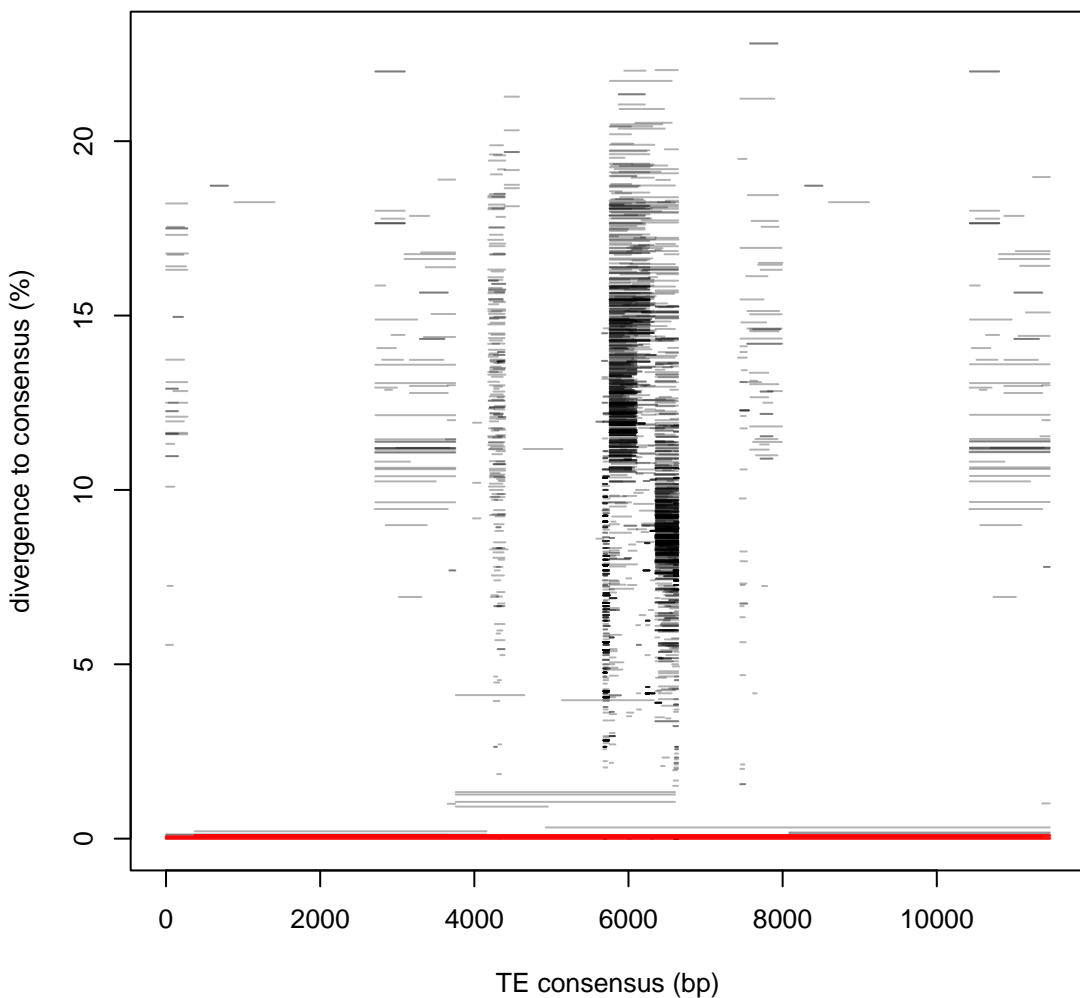
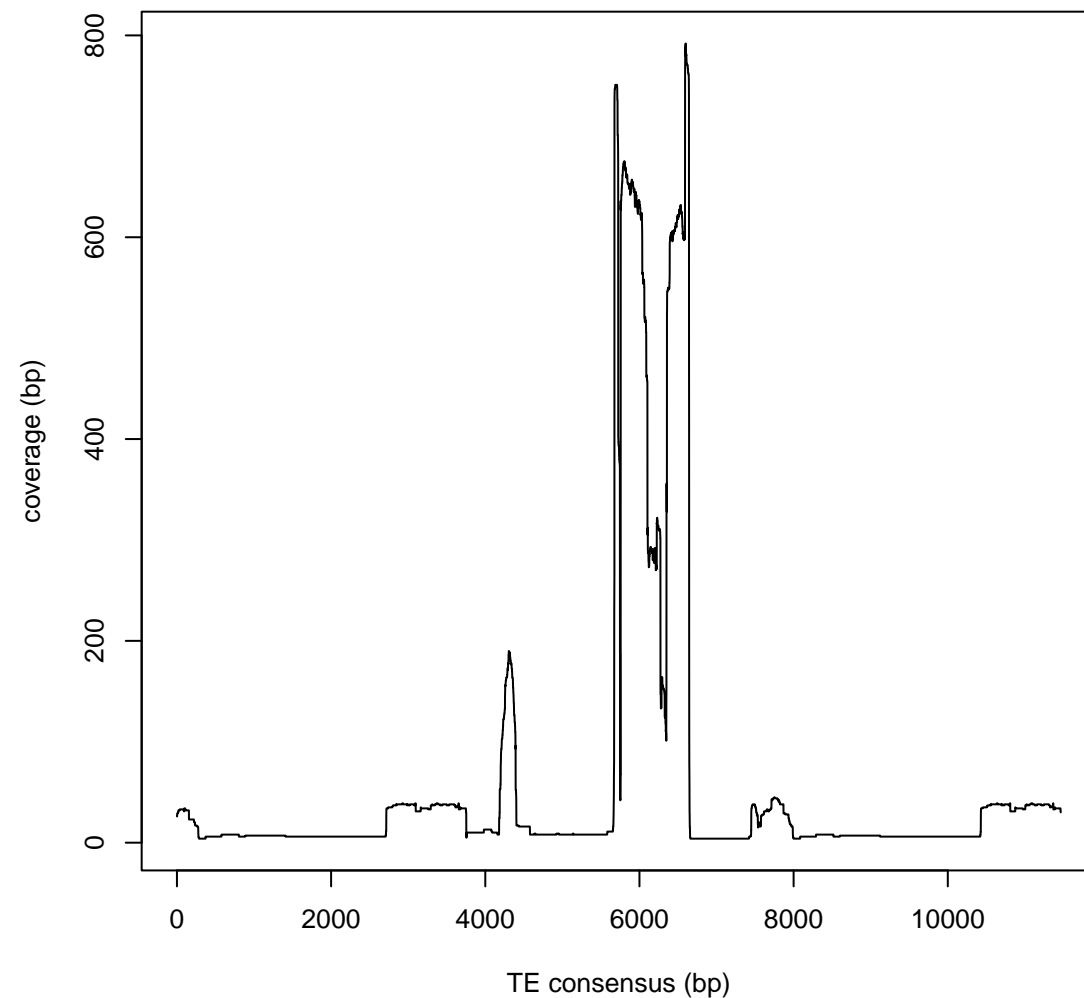


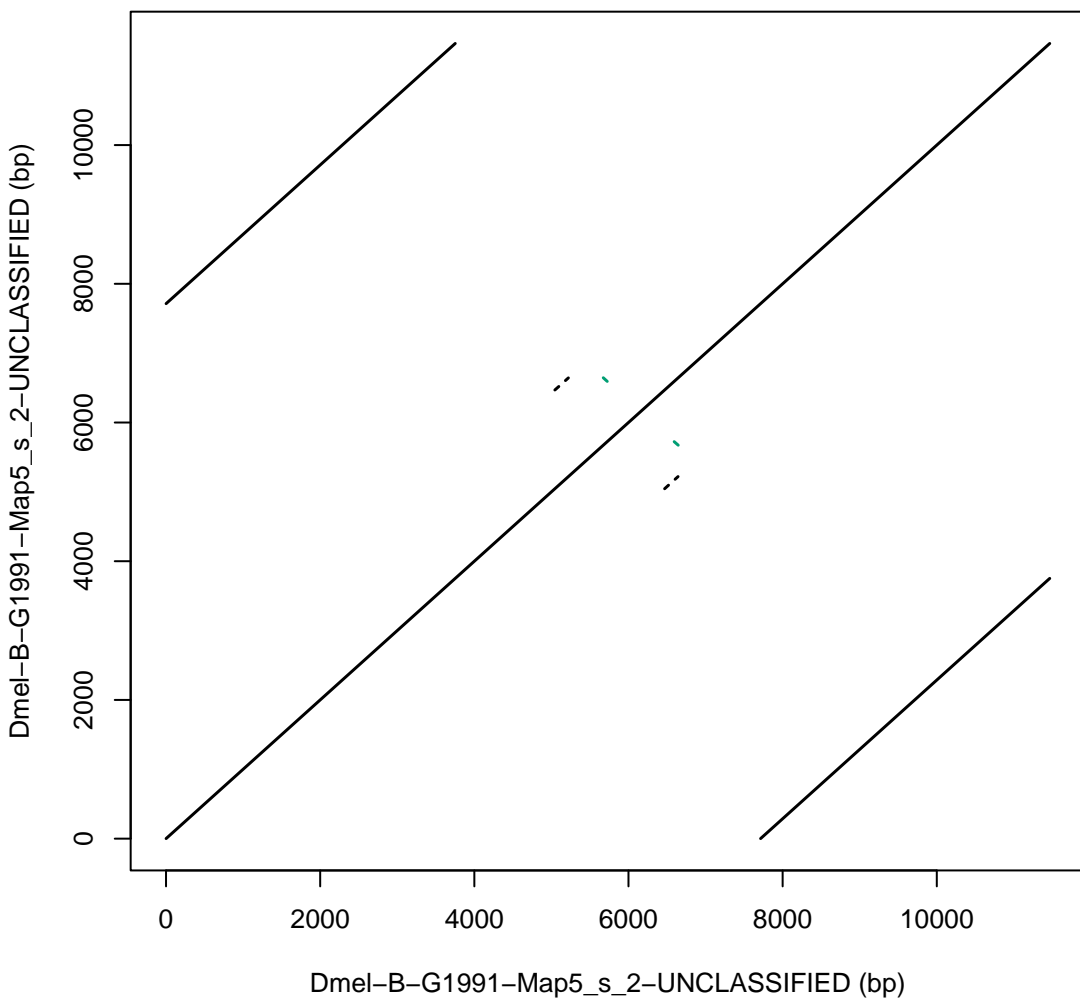
TE: Dmel-B-G1991-Map5\_s\_2-UNCLASSIFIED  
consensus size: 11465bp; fragments: 3016; full length: 3 (>=10318.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

