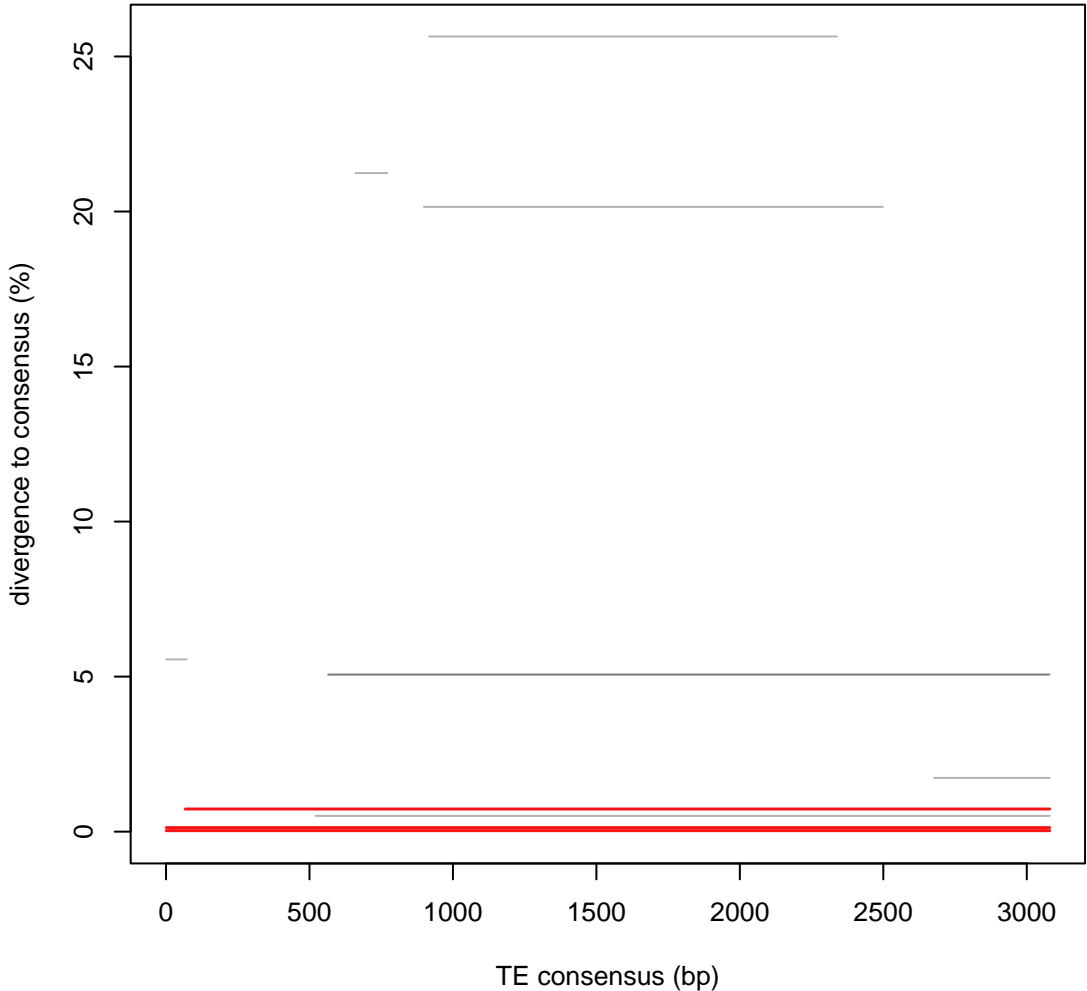
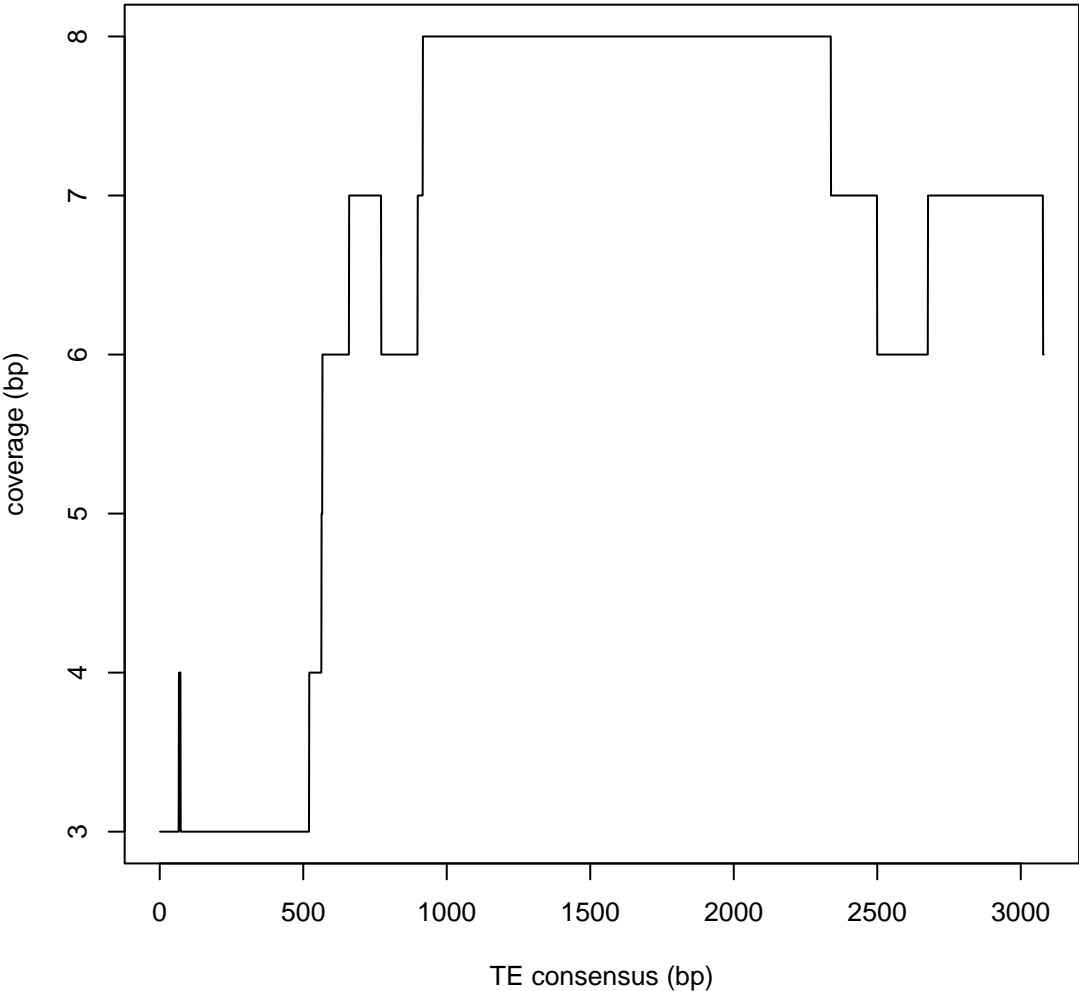


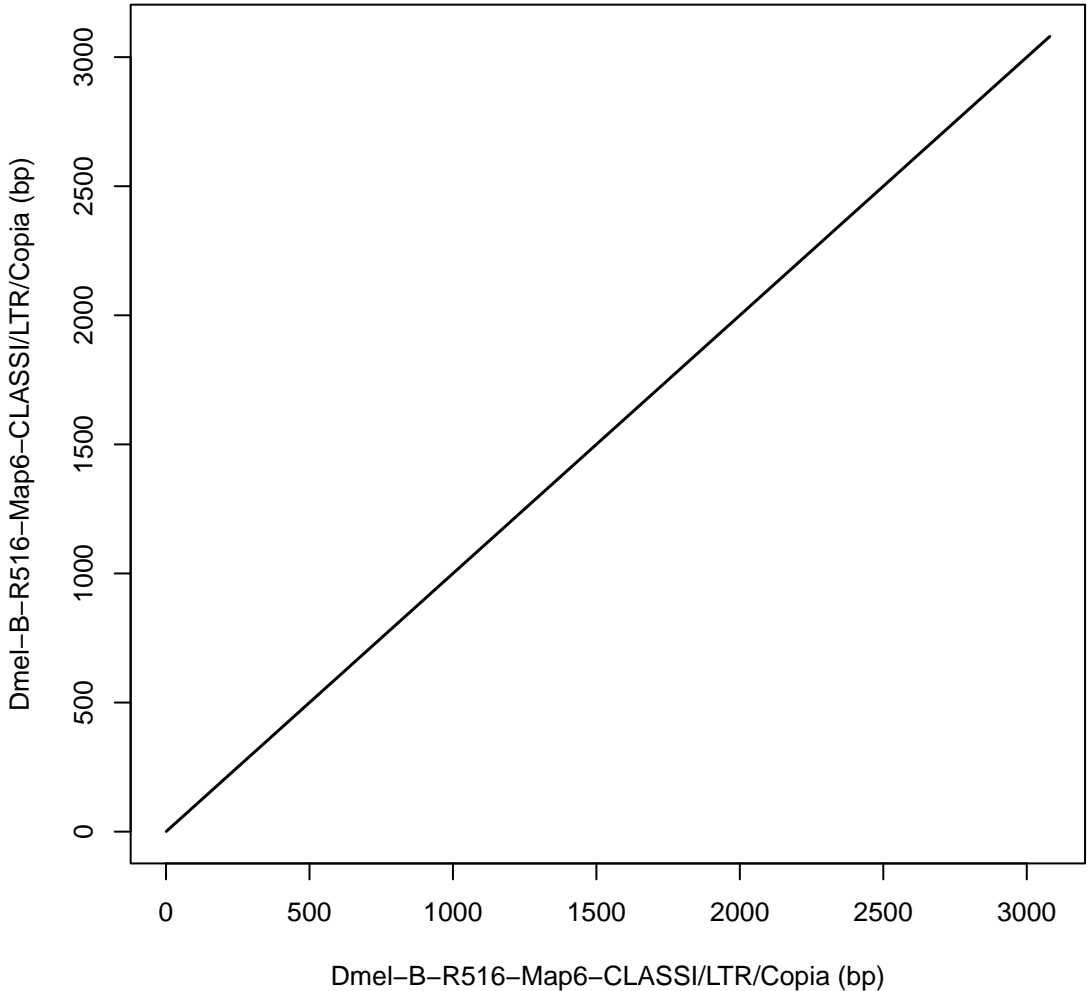
TE: Dmel-B-R516-Map6-CLASSI/LTR/Copia  
consensus size: 3080bp; fragments: 11; full length: 3 (>=2772bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

