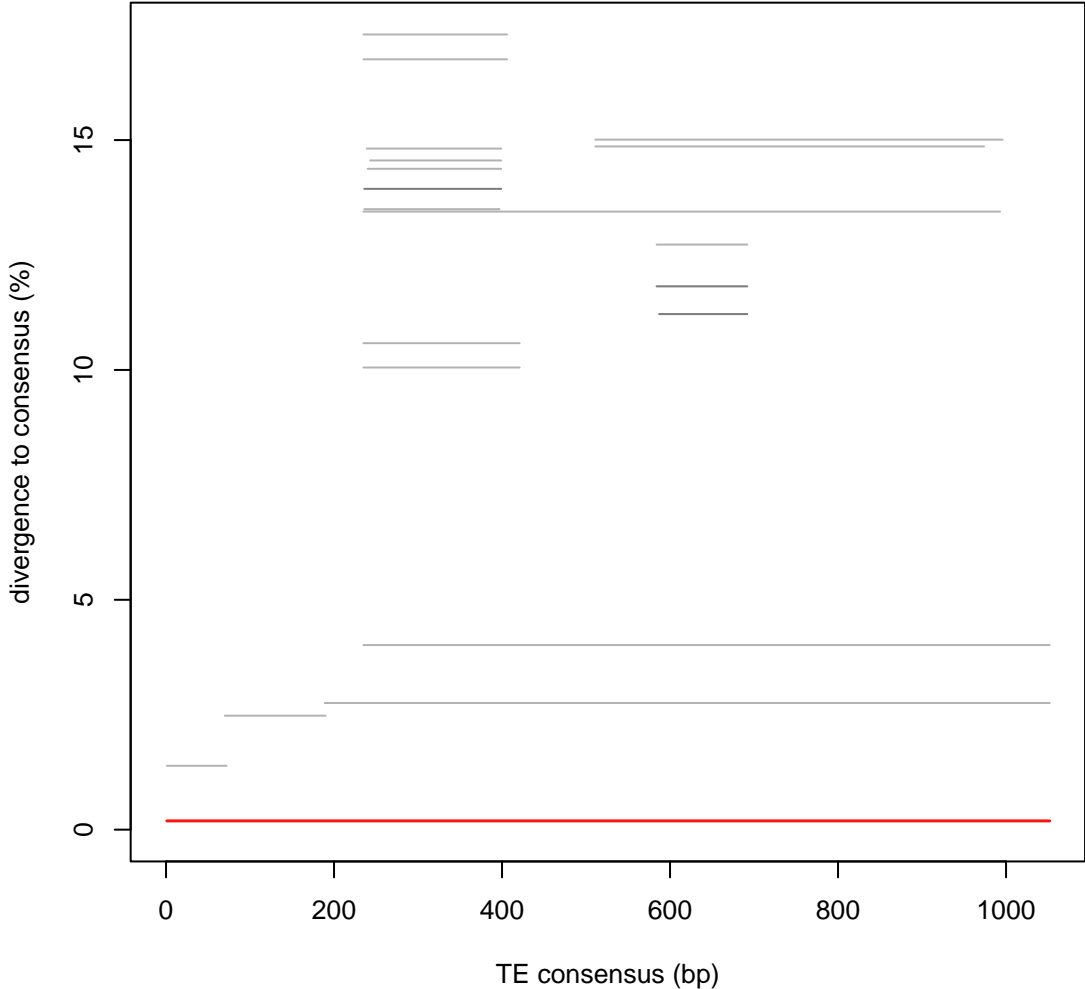
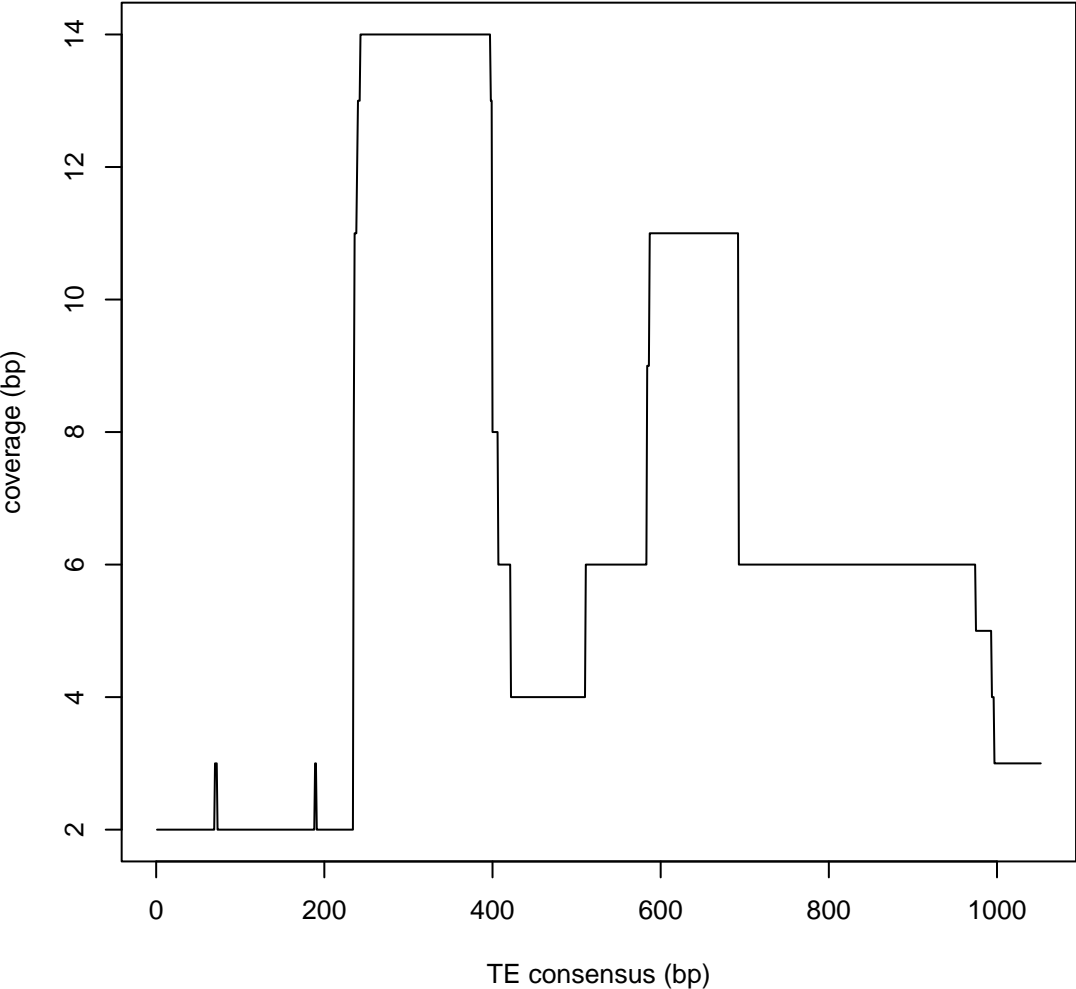


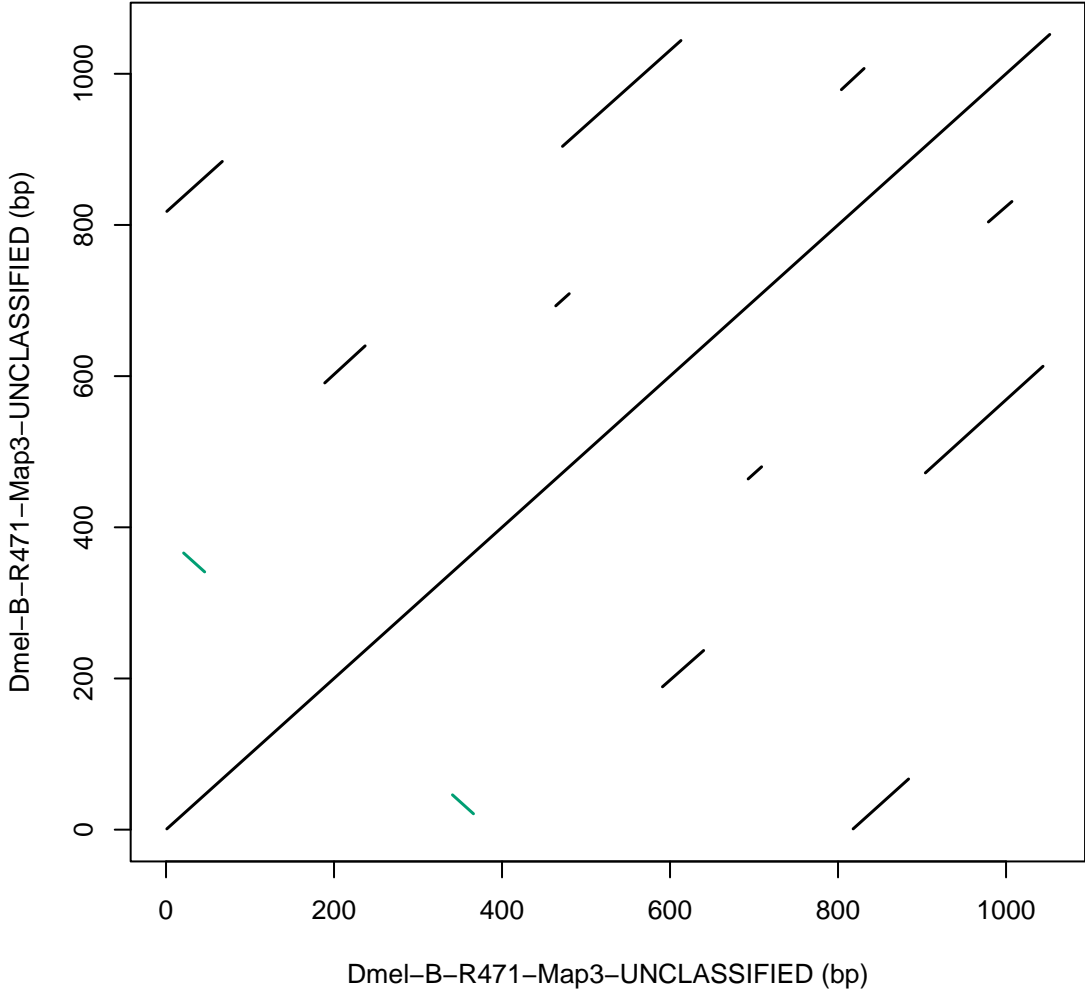
TE: Dmel-B-R471-Map3-UNCLASSIFIED
consensus size: 1052bp; fragments: 23; full length: 1 (>=946.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

