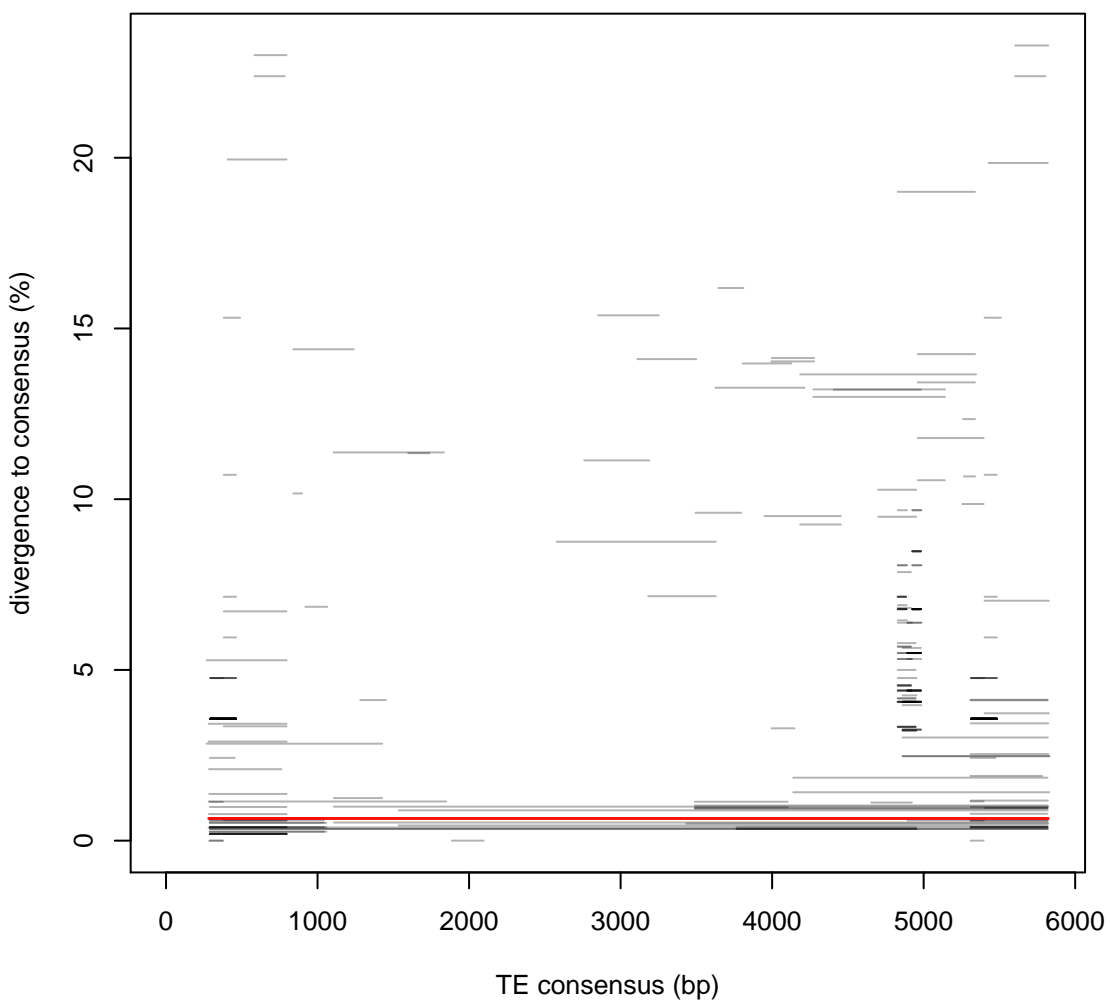
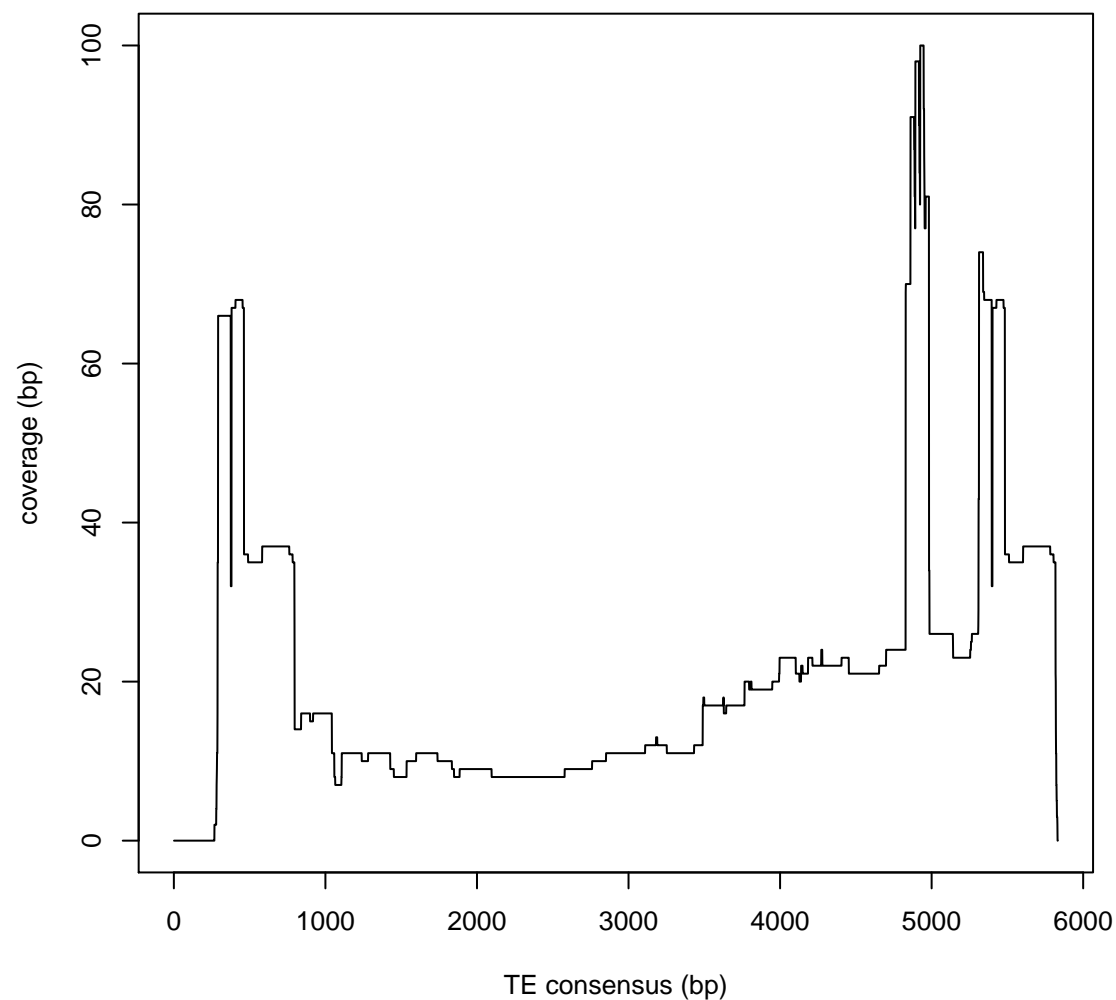


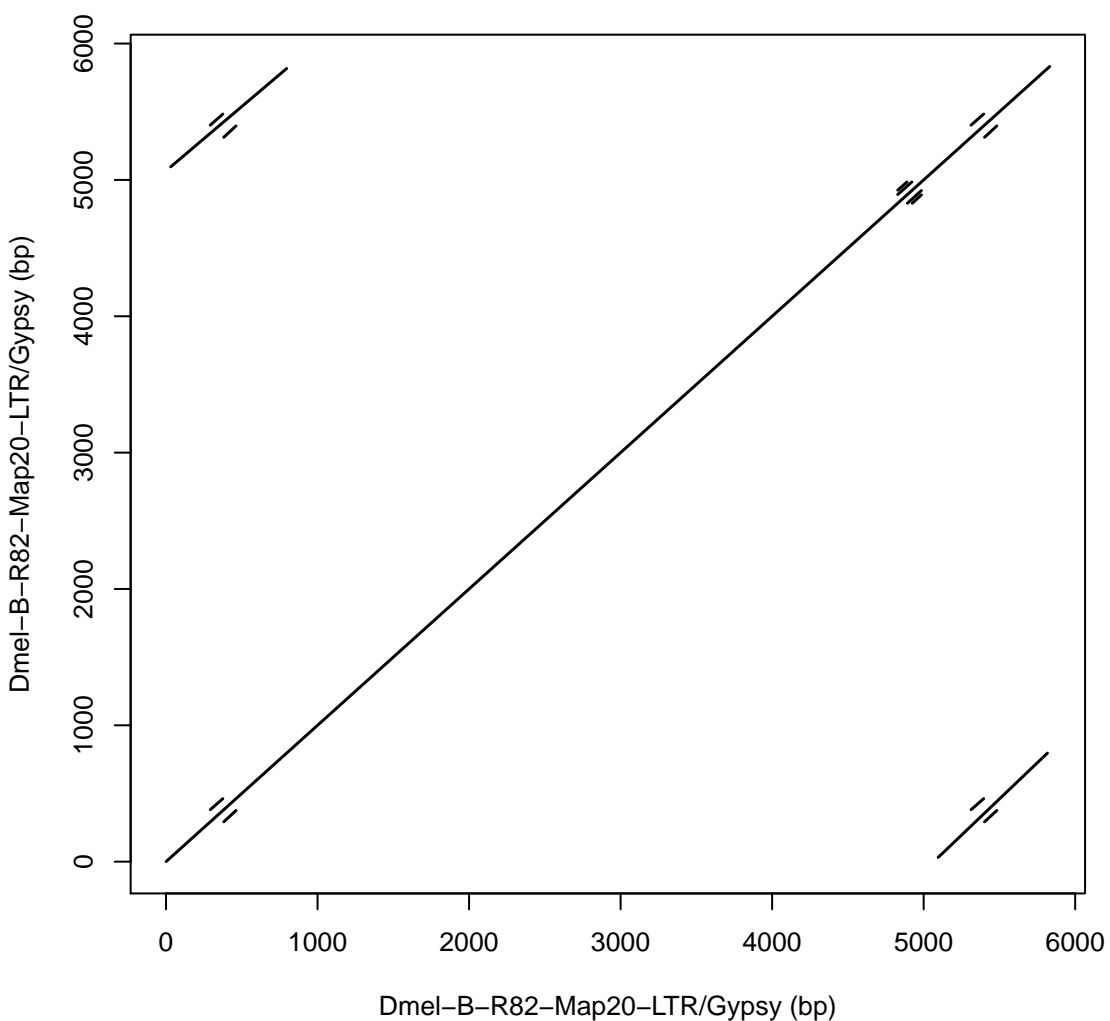
TE: Dmel-B-R82-Map20-LTR/Gypsy  
consensus size: 5832bp; fragments: 349; full length: 1 ( $\geq 5248.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

