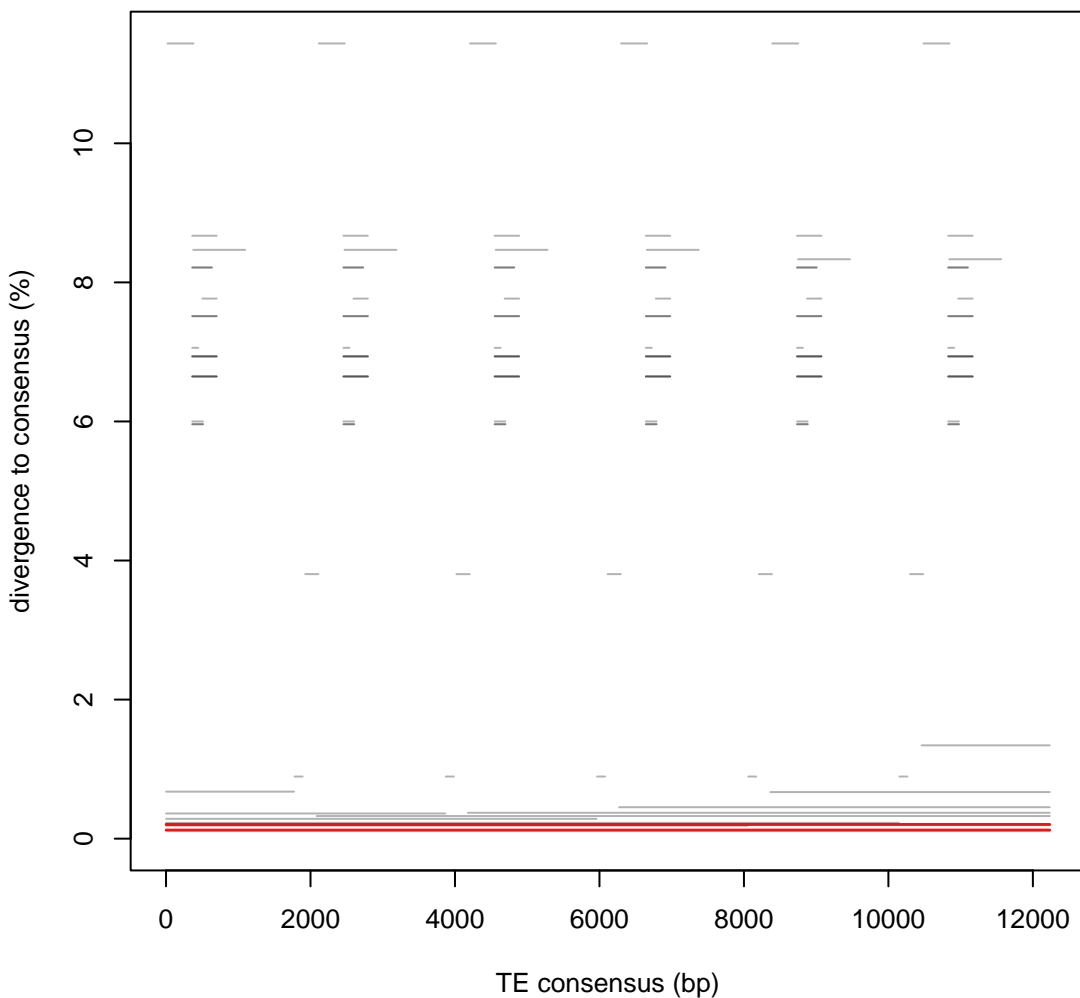
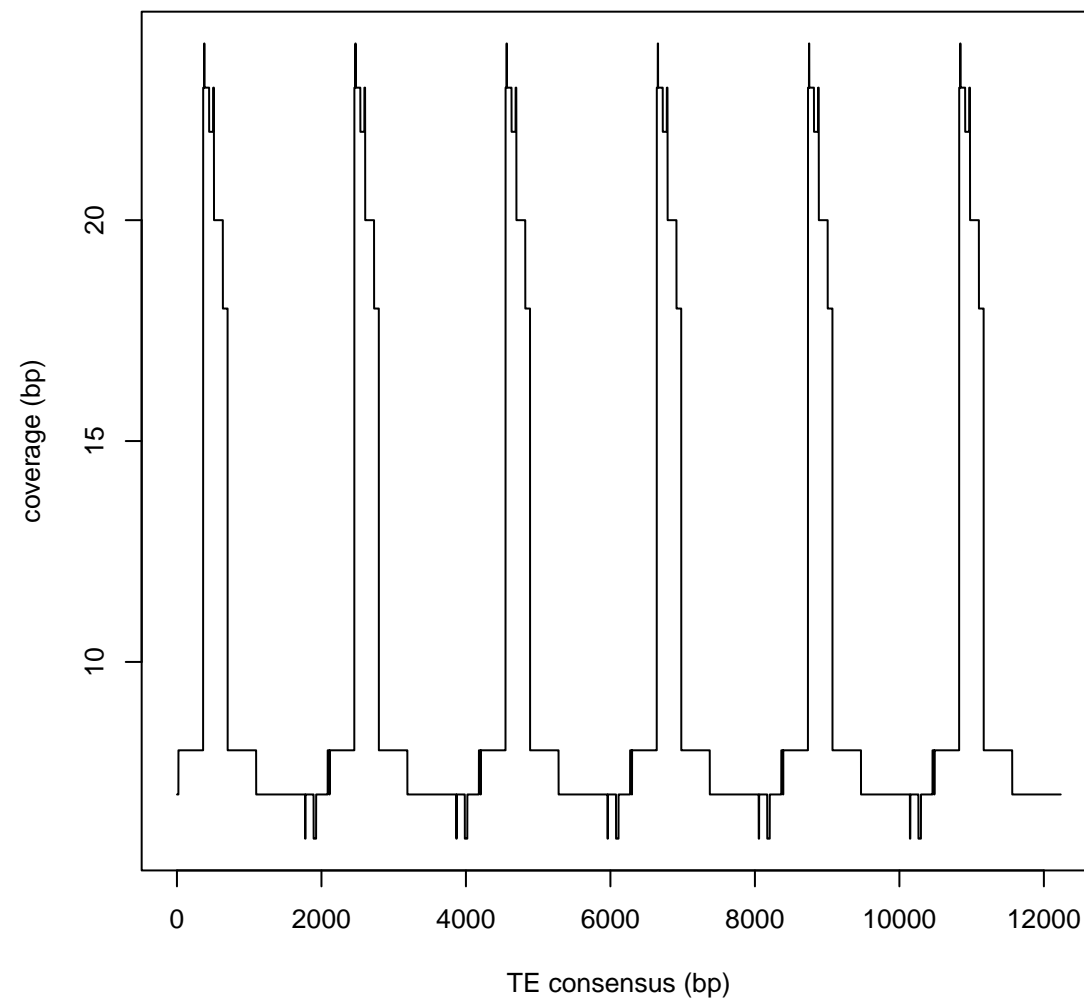


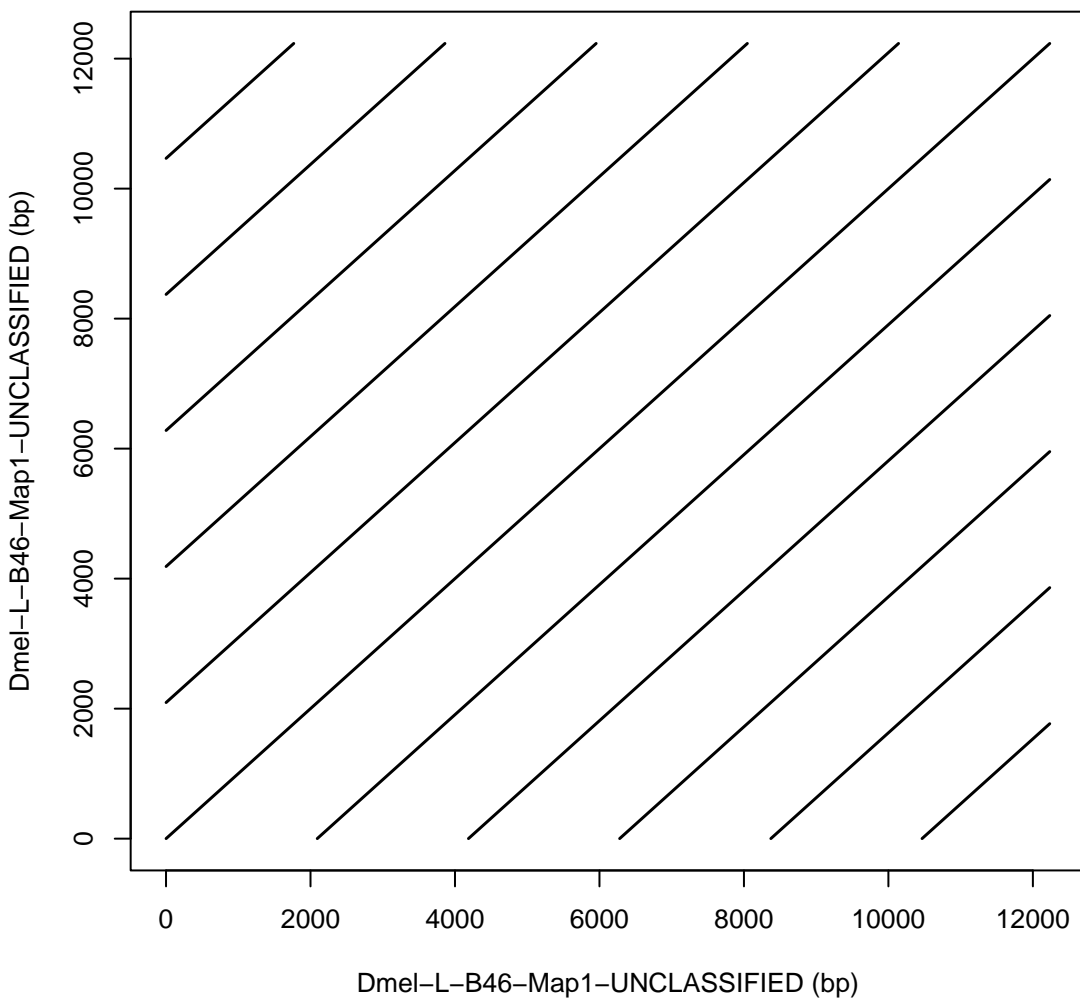
TE: Dmel-L-B46-Map1-UNCLASSIFIED  
consensus size: 12233bp; fragments: 130; full length: 2 ( $\geq 11009.7$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

