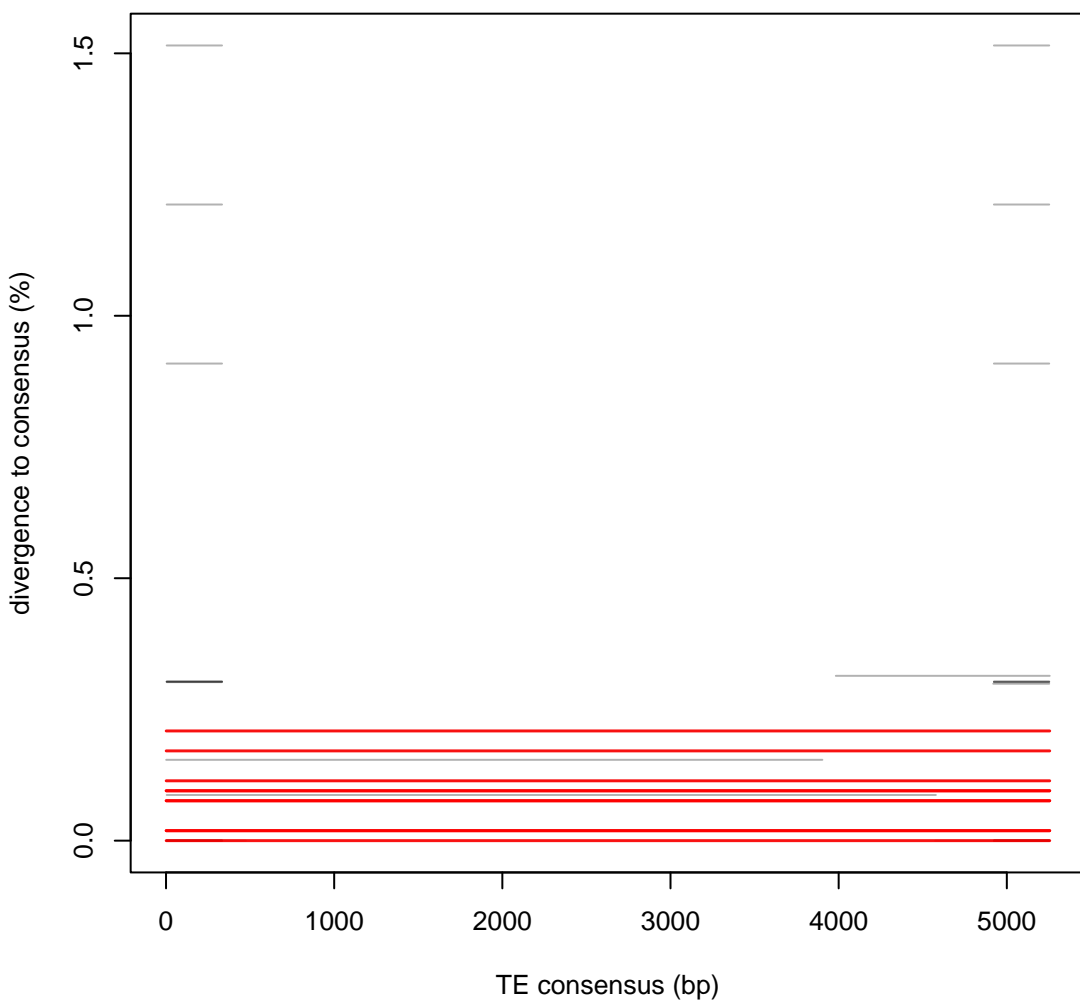
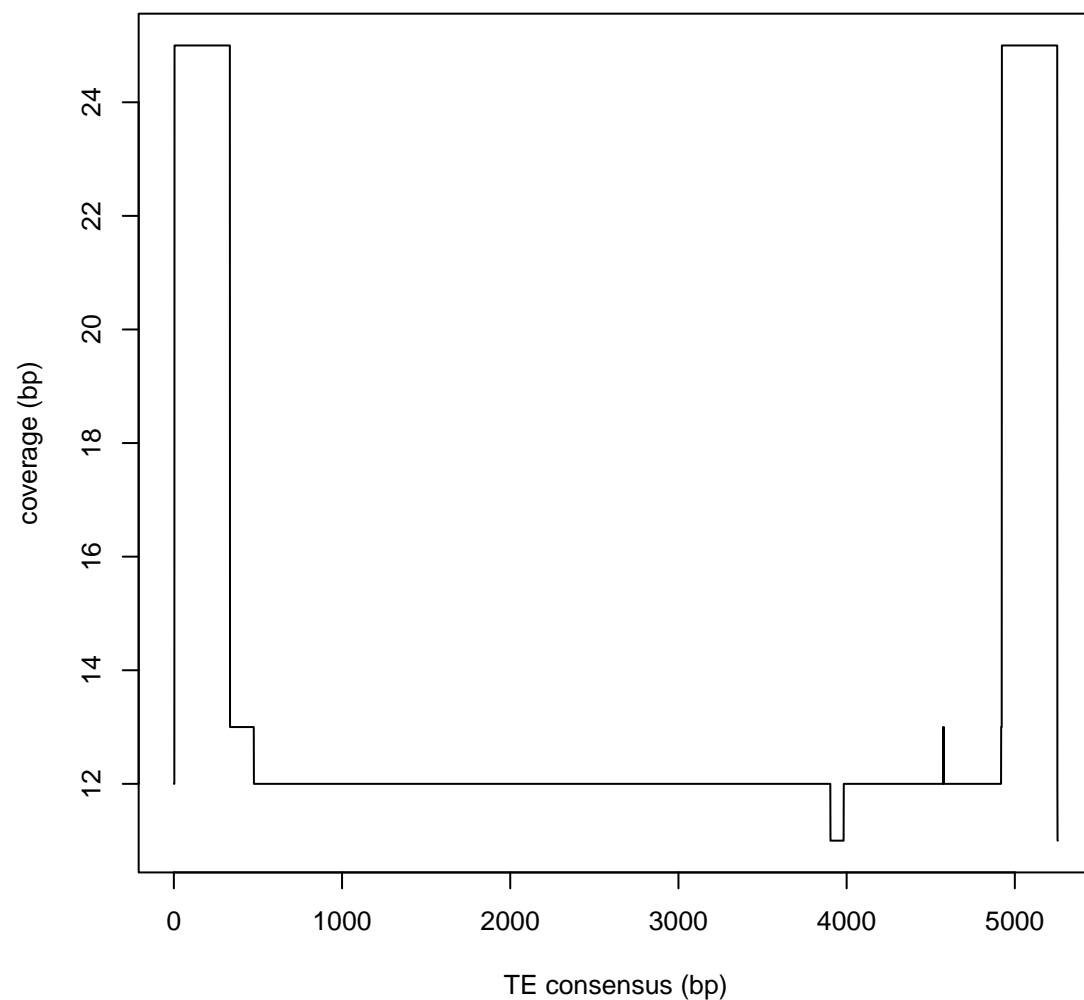


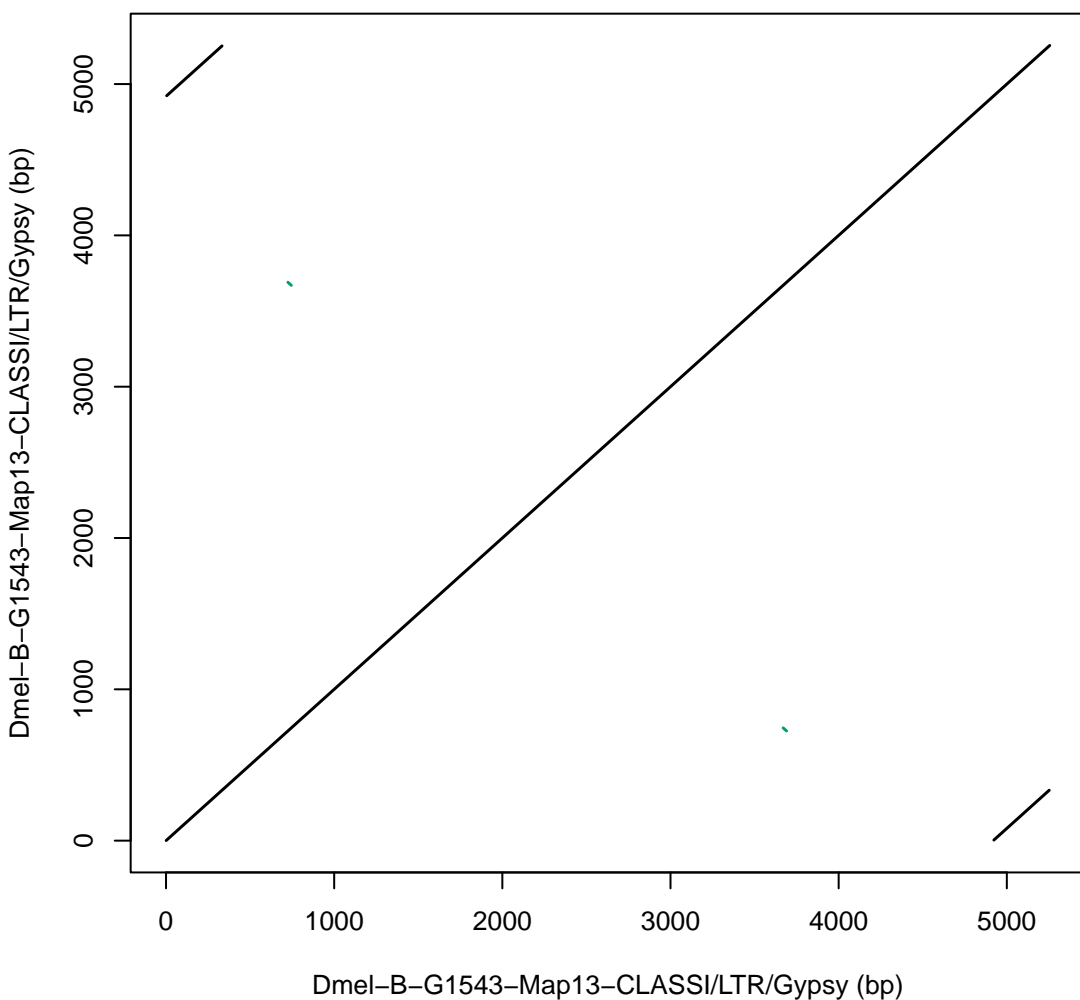
TE: Dmel-B-G1543-Map13-CLASSI/LTR/Gypsy  
consensus size: 5255bp; fragments: 40; full length: 10 (>=4729.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

