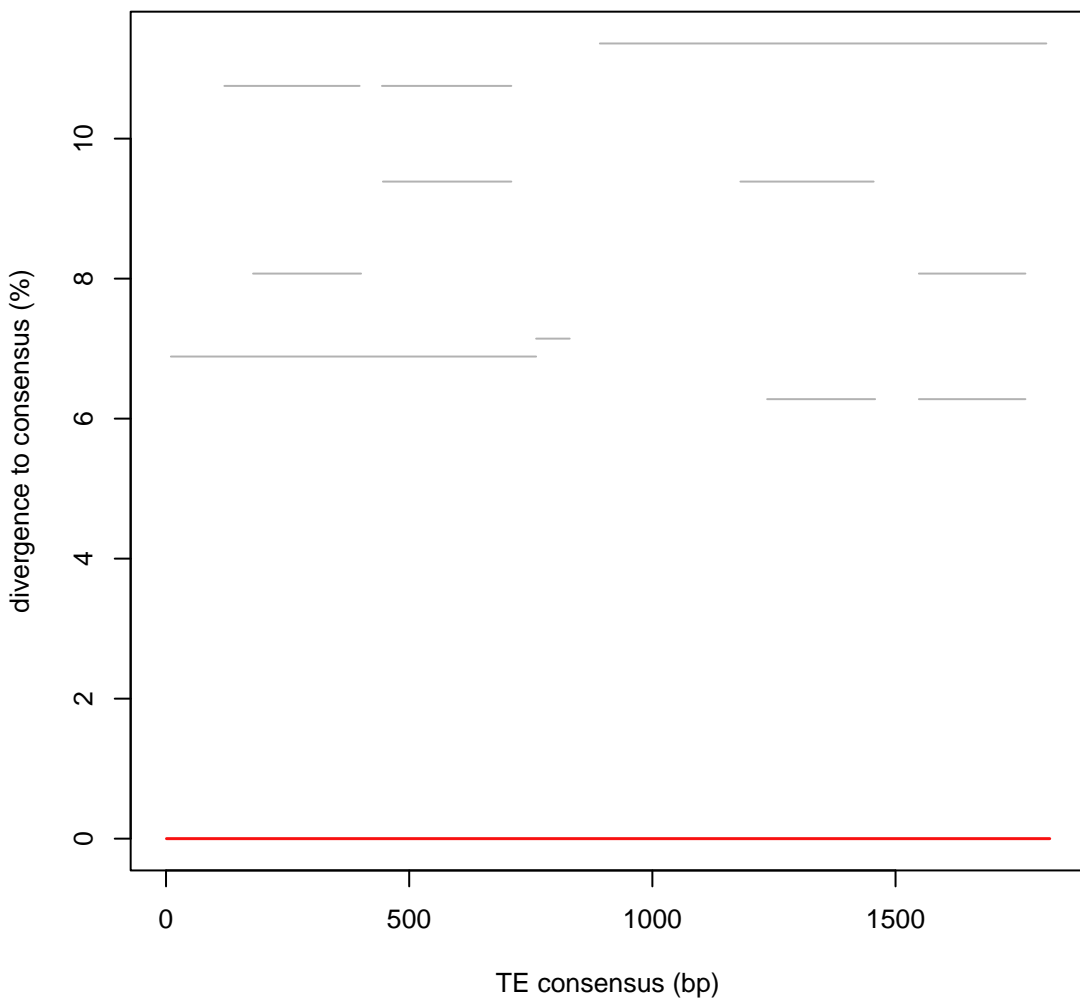
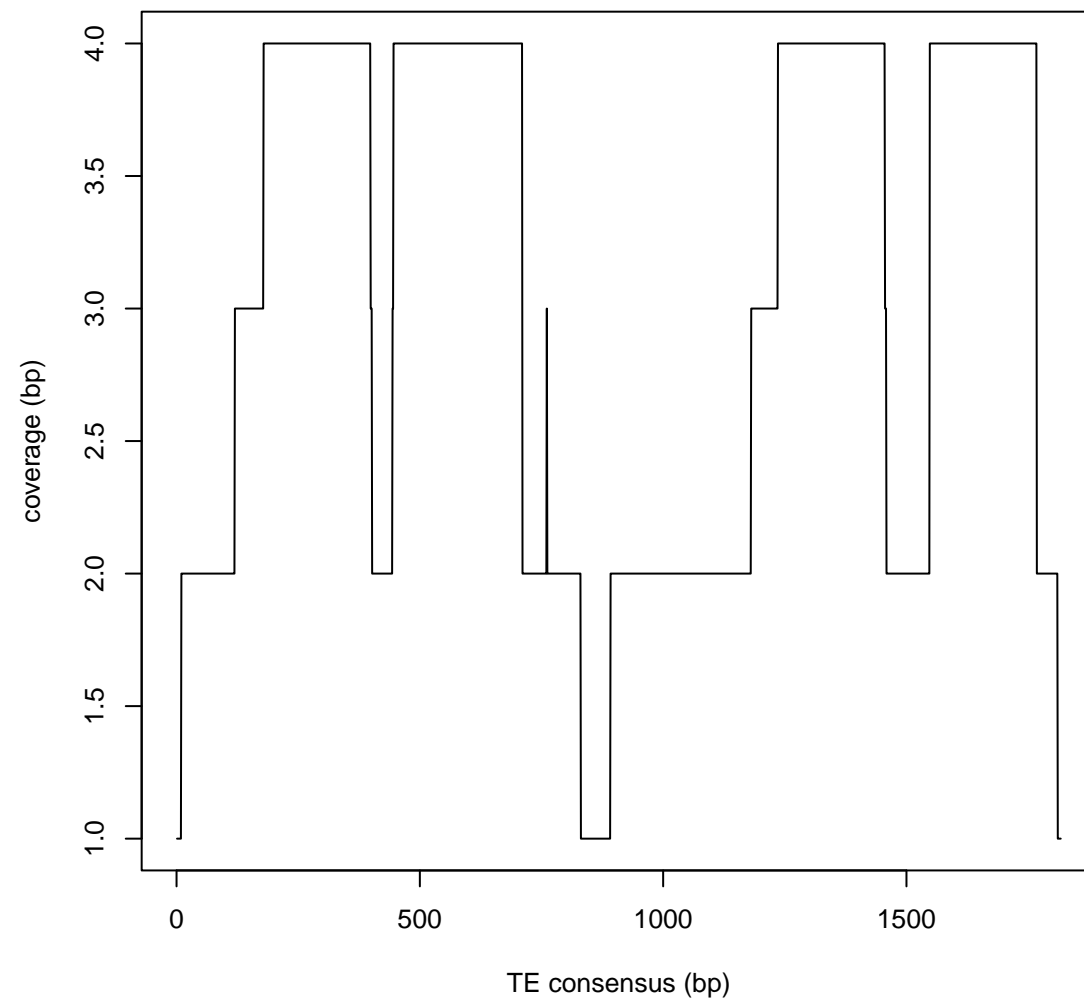


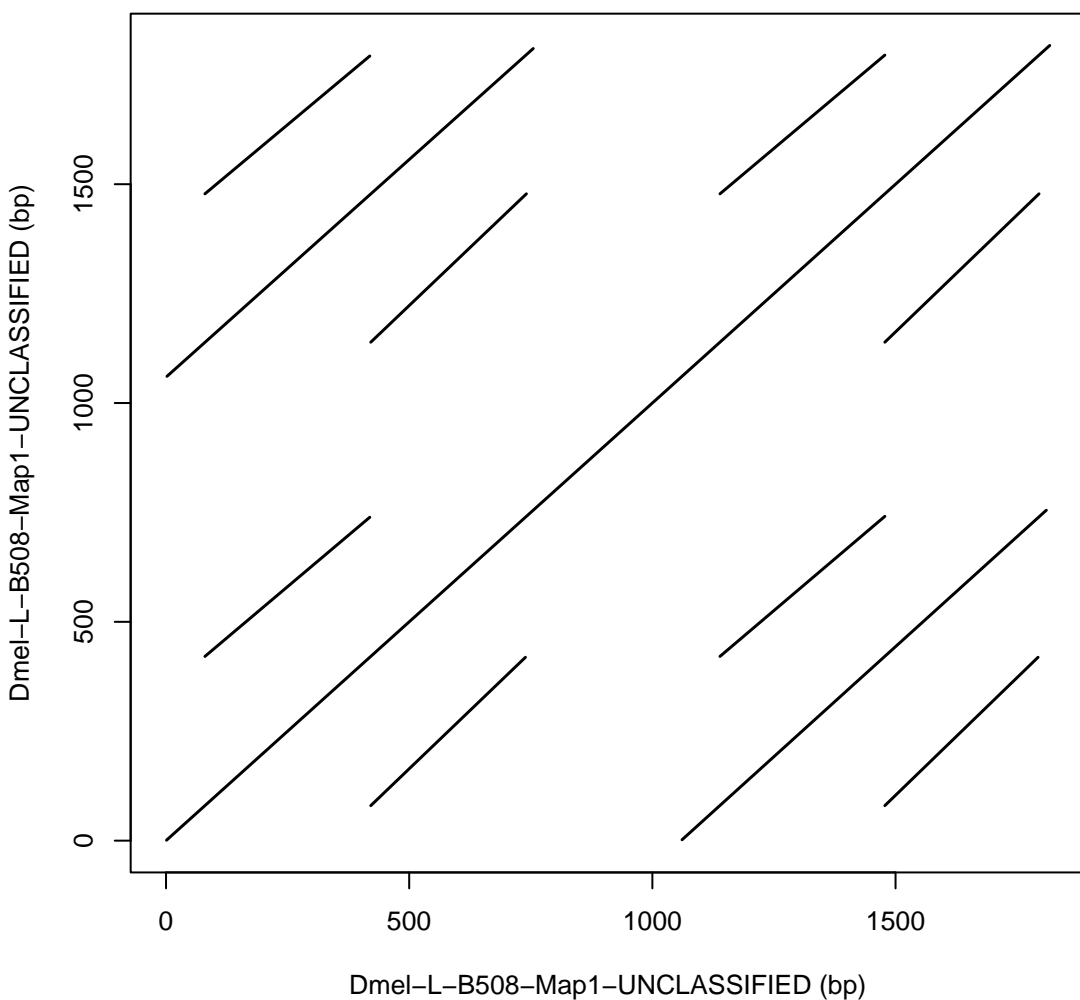
TE: Dmel-L-B508-Map1-UNCLASSIFIED  
consensus size: 1817bp; fragments: 12; full length: 1 ( $\geq 1635.3$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

