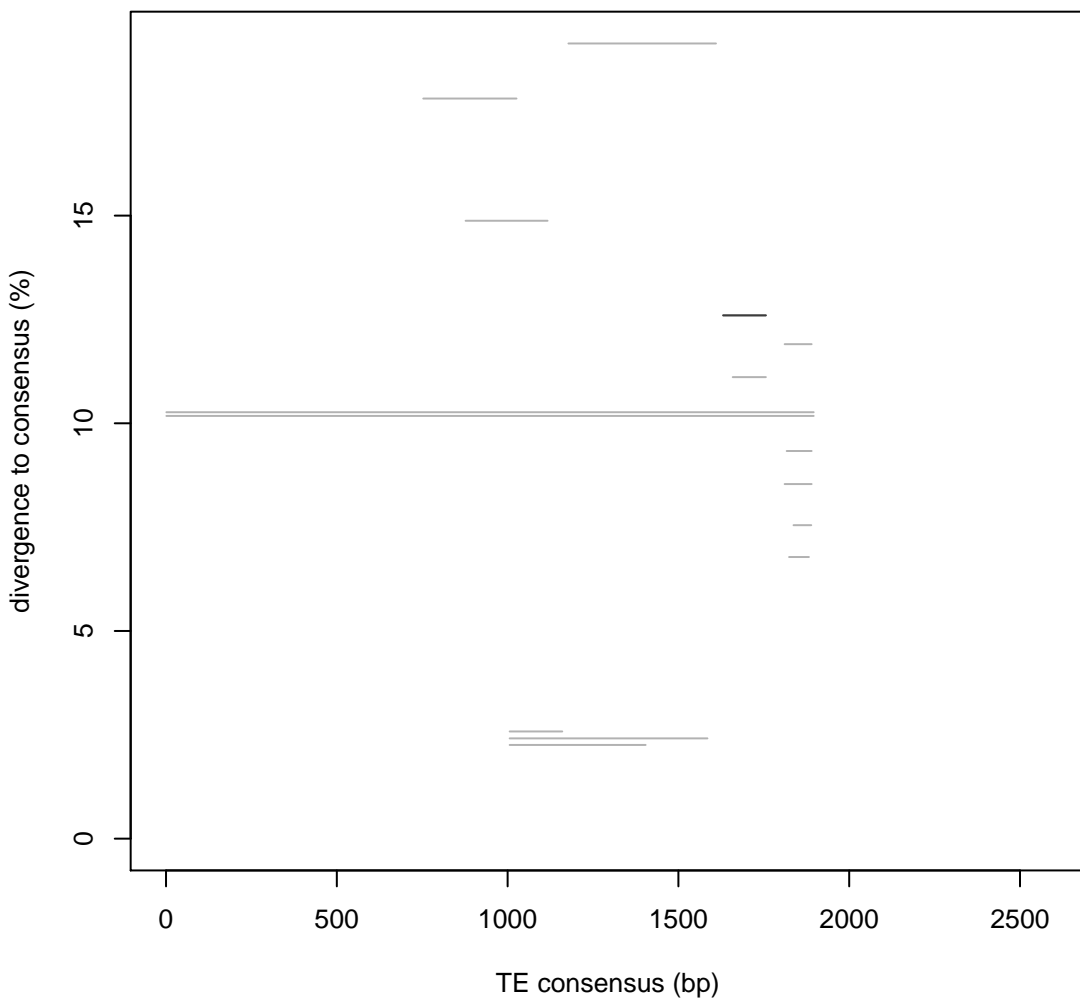
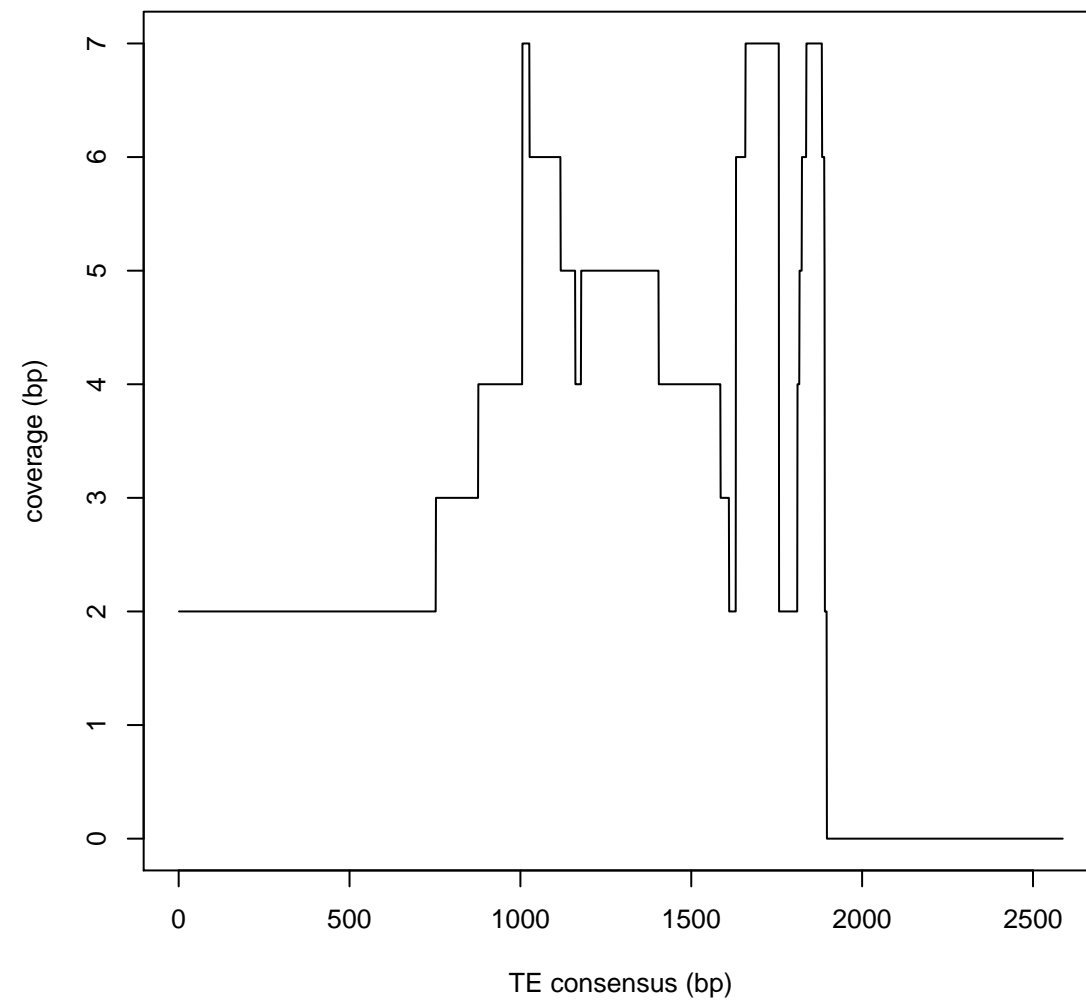


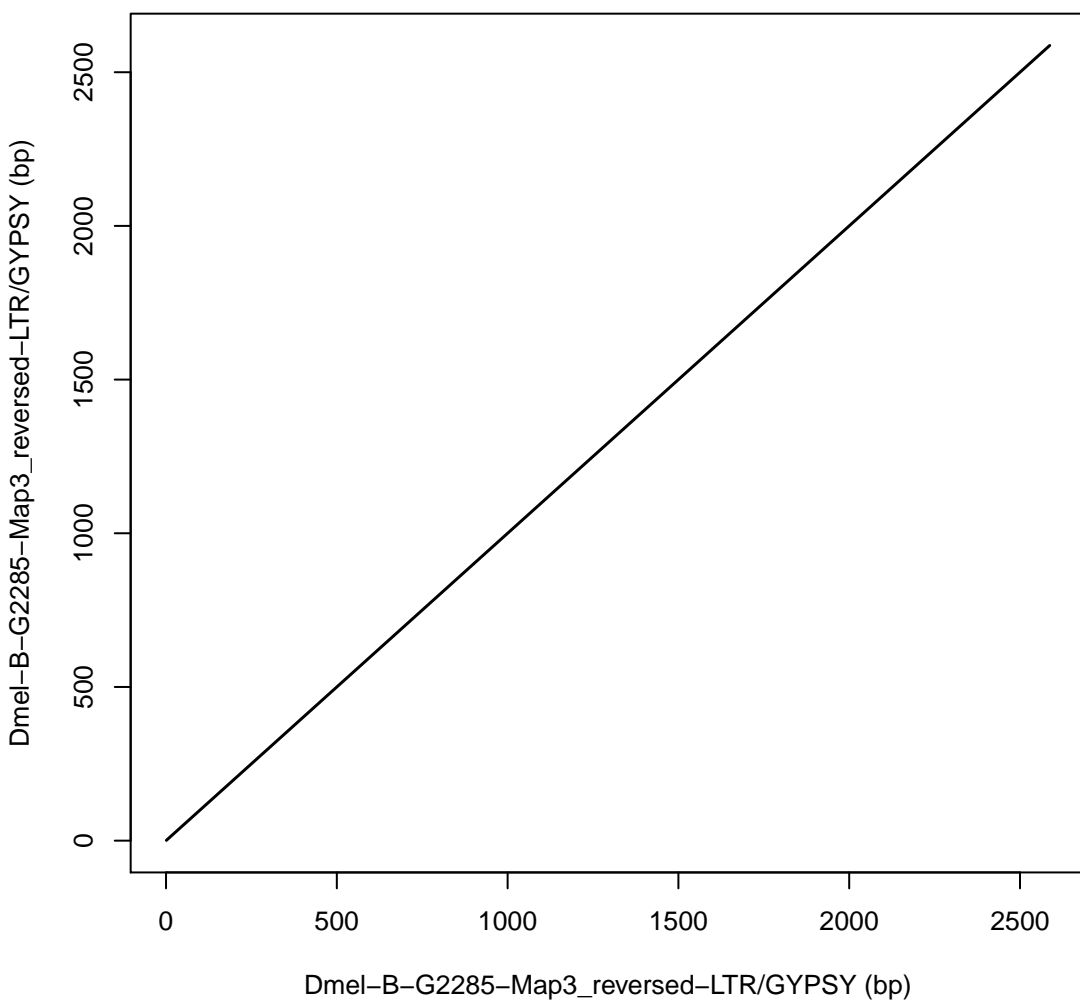
TE: Dmel-B-G2285-Map3\_reversed-LTR/GYPSY  
consensus size: 2587bp; fragments: 18; full length: 0 ( $\geq 2328.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

