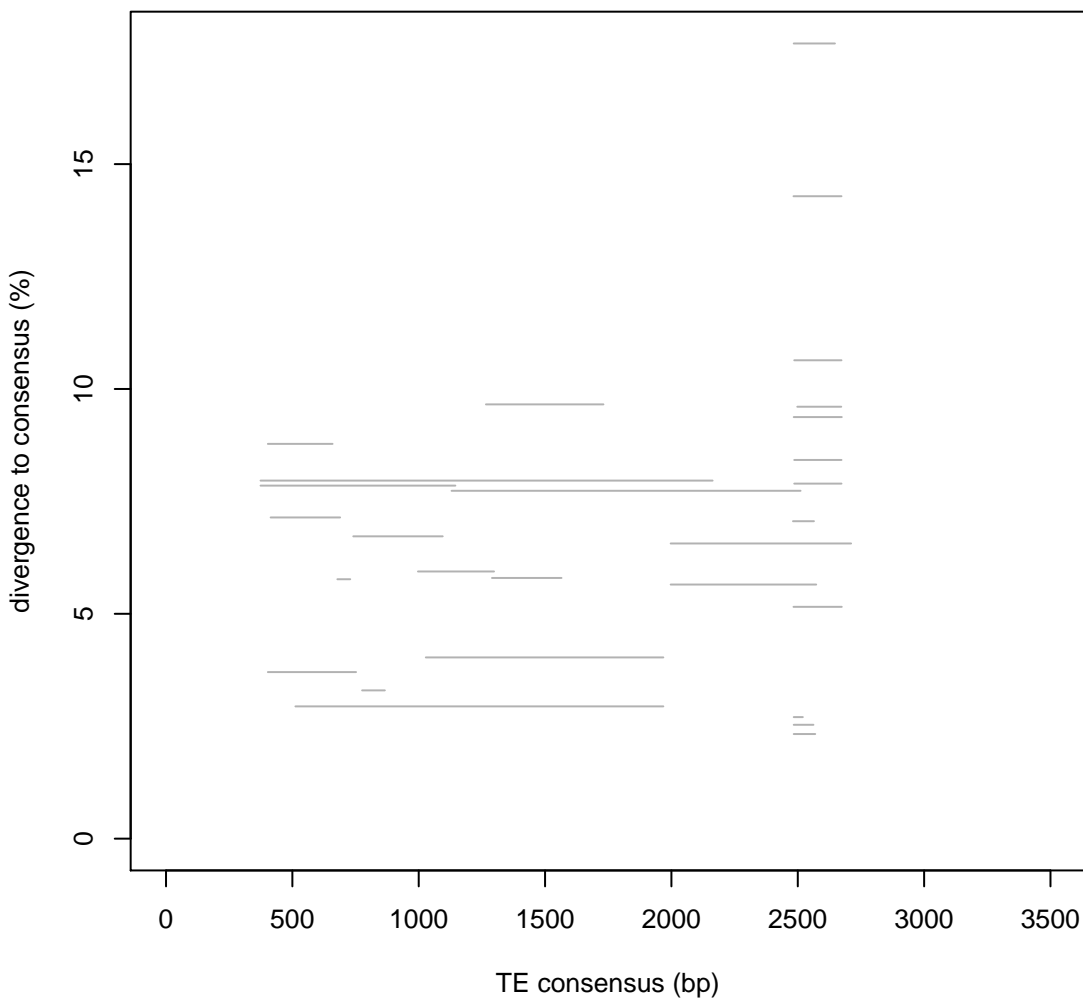
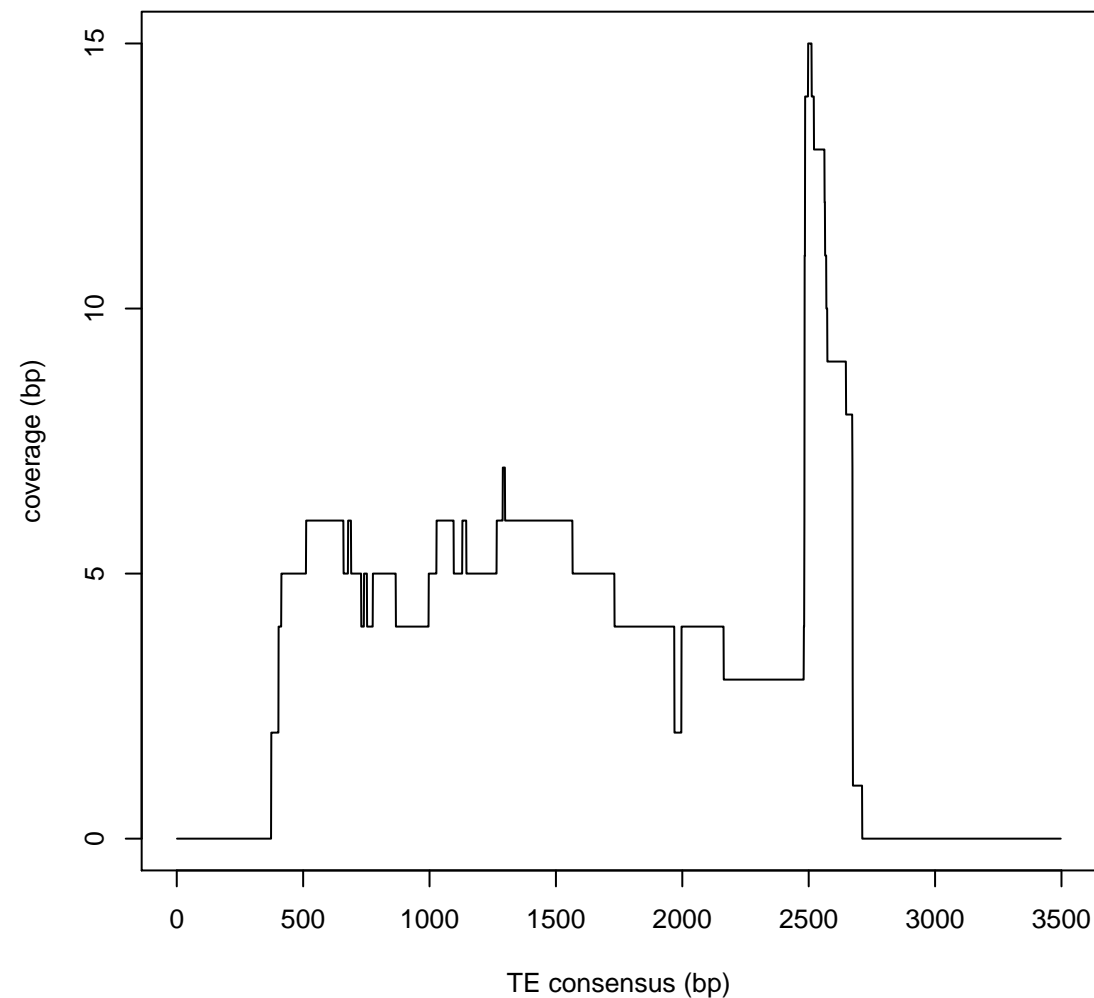


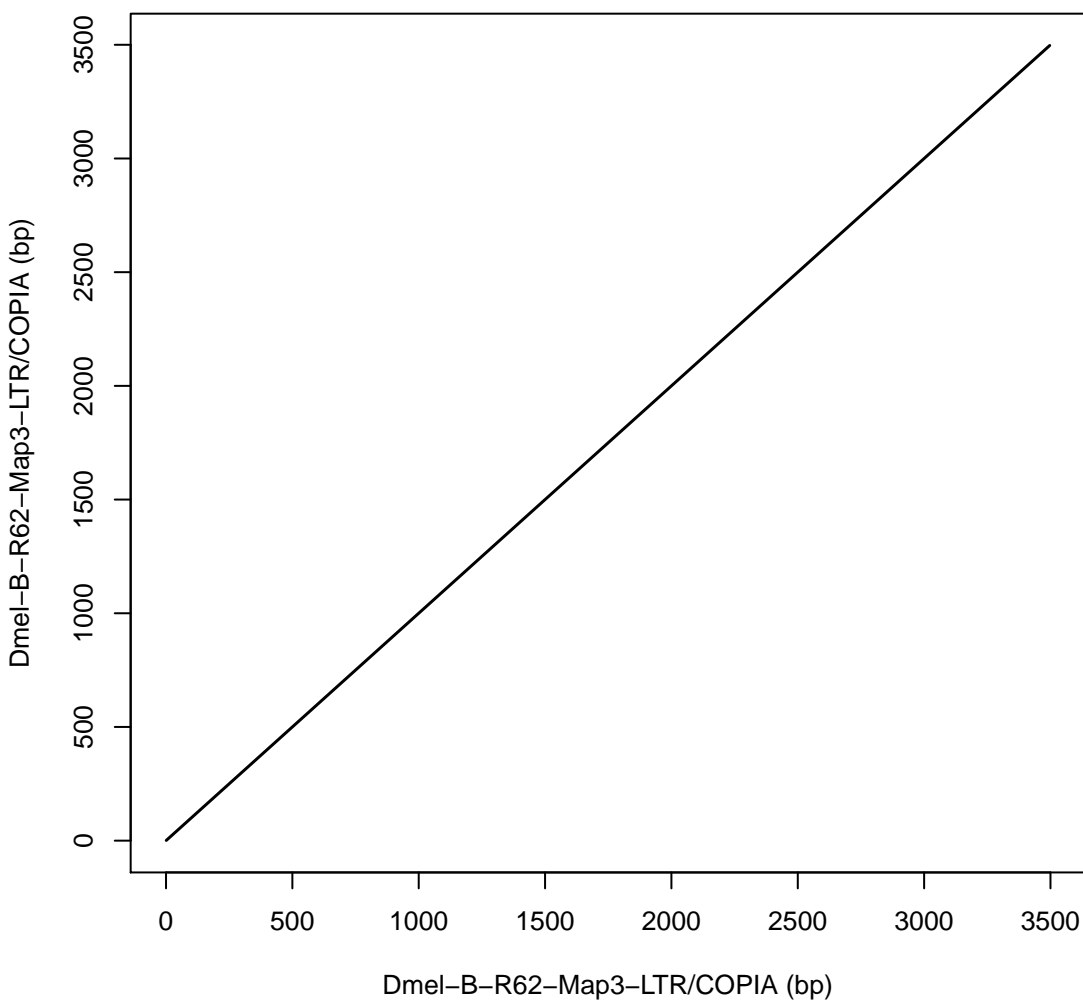
TE: Dmel-B-R62-Map3-LTR/COPIA  
consensus size: 3497bp; fragments: 28; full length: 0 ( $\geq 3147.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

no orf >400 bp detected