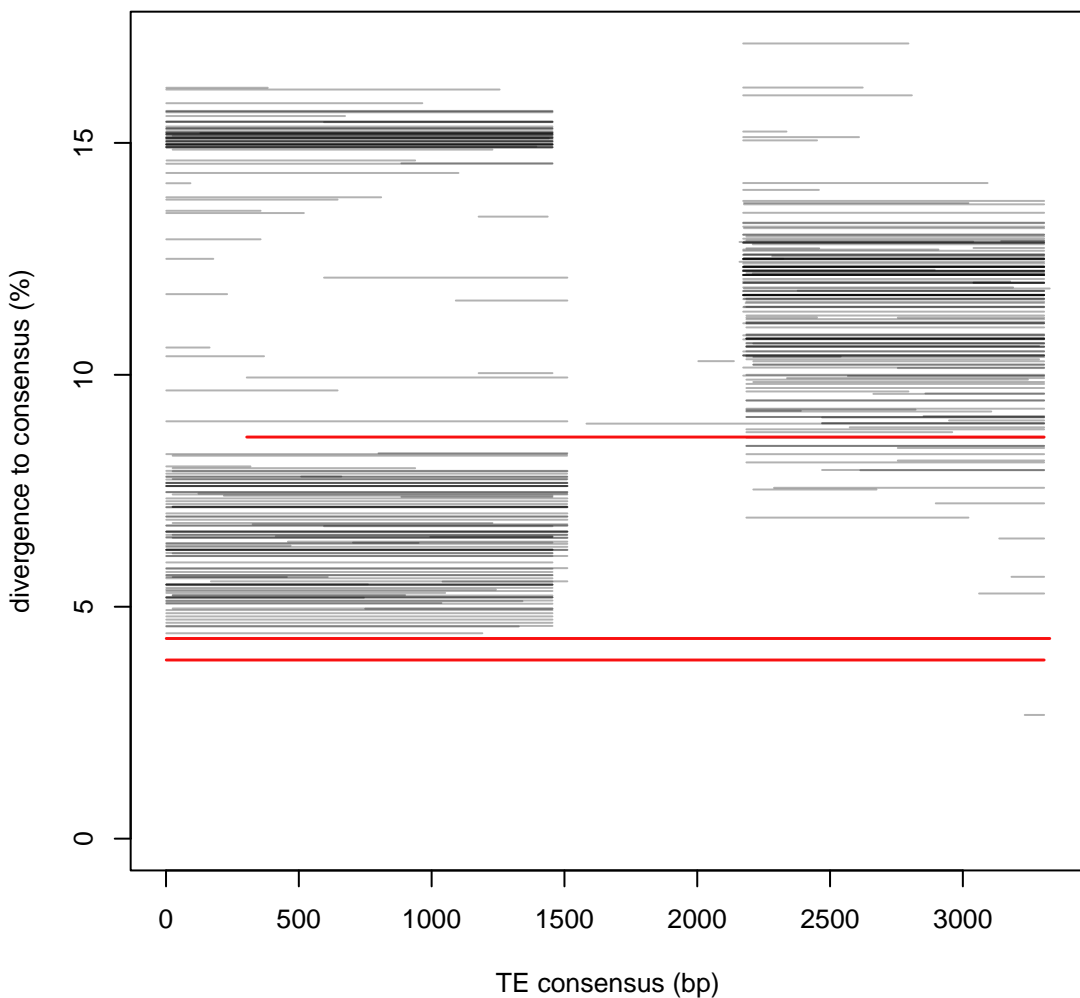
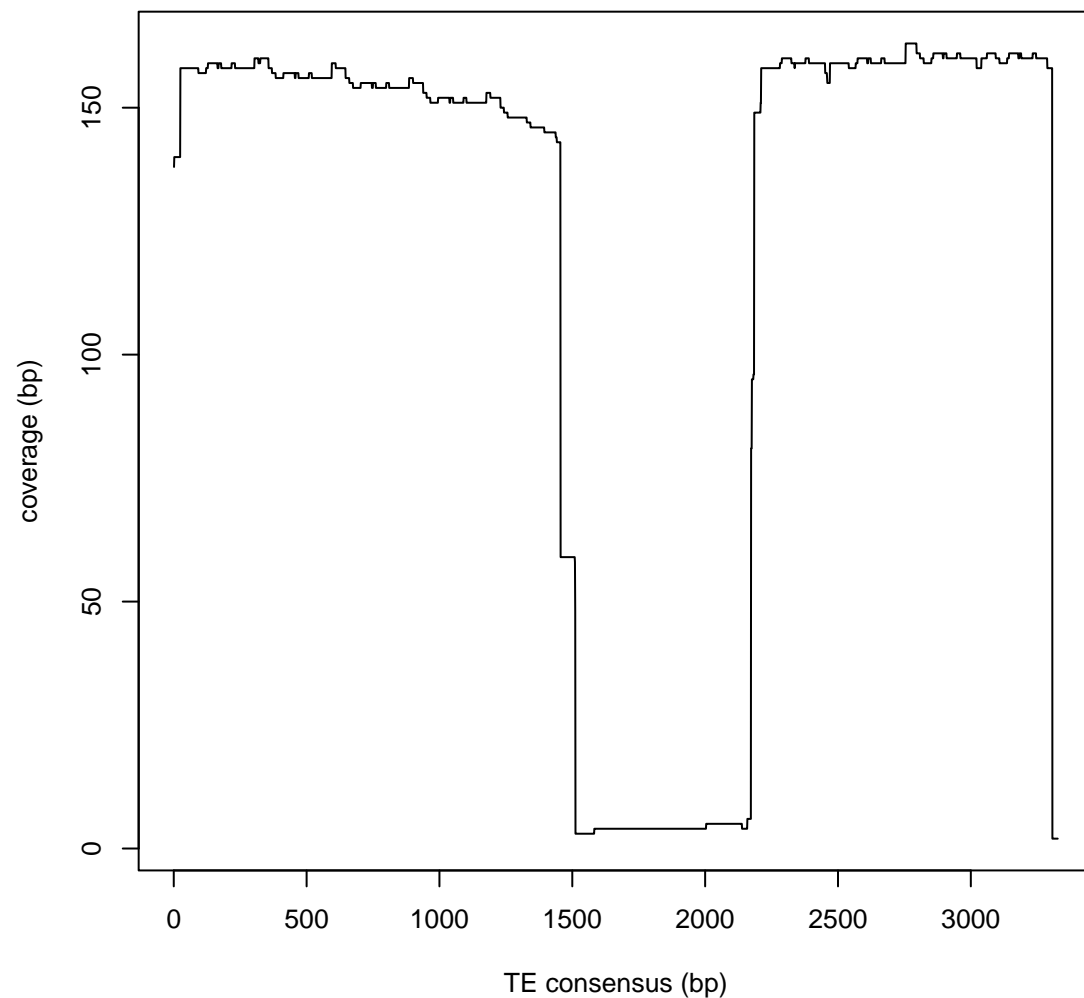


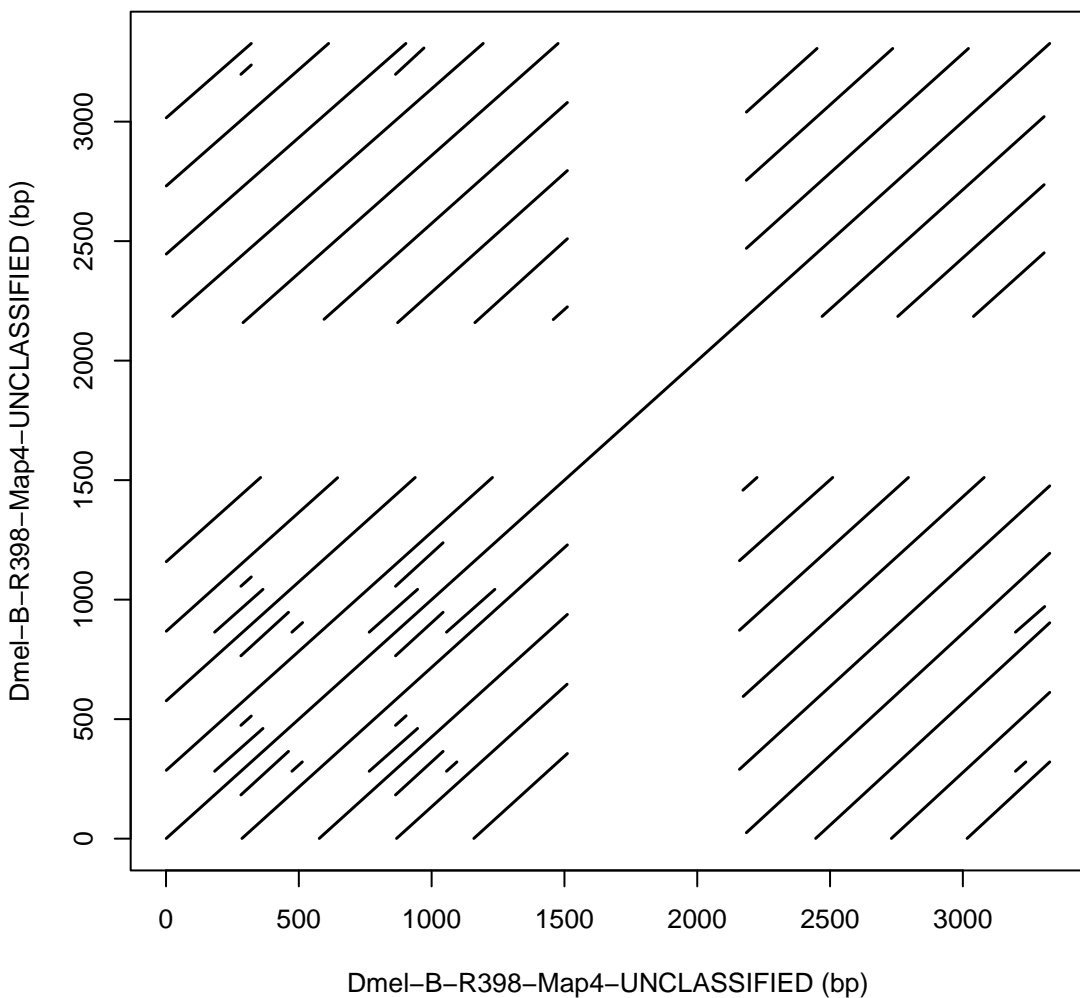
TE: Dmel-B-R398-Map4-UNCLASSIFIED  
consensus size: 3327bp; fragments: 365; full length: 3 (>=2994.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

