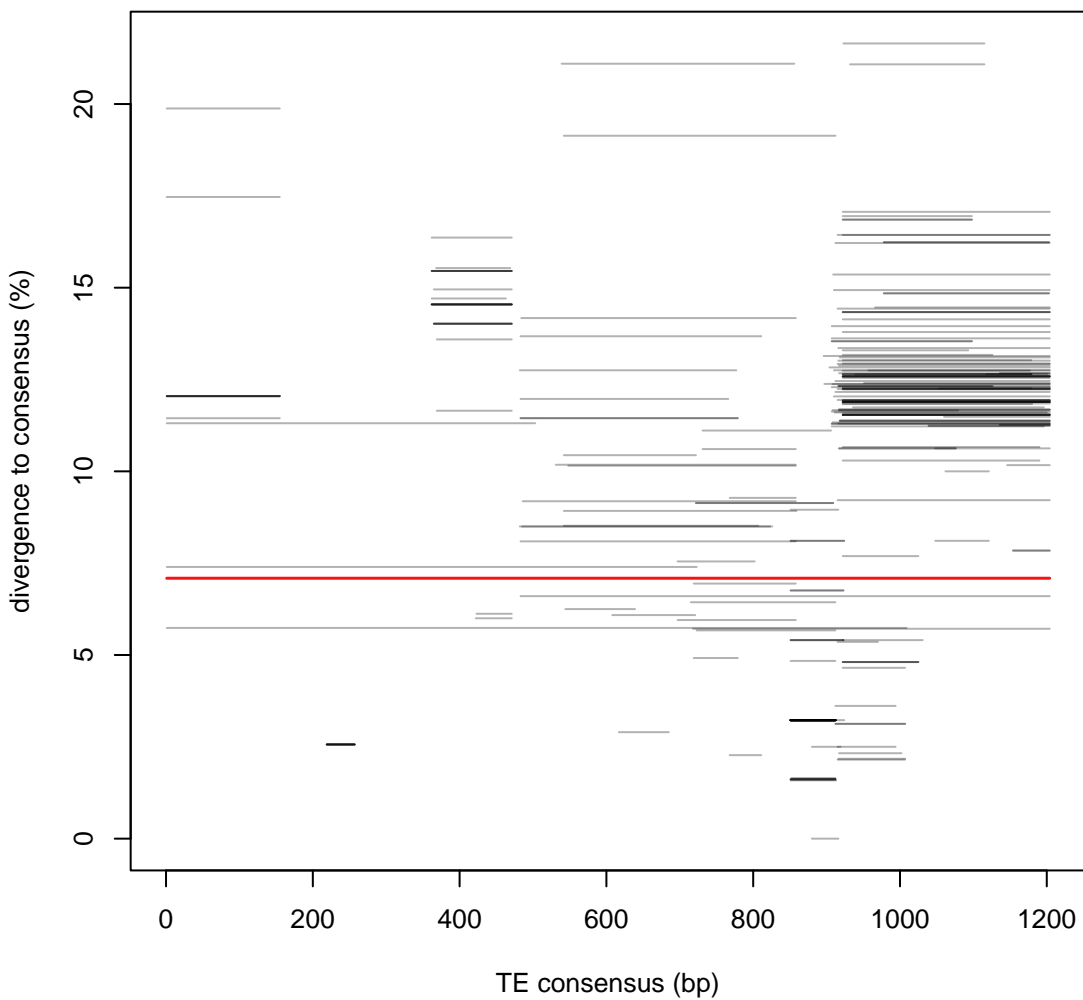
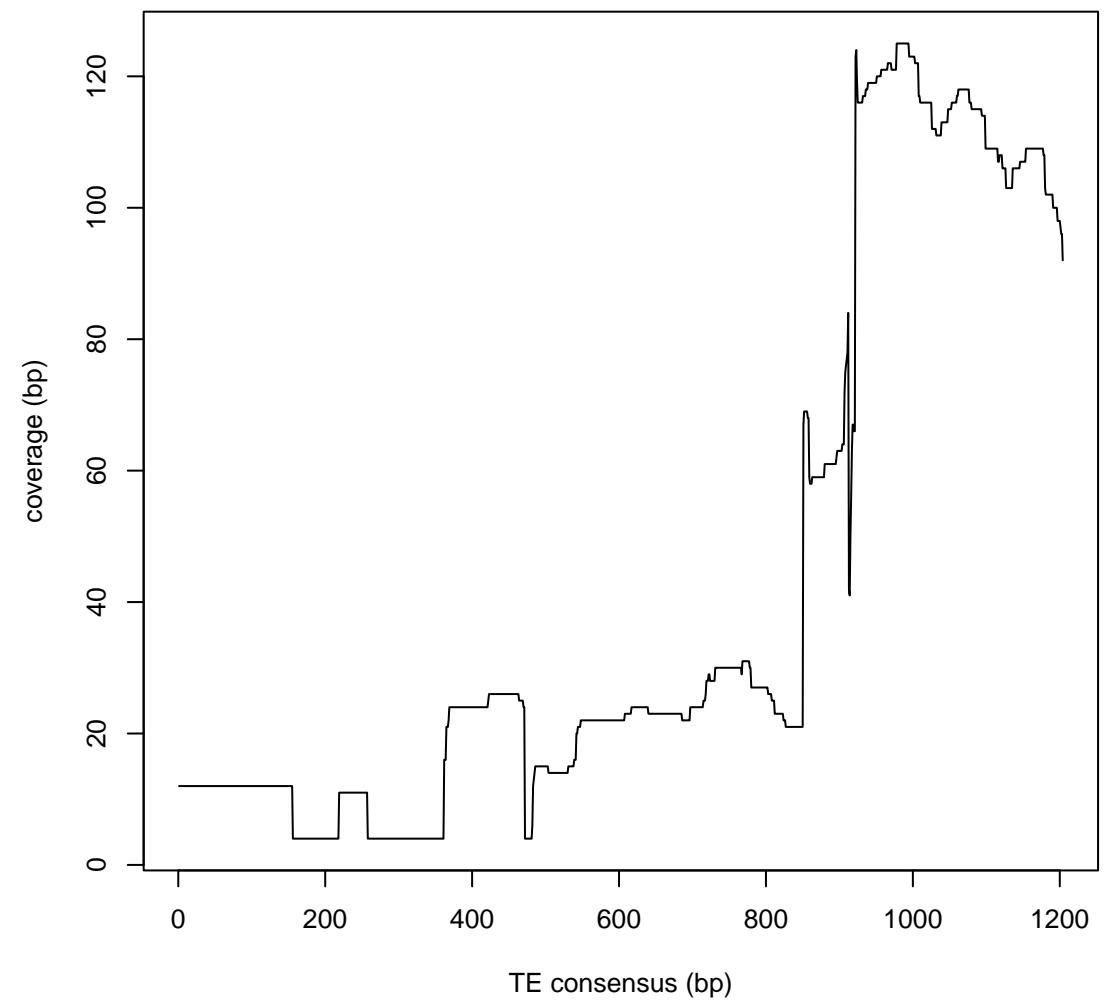


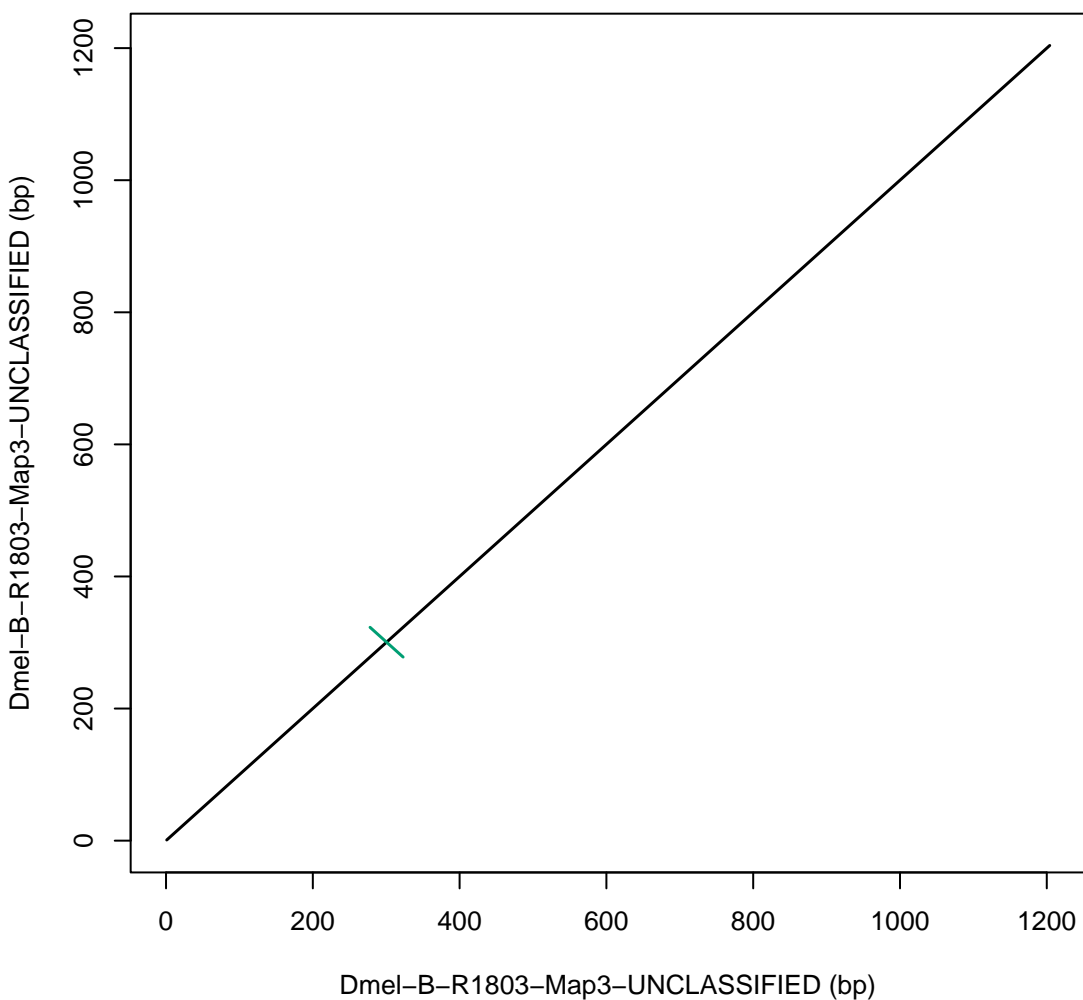
TE: Dmel-B-R1803-Map3-UNCLASSIFIED  
consensus size: 1204bp; fragments: 262; full length: 1 (>=1083.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

