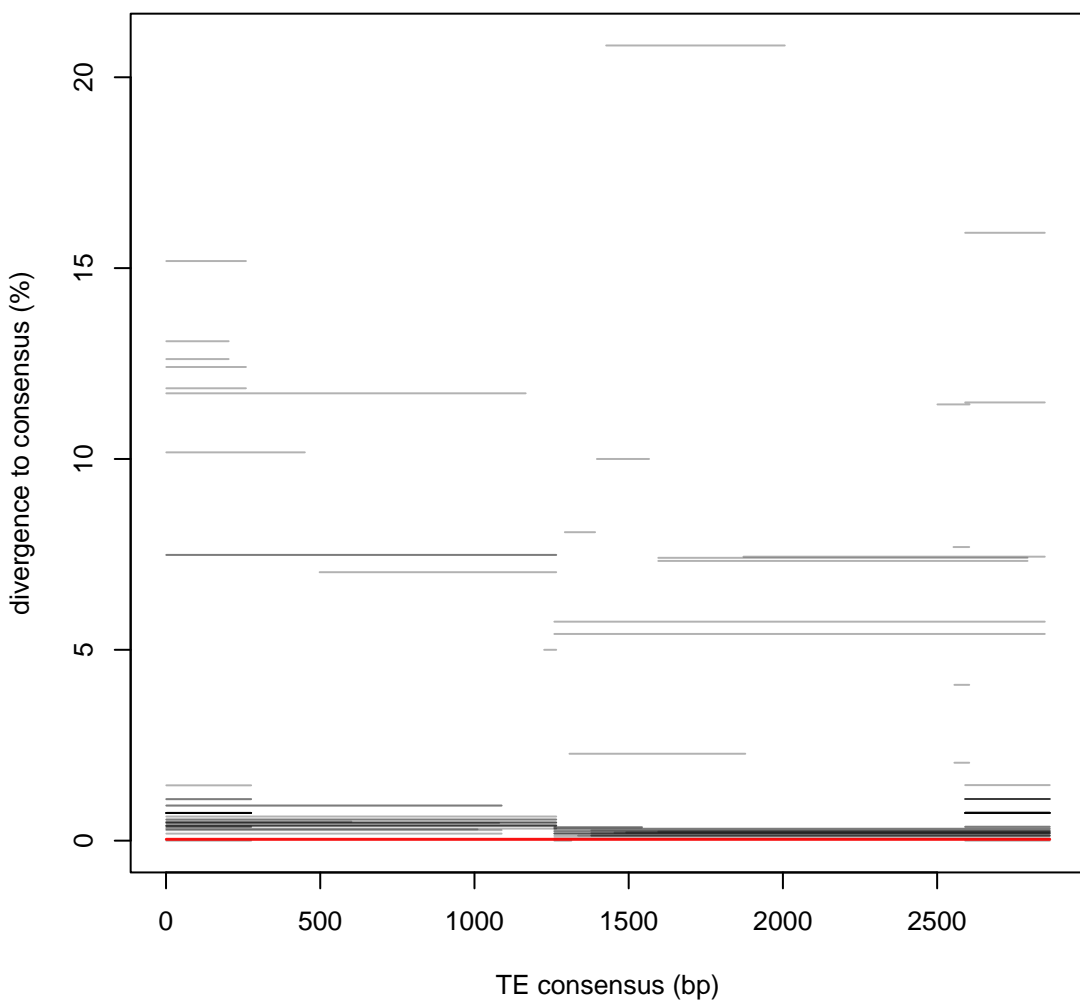
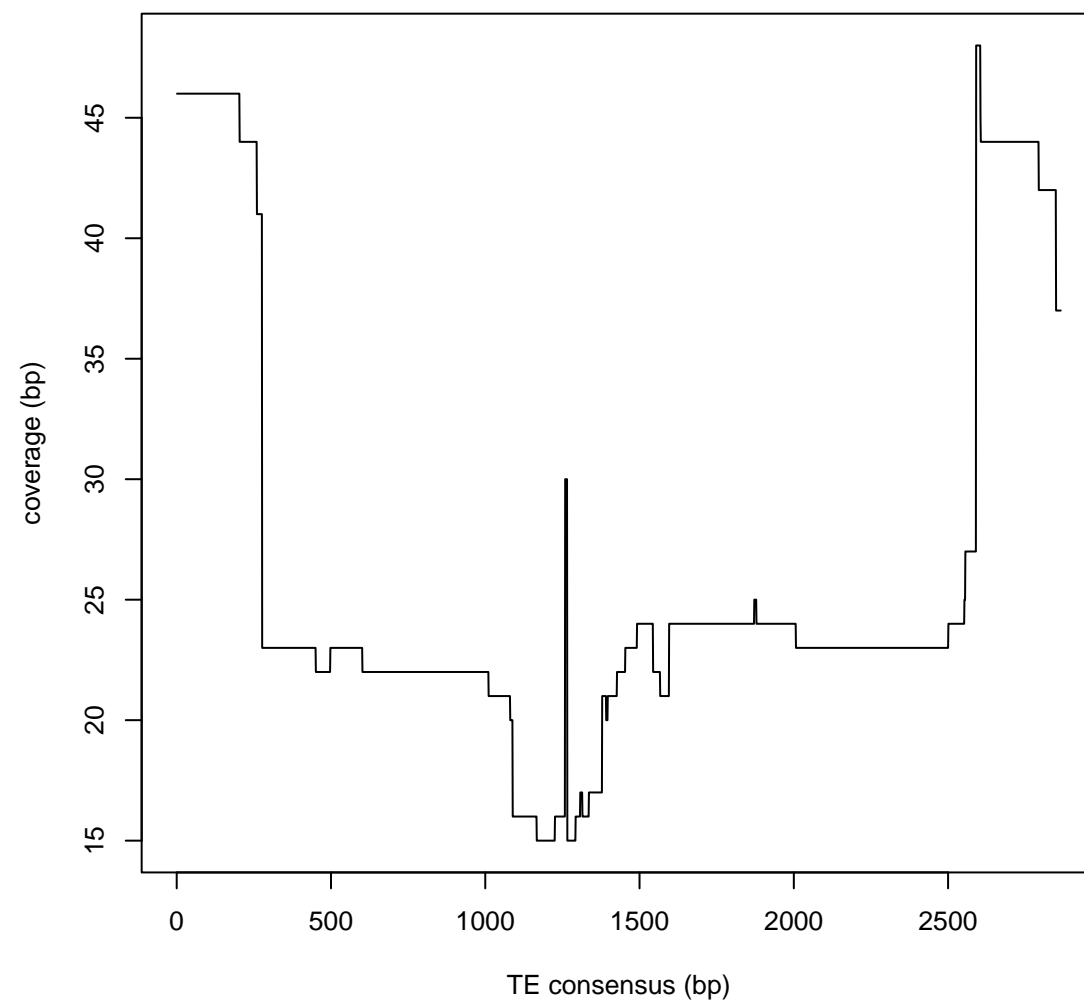


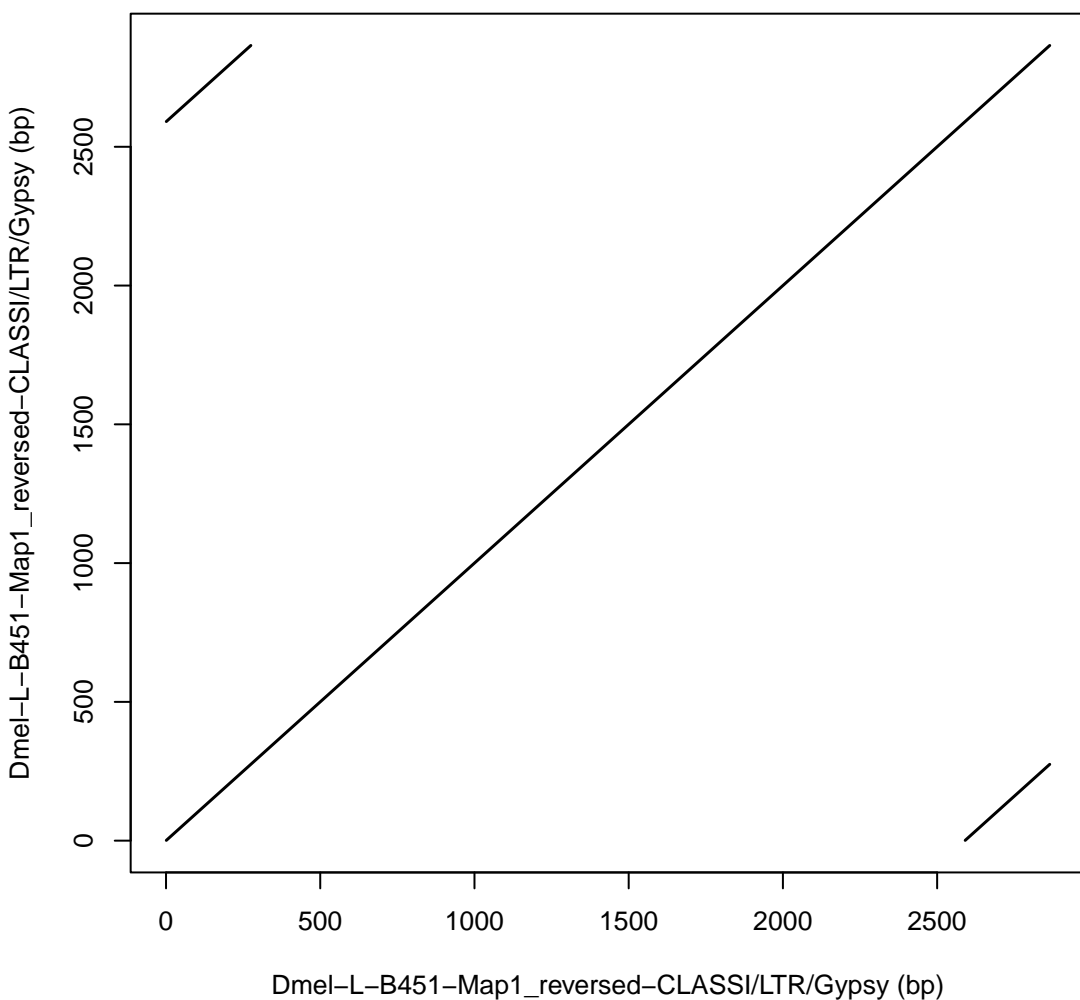
TE: Dmel-L-B451-Map1\_reversed-CLASSI/LTR/Gypsy  
consensus size: 2865bp; fragments: 102; full length: 1 (>=2578.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

