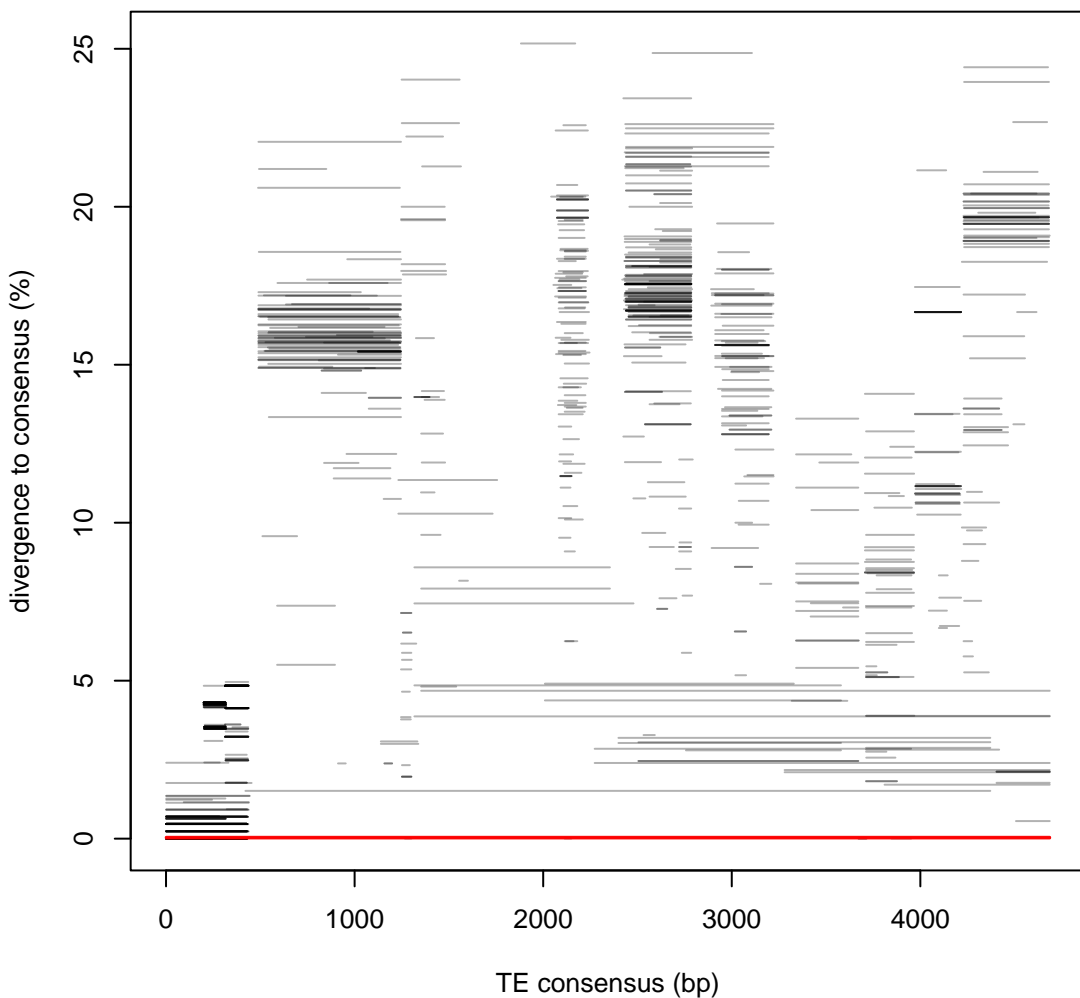
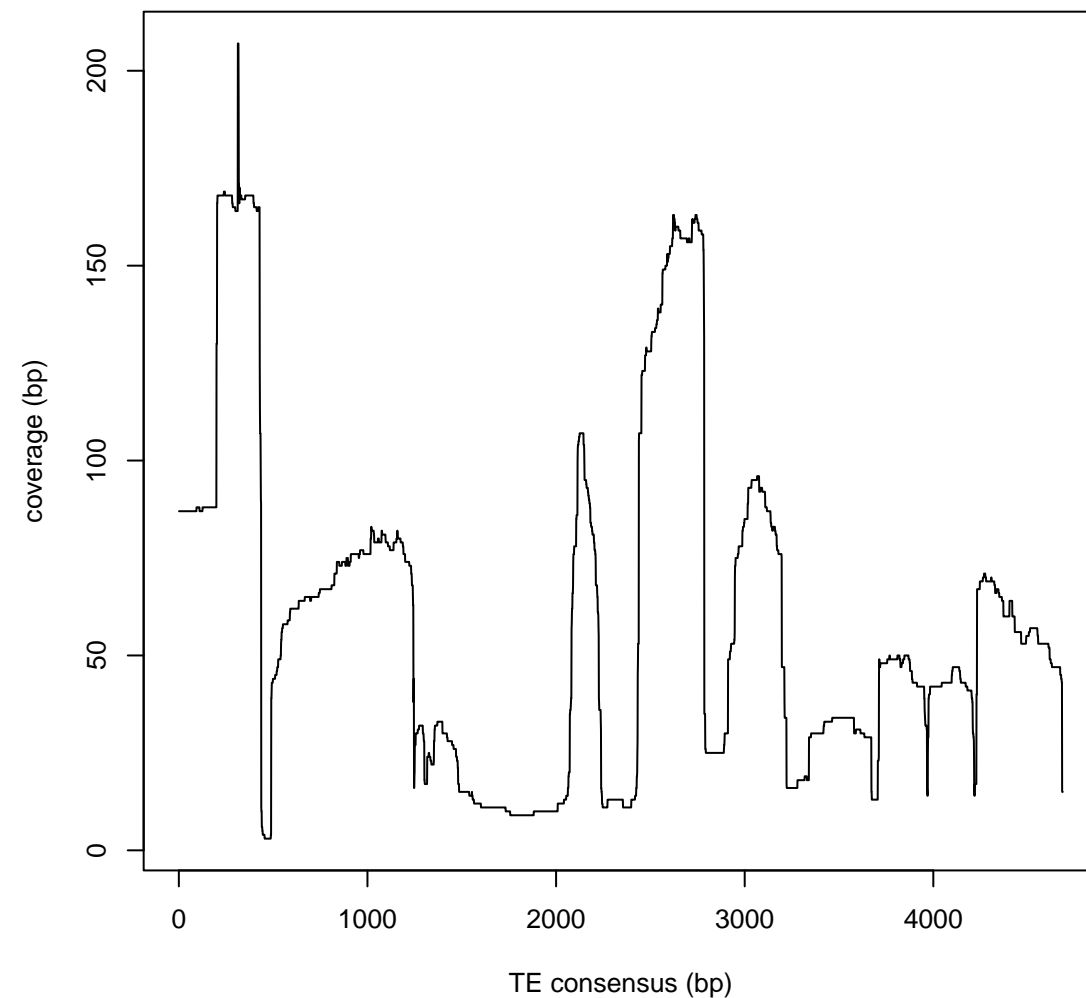


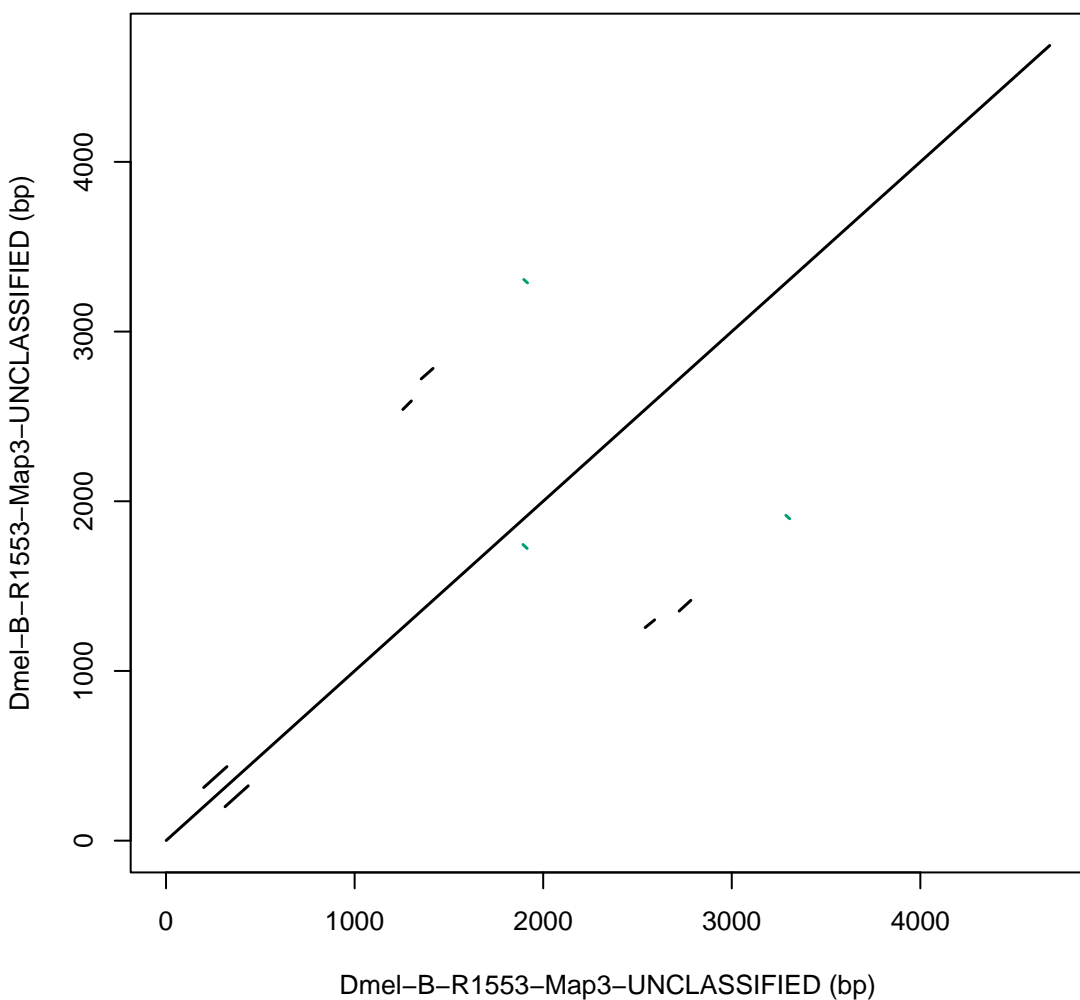
TE: Dmel-B-R1553-Map3-UNCLASSIFIED  
consensus size: 4686bp; fragments: 913; full length: 2 ( $\geq 4217.4$ bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

