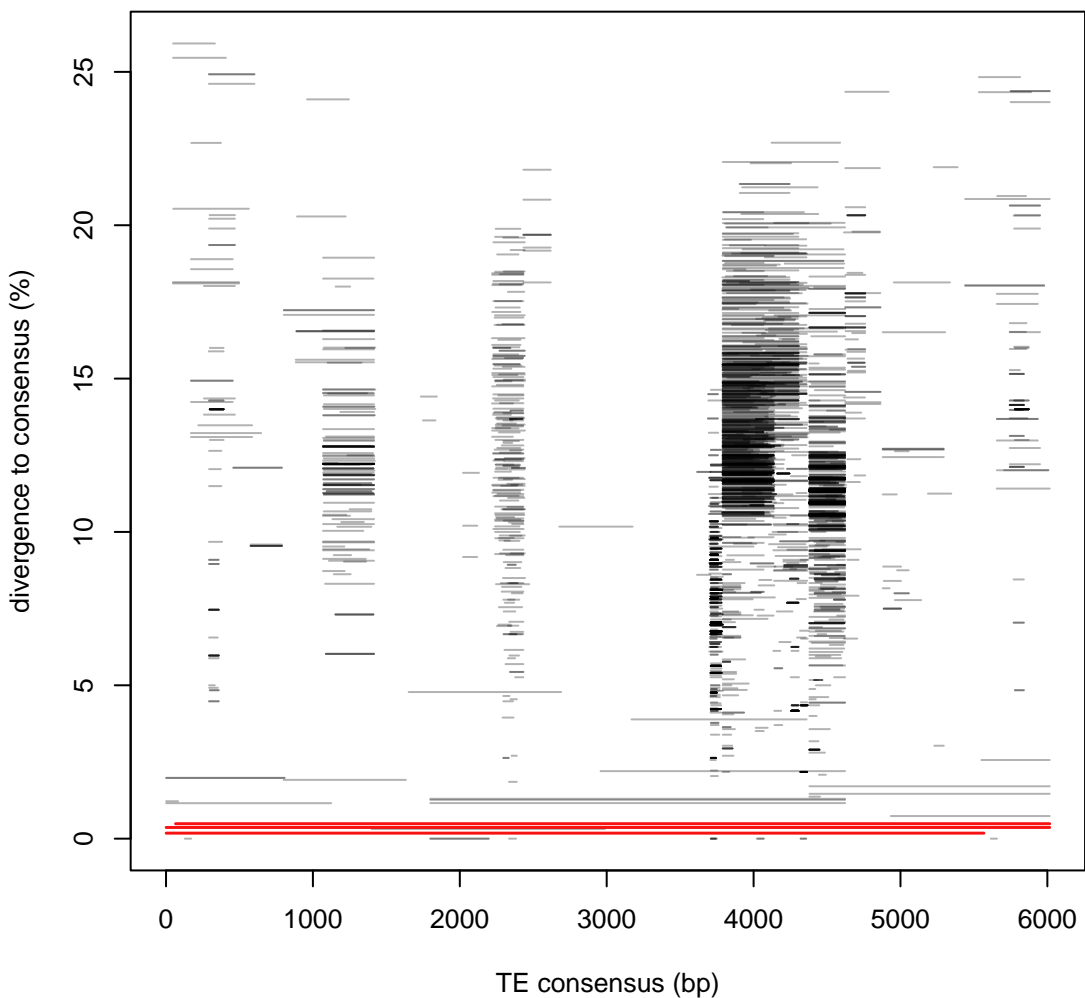
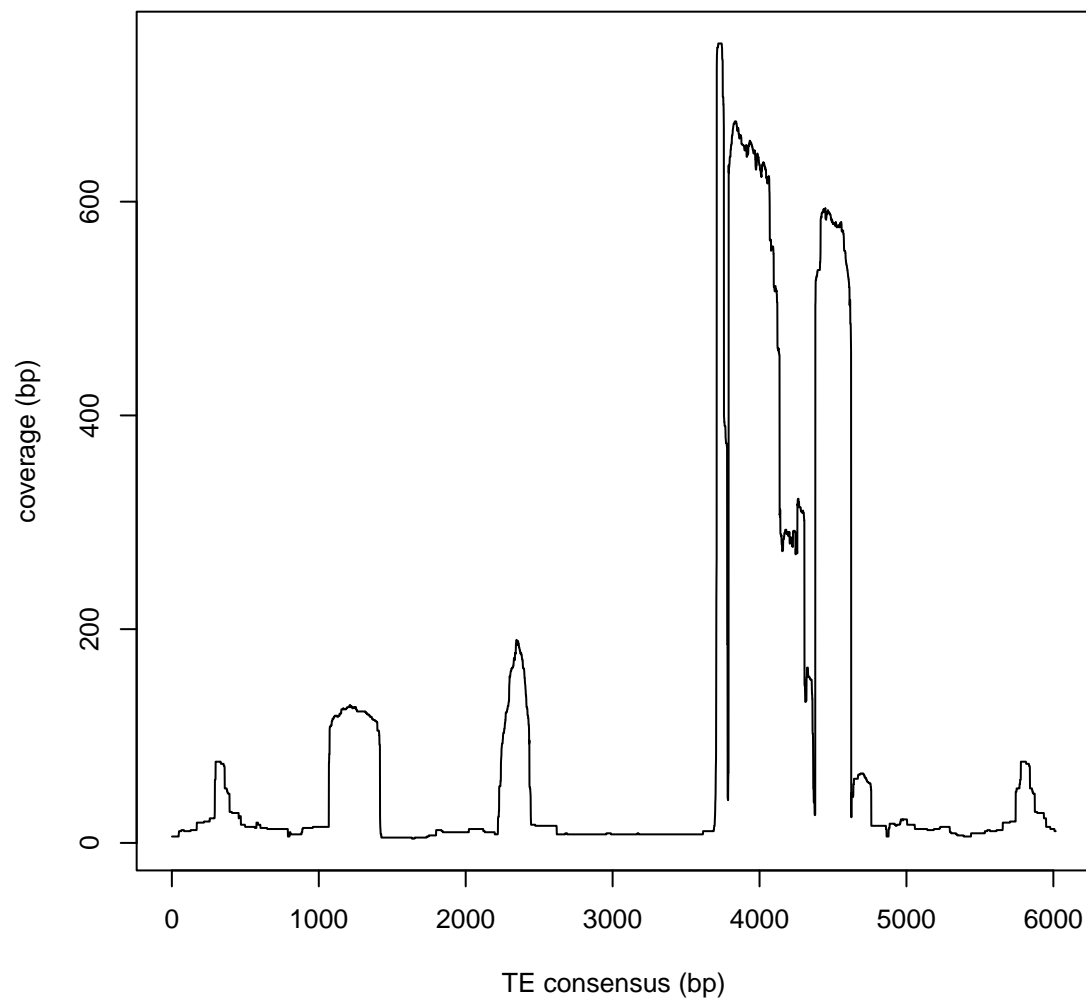


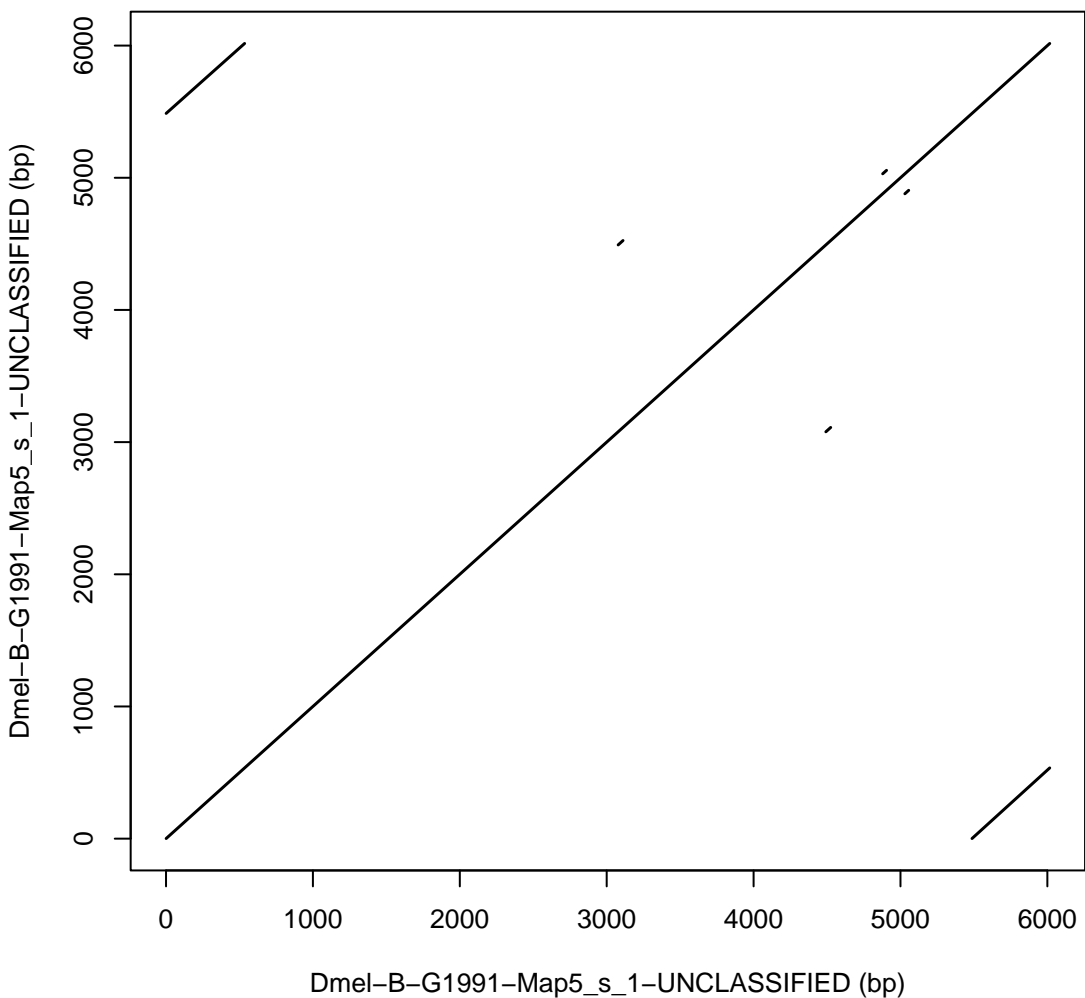
TE: Dmel-B-G1991-Map5_s_1-UNCLASSIFIED
consensus size: 6016bp; fragments: 2903; full length: 3 (>=5414.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

