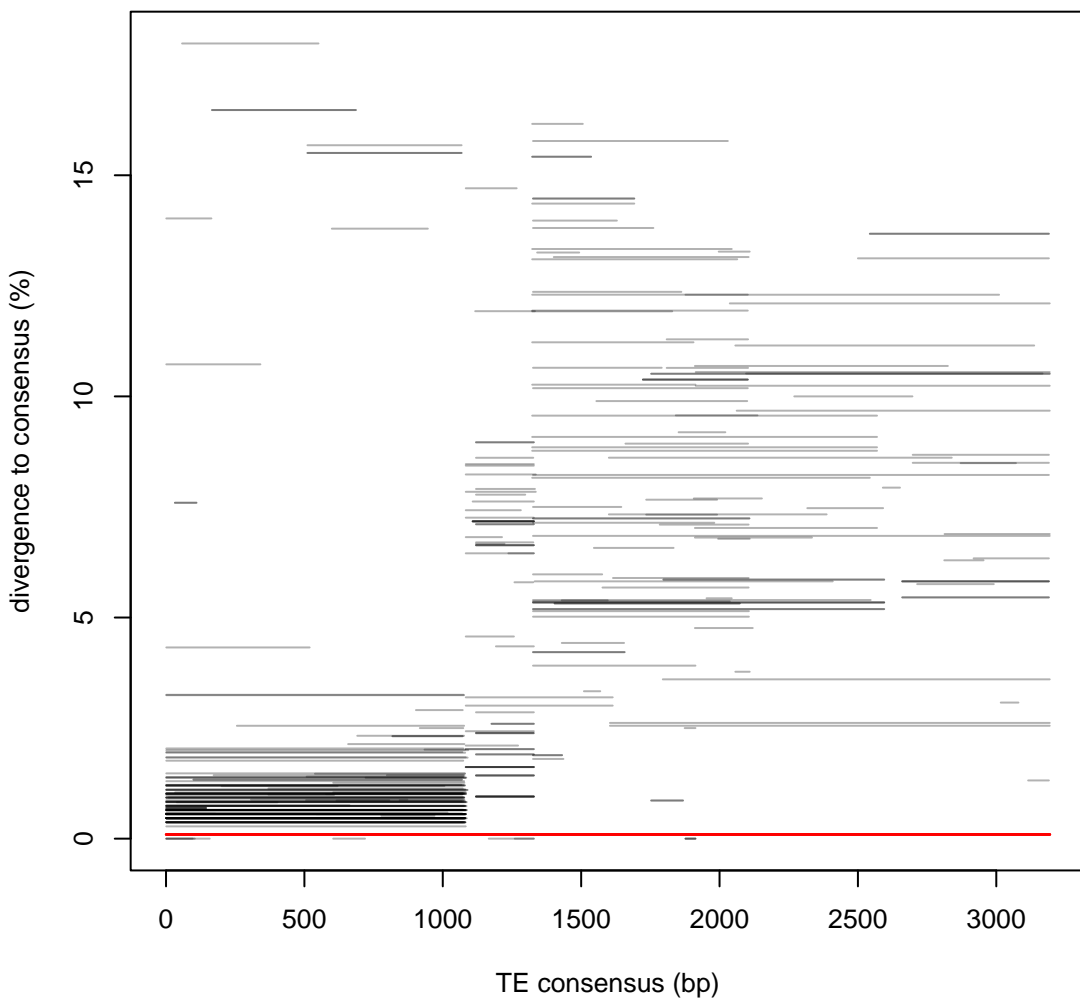
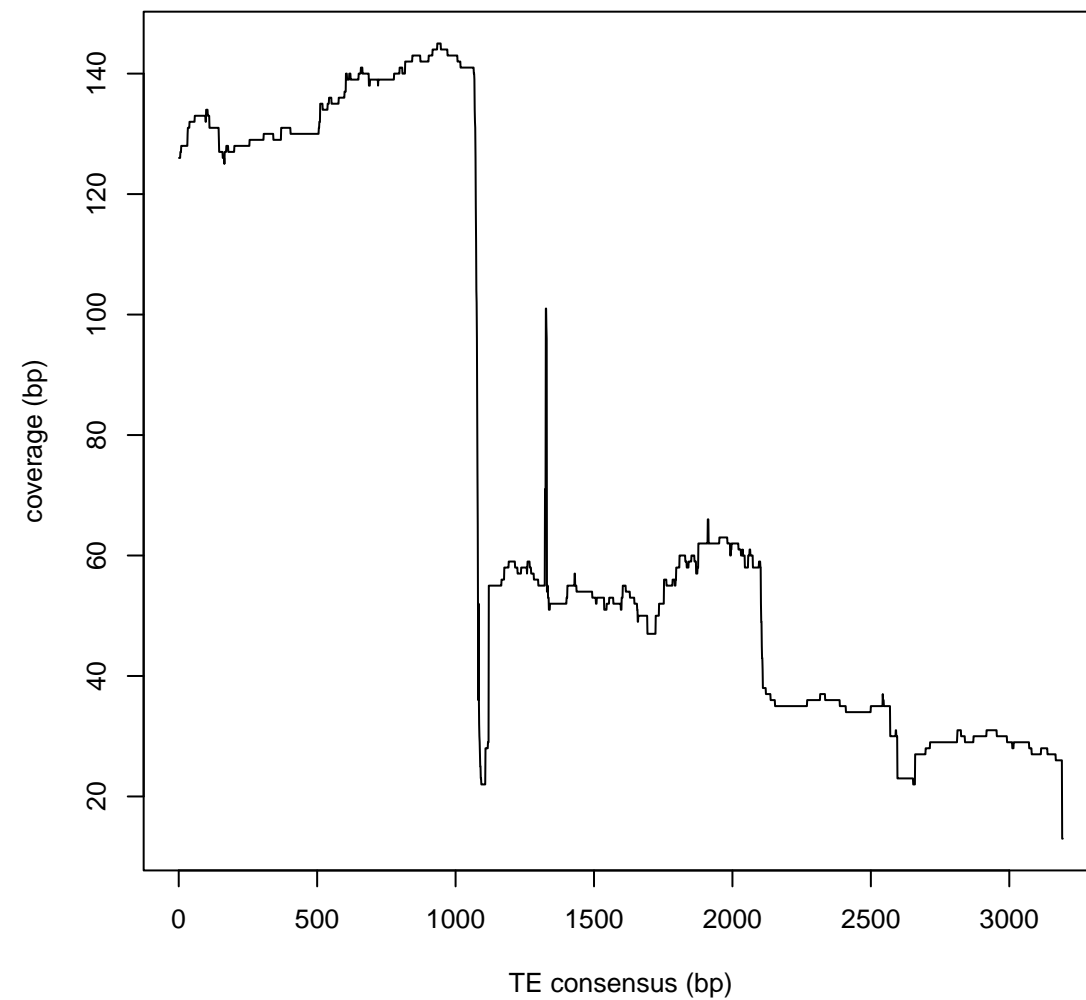


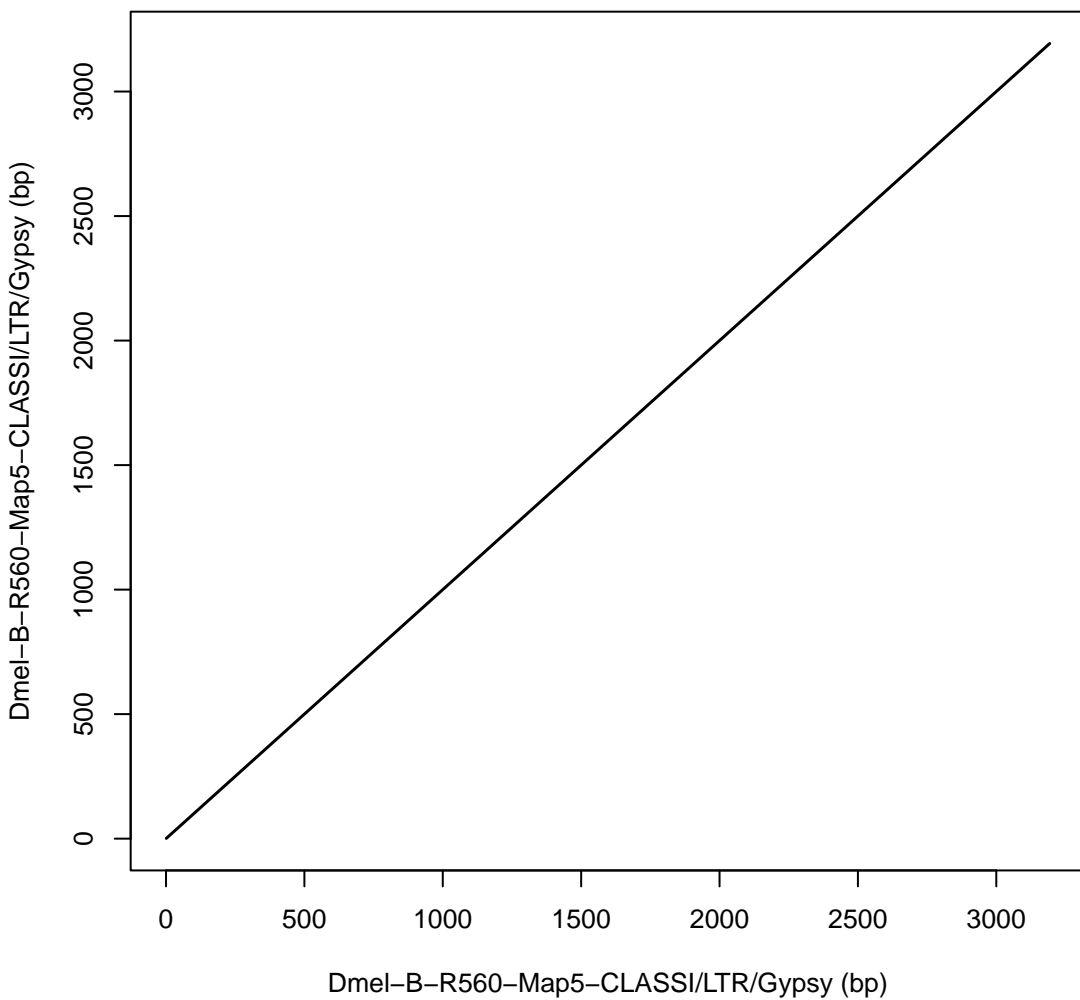
TE: Dmel-B-R560-Map5-CLASSI/LTR/Gypsy  
consensus size: 3193bp; fragments: 348; full length: 2 ( $\geq 2873.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

