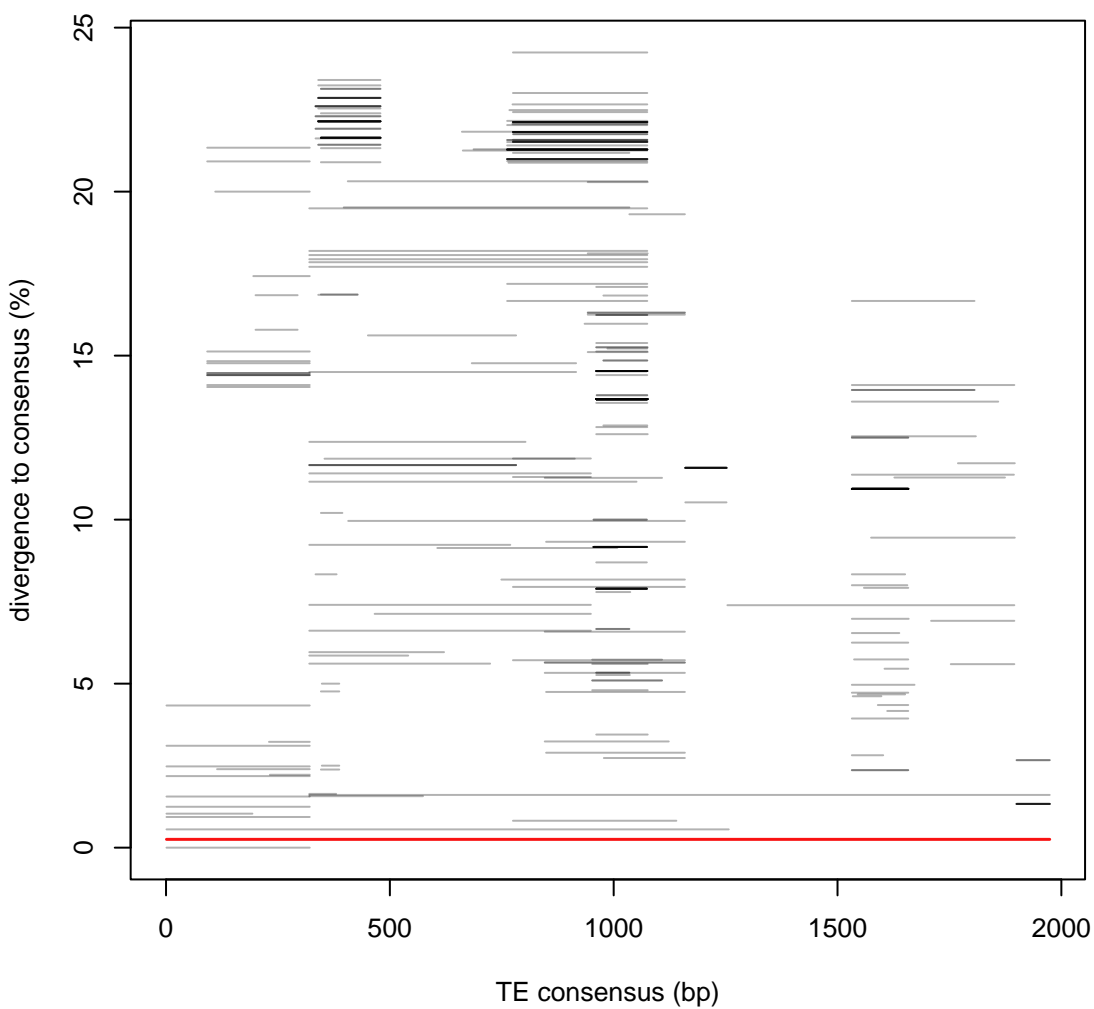
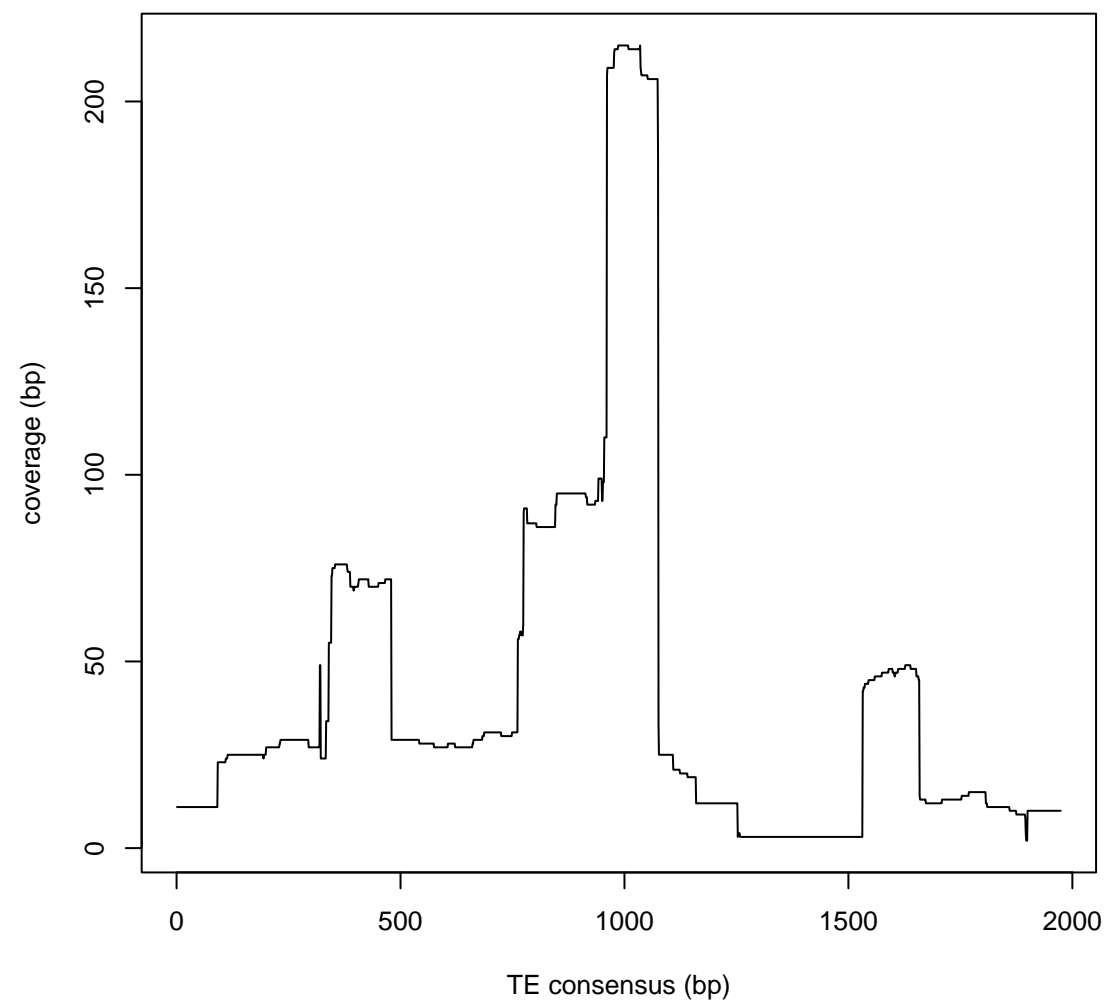


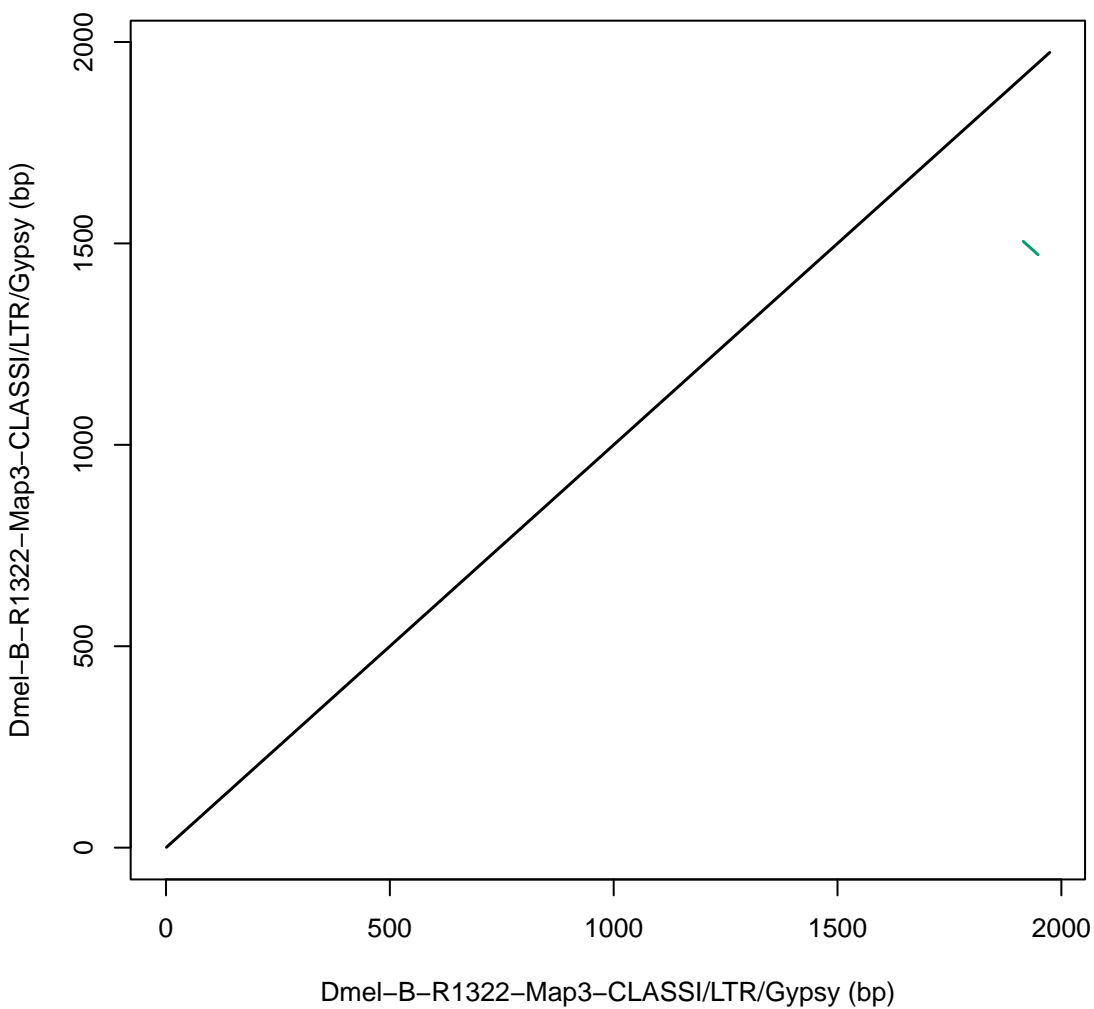
TE: Dmel-B-R1322-Map3-CLASSI/LTR/Gypsy  
consensus size: 1974bp; fragments: 384; full length: 1 (>=1776.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

