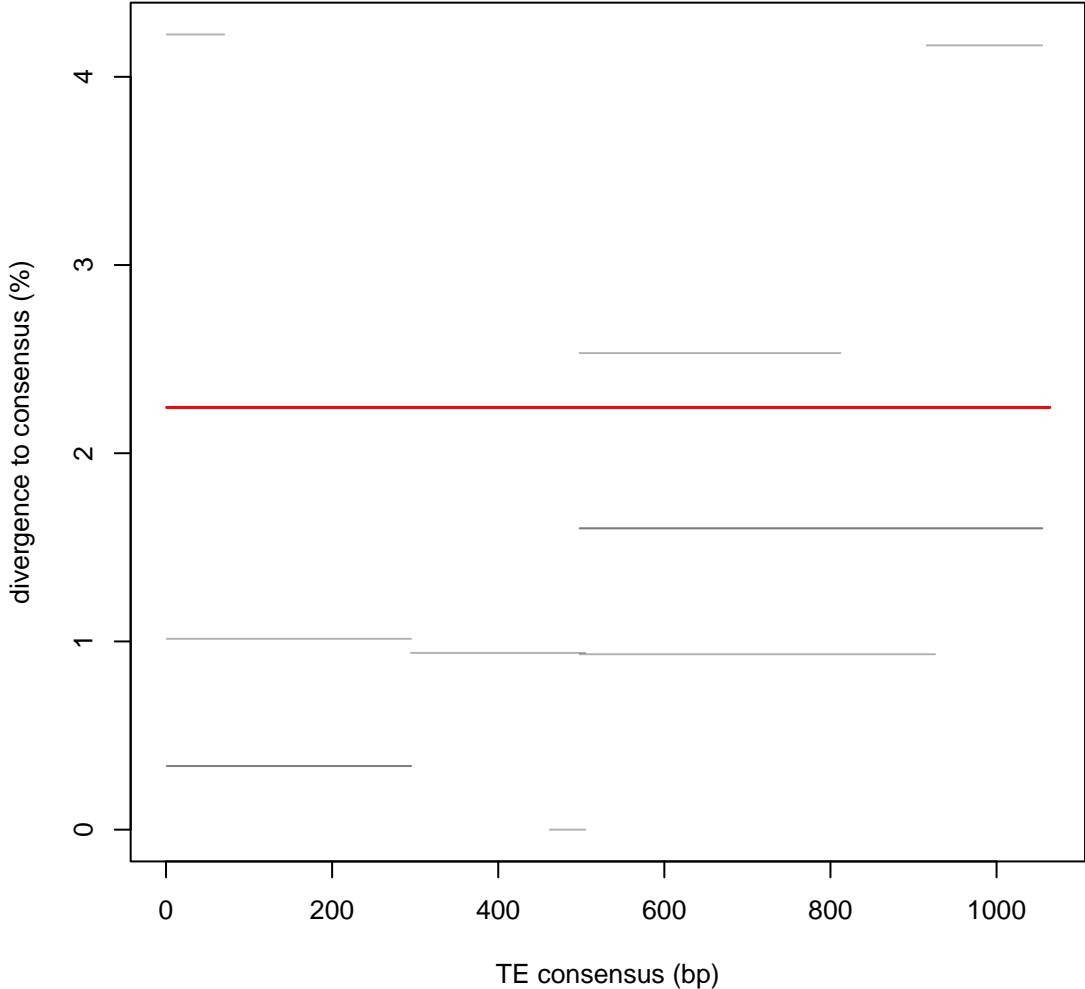
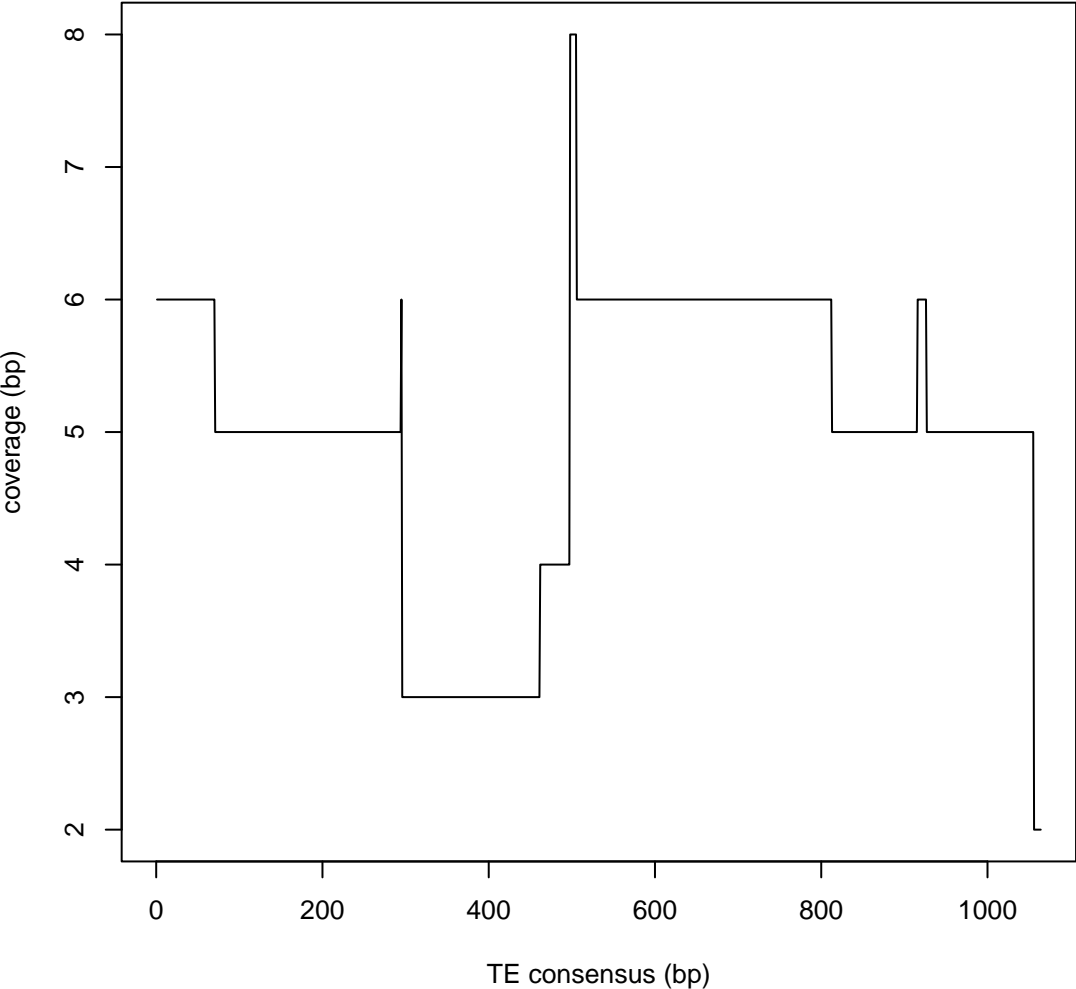


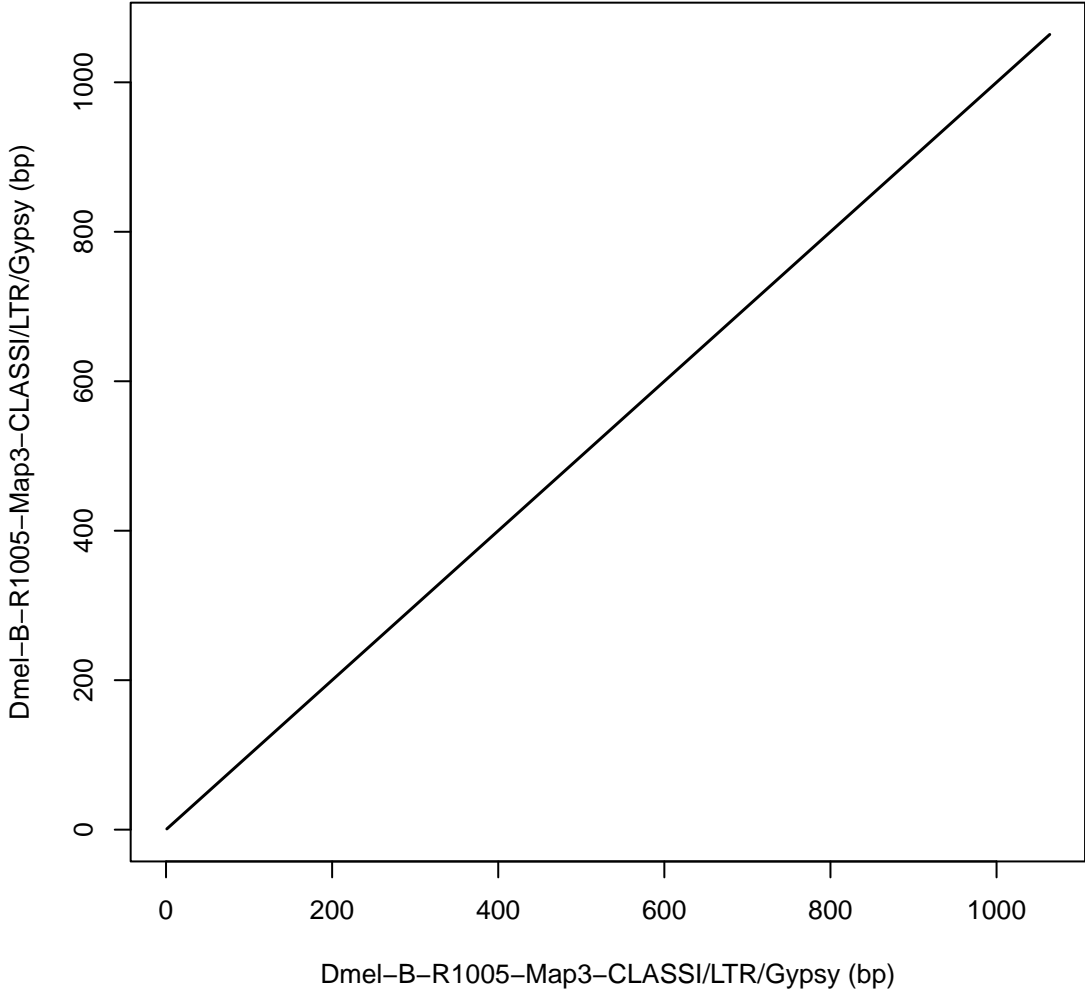
TE: Dmel-B-R1005-Map3-CLASSI/LTR/Gypsy
consensus size: 1064bp; fragments: 13; full length: 2 (>=957.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

