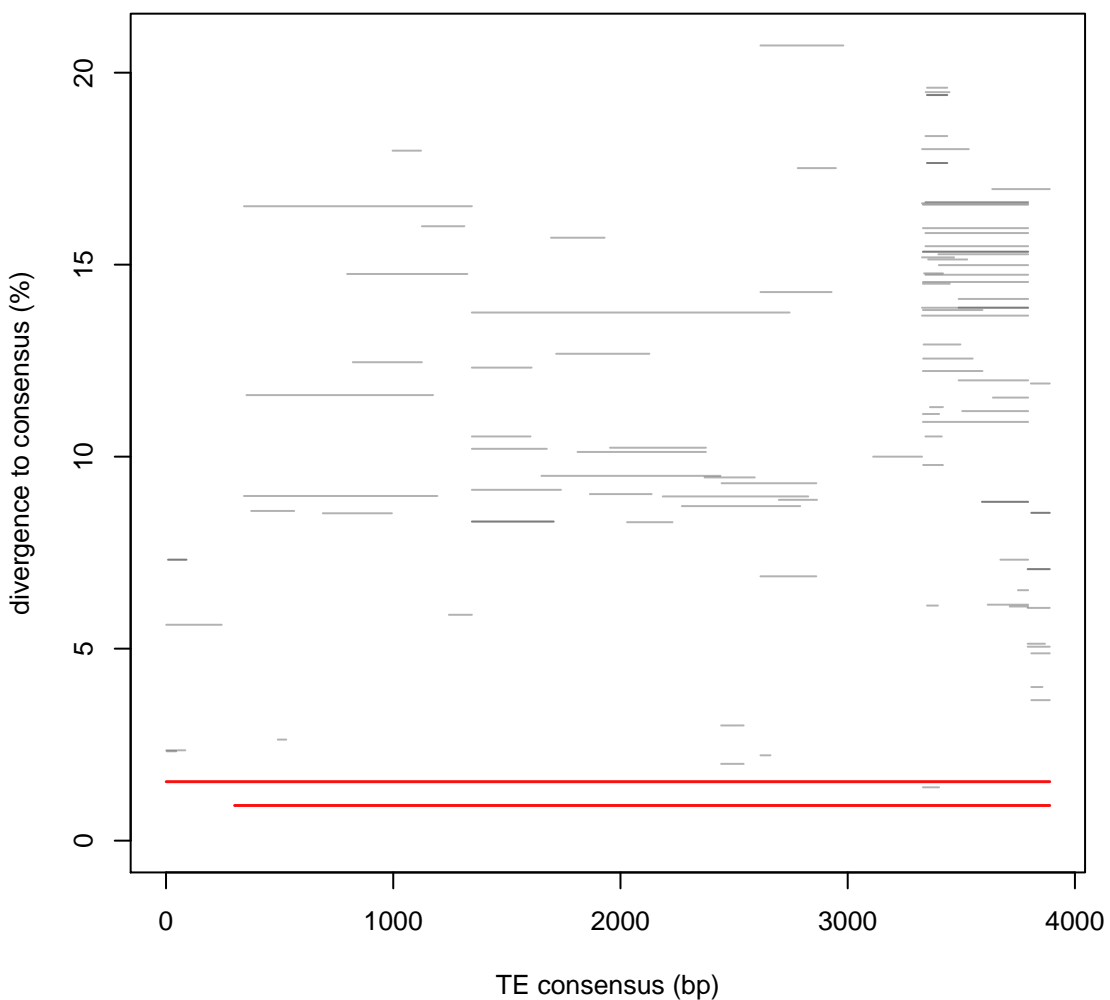
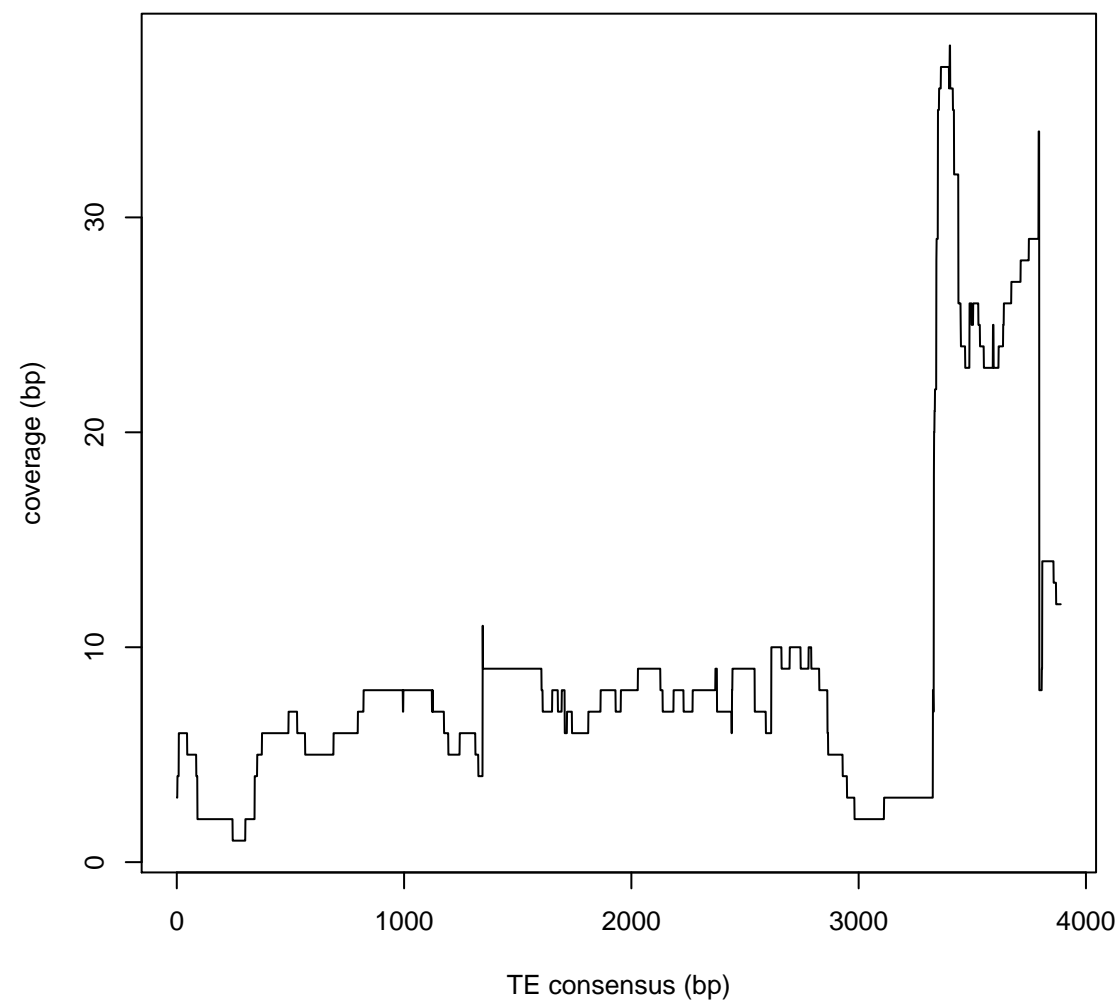


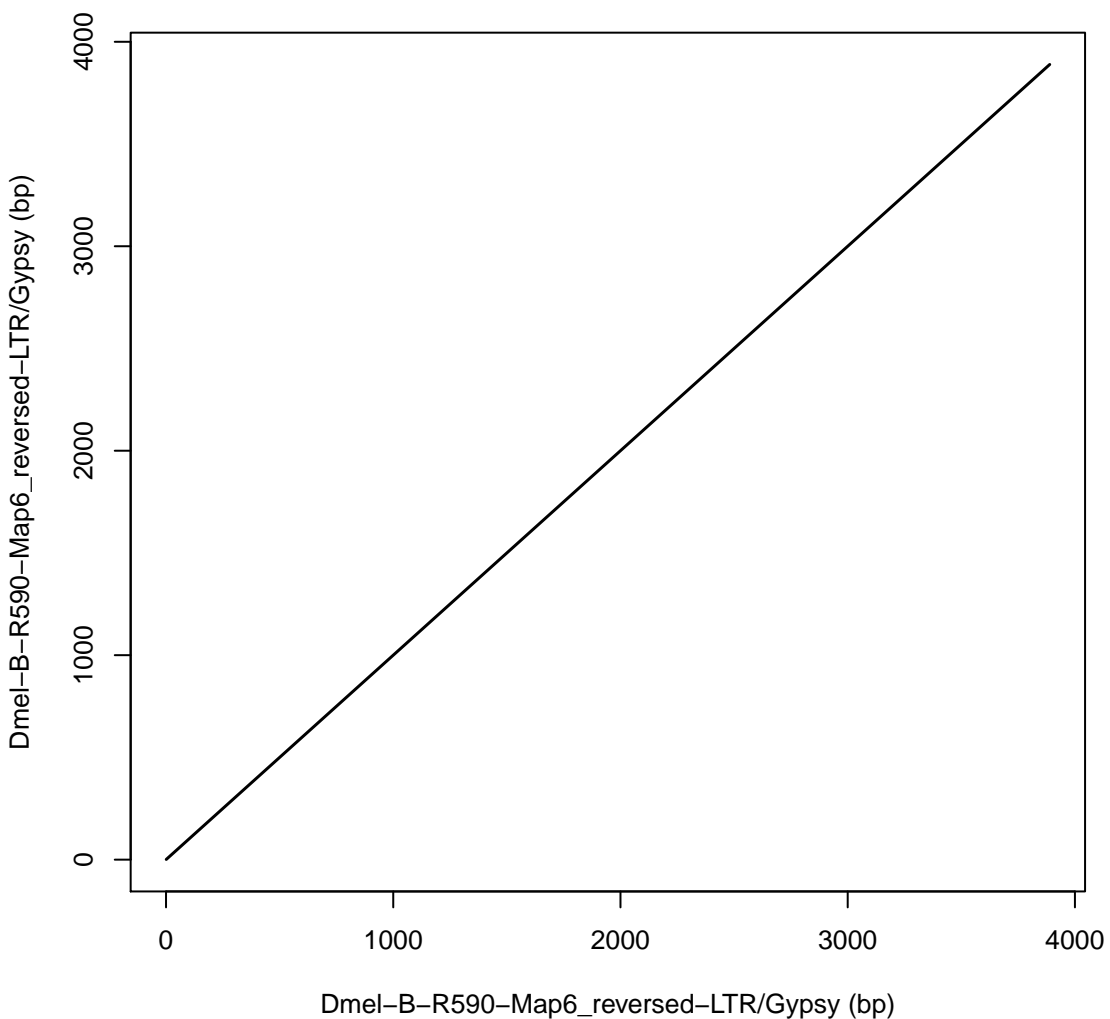
TE: Dmel-B-R590-Map6\_reversed-LTR/Gypsy  
consensus size: 3889bp; fragments: 105; full length: 2 ( $\geq 3500.1$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

