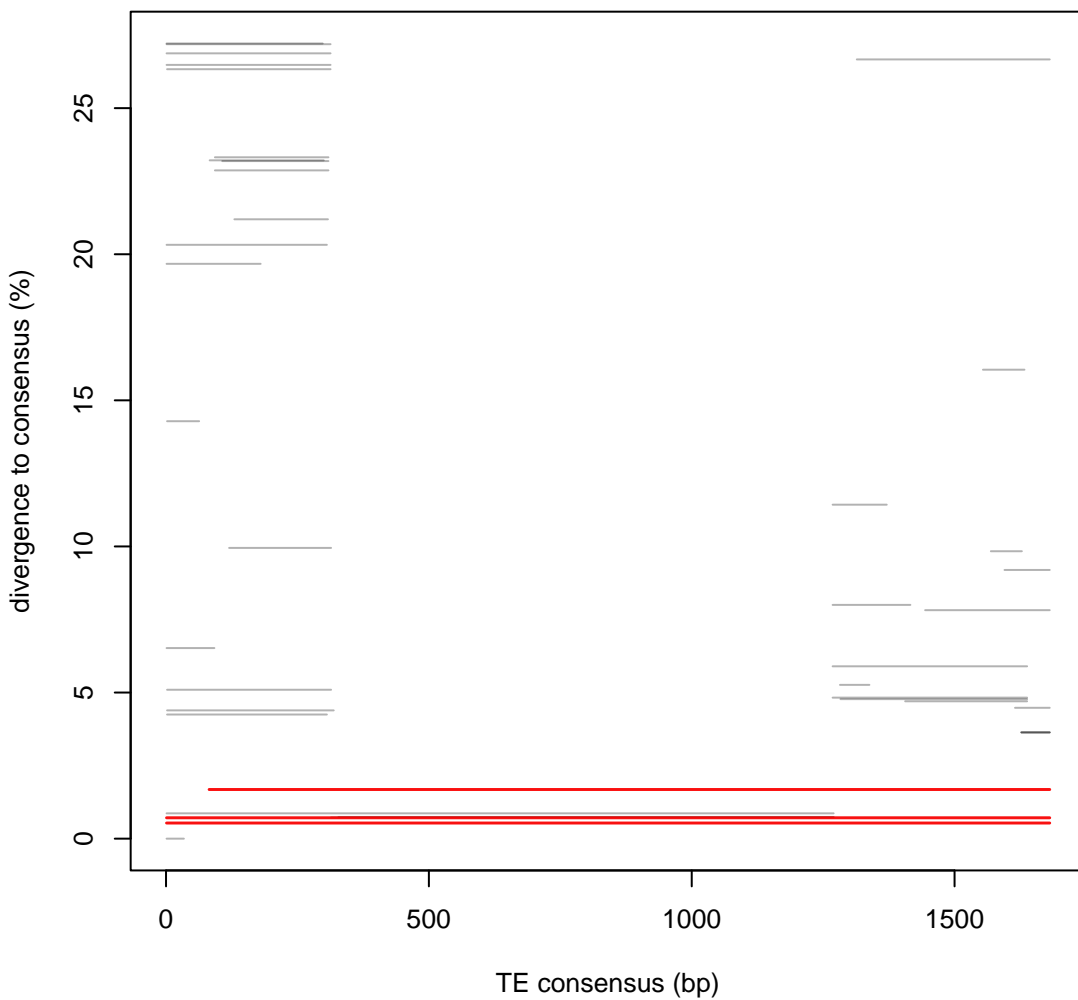
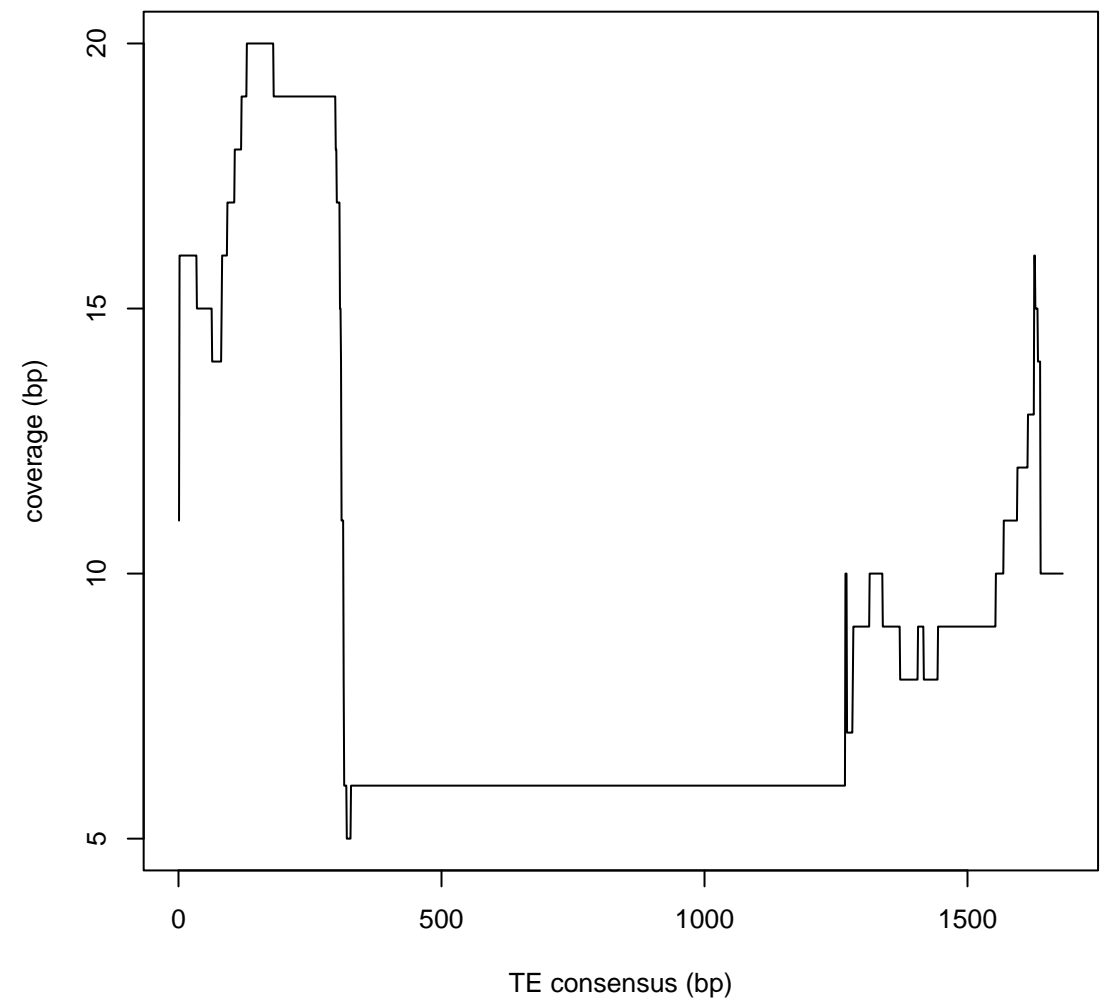


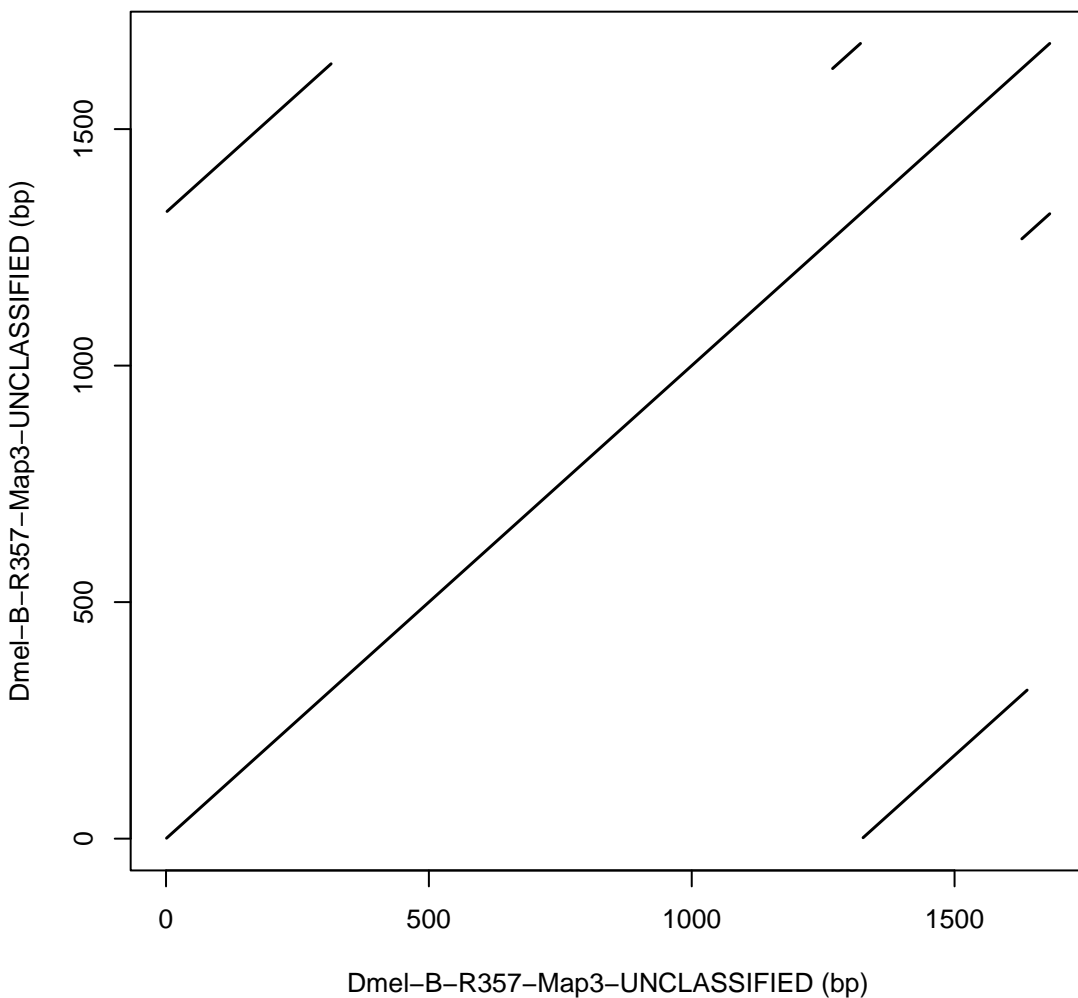
TE: Dmel-B-R357-Map3-UNCLASSIFIED  
consensus size: 1681bp; fragments: 41; full length: 3 ( $\geq 1512.9$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

