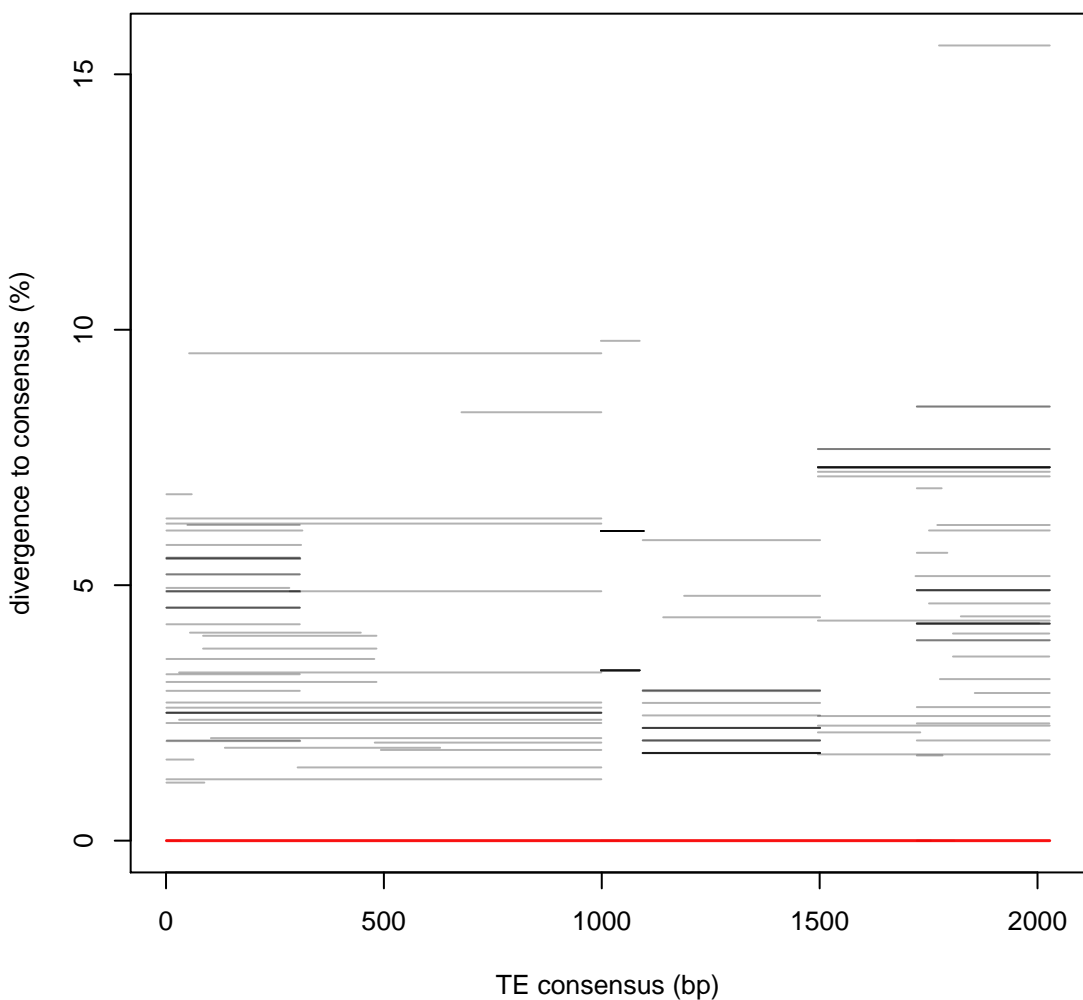
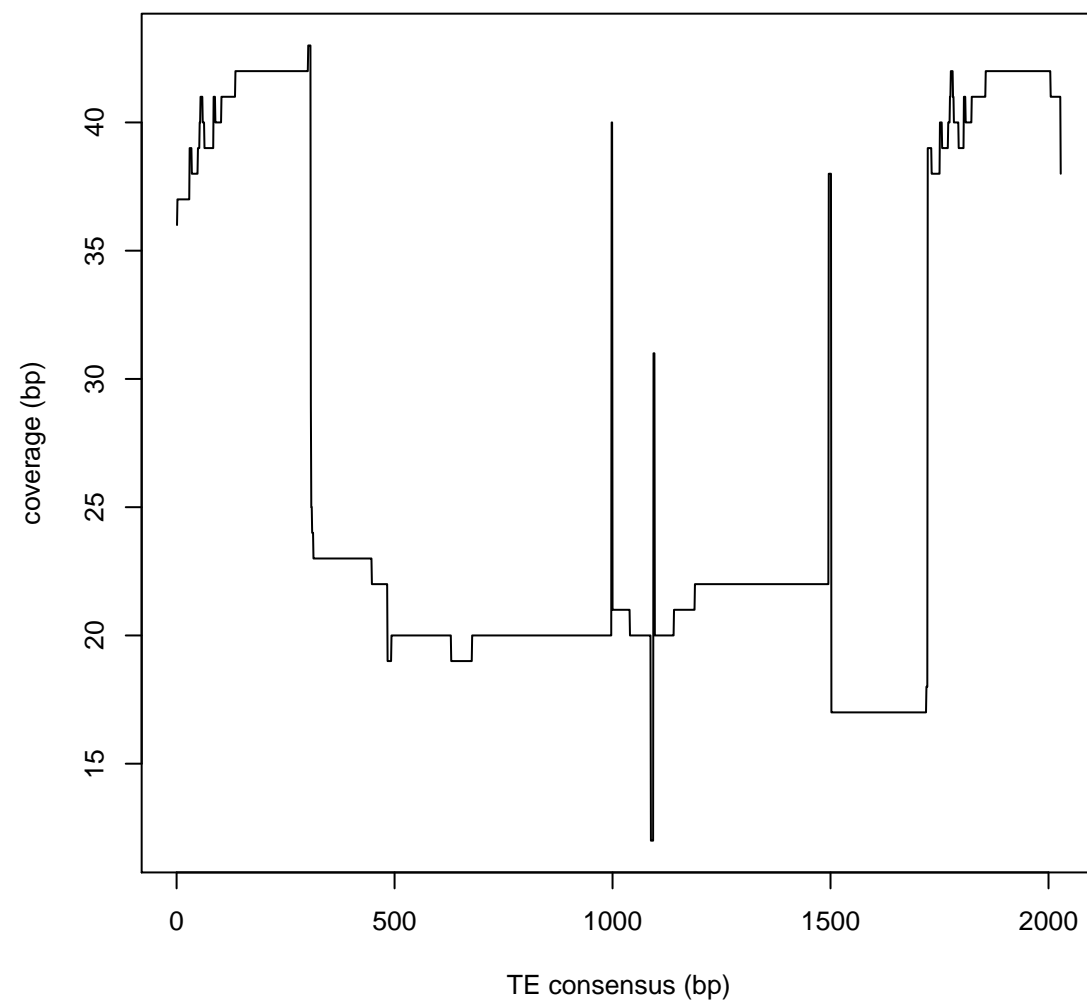


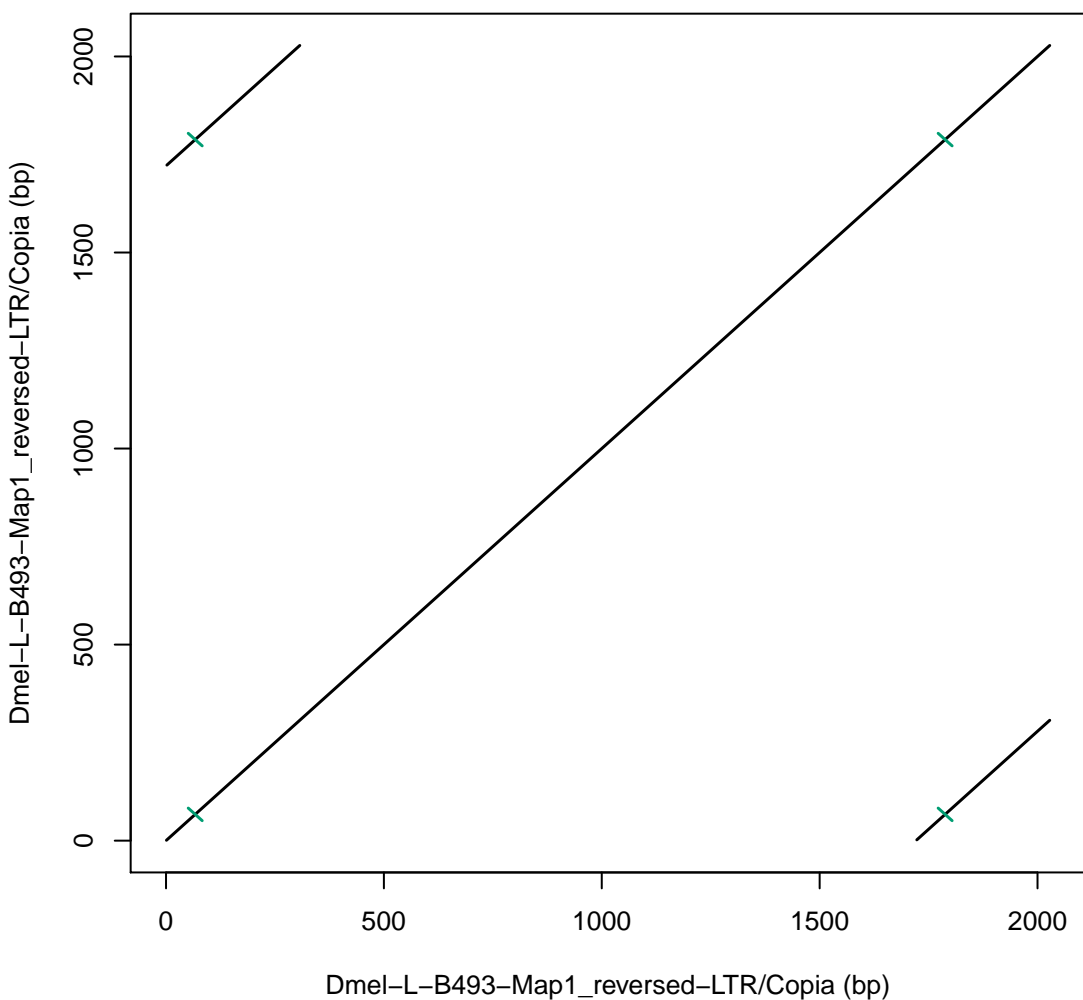
TE: Dmel-L-B493-Map1_reversed-LTR/Copia
consensus size: 2028bp; fragments: 139; full length: 1 (>=1825.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

