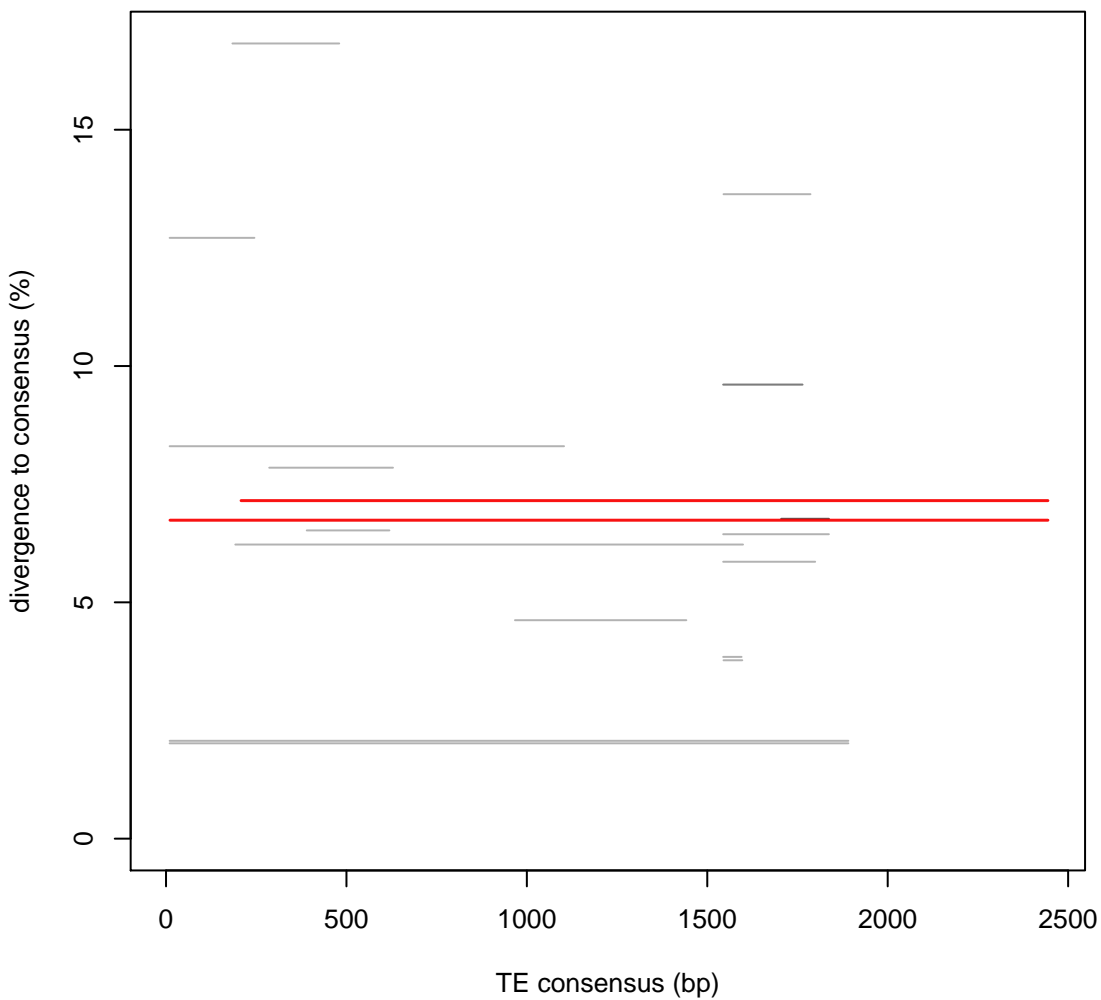
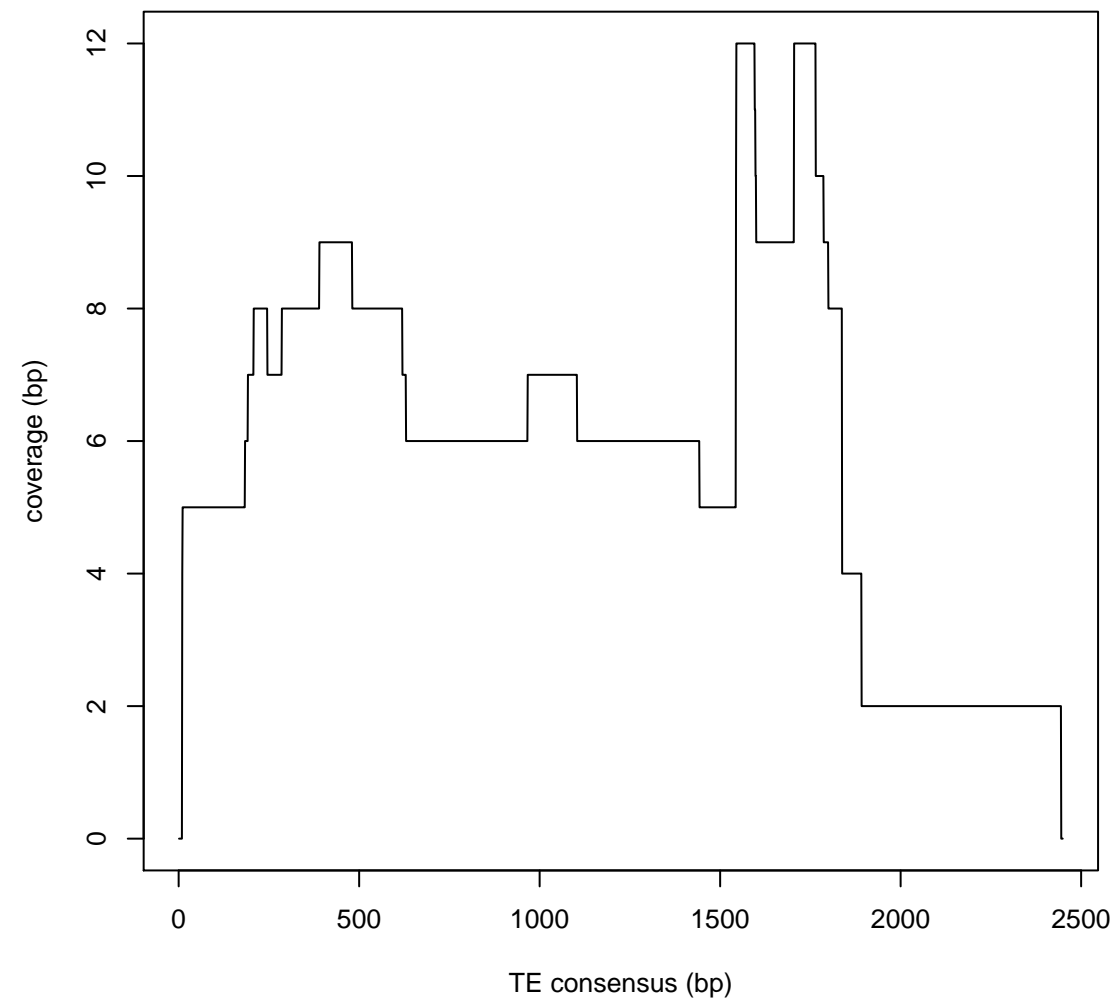


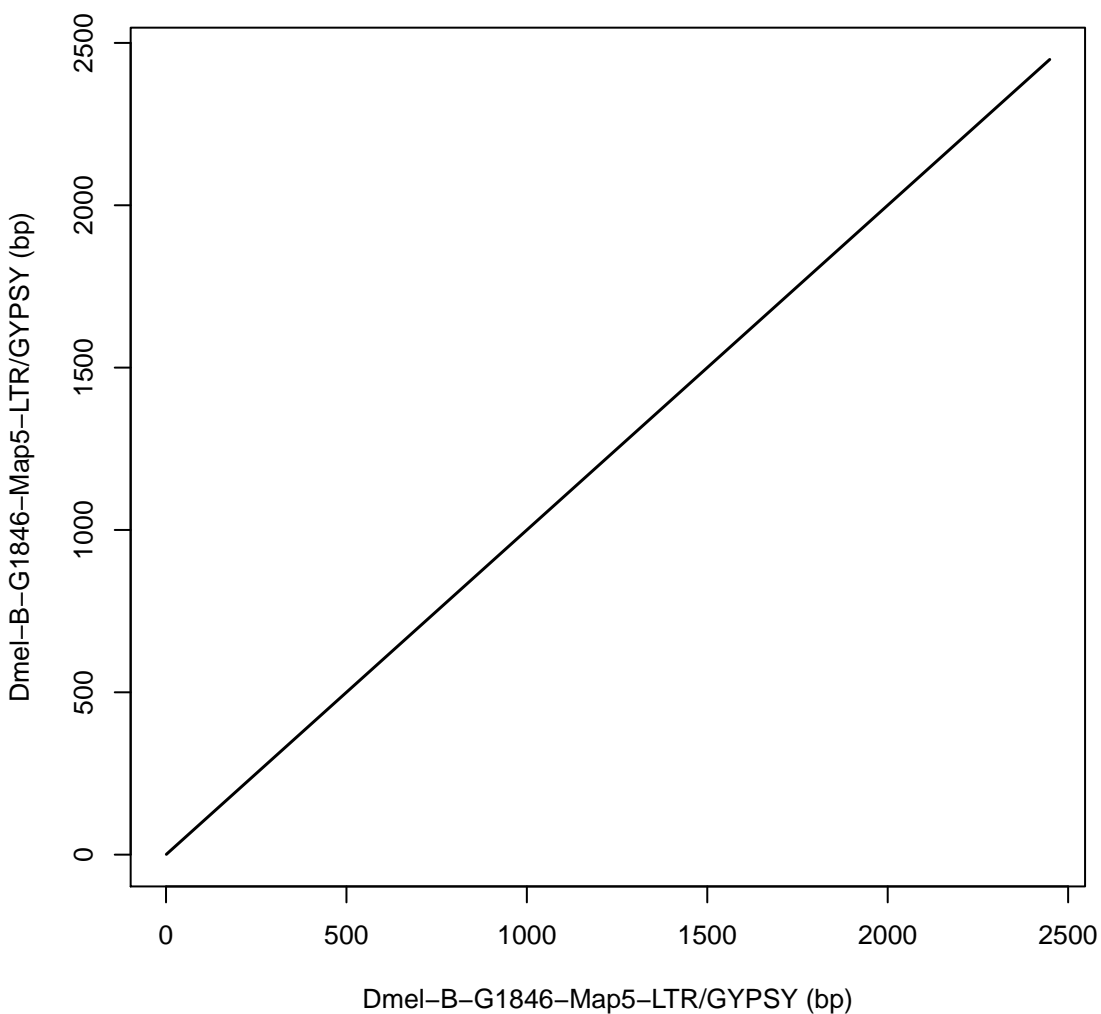
TE: Dmel-B-G1846-Map5-LTR/GYPSY  
consensus size: 2449bp; fragments: 21; full length: 2 ( $\geq 2204.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

