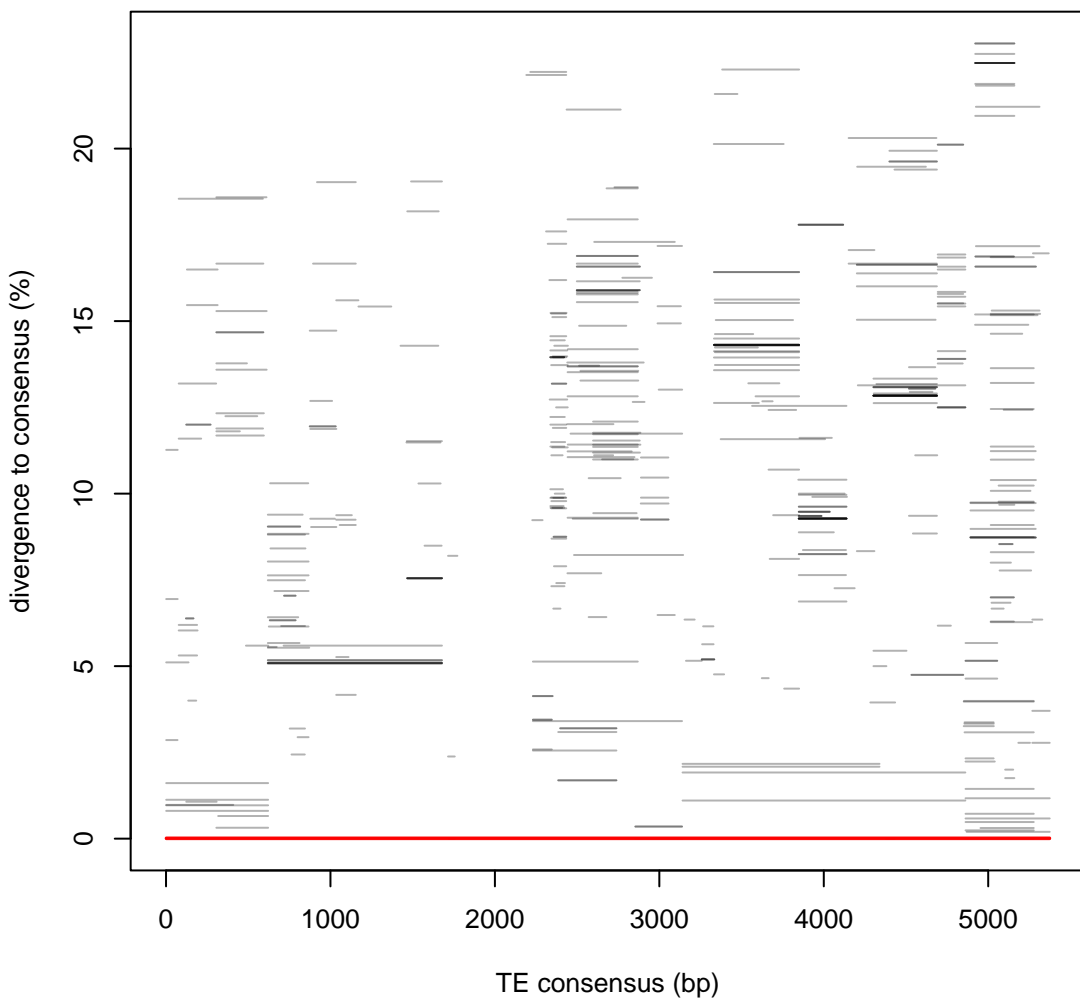
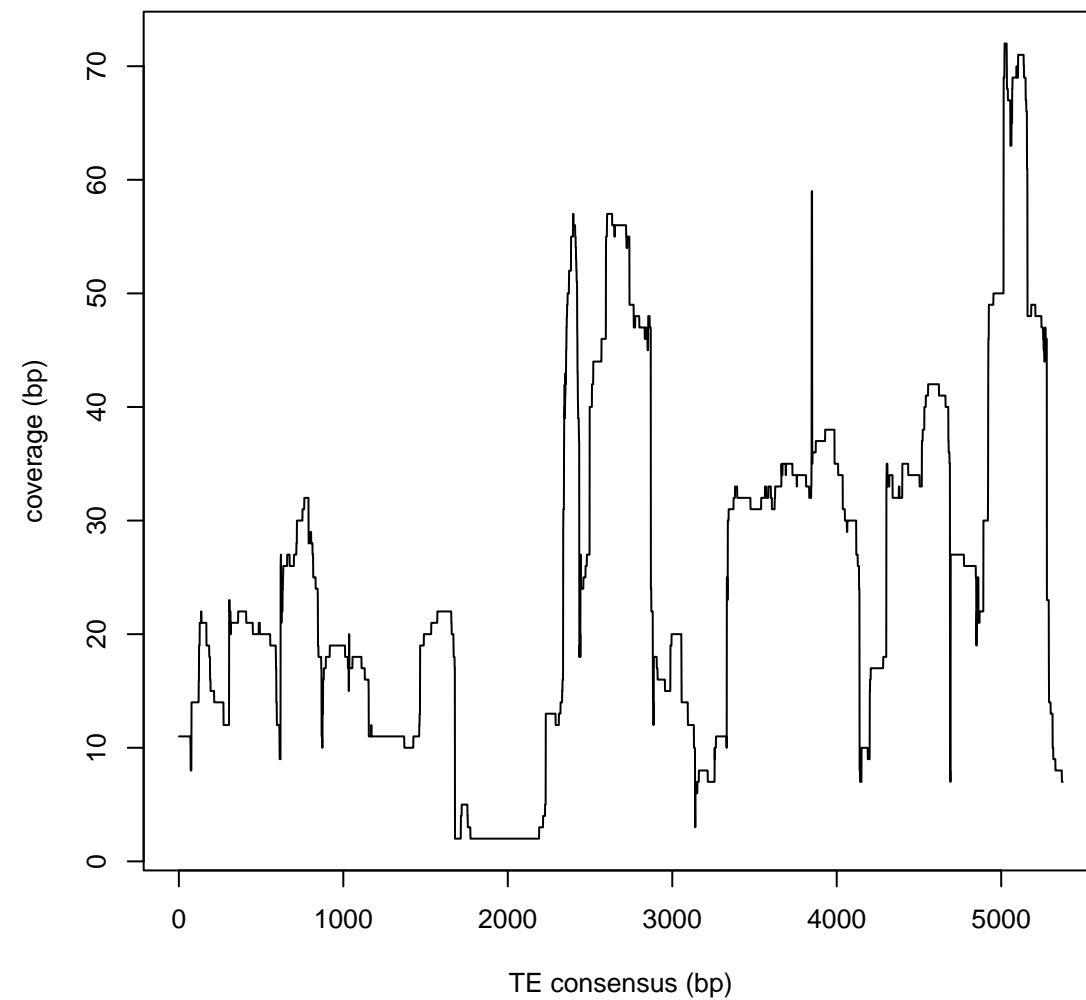


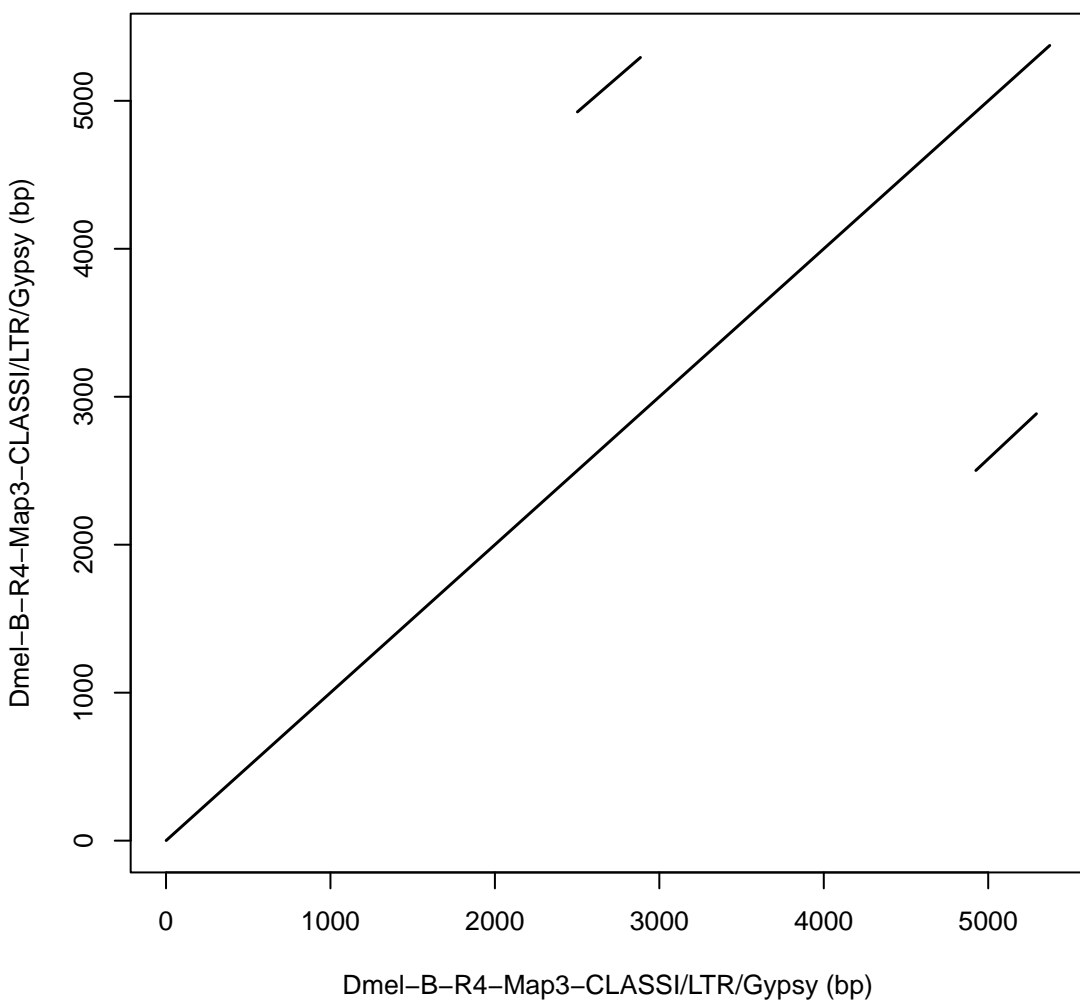
TE: Dmel-B-R4-Map3-CLASSI/LTR/Gypsy
consensus size: 5374bp; fragments: 457; full length: 2 (≥ 4836.6 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

