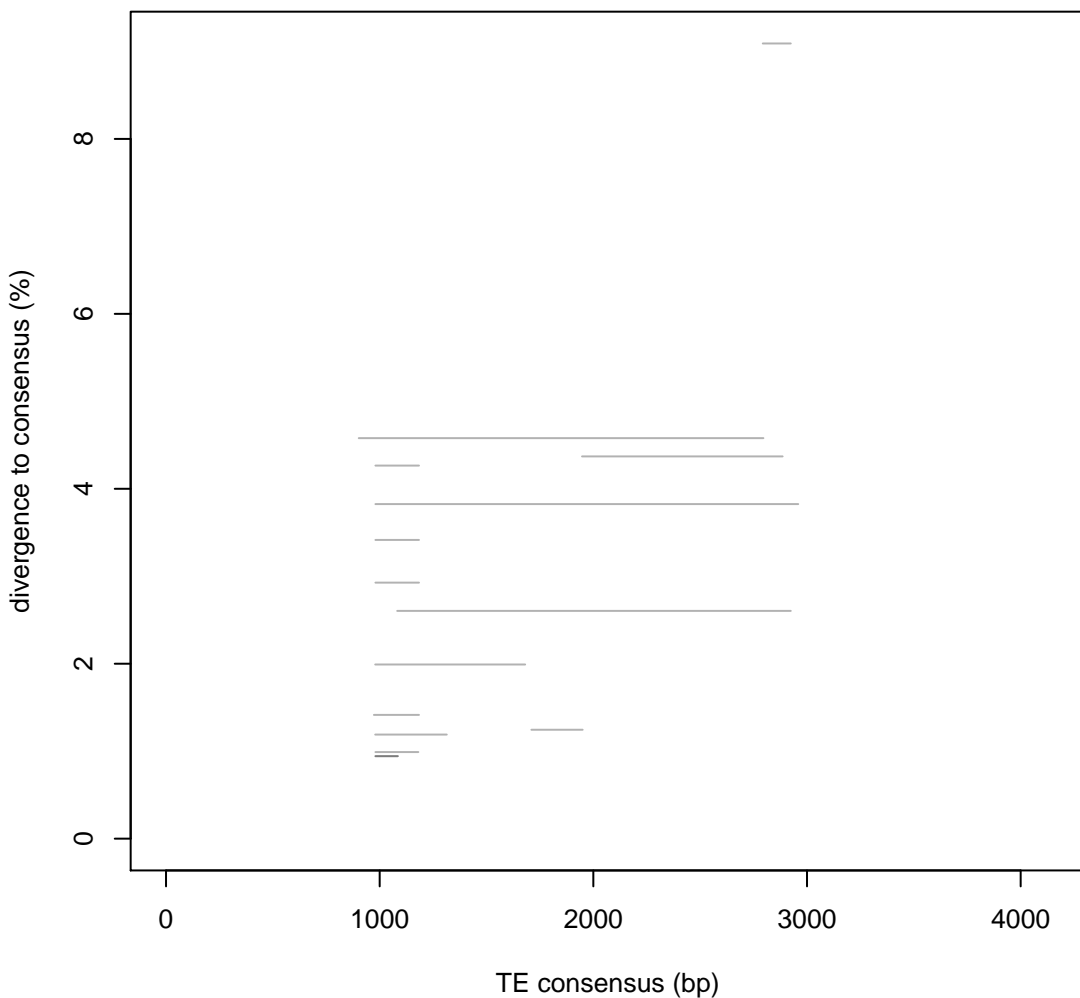
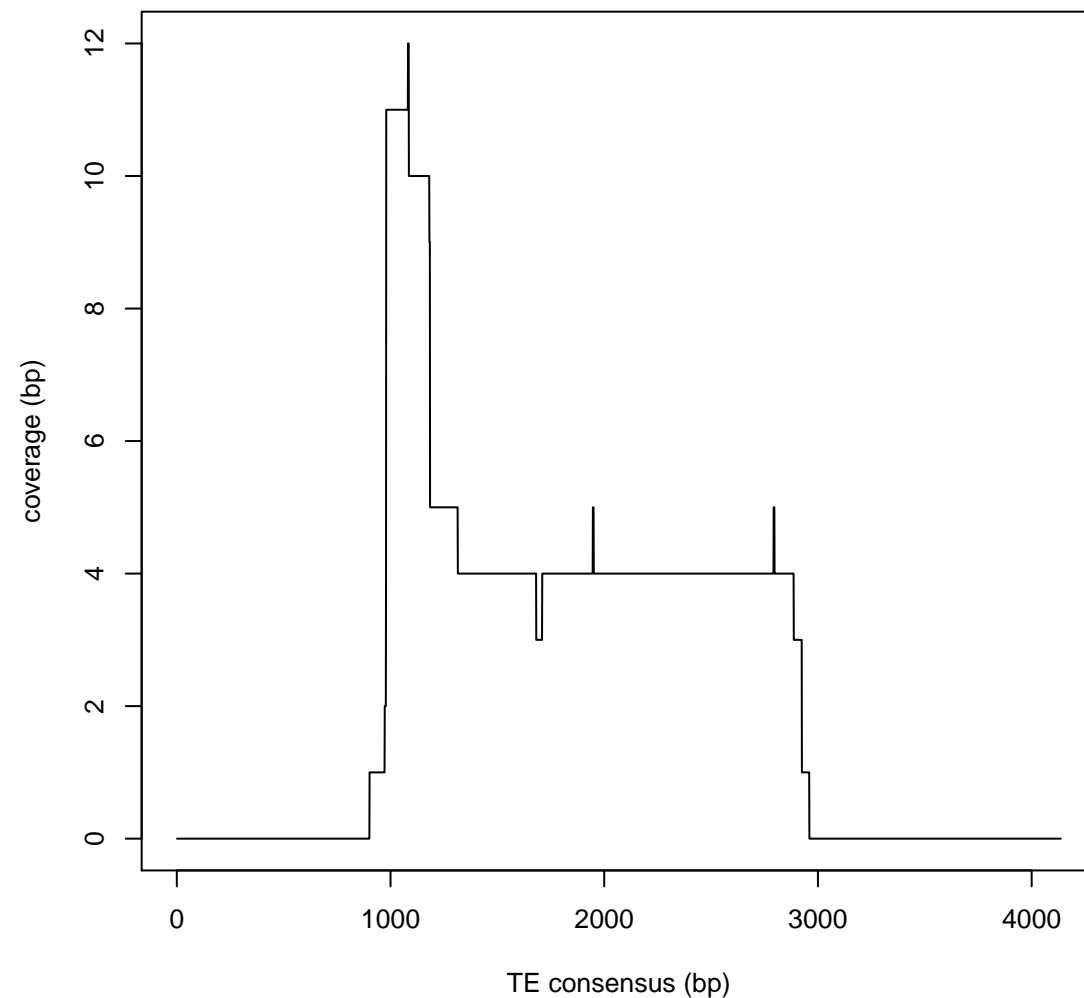


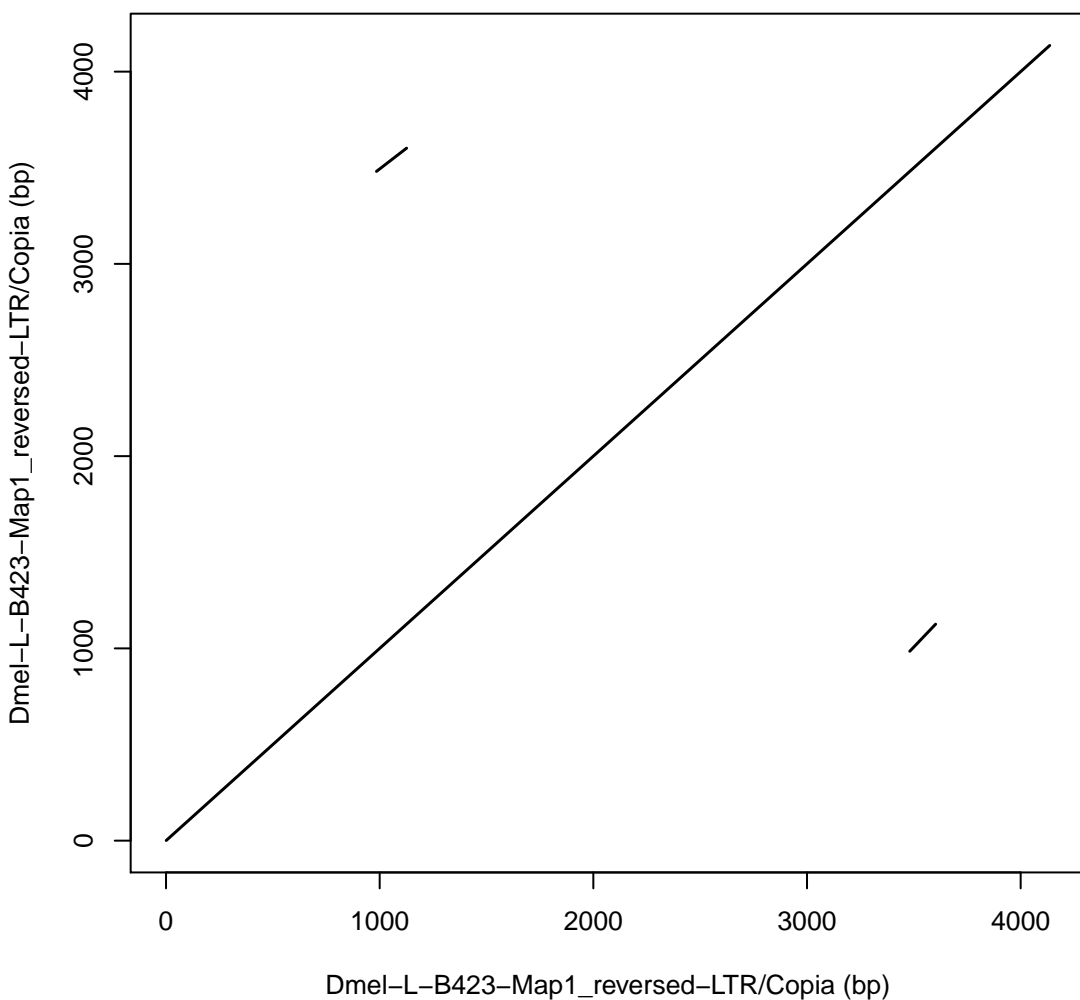
TE: Dmel-L-B423-Map1_reversed-LTR/Copia
consensus size: 4136bp; fragments: 15; full length: 0 (>=3722.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

