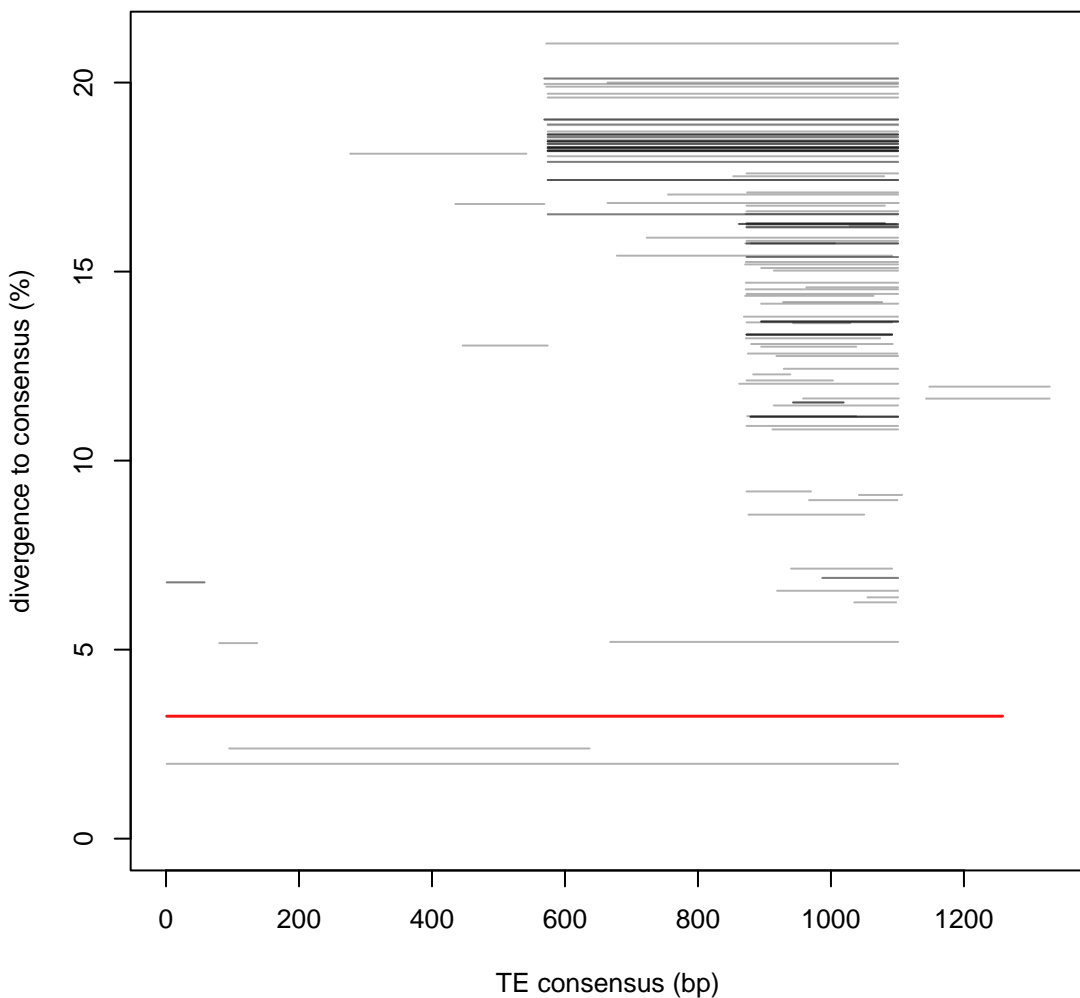
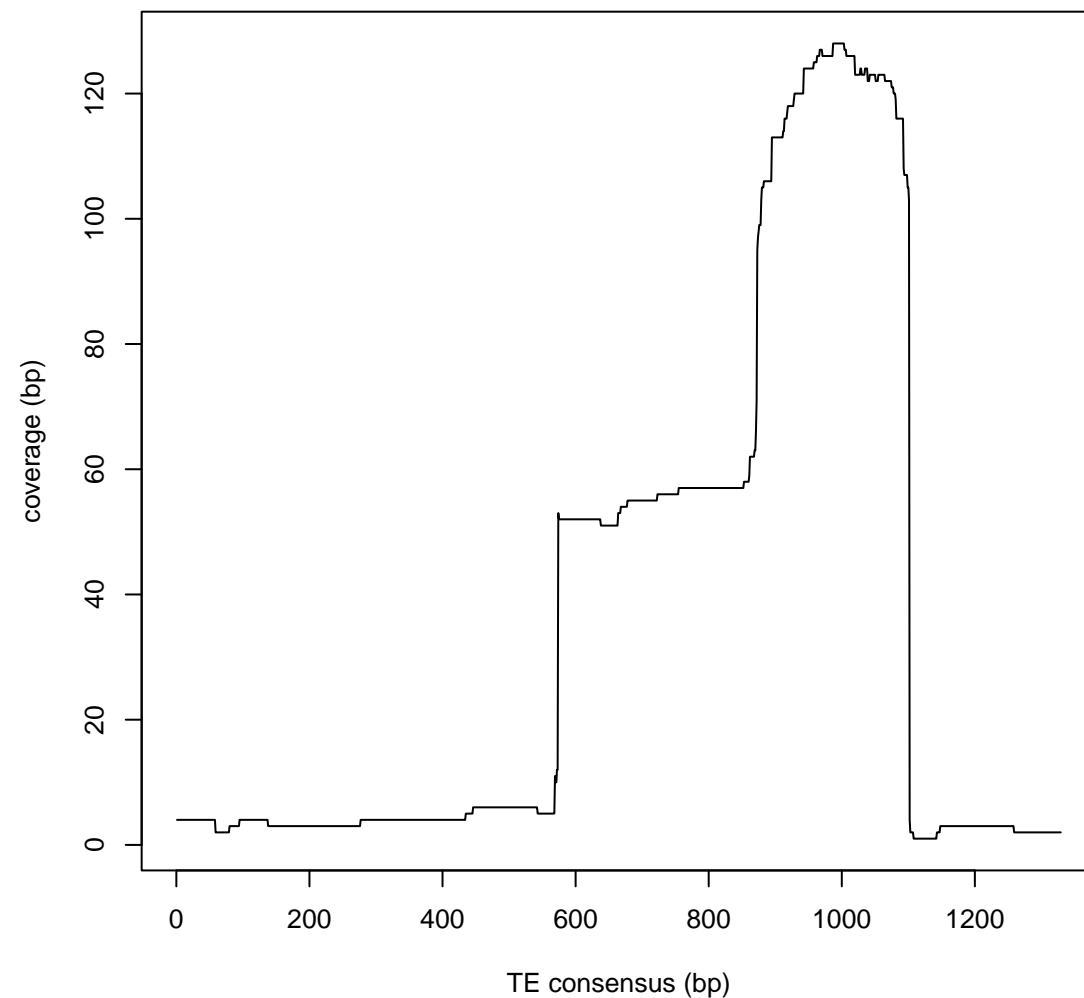


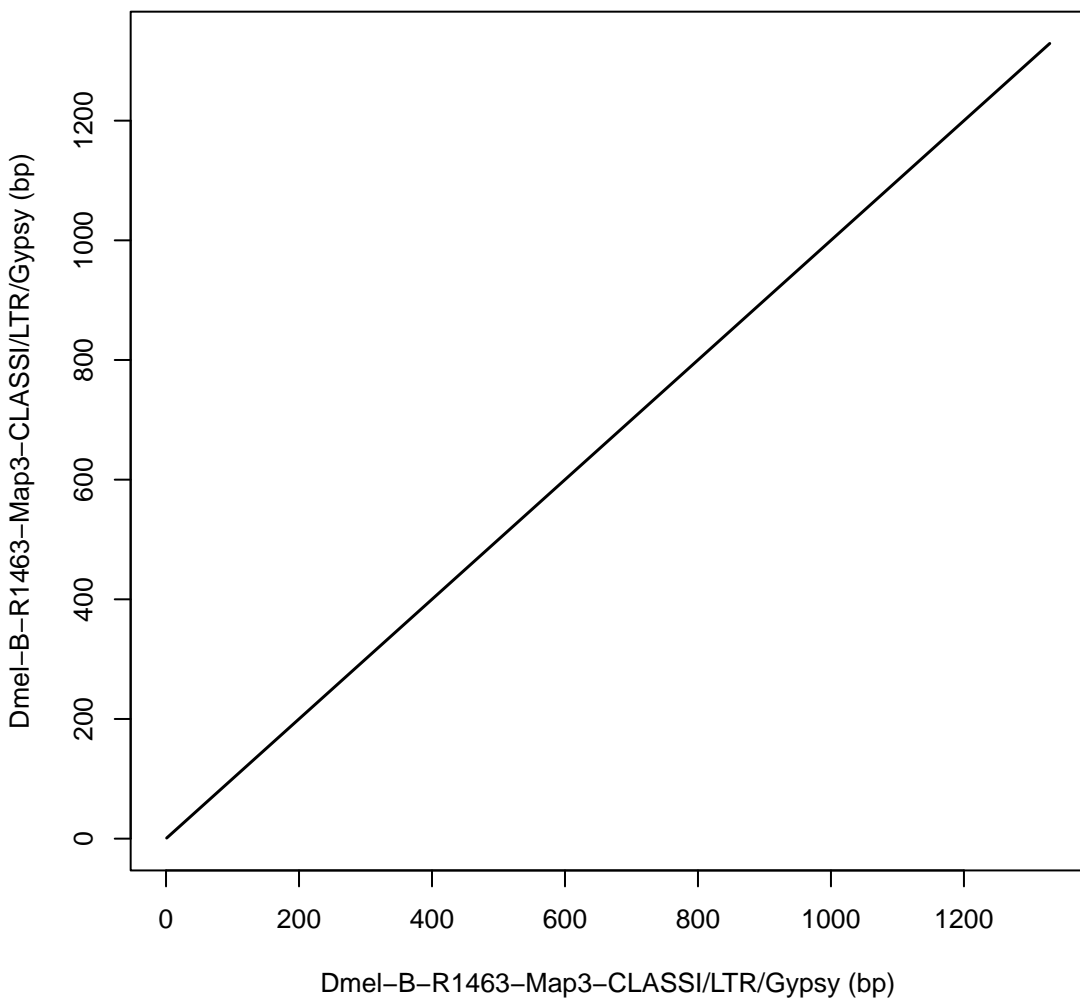
TE: Dmel-B-R1463-Map3-CLASSI/LTR/Gypsy
consensus size: 1329bp; fragments: 143; full length: 1 (>=1196.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

