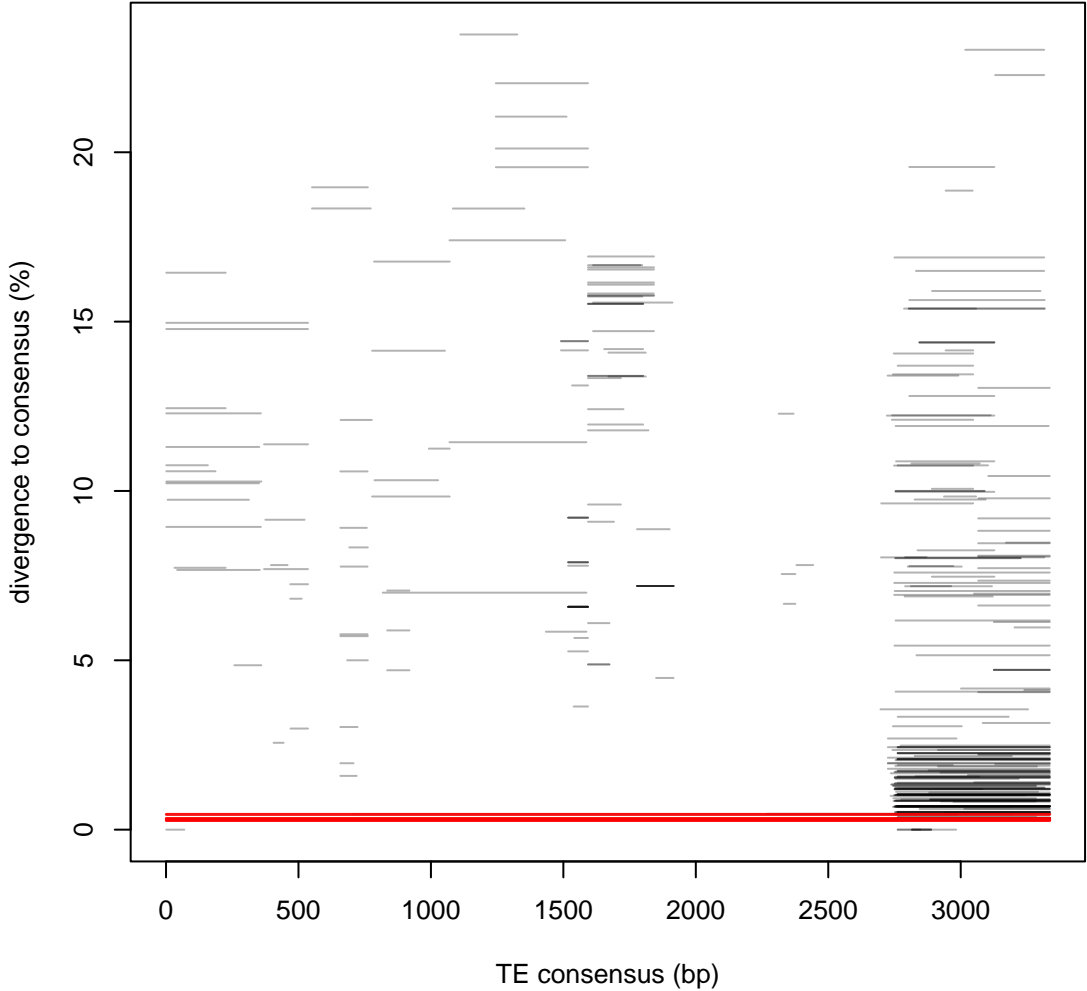
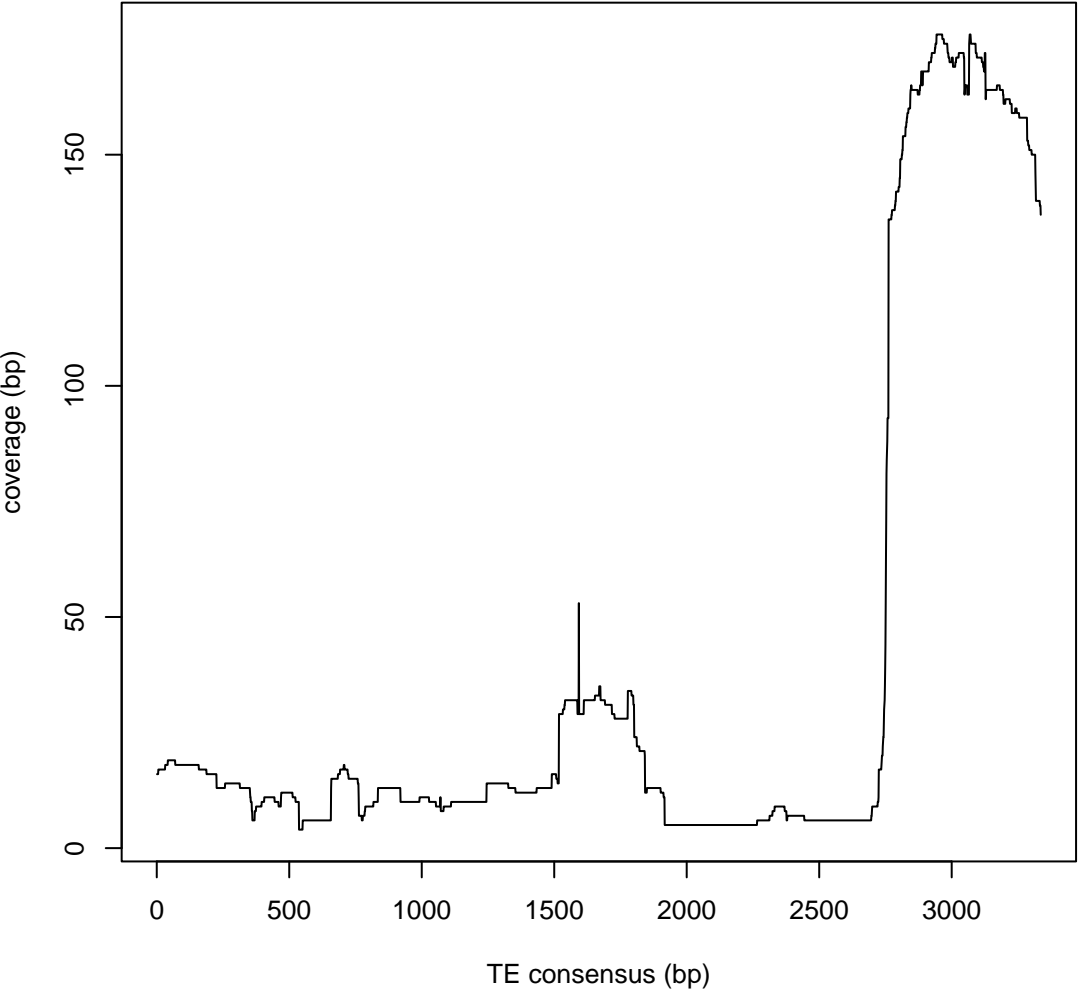


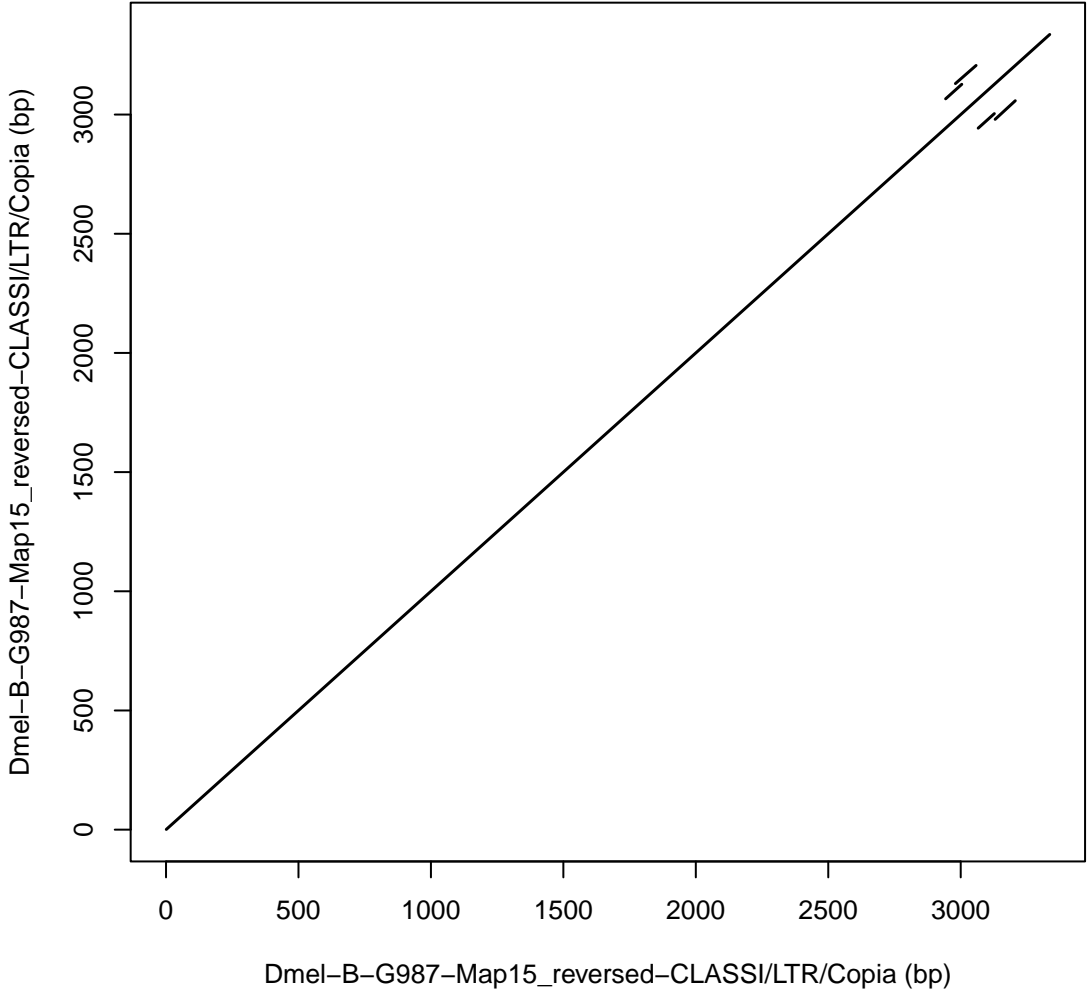
TE: Dmel-B-G987-Map15\_reversed-CLASSI/LTR/Copia  
consensus size: 3336bp; fragments: 329; full length: 4 (>=3002.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

