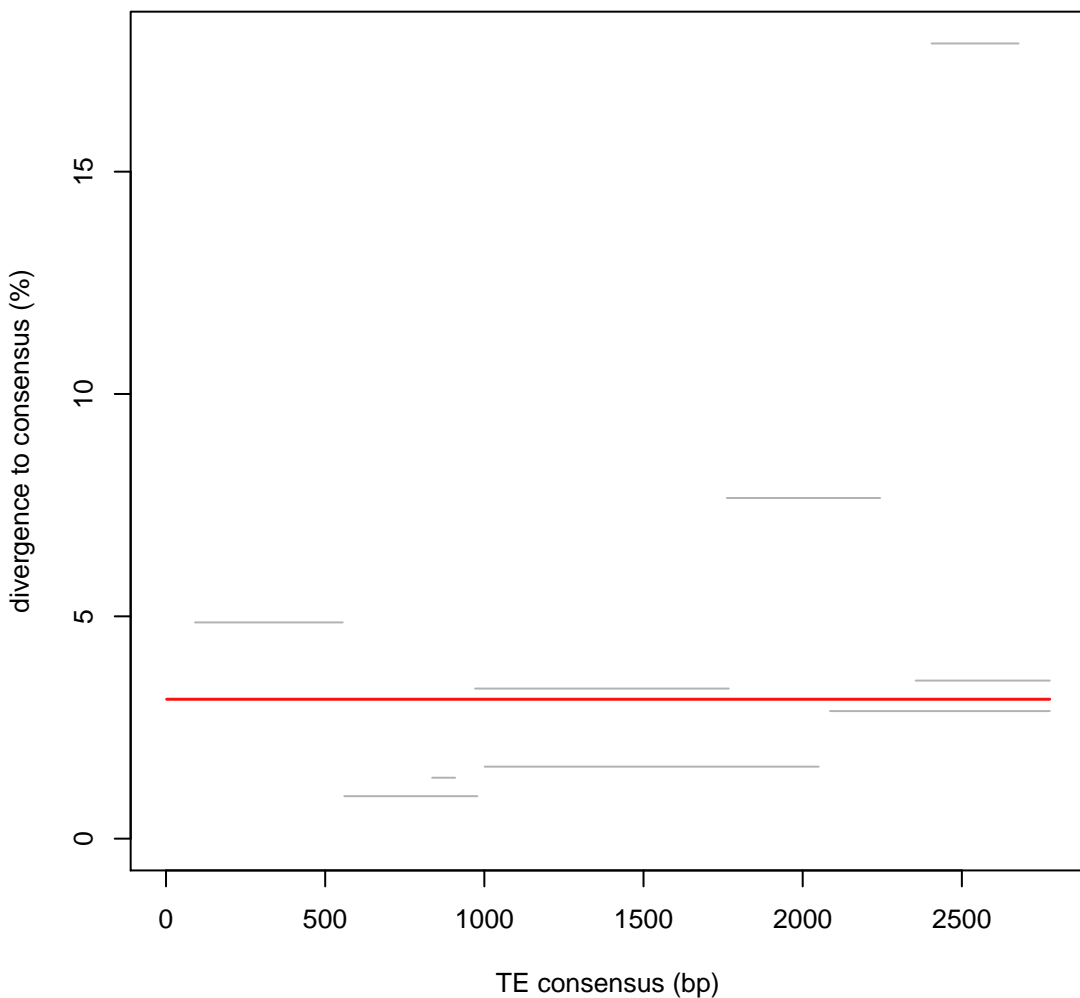
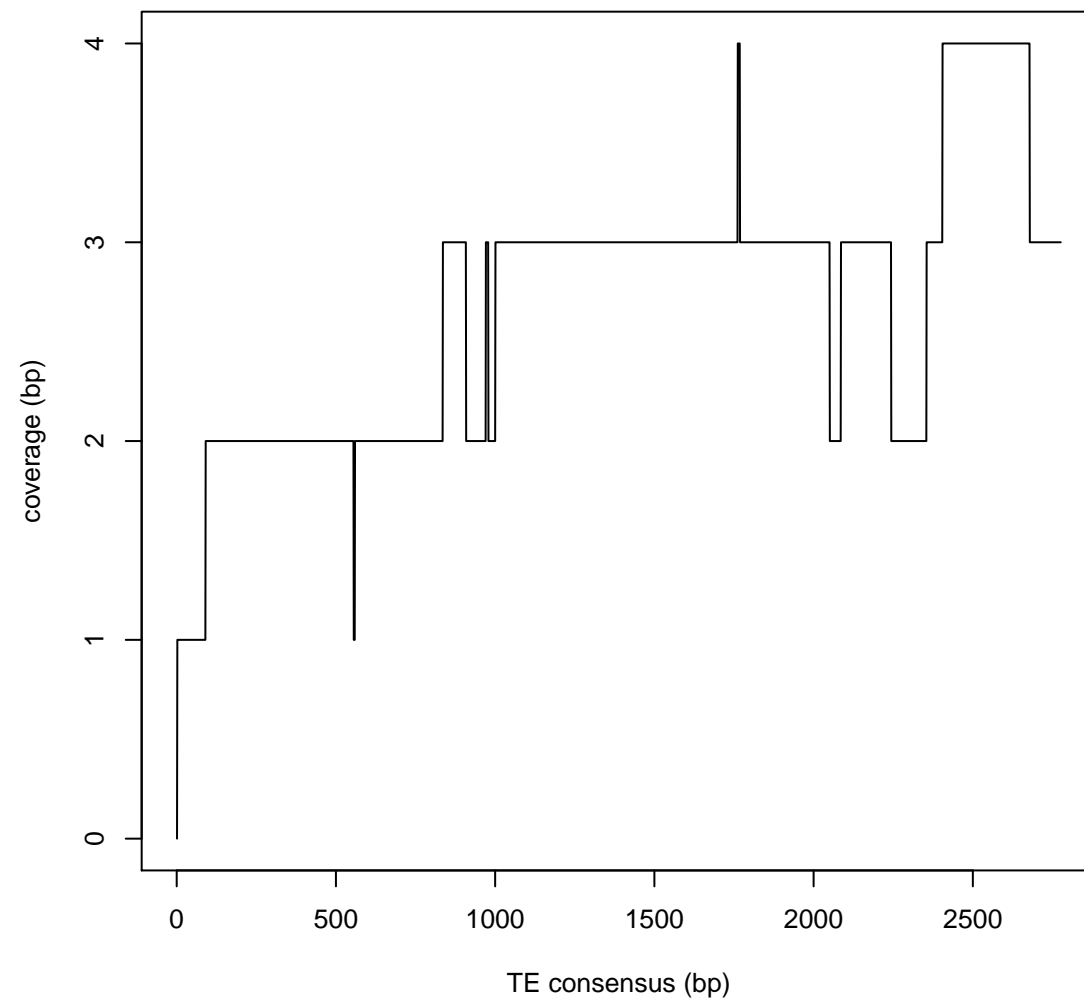


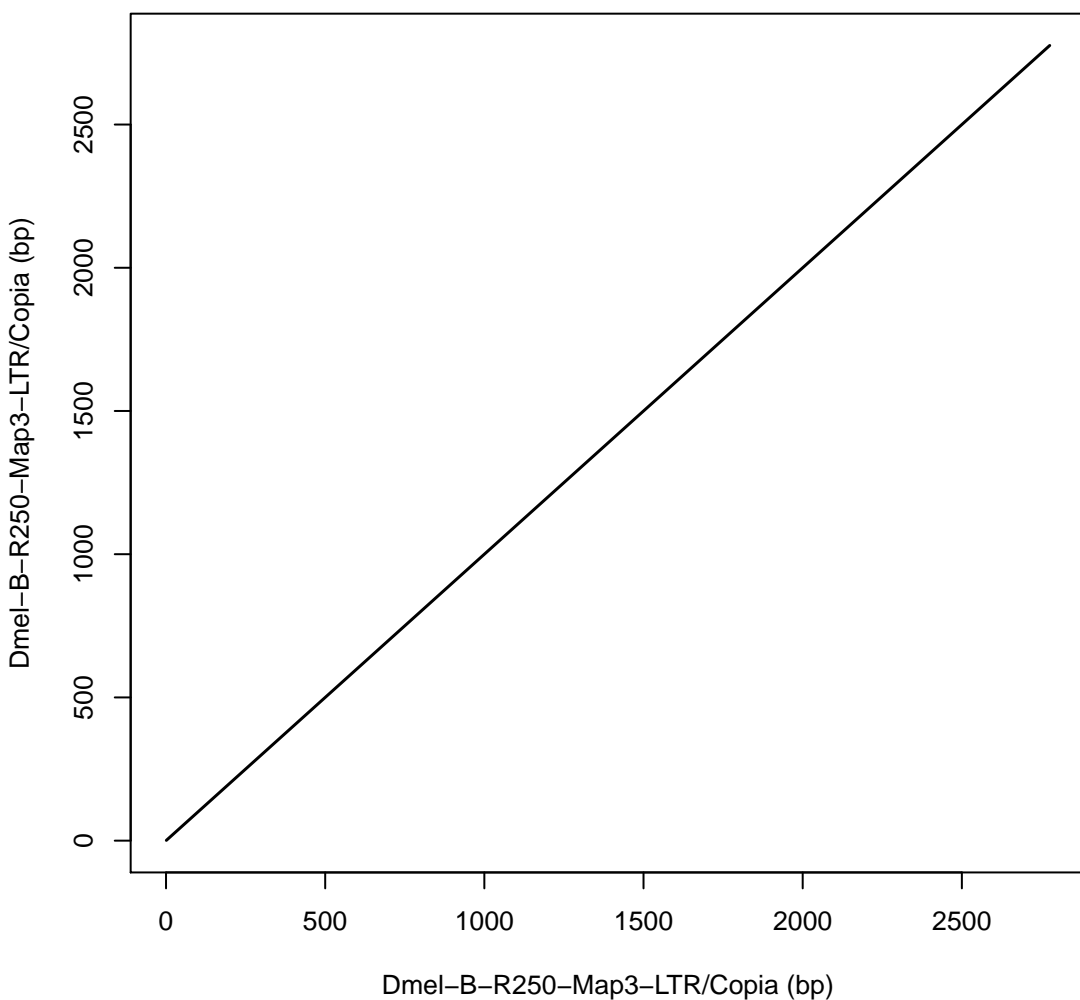
TE: Dmel-B-R250-Map3-LTR/Copia
consensus size: 2776bp; fragments: 10; full length: 1 (>=2498.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

