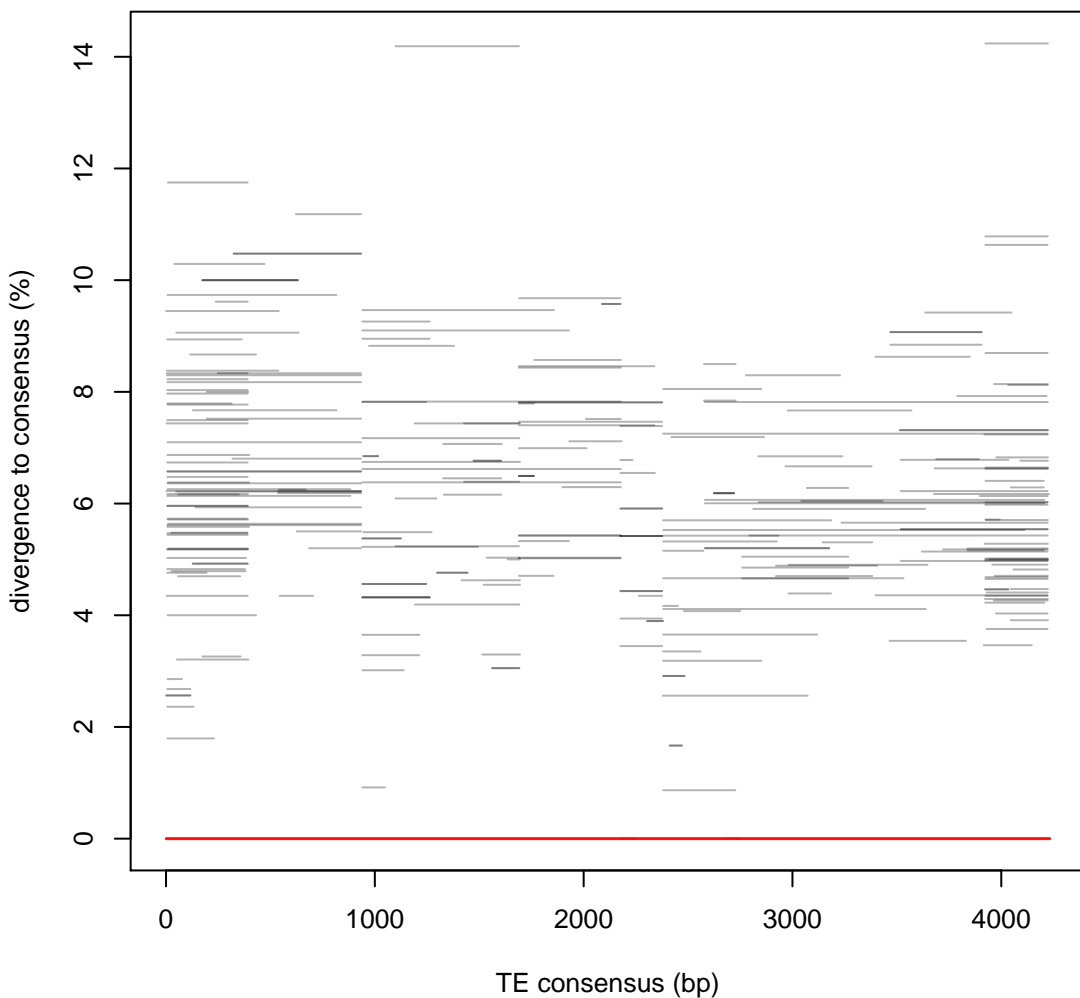
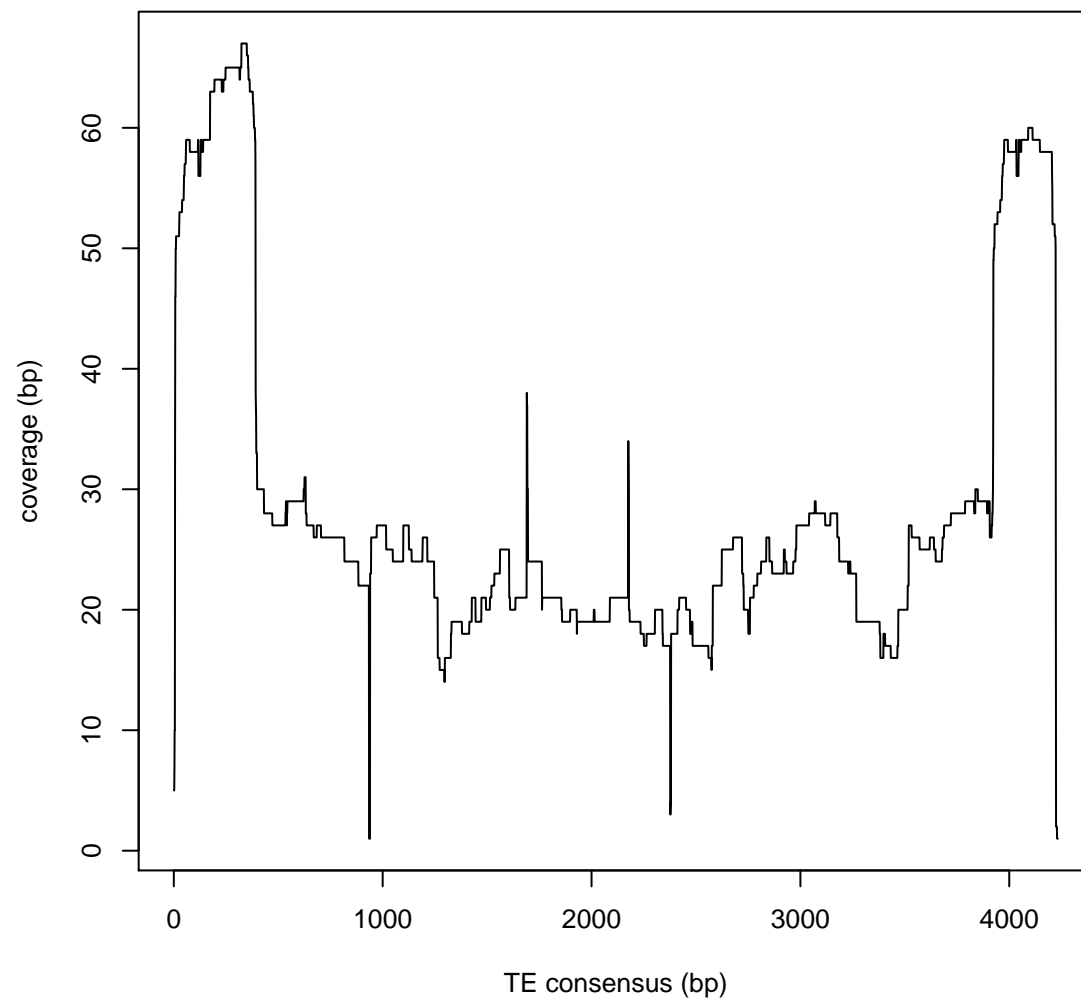


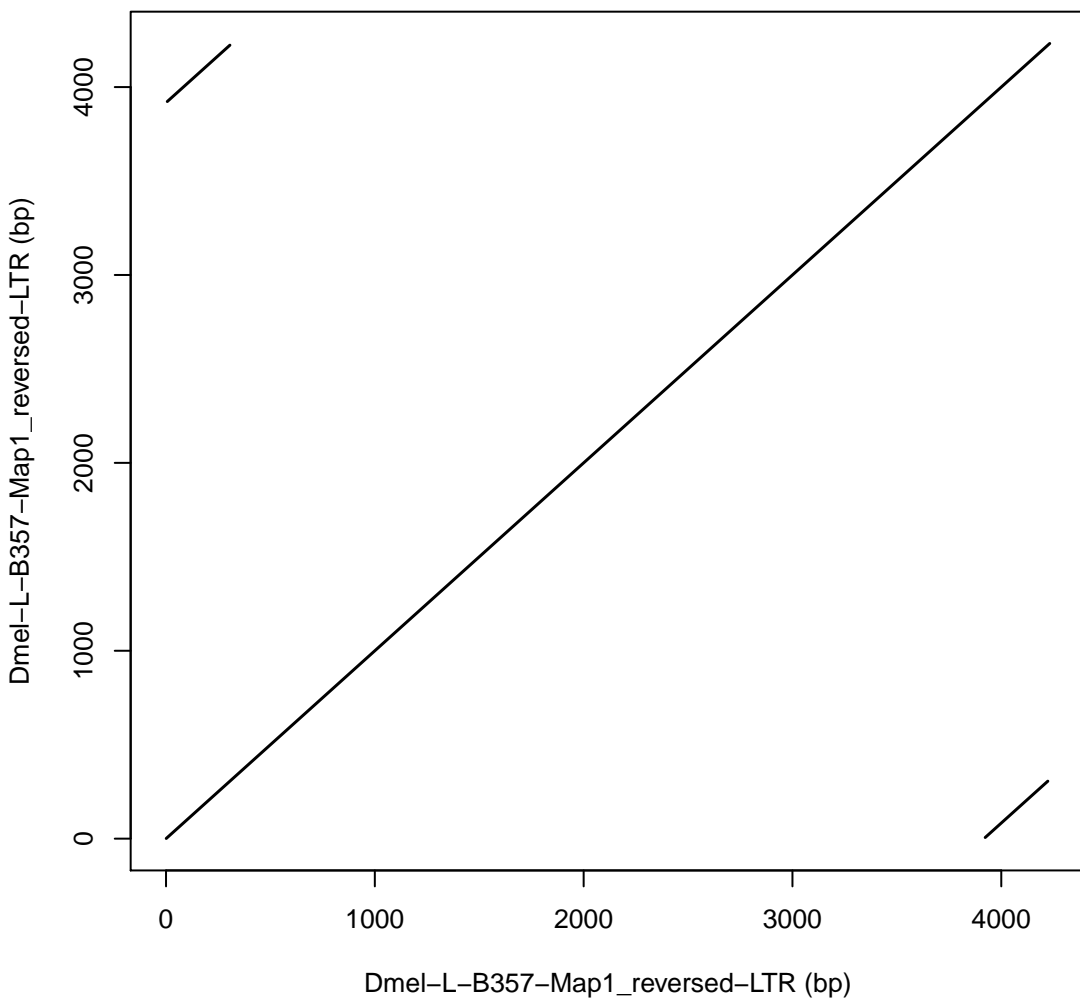
TE: Dmel-L-B357-Map1\_reversed-LTR  
consensus size: 4232bp; fragments: 281; full length: 1 ( $\geq 3808.8$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

