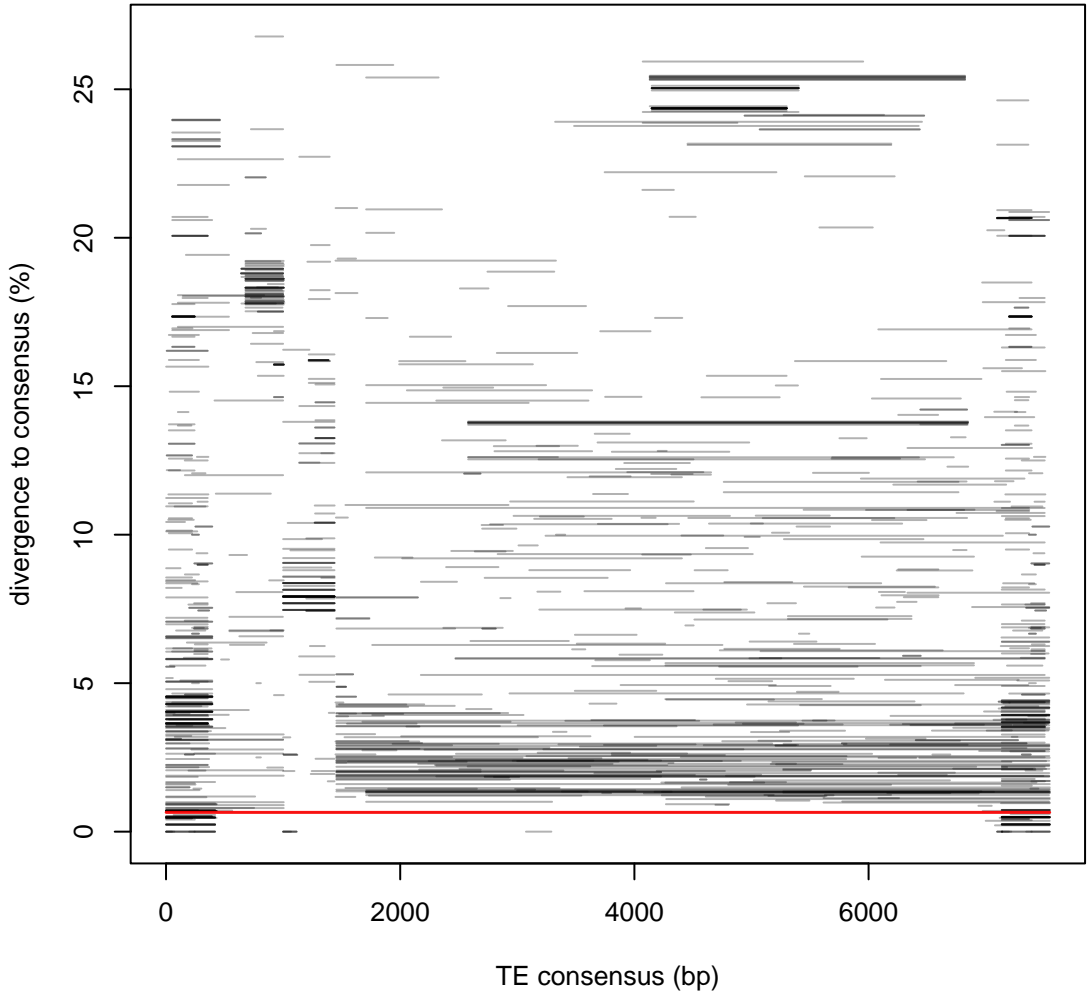
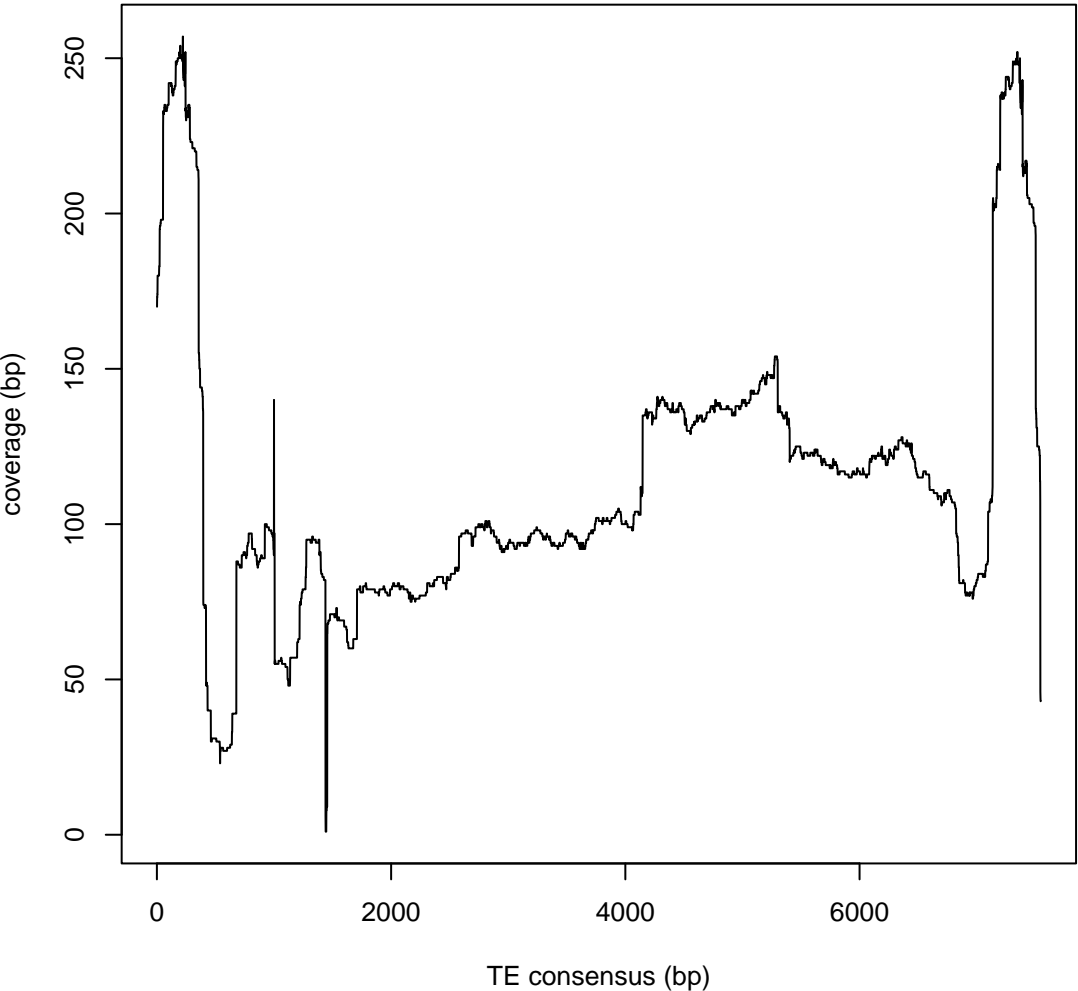


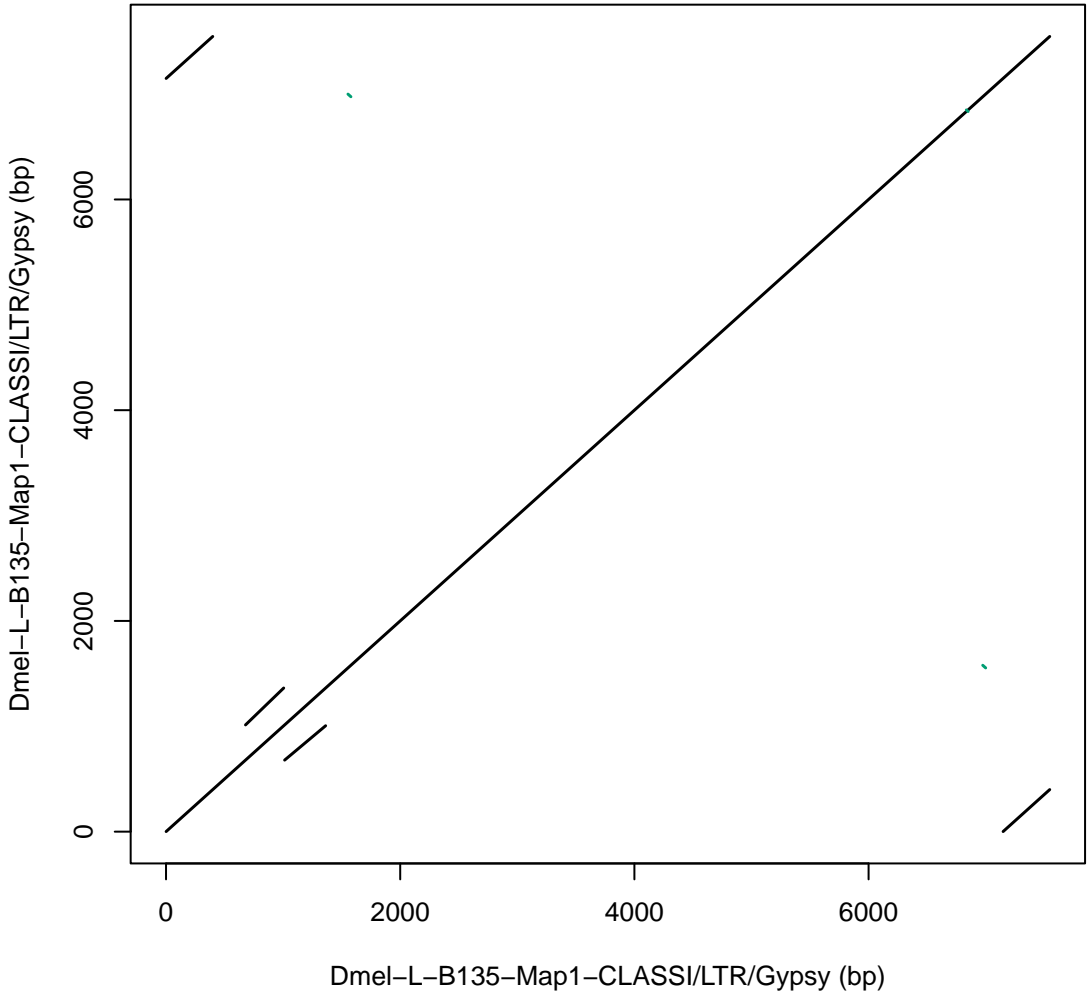
TE: Dmel-L-B135-Map1-CLASSI/LTR/Gypsy
consensus size: 7547bp; fragments: 1175; full length: 1 (>=6792.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

