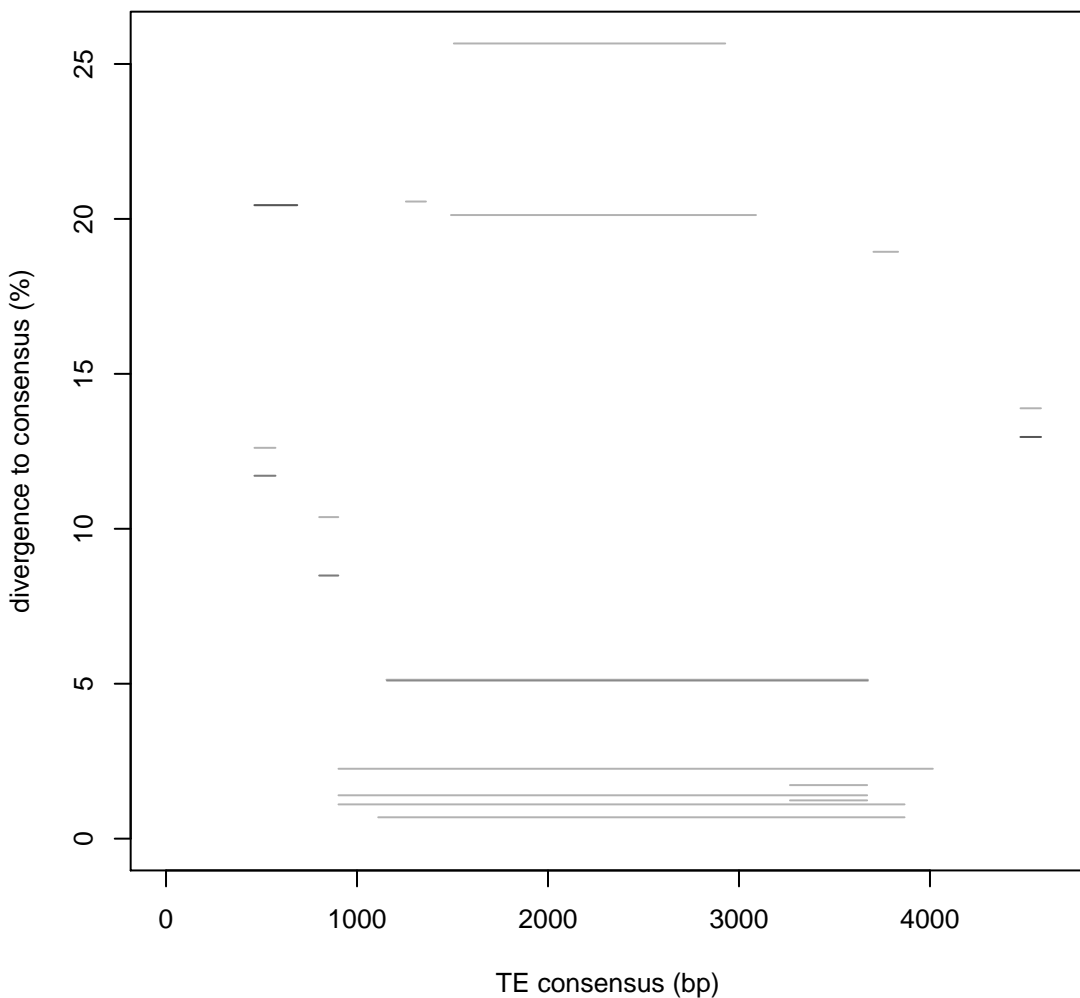
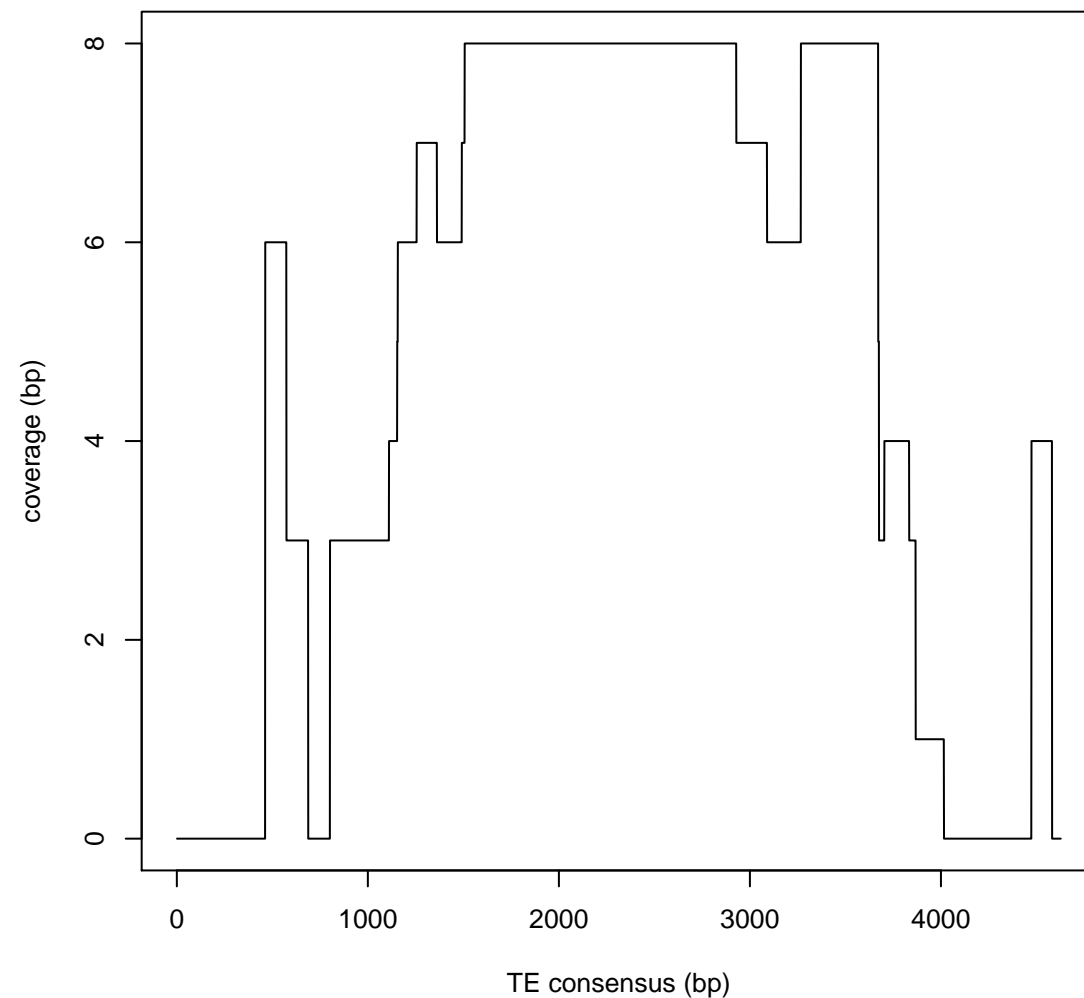


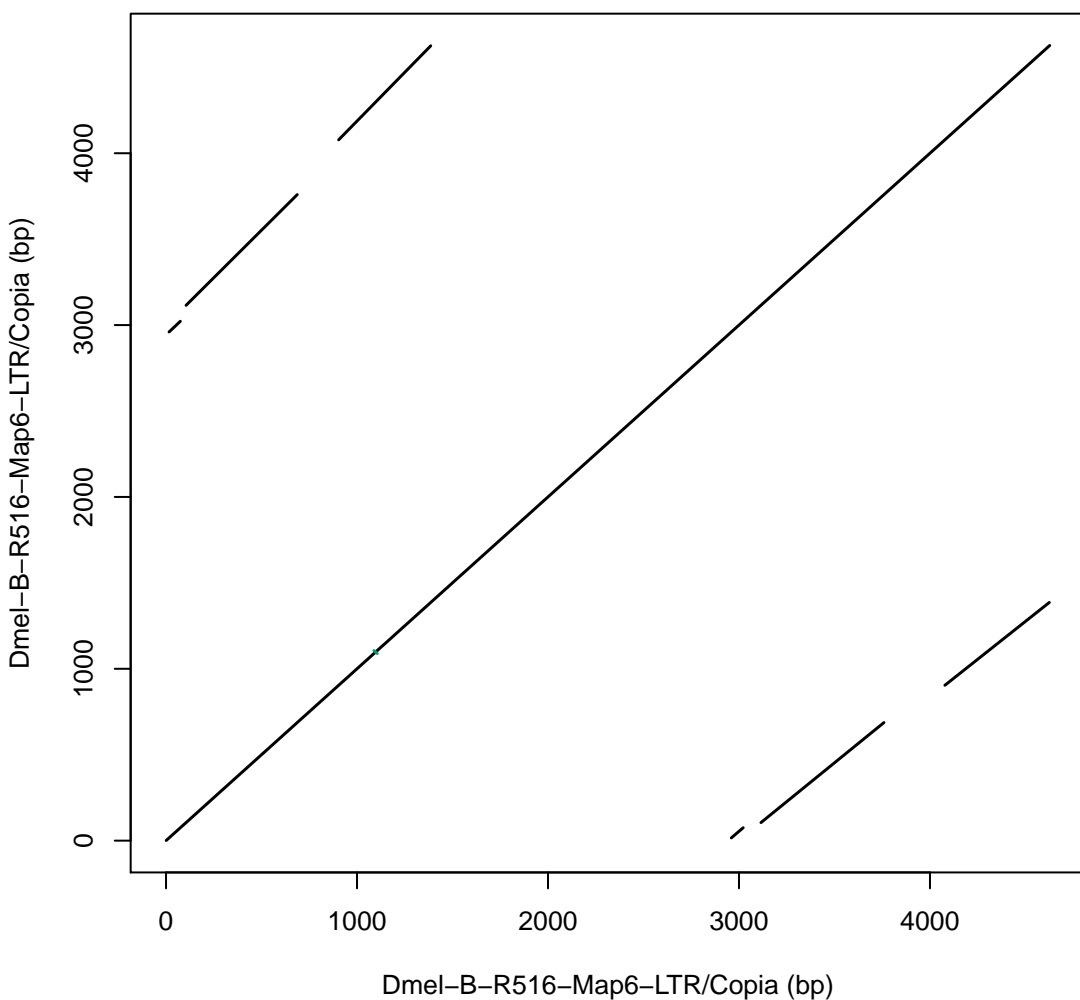
TE: Dmel-B-R516-Map6-LTR/Copia
consensus size: 4627bp; fragments: 25; full length: 0 (≥ 4164.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

