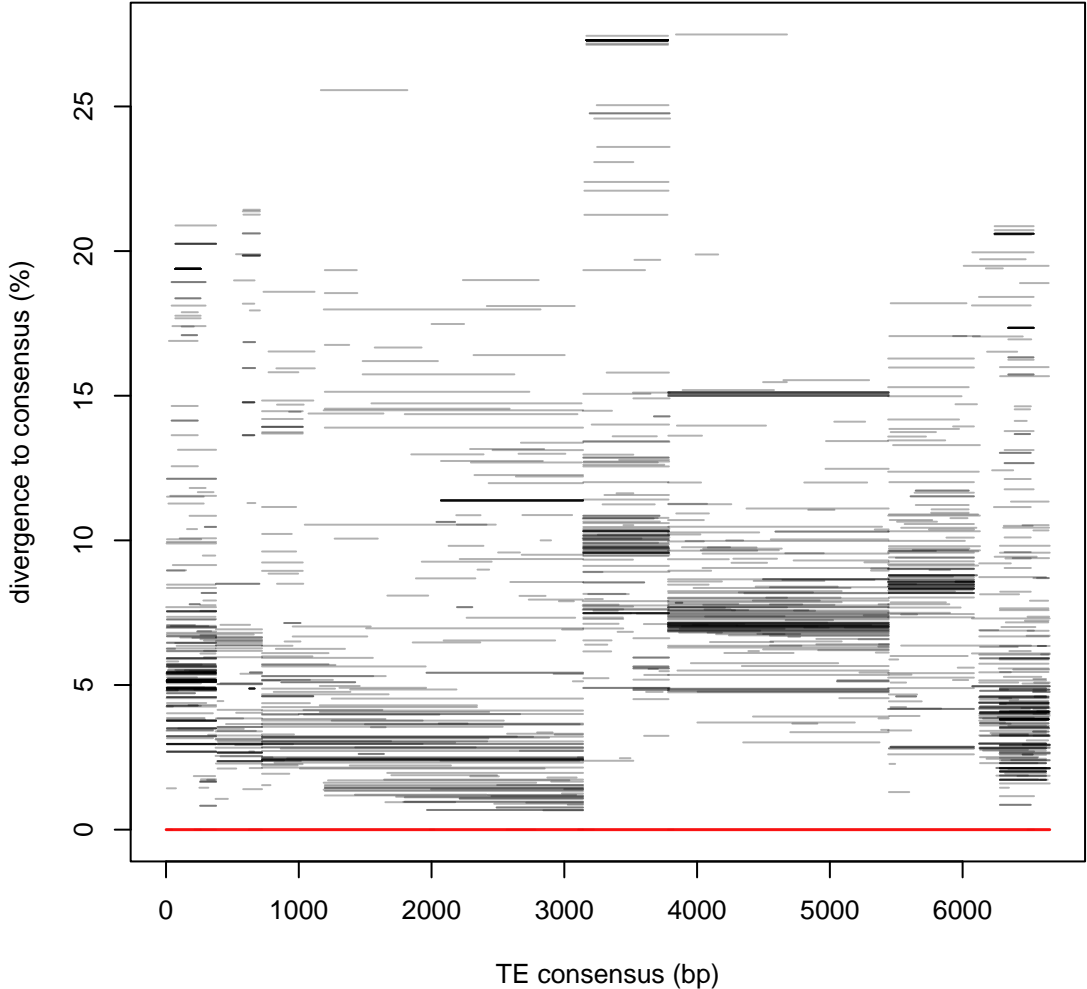
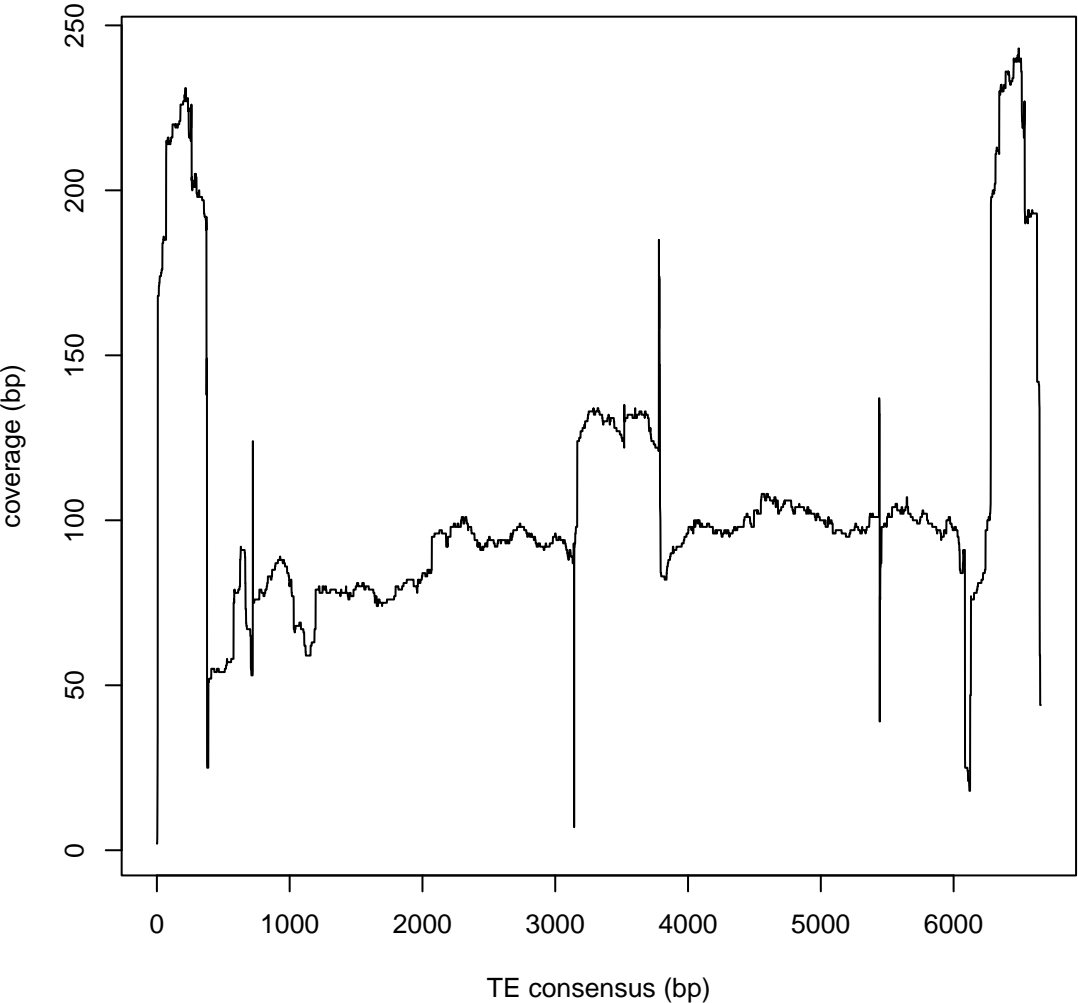


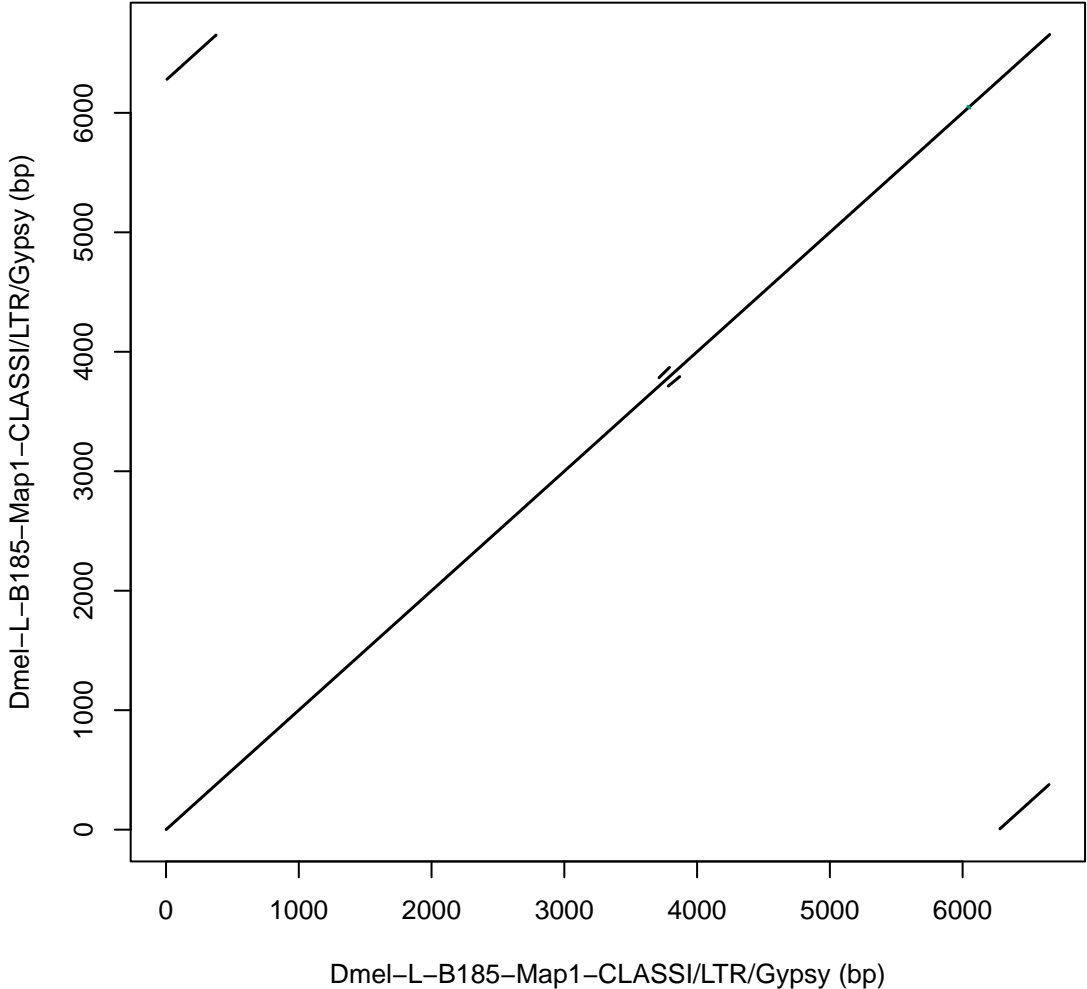
TE: Dmel-L-B185-Map1-CLASSI/LTR/Gypsy
consensus size: 6656bp; fragments: 1342; full length: 1 (>=5990.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

