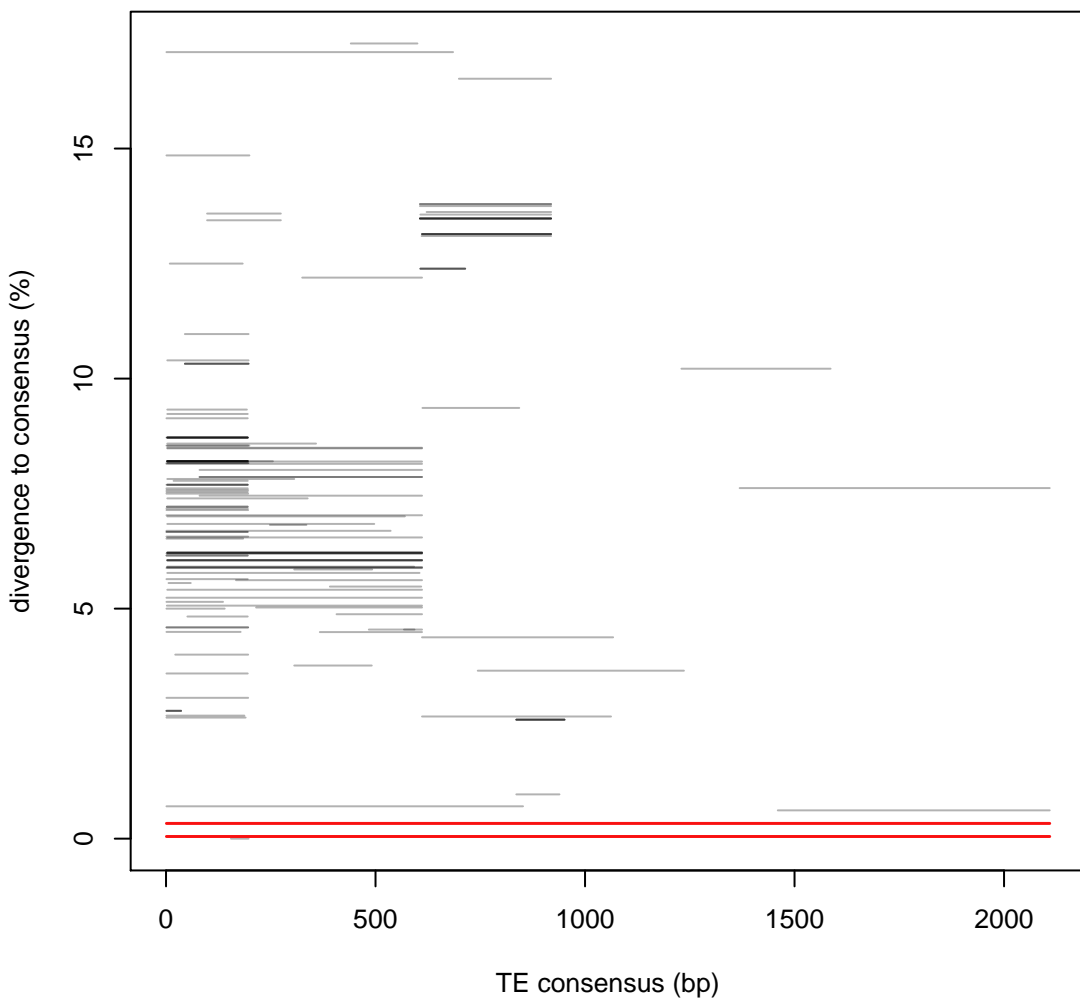
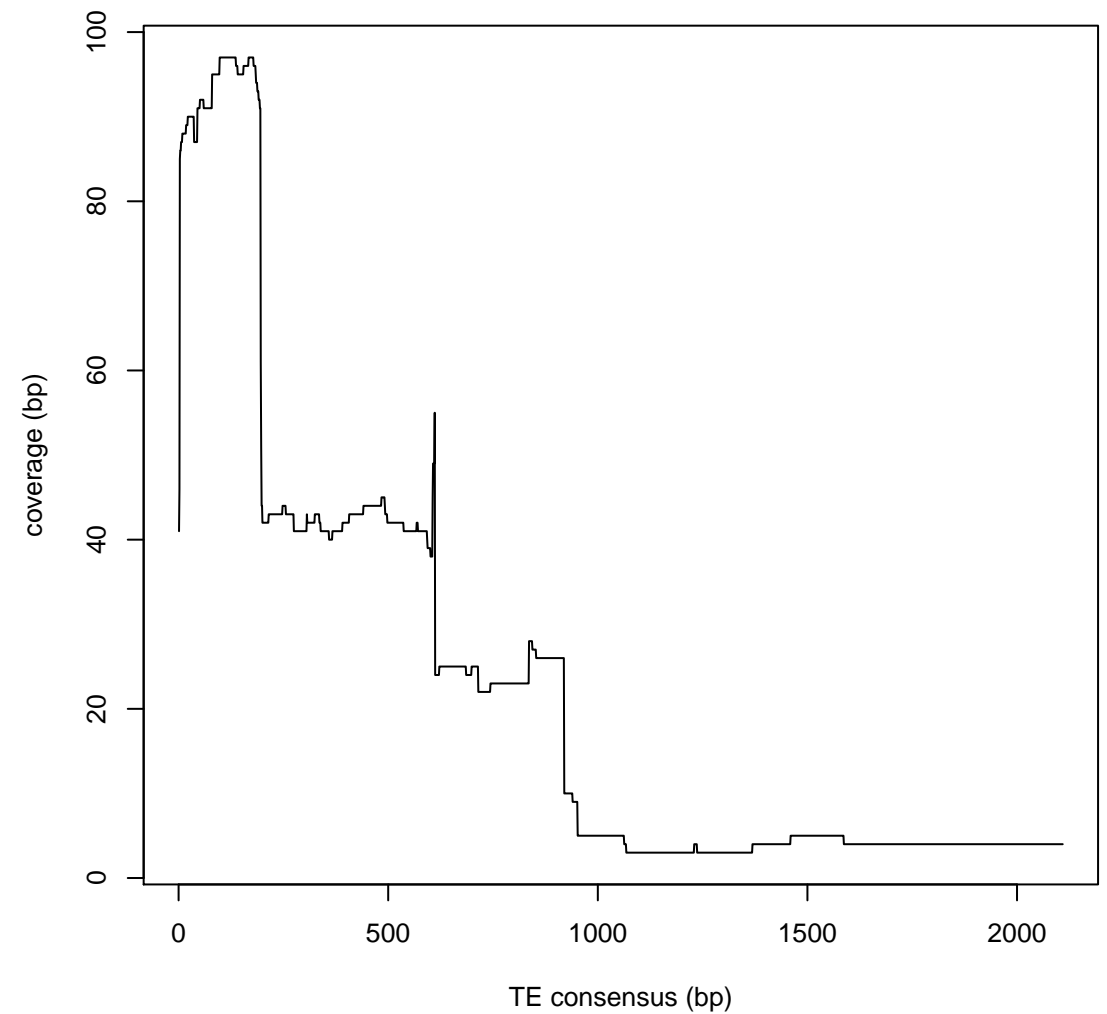


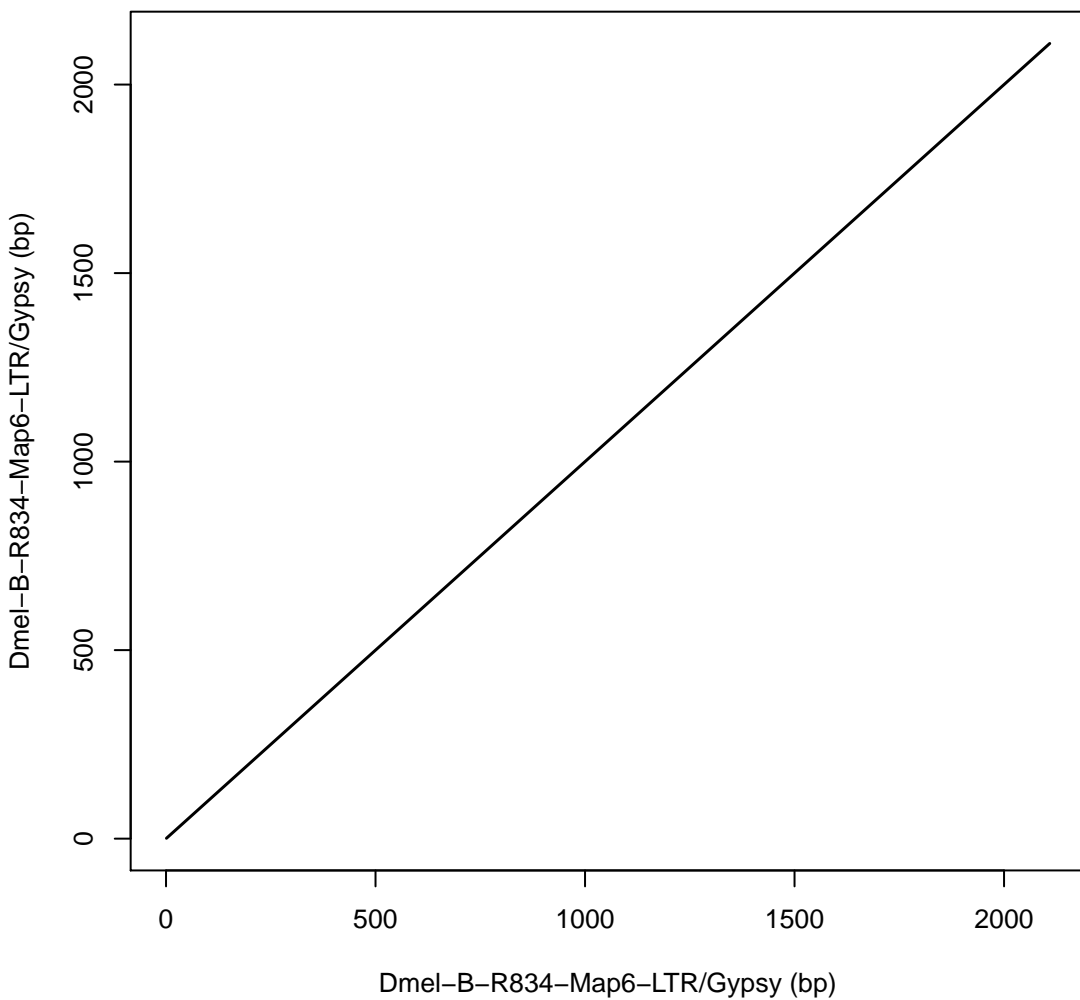
TE: Dmel-B-R834-Map6-LTR/Gypsy  
consensus size: 2109bp; fragments: 145; full length: 2 ( $\geq 1898.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

