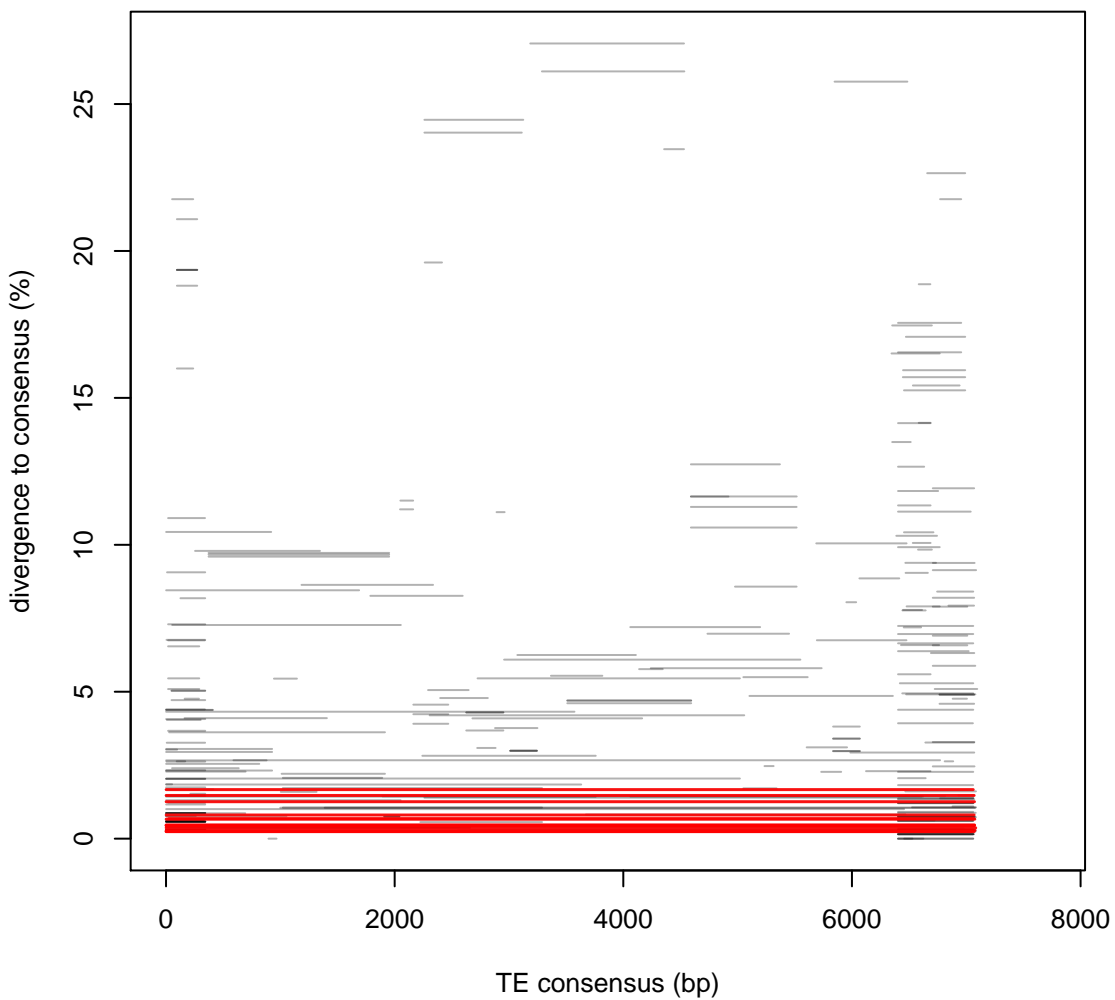
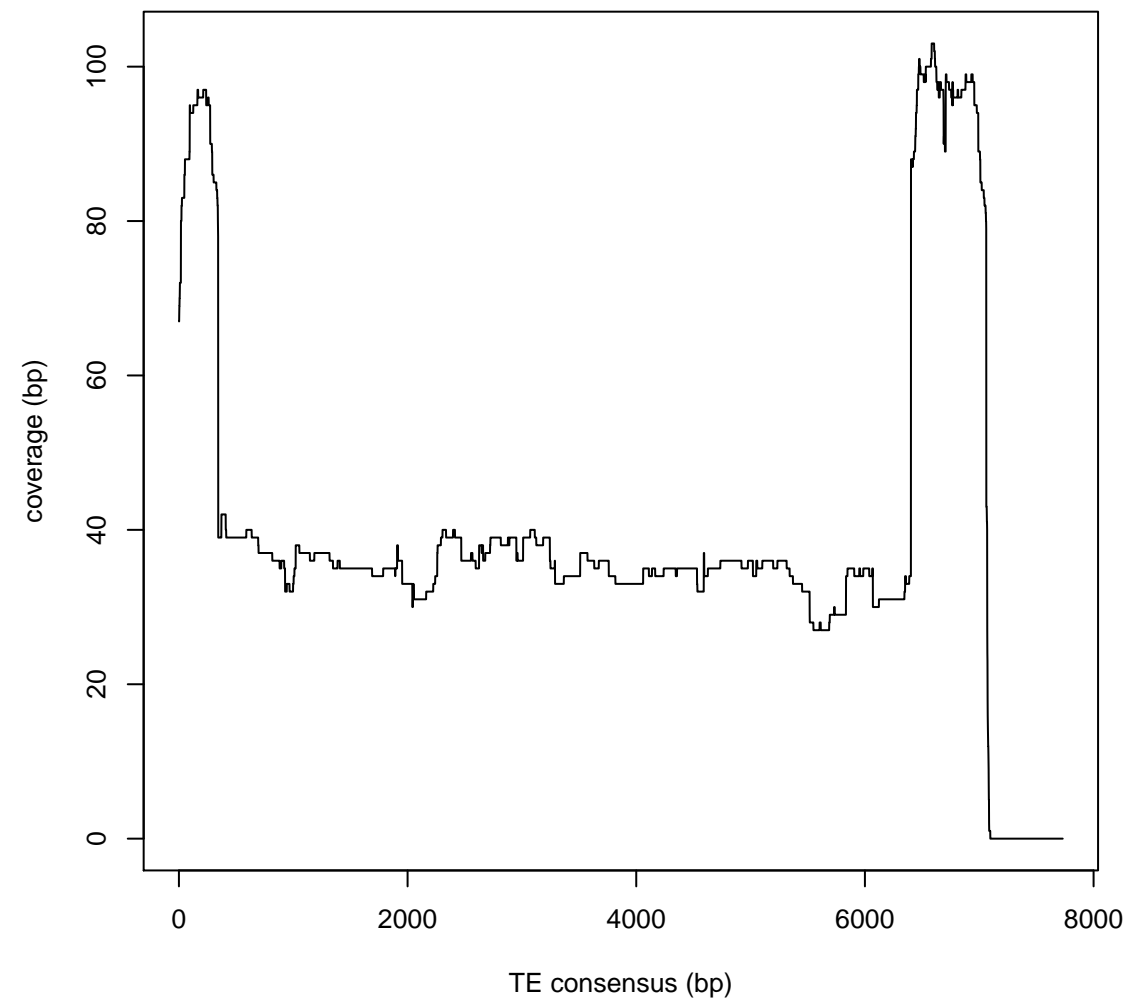


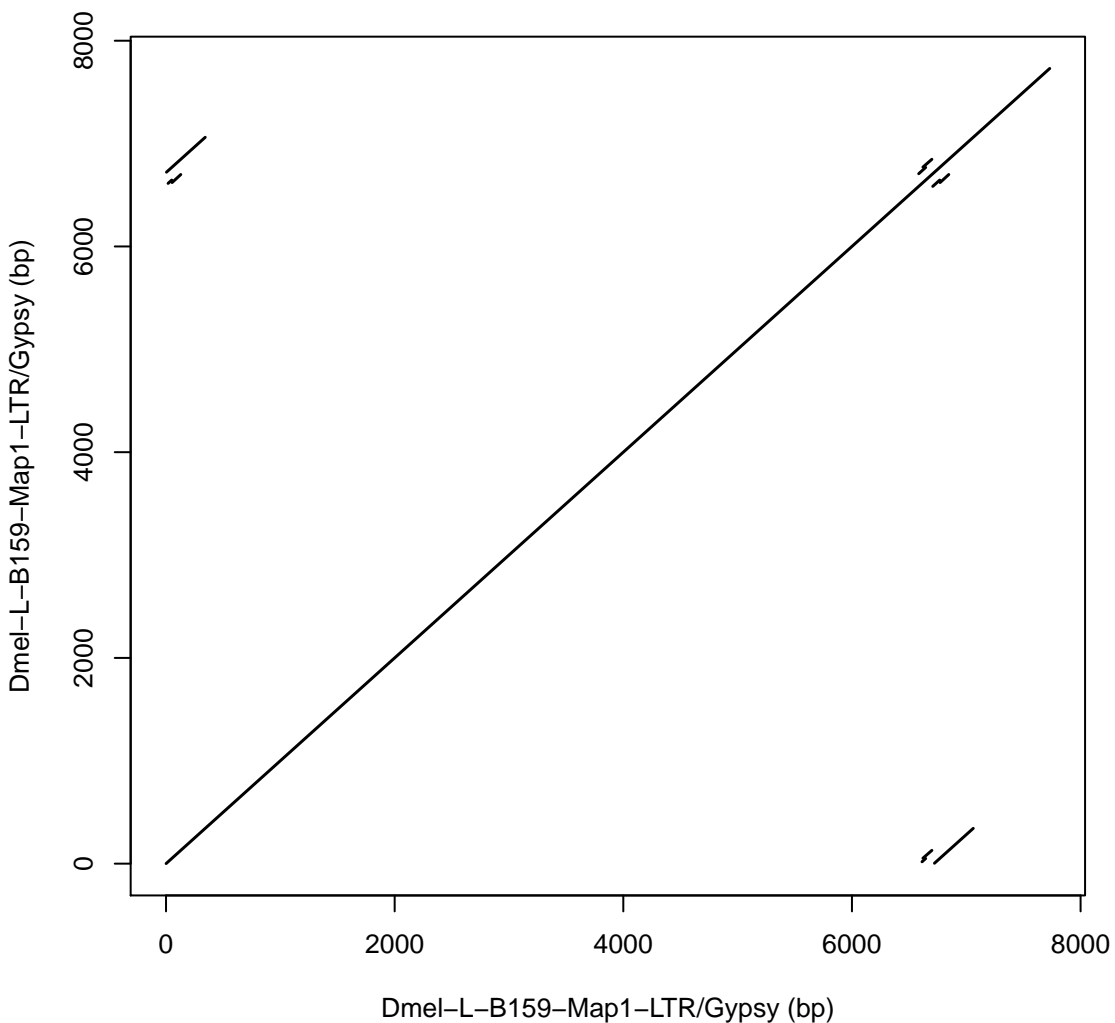
TE: Dmel-L-B159-Map1-LTR/Gypsy
consensus size: 7730bp; fragments: 295; full length: 13 (>=6957bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

