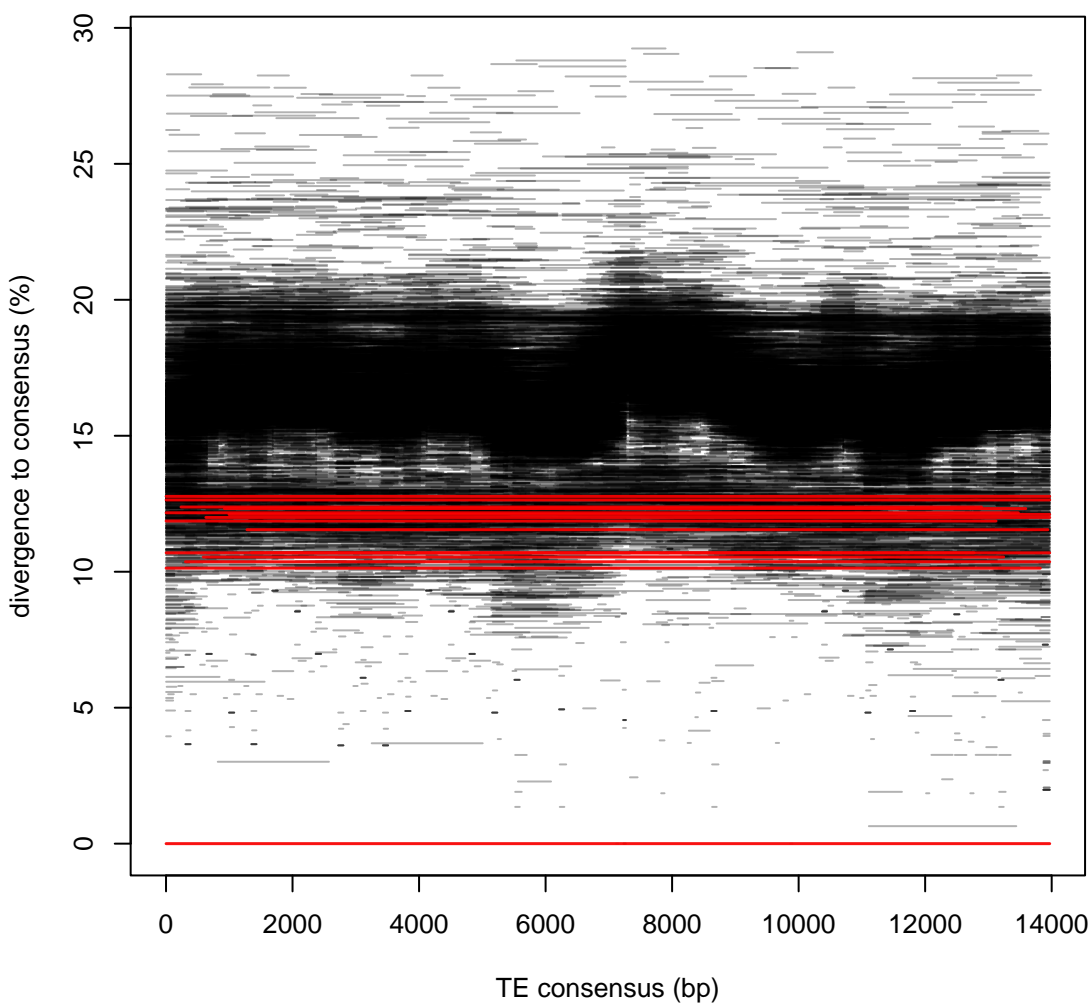
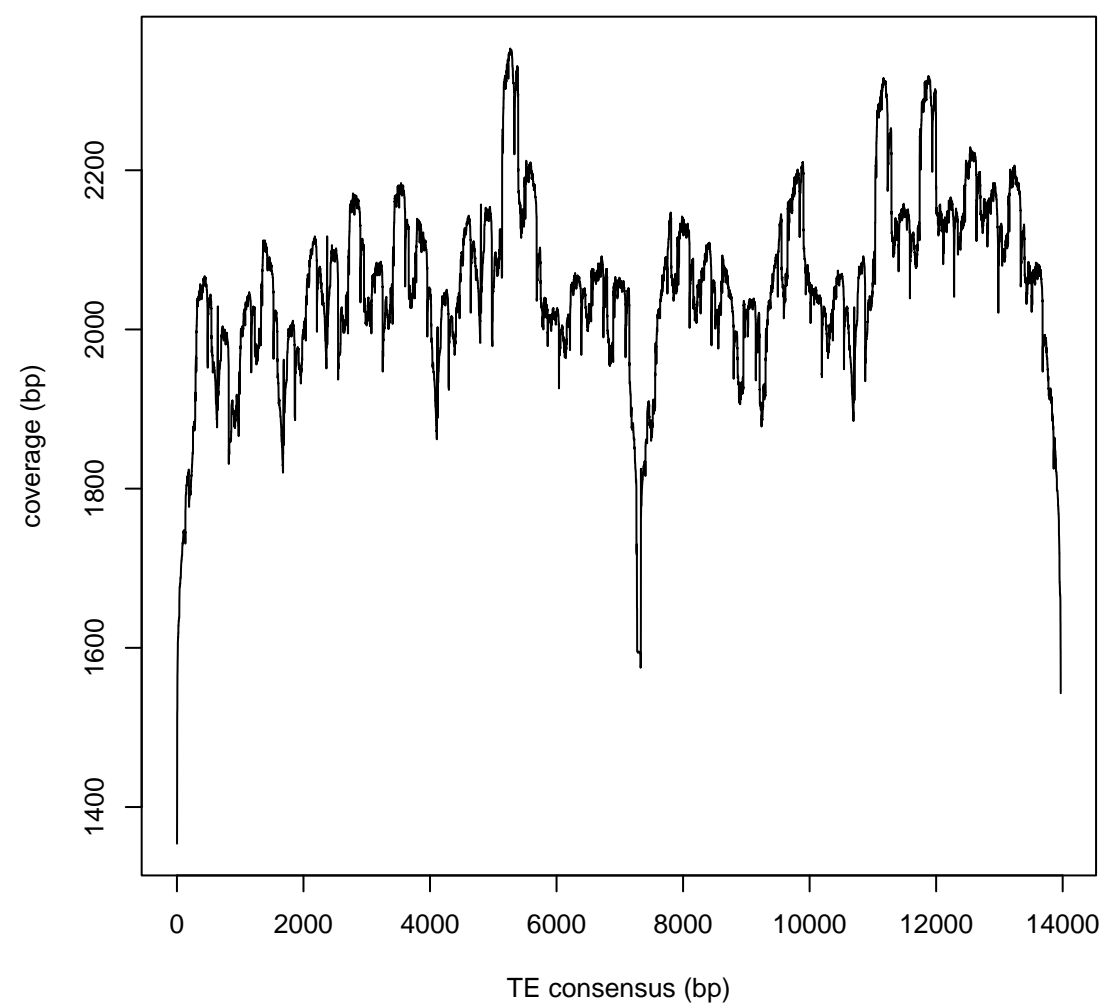


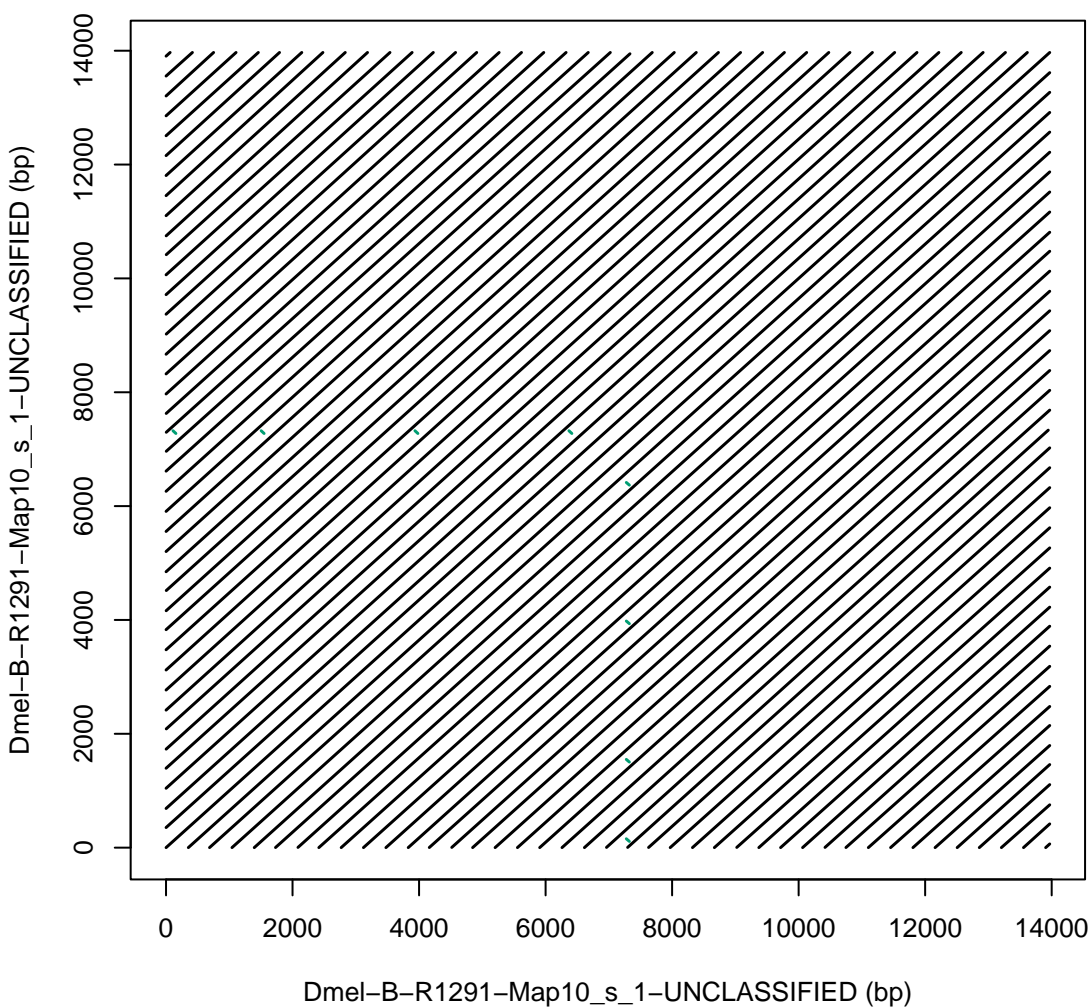
TE: Dmel-B-R1291-Map10\_s\_1-UNCLASSIFIED  
consensus size: 13969bp; fragments: 26045; full length: 15 (>=12572.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

