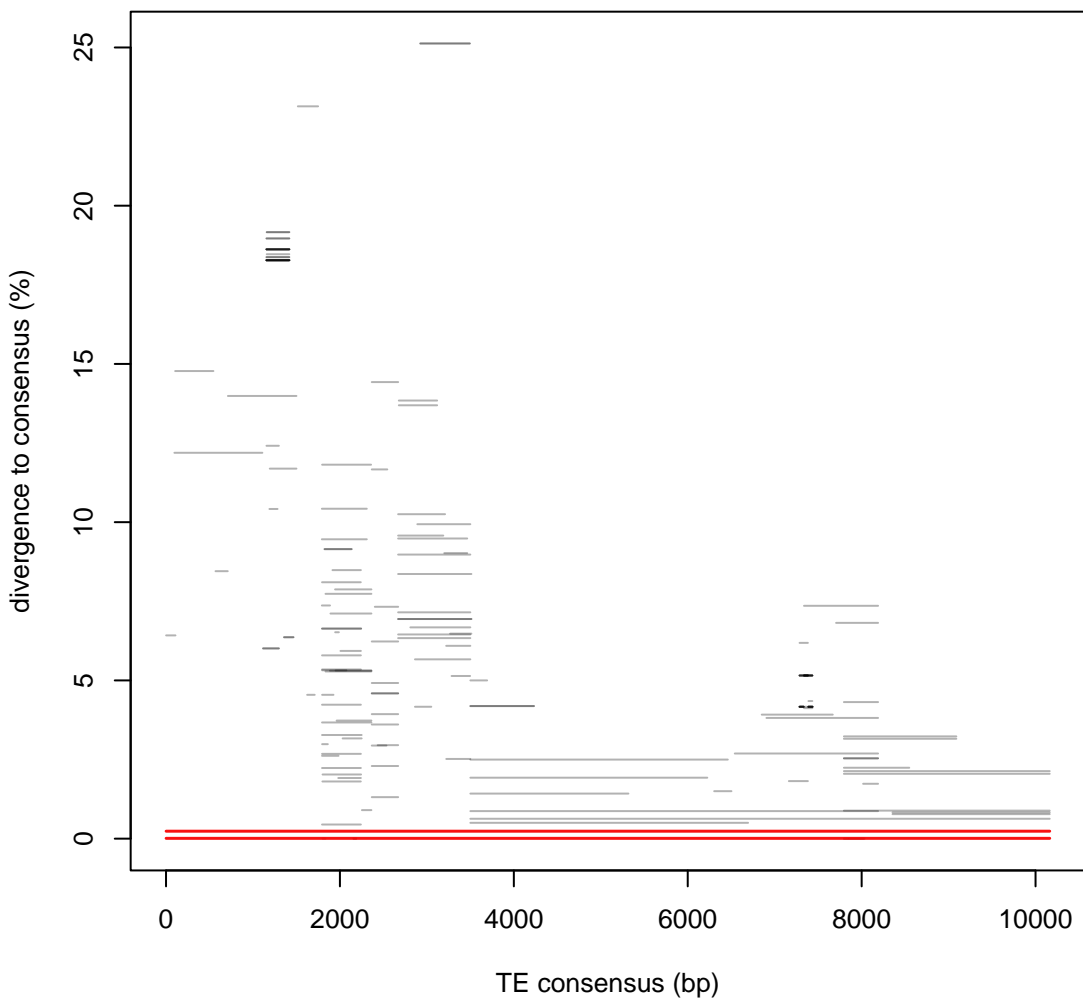
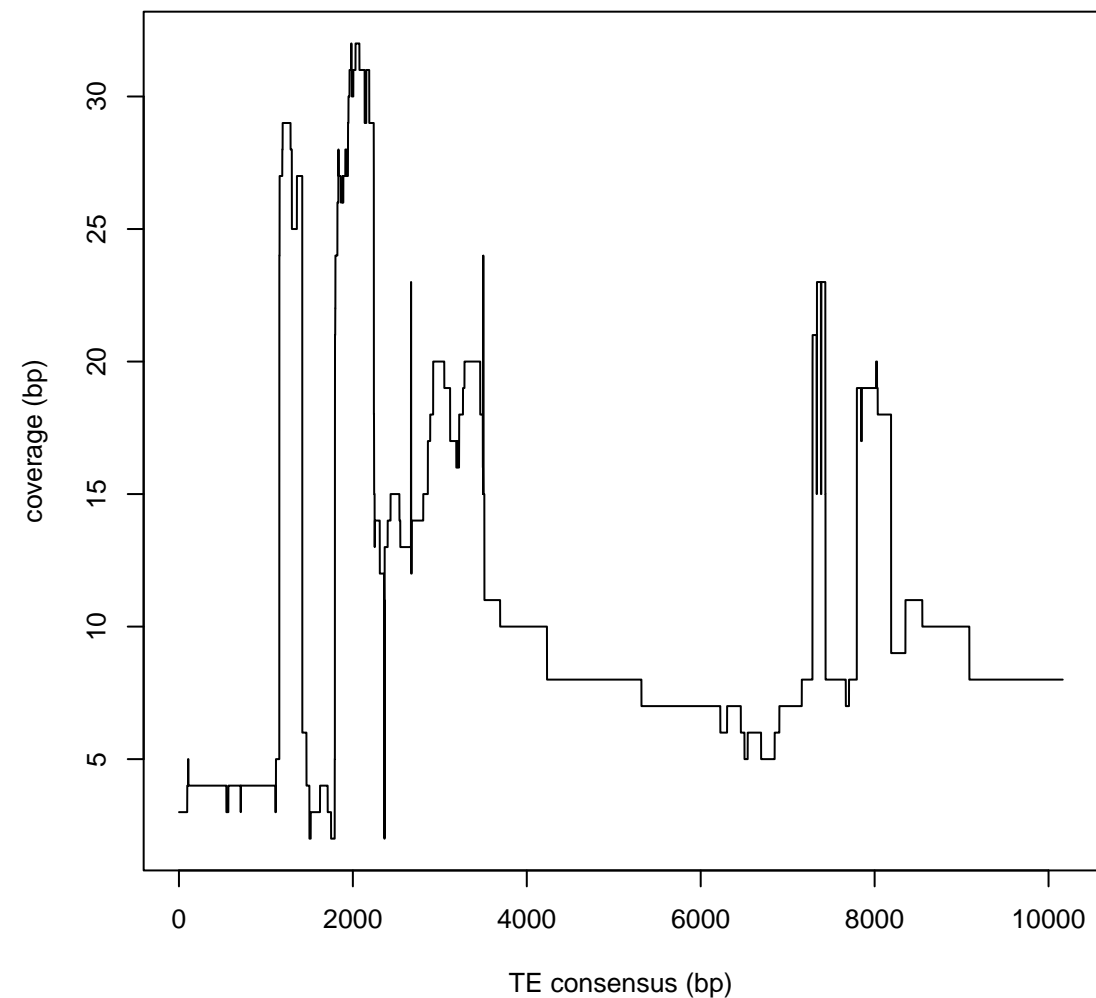


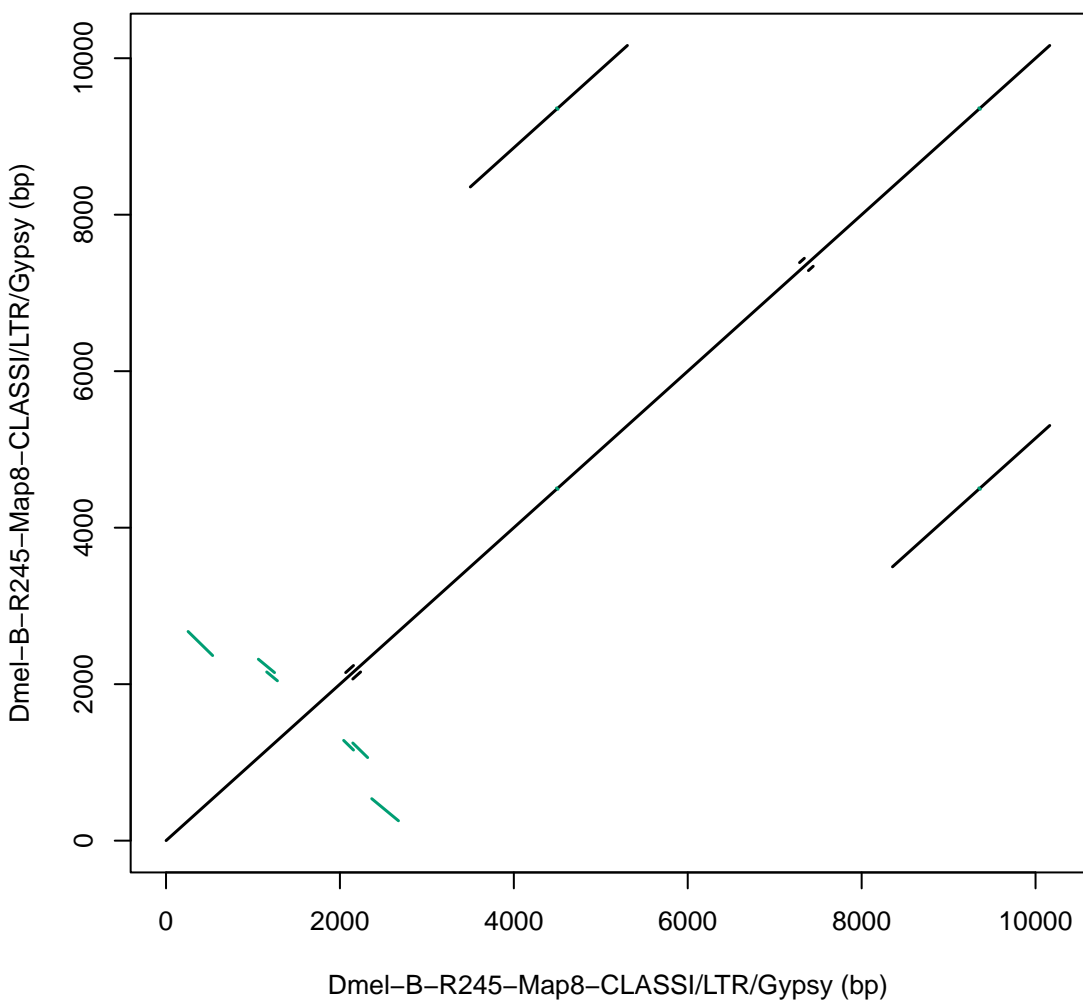
TE: Dmel-B-R245-Map8-CLASSI/LTR/Gypsy
consensus size: 10163bp; fragments: 172; full length: 2 (>=9146.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

