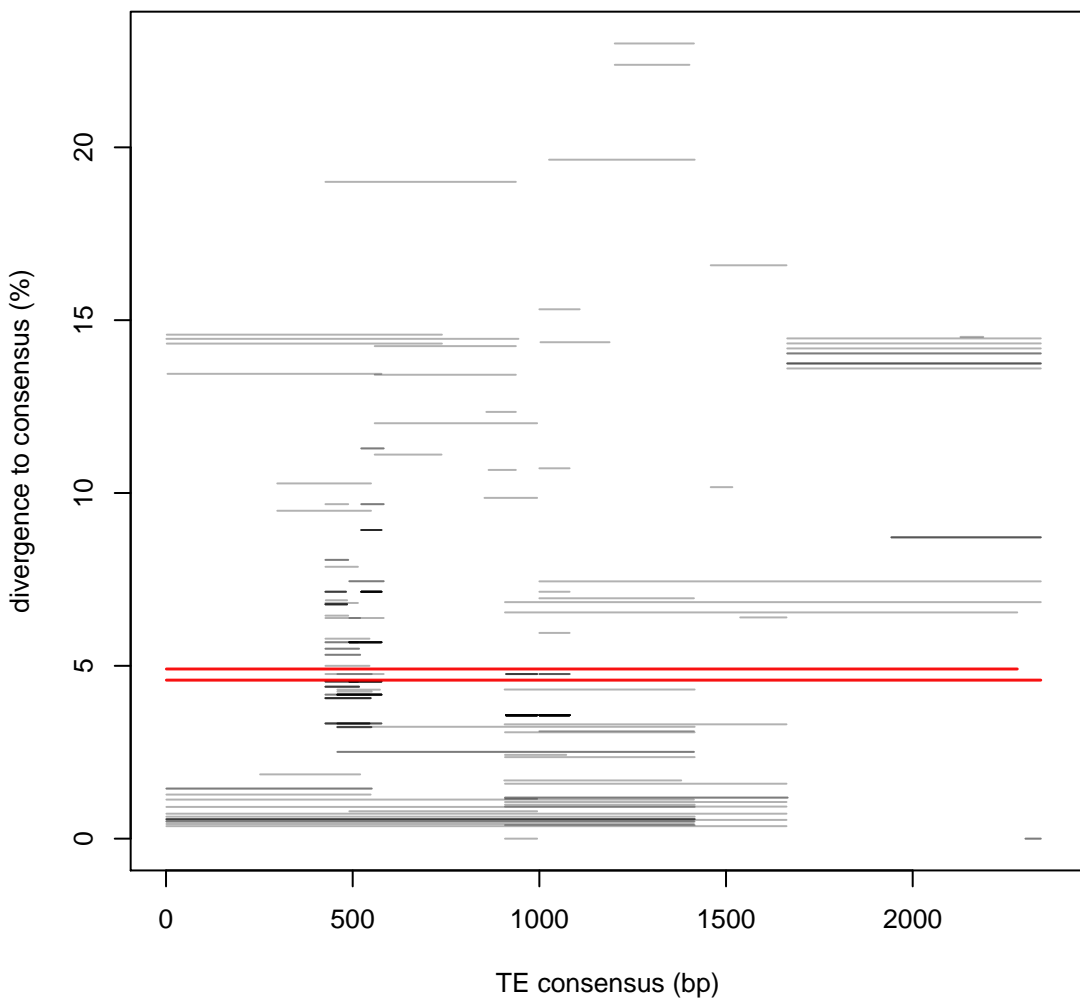
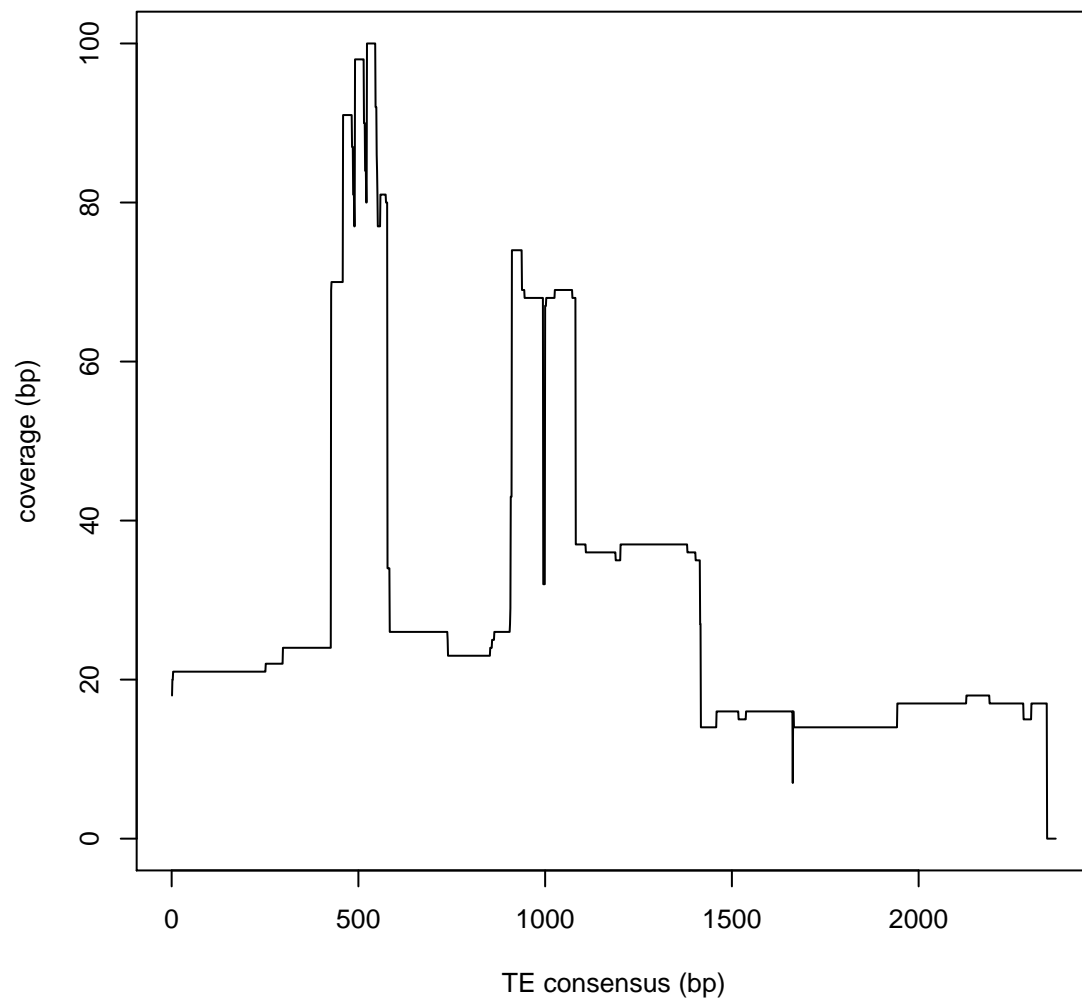


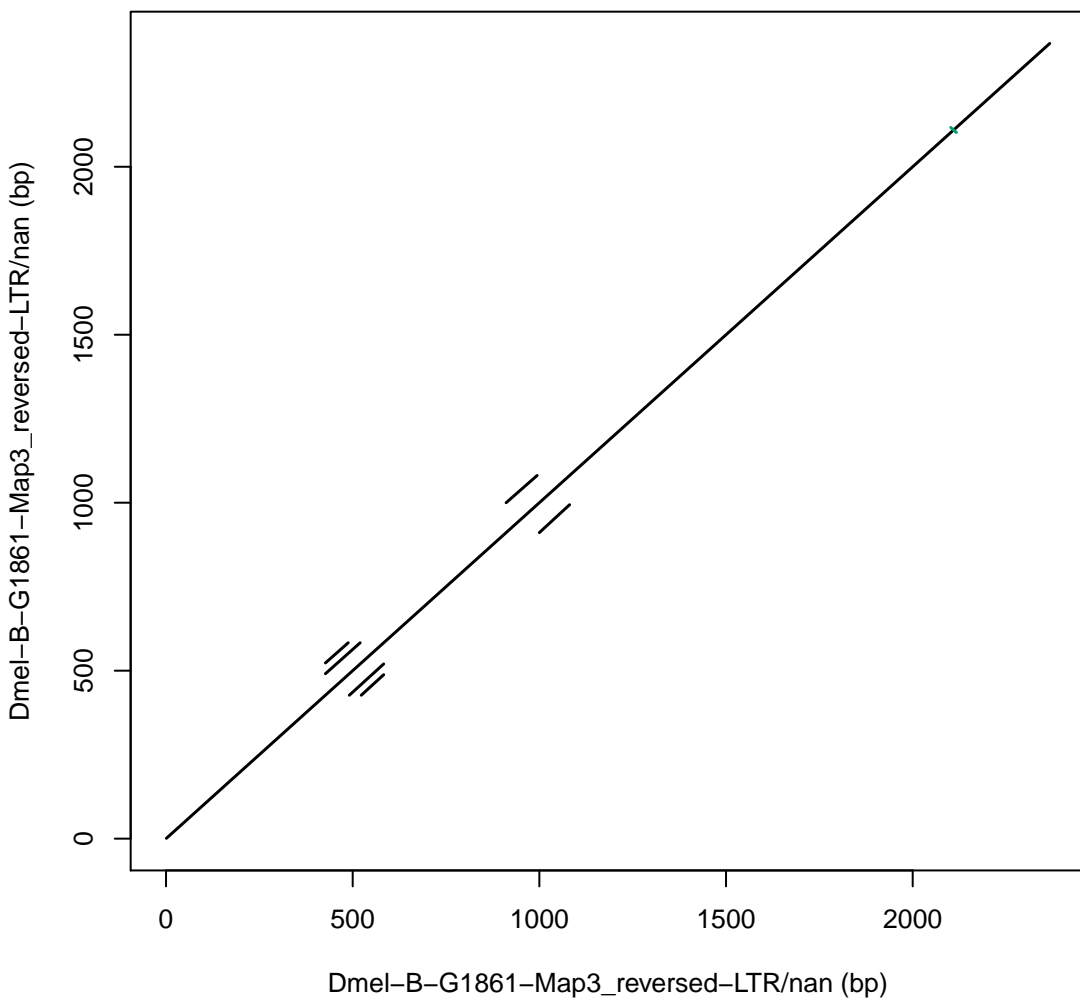
TE: Dmel-B-G1861-Map3\_reversed-LTR/nan  
consensus size: 2367bp; fragments: 244; full length: 2 ( $\geq 2130.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

