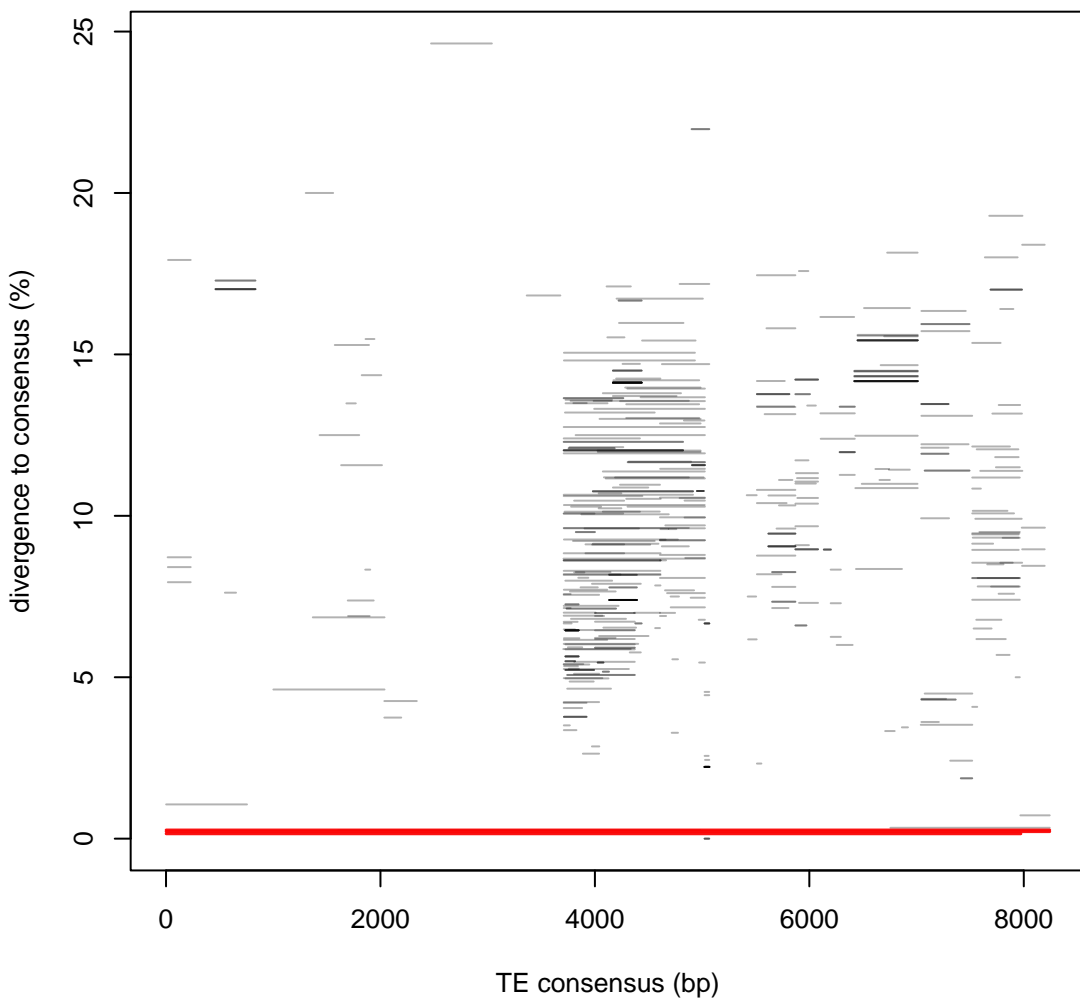
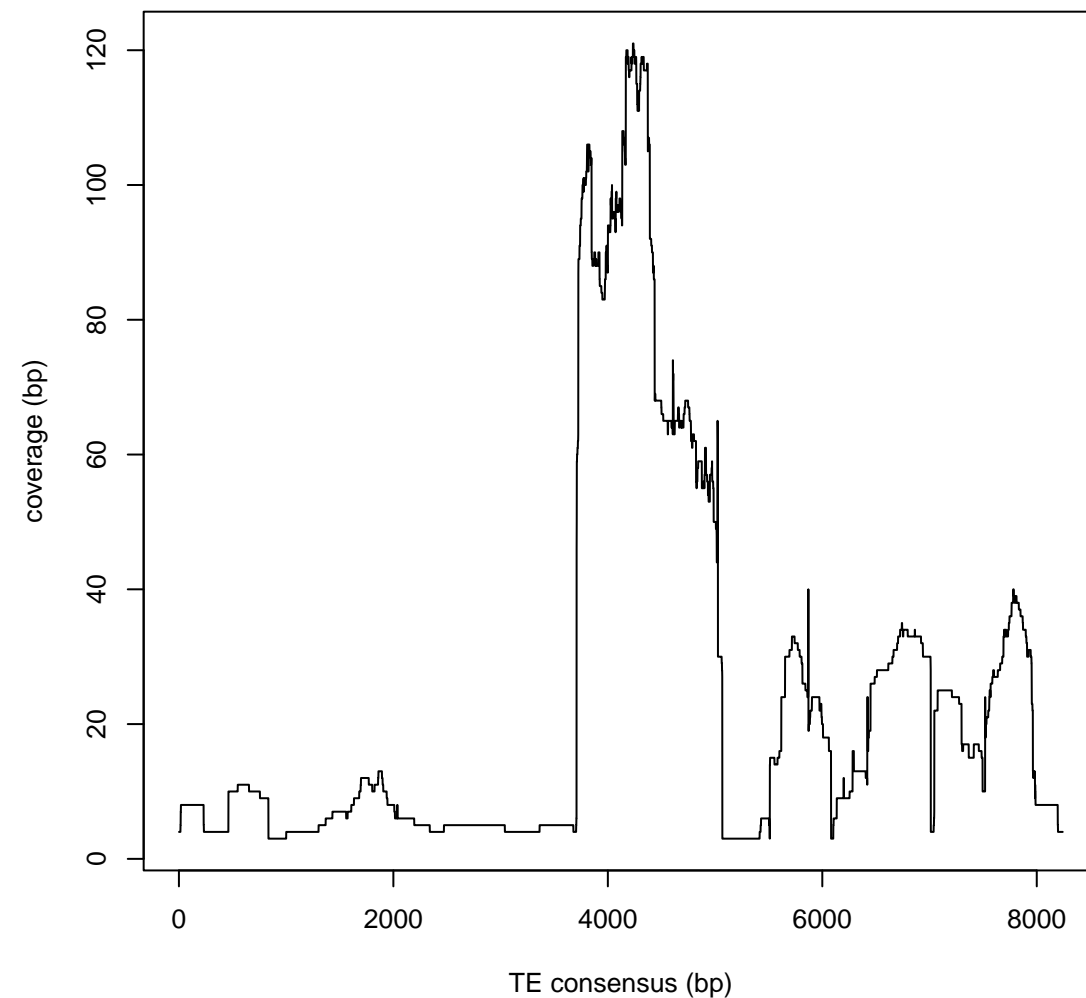


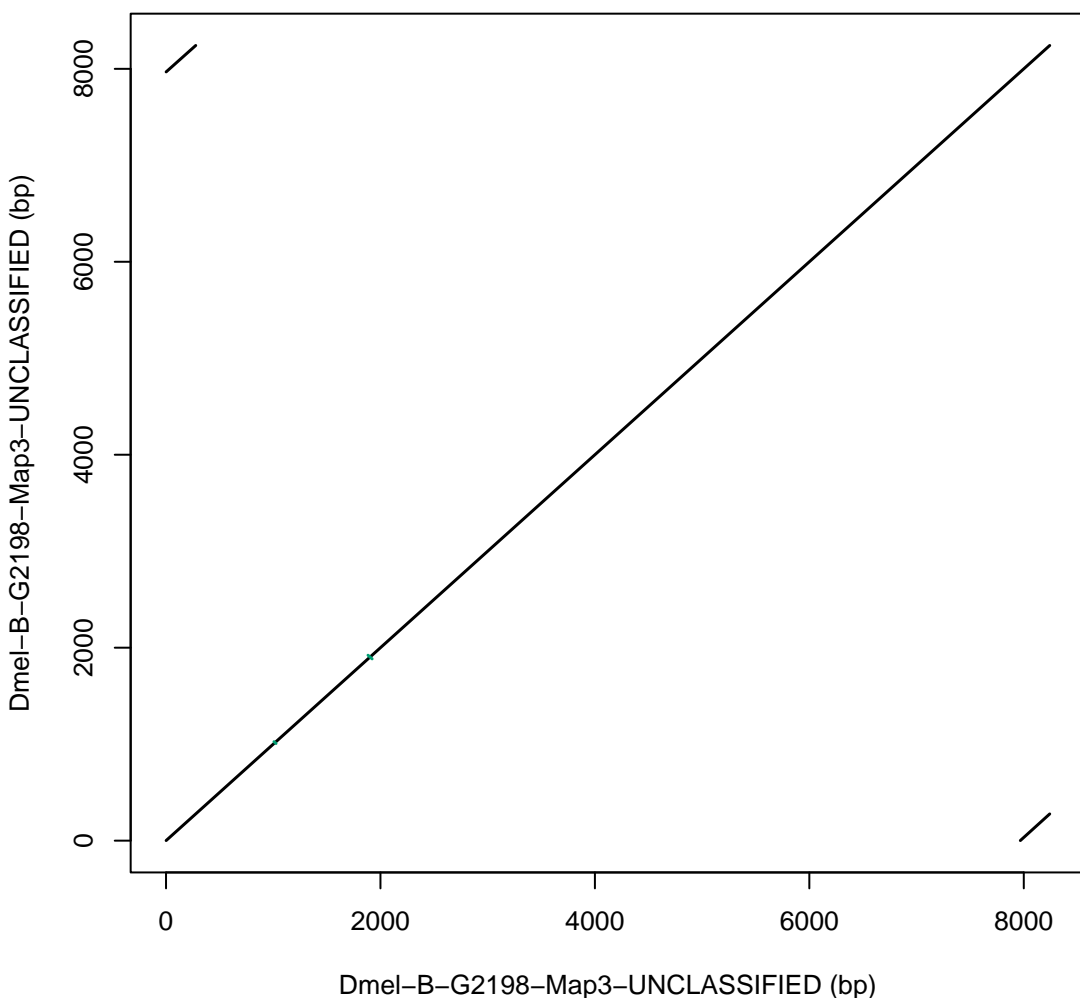
TE: Dmel-B-G2198-Map3-UNCLASSIFIED
consensus size: 8242bp; fragments: 486; full length: 3 (>=7417.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

