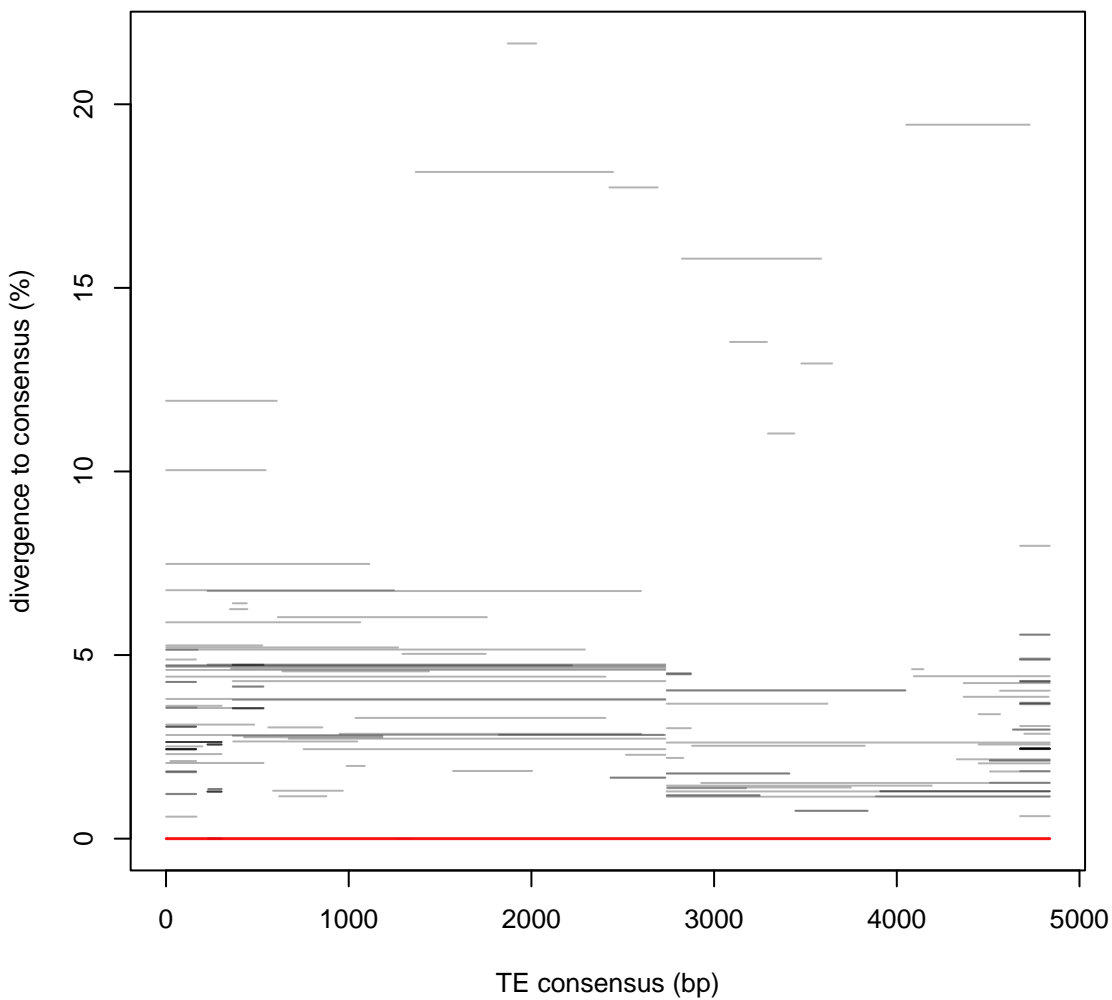
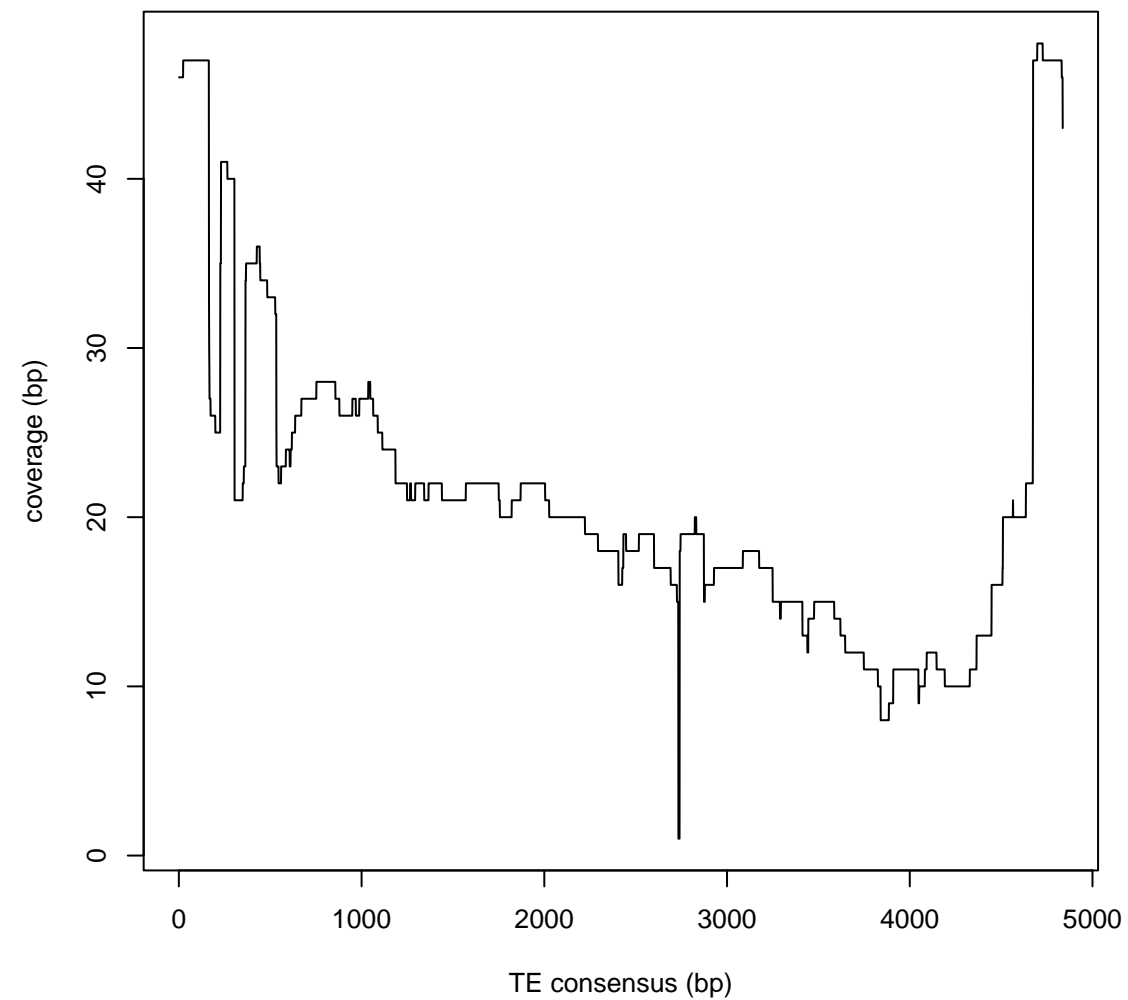


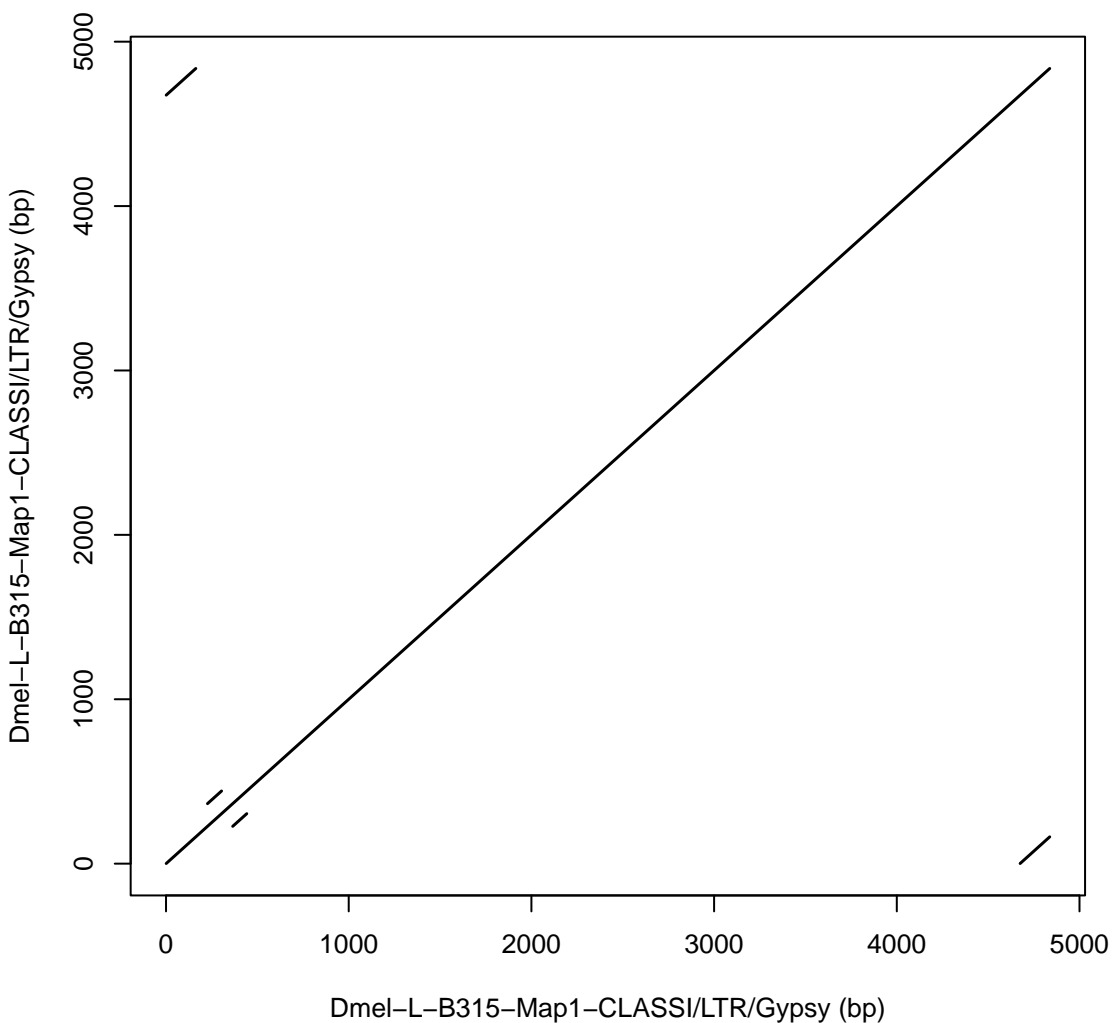
TE: Dmel-L-B315-Map1-CLASSI/LTR/Gypsy
consensus size: 4837bp; fragments: 169; full length: 1 (≥ 4353.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

