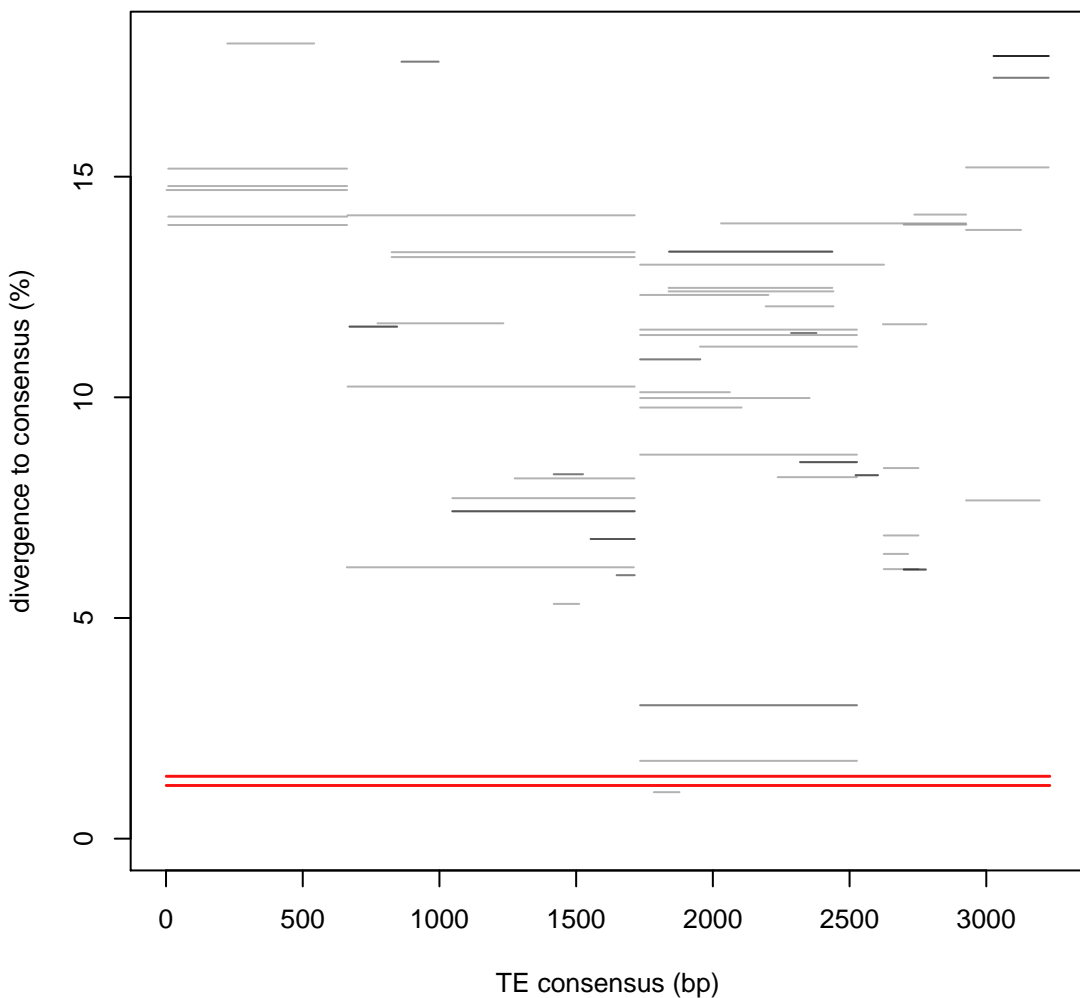
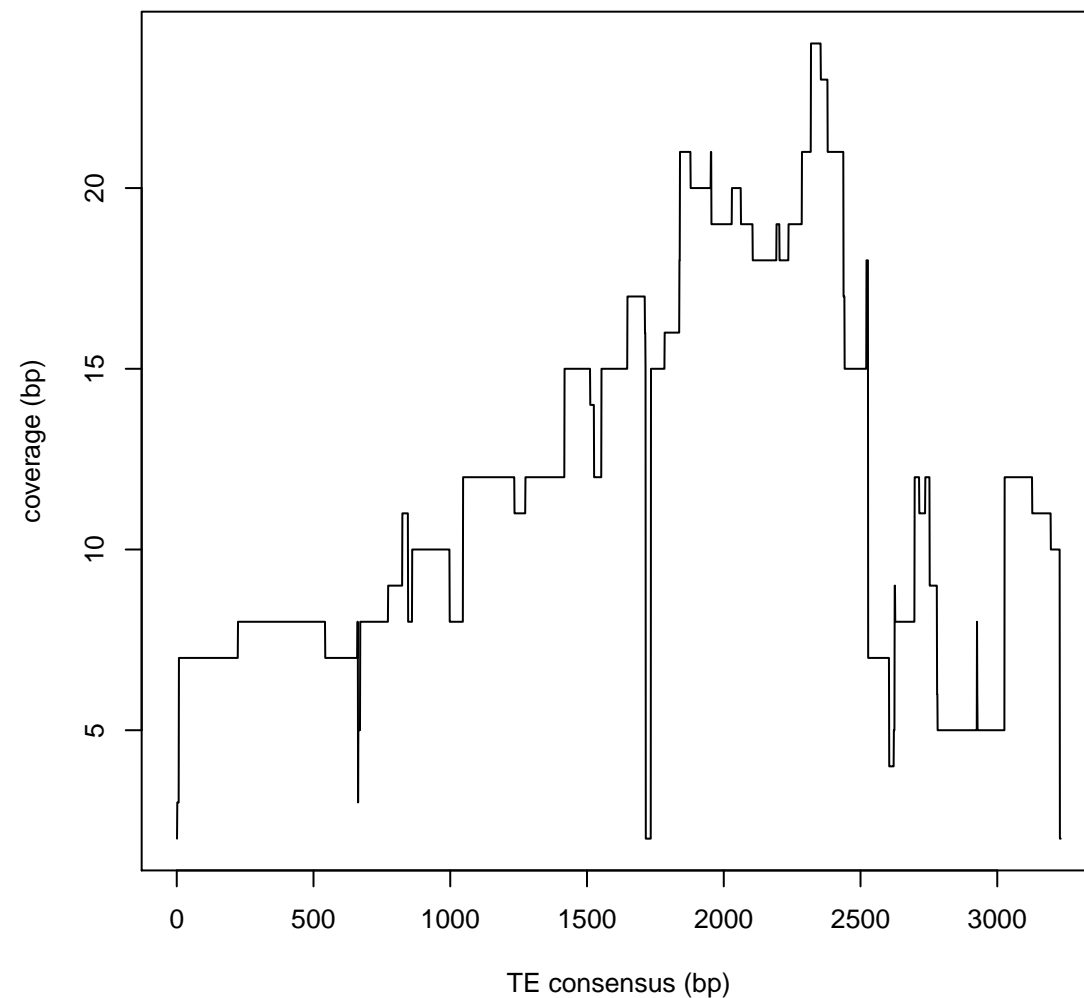


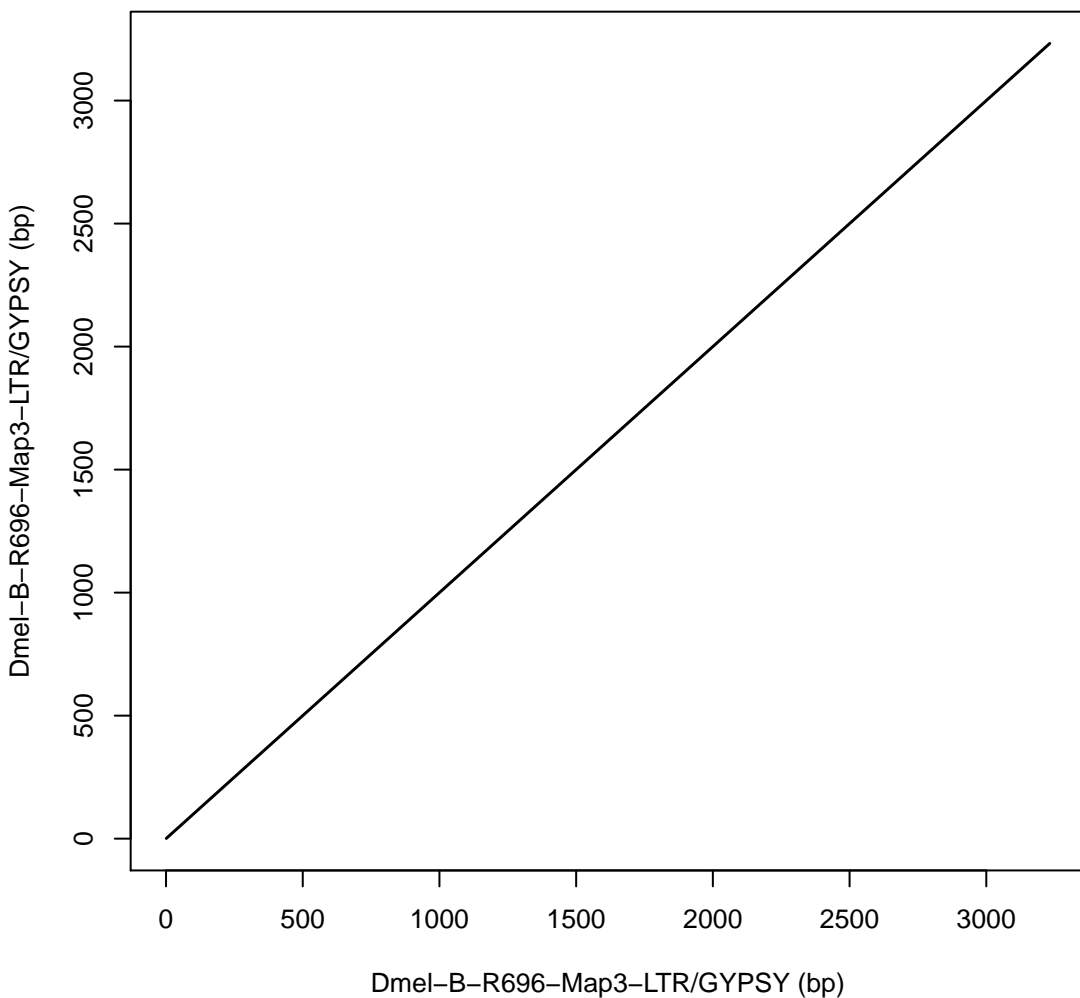
TE: Dmel-B-R696-Map3-LTR/GYPSY  
consensus size: 3232bp; fragments: 83; full length: 2 ( $\geq 2908.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

