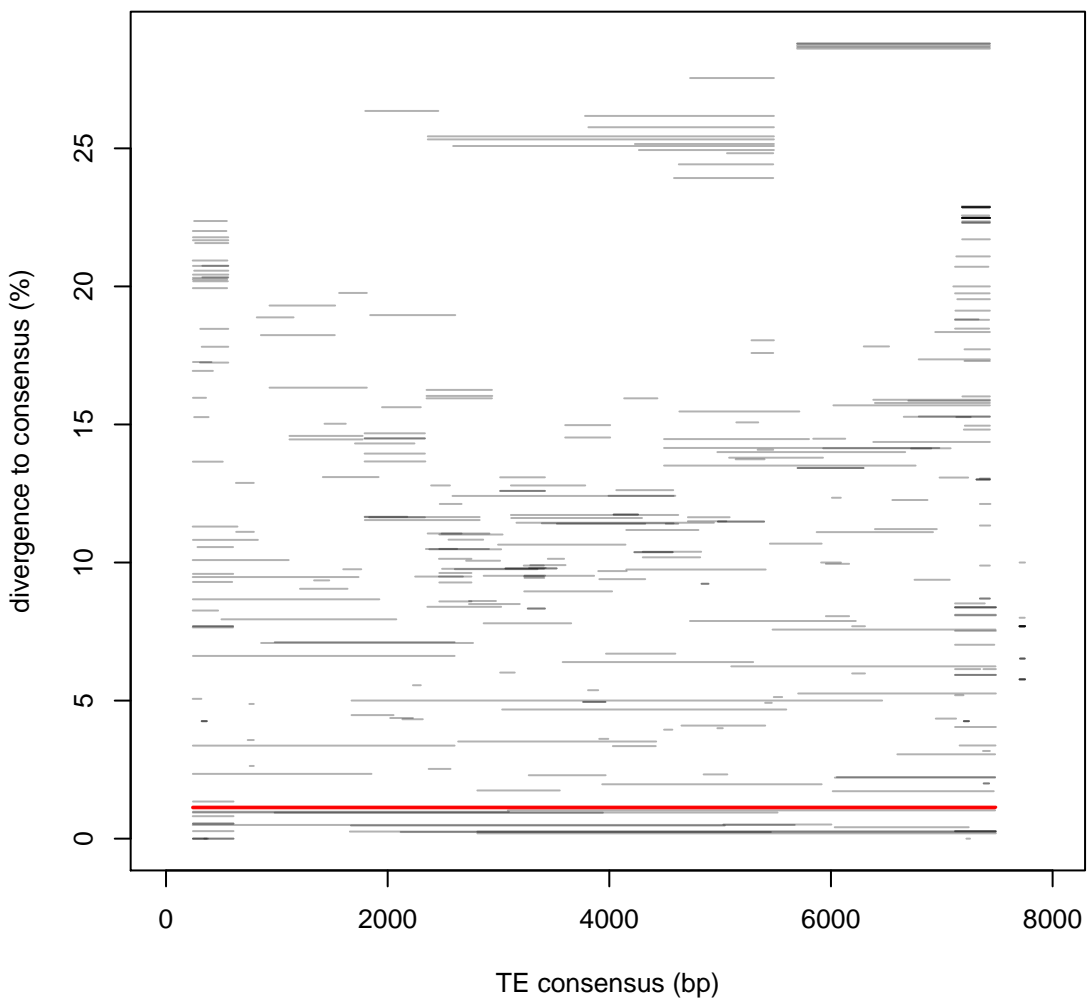
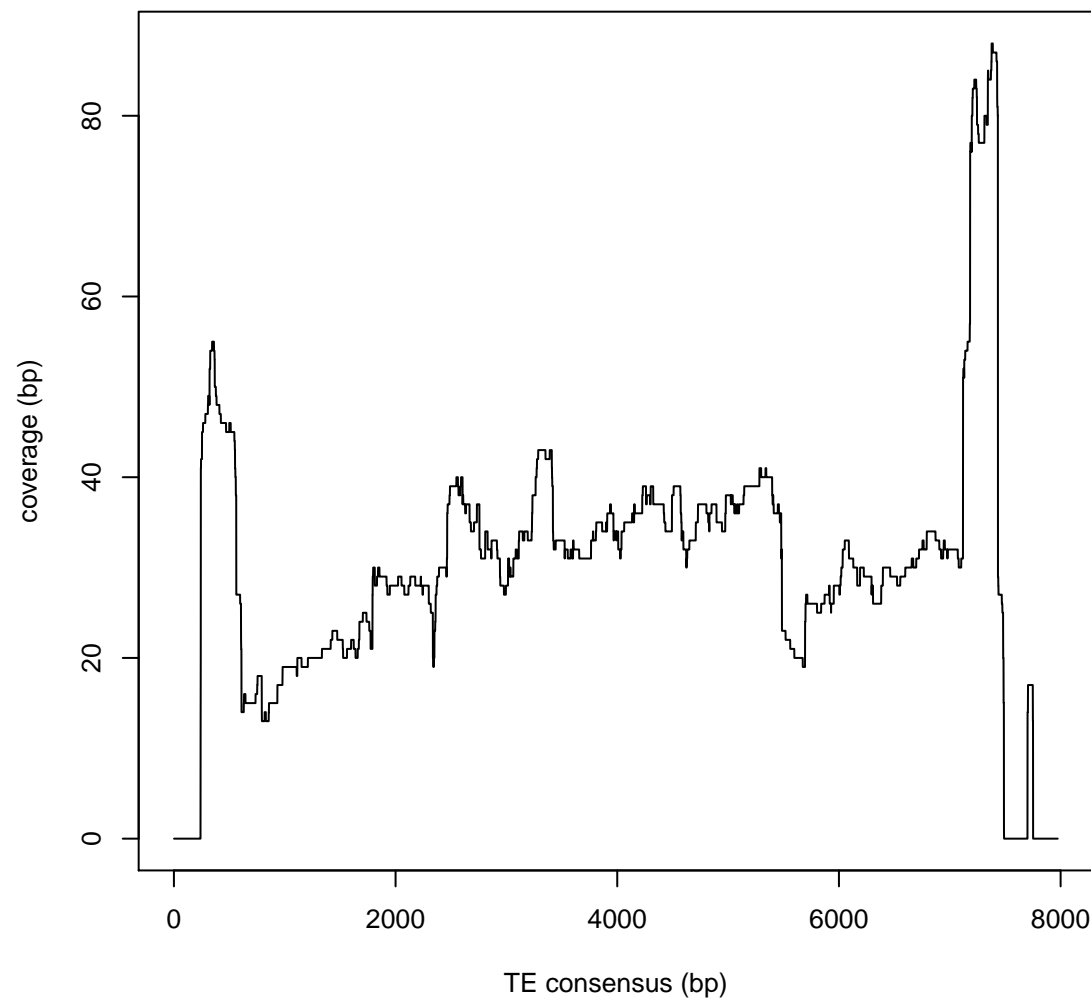


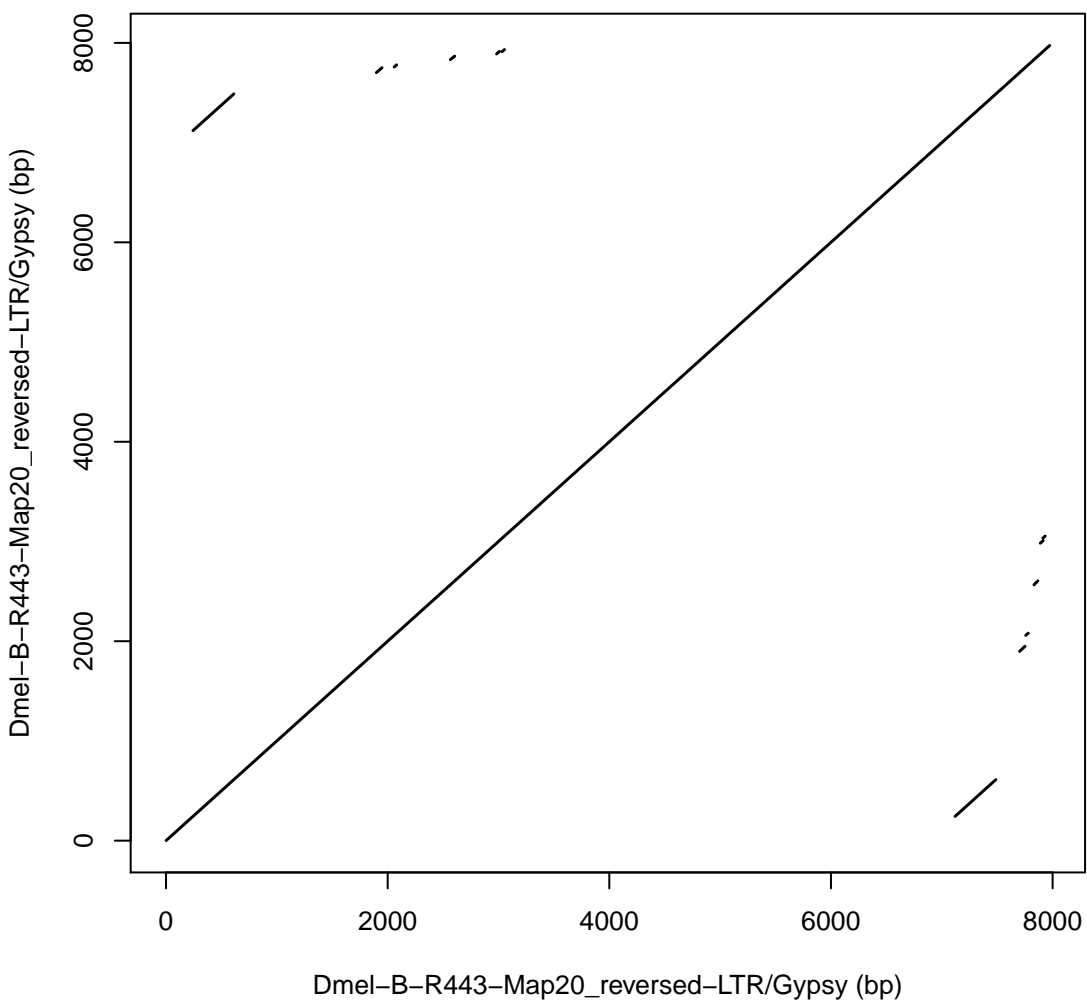
TE: Dmel-B-R443-Map20\_reversed-LTR/Gypsy  
consensus size: 7974bp; fragments: 343; full length: 2 ( $\geq 7176.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

