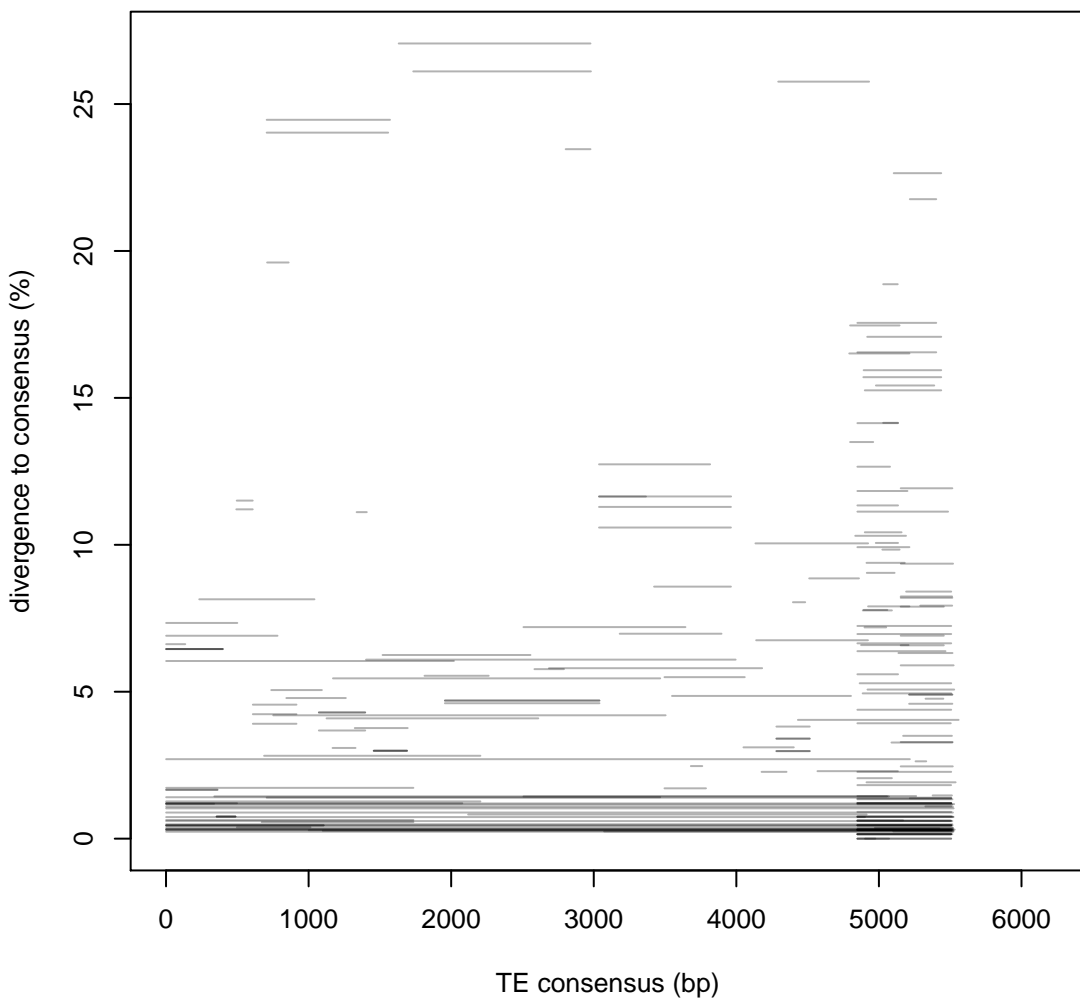
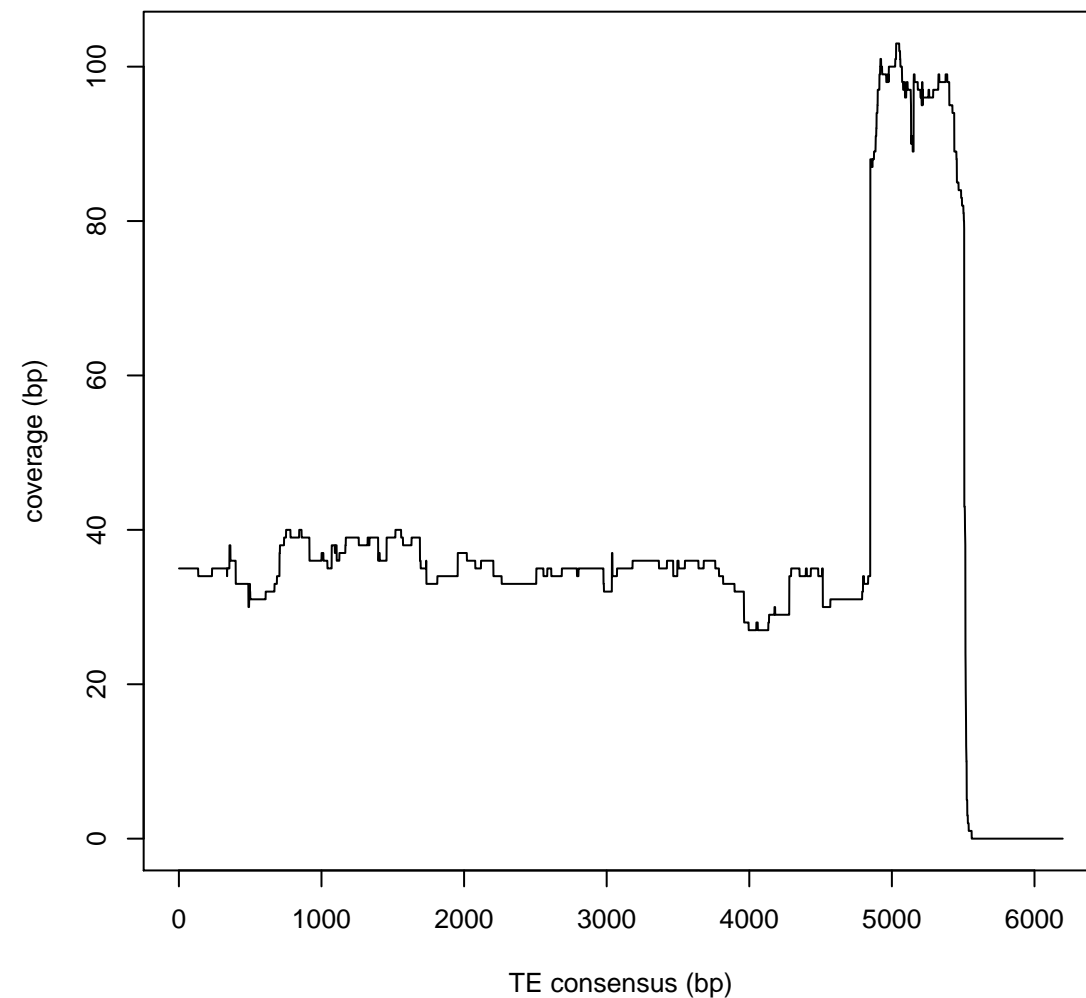


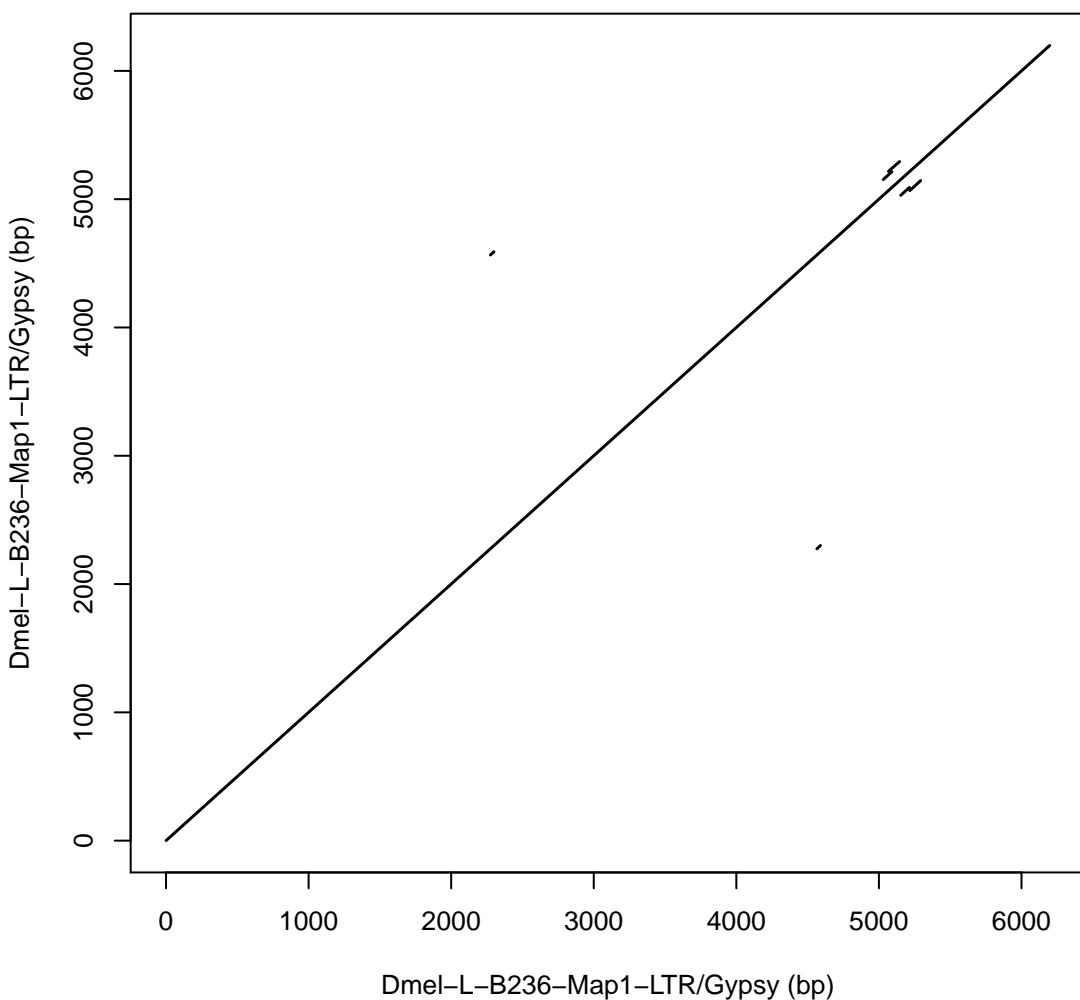
TE: Dmel-L-B236-Map1-LTR/Gypsy  
consensus size: 6198bp; fragments: 215; full length: 0 ( $\geq 5578.2$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

