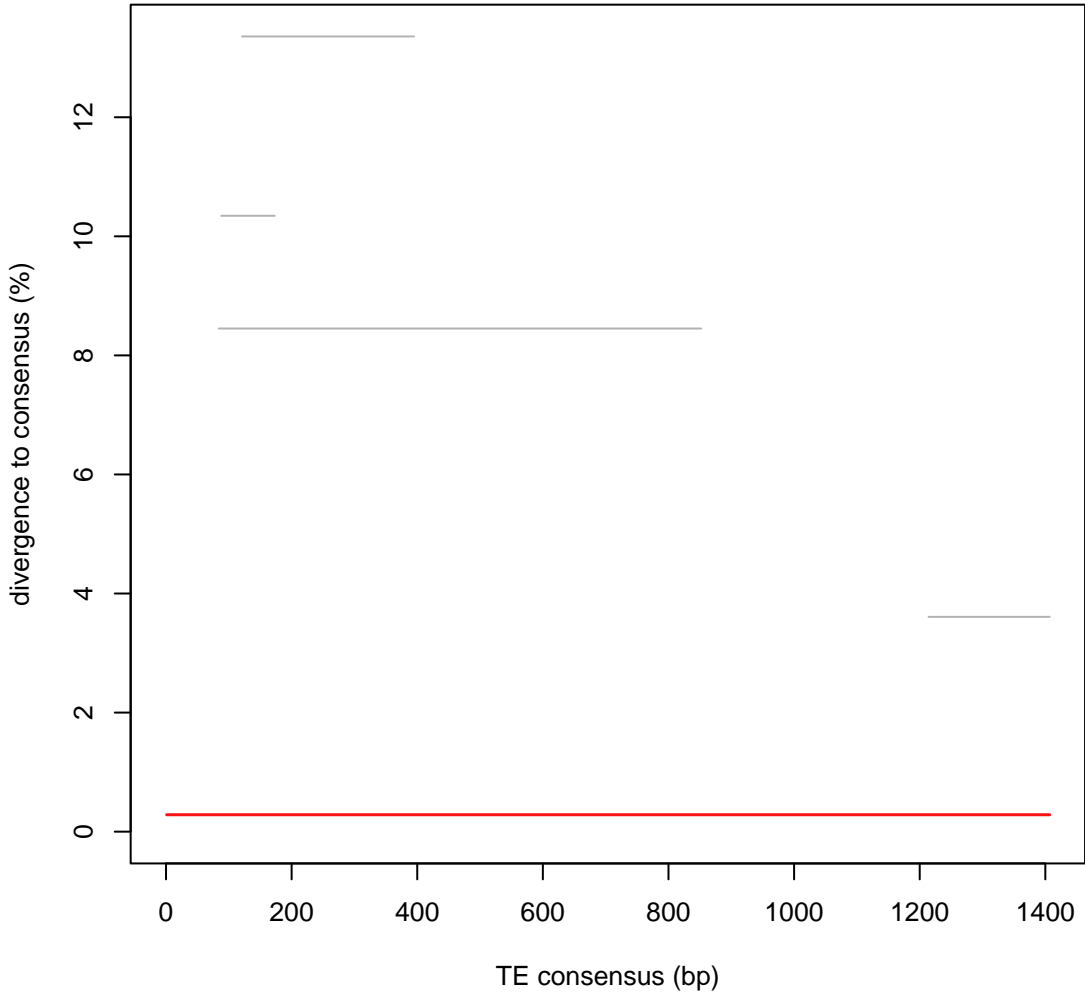
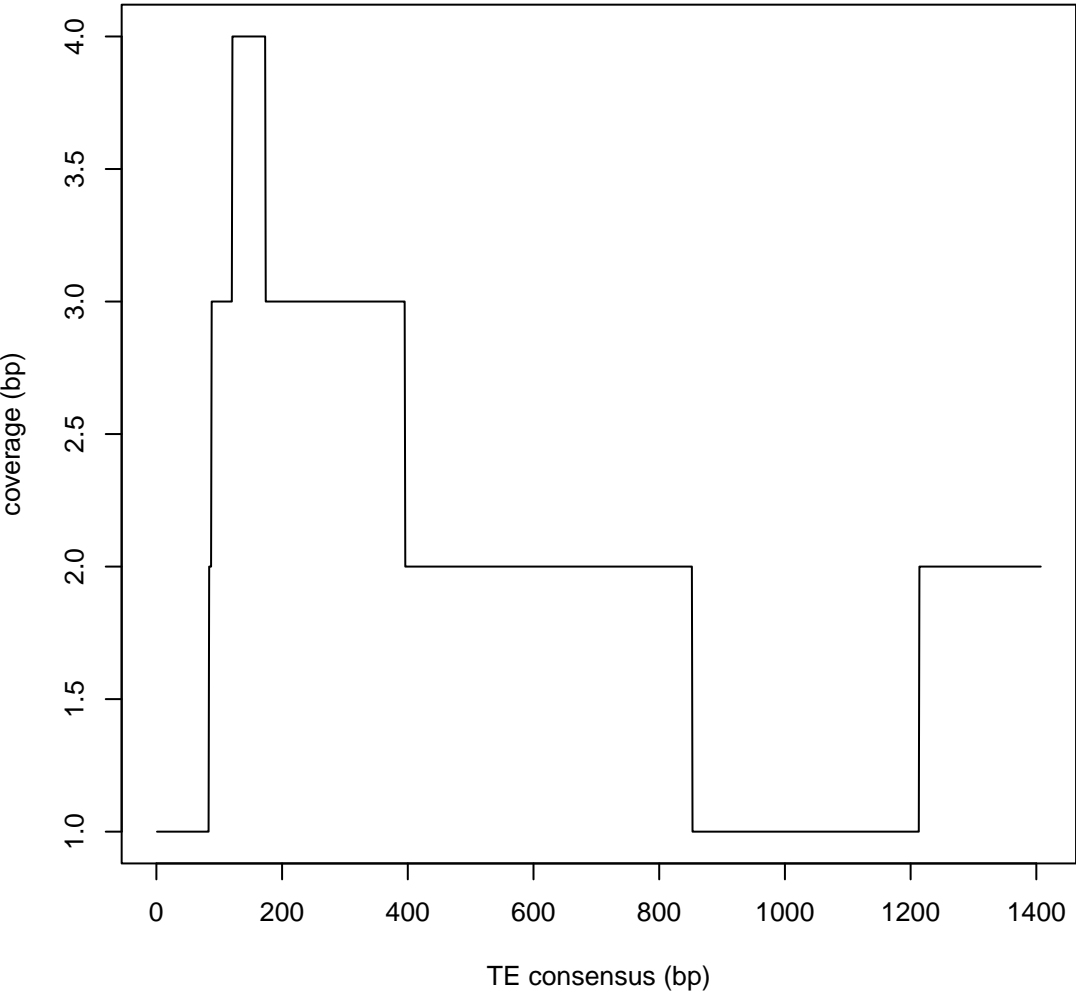


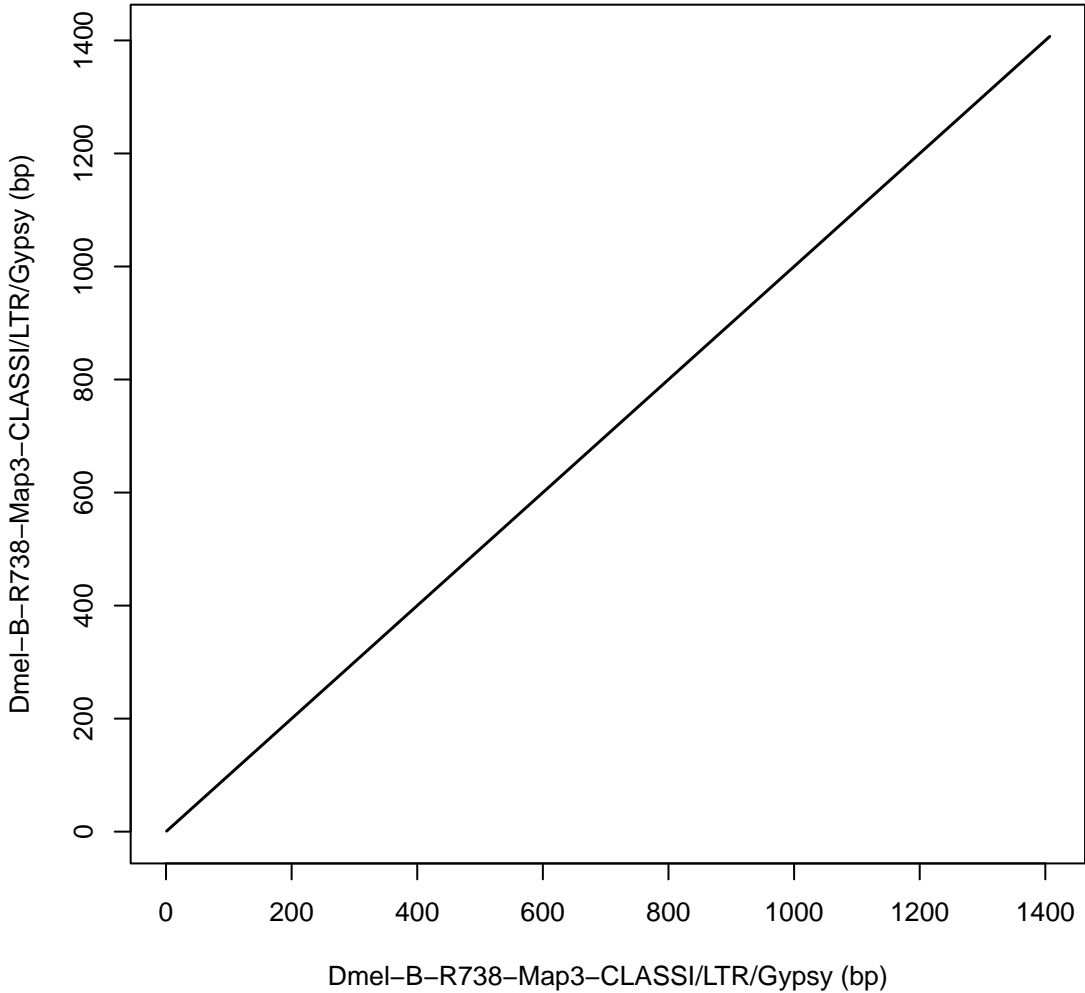
TE: Dmel-B-R738-Map3-CLASSI/LTR/Gypsy
consensus size: 1407bp; fragments: 5; full length: 1 (>=1266.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

