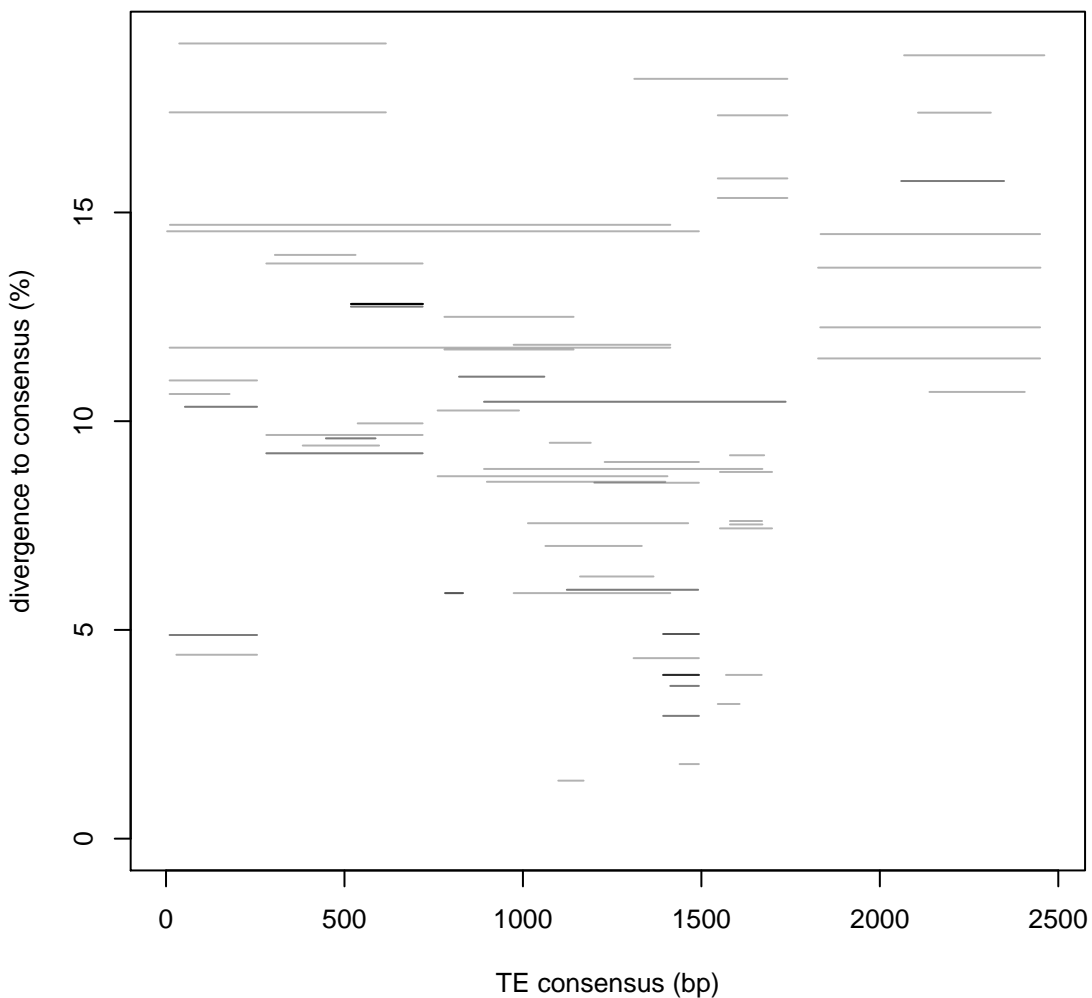
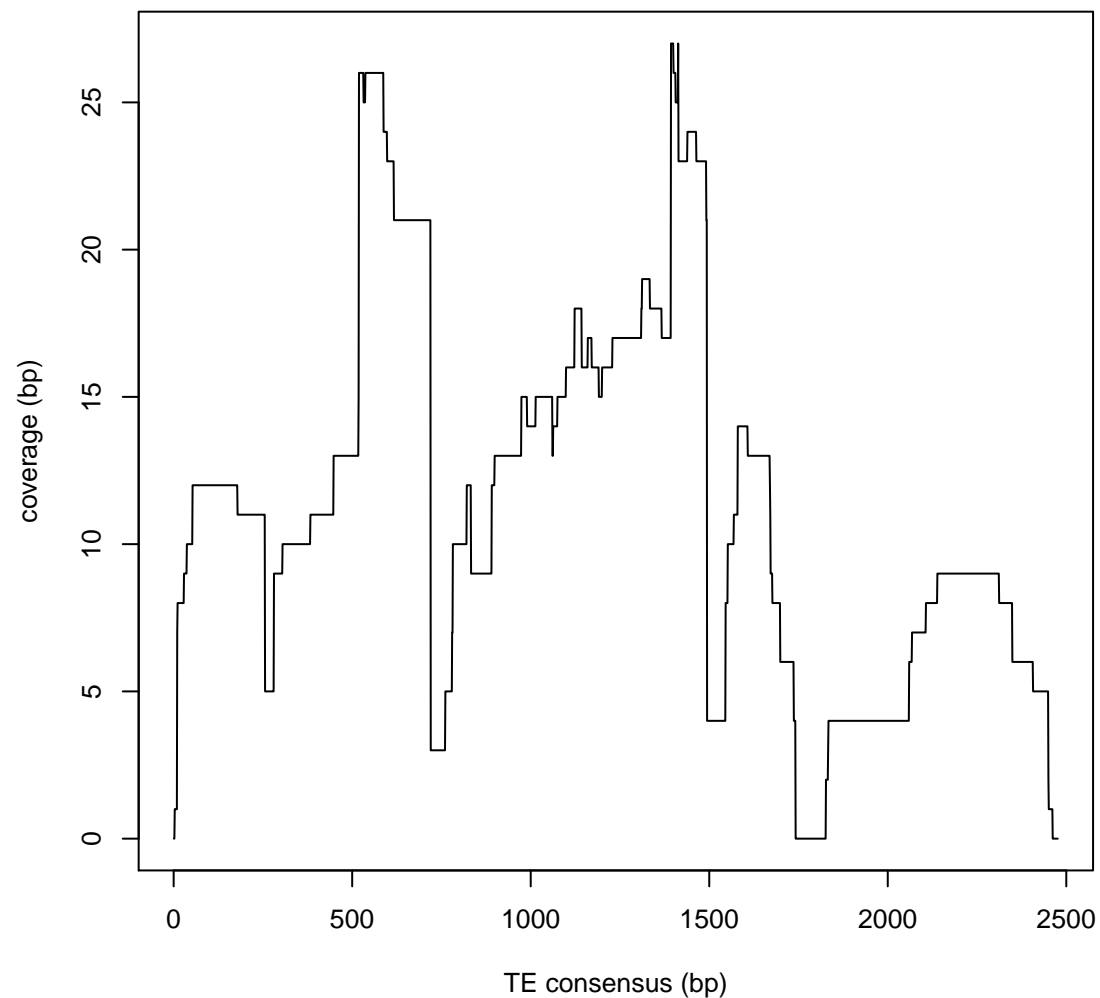


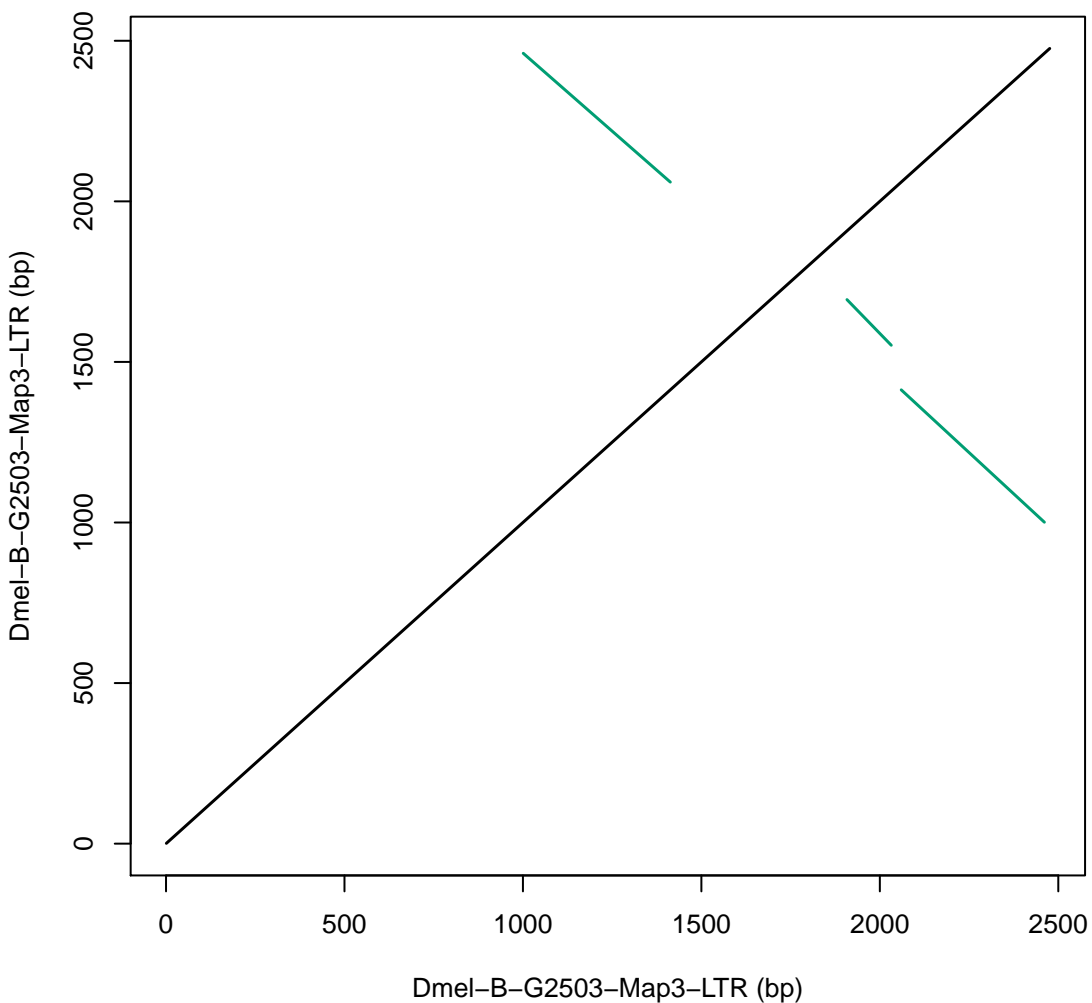
TE: Dmel-B-G2503-Map3-LTR  
consensus size: 2476bp; fragments: 92; full length: 0 ( $\geq 2228.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

