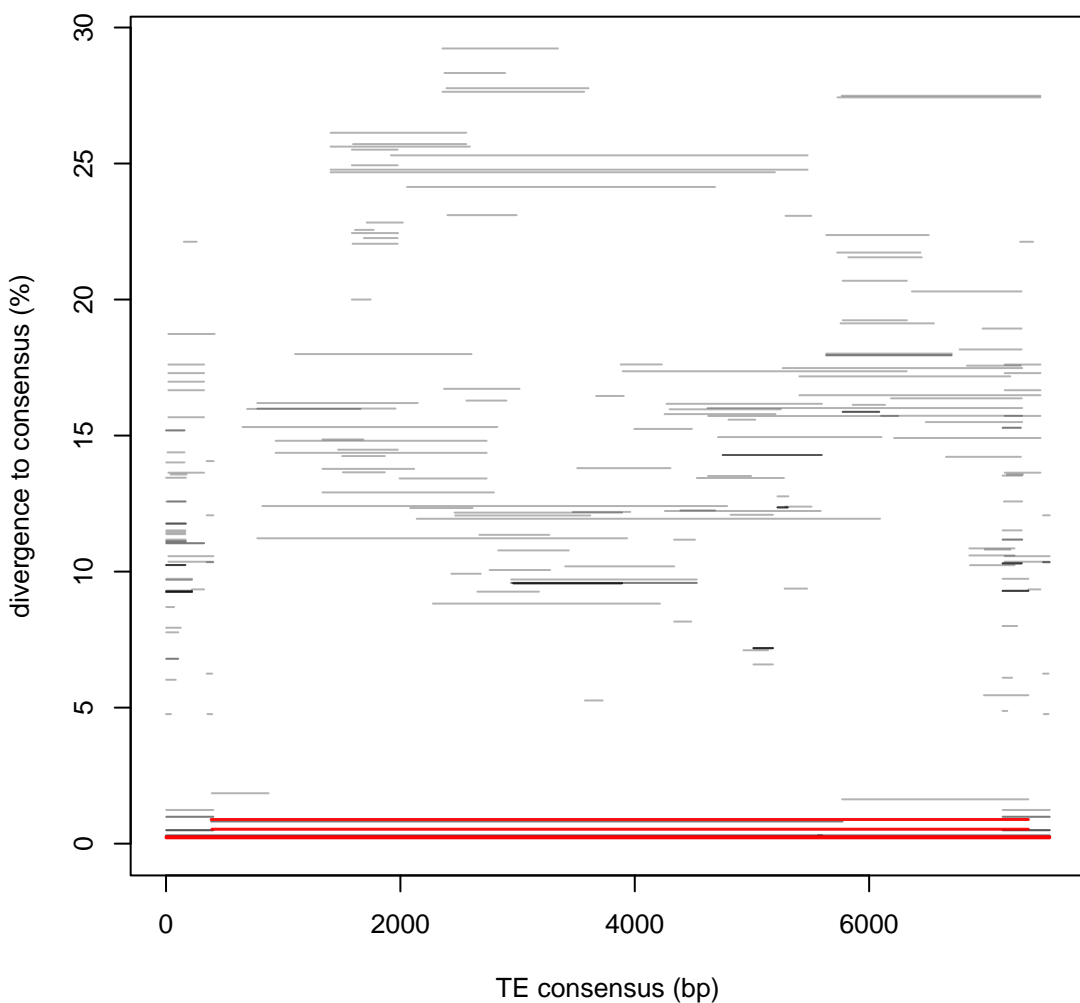
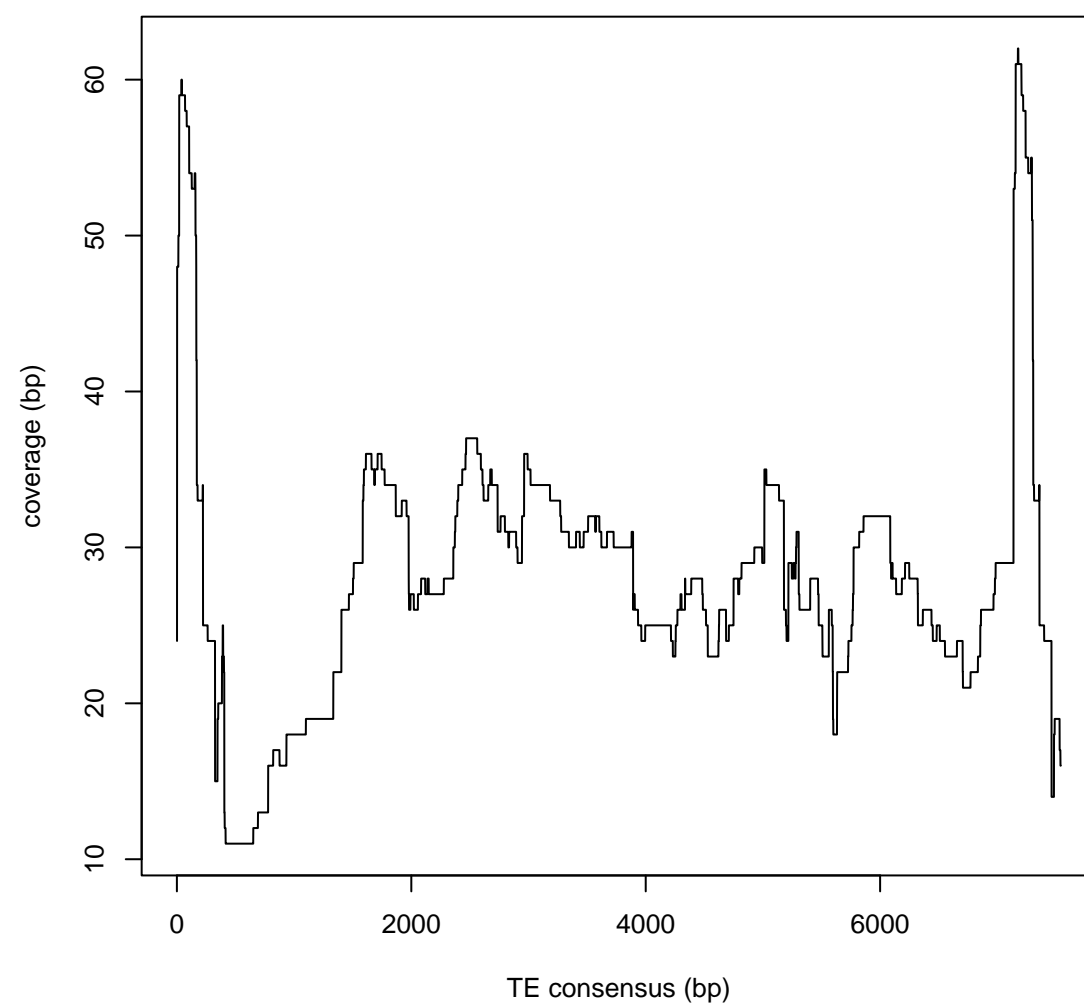


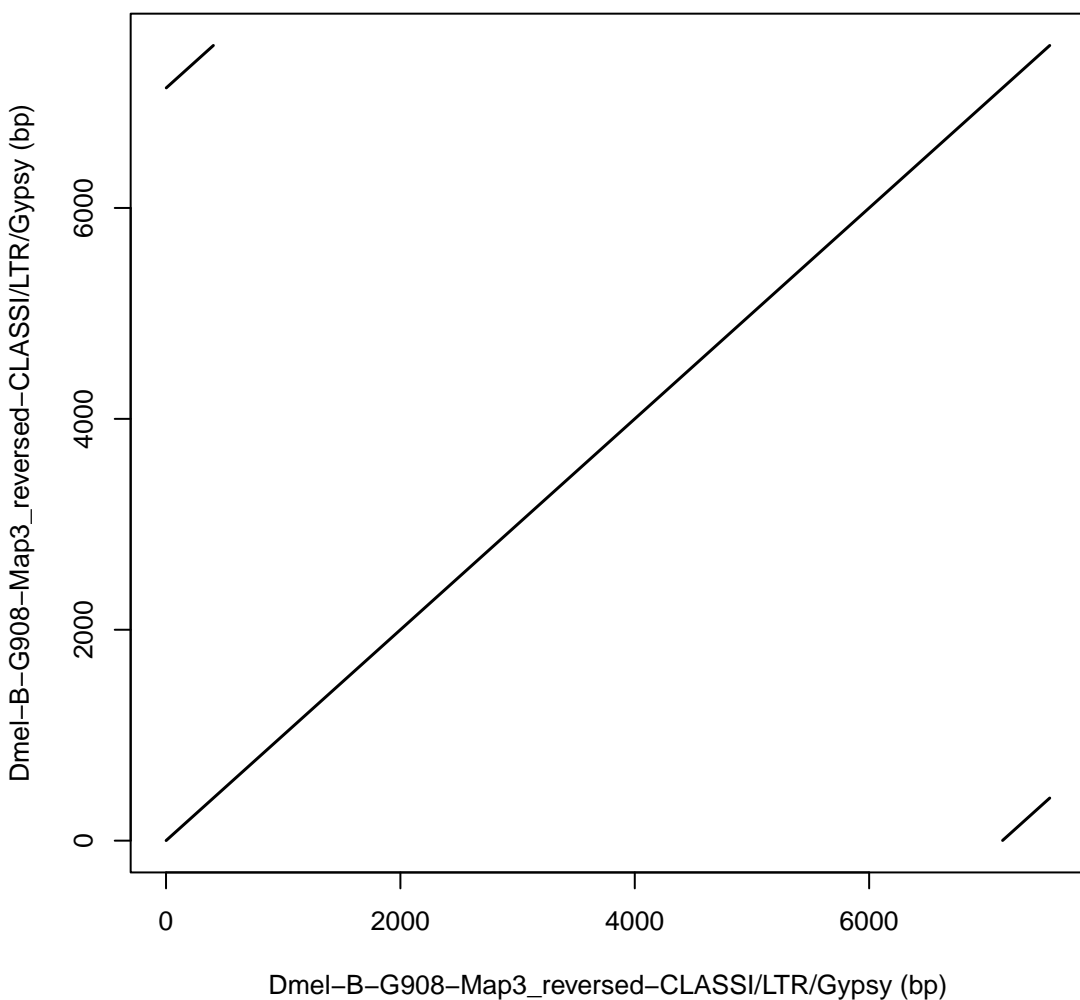
TE: Dmel-B-G908-Map3\_reversed-CLASSI/LTR/Gypsy  
consensus size: 7541bp; fragments: 243; full length: 5 ( $\geq 6786.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

