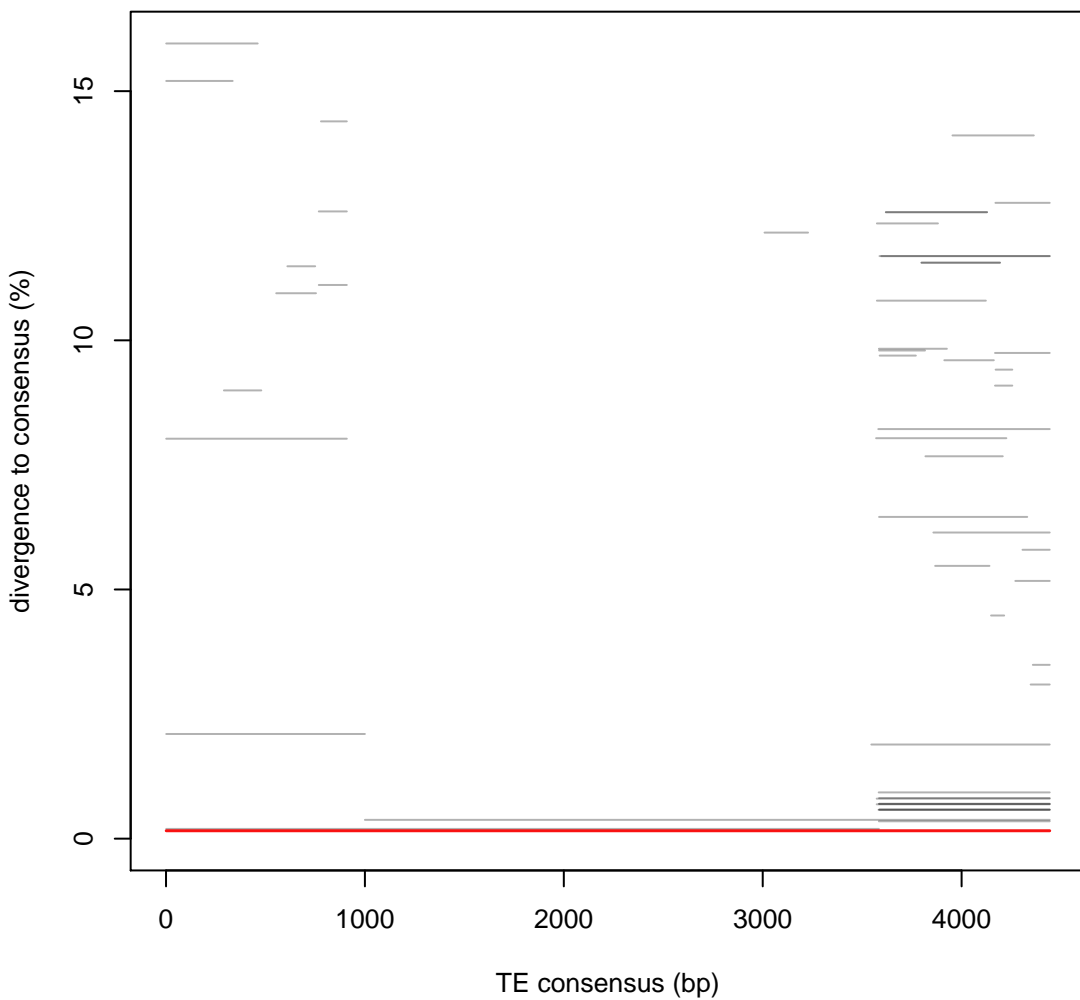
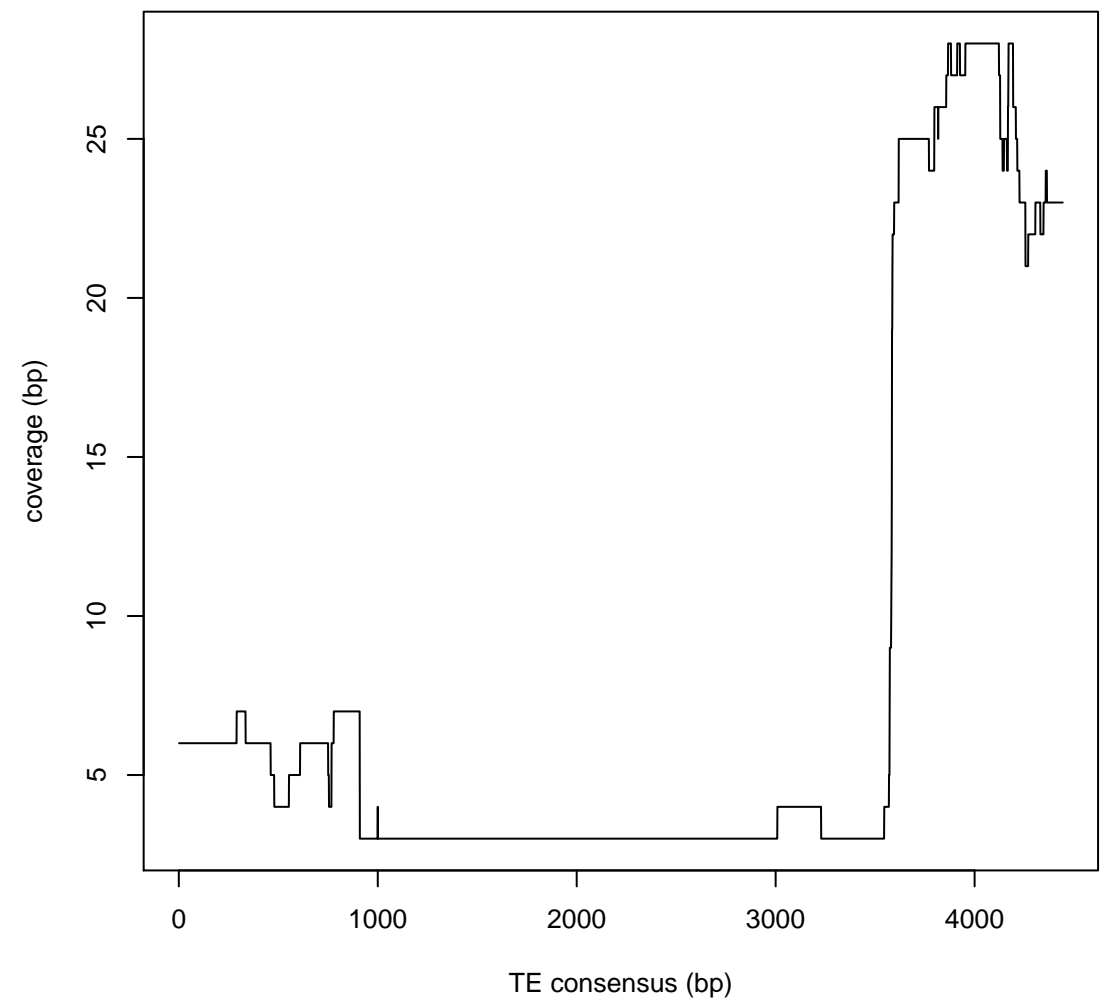


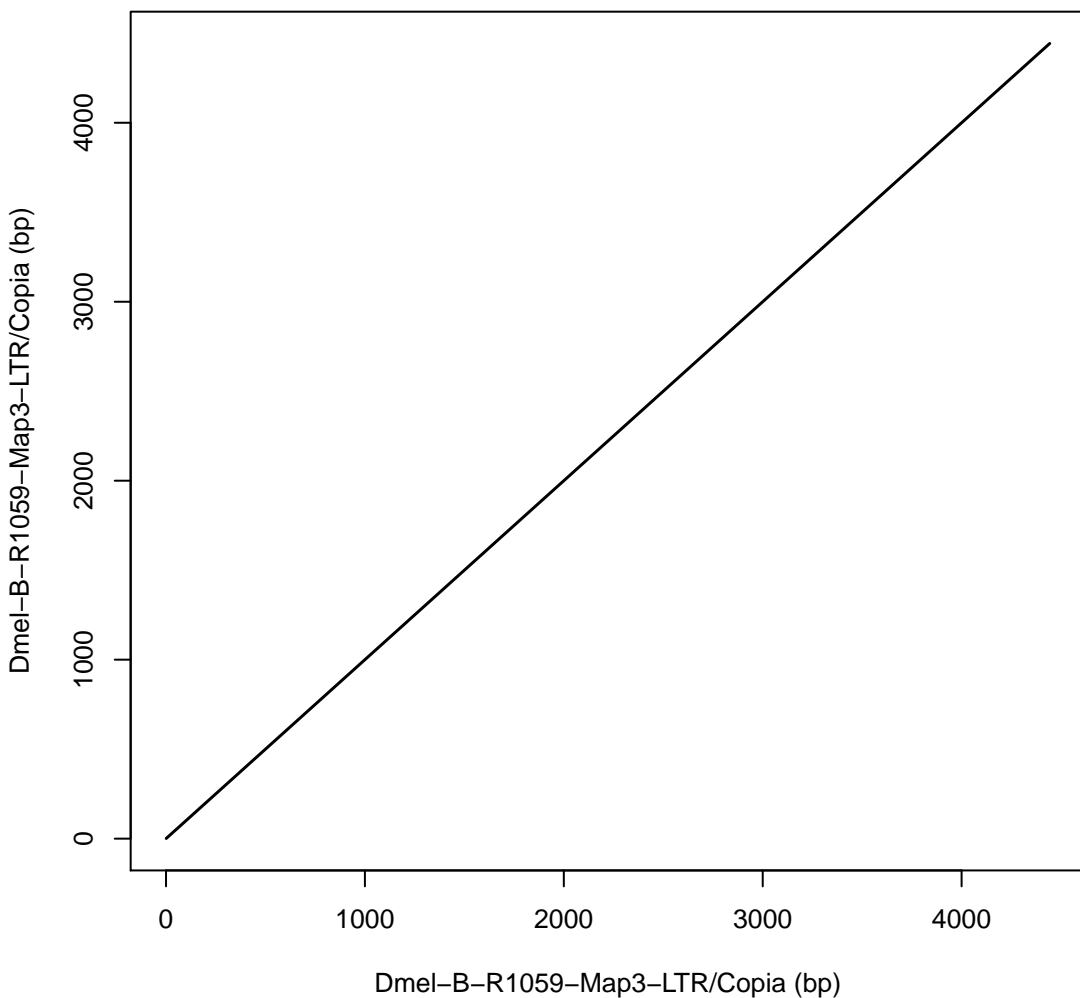
TE: Dmel-B-R1059-Map3-LTR/Copia
consensus size: 4443bp; fragments: 53; full length: 1 (>=3998.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

