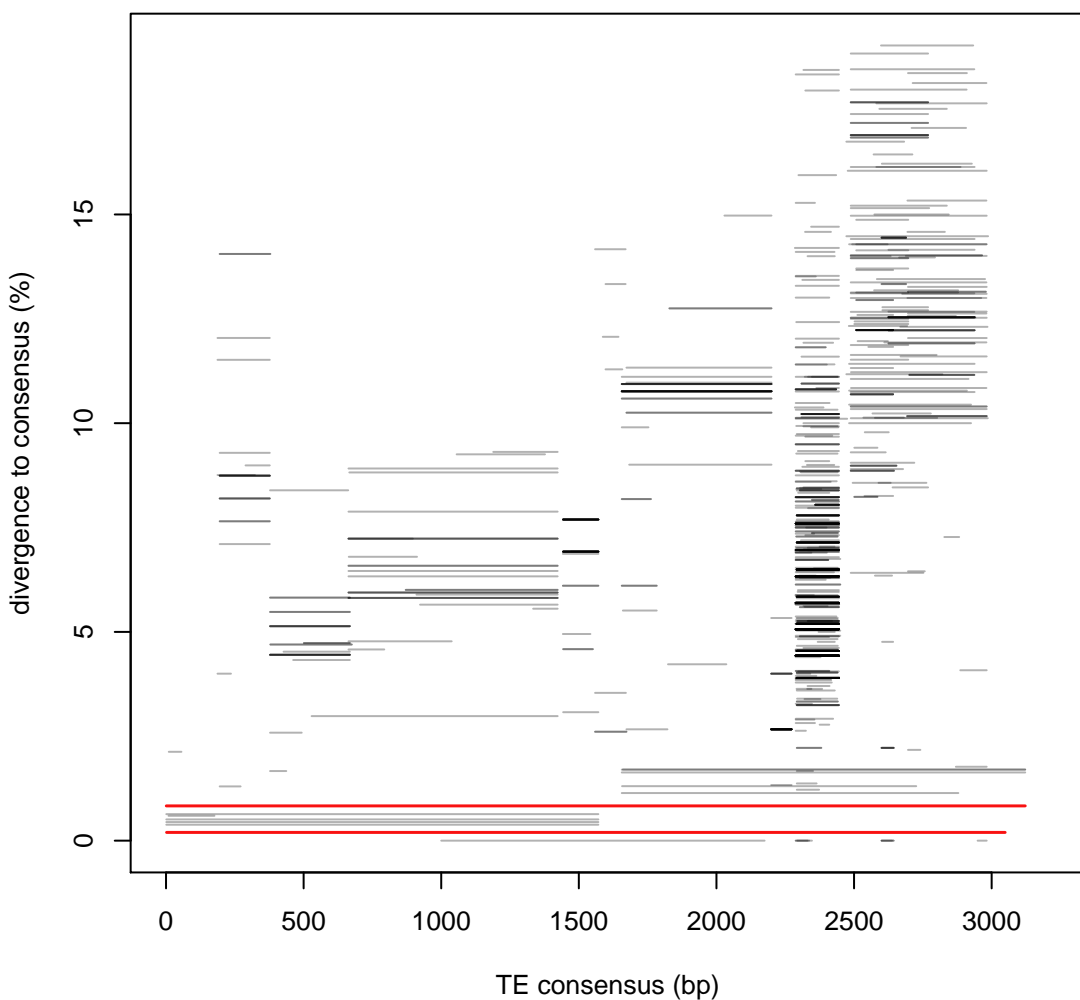
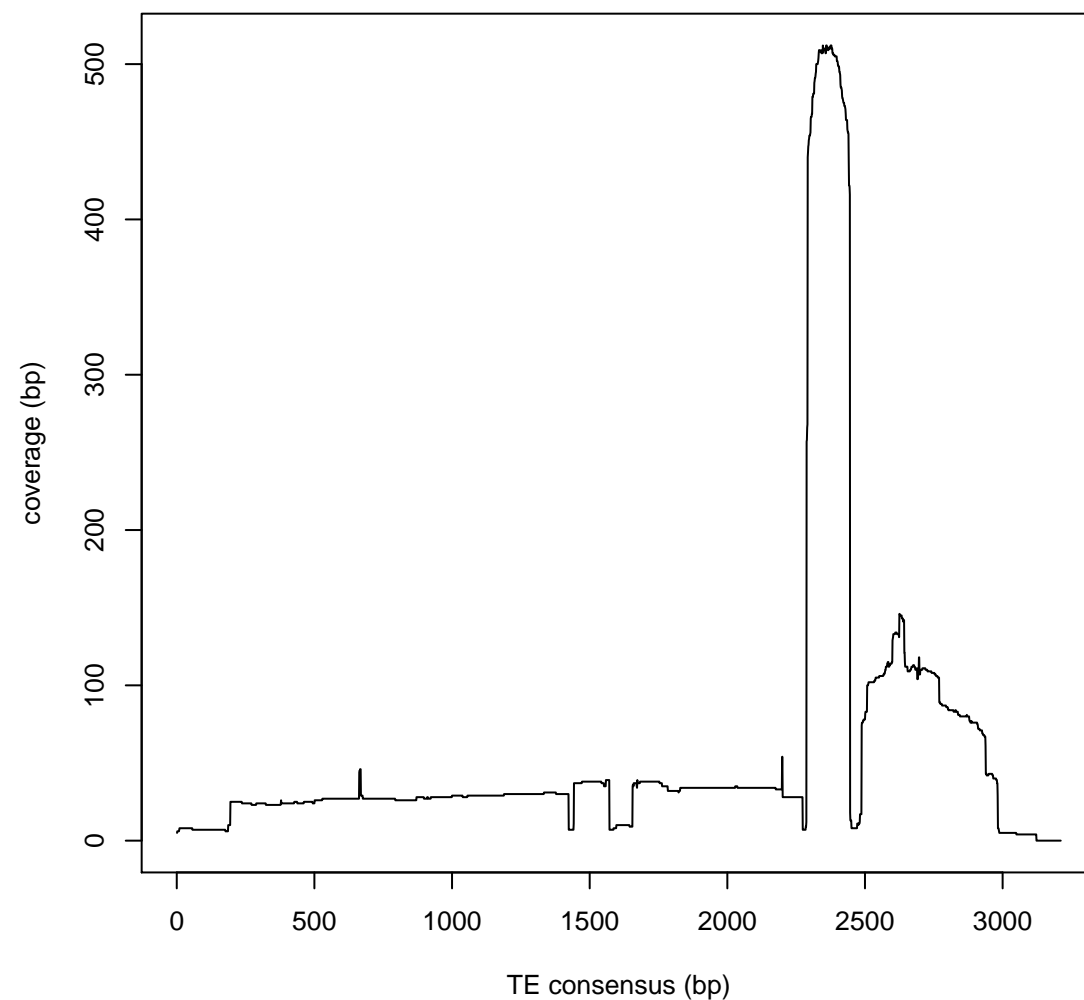


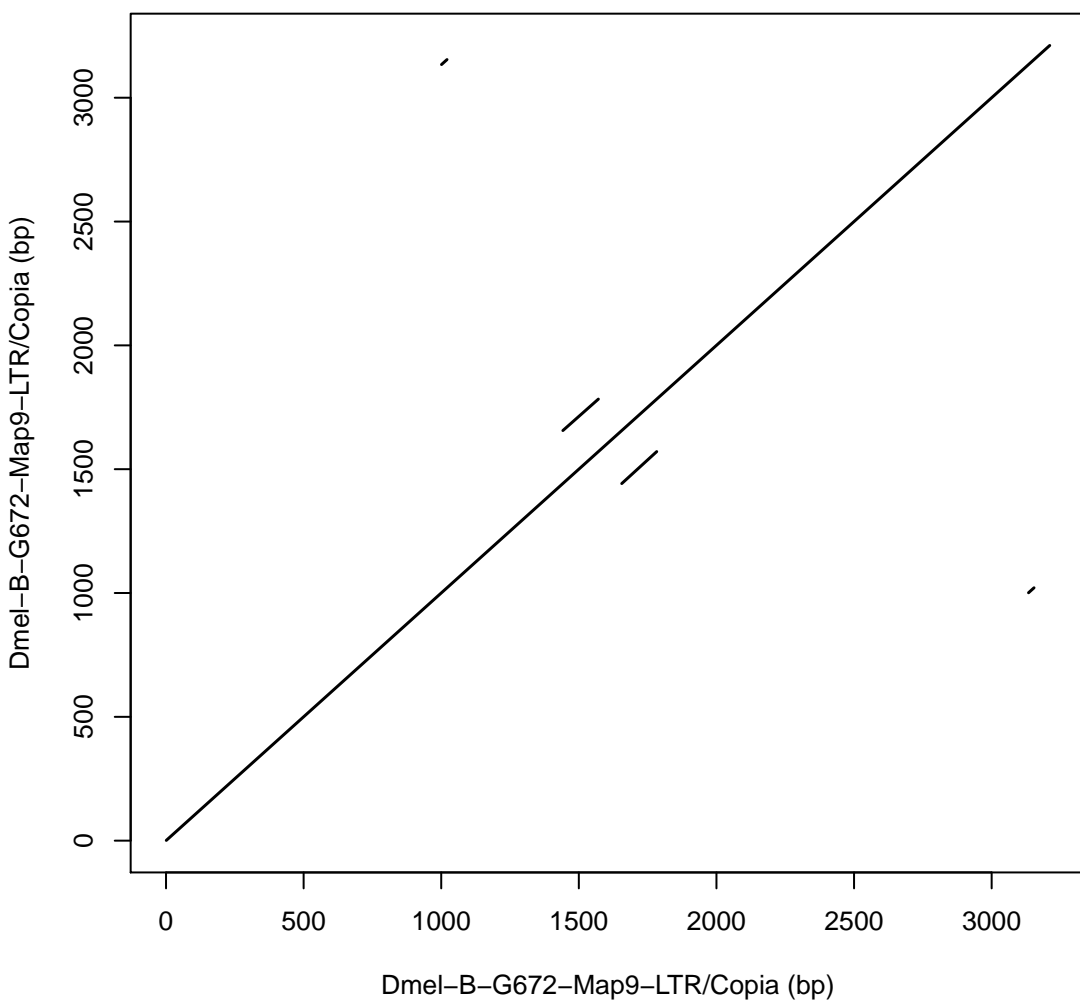
TE: Dmel-B-G672-Map9-LTR/Copia  
consensus size: 3211bp; fragments: 876; full length: 2 ( $\geq 2889.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

