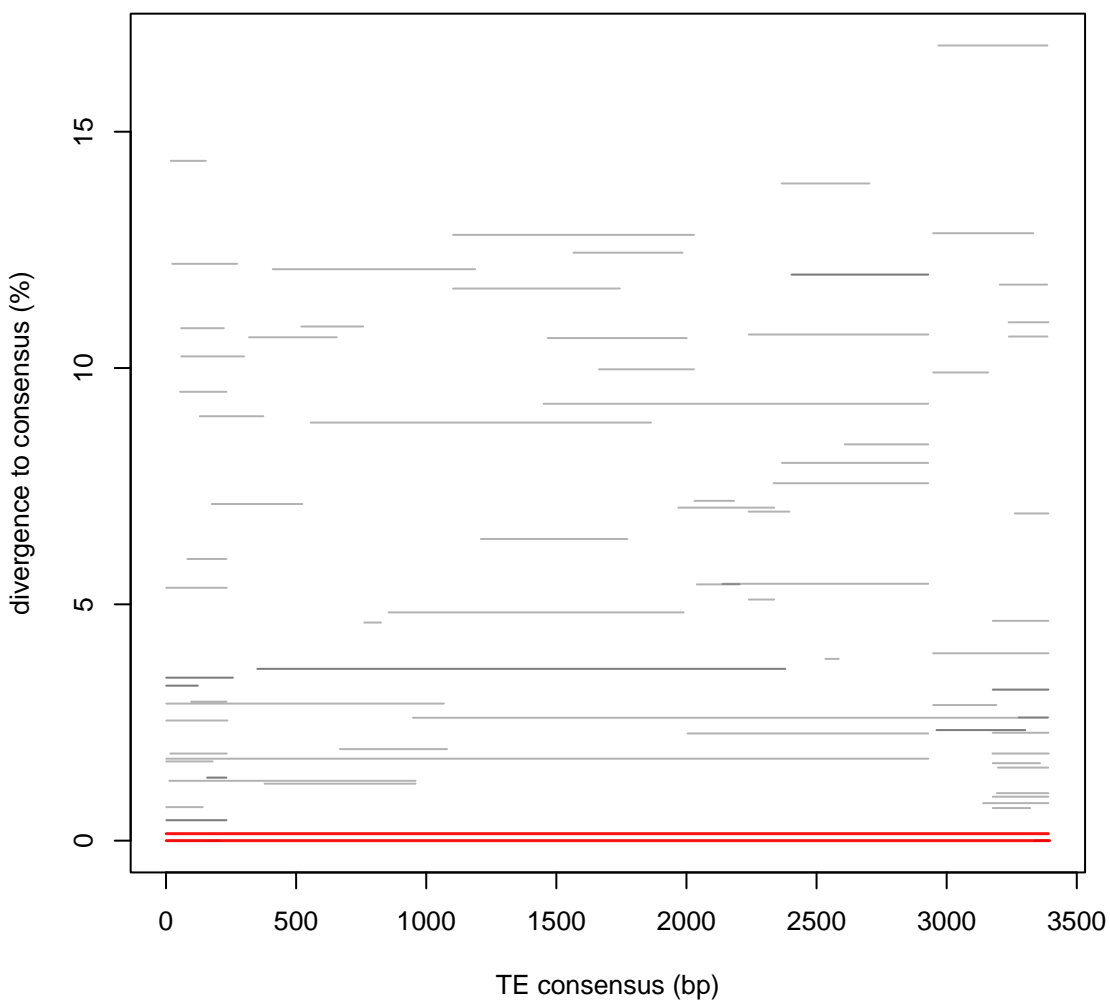
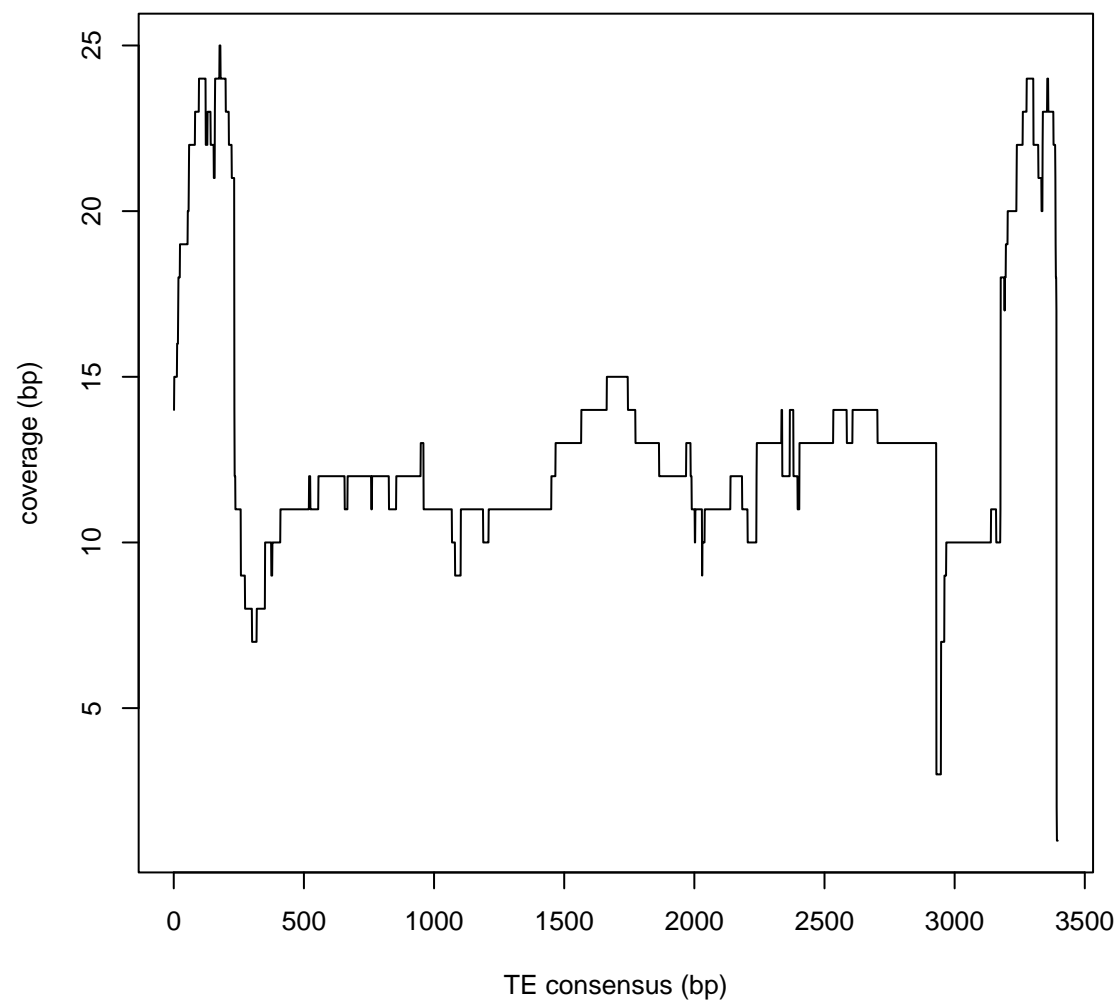


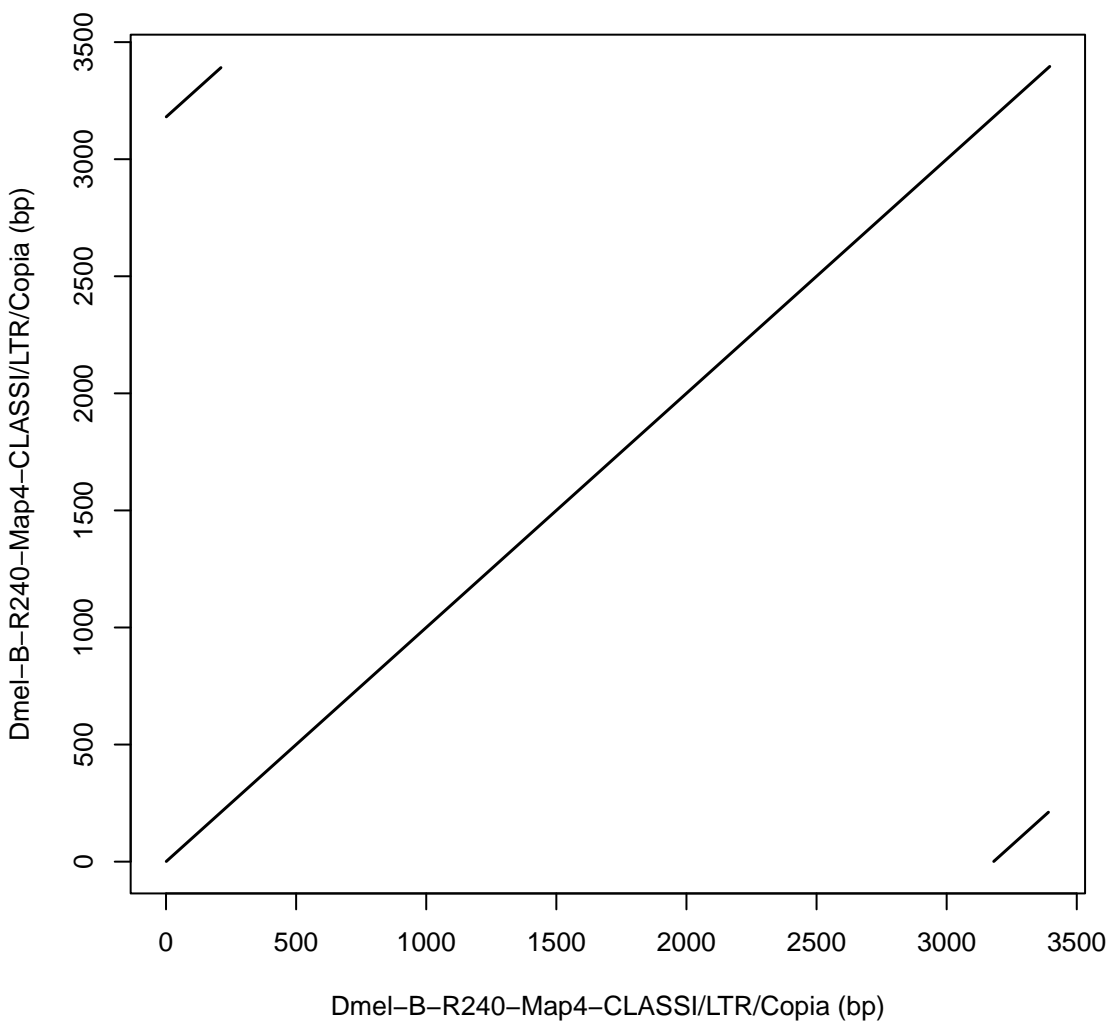
TE: Dmel-B-R240-Map4-CLASSI/LTR/Copia
consensus size: 3396bp; fragments: 89; full length: 2 (≥ 3056.4 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

