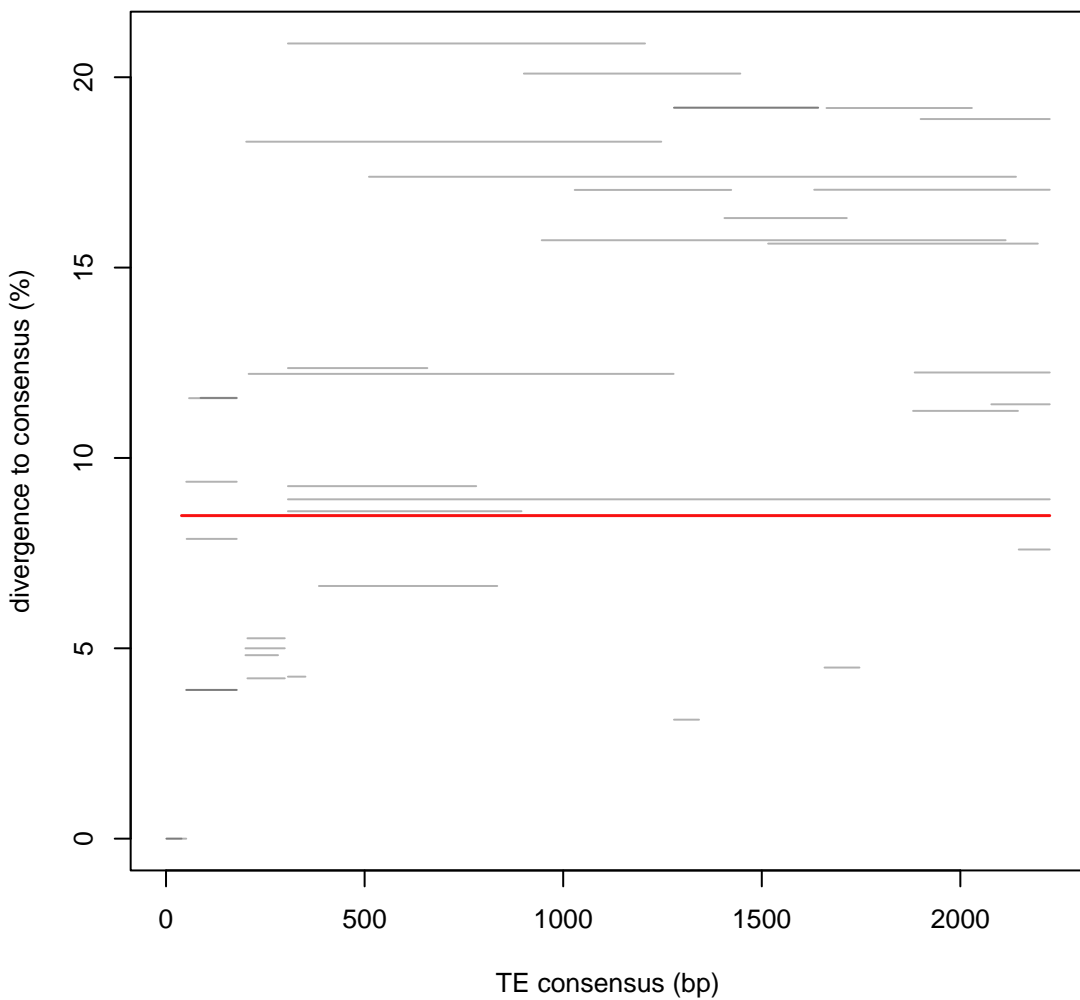
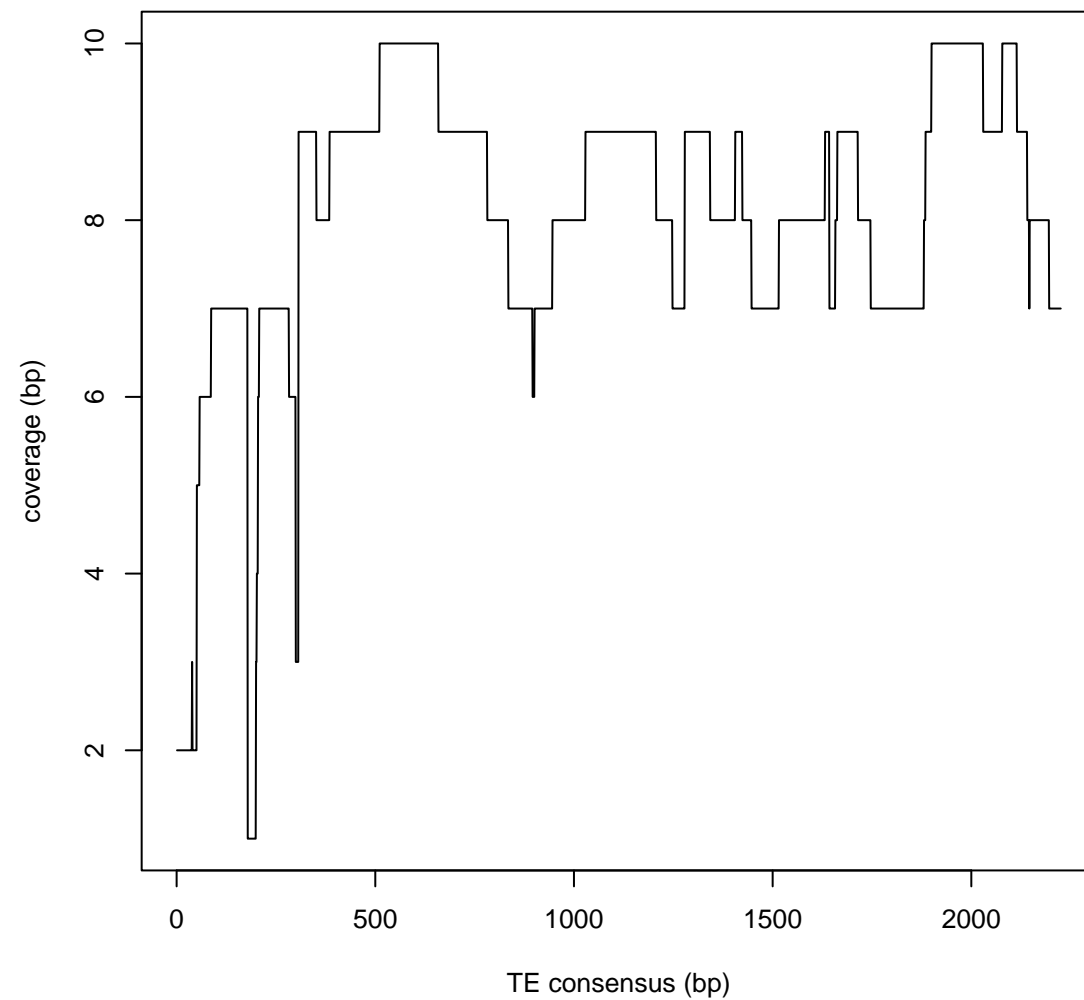


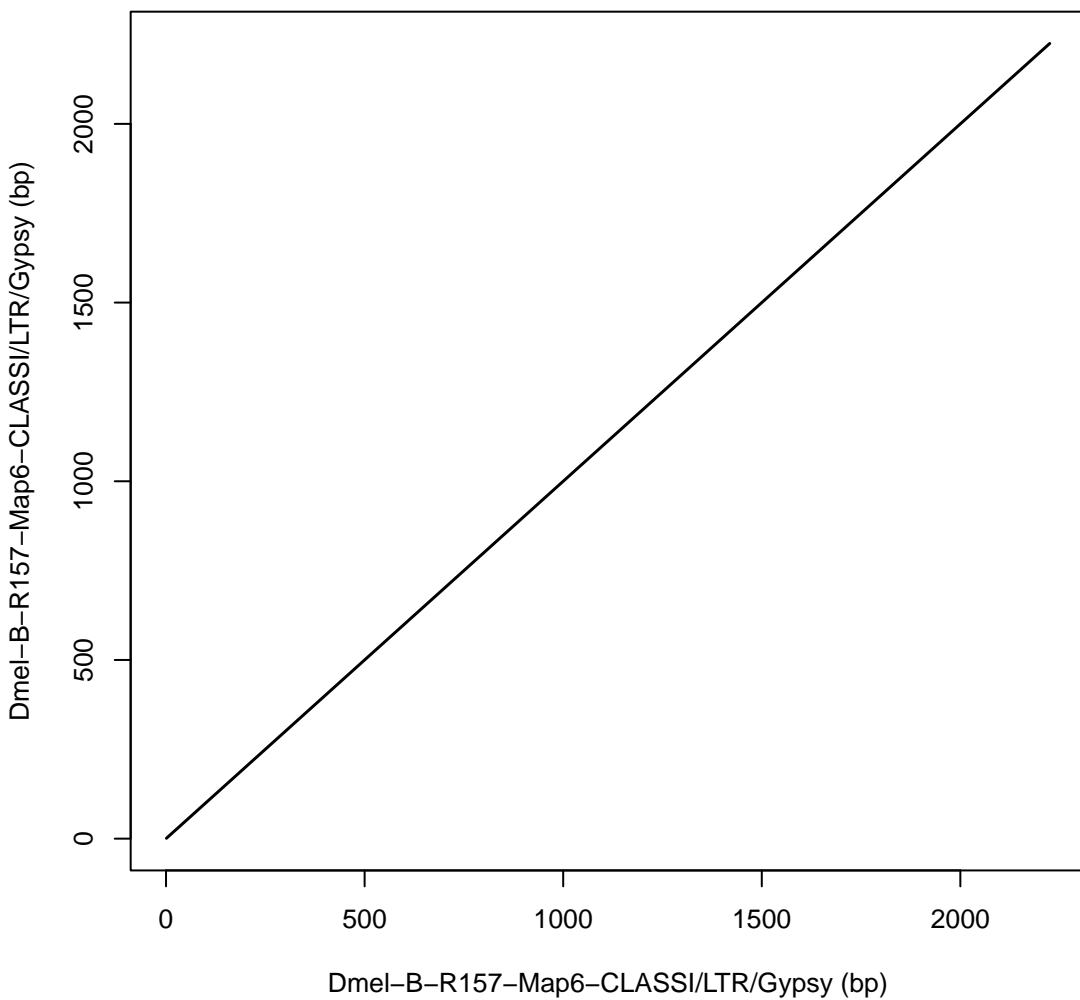
TE: Dmel-B-R157-Map6-CLASSI/LTR/Gypsy  
consensus size: 2225bp; fragments: 39; full length: 1 (>=2002.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

