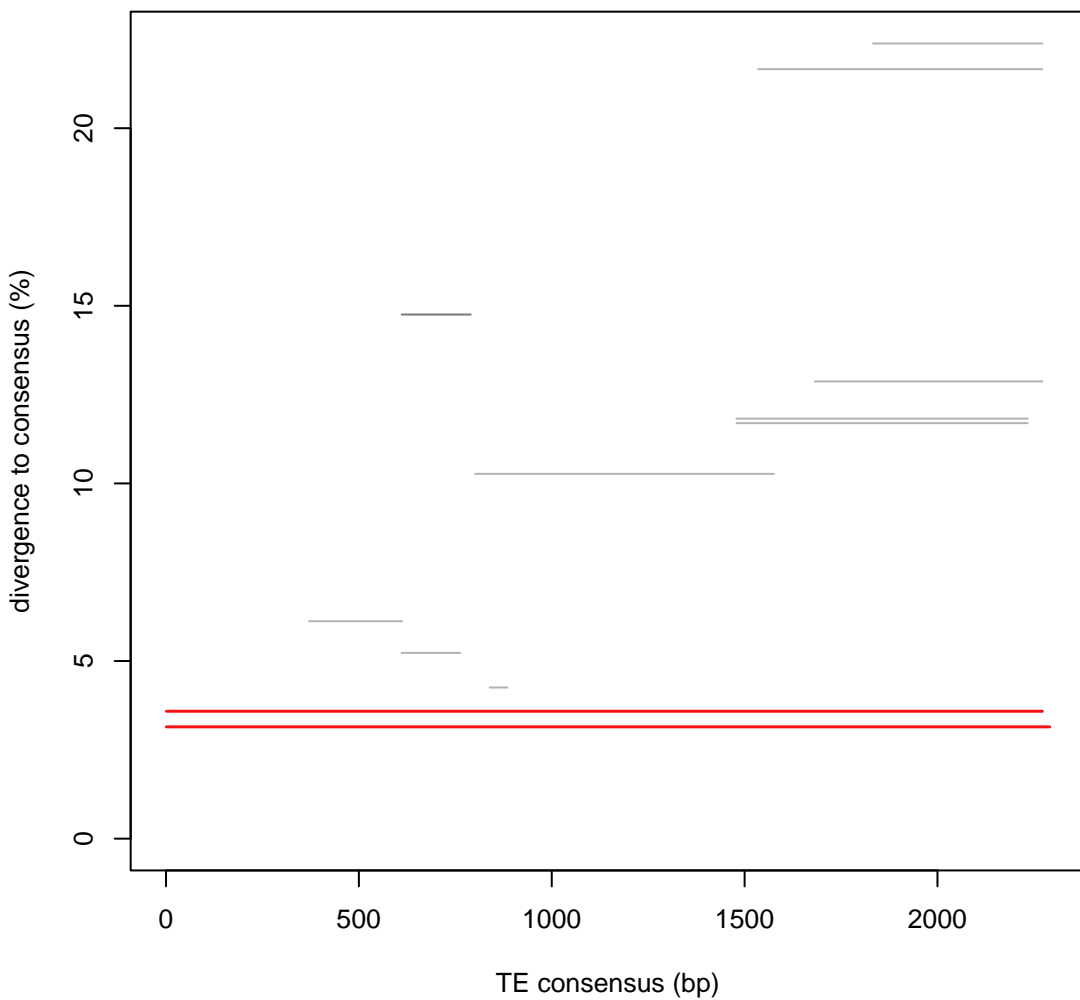
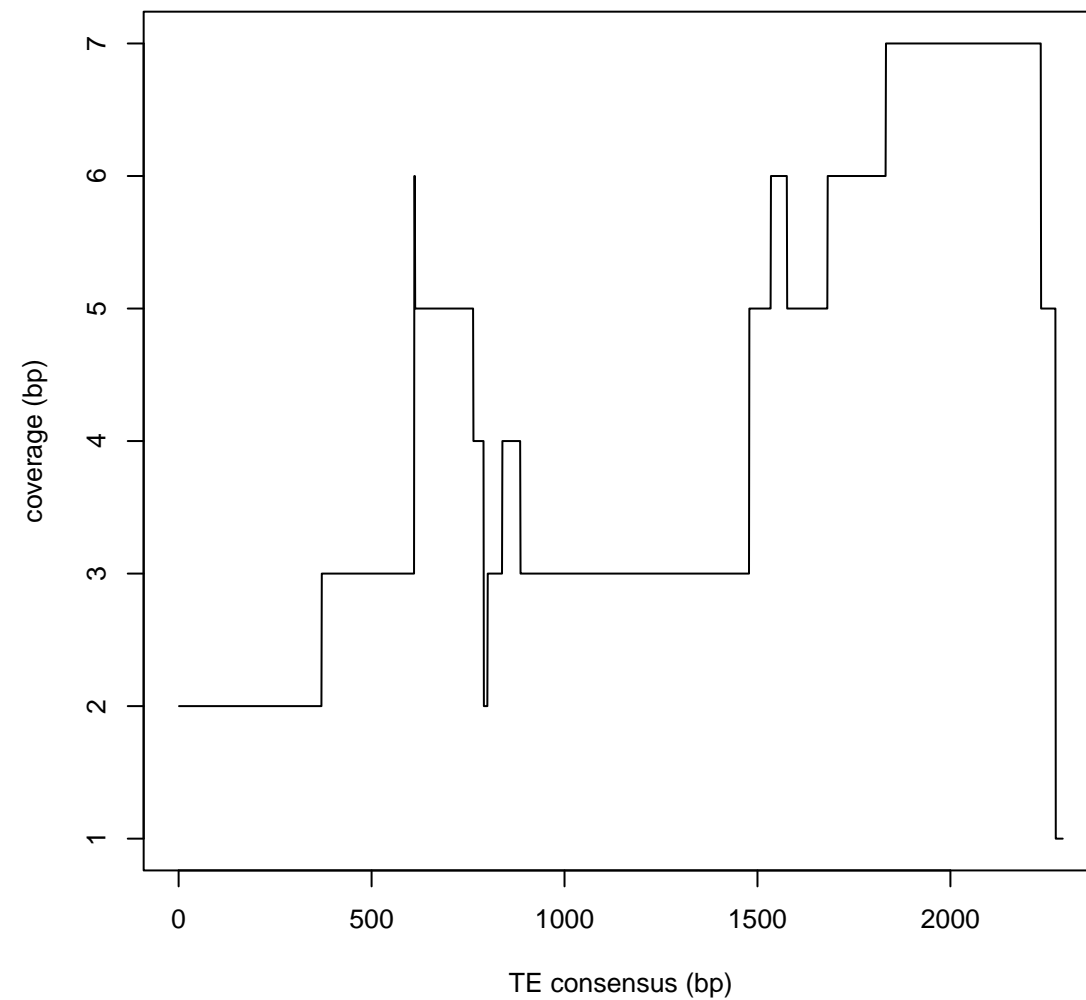


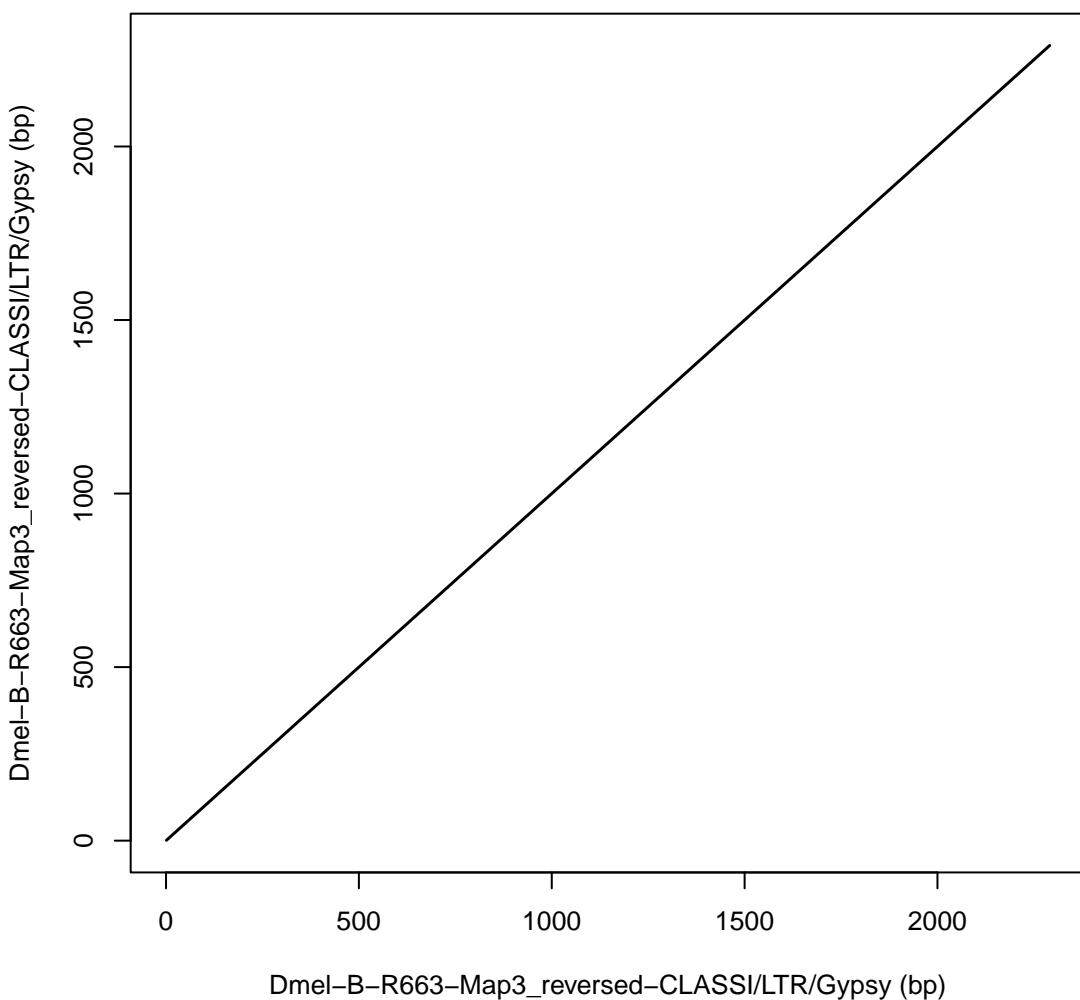
TE: Dmel-B-R663-Map3\_reversed-CLASSI/LTR/Gypsy  
consensus size: 2291bp; fragments: 13; full length: 2 (>=2061.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

