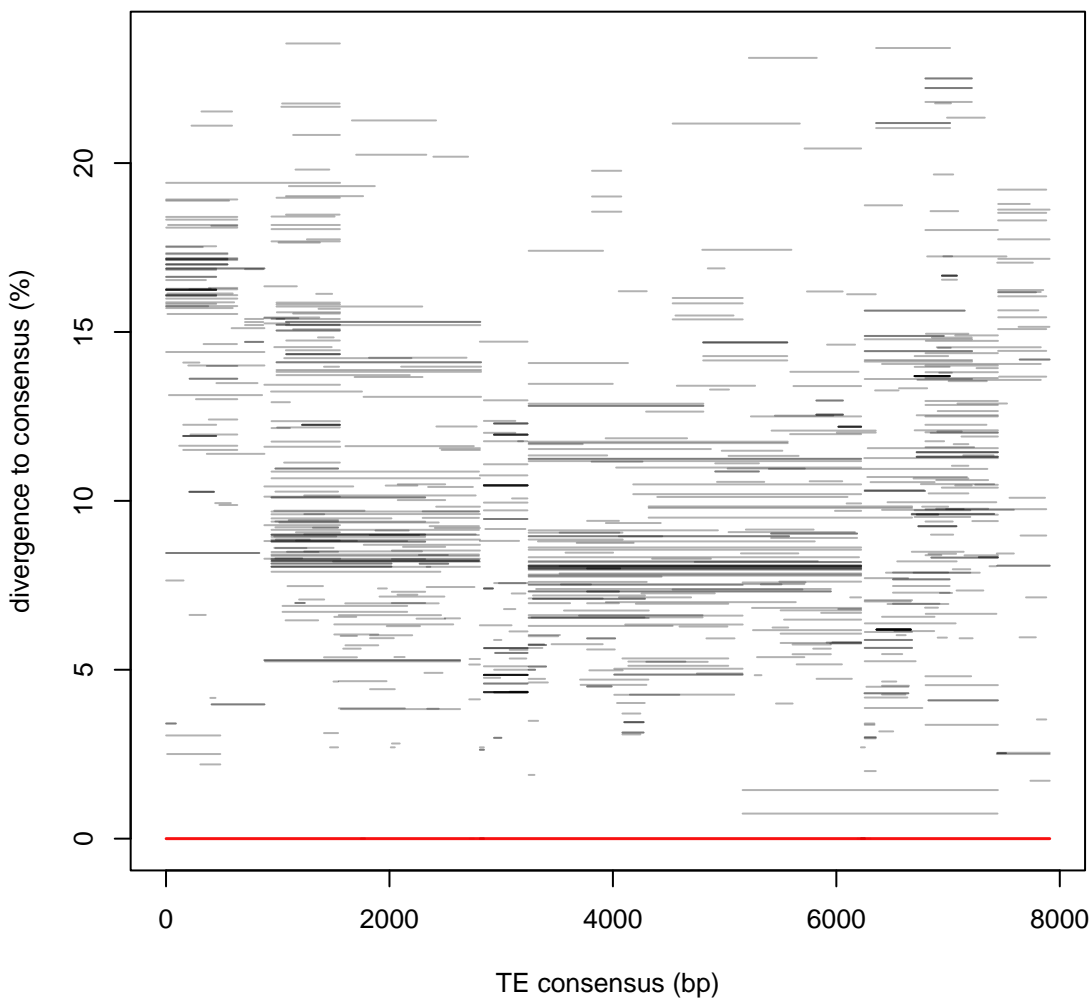
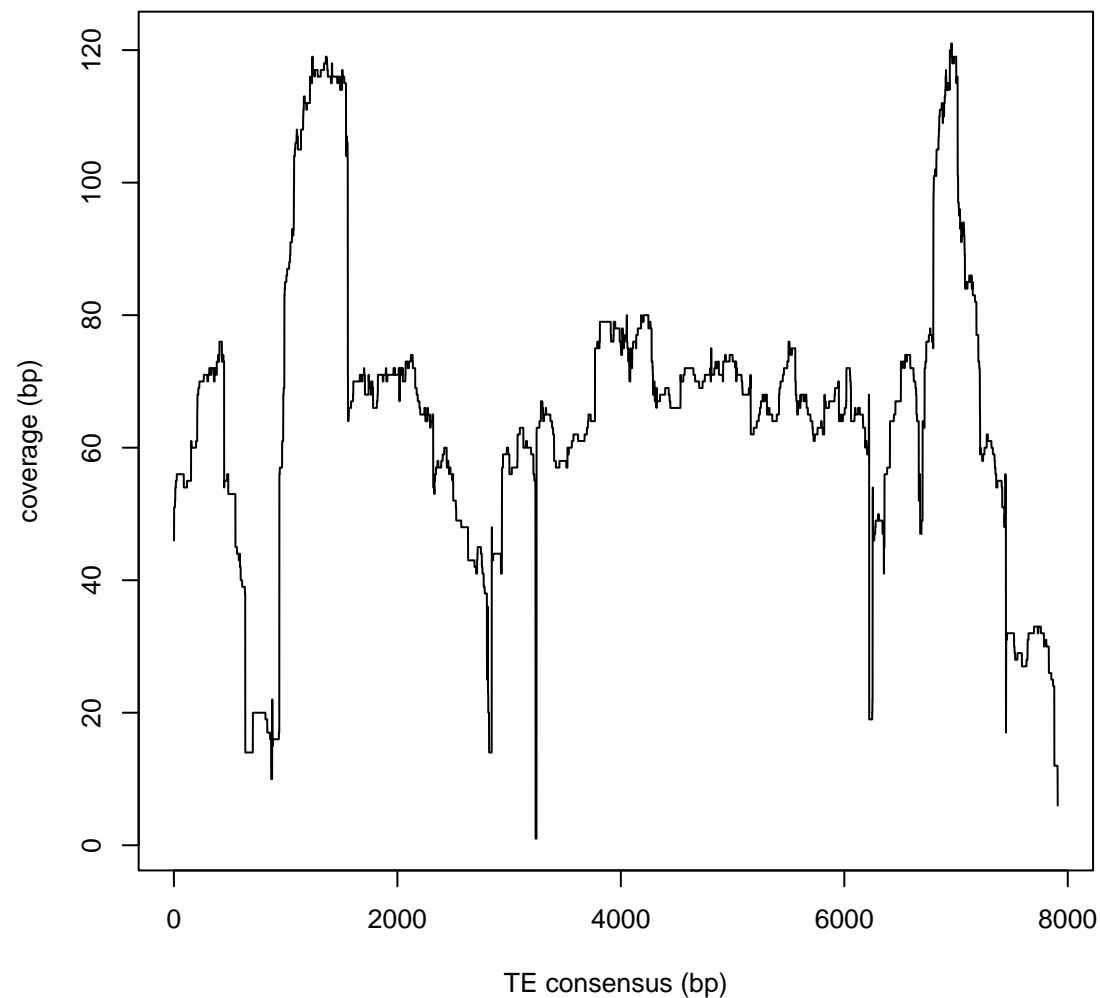


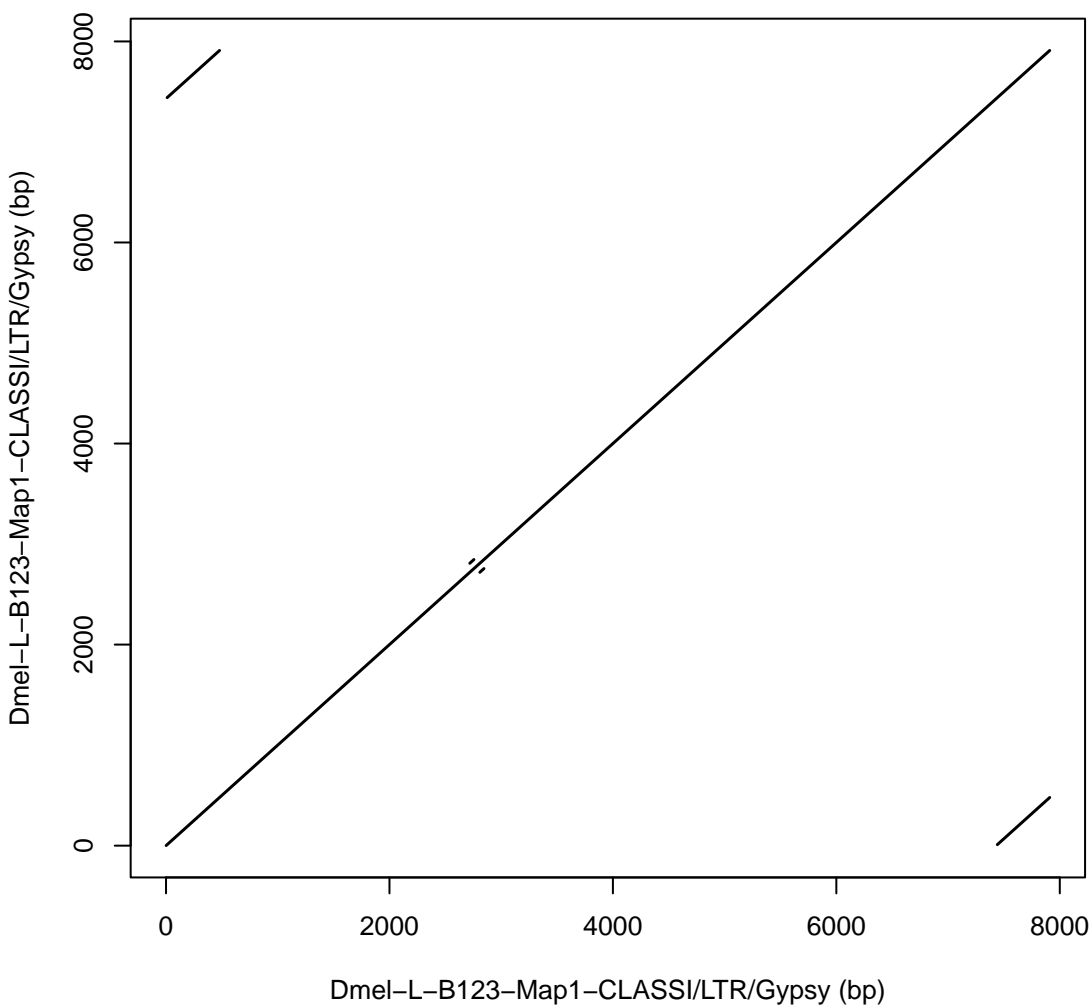
TE: Dmel-L-B123-Map1-CLASSI/LTR/Gypsy
consensus size: 7910bp; fragments: 821; full length: 1 (>=7119bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

