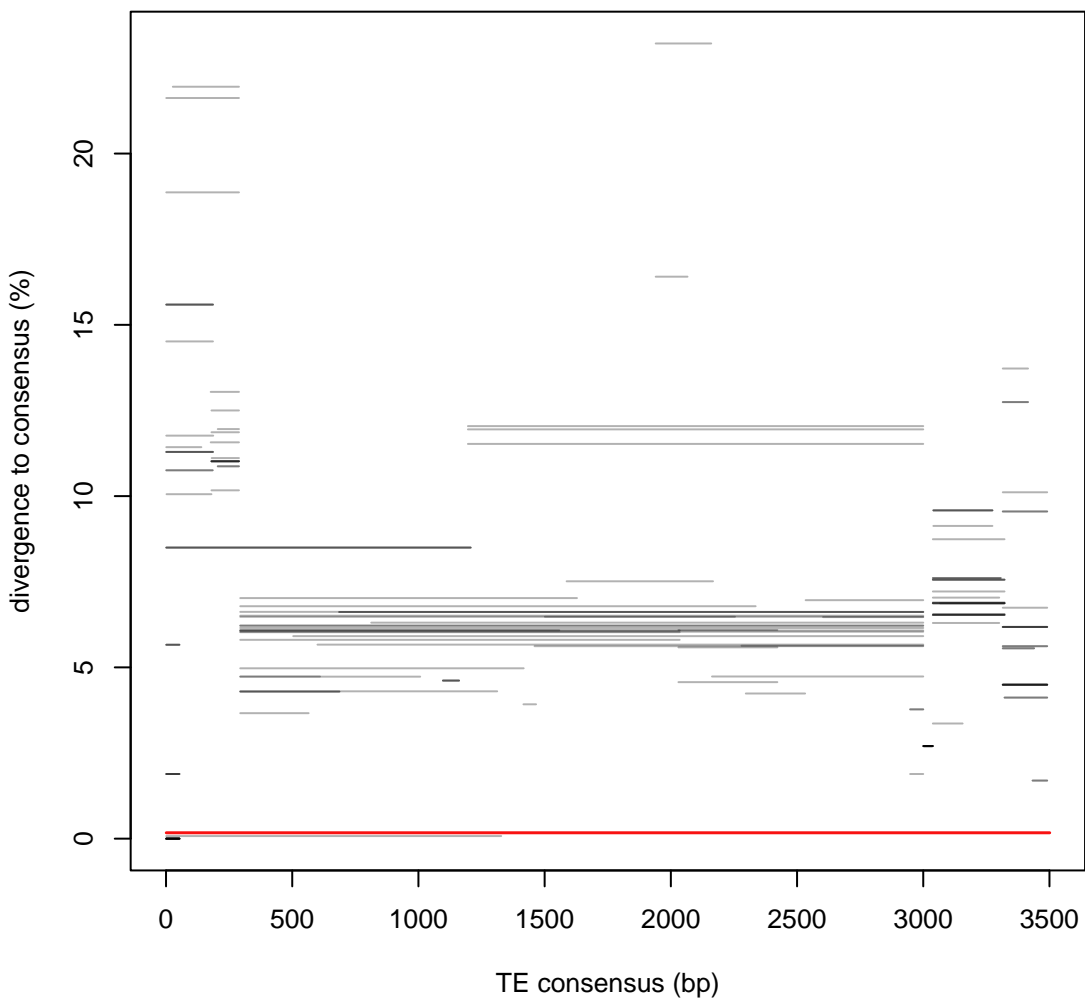
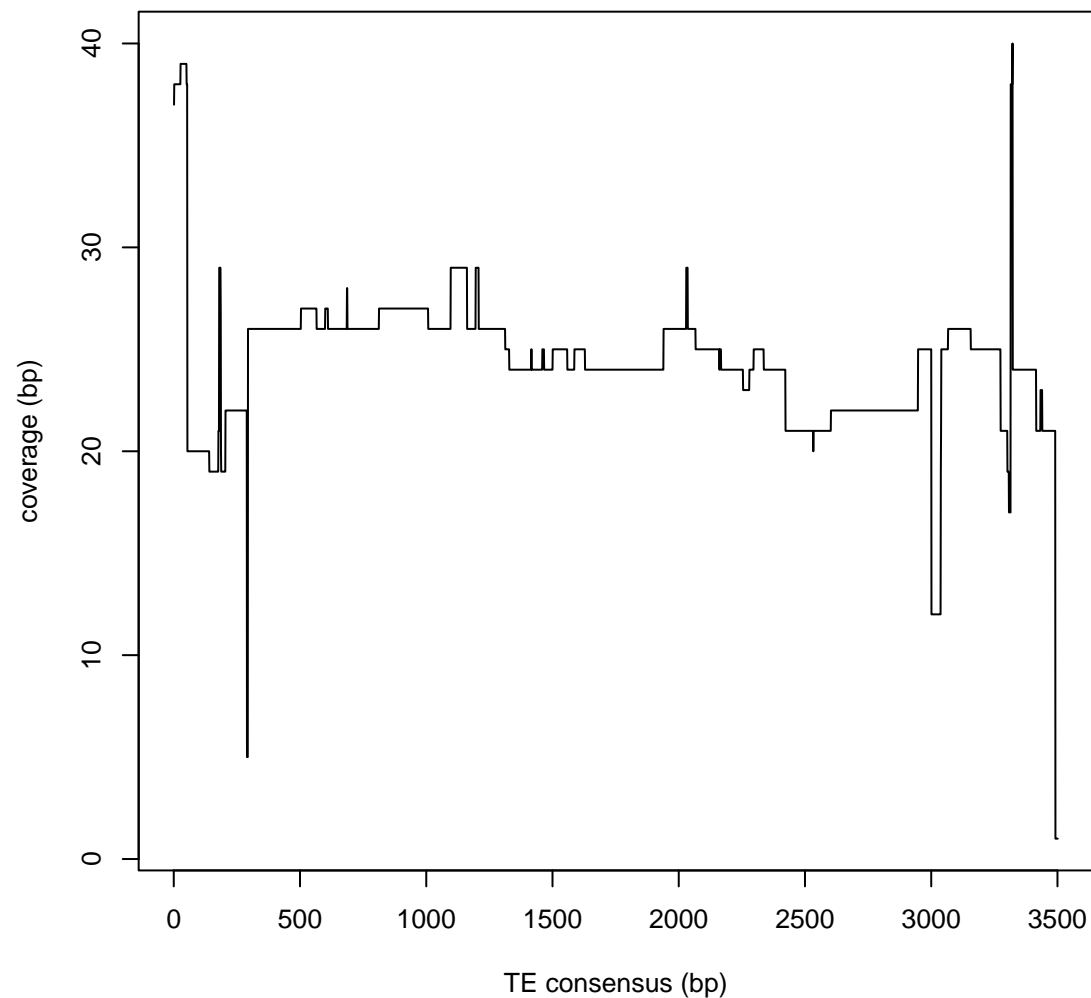


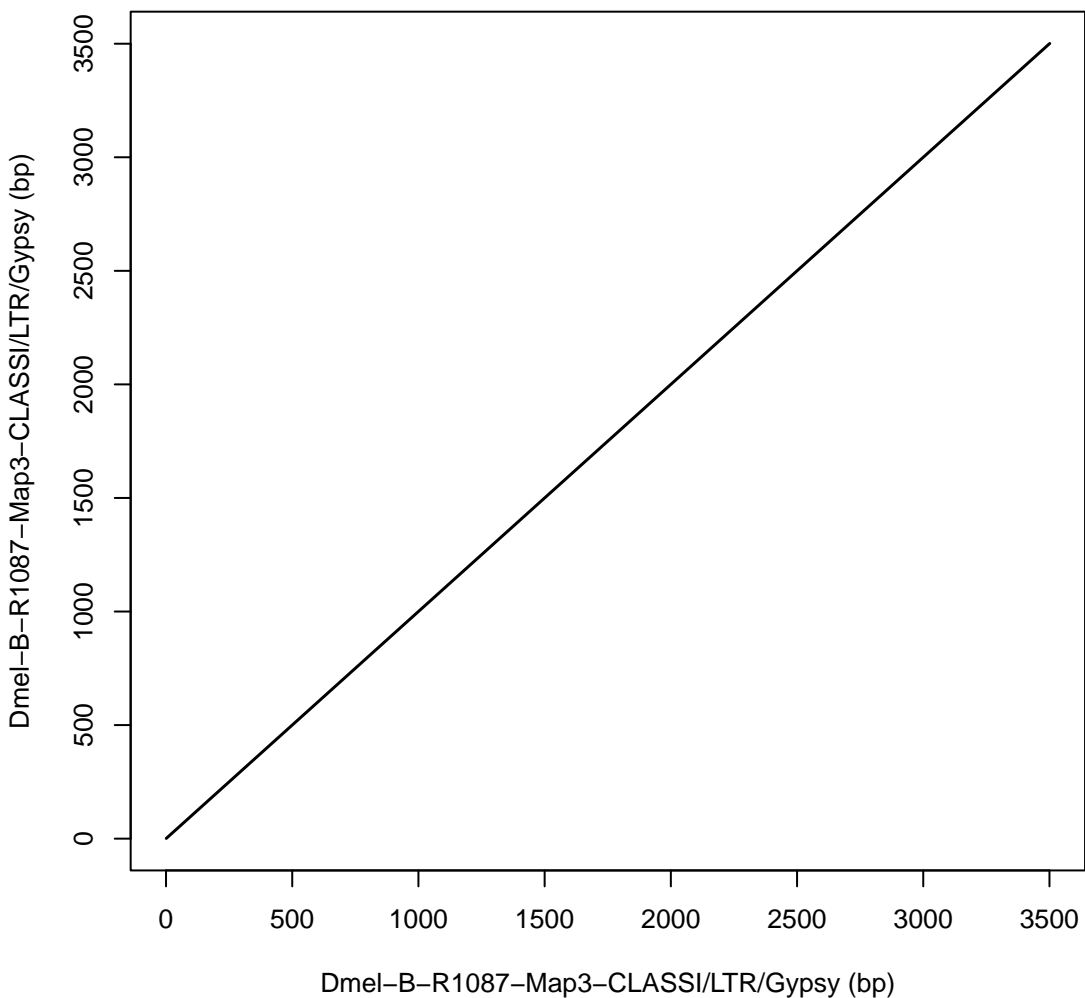
TE: Dmel-B-R1087-Map3-CLASSI/LTR/Gypsy
consensus size: 3501bp; fragments: 163; full length: 1 (>=3150.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

