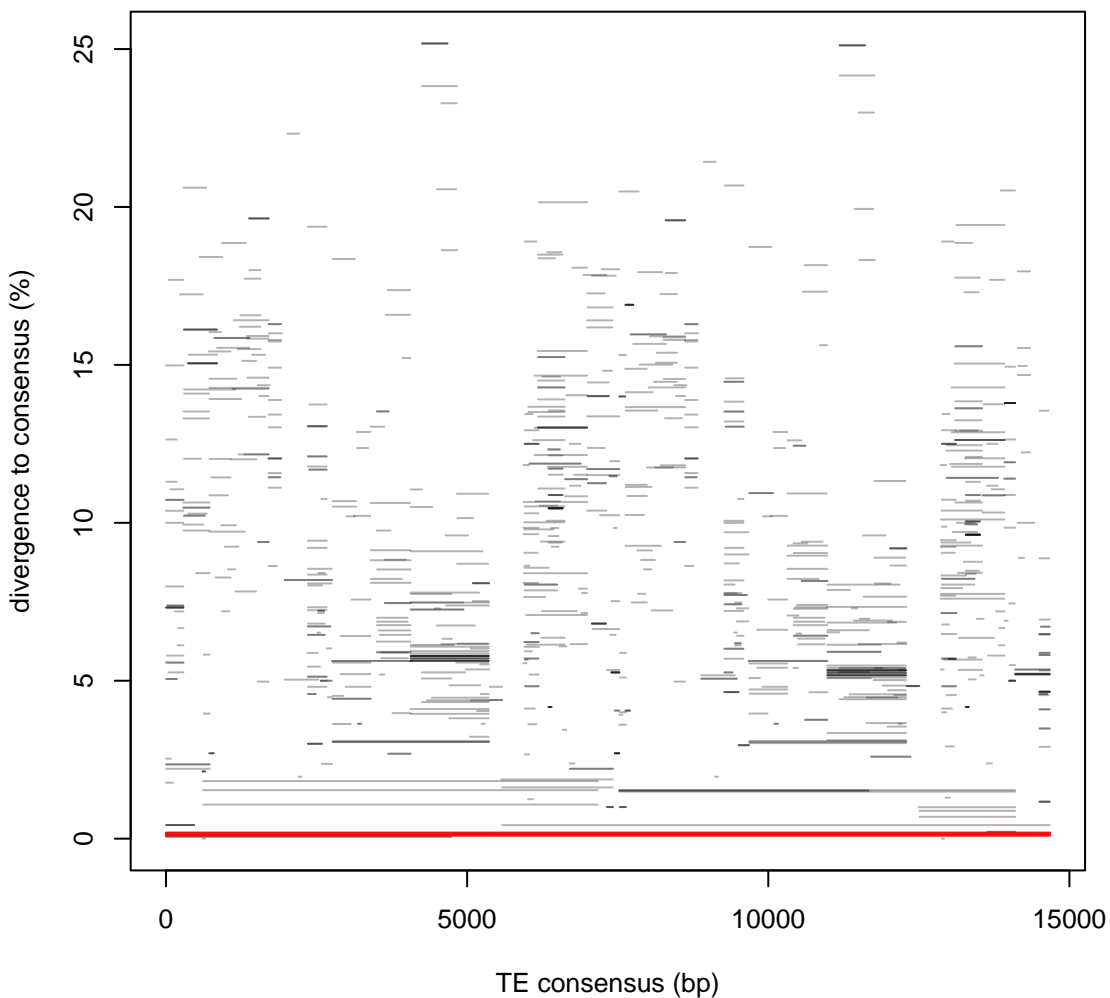
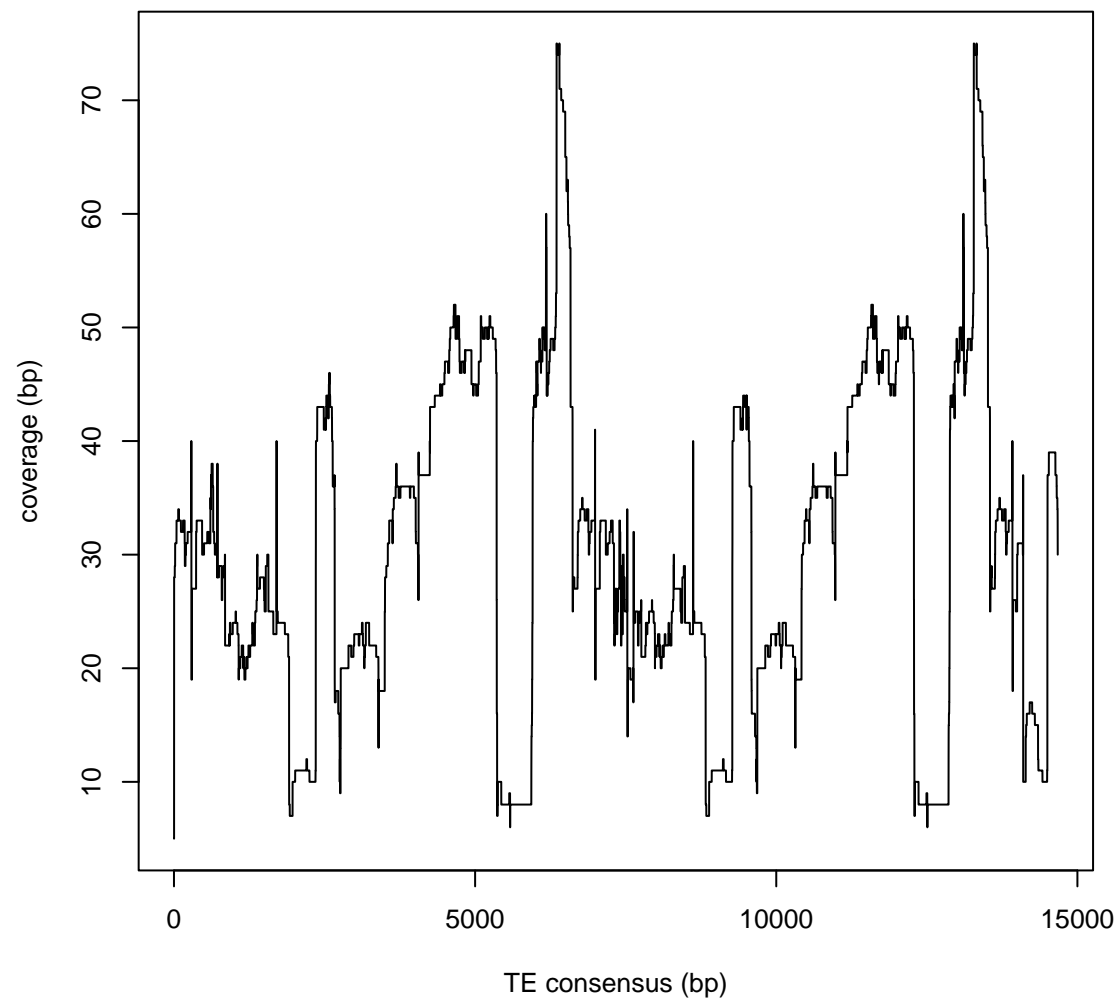


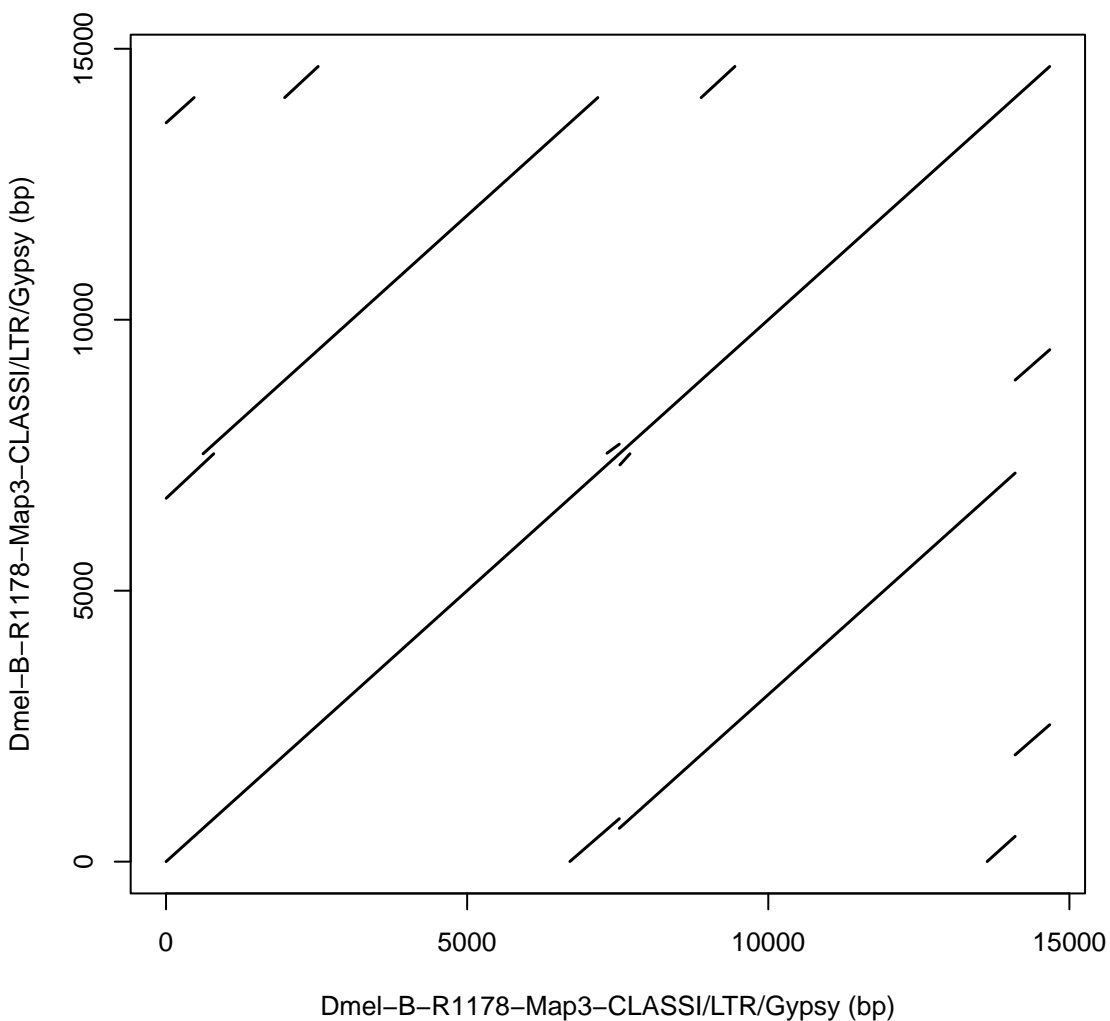
TE: Dmel-B-R1178-Map3-CLASSI/LTR/Gypsy  
consensus size: 14673bp; fragments: 918; full length: 2 ( $\geq 13205.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

