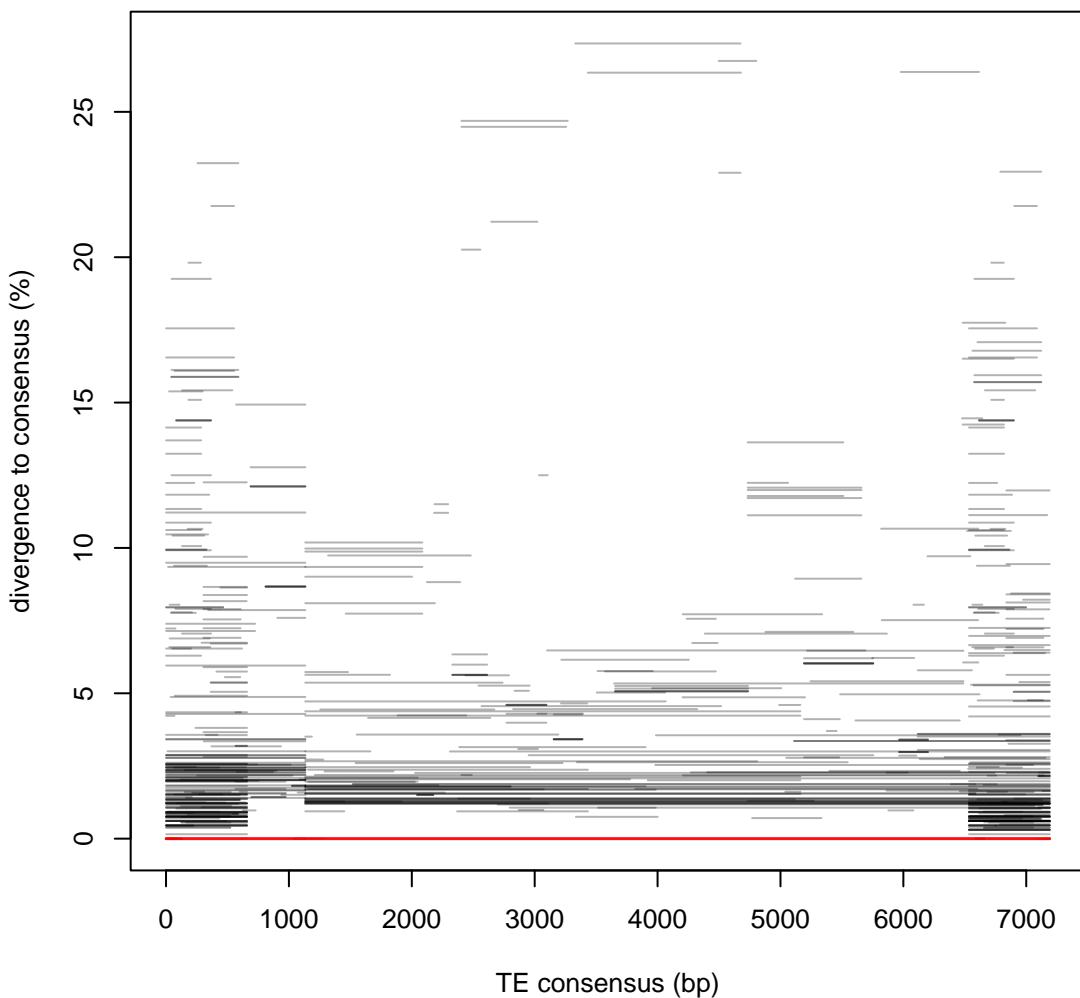
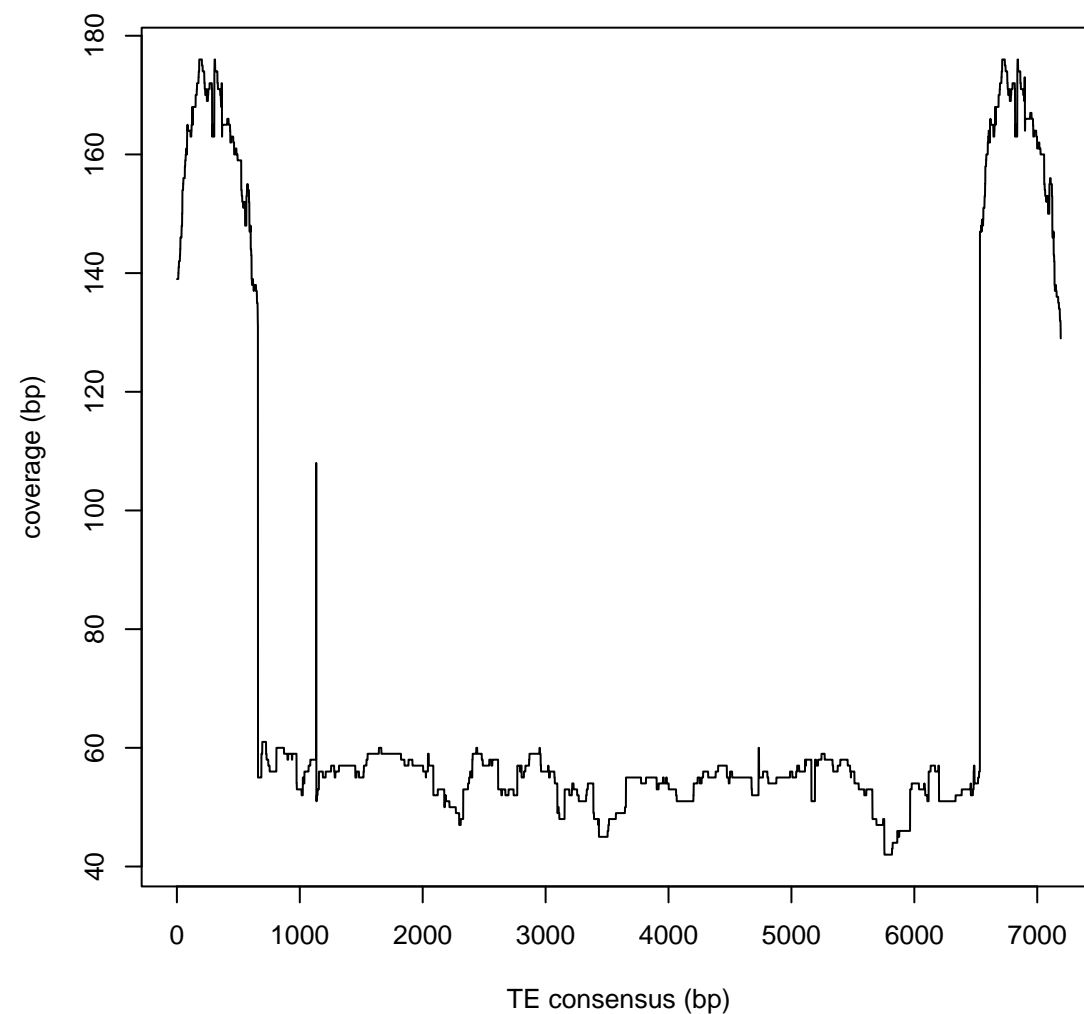


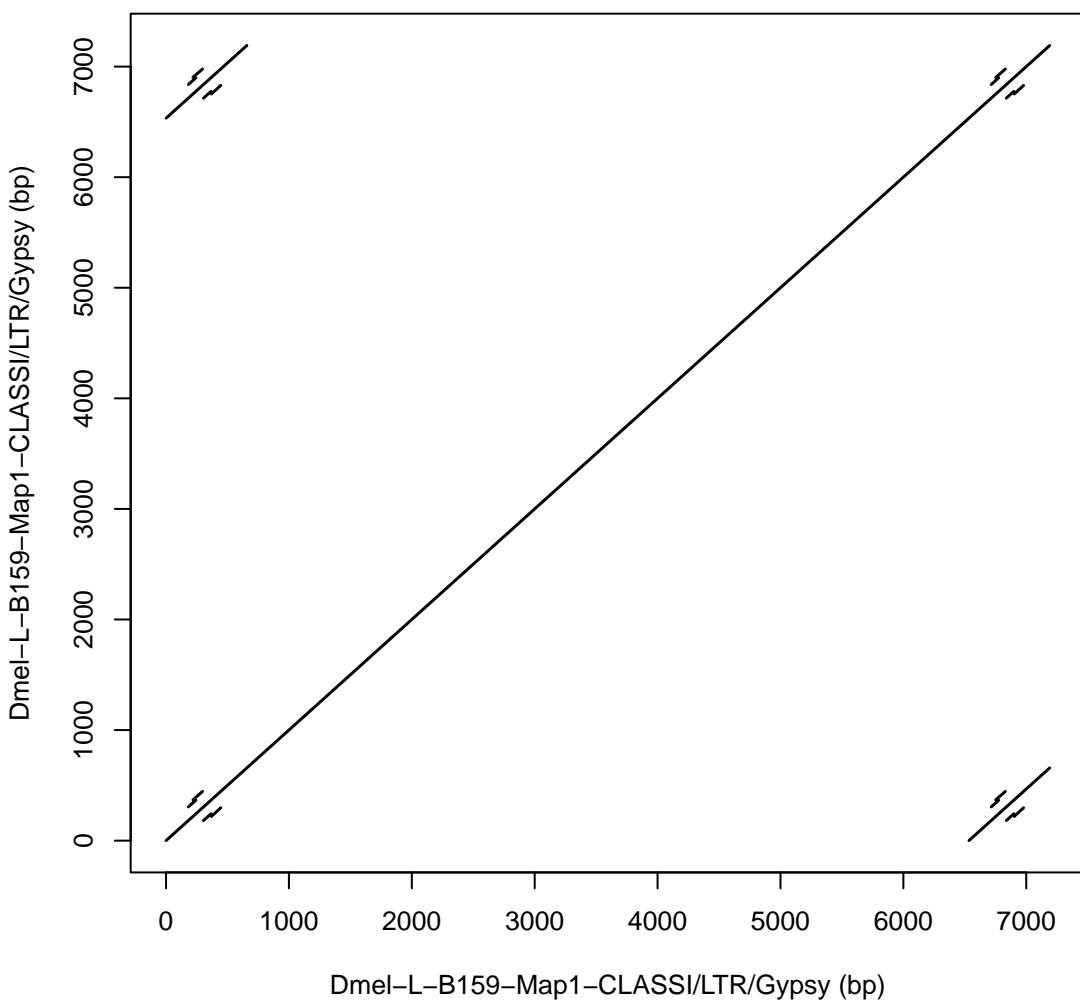
TE: Dmel-L-B159-Map1-CLASSI/LTR/Gypsy
consensus size: 7191bp; fragments: 625; full length: 1 (>=6471.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

