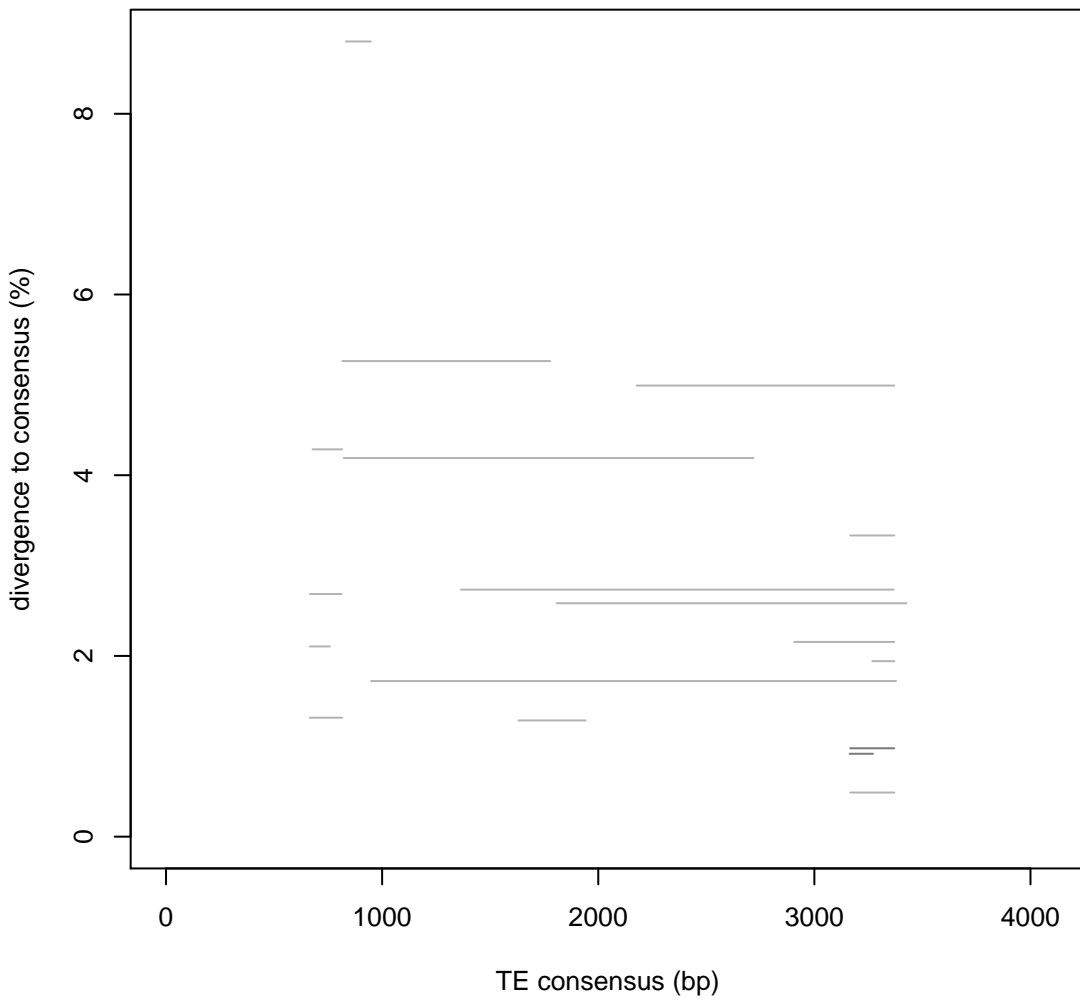
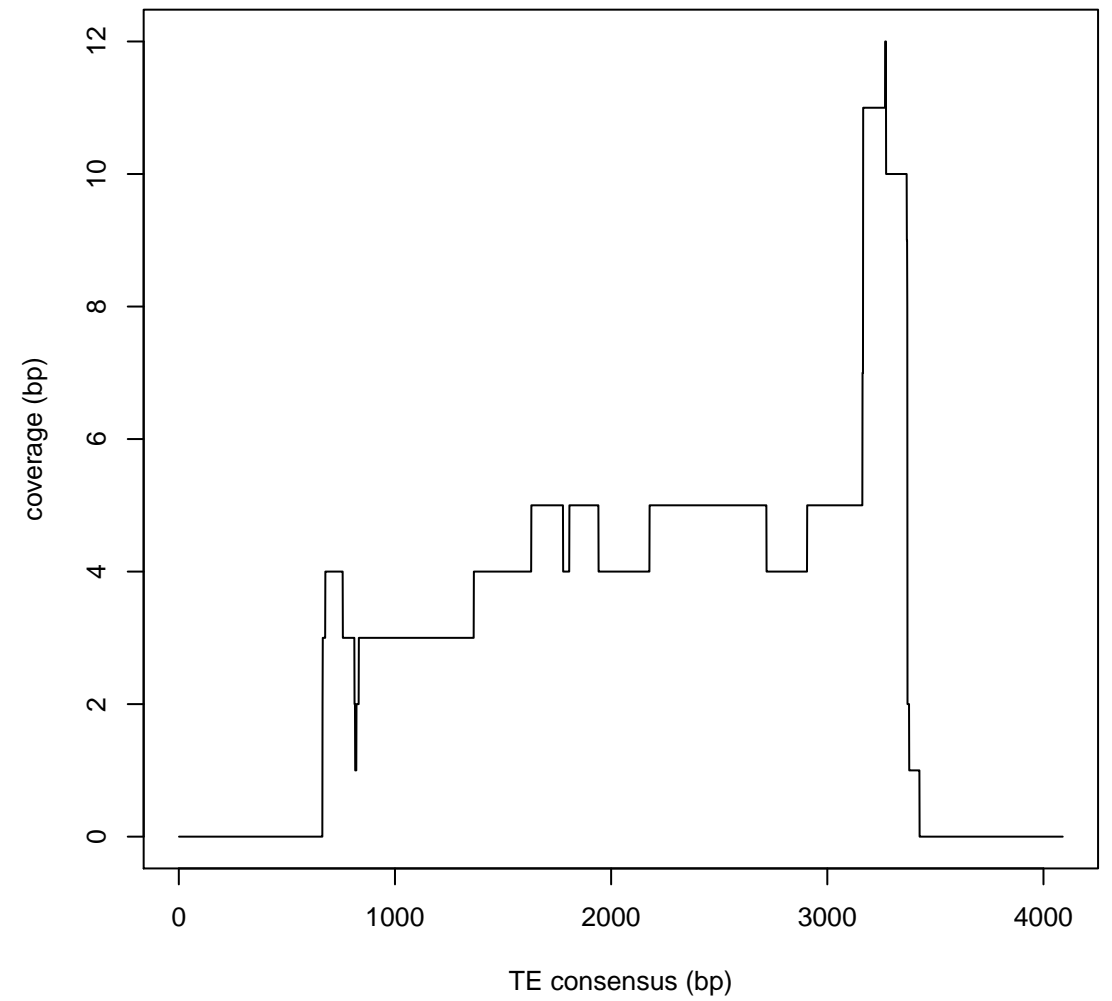


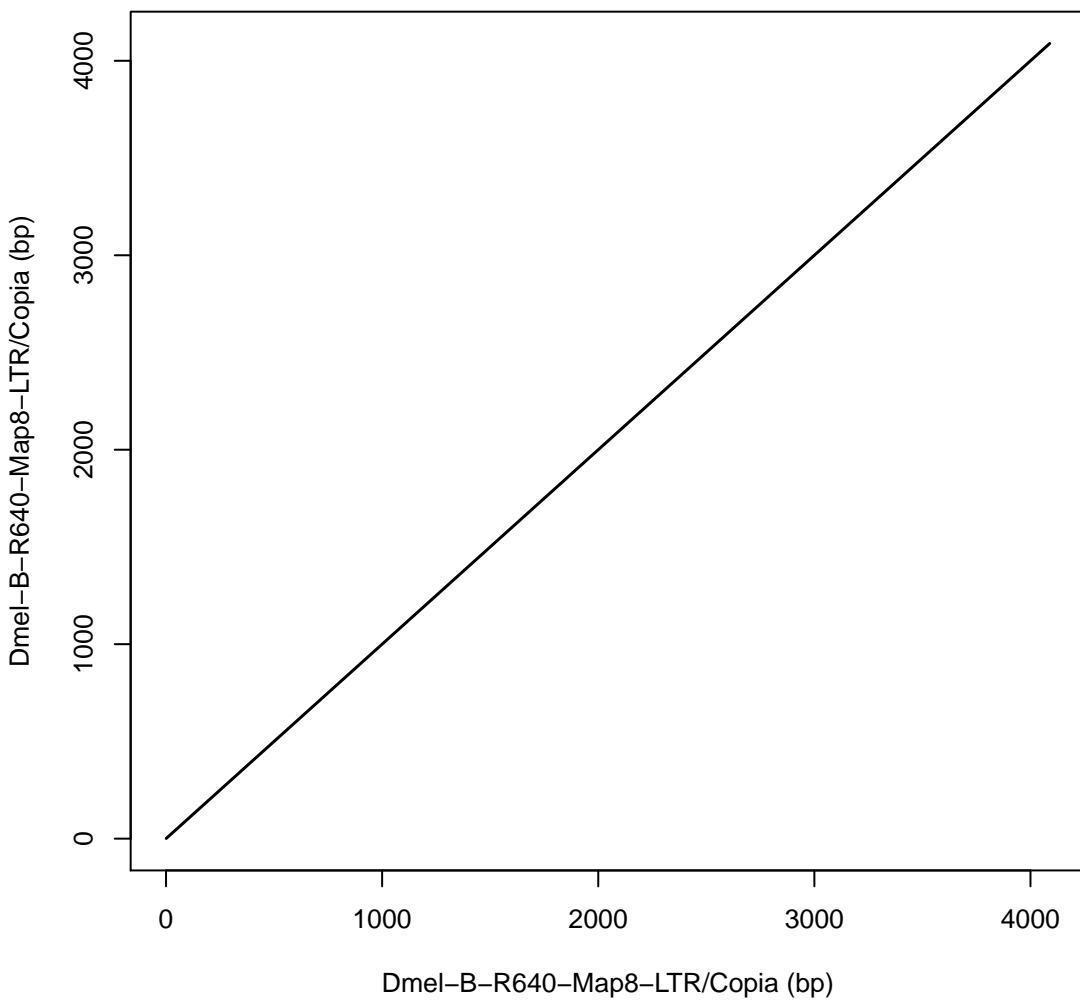
TE: Dmel-B-R640-Map8-LTR/Copia
consensus size: 4089bp; fragments: 20; full length: 0 (>=3680.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

