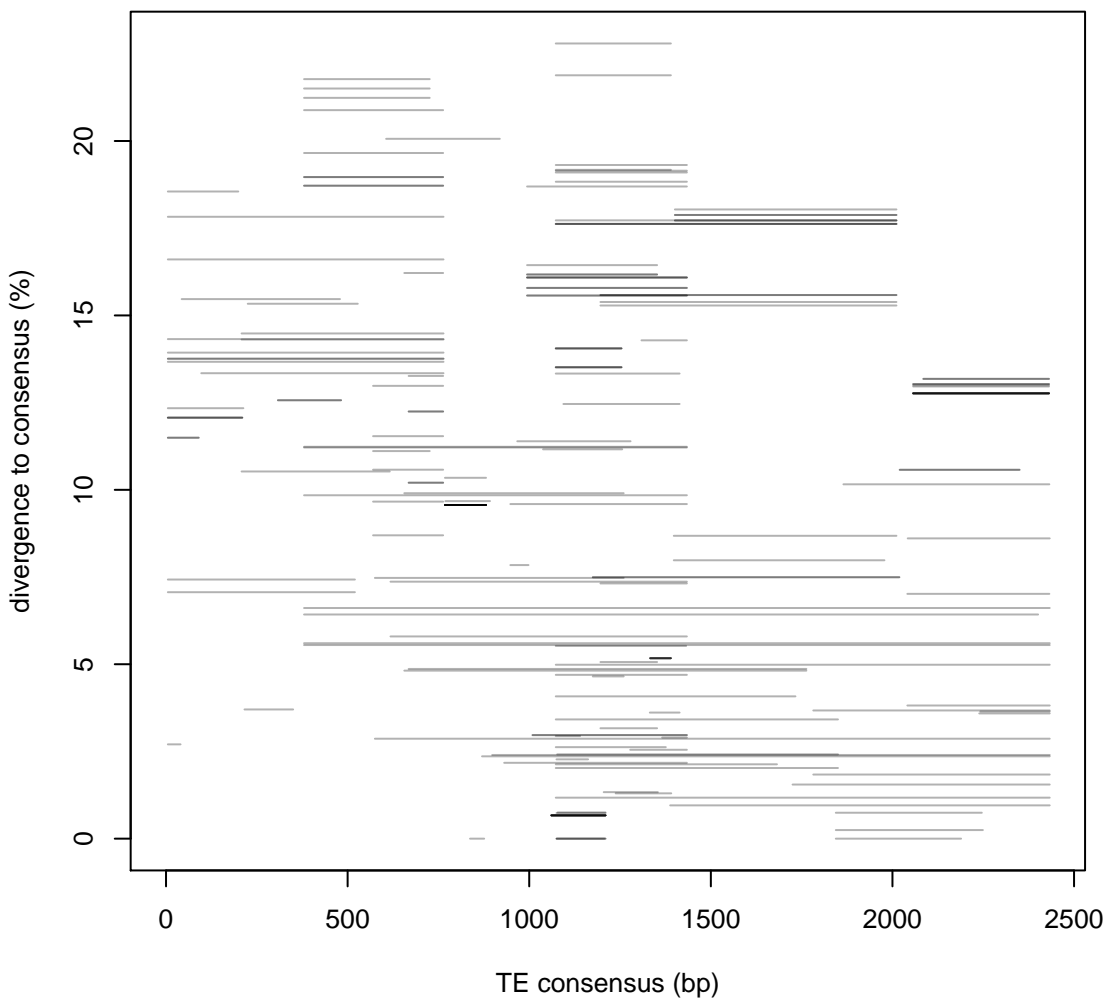
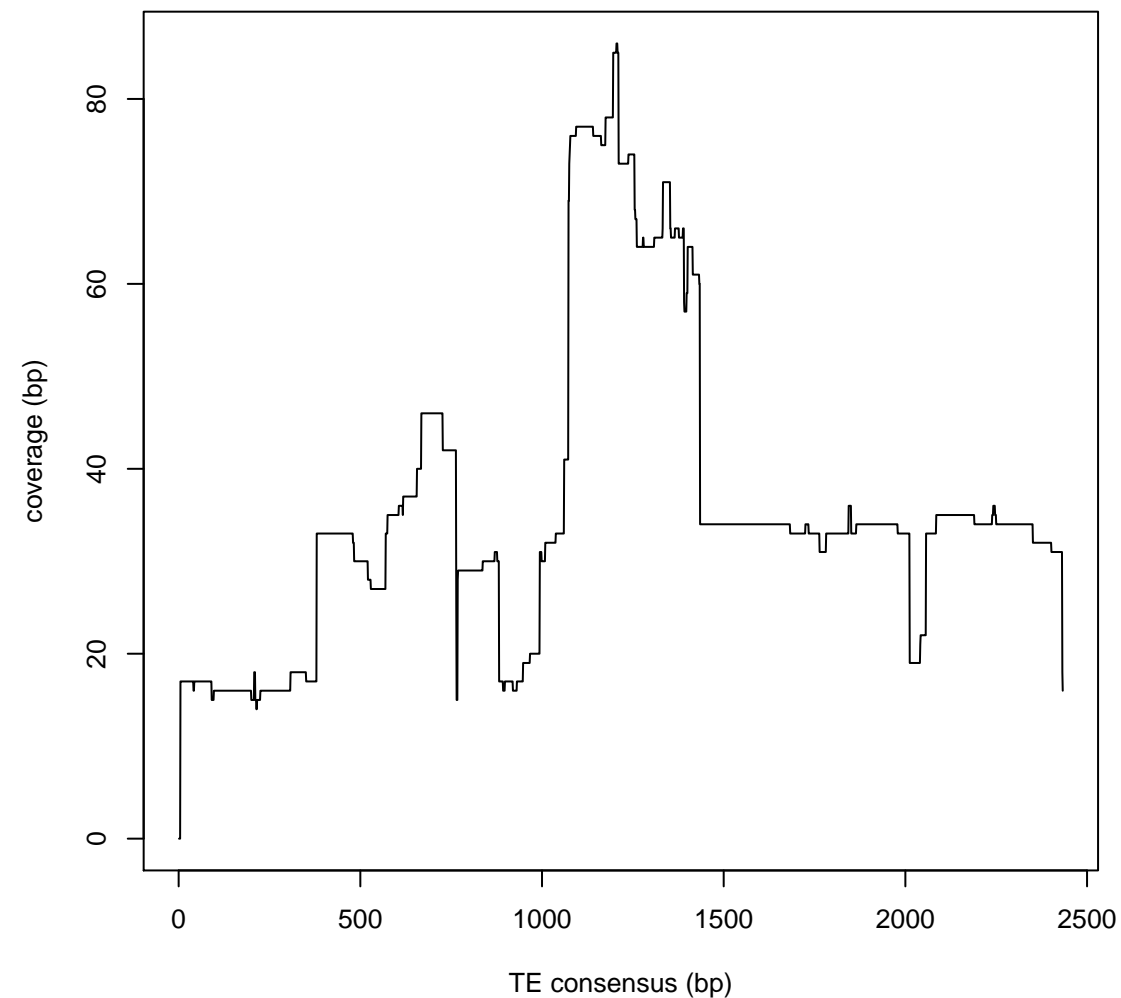


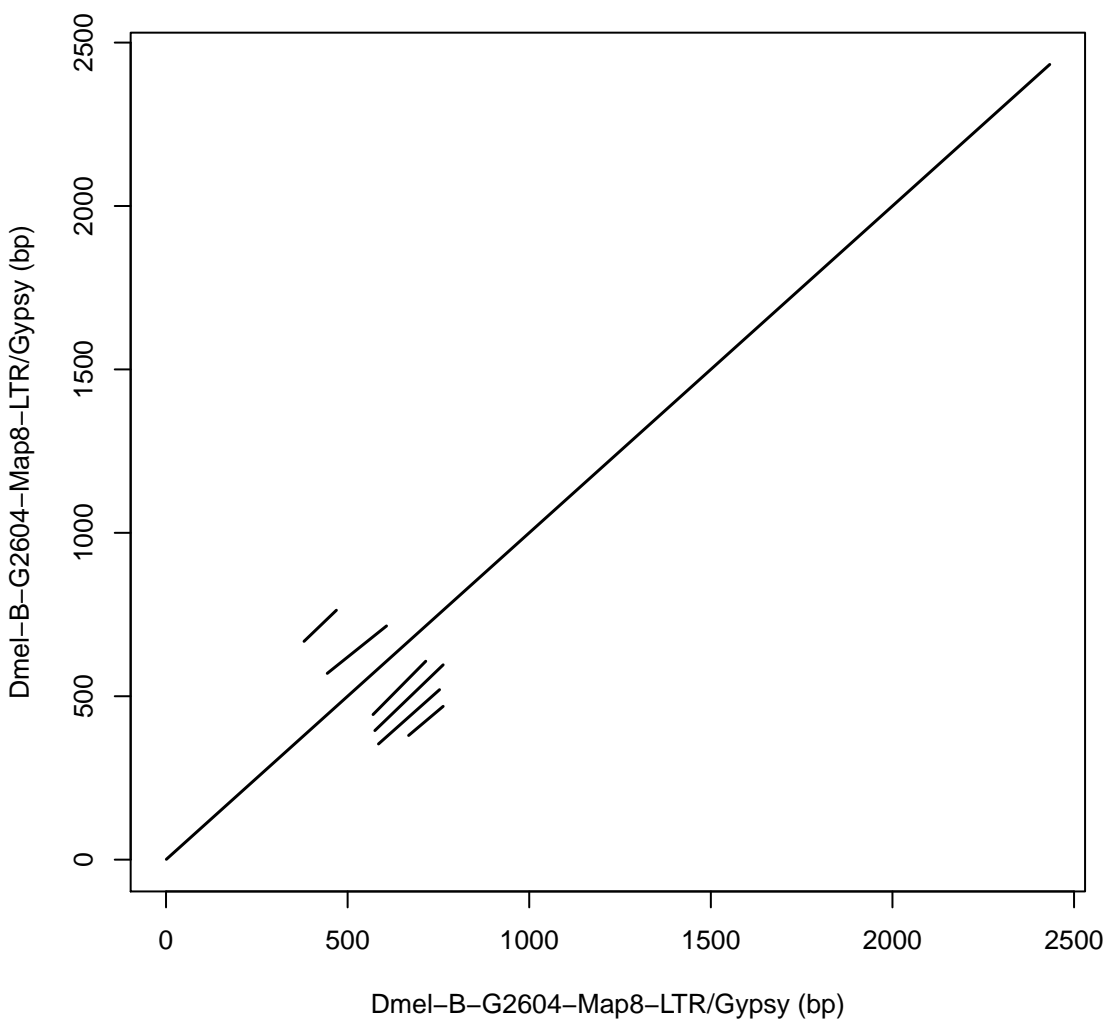
TE: Dmel-B-G2604-Map8-LTR/Gypsy  
consensus size: 2433bp; fragments: 197; full length: 0 ( $\geq 2189.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

