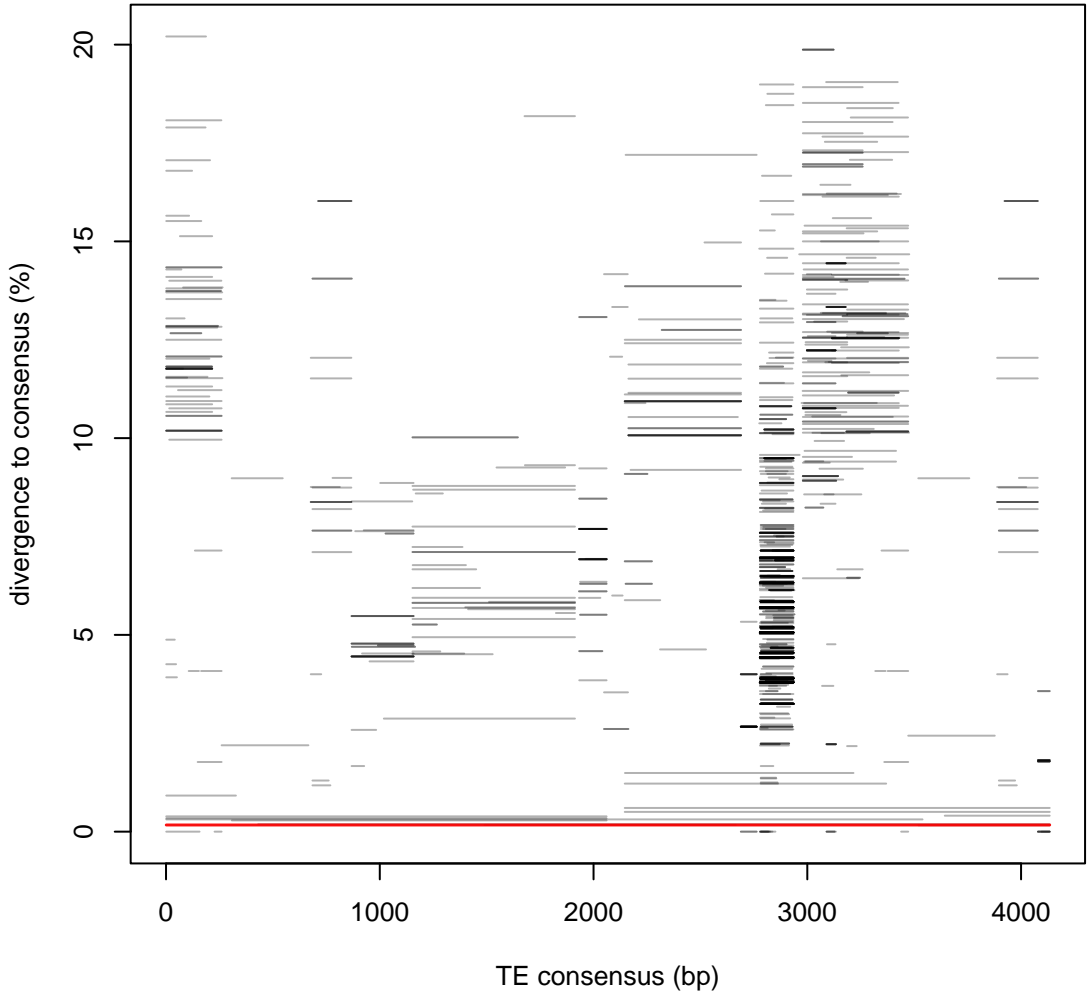
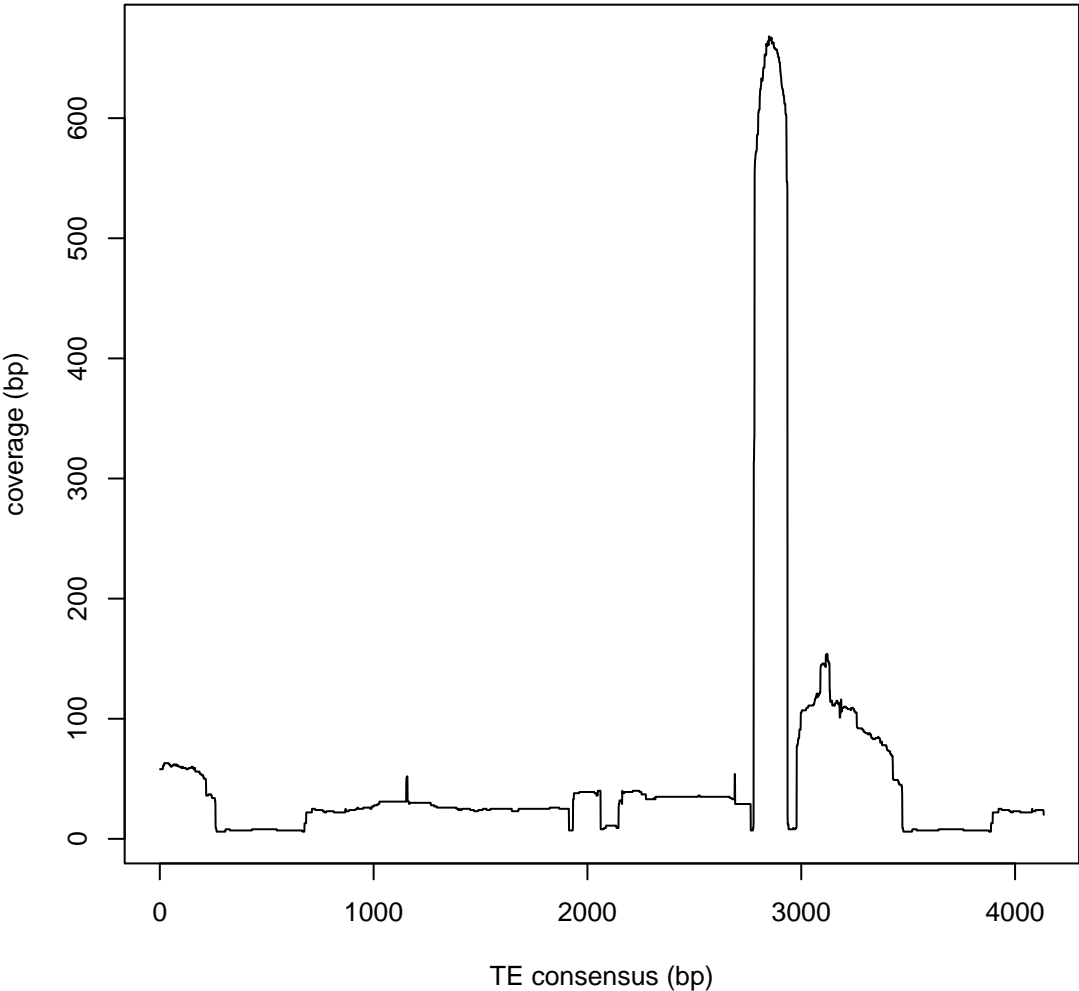


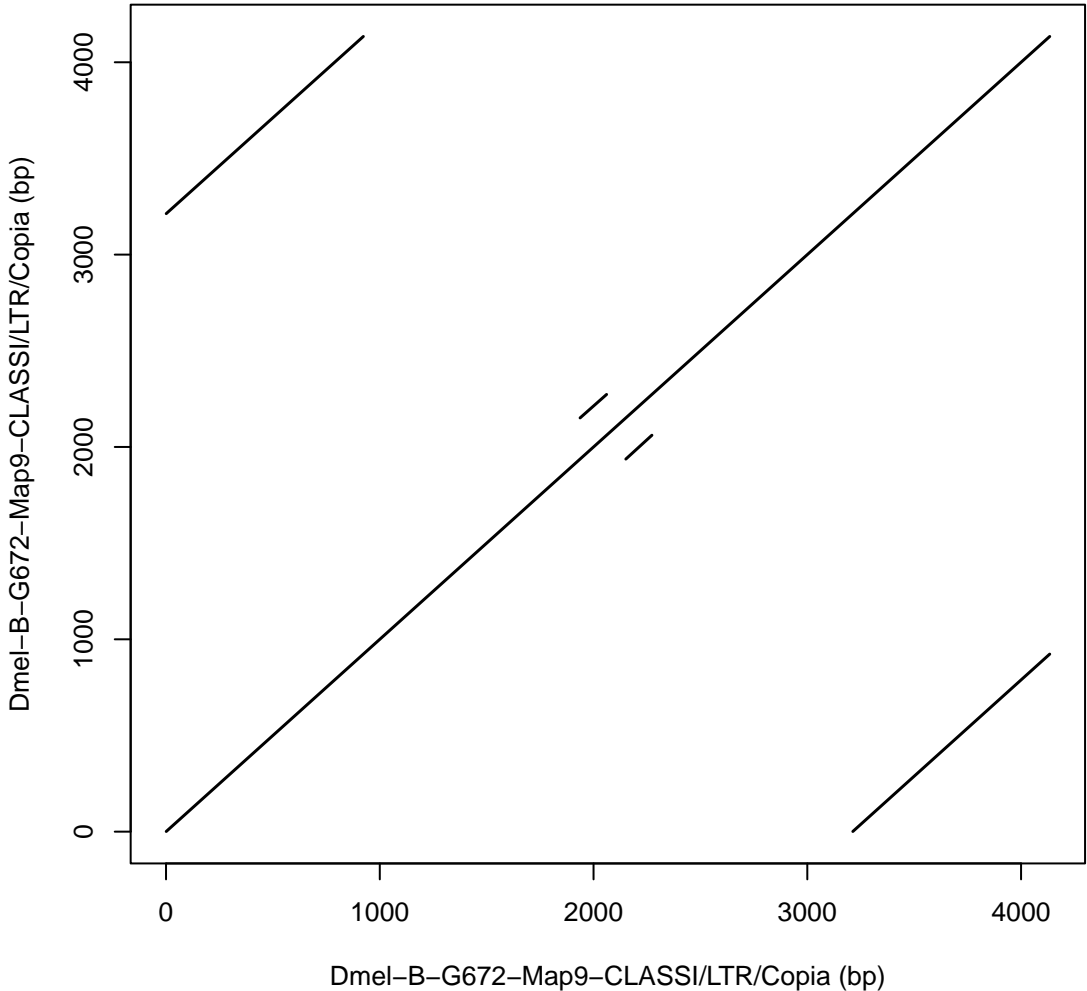
TE: Dmel-B-G672-Map9-CLASSI/LTR/Copia
consensus size: 4134bp; fragments: 1165; full length: 1 (>=3720.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

