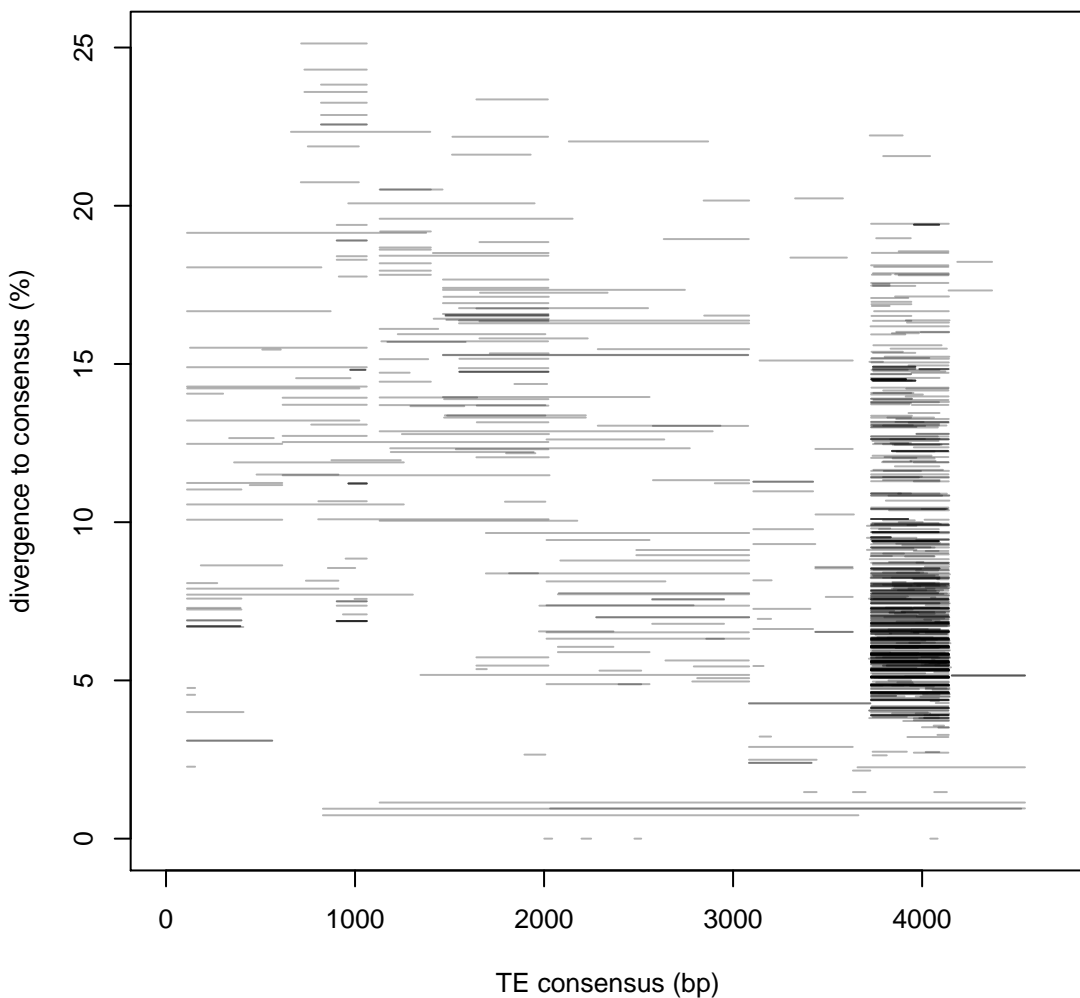
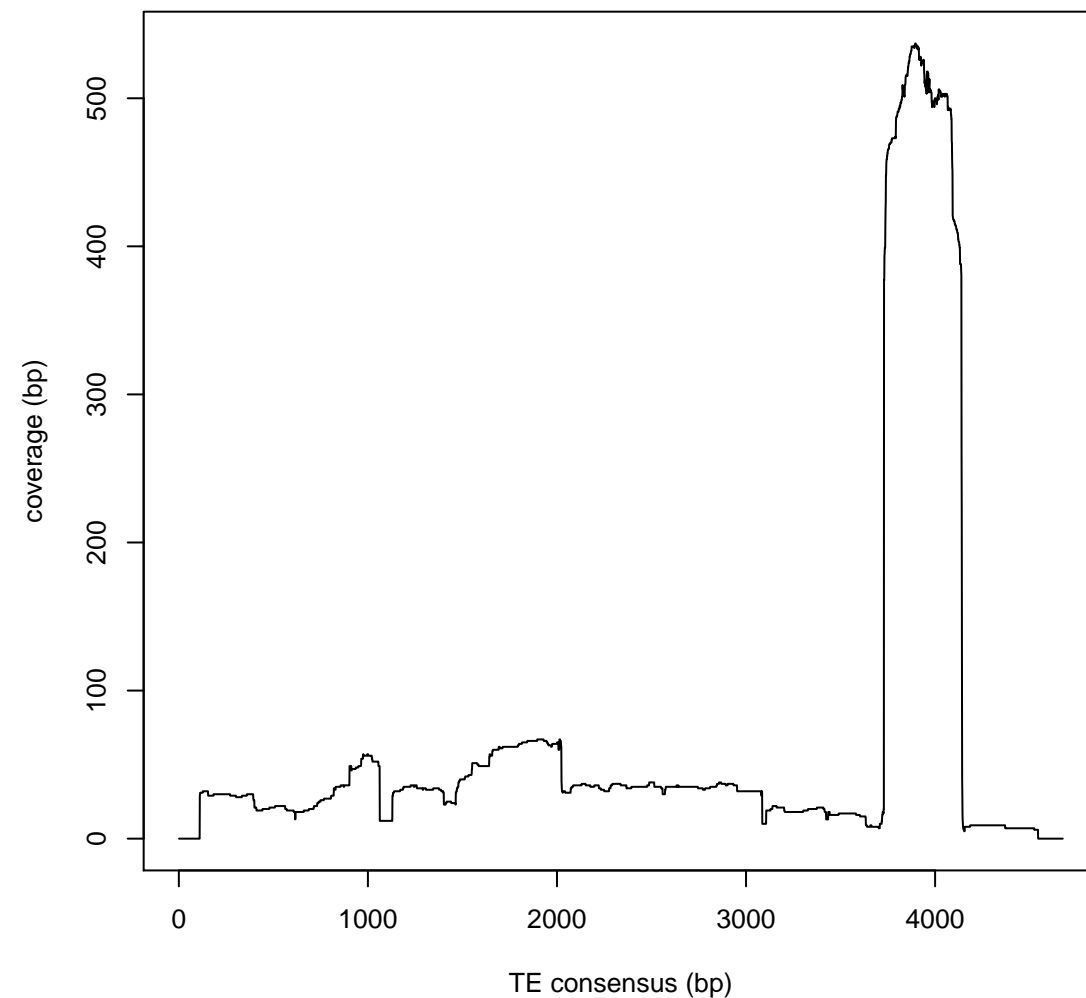


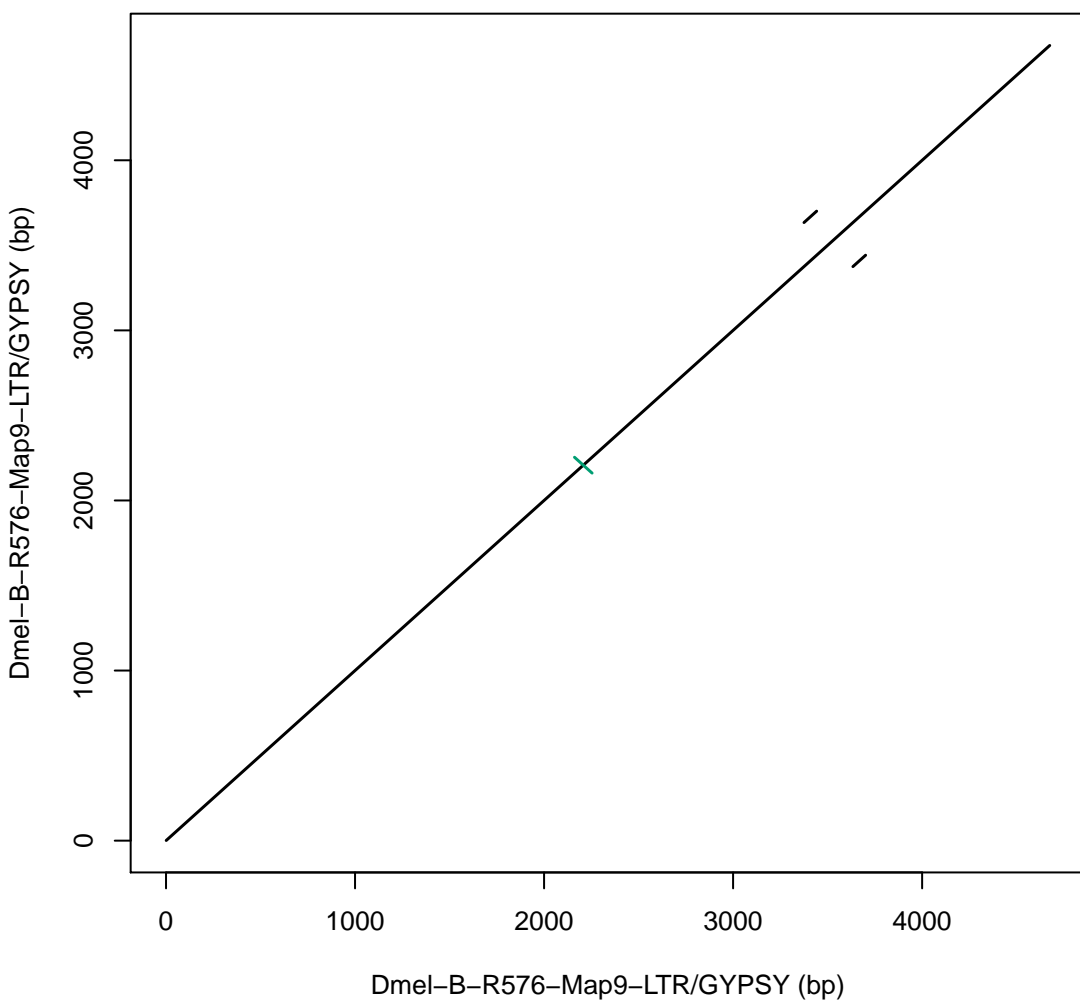
TE: Dmel-B-R576-Map9-LTR/GYPSY  
consensus size: 4675bp; fragments: 891; full length: 0 ( $\geq 4207.5$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

