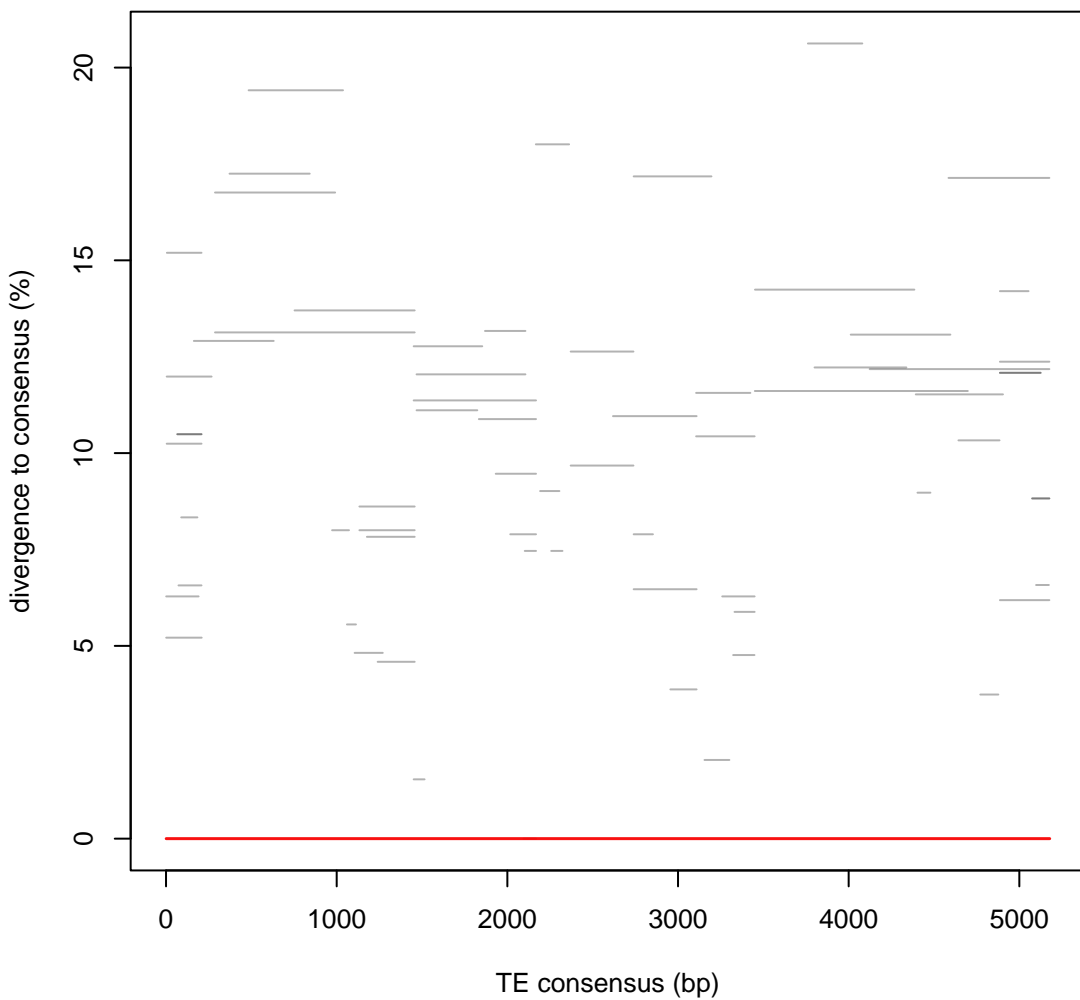
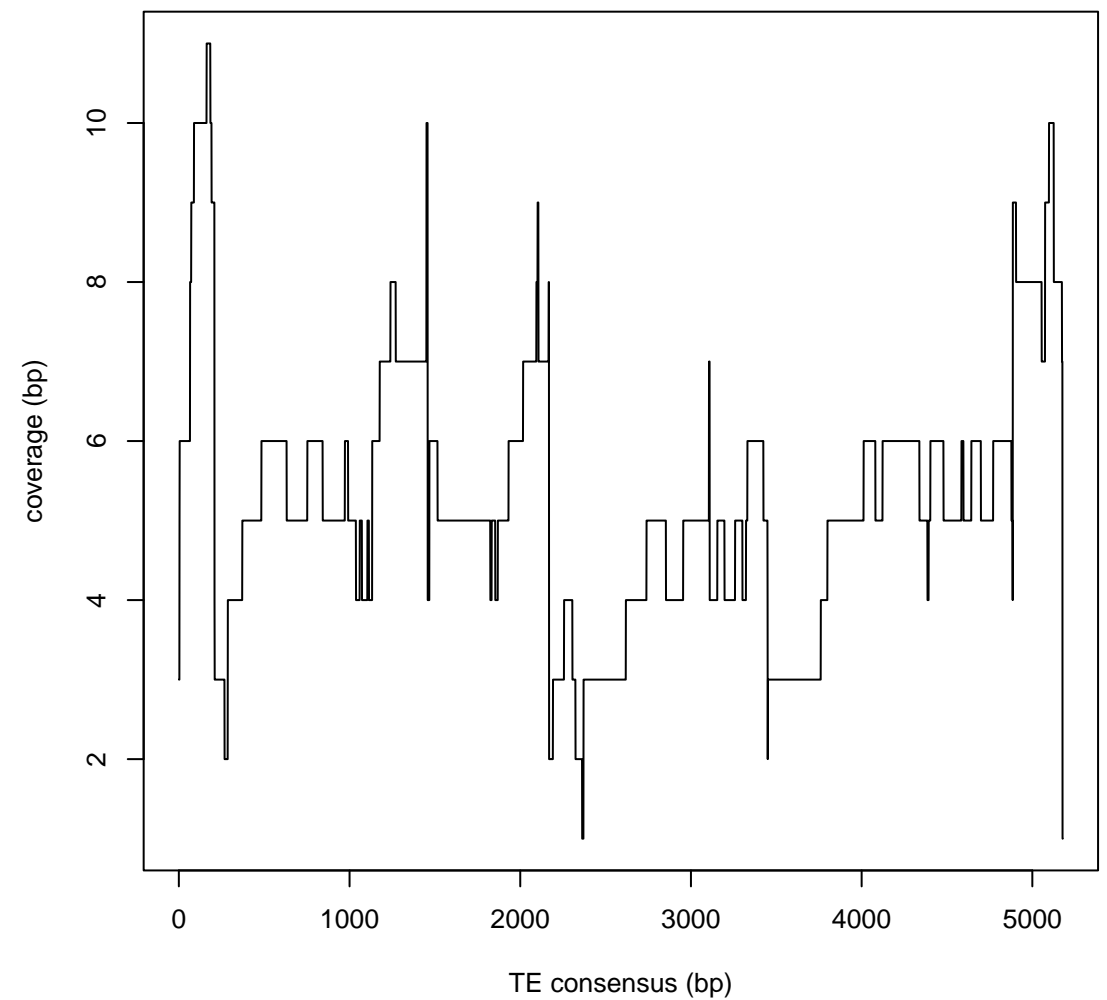


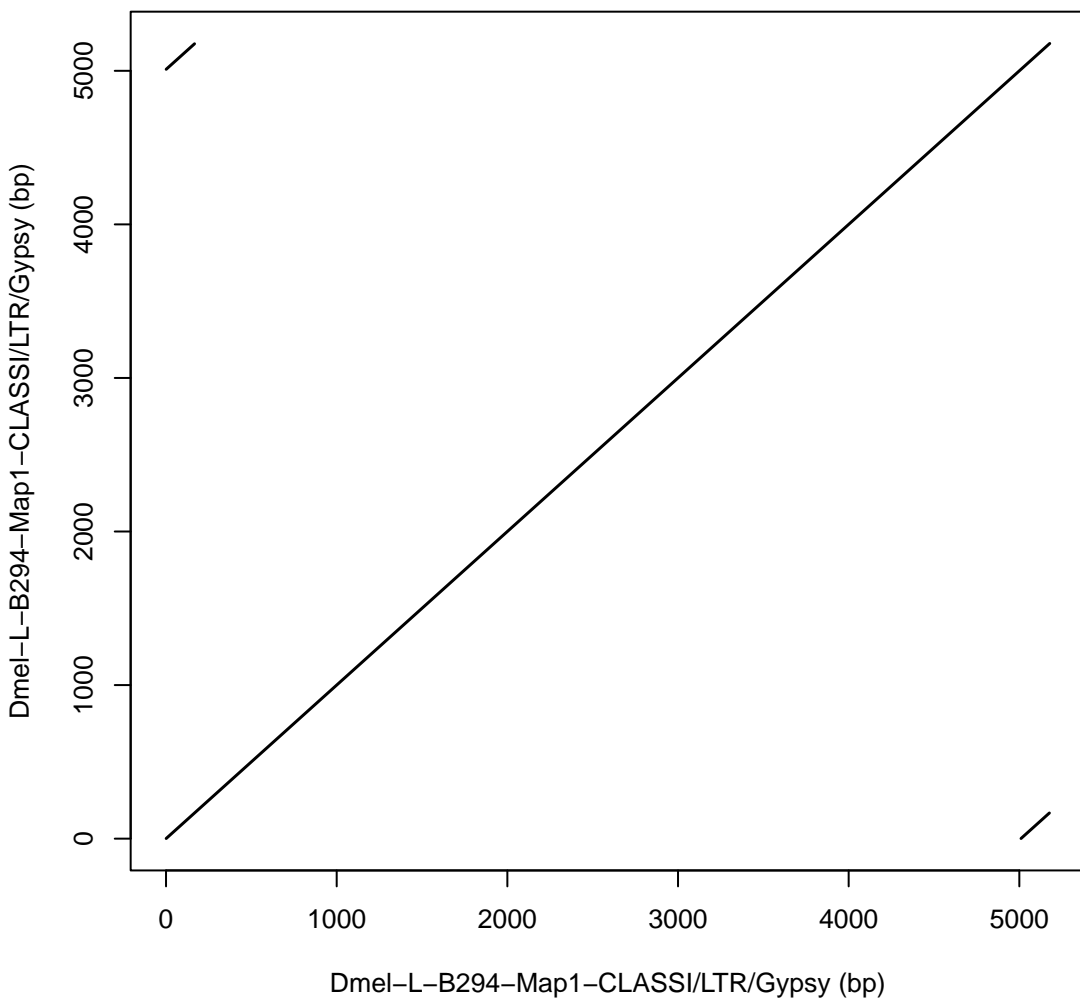
TE: Dmel-L-B294-Map1-CLASSI/LTR/Gypsy  
consensus size: 5178bp; fragments: 69; full length: 1 ( $\geq 4660.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

