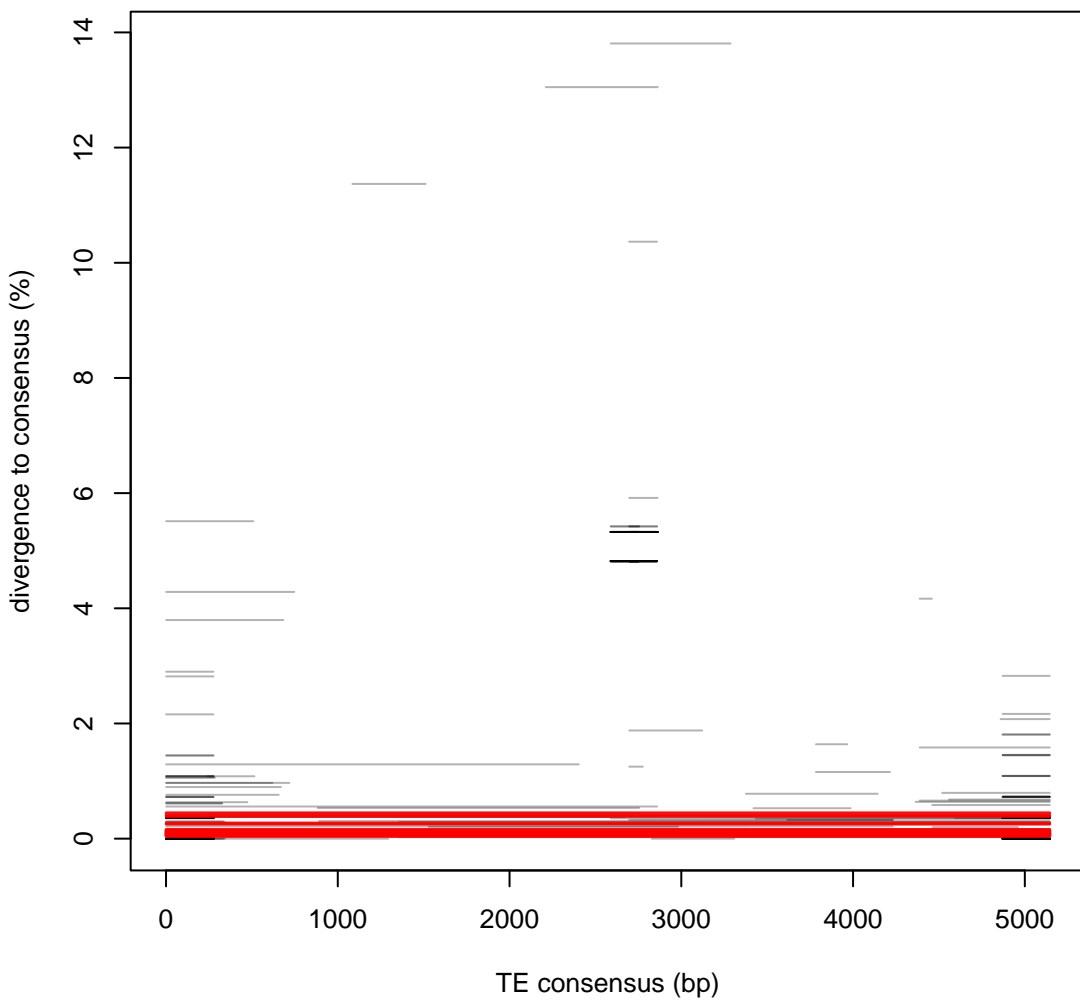
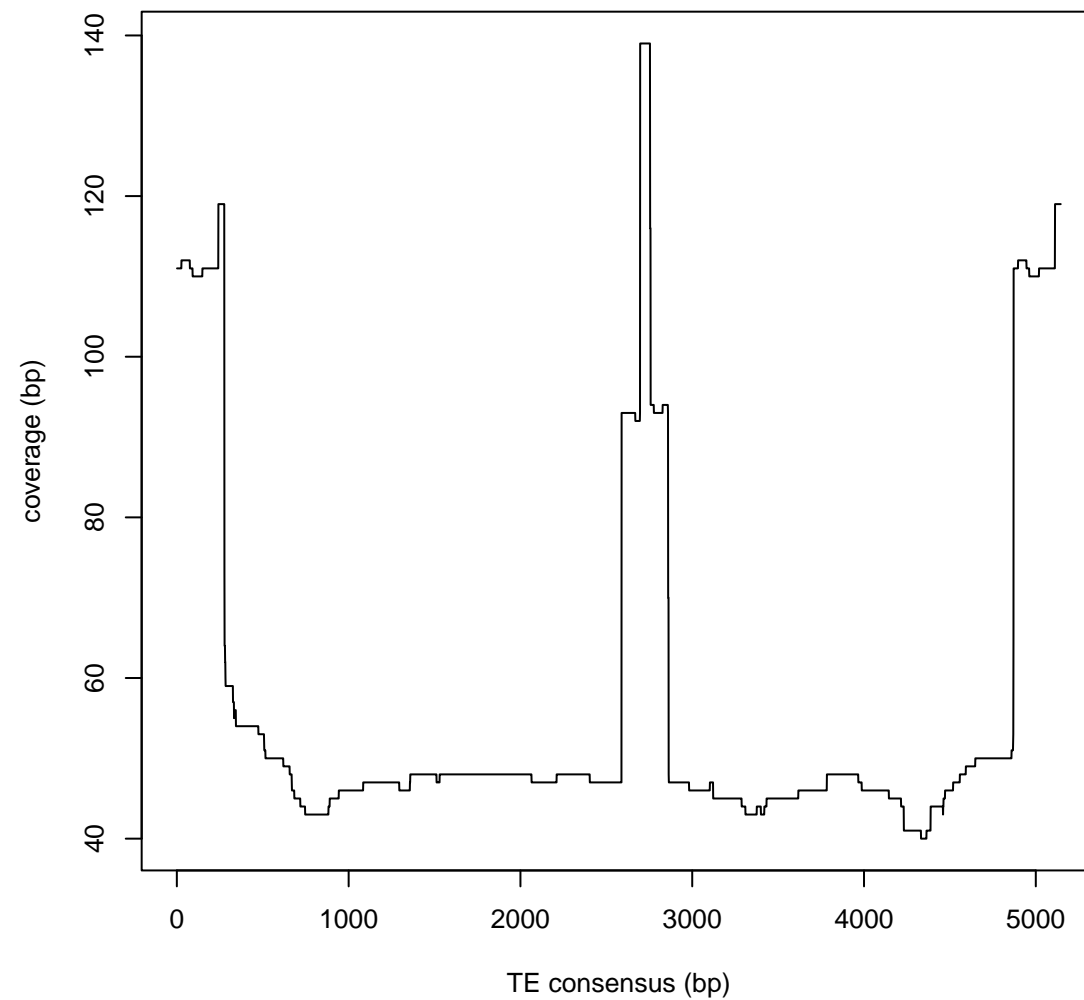


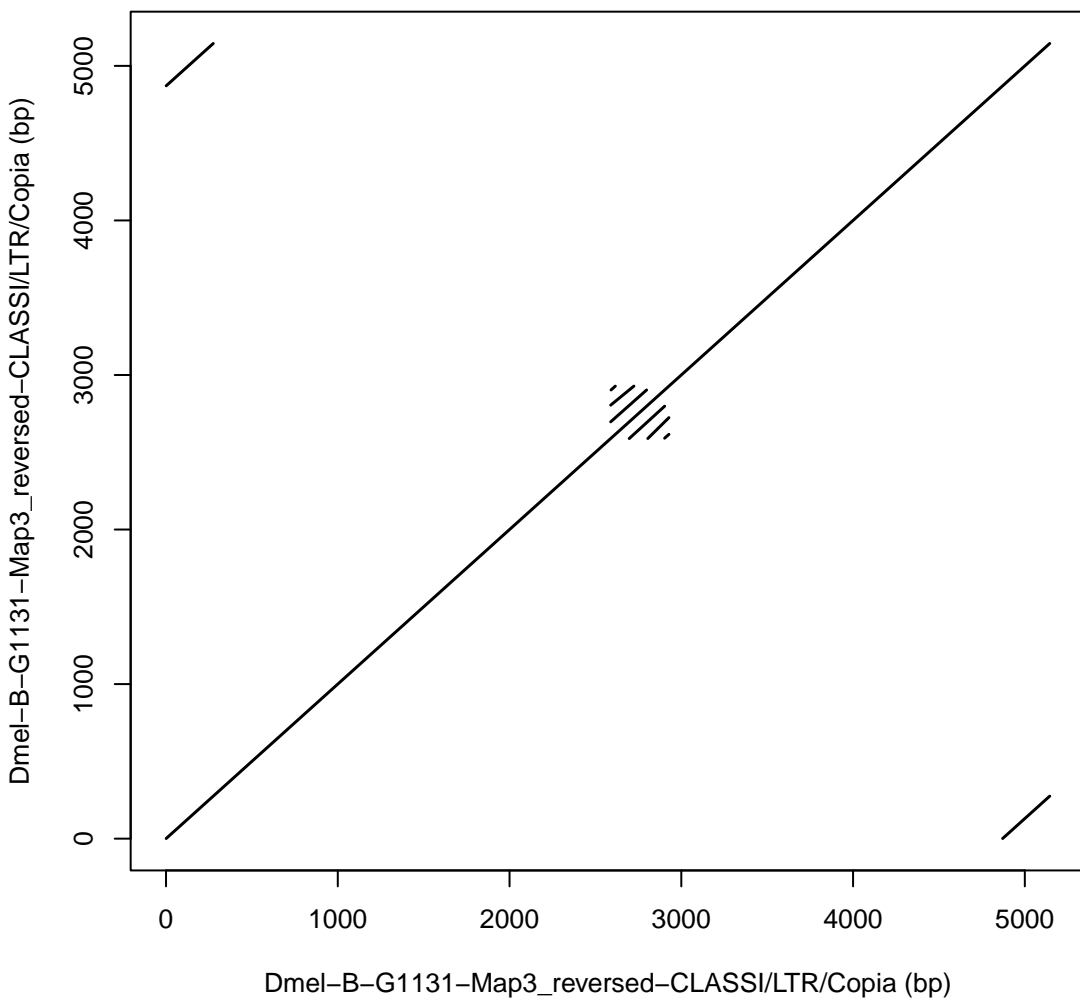
TE: Dmel-B-G1131-Map3_reversed-CLASSI/LTR/Copia
consensus size: 5145bp; fragments: 313; full length: 32 (≥ 4630.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

