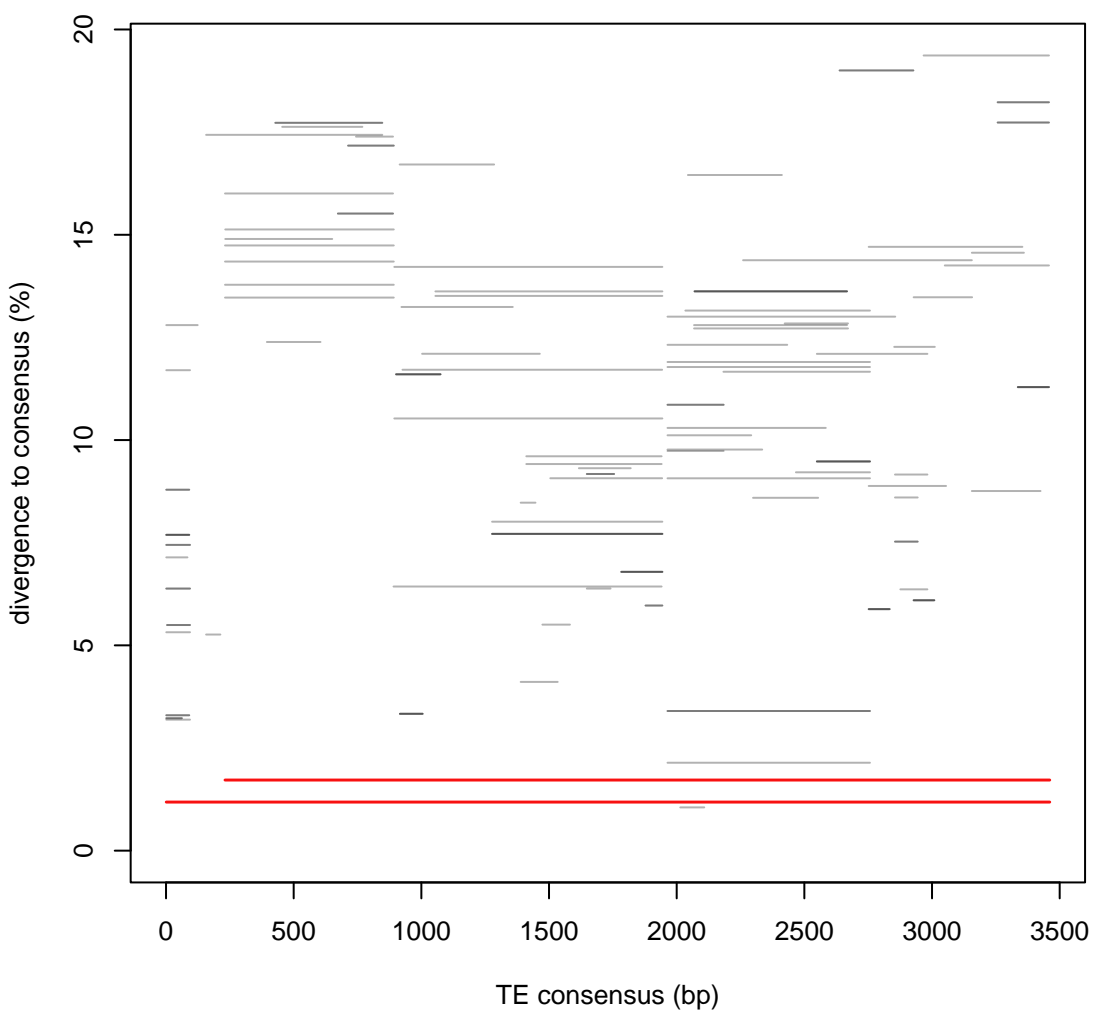
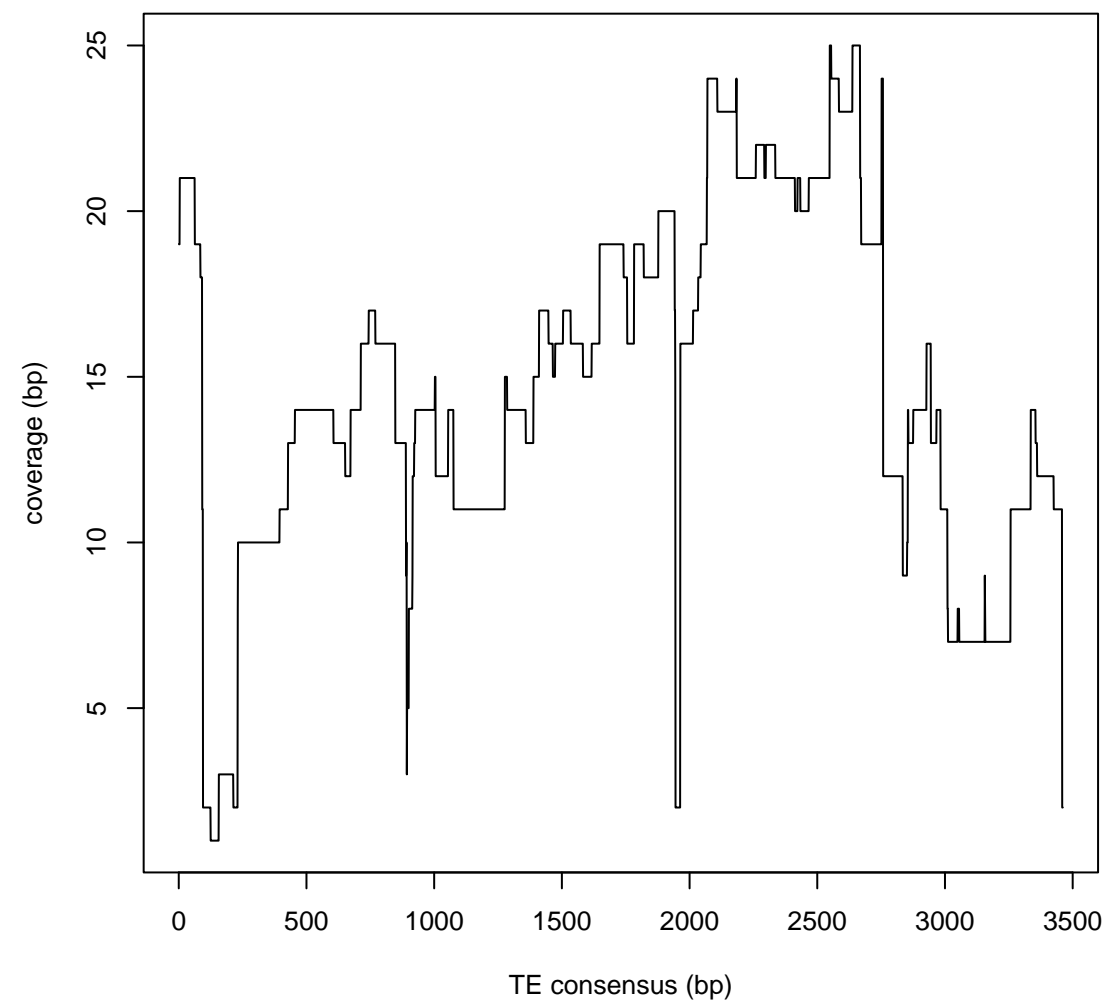


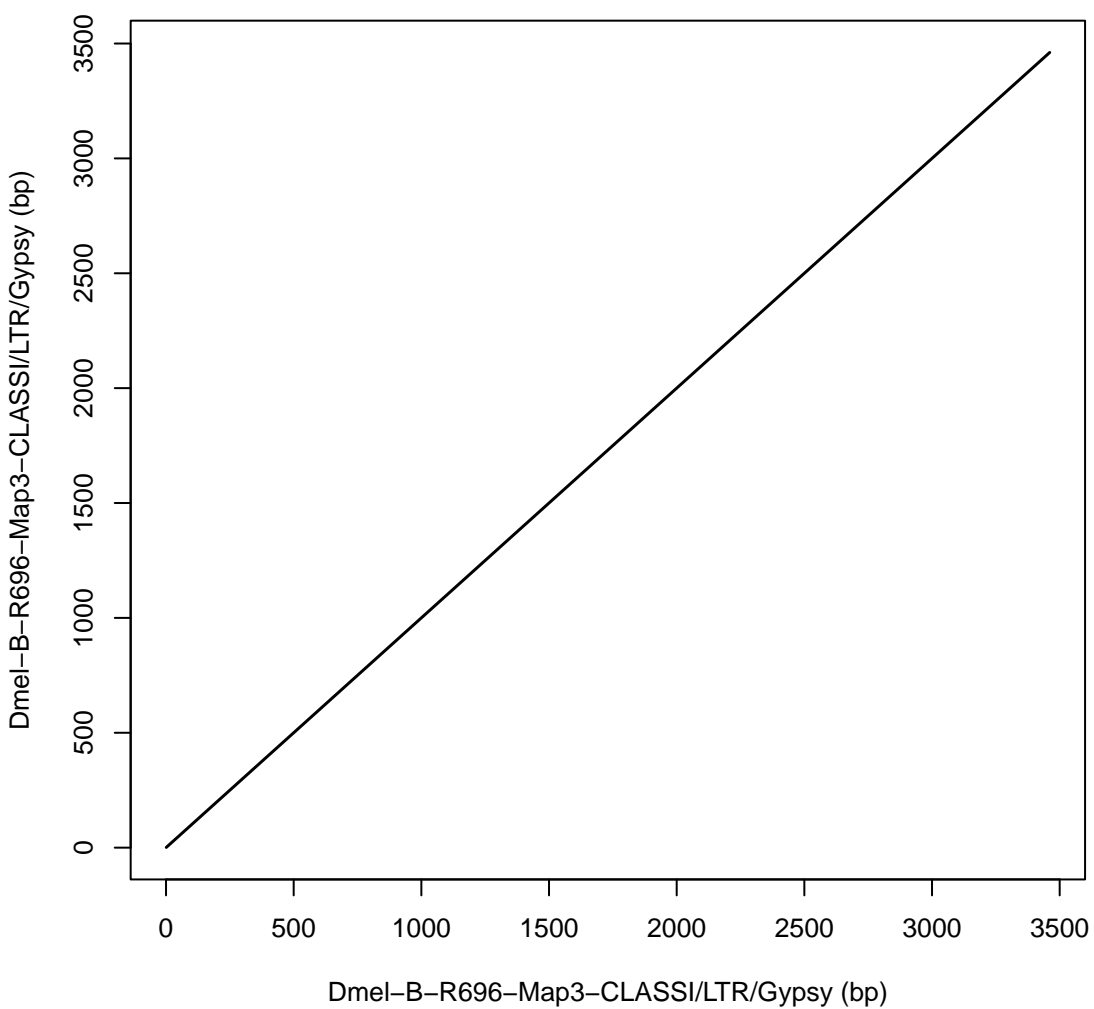
TE: Dmel-B-R696-Map3-CLASSI/LTR/Gypsy  
consensus size: 3461bp; fragments: 133; full length: 2 ( $\geq 3114.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

