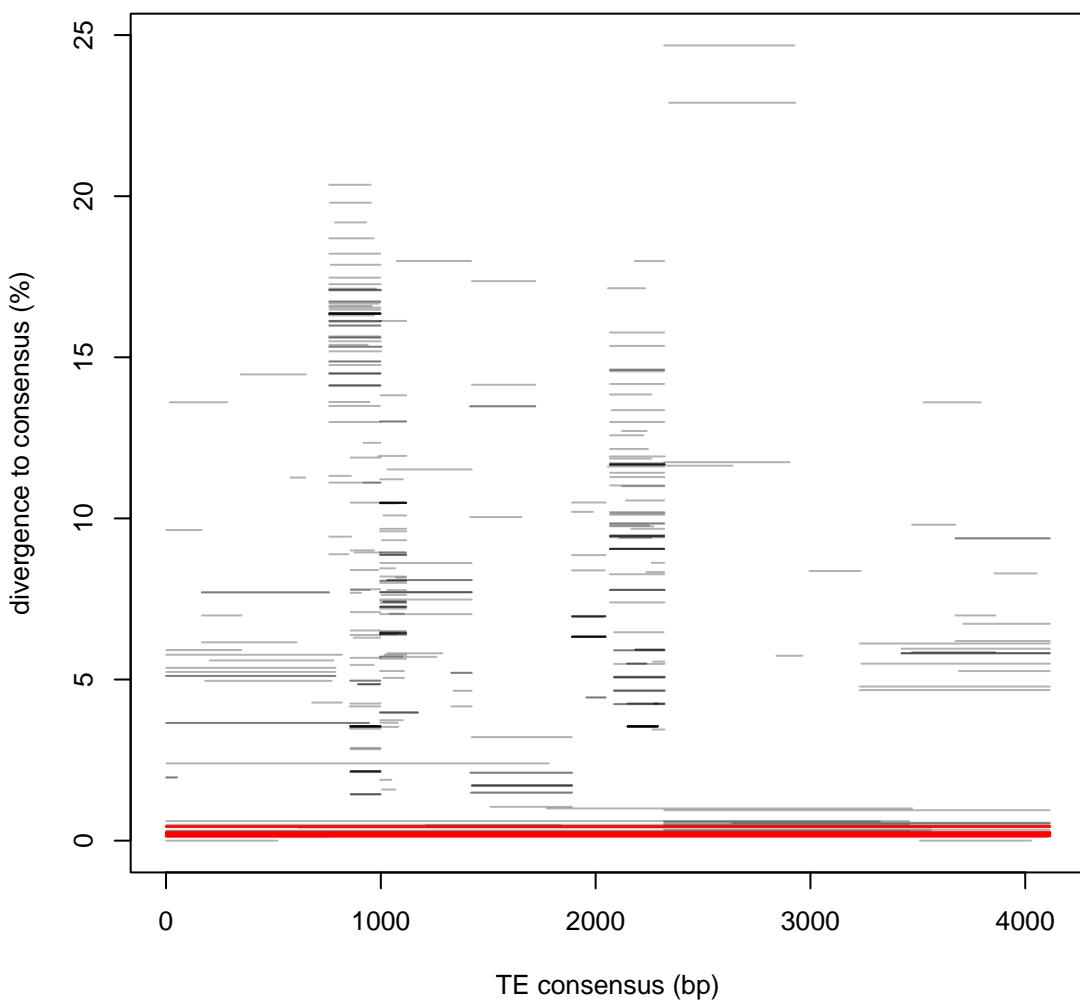
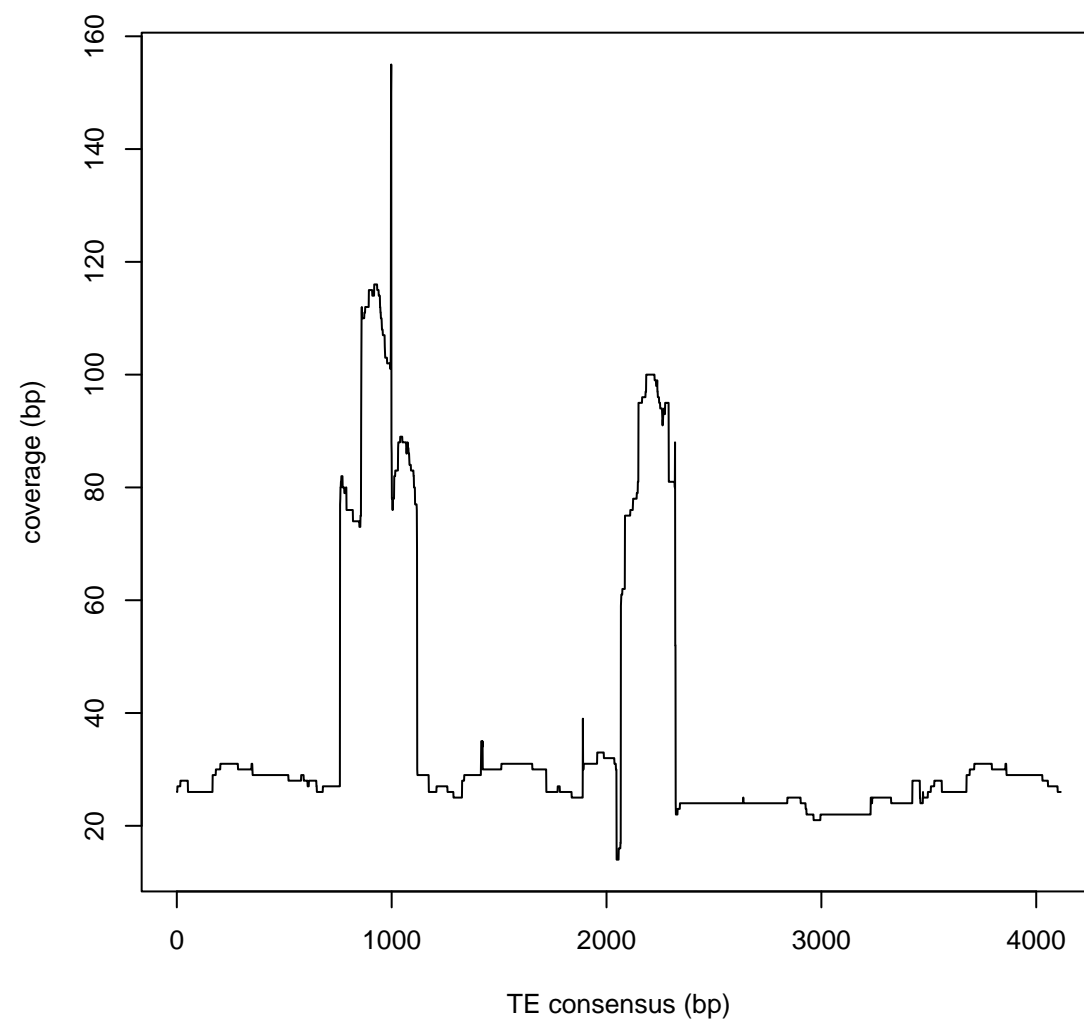


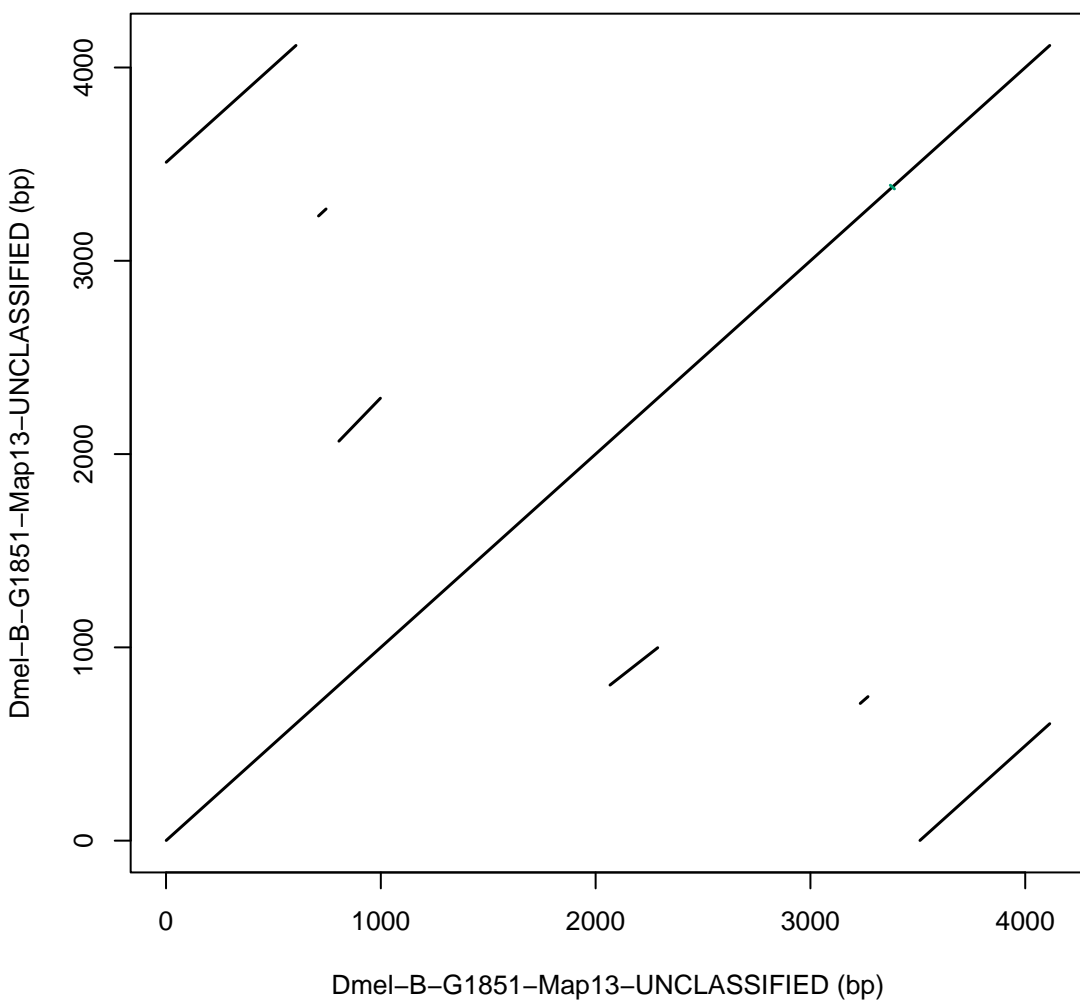
TE: Dmel-B-G1851-Map13-UNCLASSIFIED  
consensus size: 4114bp; fragments: 385; full length: 7 ( $\geq 3702.6$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

