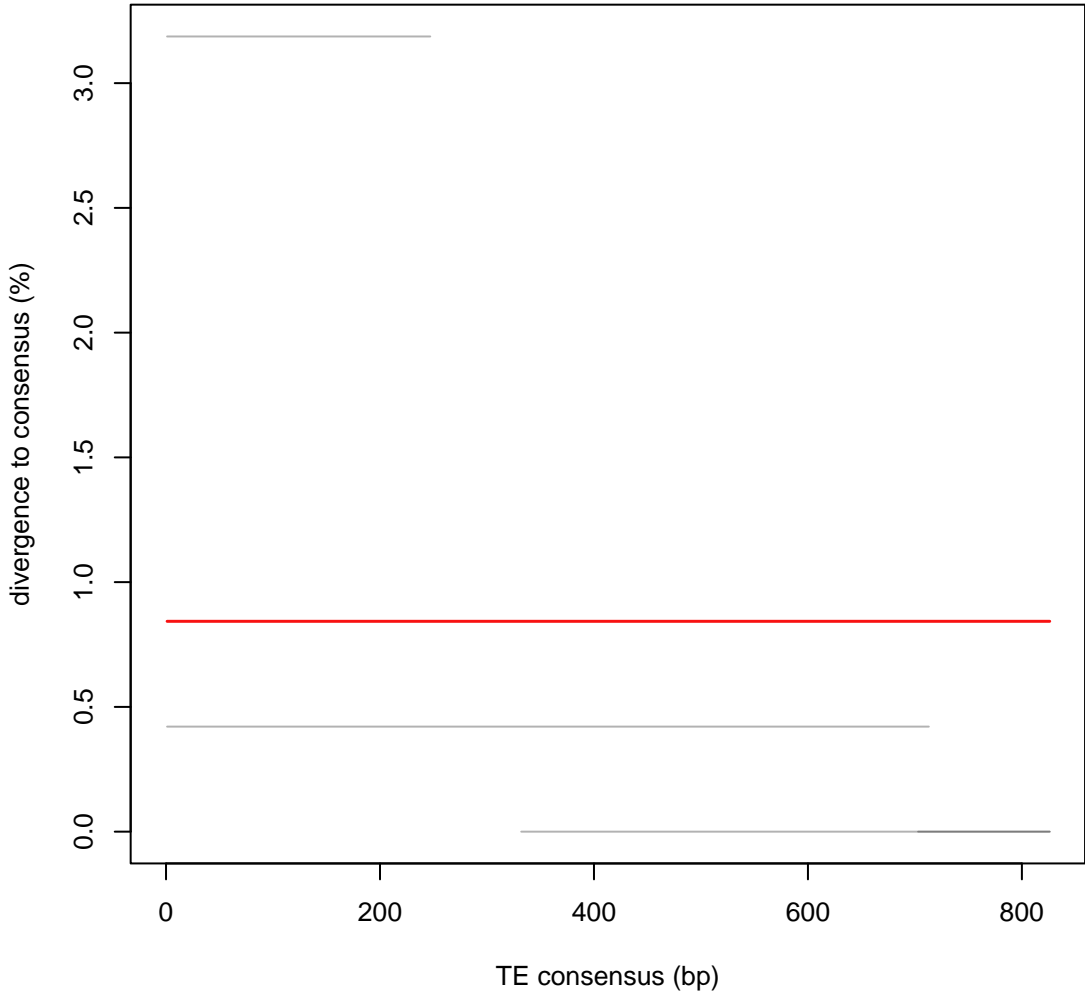
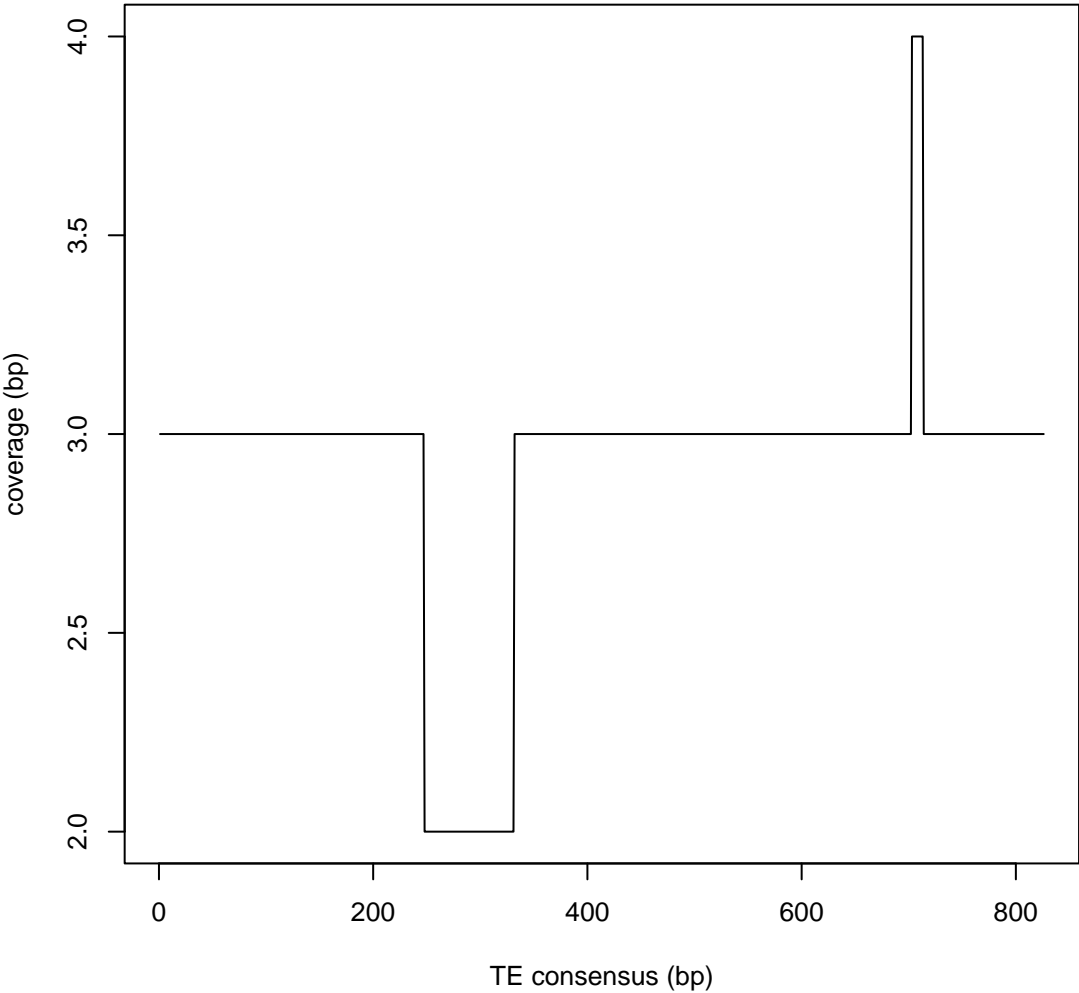


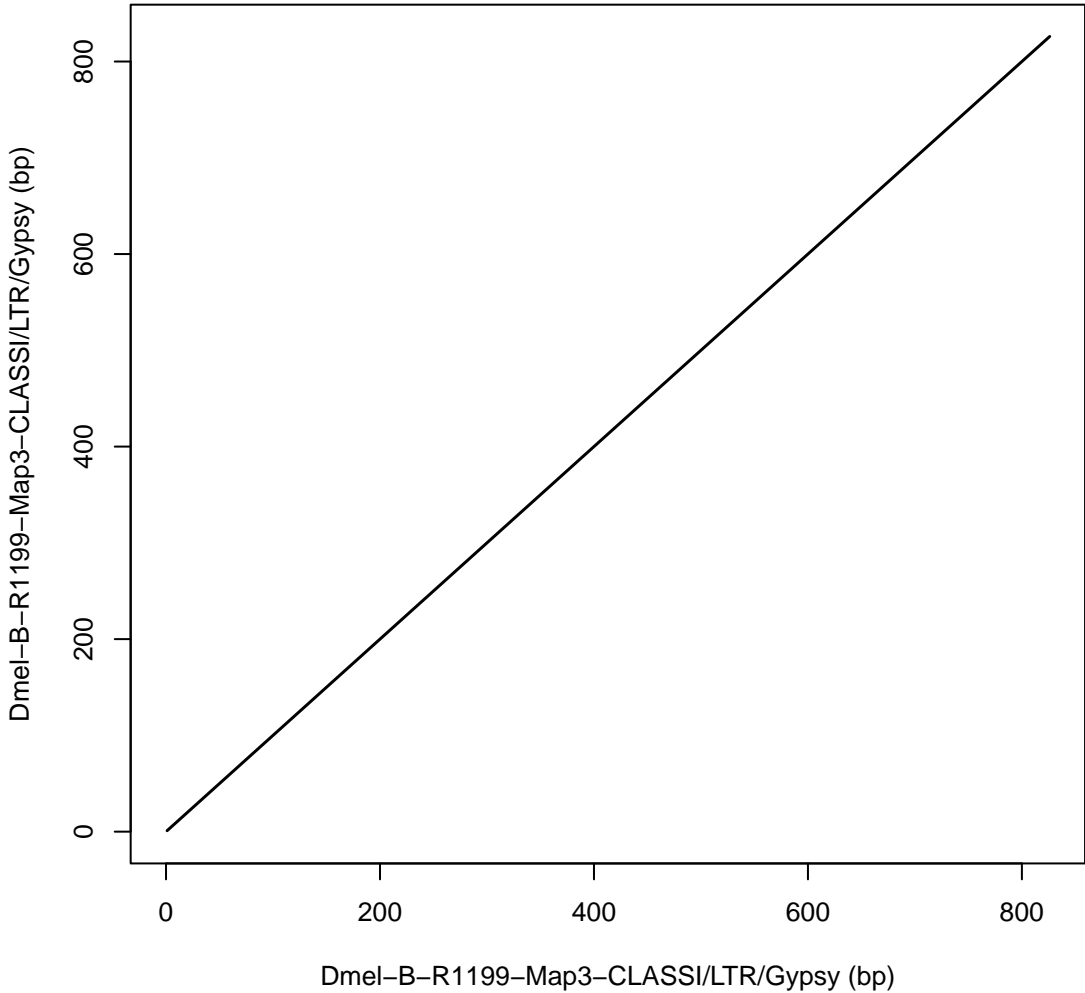
TE: Dmel-B-R1199-Map3-CLASSI/LTR/Gypsy  
consensus size: 826bp; fragments: 5; full length: 1 (>=743.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

