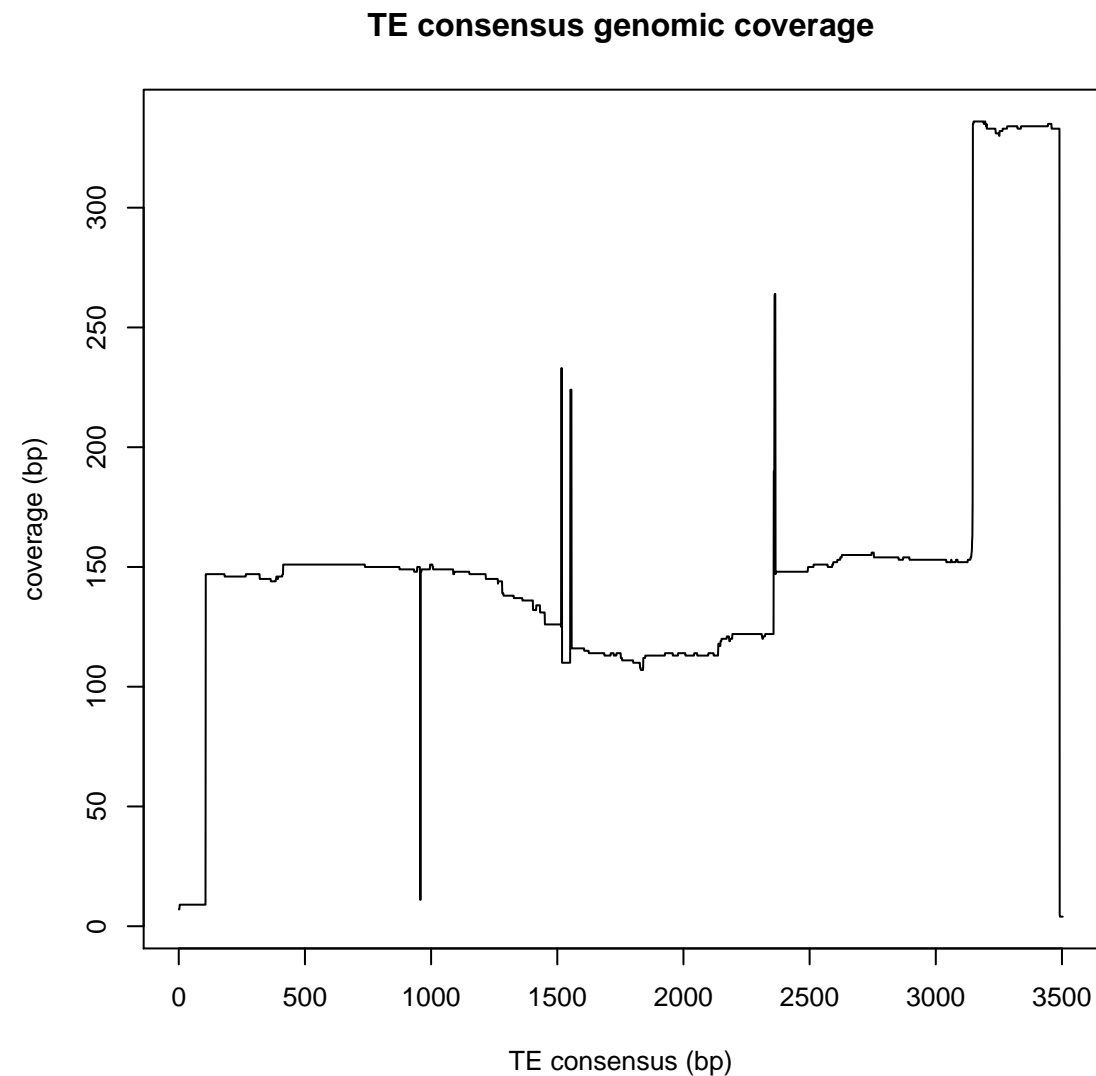
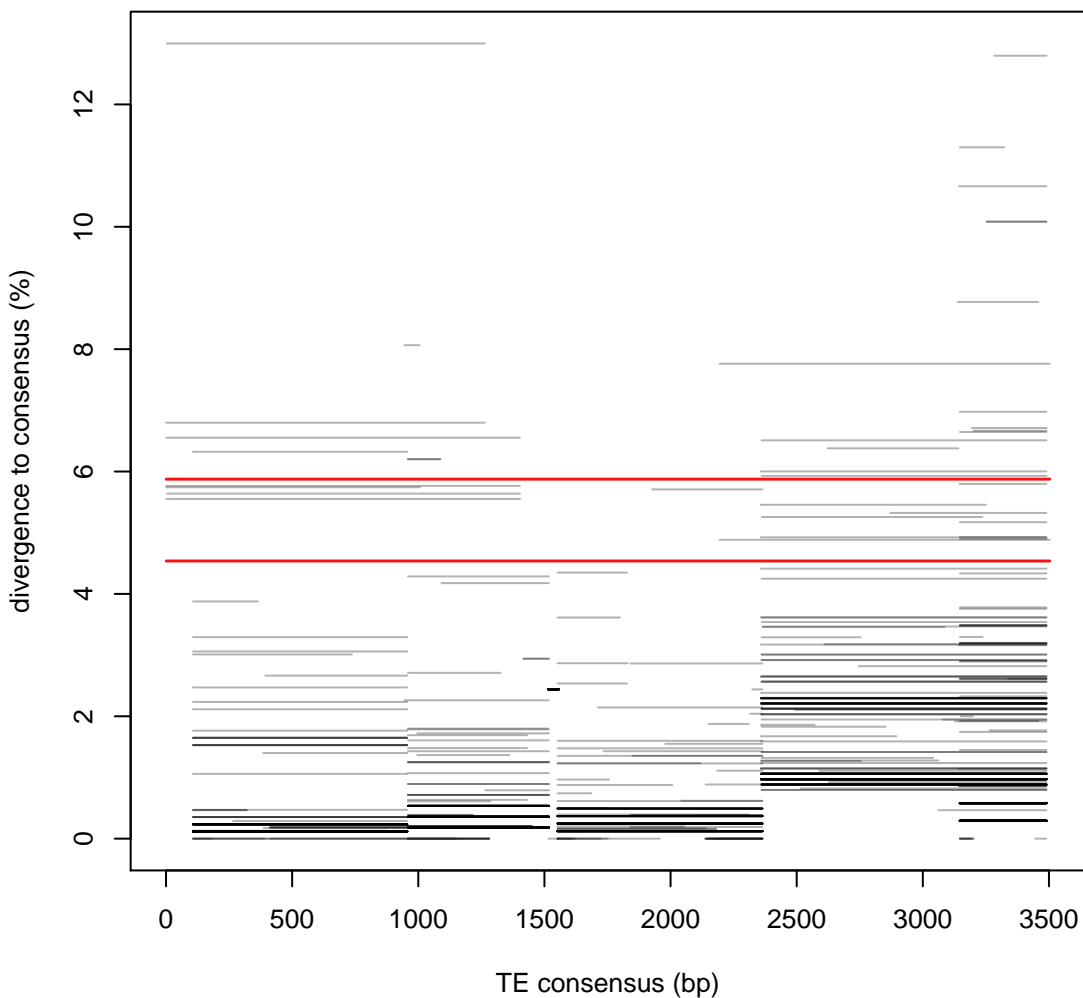
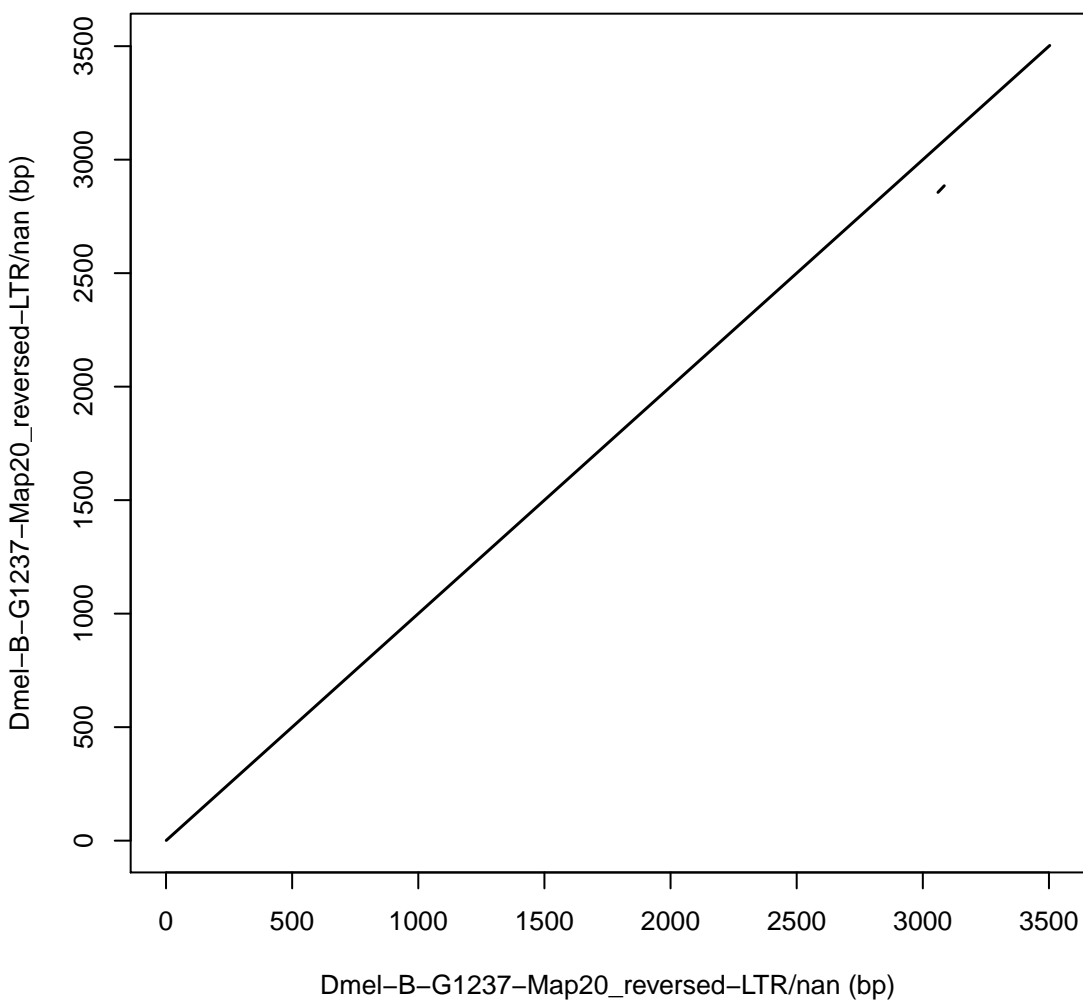


TE: Dmel-B-G1237-Map20\_reversed-LTR/nan  
consensus size: 3503bp; fragments: 899; full length: 2 ( $\geq 3152.7$ bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

