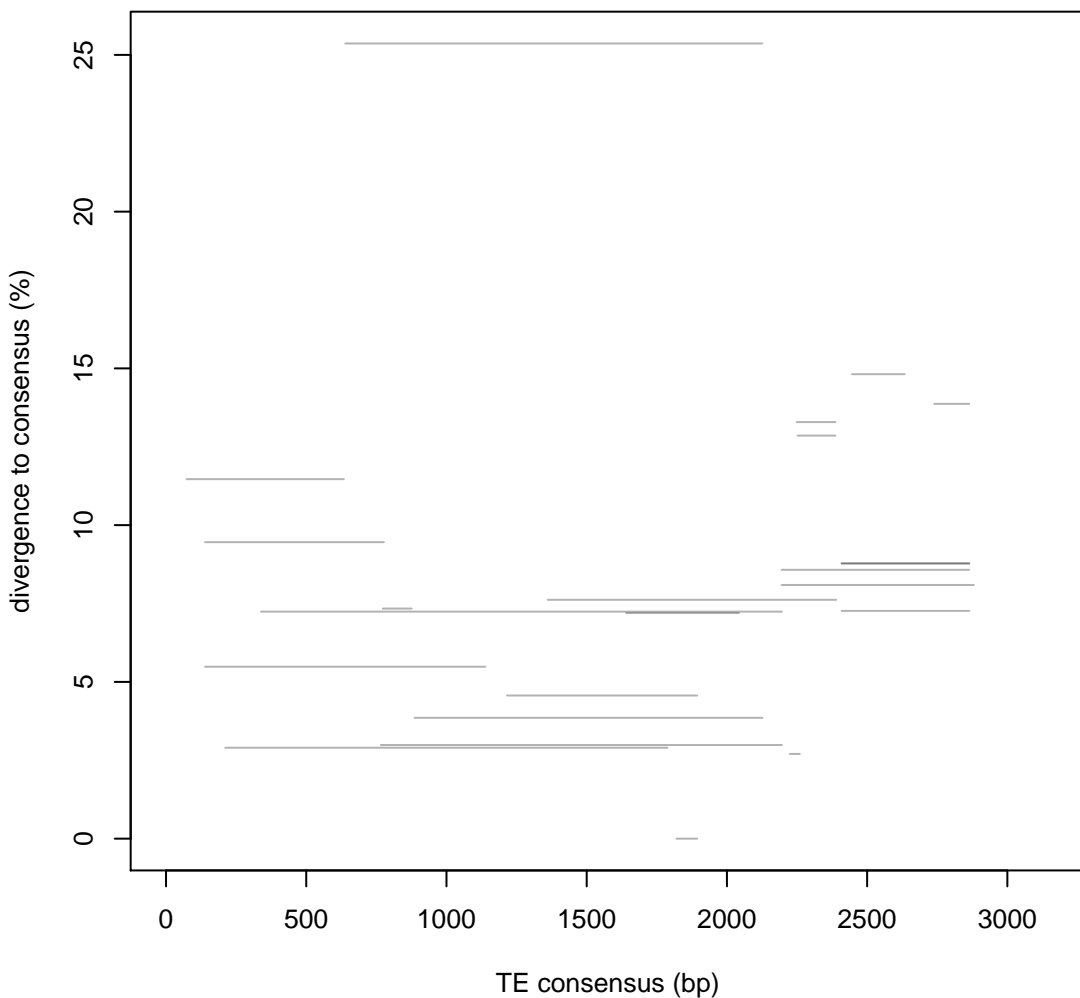
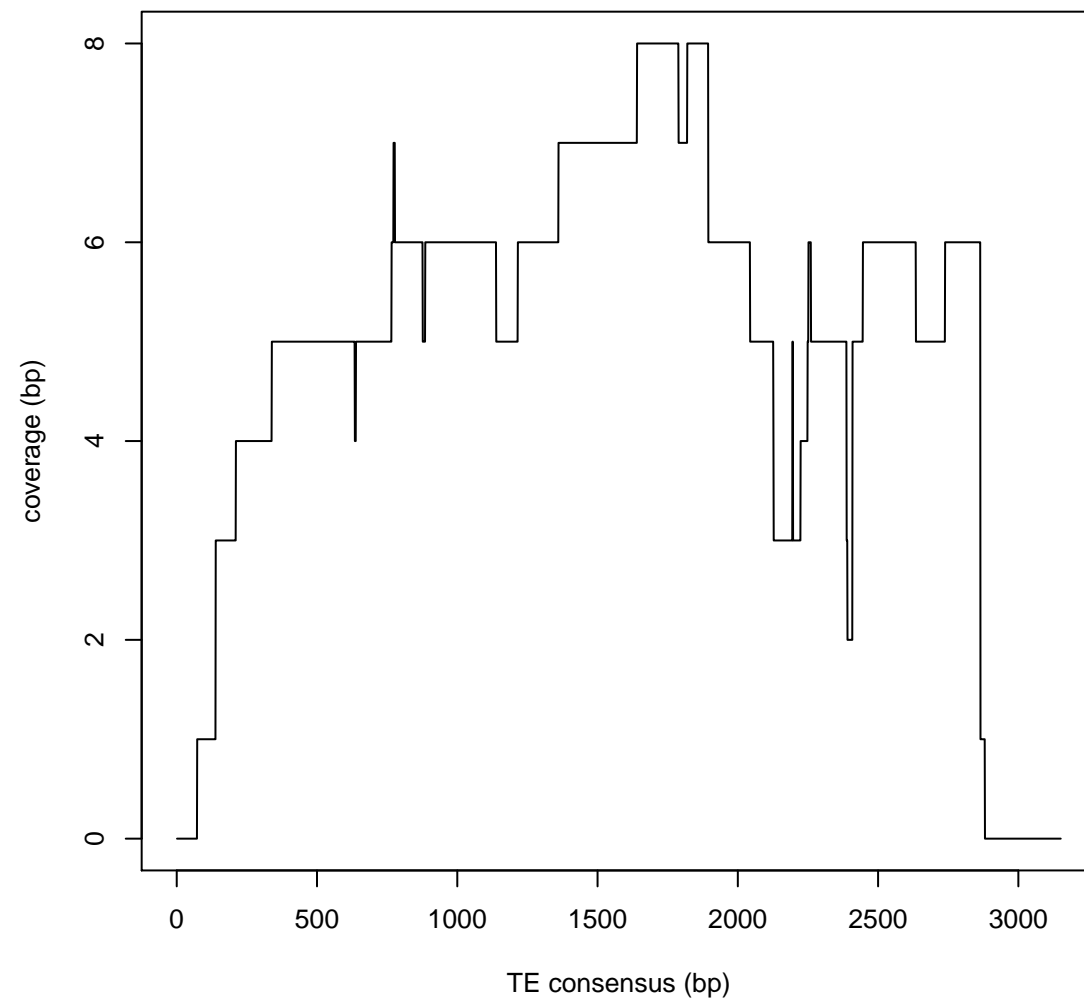


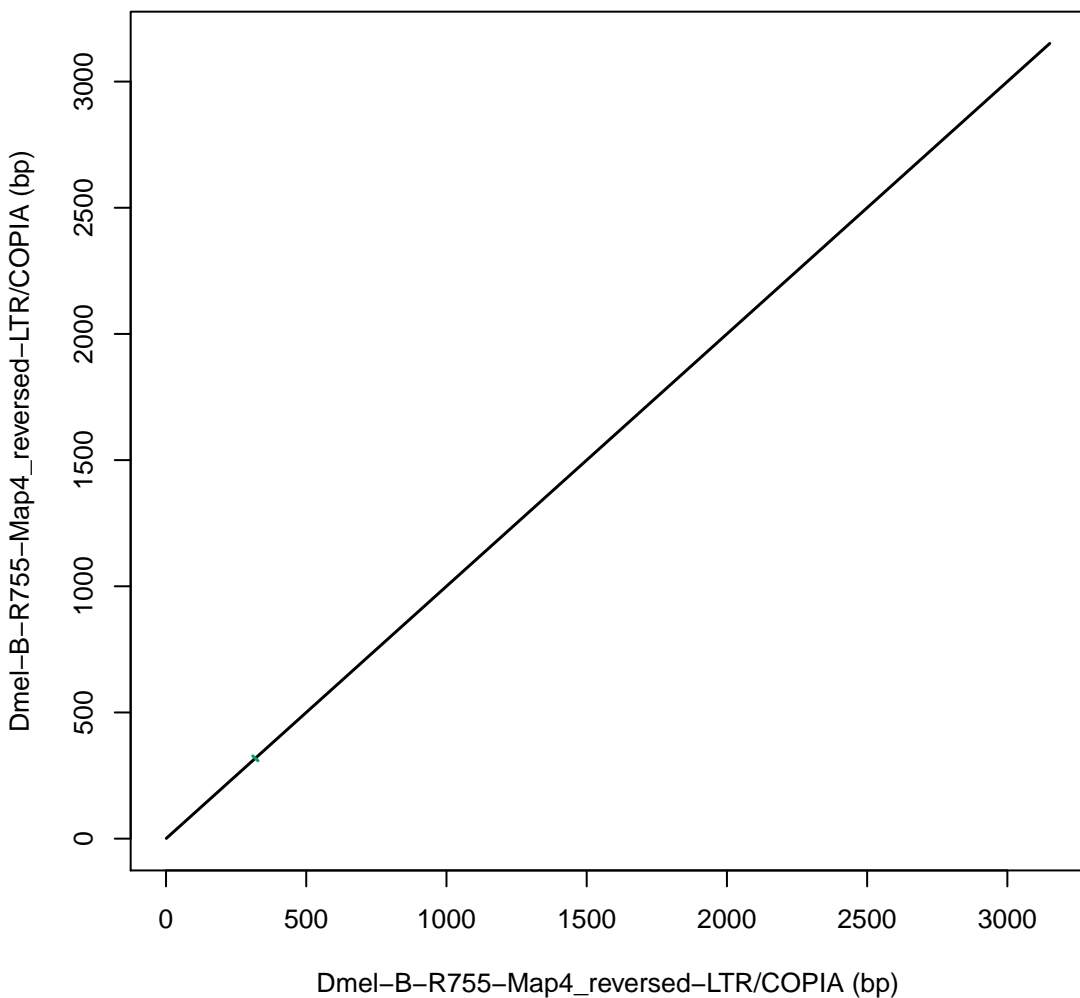
TE: Dmel-B-R755-Map4_reversed-LTR/COPIA
consensus size: 3151bp; fragments: 23; full length: 0 (>=2835.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

