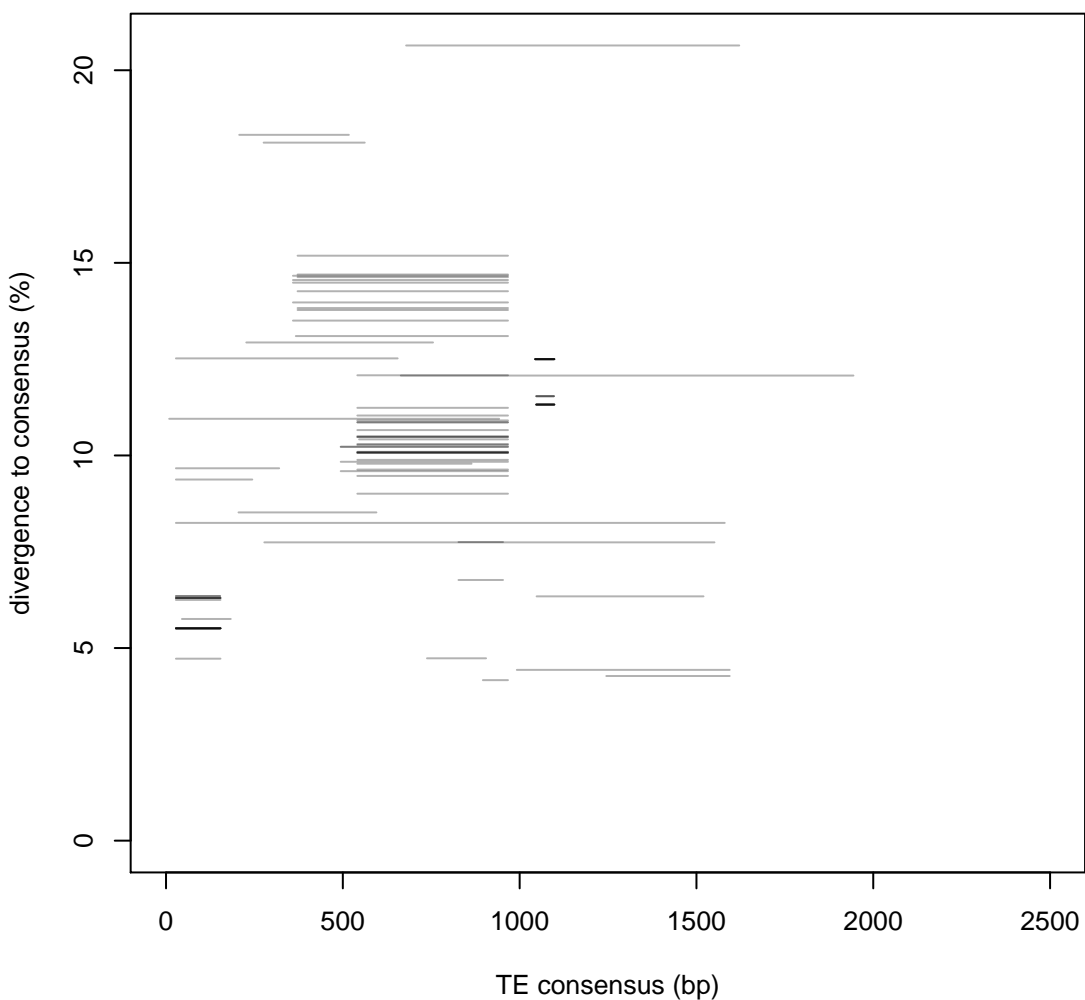
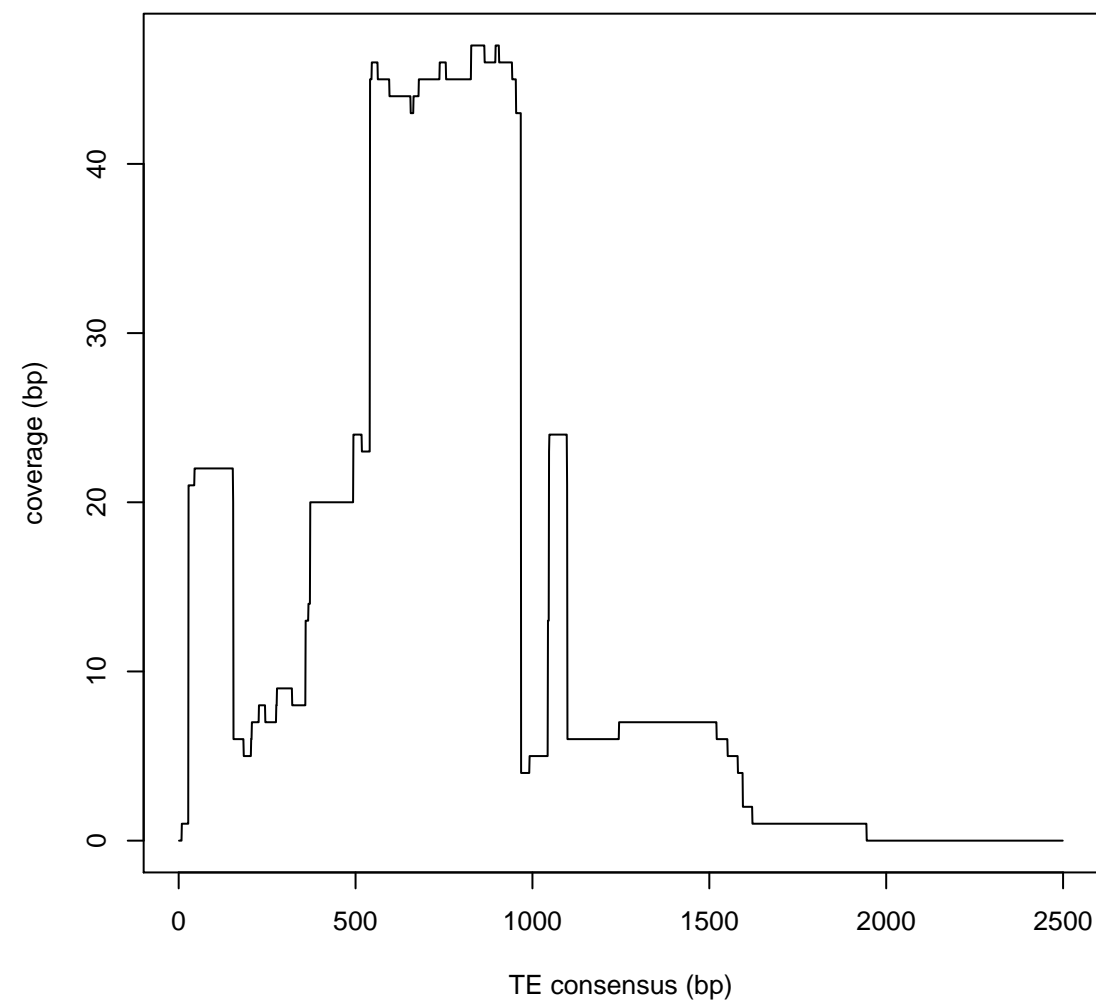


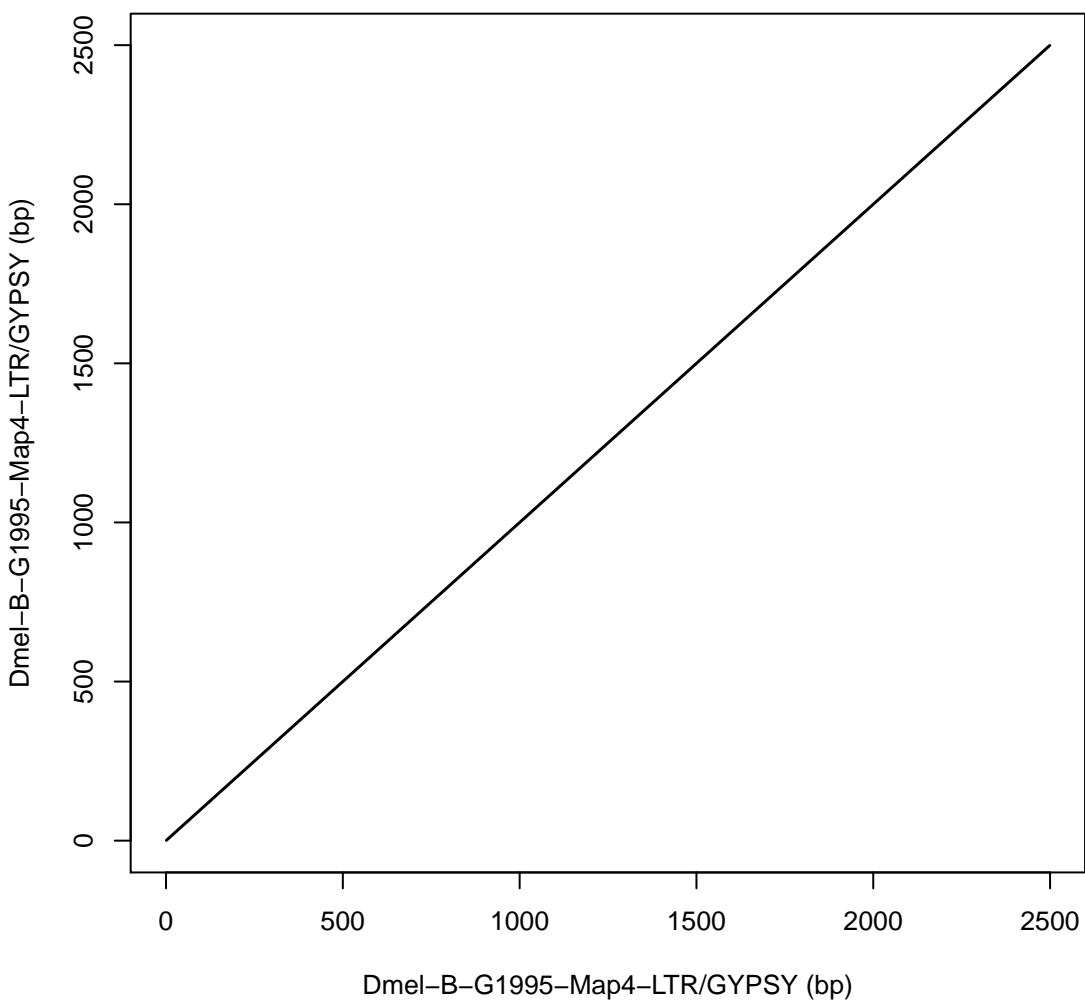
TE: Dmel-B-G1995-Map4-LTR/GYPSY  
consensus size: 2499bp; fragments: 93; full length: 0 ( $\geq 2249.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

