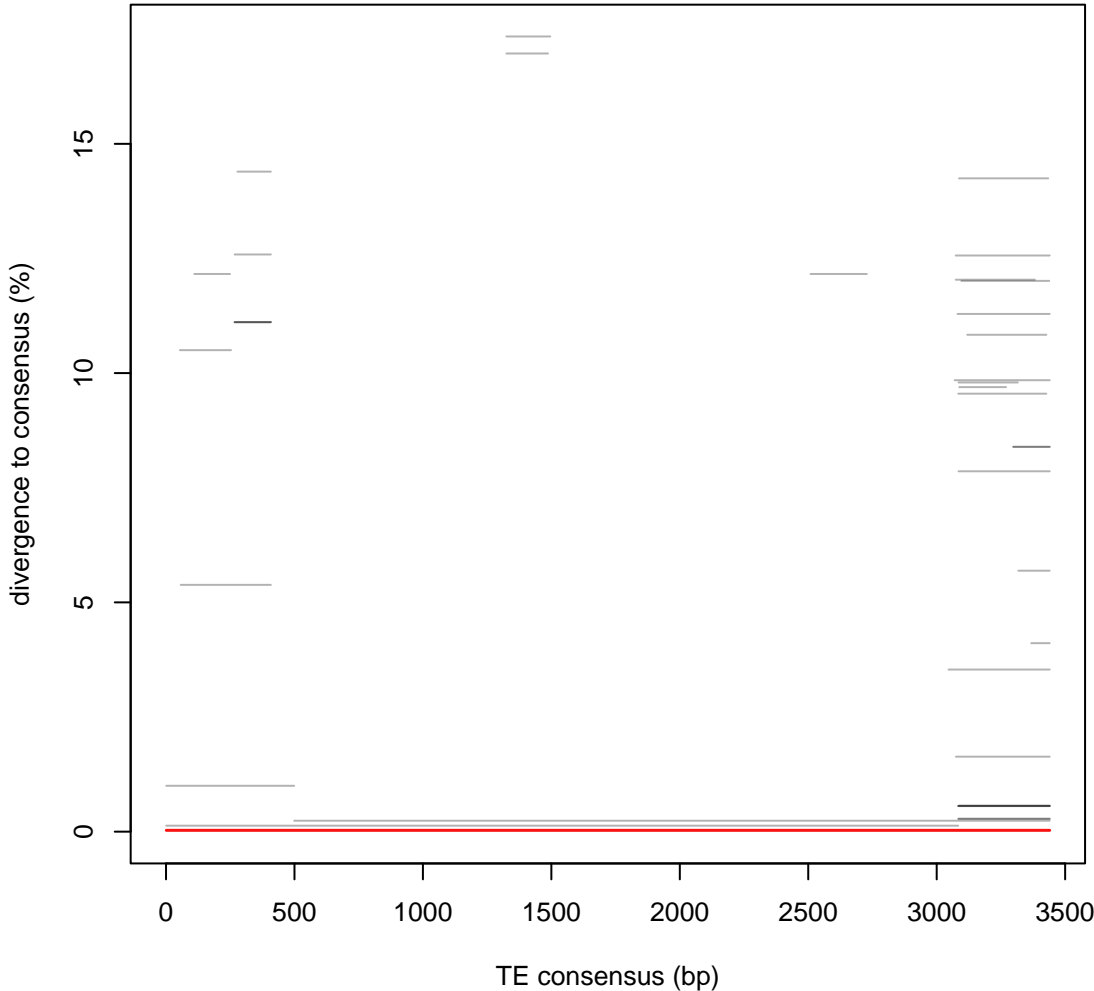
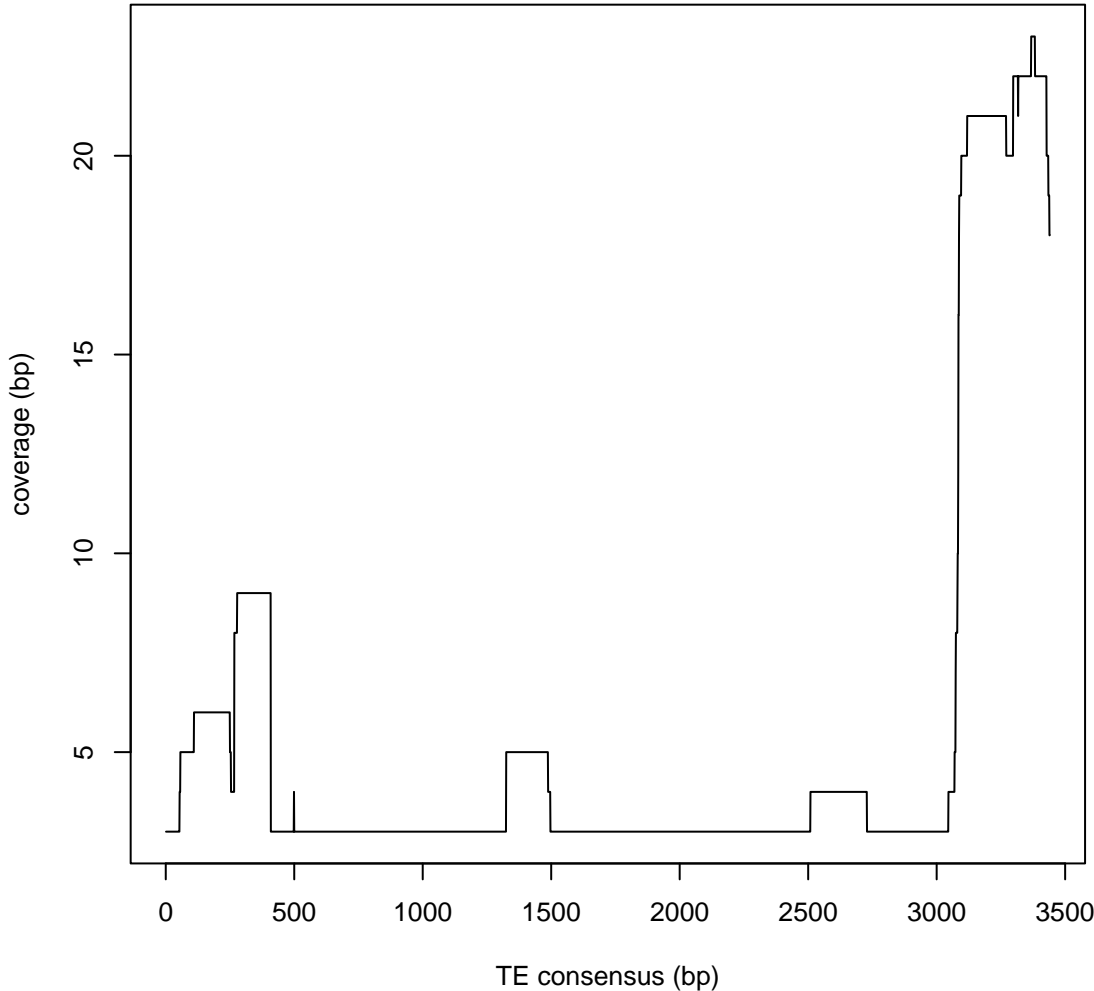


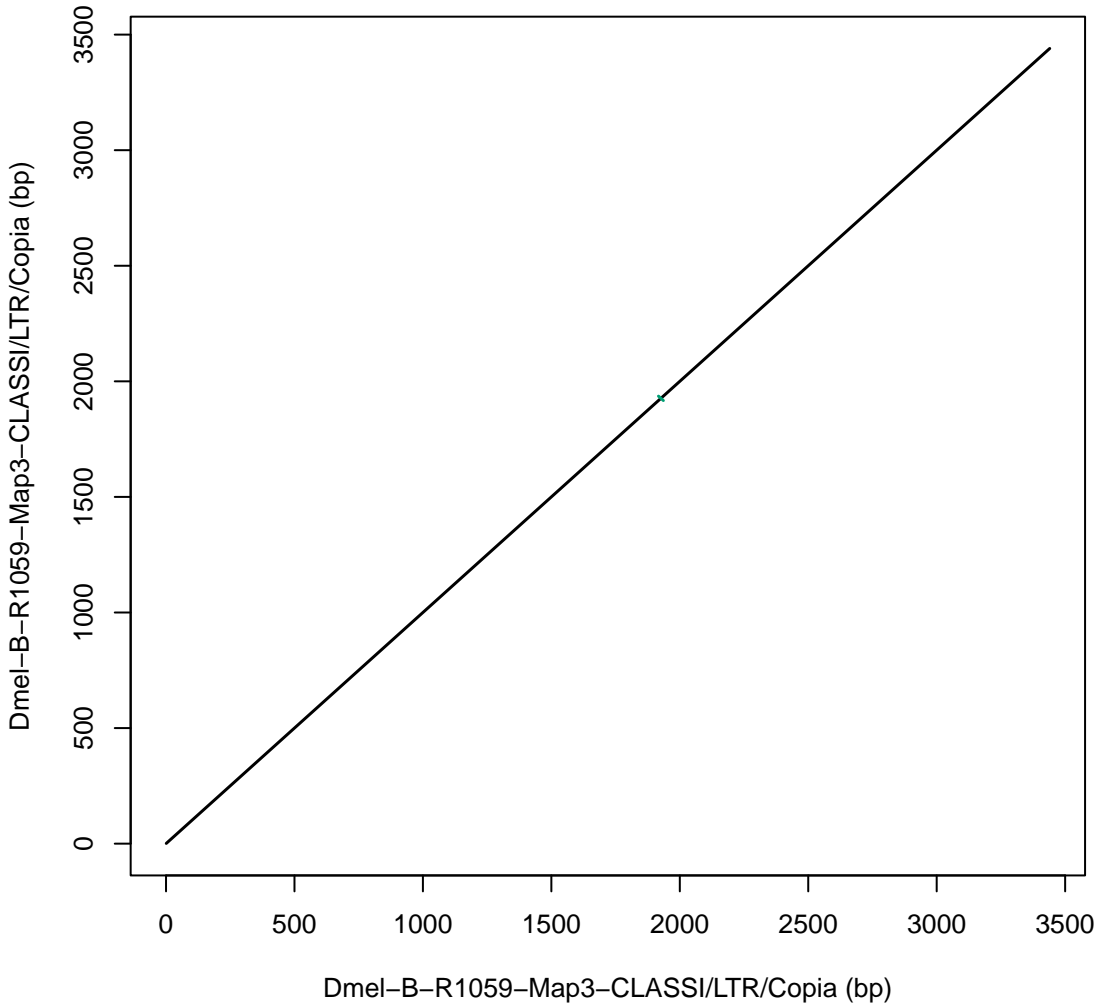
TE: Dmel-B-R1059-Map3-CLASSI/LTR/Copia  
consensus size: 3440bp; fragments: 38; full length: 1 (>=3096bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

