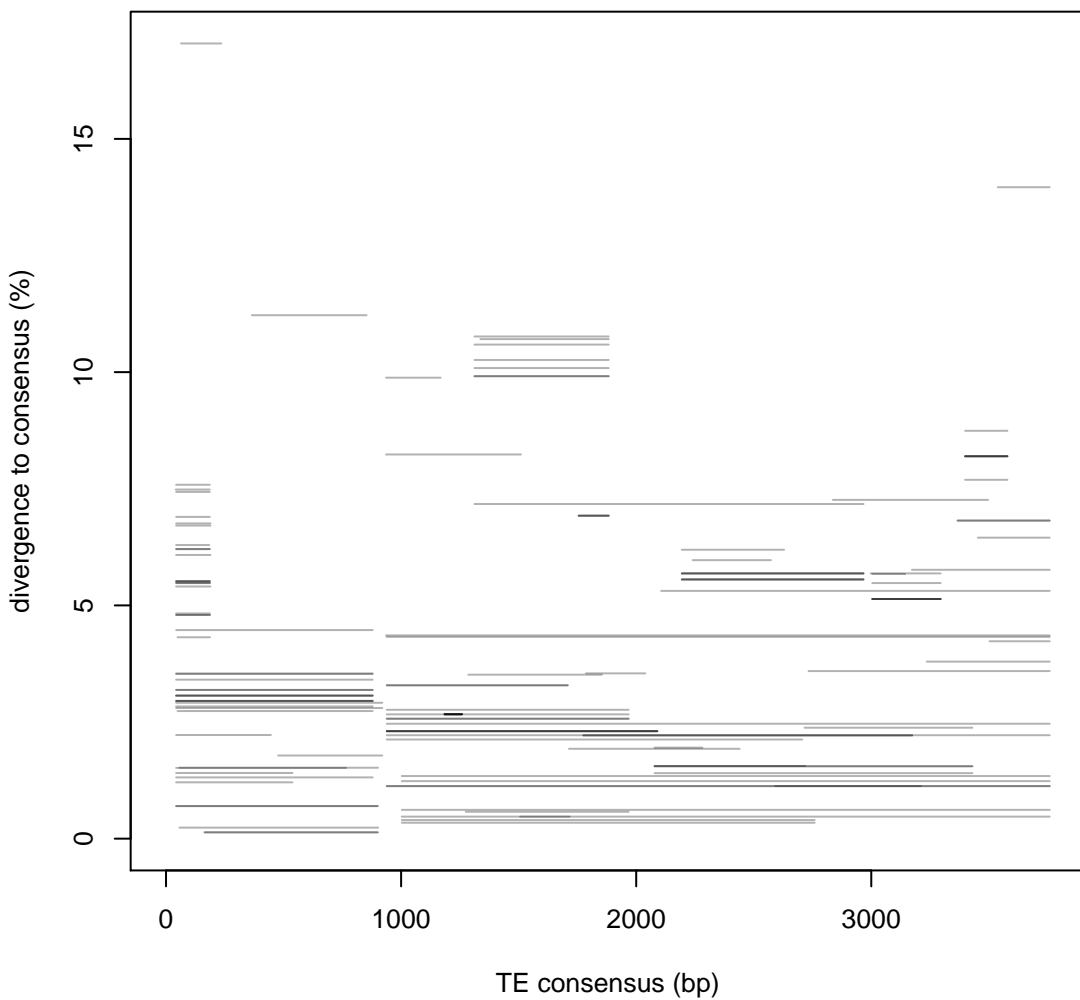
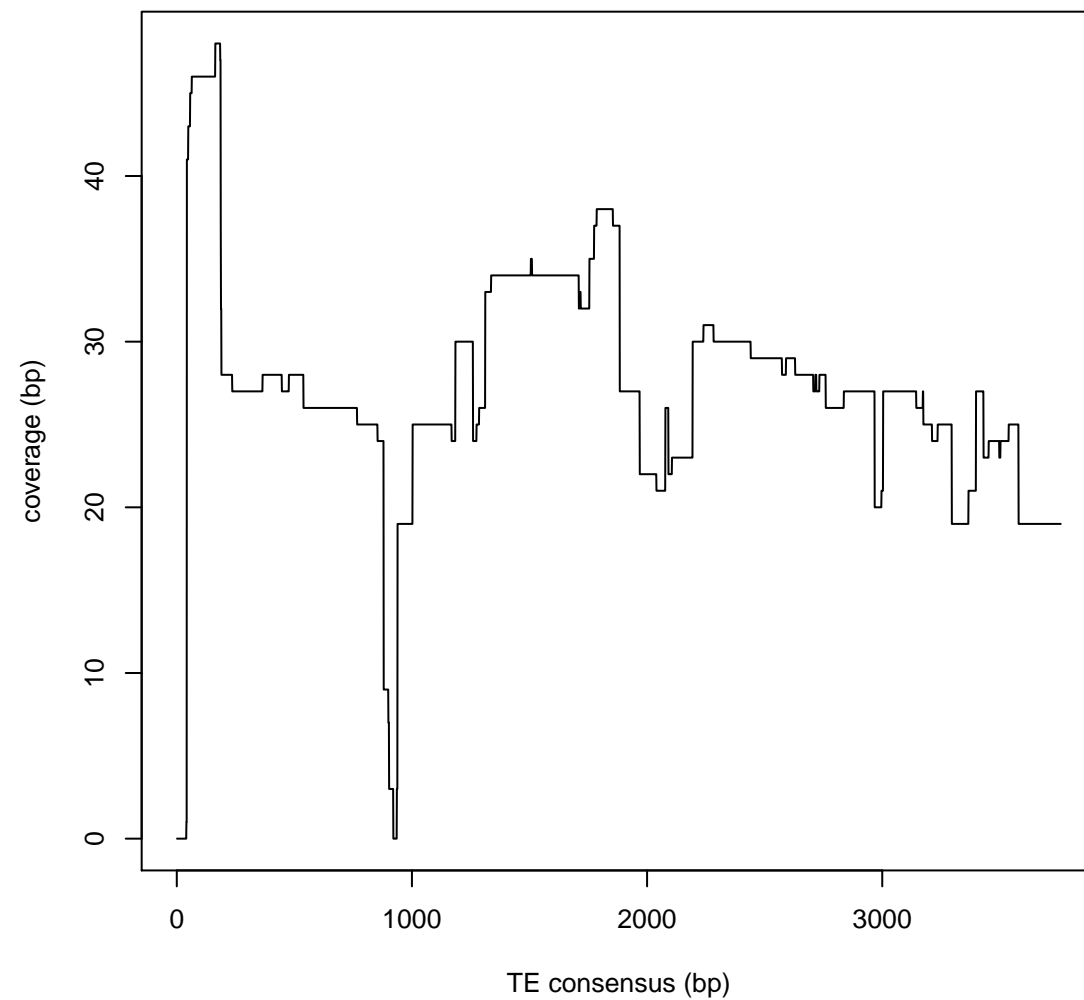


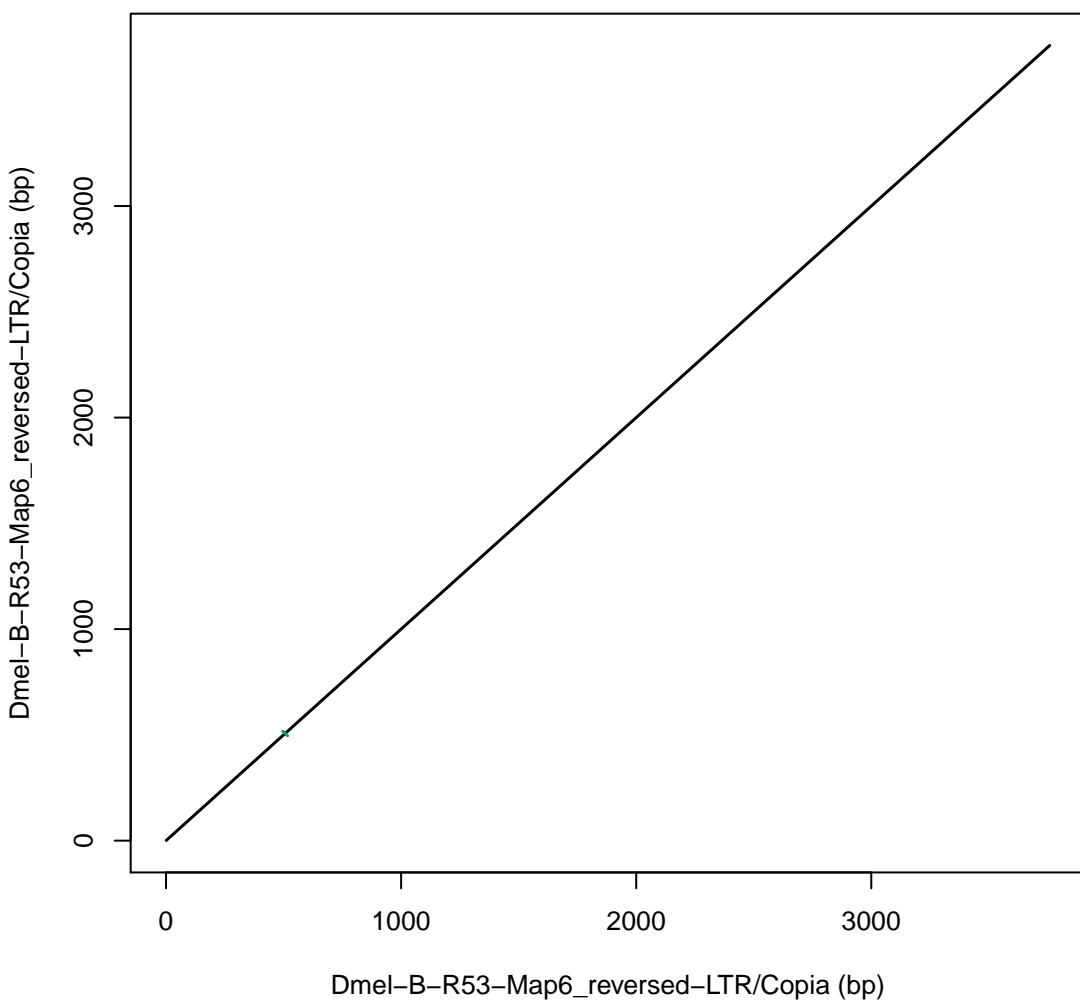
TE: Dmel-B-R53-Map6\_reversed-LTR/Copia  
consensus size: 3759bp; fragments: 137; full length: 0 (>=3383.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

