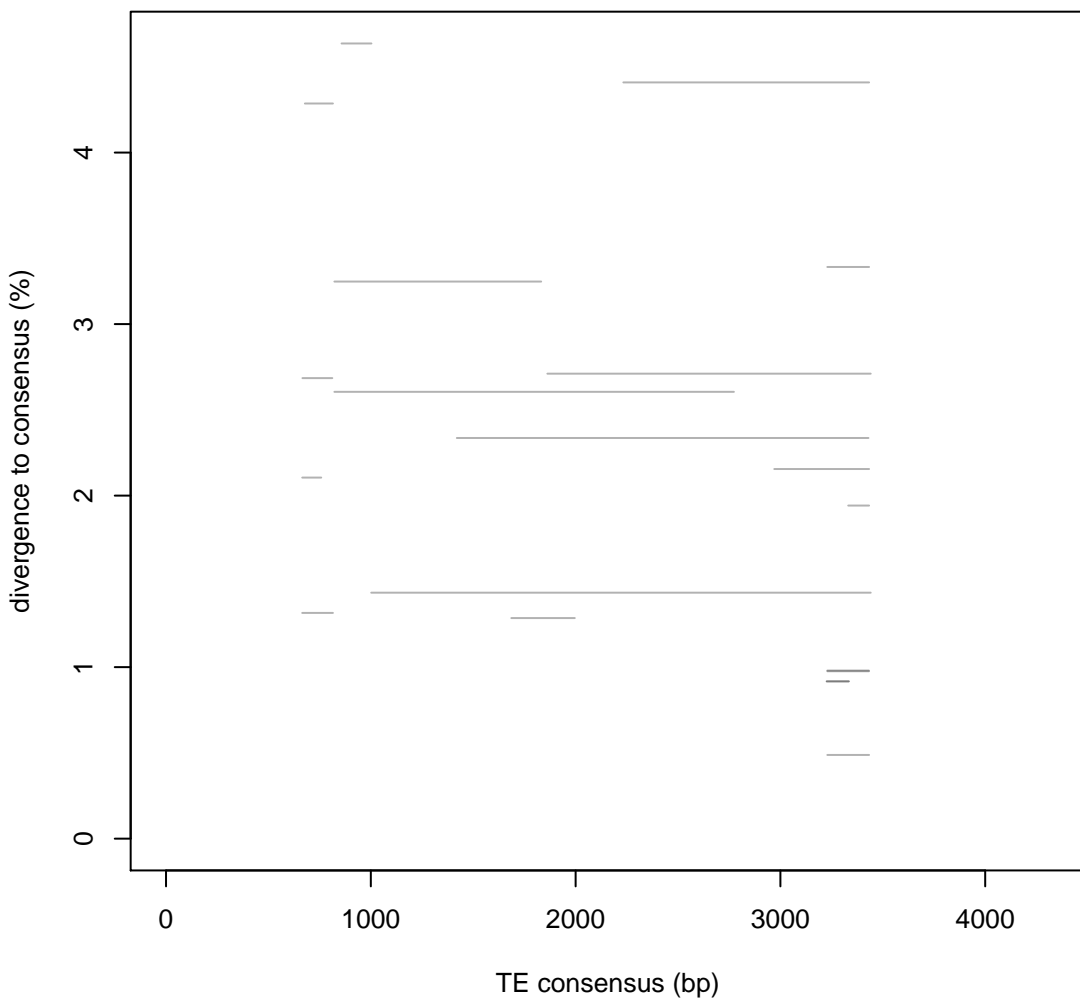
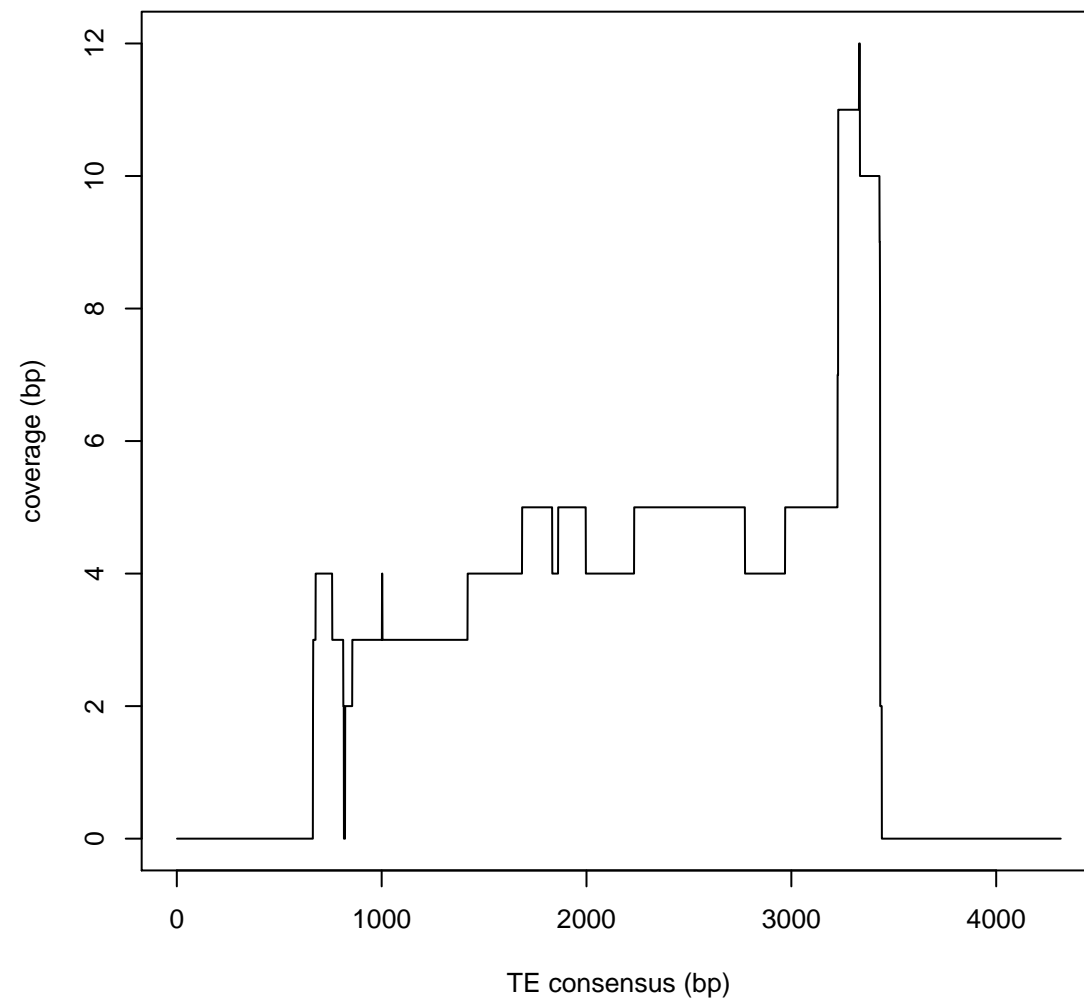


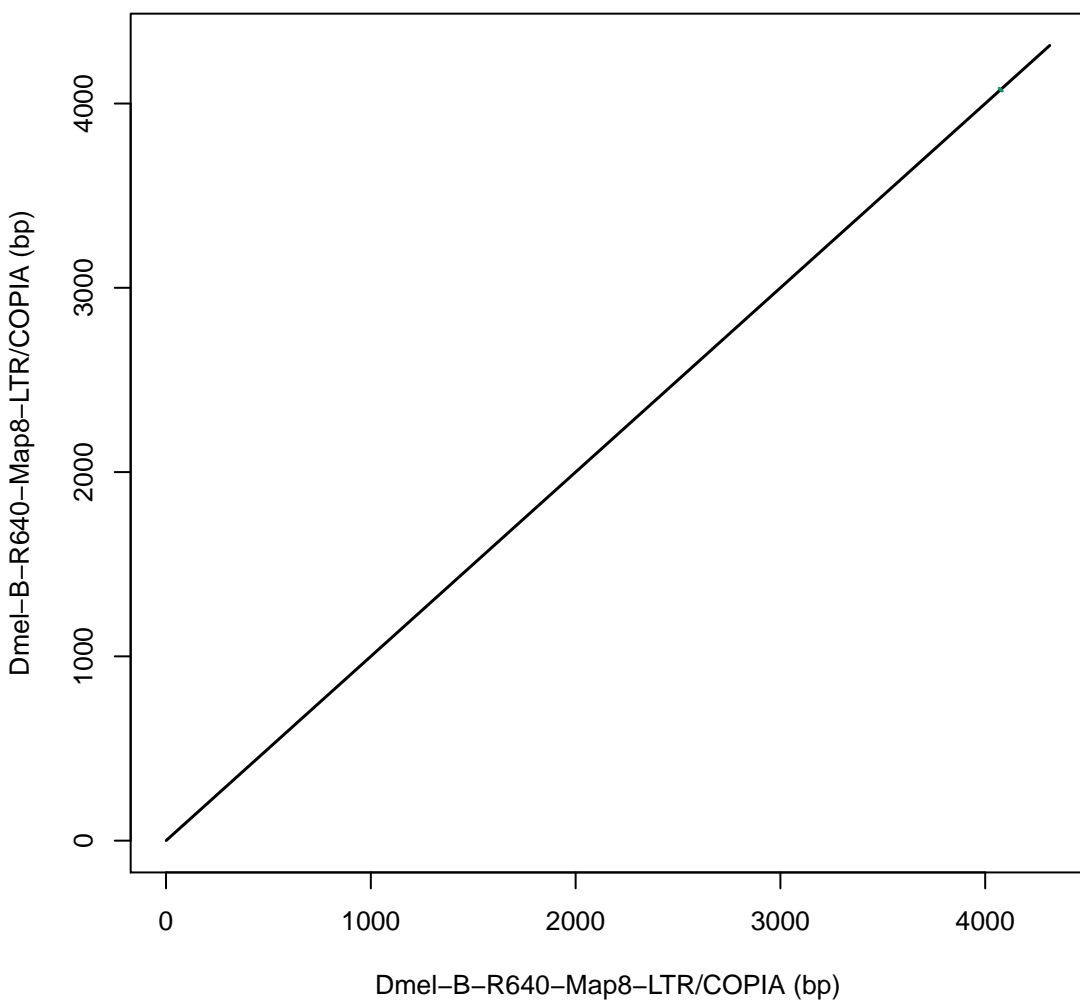
TE: Dmel-B-R640-Map8-LTR/COPIA
consensus size: 4315bp; fragments: 20; full length: 0 (>=3883.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

