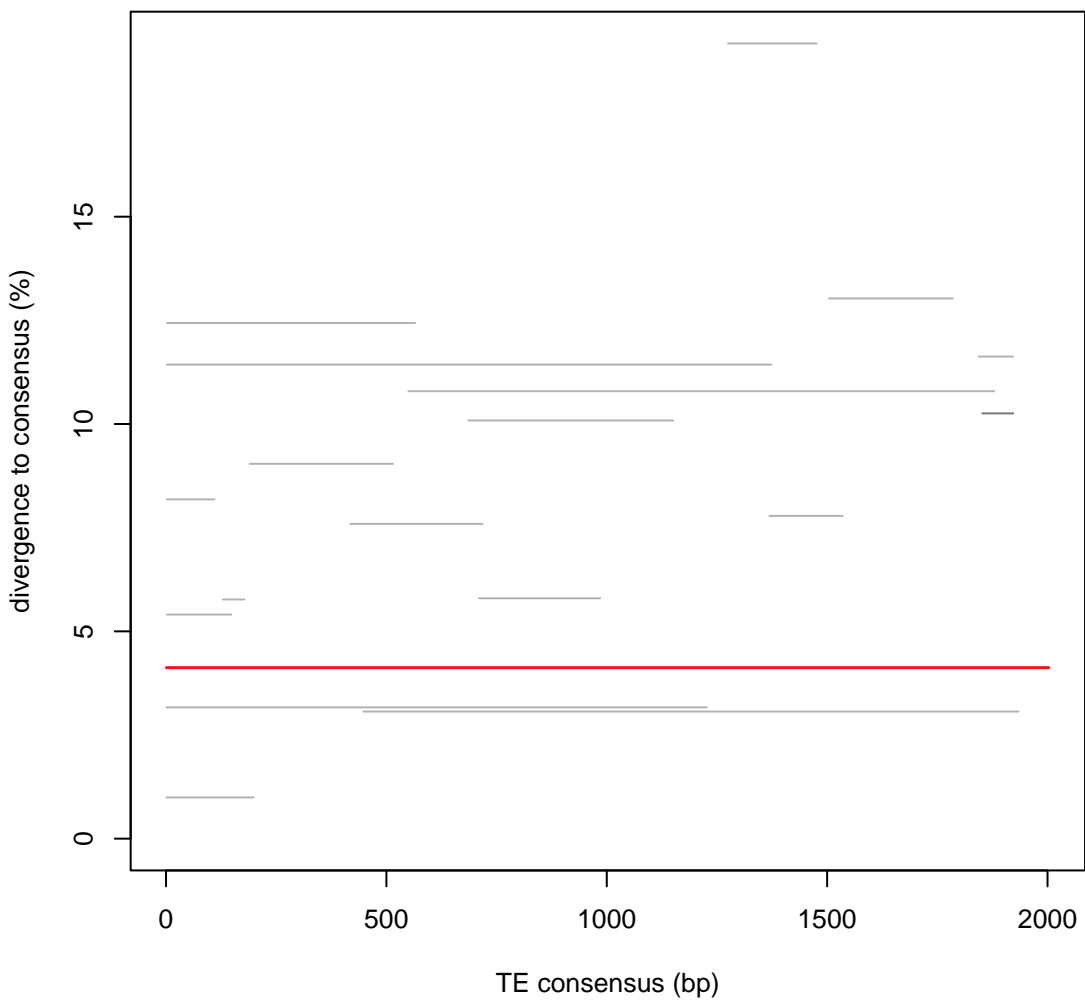
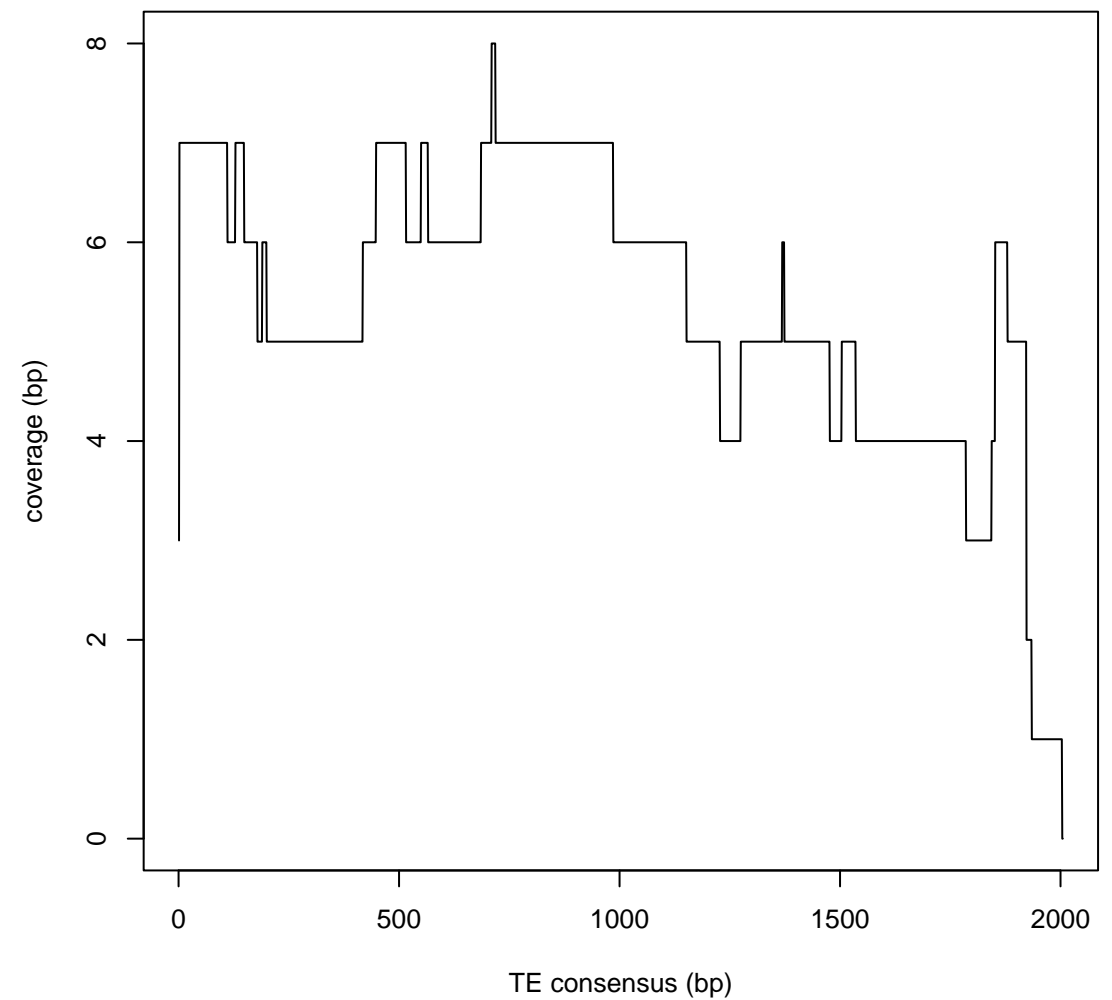


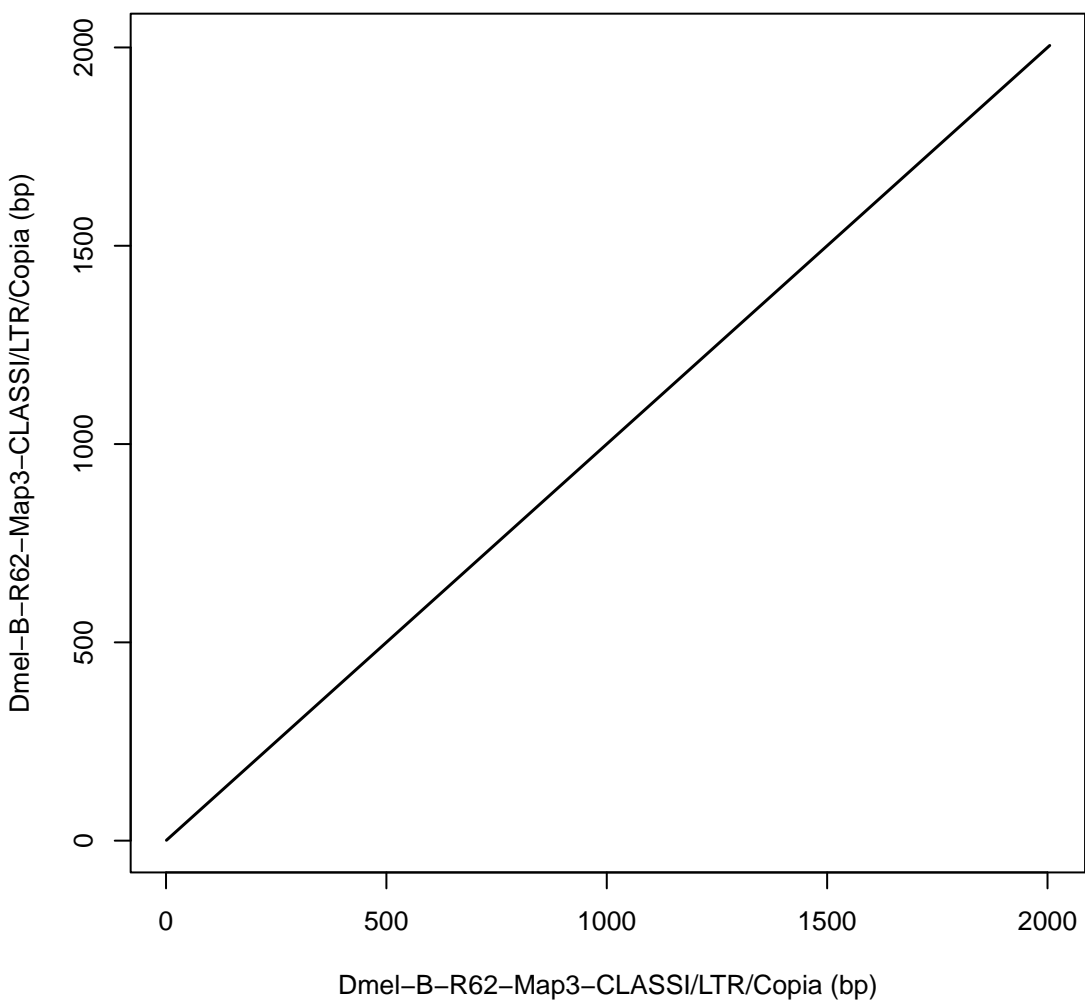
TE: Dmel-B-R62-Map3-CLASSI/LTR/Copia  
consensus size: 2005bp; fragments: 20; full length: 1 ( $\geq 1804.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

