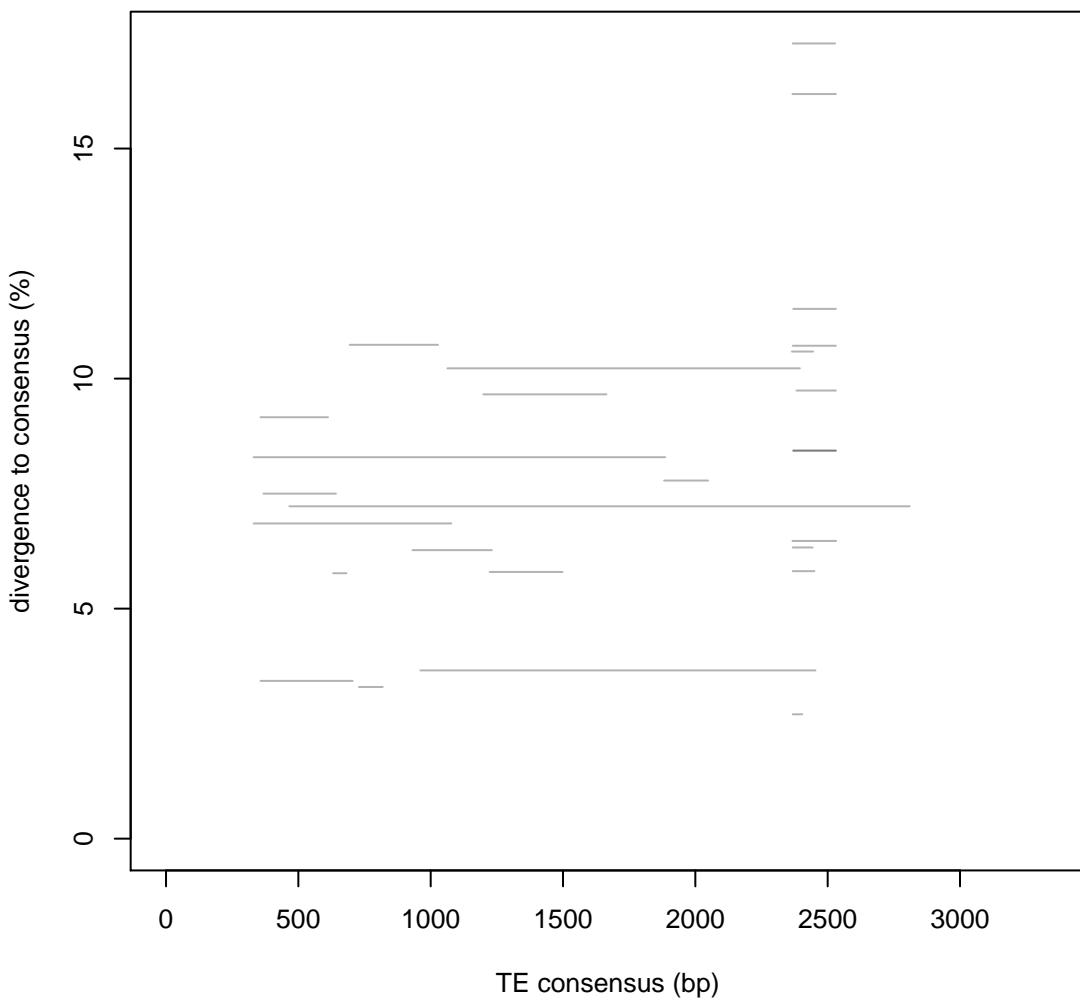
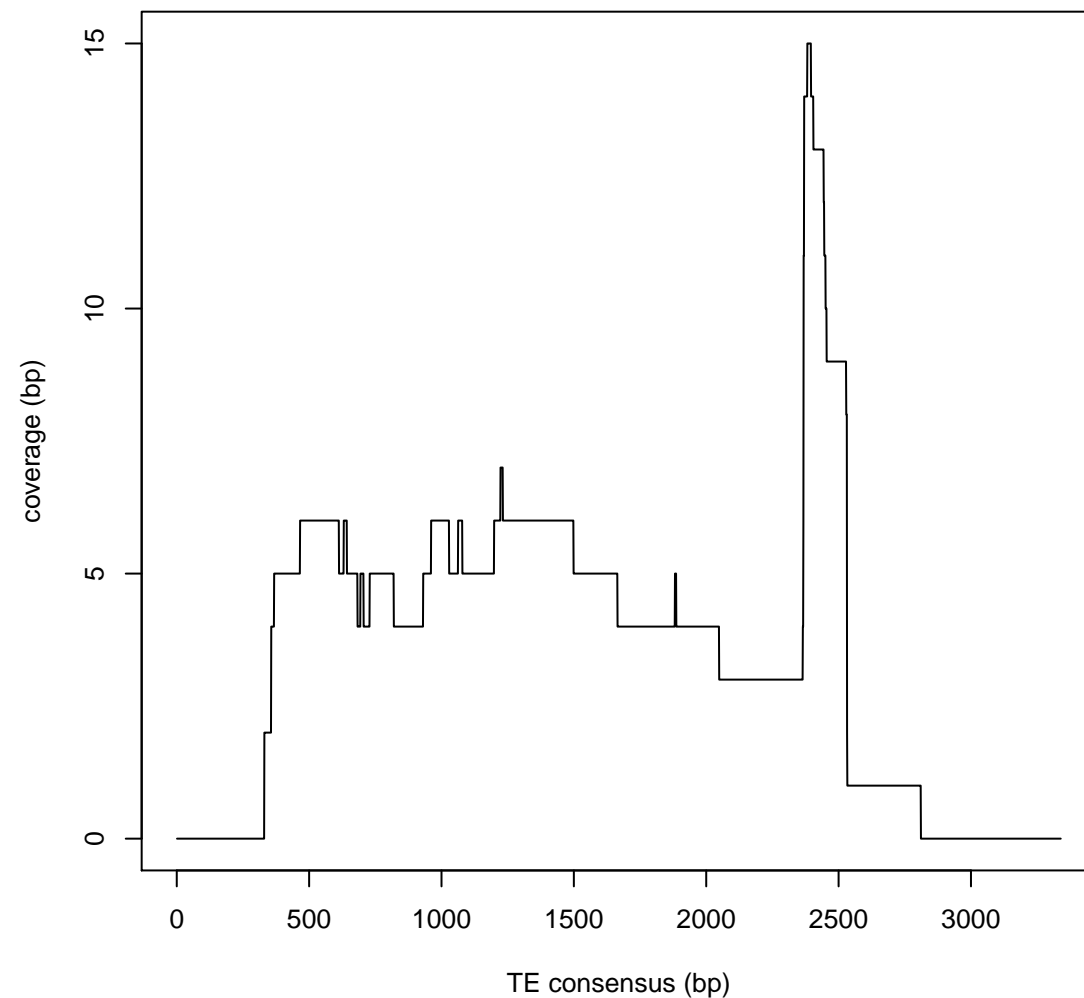


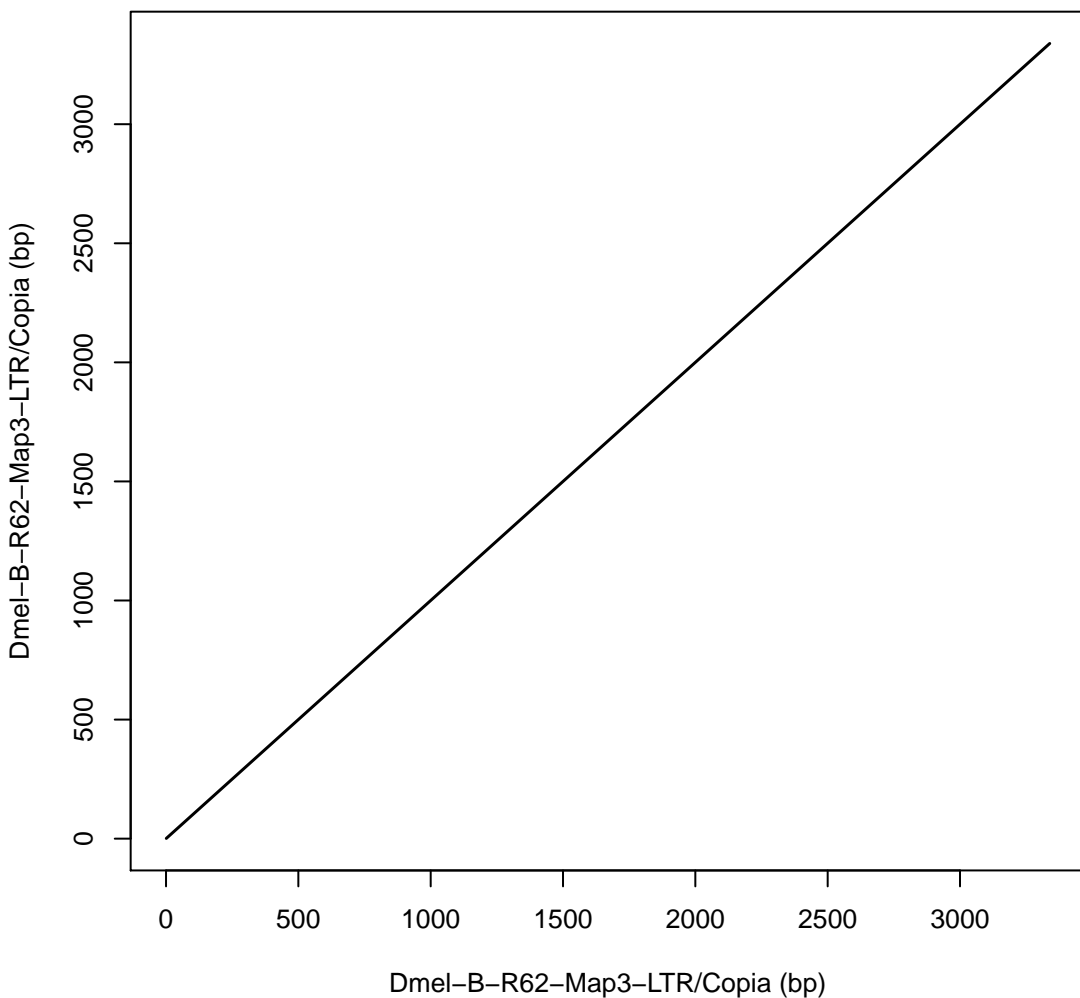
TE: Dmel-B-R62-Map3-LTR/Copia
consensus size: 3339bp; fragments: 27; full length: 0 (>=3005.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

