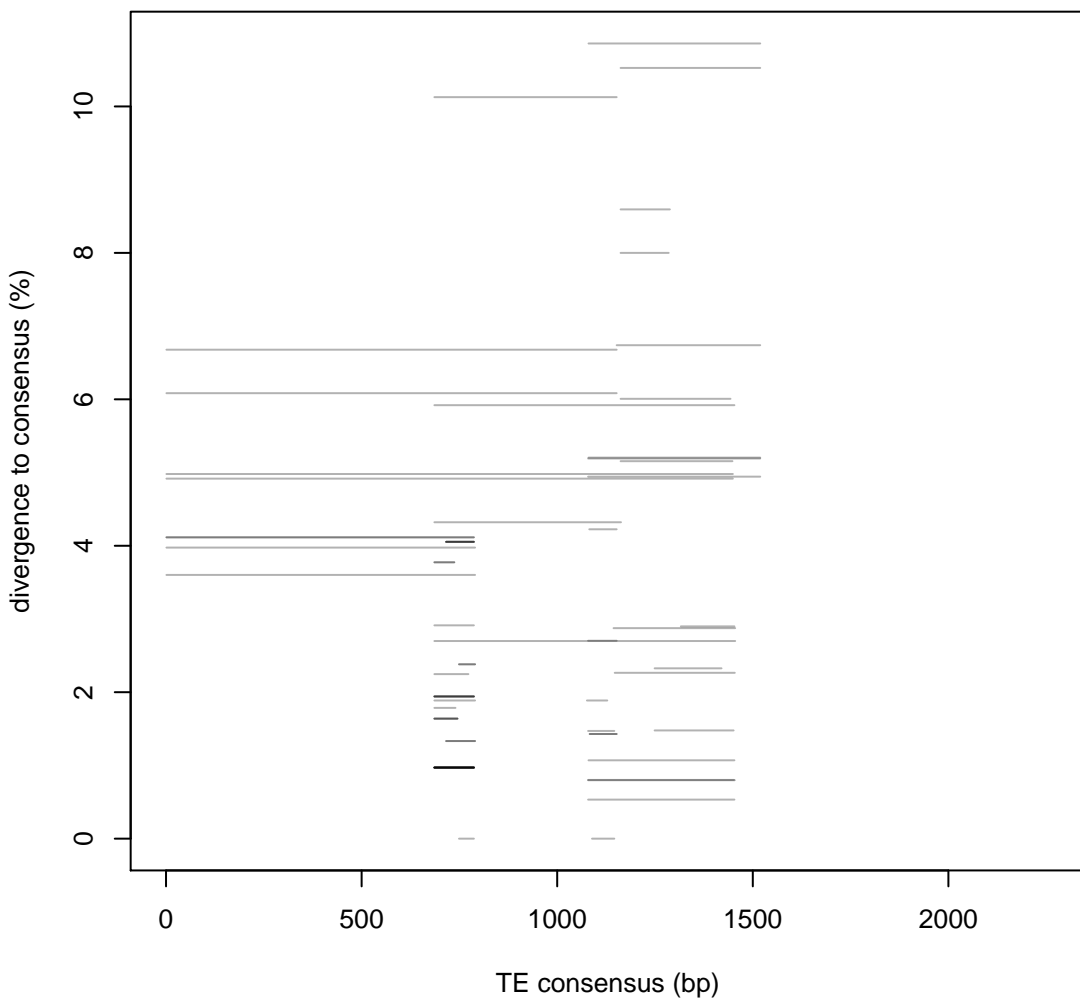
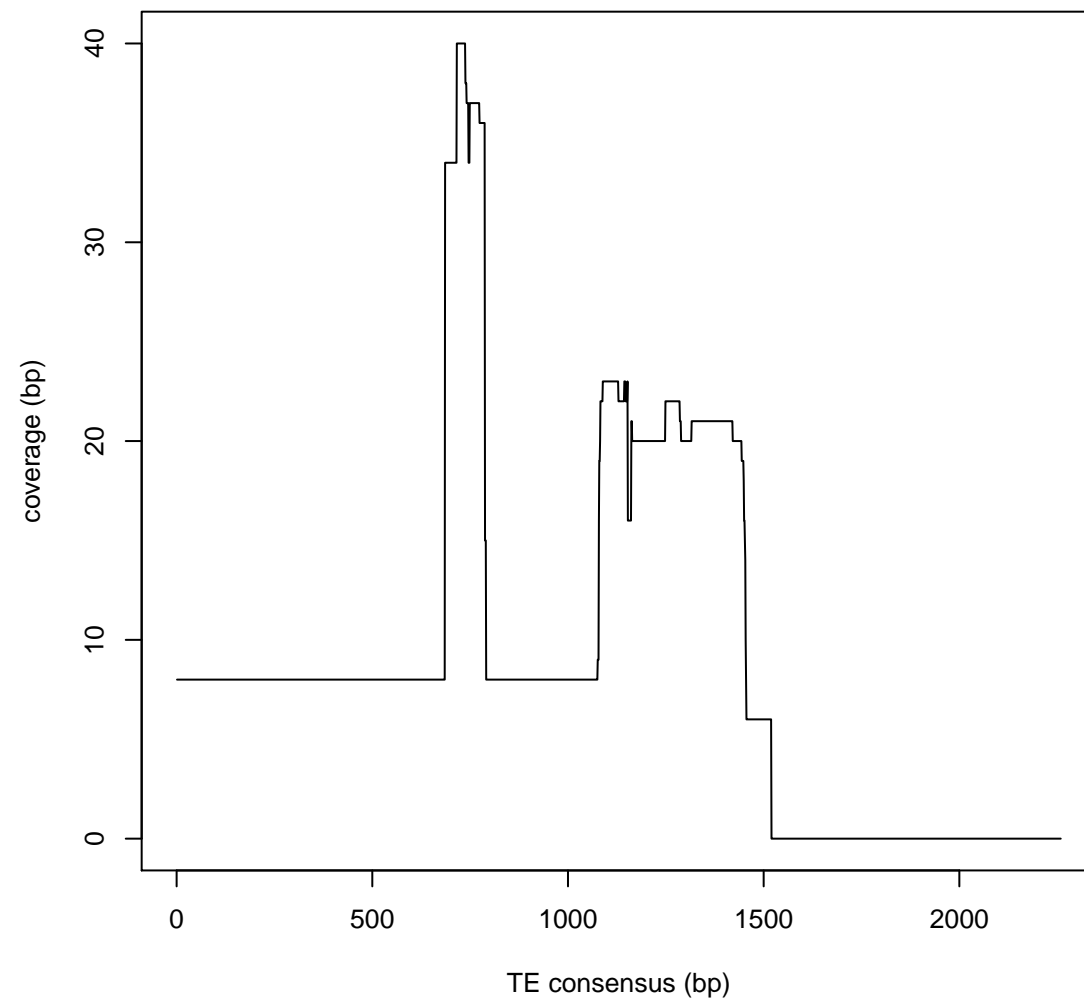


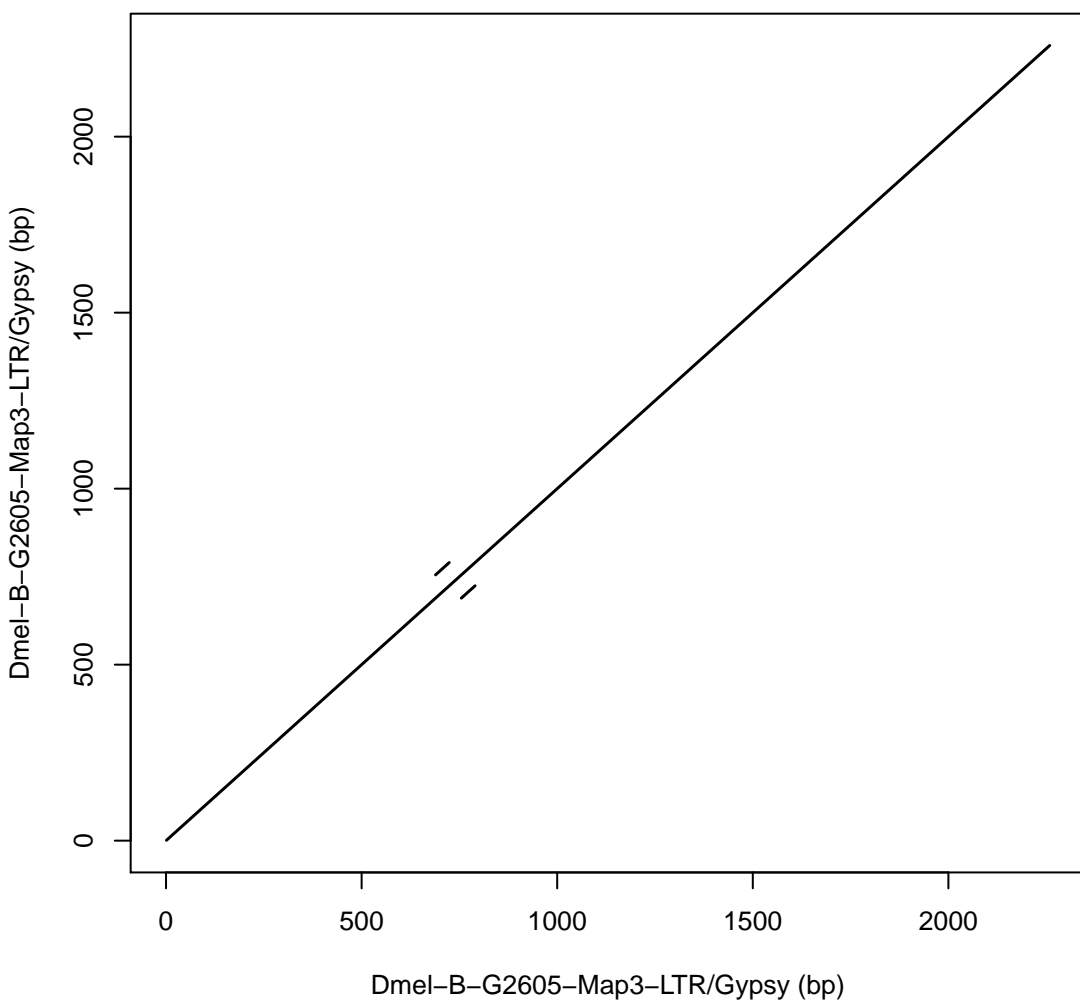
TE: Dmel-B-G2605-Map3-LTR/Gypsy  
consensus size: 2259bp; fragments: 69; full length: 0 ( $\geq 2033.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

