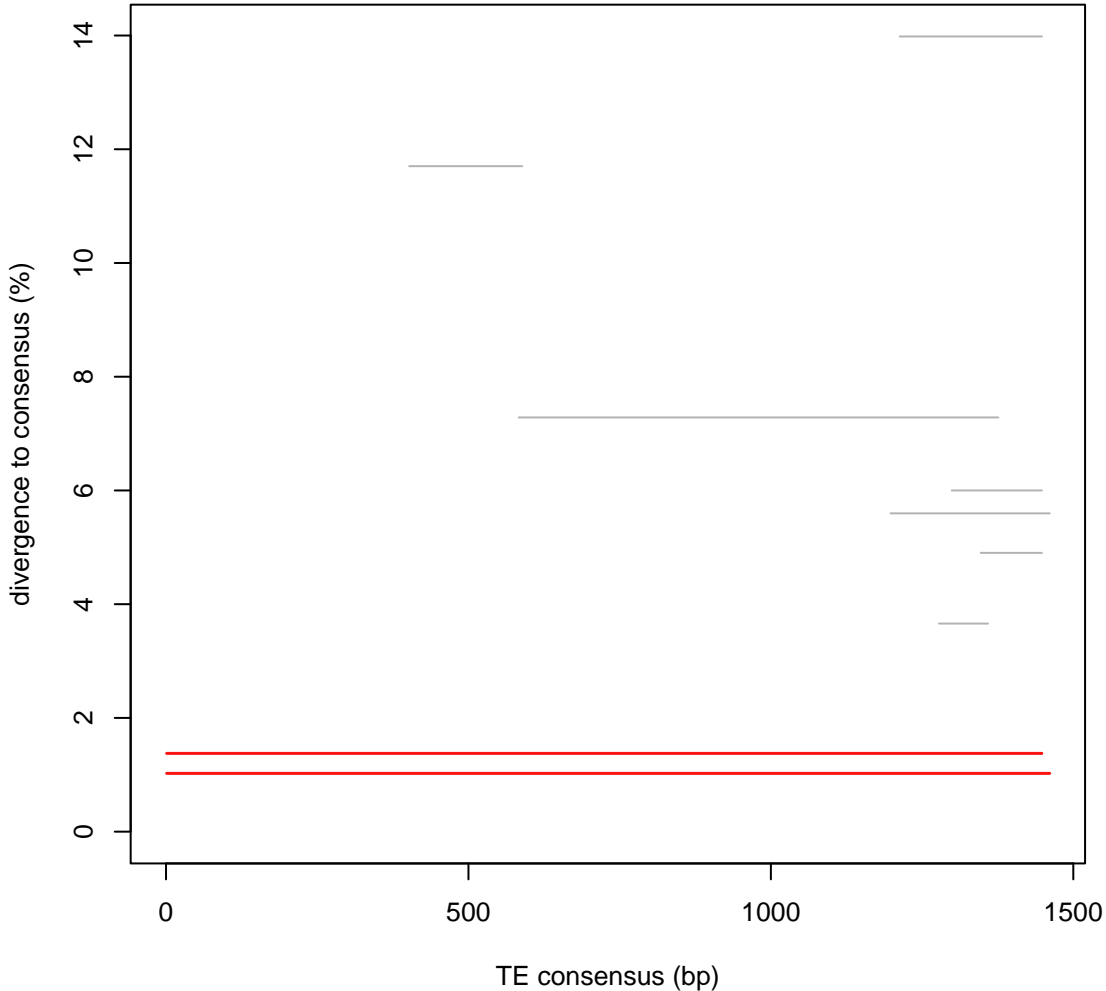
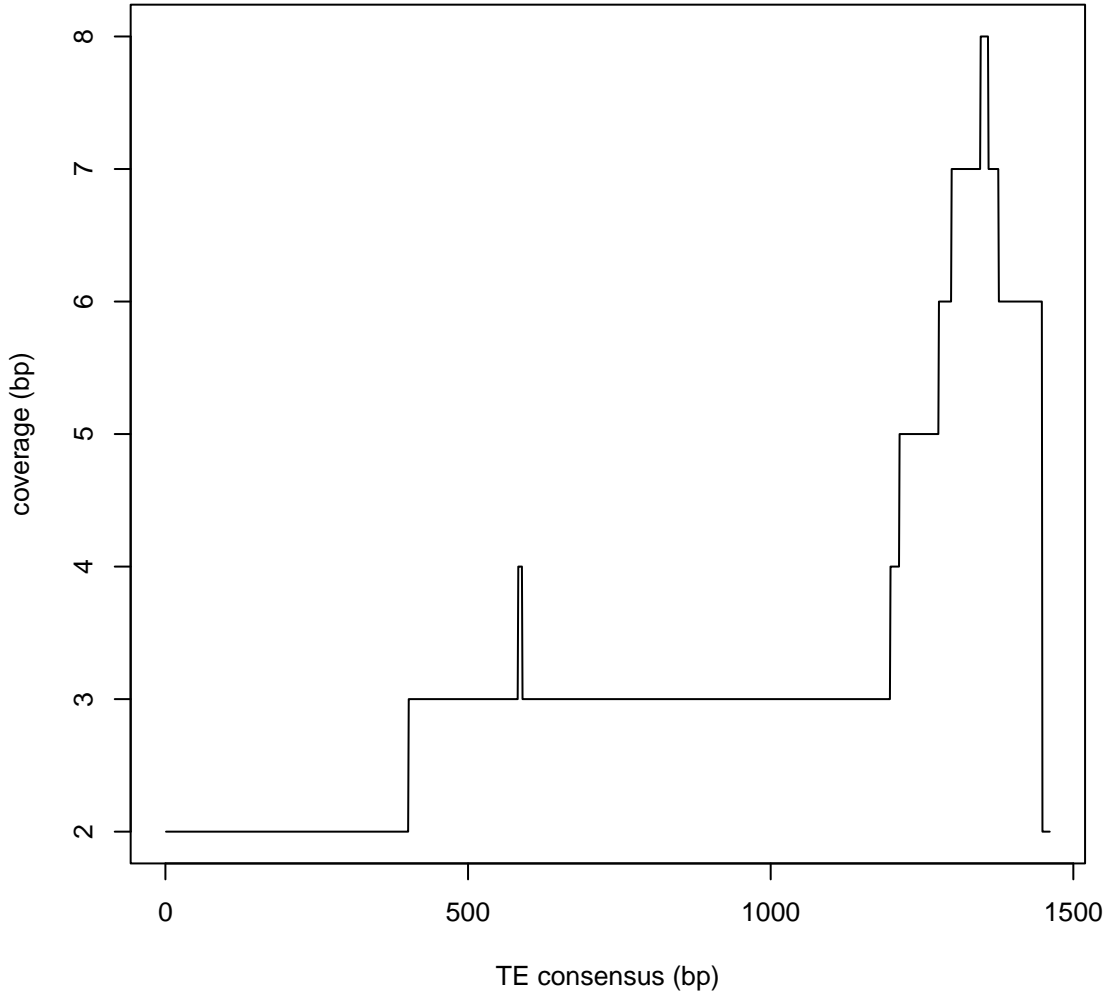


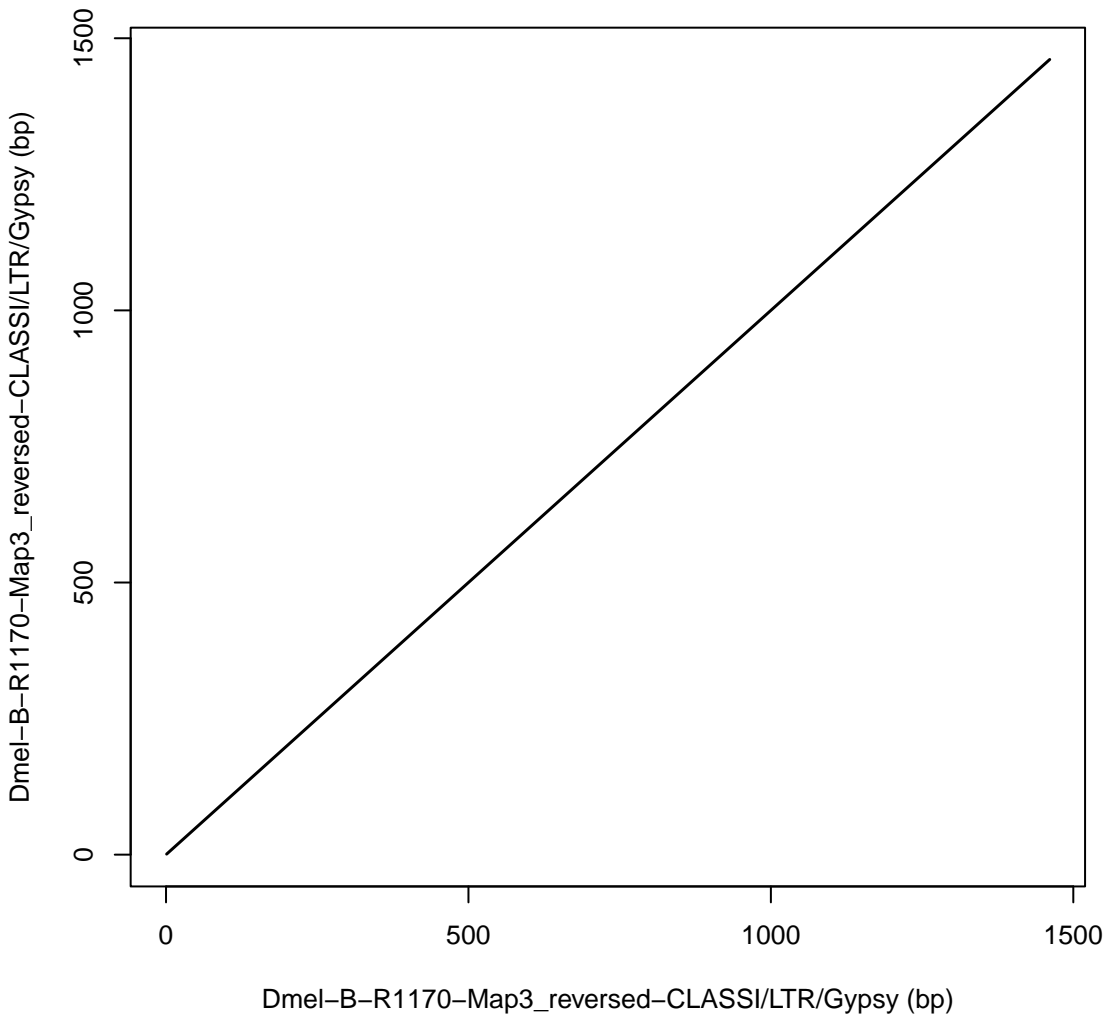
TE: Dmel-B-R1170-Map3_reversed-CLASSI/LTR/Gypsy
consensus size: 1461bp; fragments: 9; full length: 2 (>=1314.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

