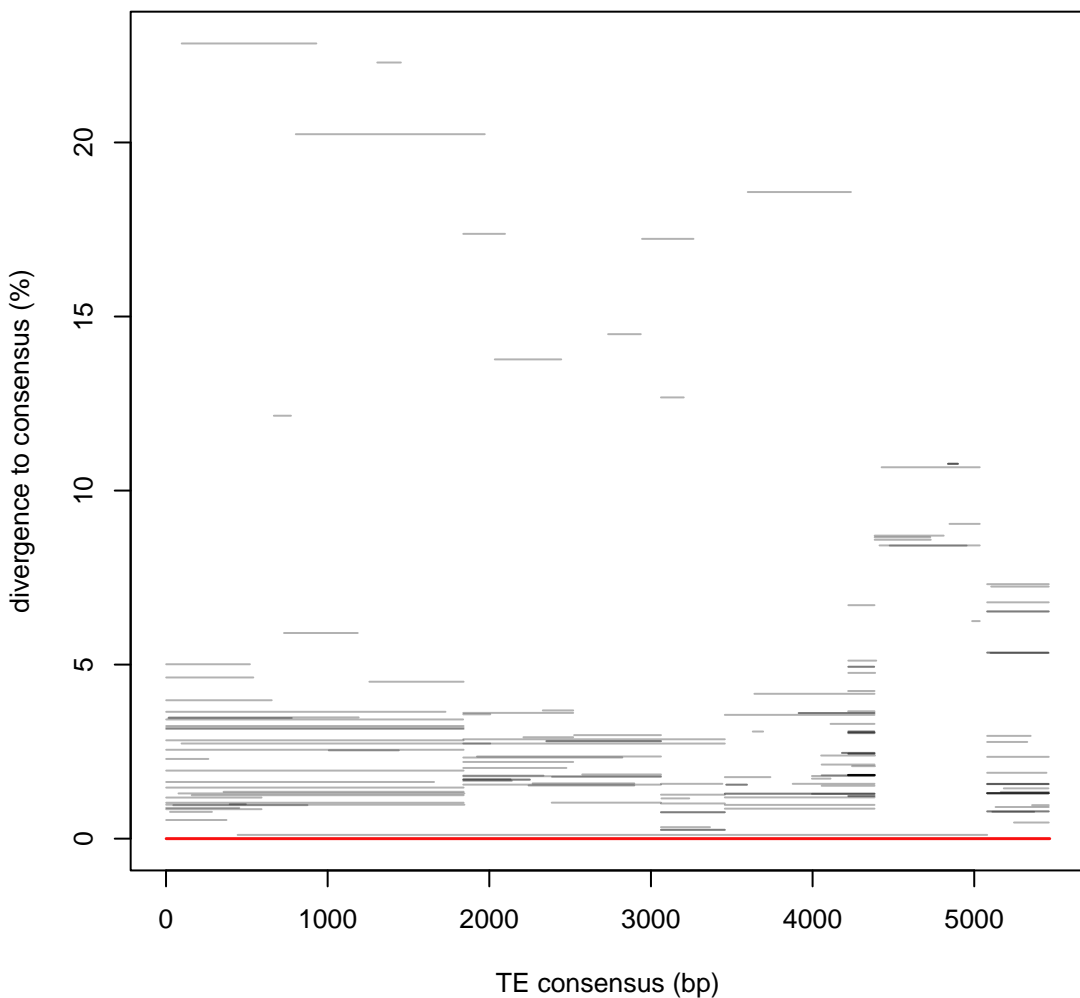
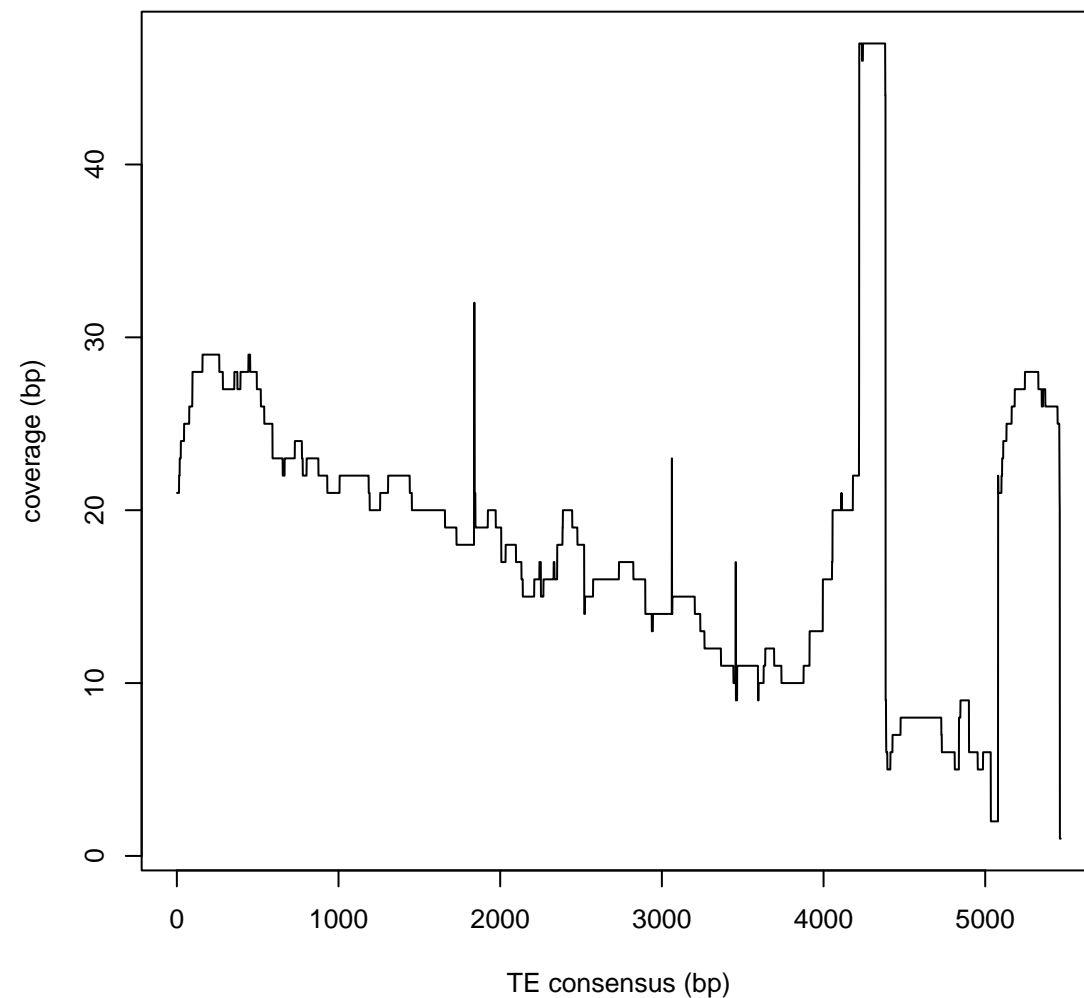


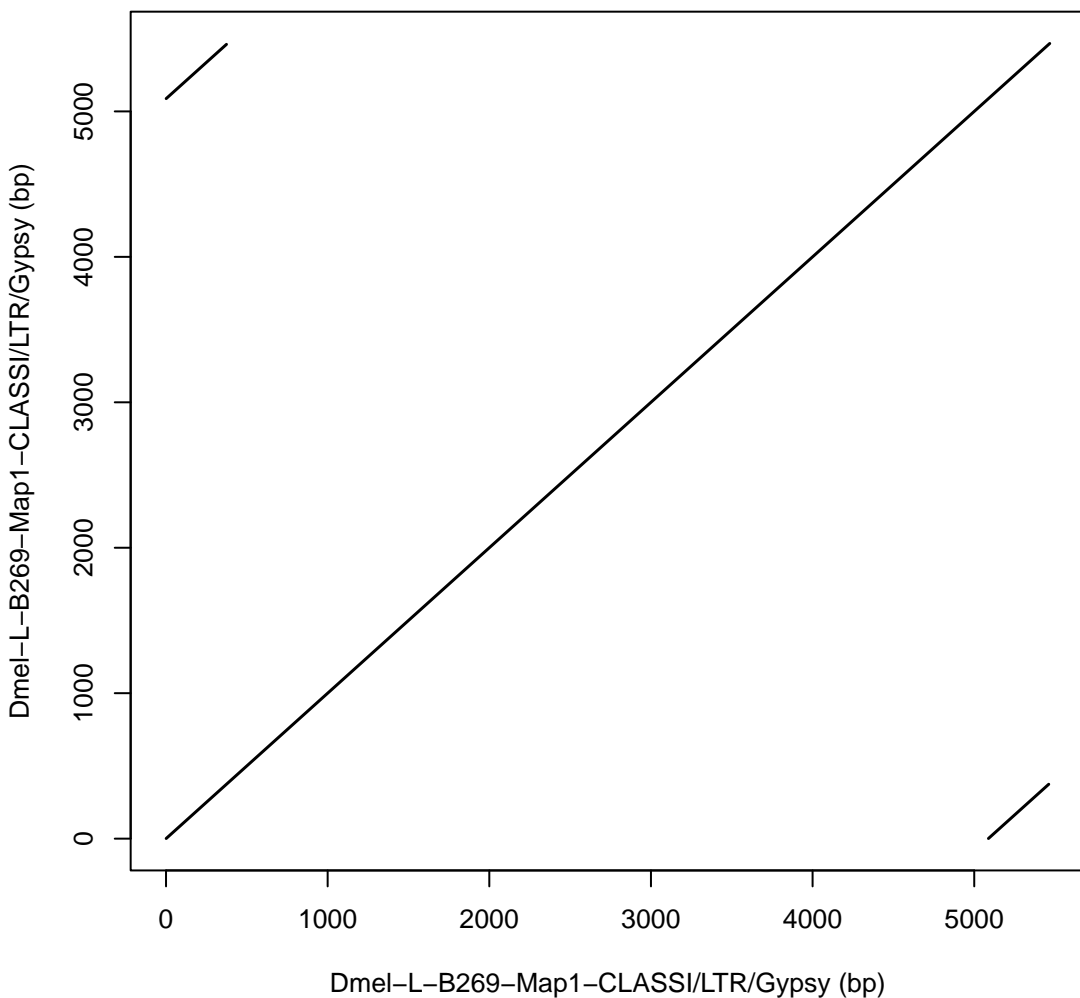
TE: Dmel-L-B269-Map1-CLASSI/LTR/Gypsy  
consensus size: 5467bp; fragments: 167; full length: 1 ( $\geq 4920.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

