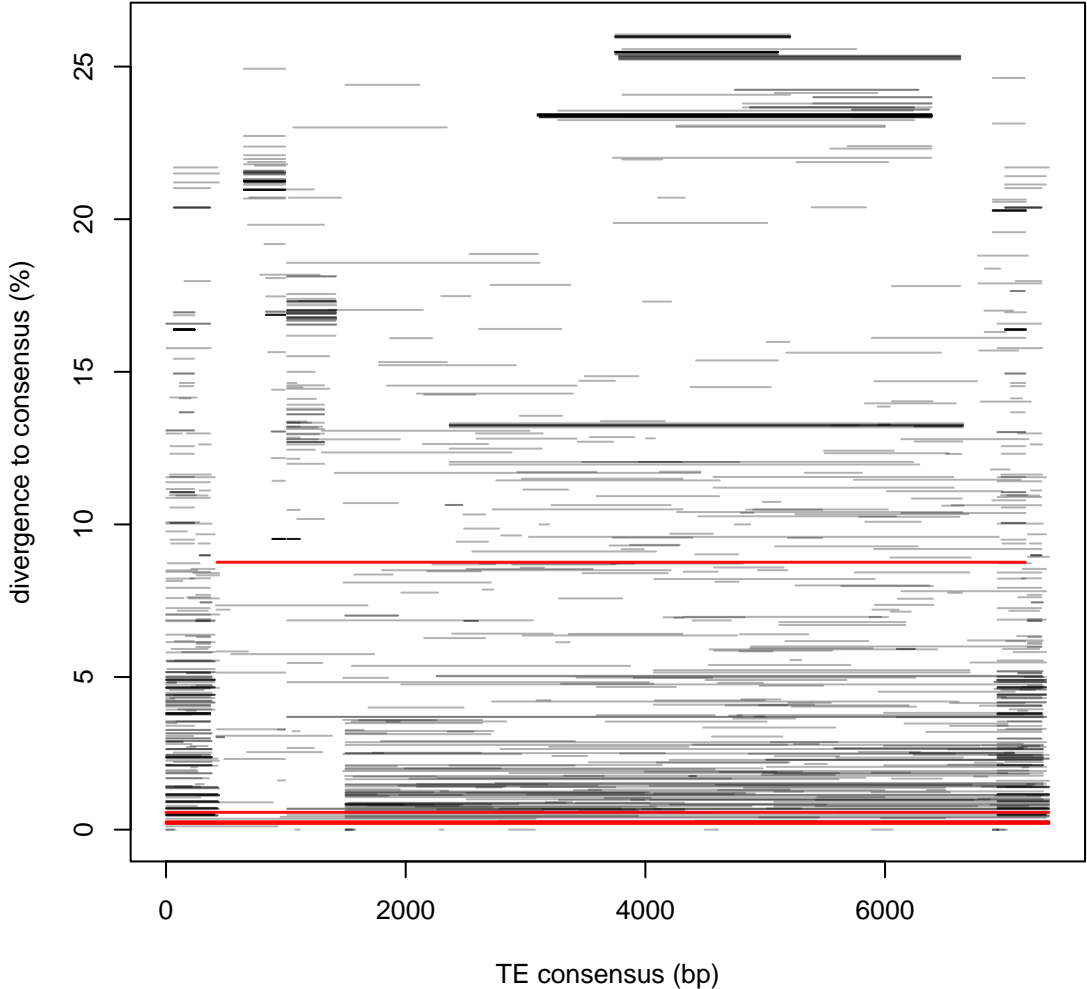
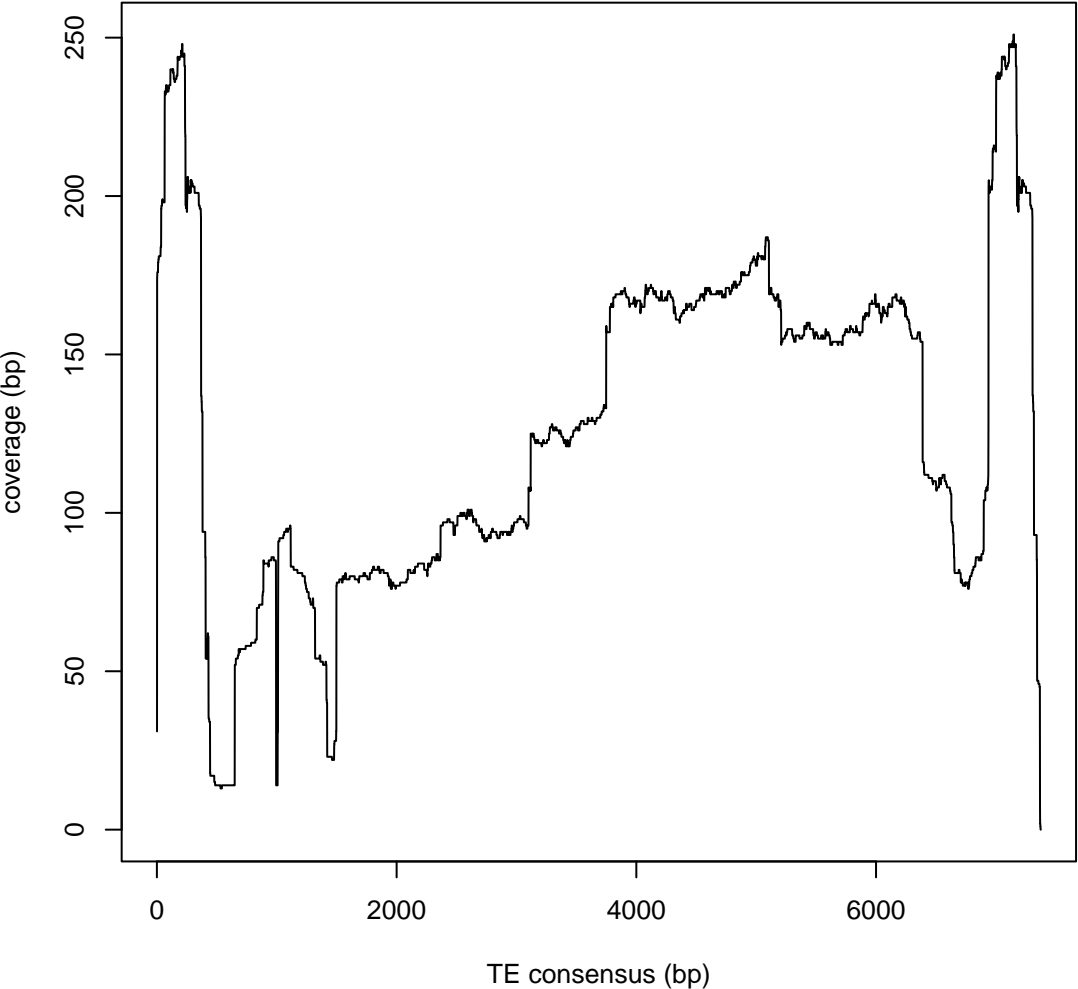


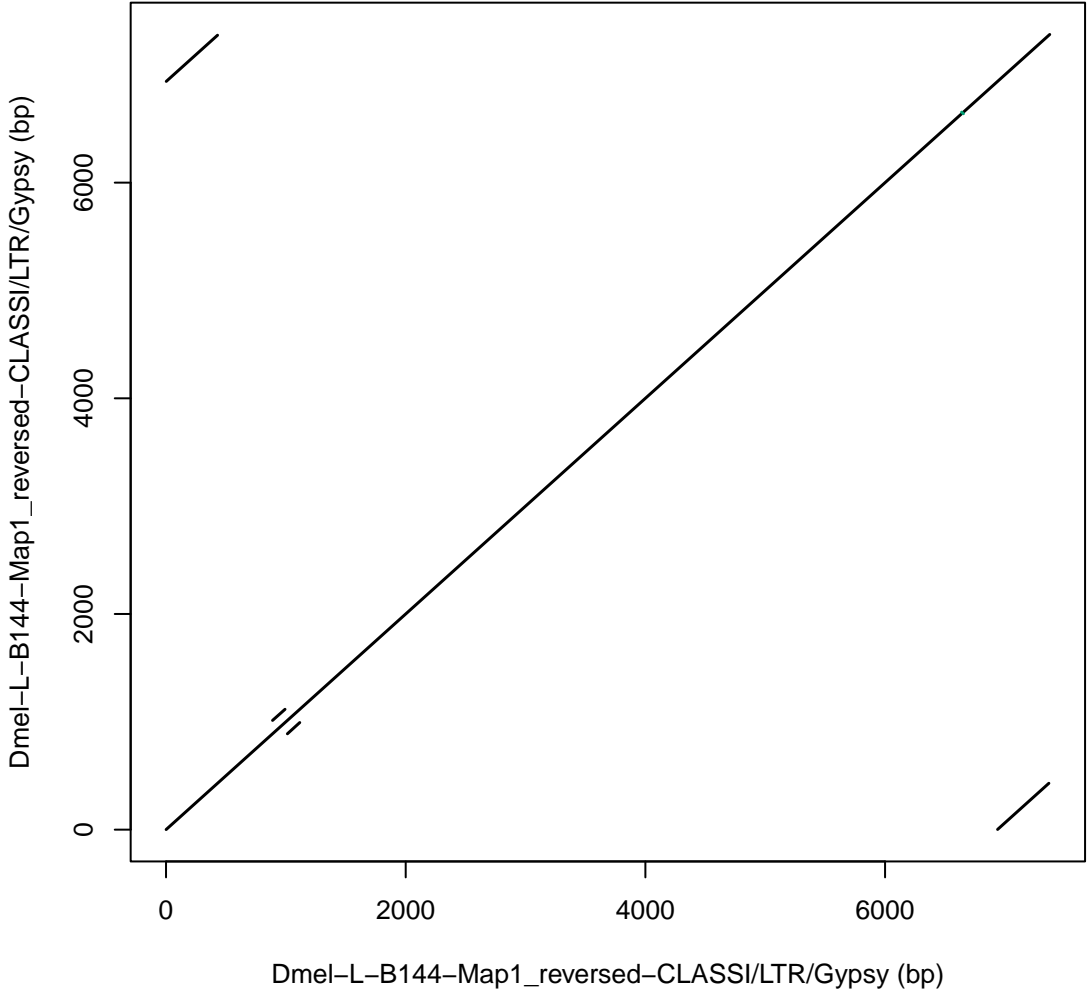
TE: Dmel-L-B144-Map1\_reversed-CLASSI/LTR/Gypsy  
consensus size: 7374bp; fragments: 1128; full length: 4 (>=6636.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

