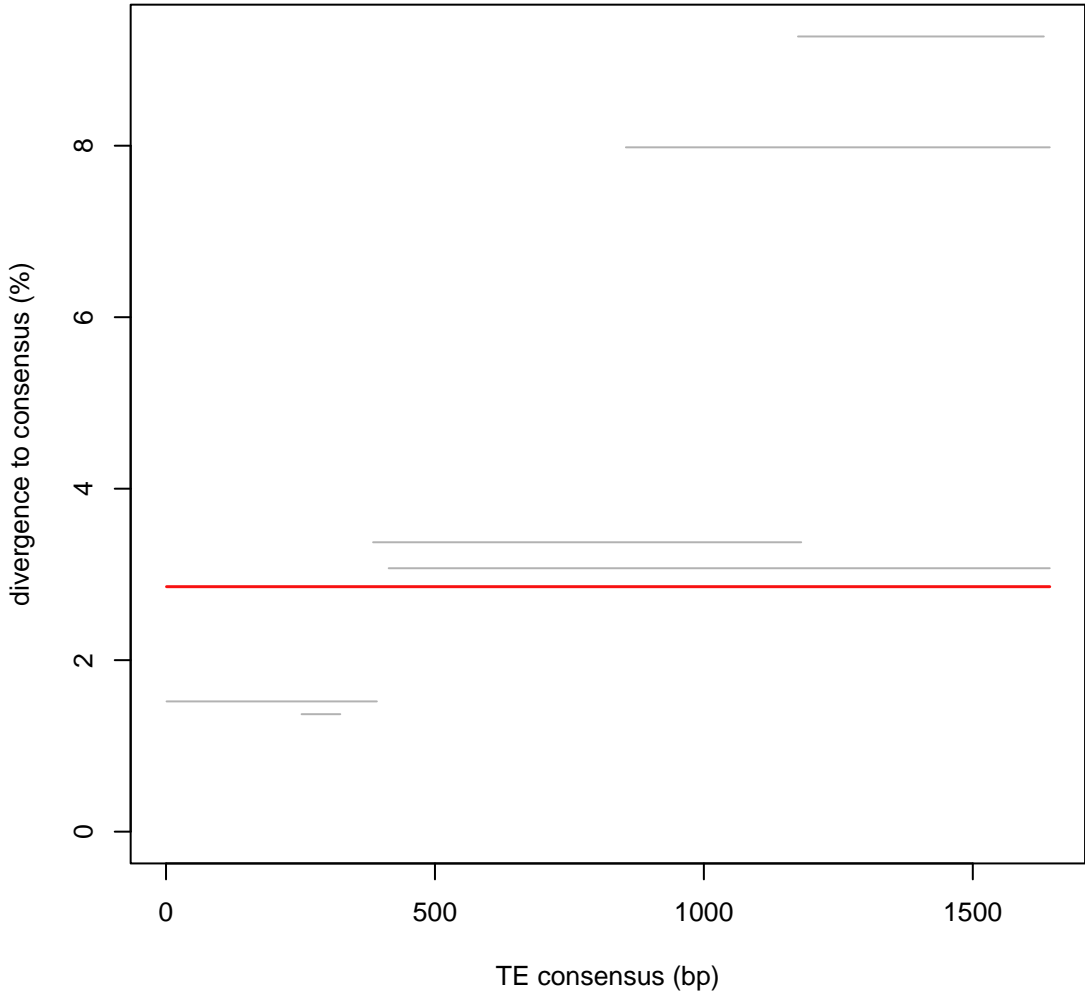
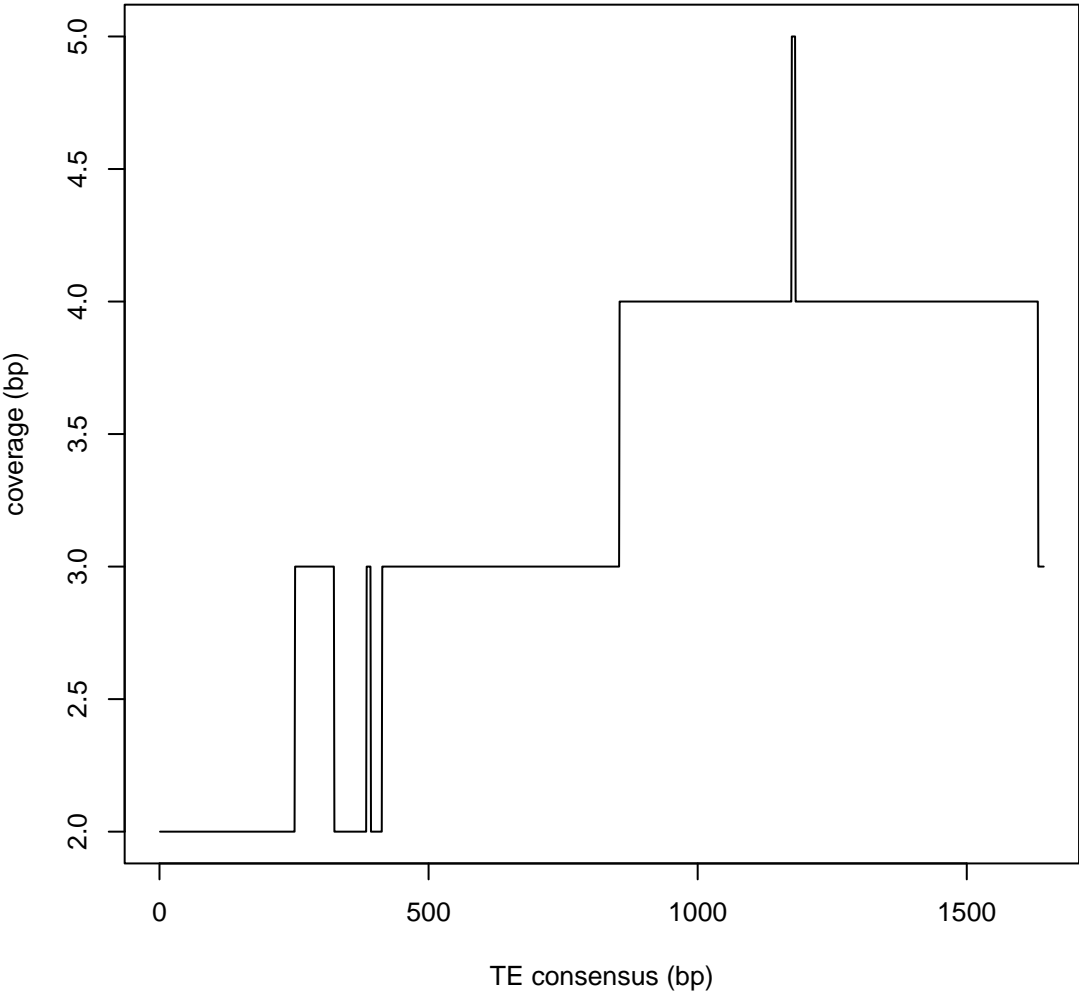


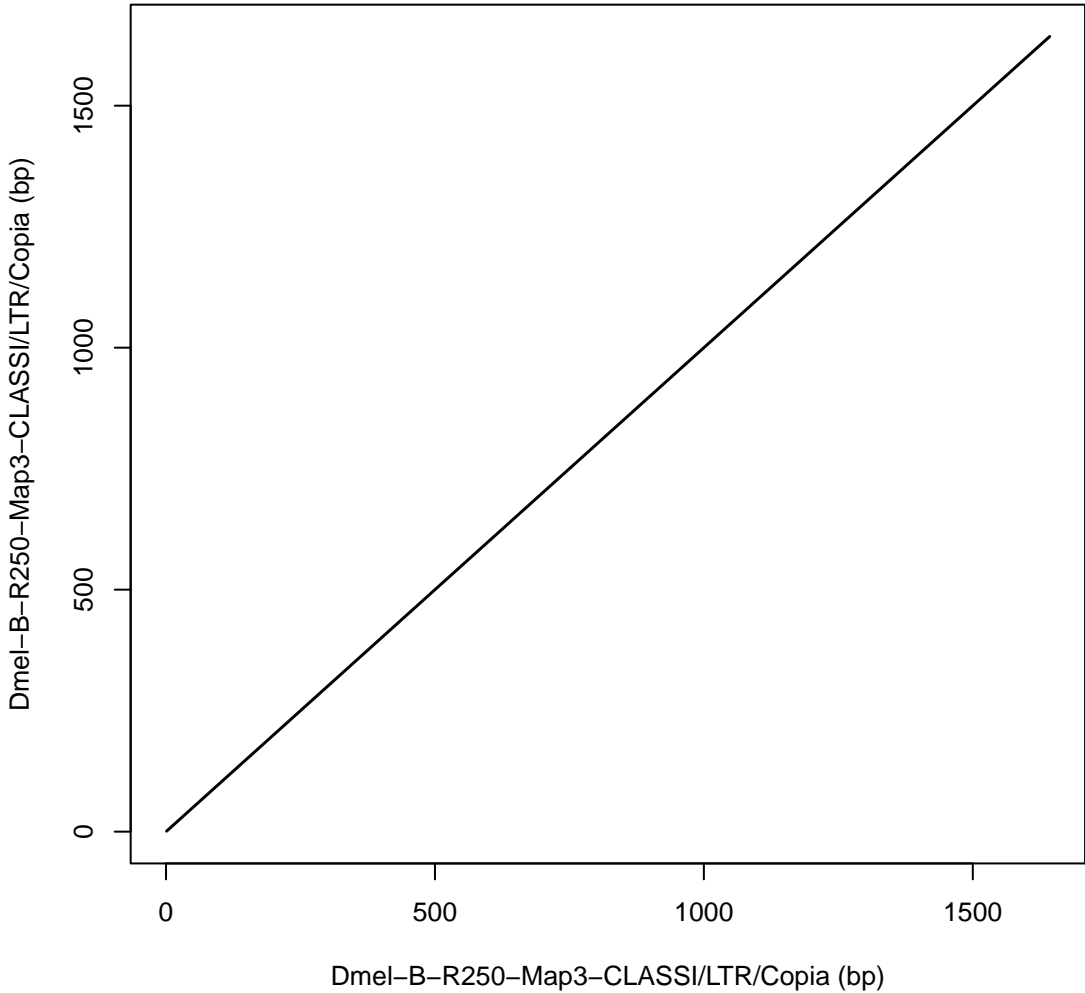
TE: Dmel-B-R250-Map3-CLASSI/LTR/Copia
consensus size: 1643bp; fragments: 7; full length: 1 (>=1478.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

