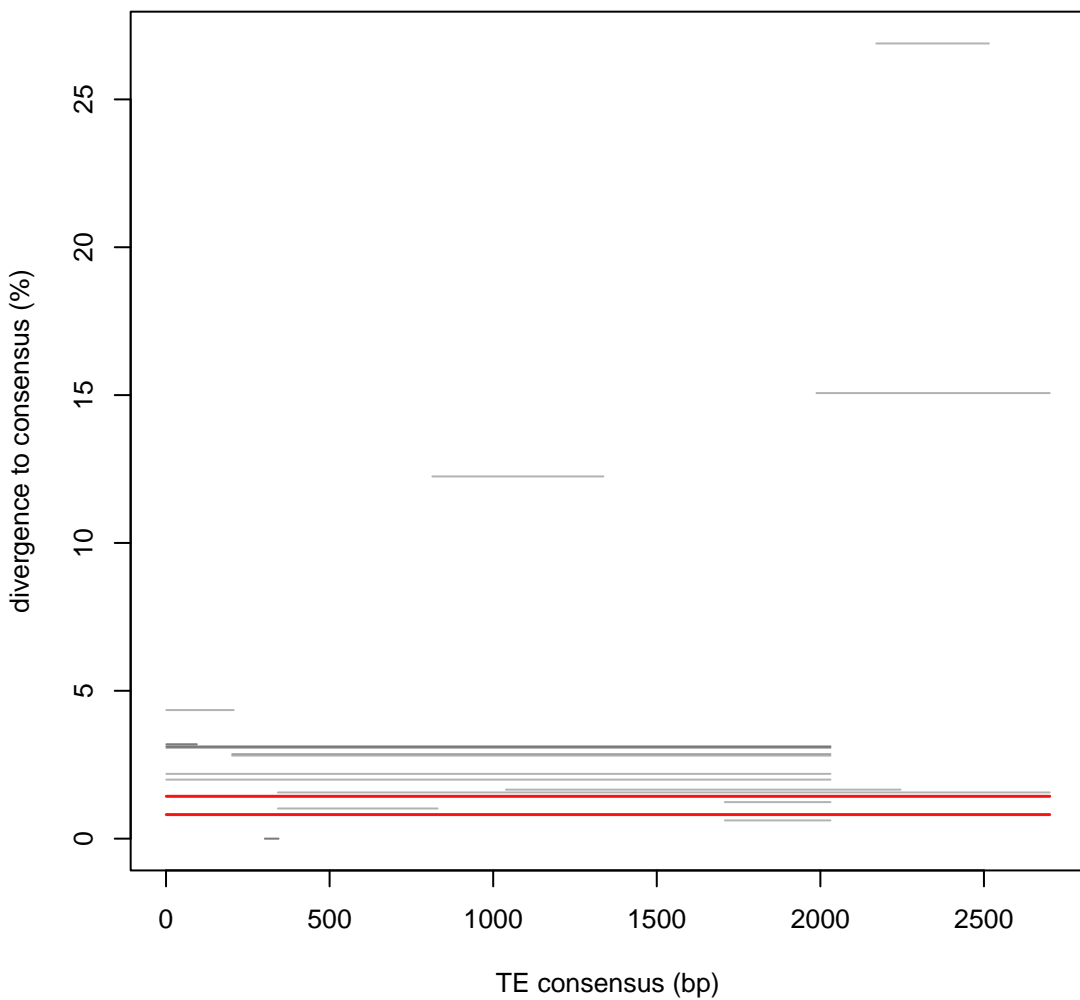
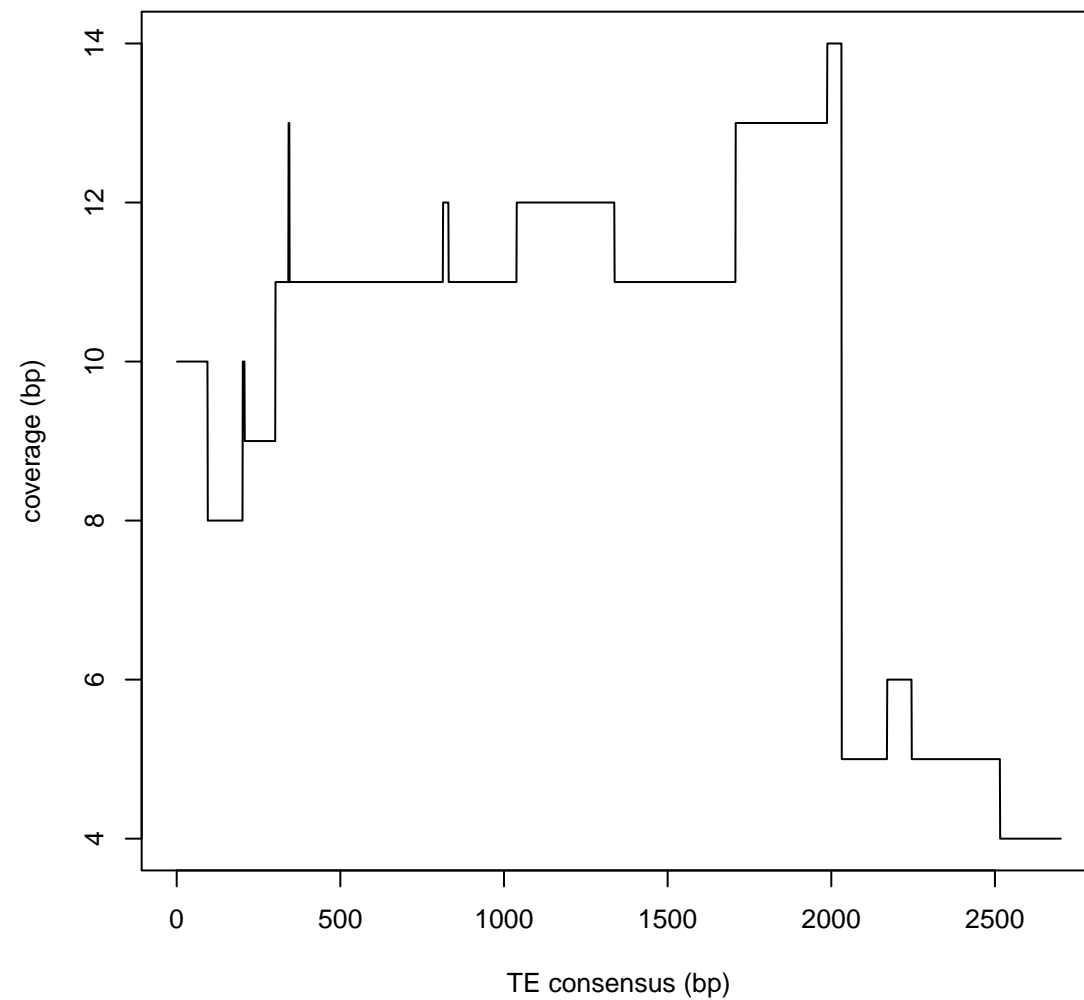


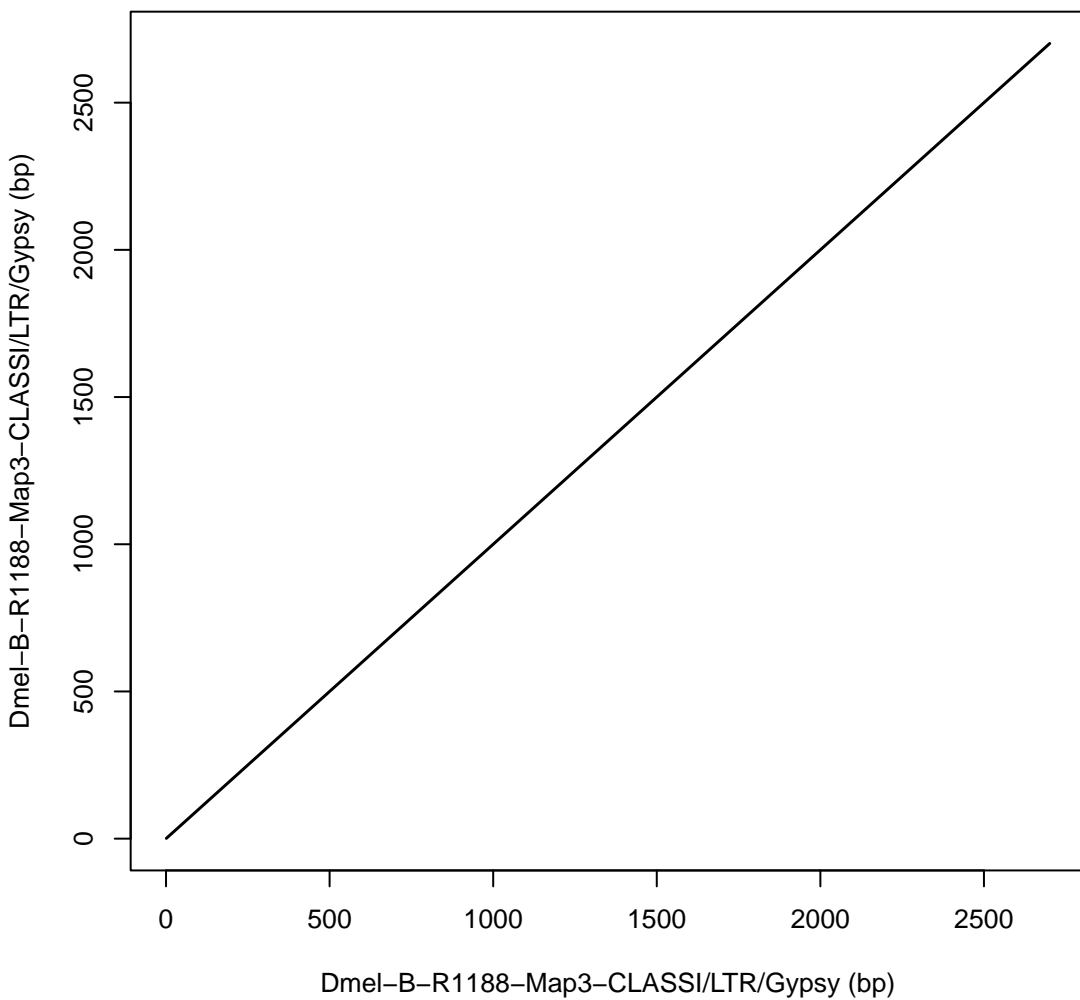
TE: Dmel-B-R1188-Map3-CLASSI/LTR/Gypsy
consensus size: 2701bp; fragments: 22; full length: 2 (>=2430.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

