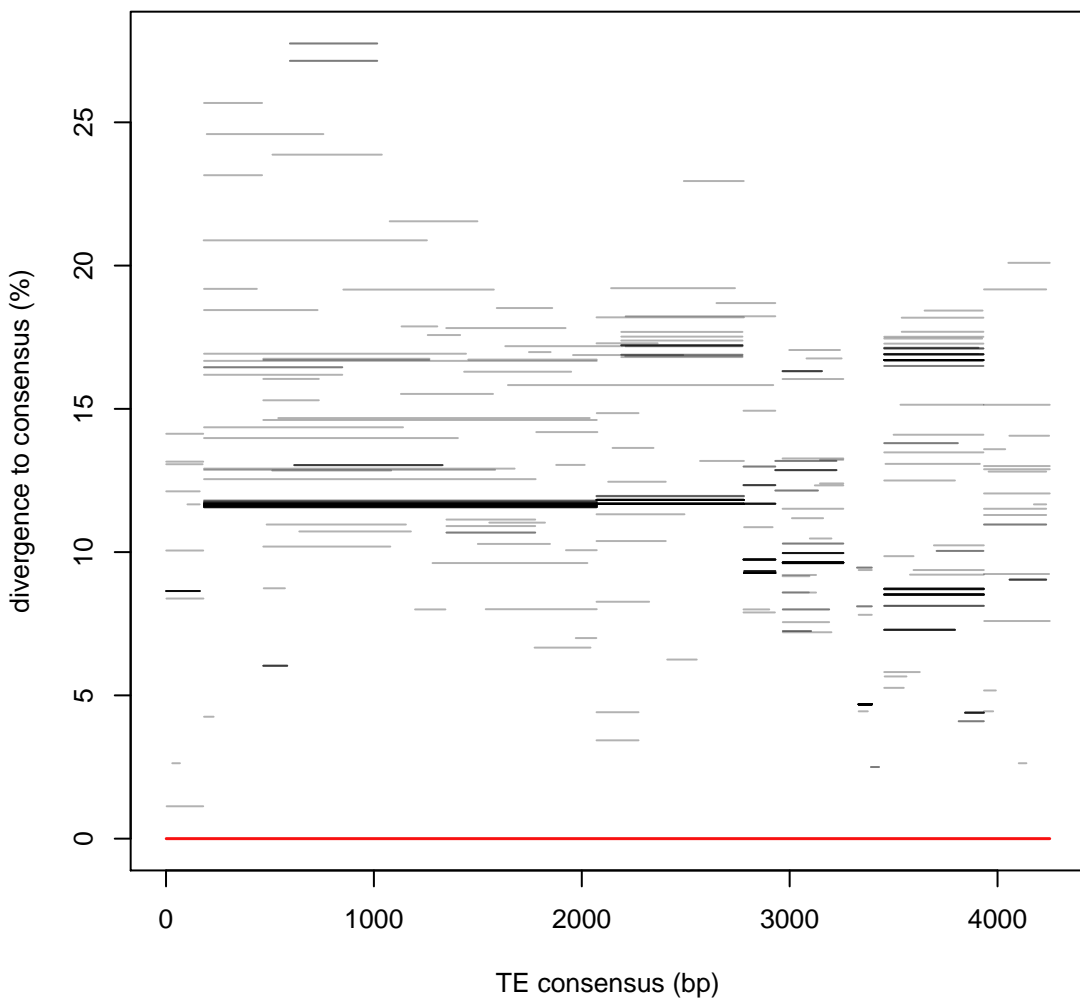
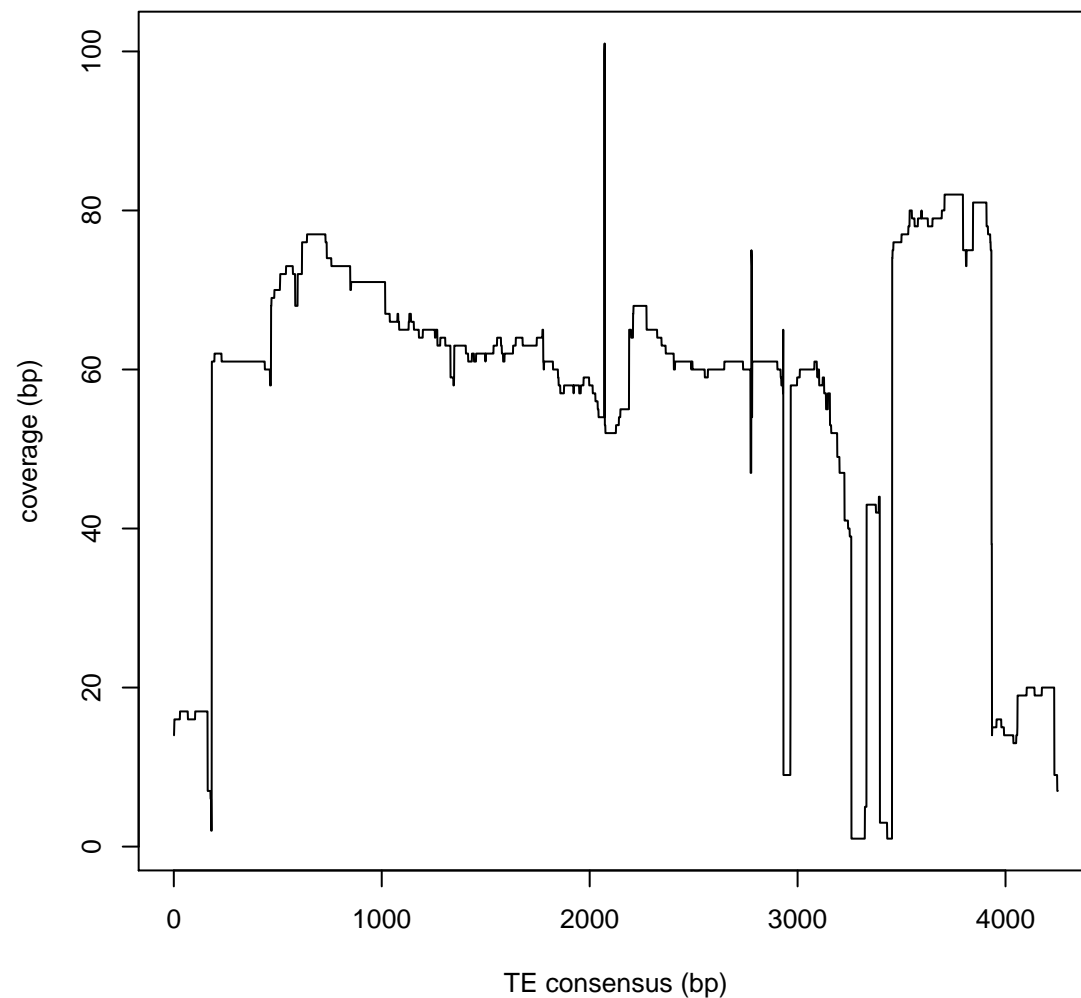


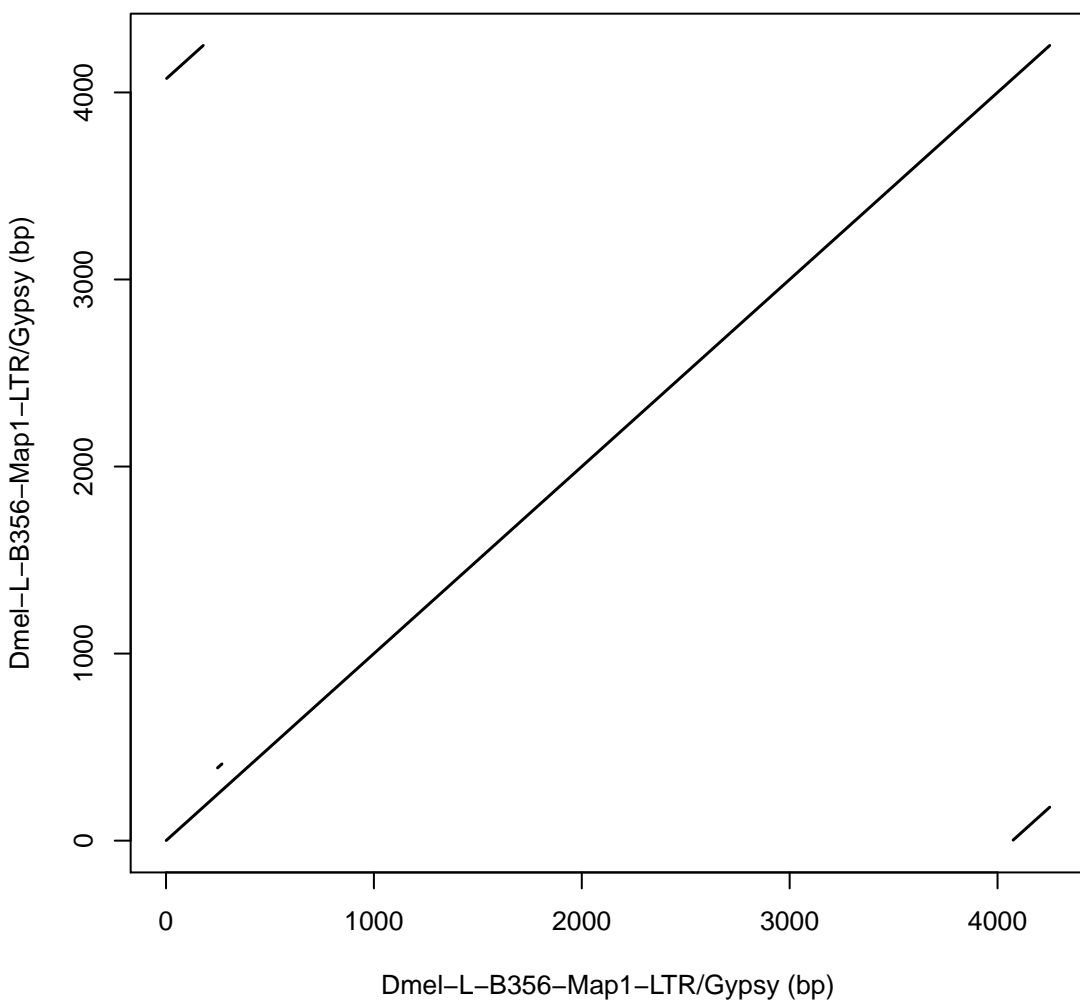
TE: Dmel-L-B356-Map1-LTR/Gypsy  
consensus size: 4251bp; fragments: 480; full length: 1 ( $\geq 3825.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

