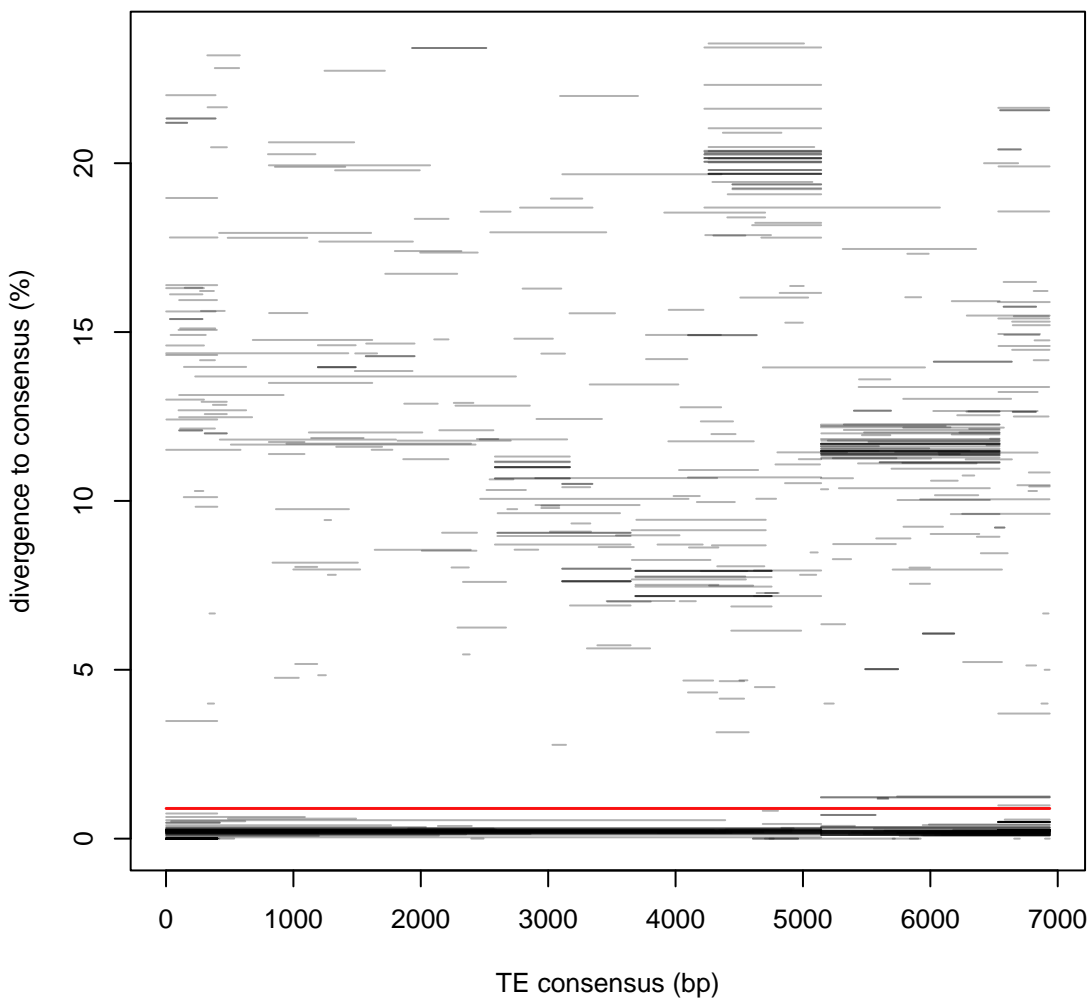
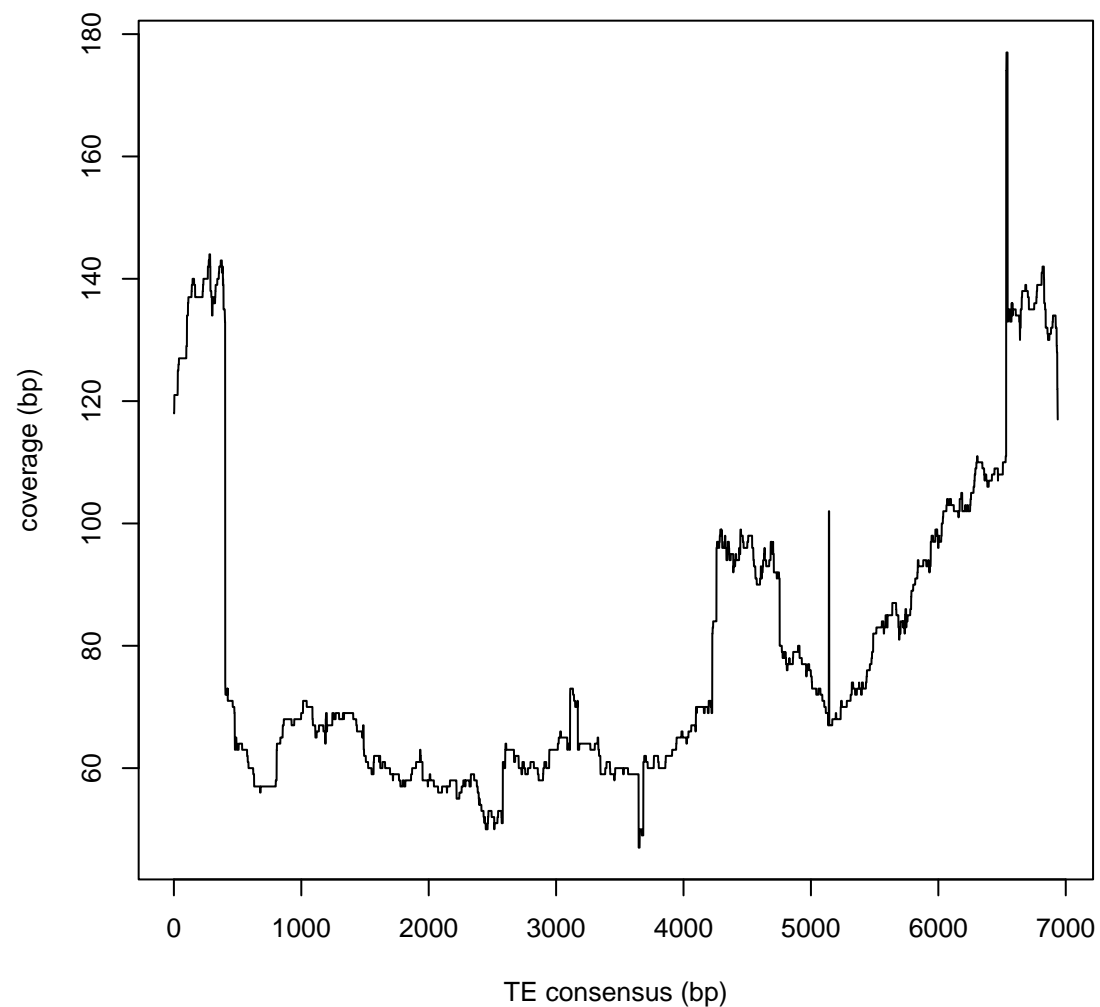


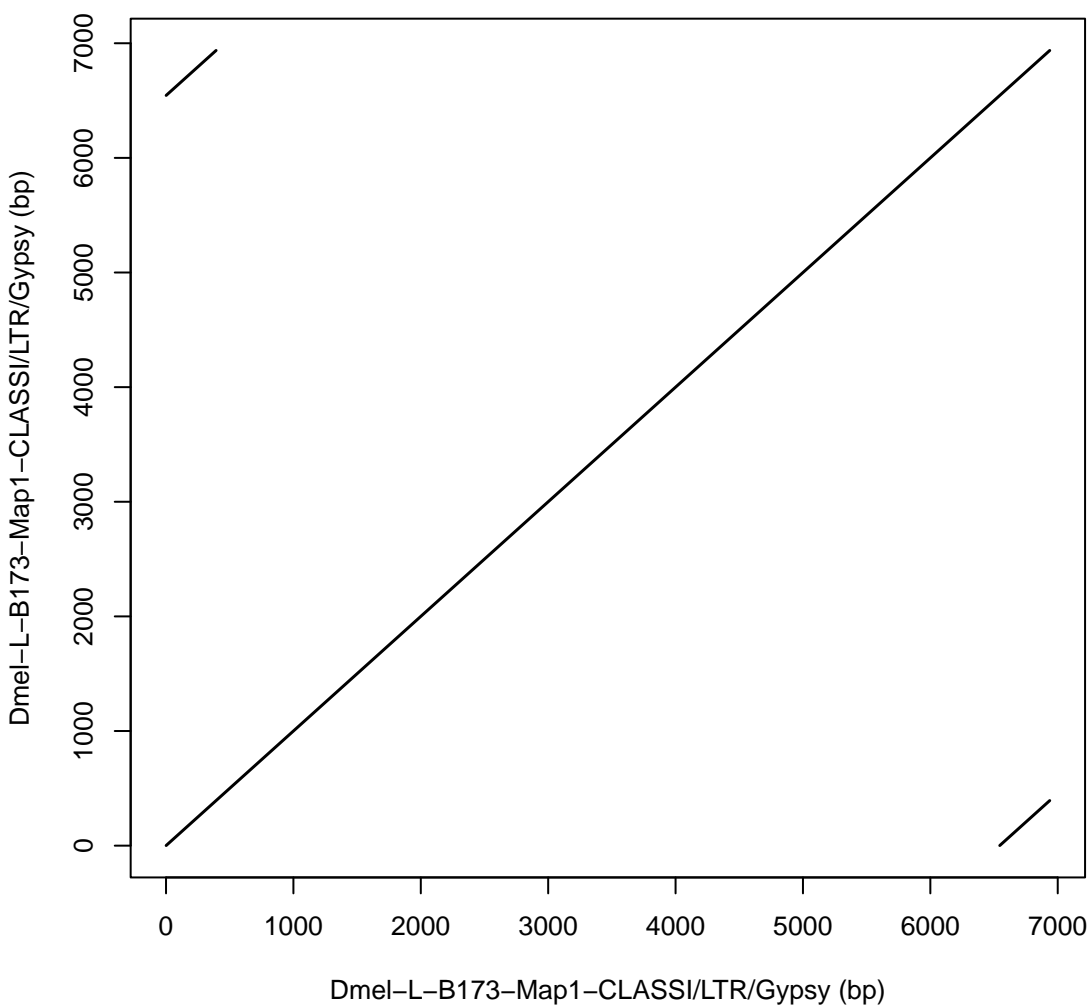
TE: Dmel-L-B173-Map1-CLASSI/LTR/Gypsy  
consensus size: 6937bp; fragments: 638; full length: 1 (>=6243.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

