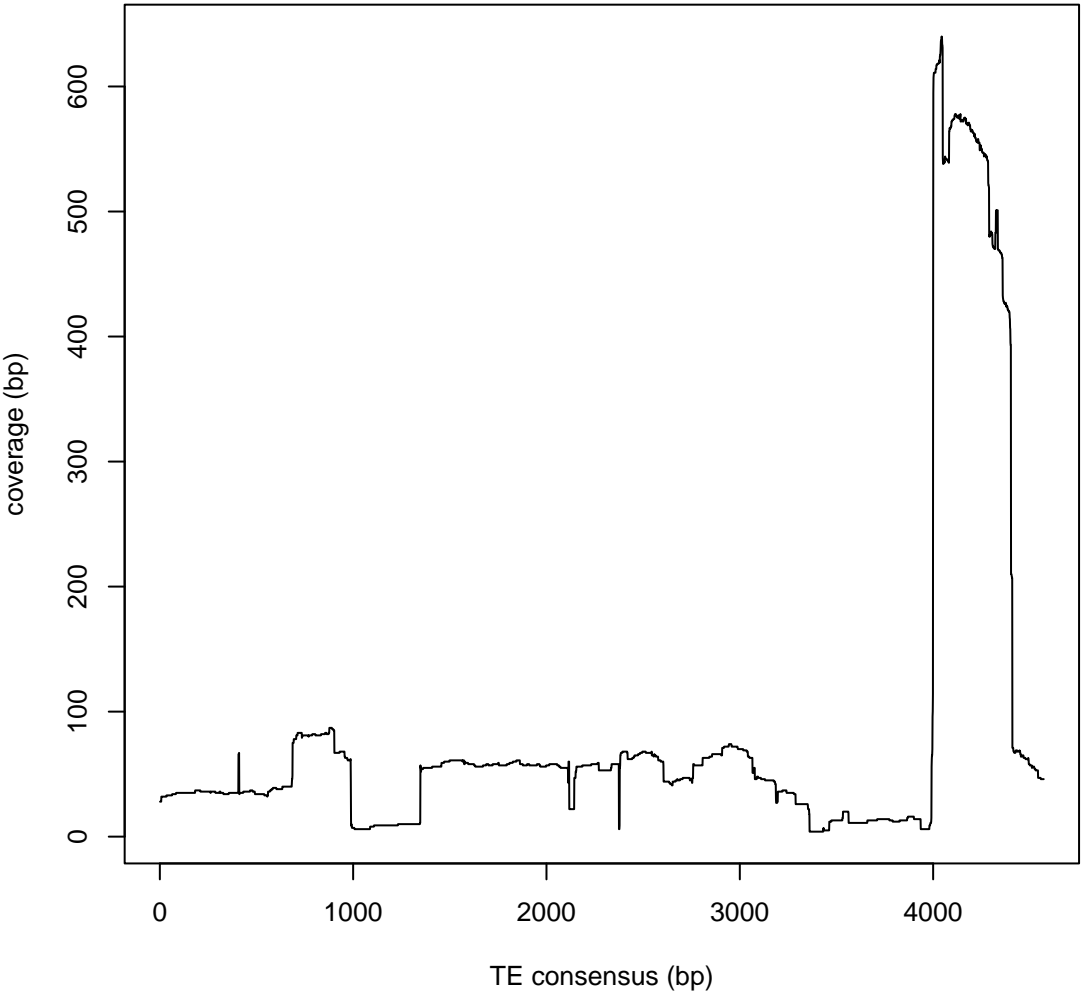


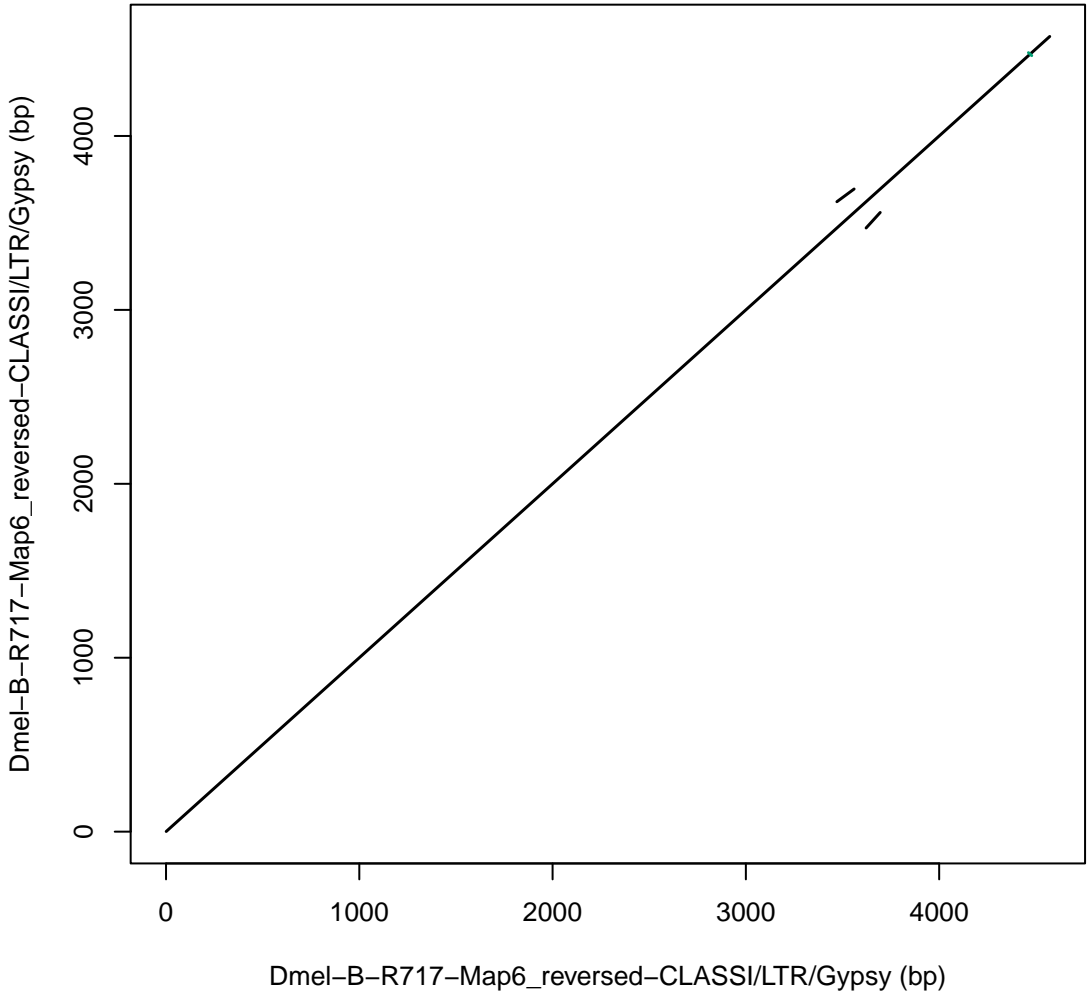
TE: Dmel-B-R717-Map6_reversed-CLASSI/LTR/Gypsy
consensus size: 4572bp; fragments: 1231; full length: 2 (>=4114.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

