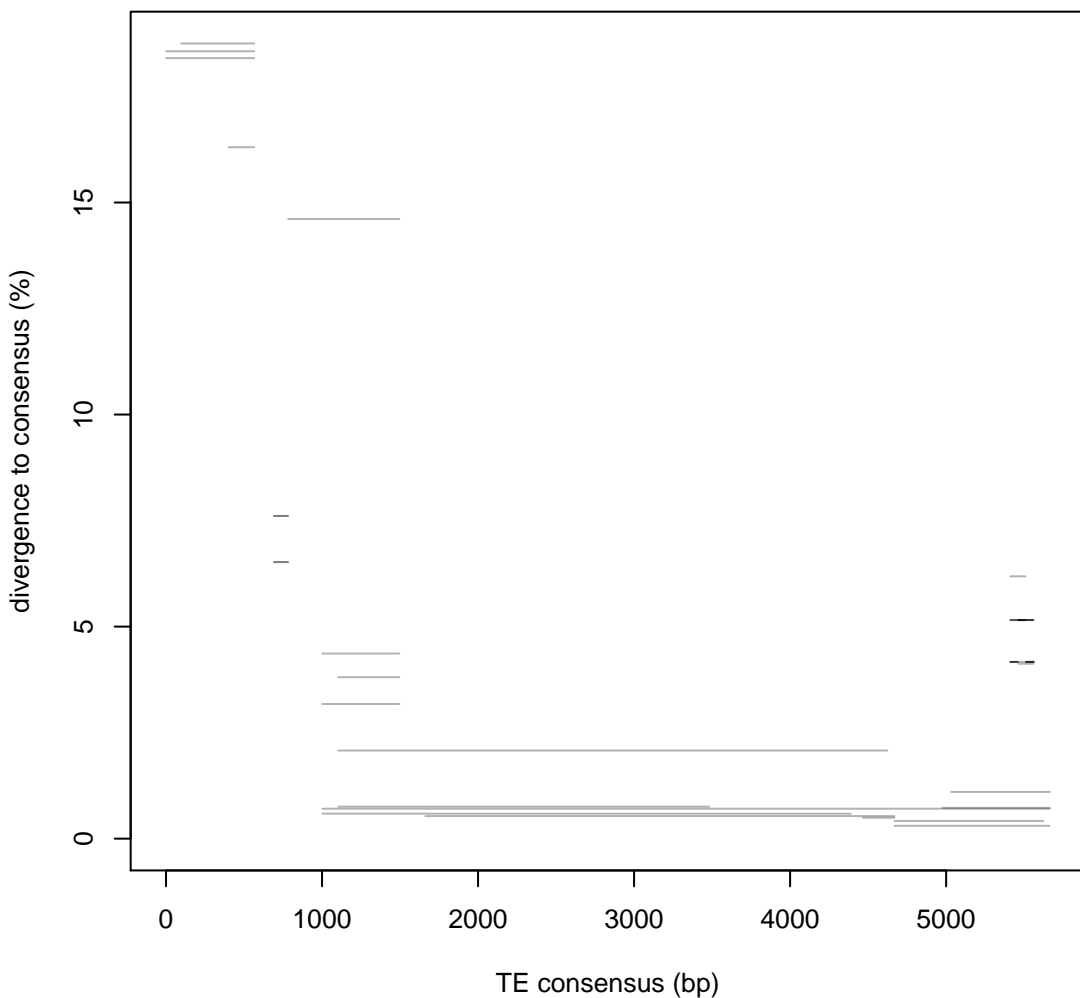
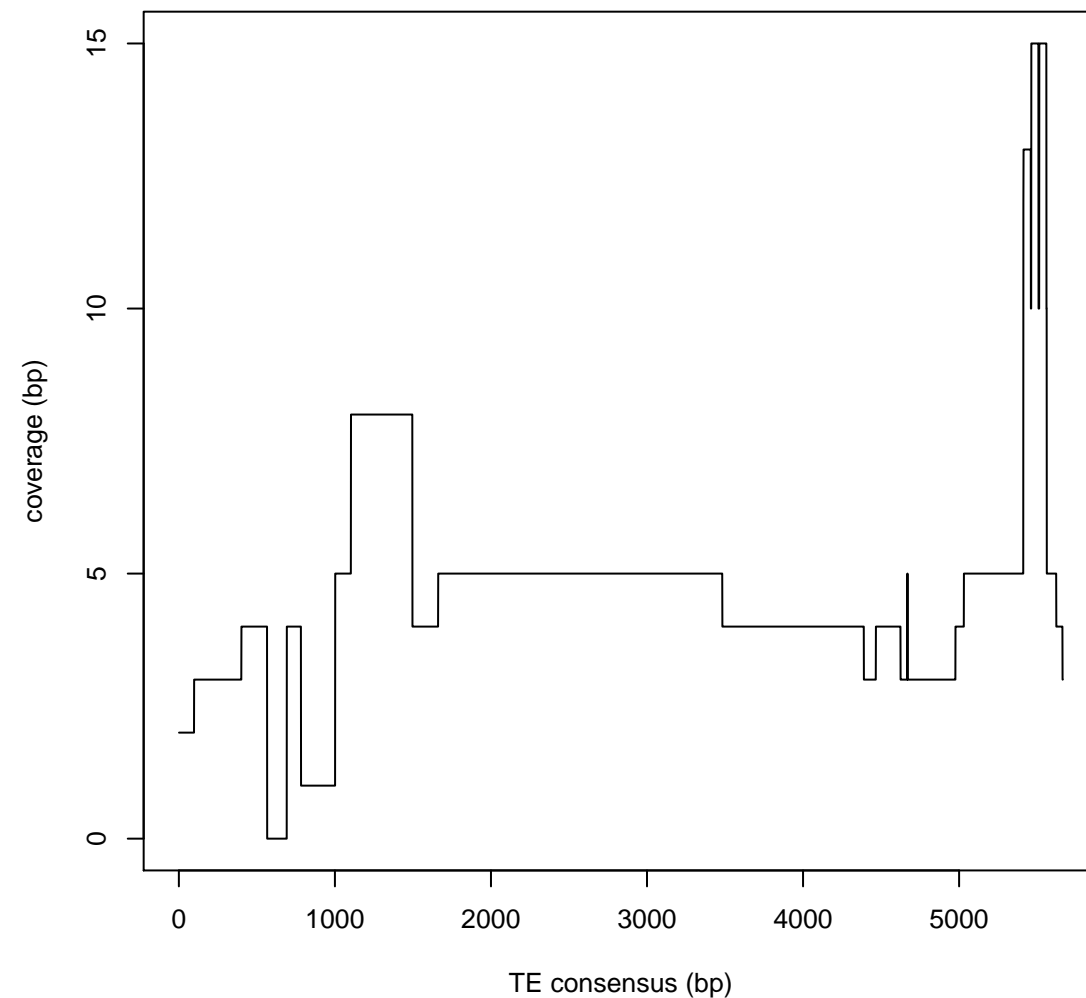


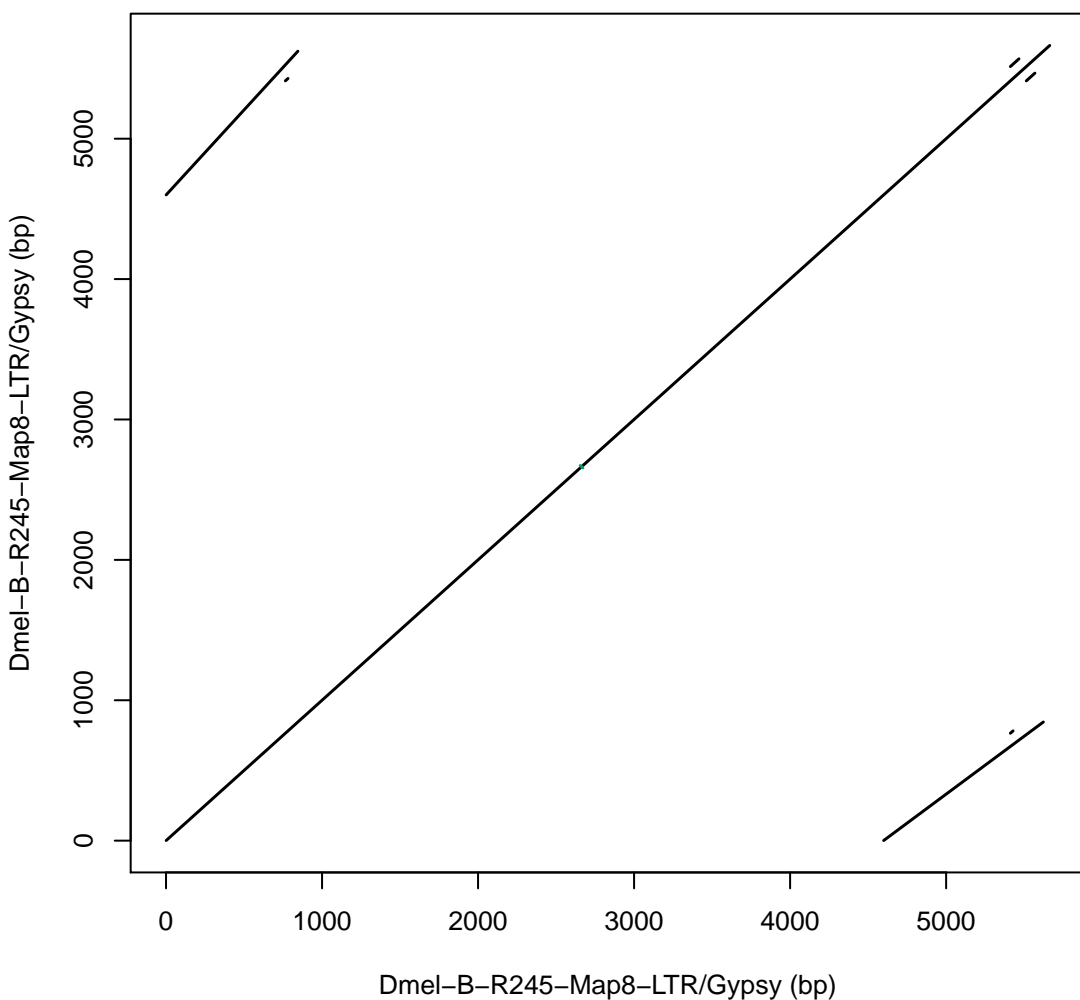
TE: Dmel-B-R245-Map8-LTR/Gypsy  
consensus size: 5664bp; fragments: 40; full length: 0 ( $\geq 5097.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

