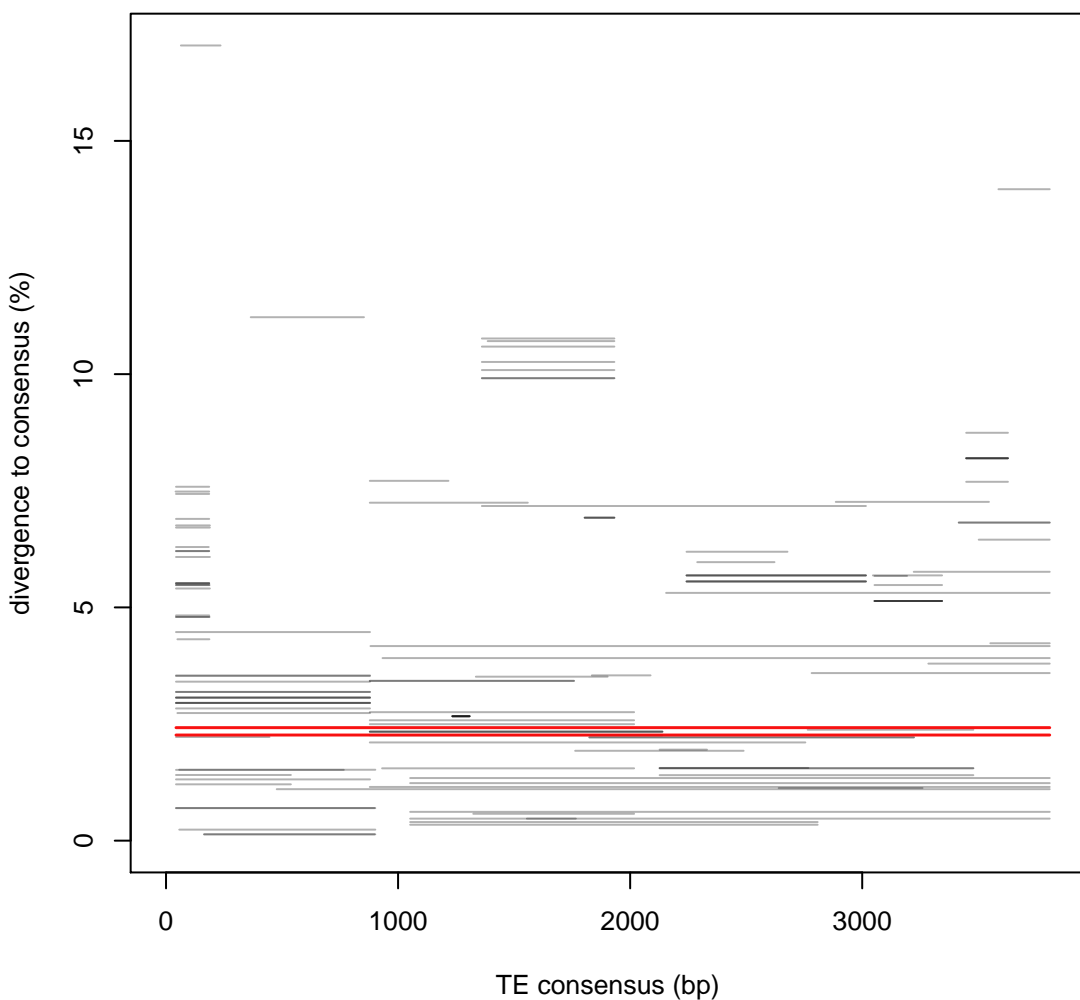
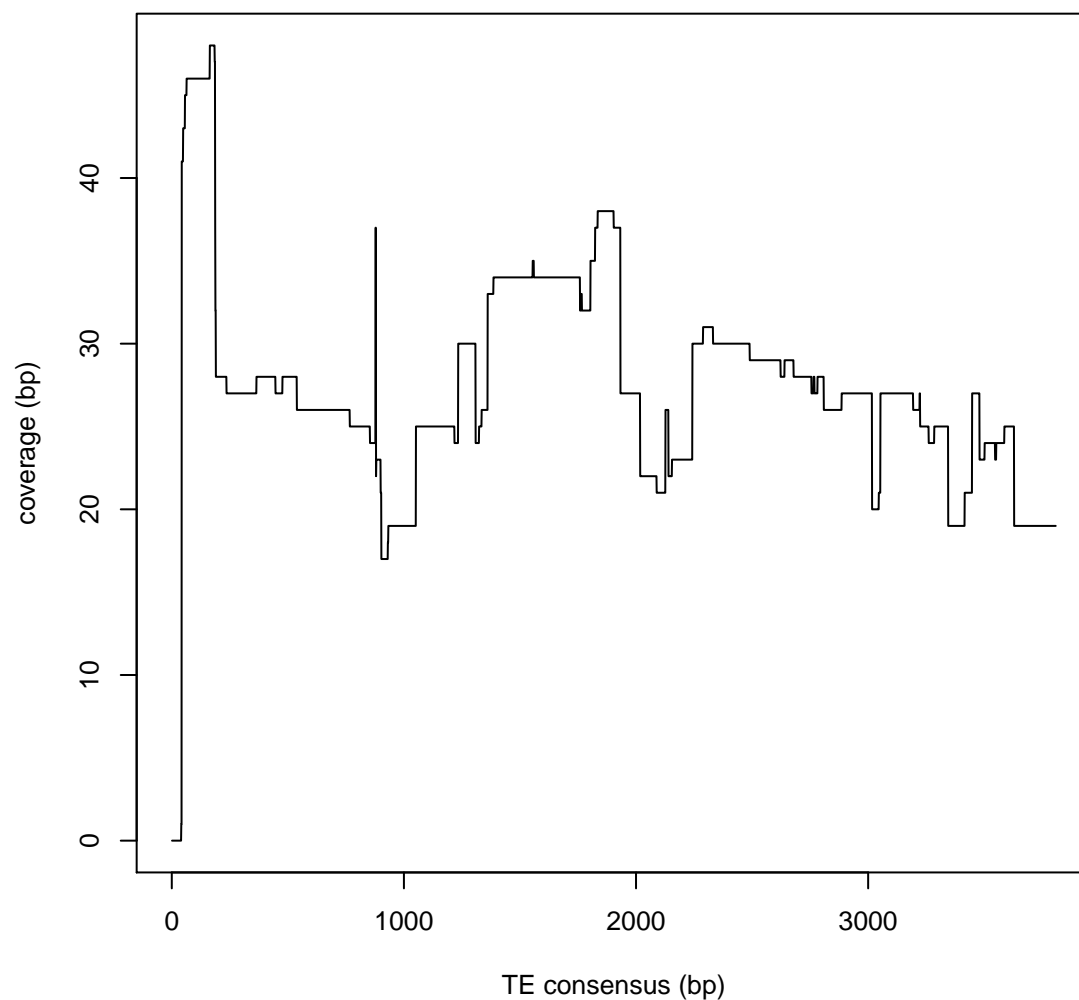


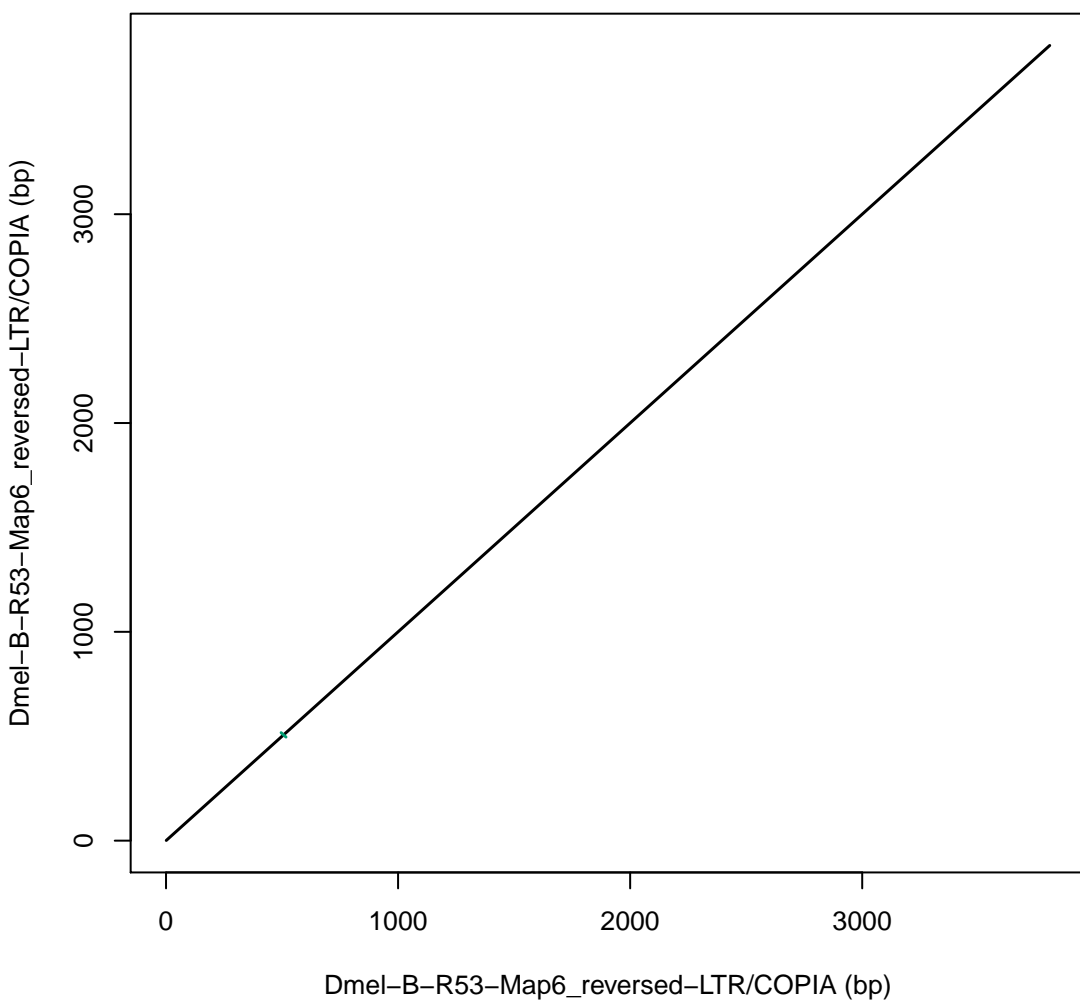
TE: Dmel-B-R53-Map6\_reversed-LTR/COPIA  
consensus size: 3808bp; fragments: 134; full length: 2 (>=3427.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

