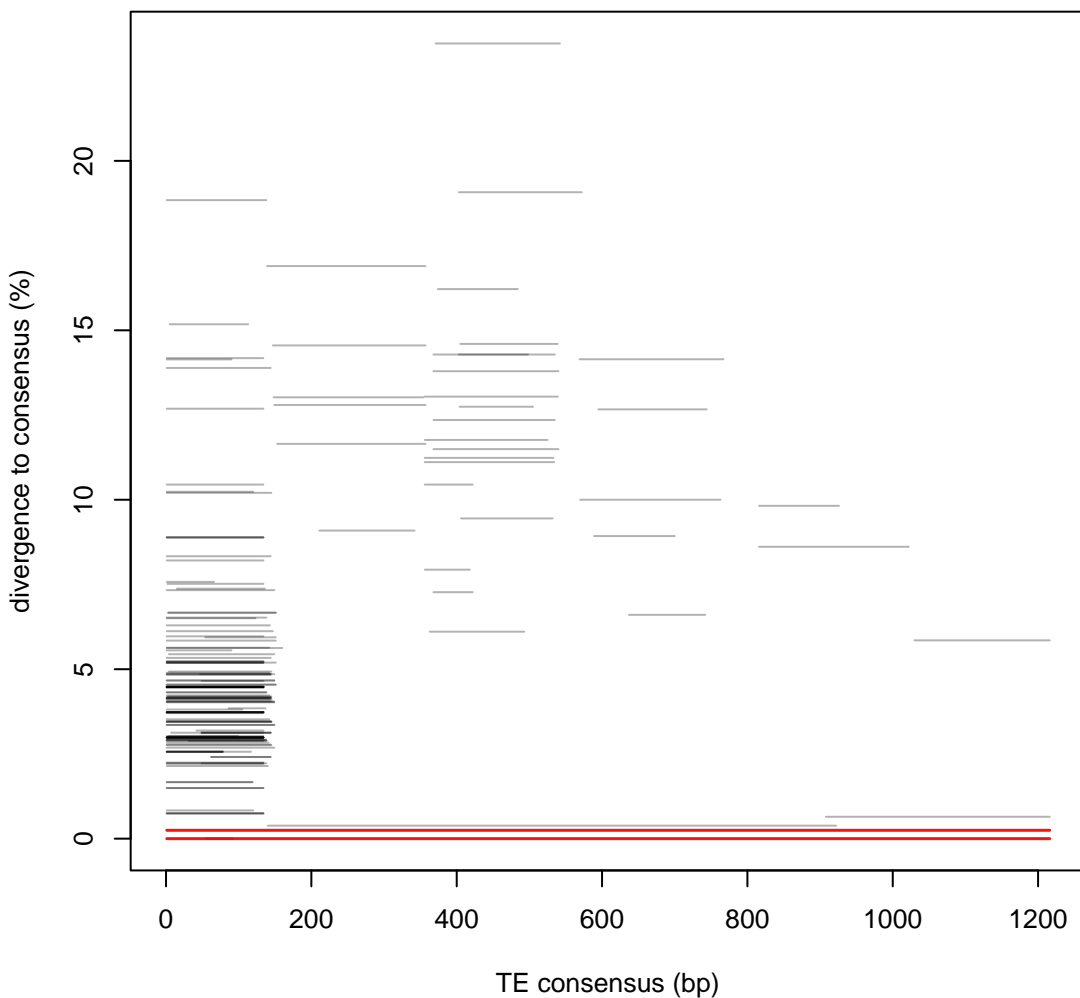
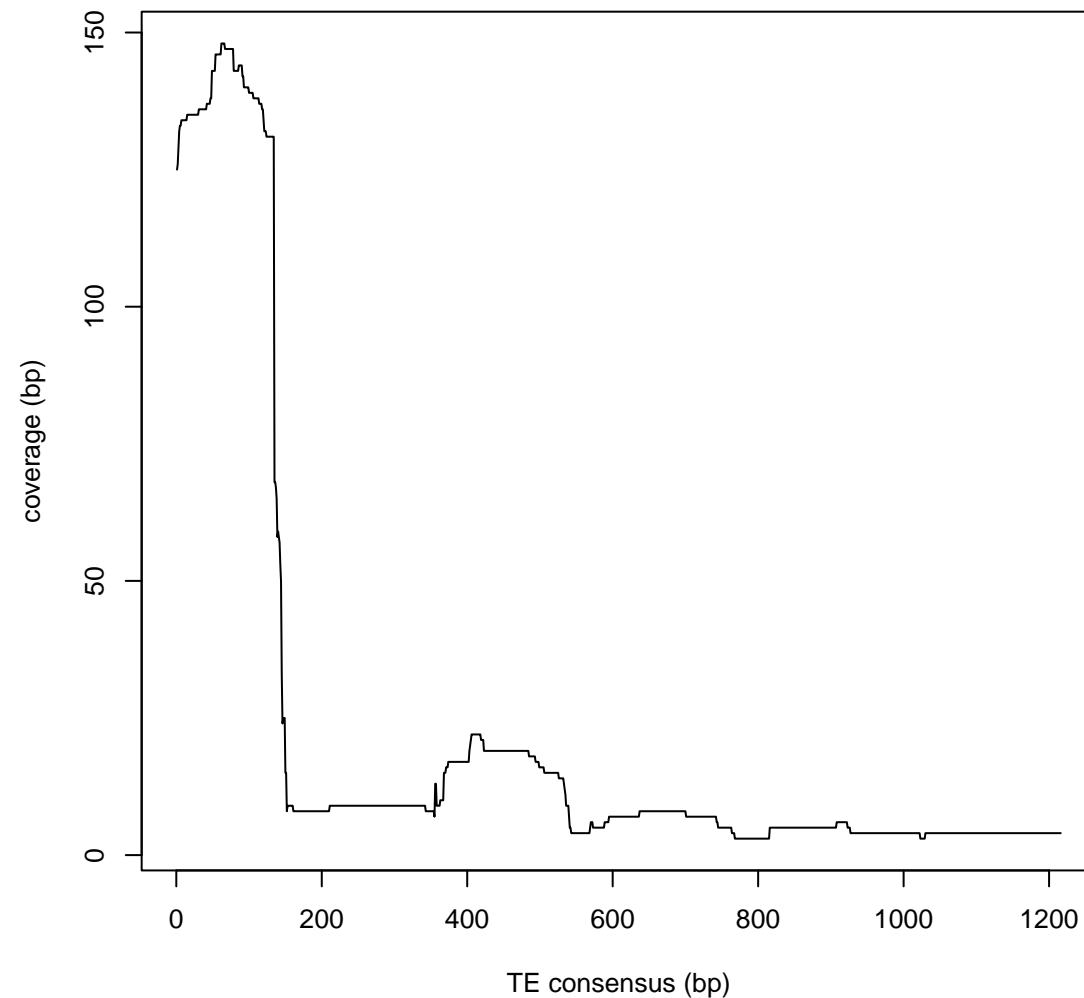


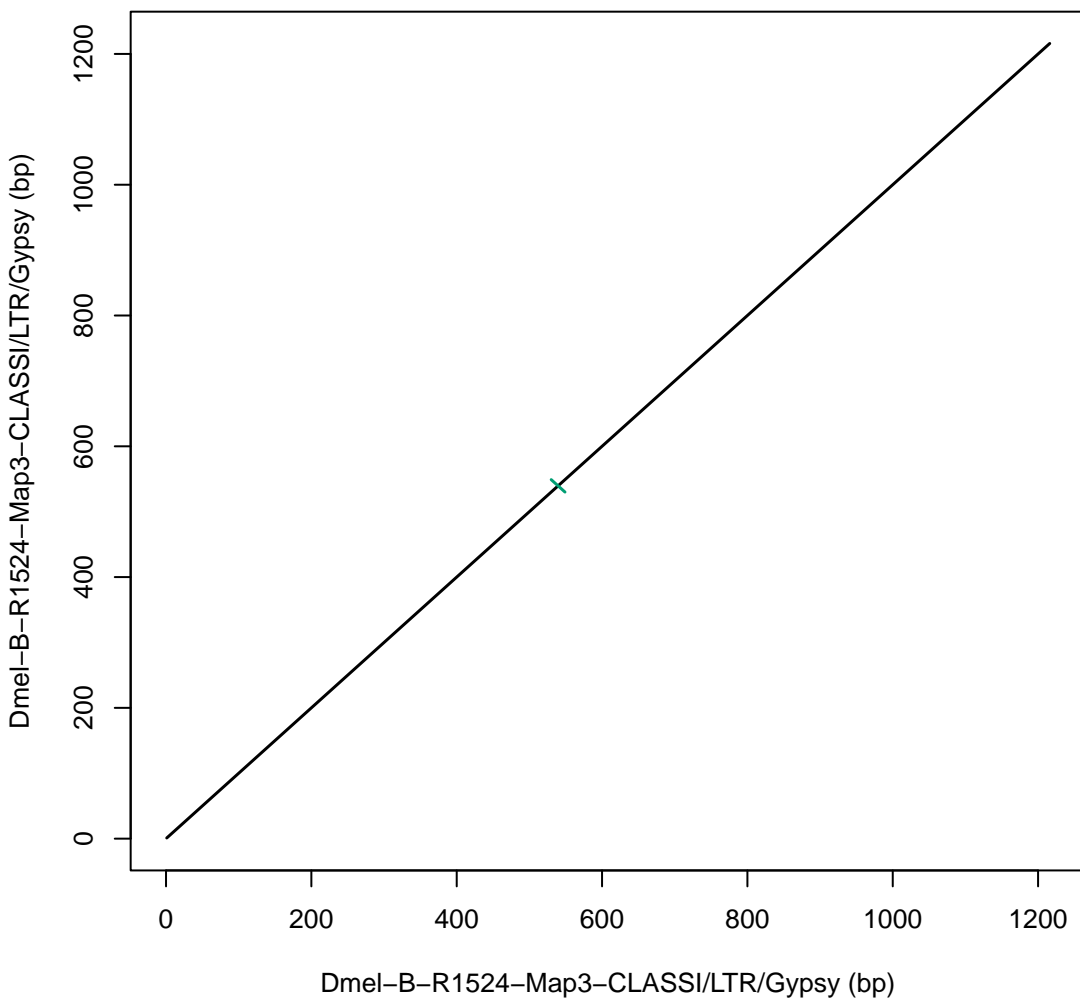
TE: Dmel-B-R1524-Map3-CLASSI/LTR/Gypsy  
consensus size: 1216bp; fragments: 184; full length: 2 ( $\geq 1094.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

