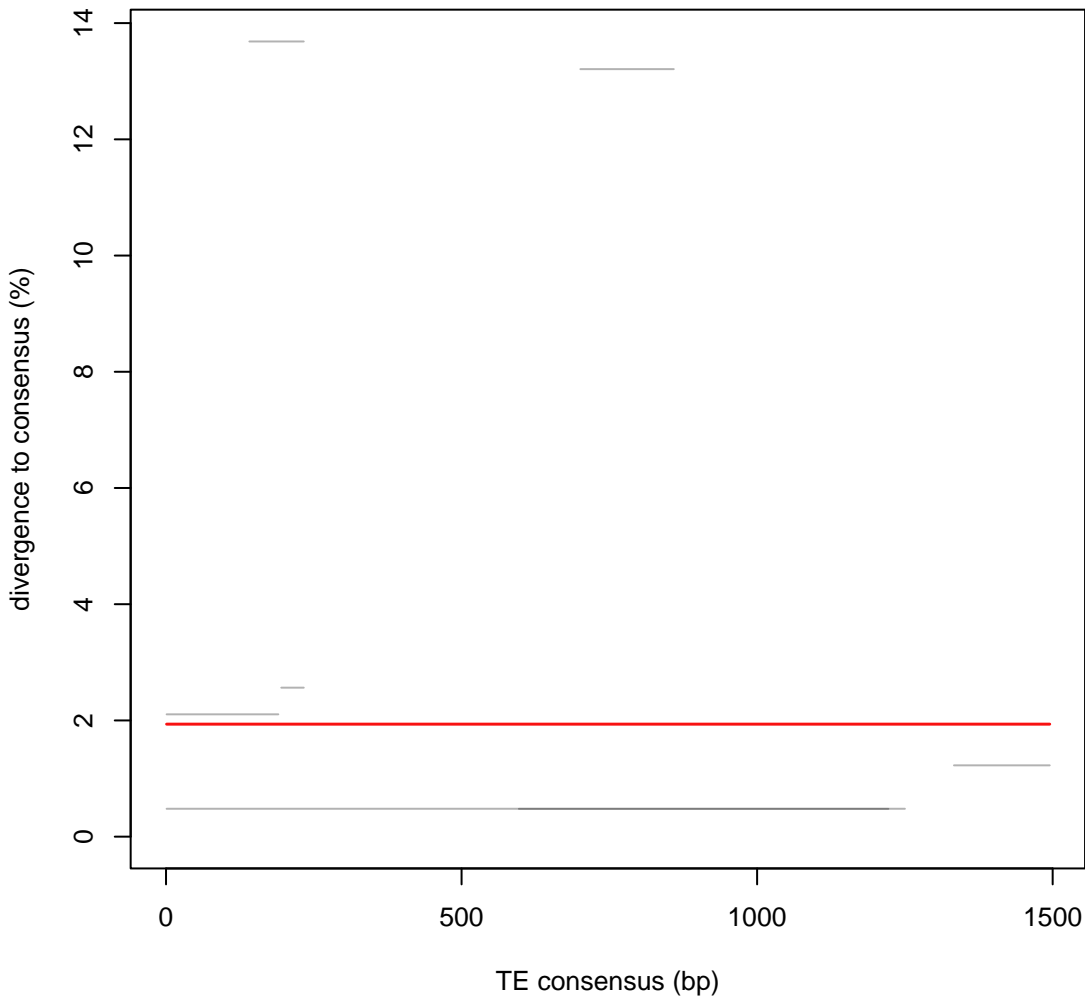
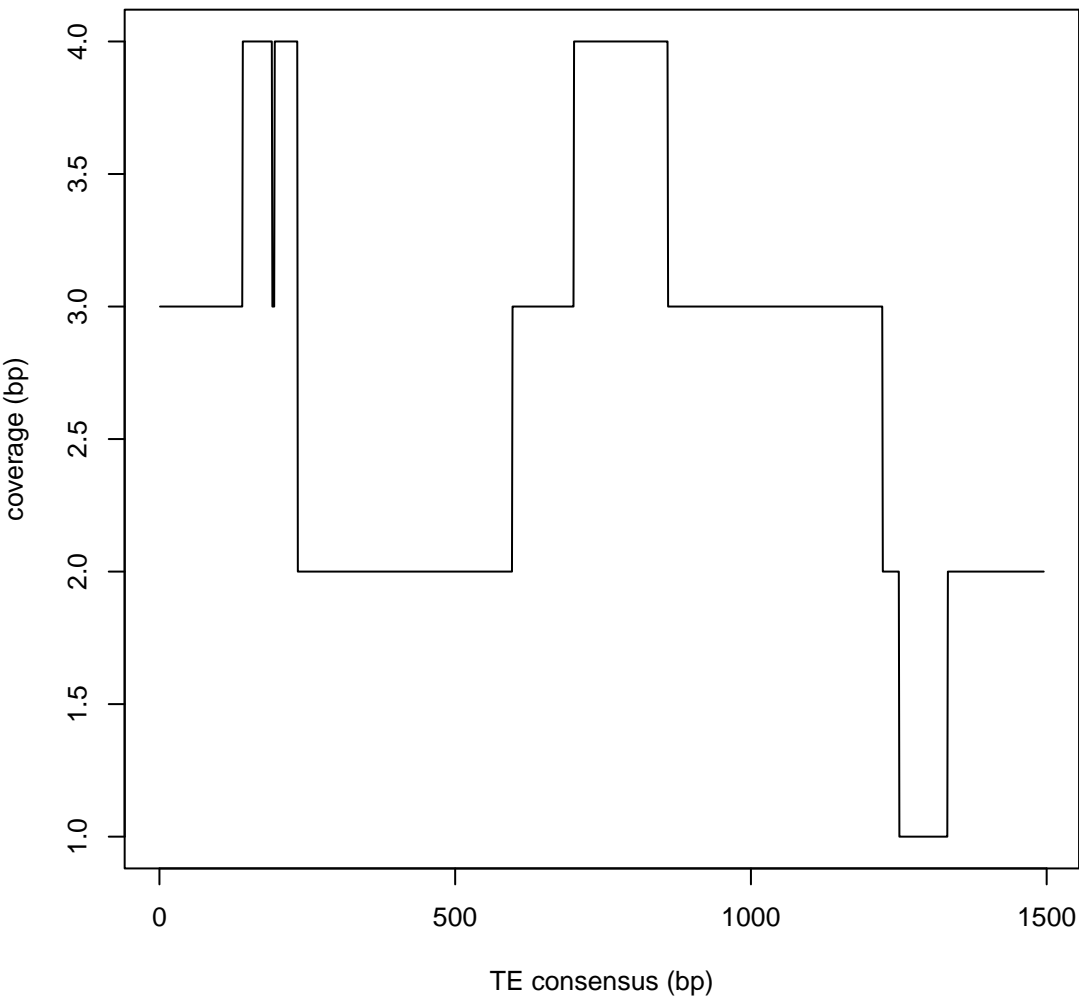


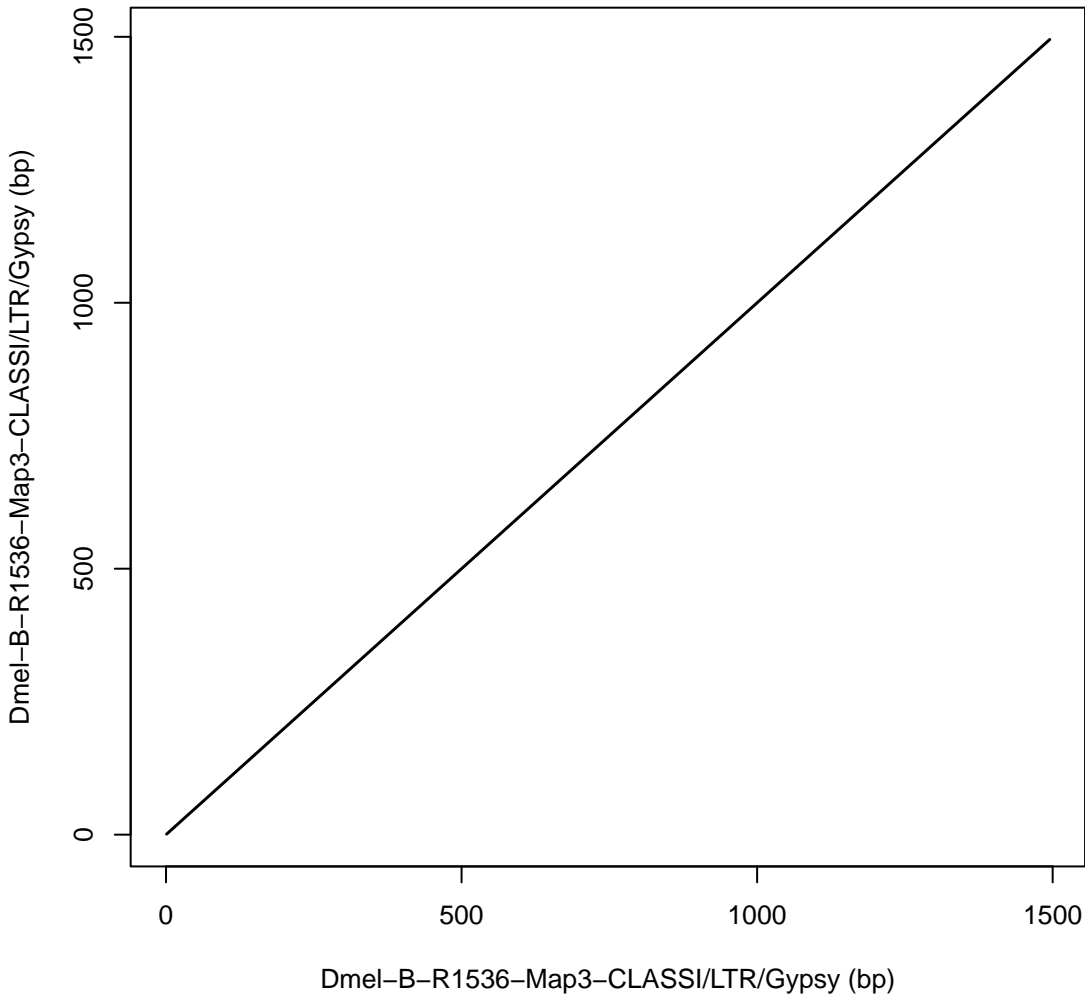
TE: Dmel-B-R1536-Map3-CLASSI/LTR/Gypsy  
consensus size: 1495bp; fragments: 8; full length: 1 (>=1345.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

