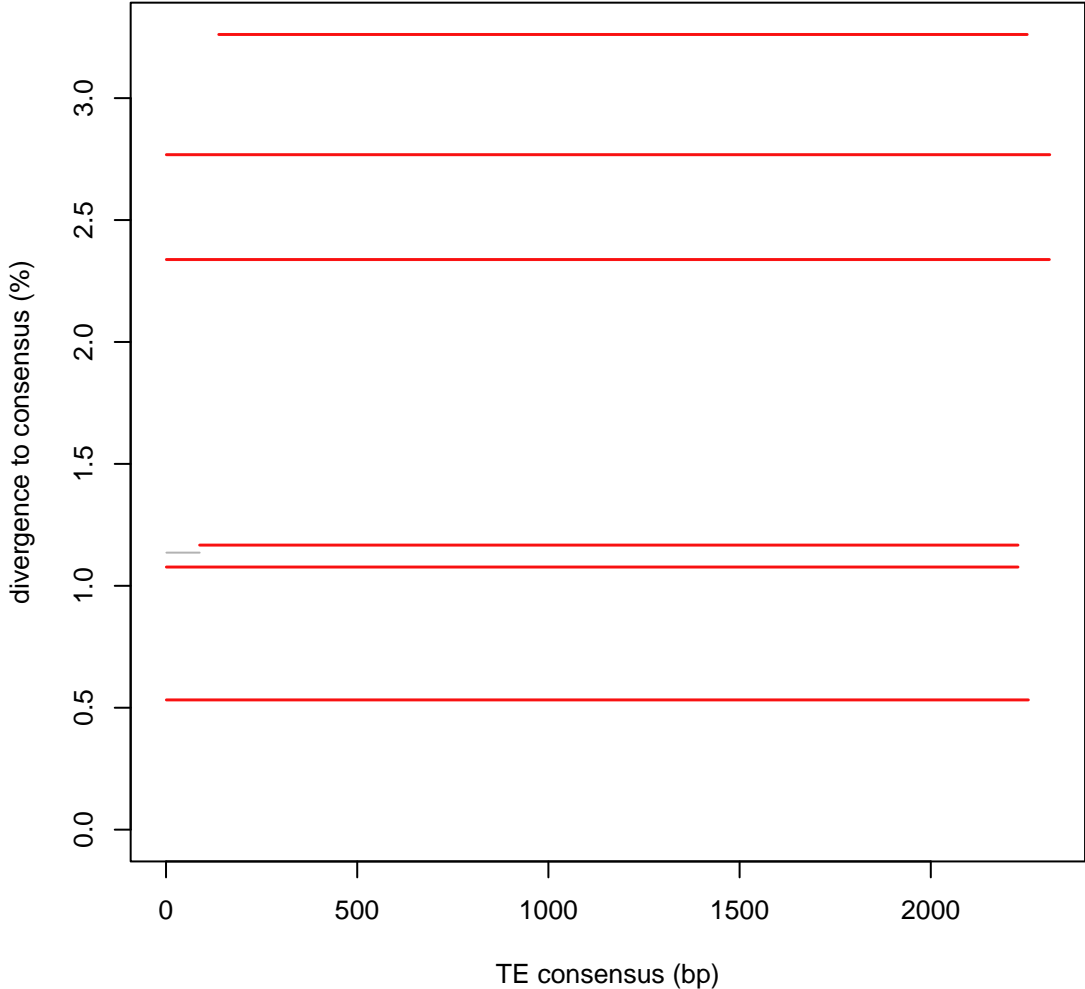
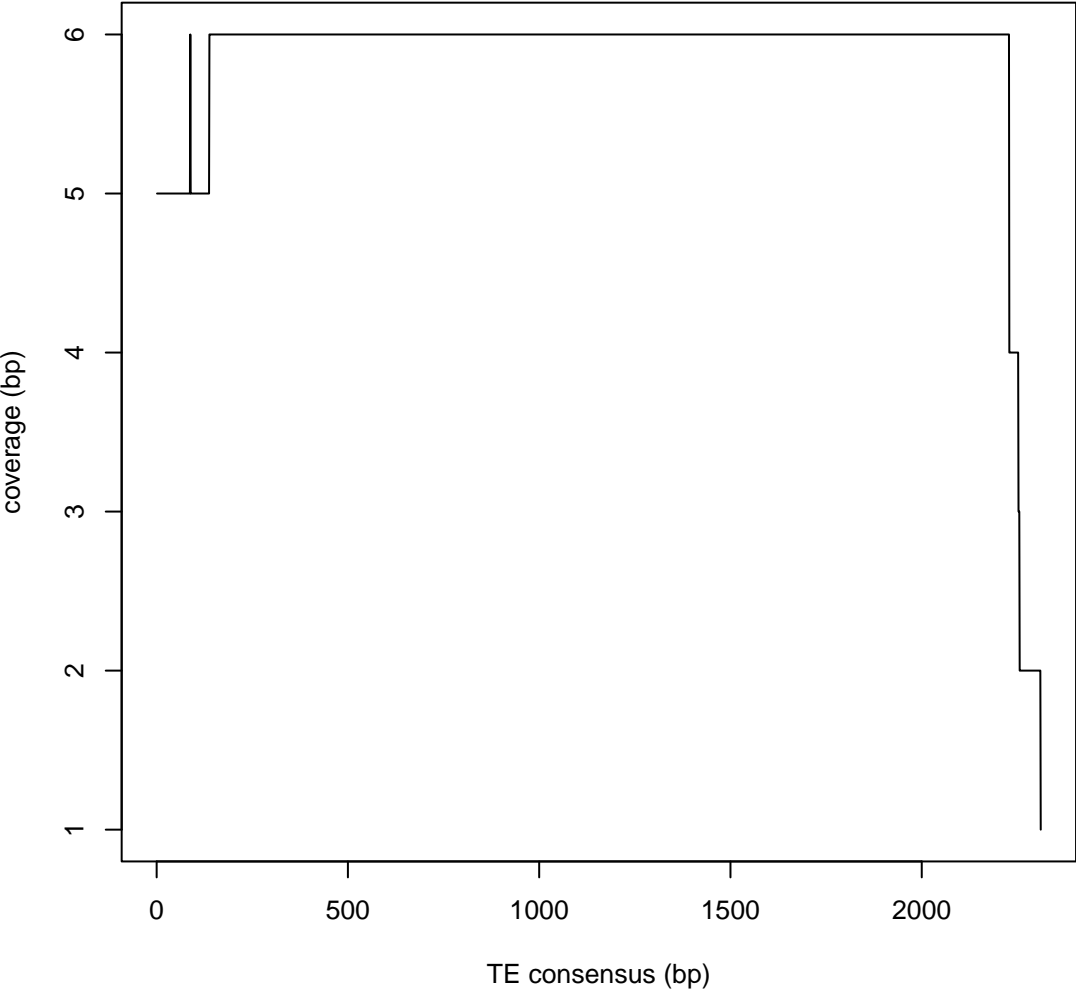


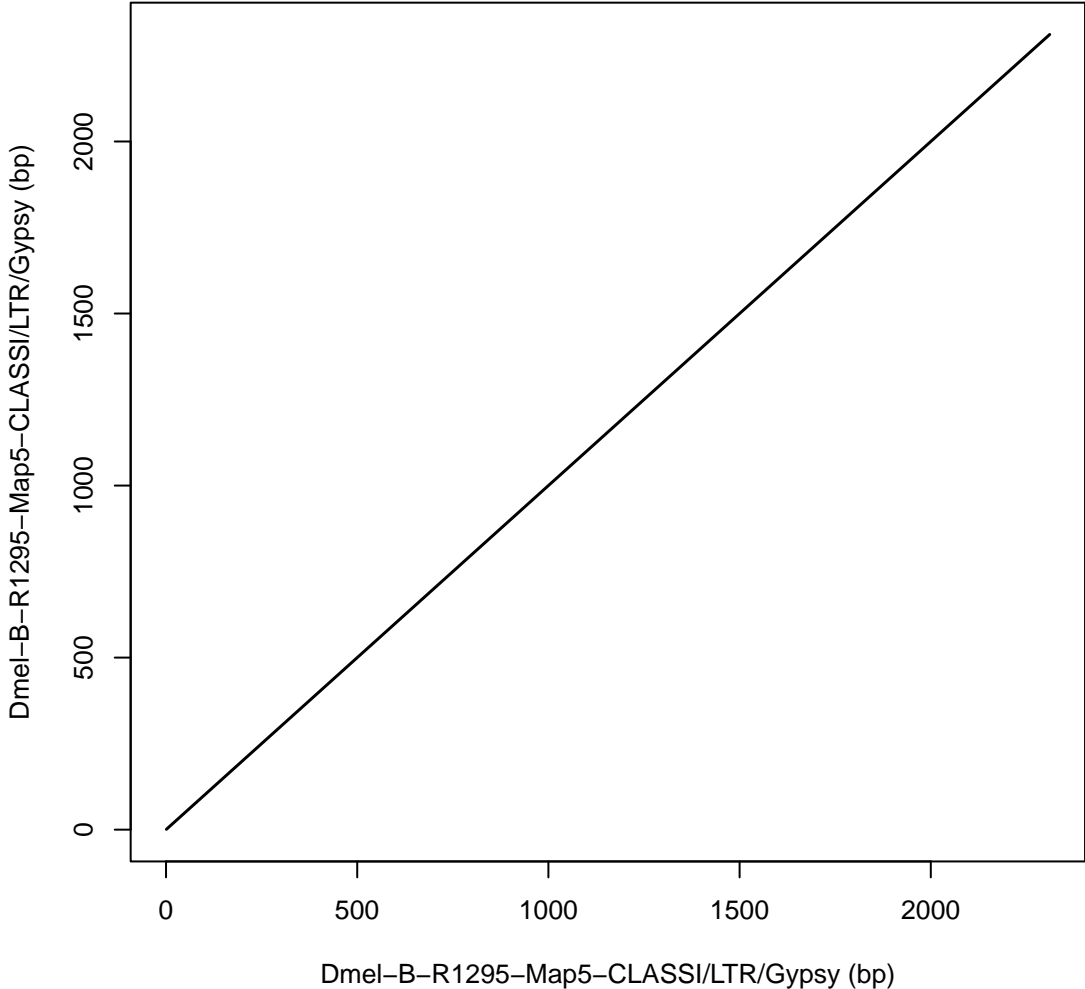
TE: Dmel-B-R1295-Map5-CLASSI/LTR/Gypsy  
consensus size: 2311bp; fragments: 7; full length: 6 (>=2079.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

