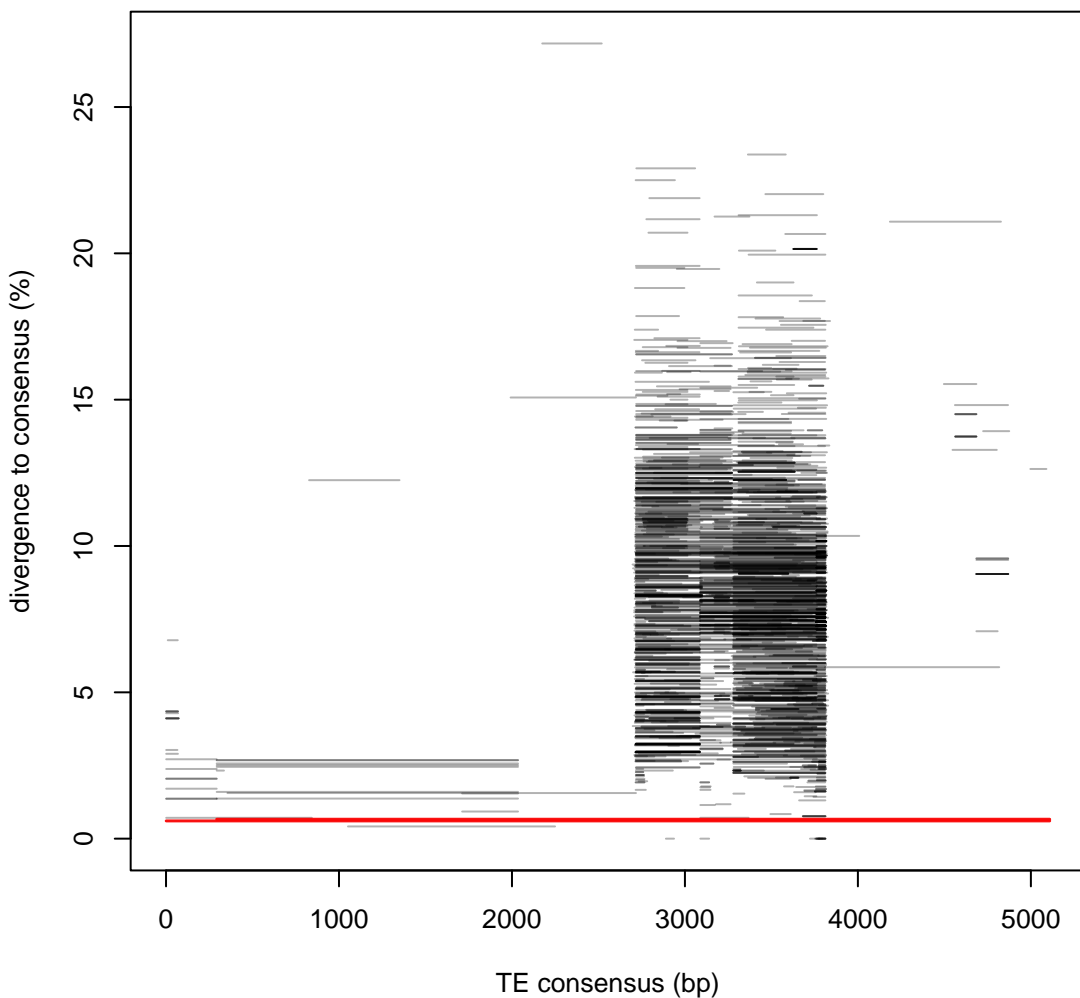
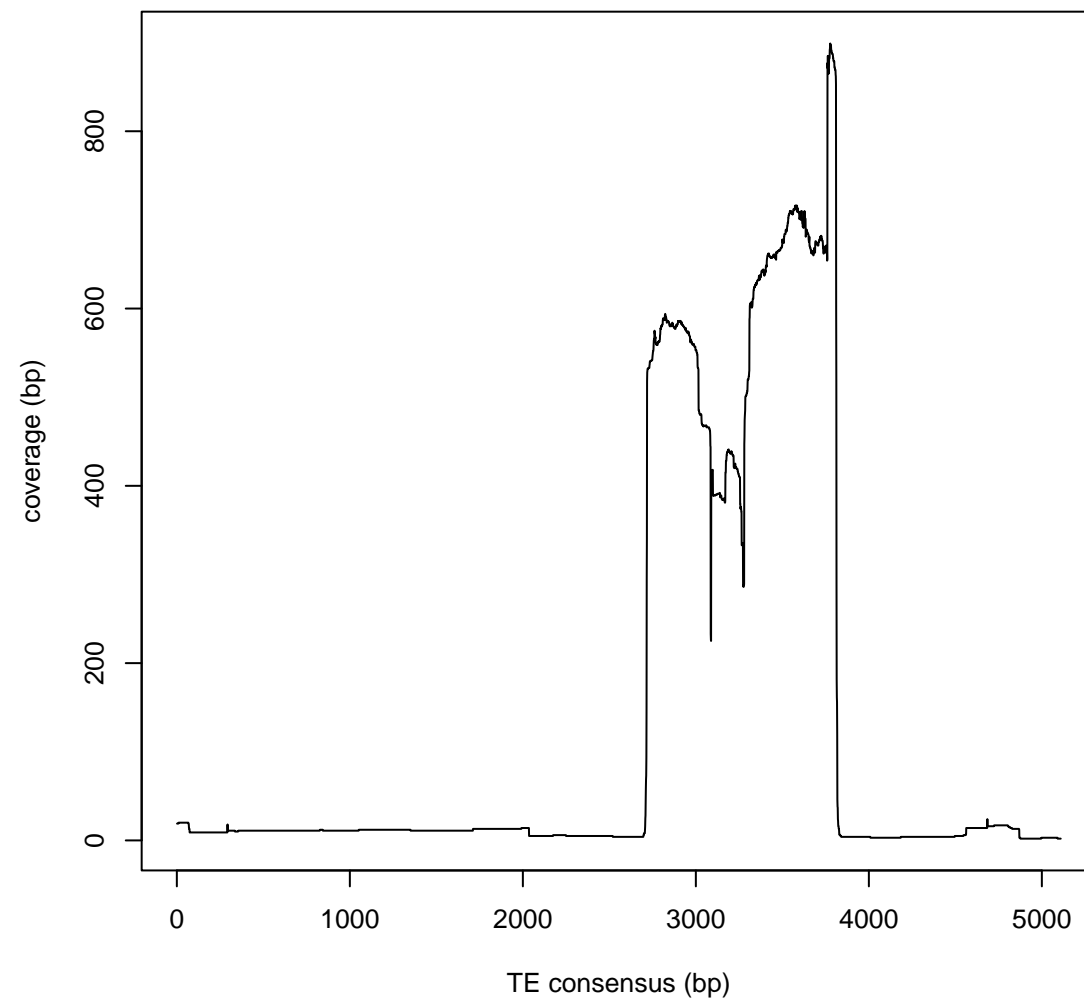


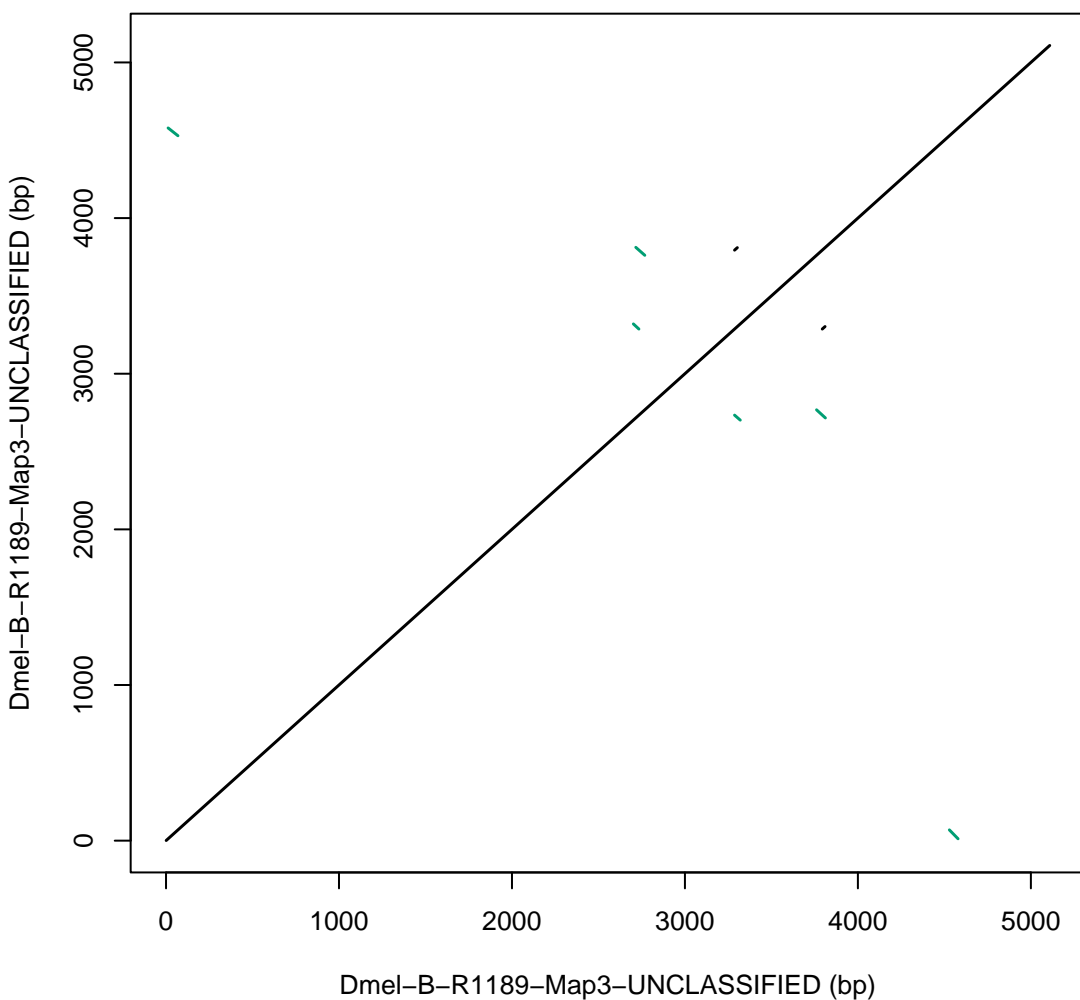
TE: Dmel-B-R1189-Map3-UNCLASSIFIED
consensus size: 5109bp; fragments: 2094; full length: 2 (>=4598.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

