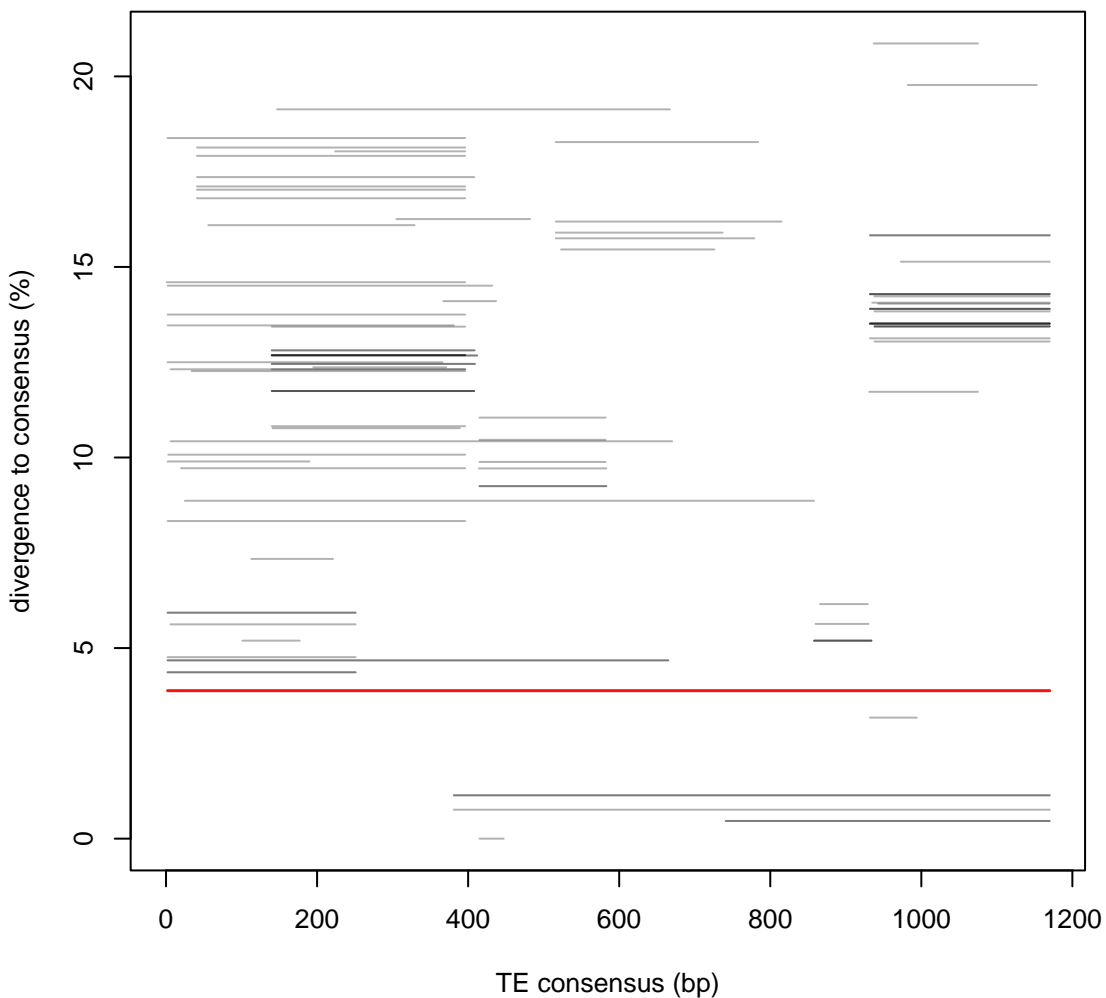
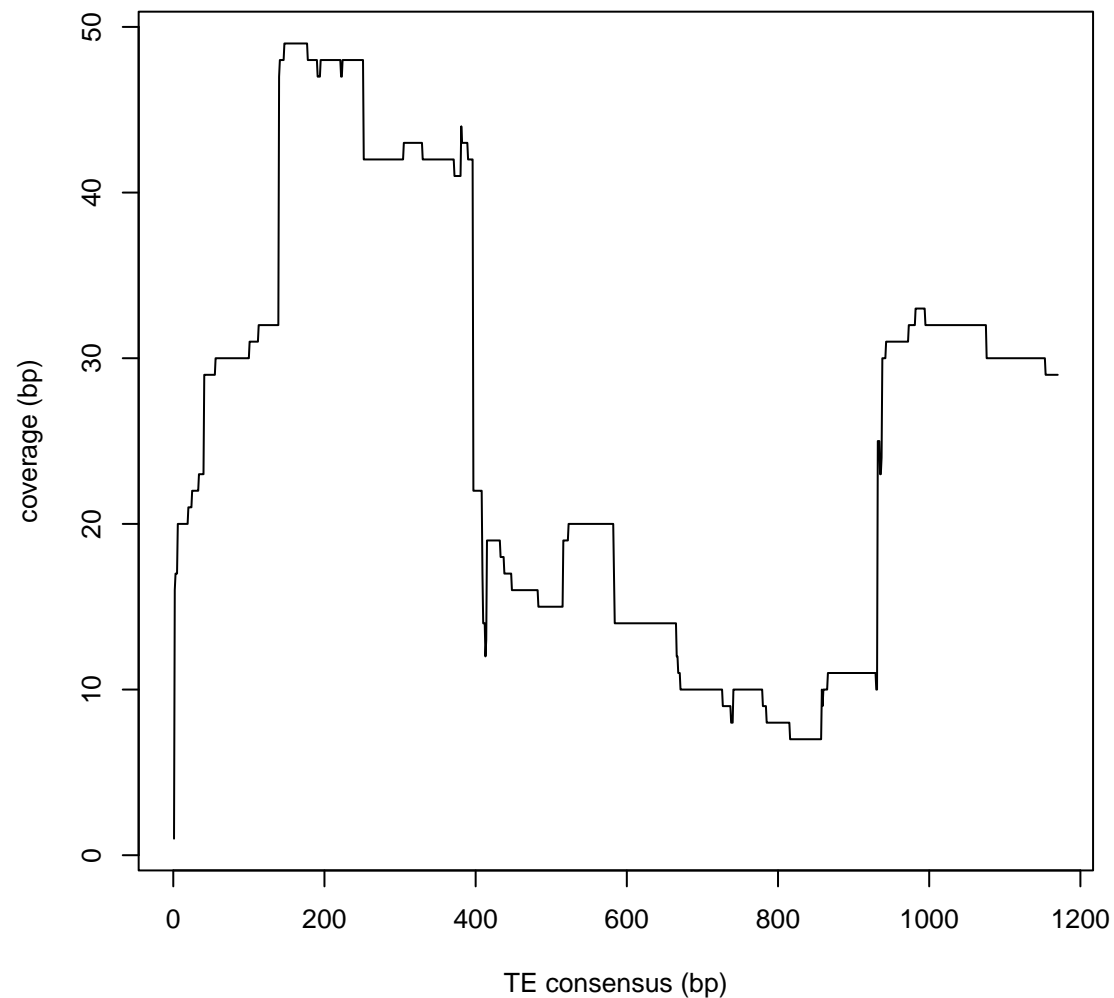


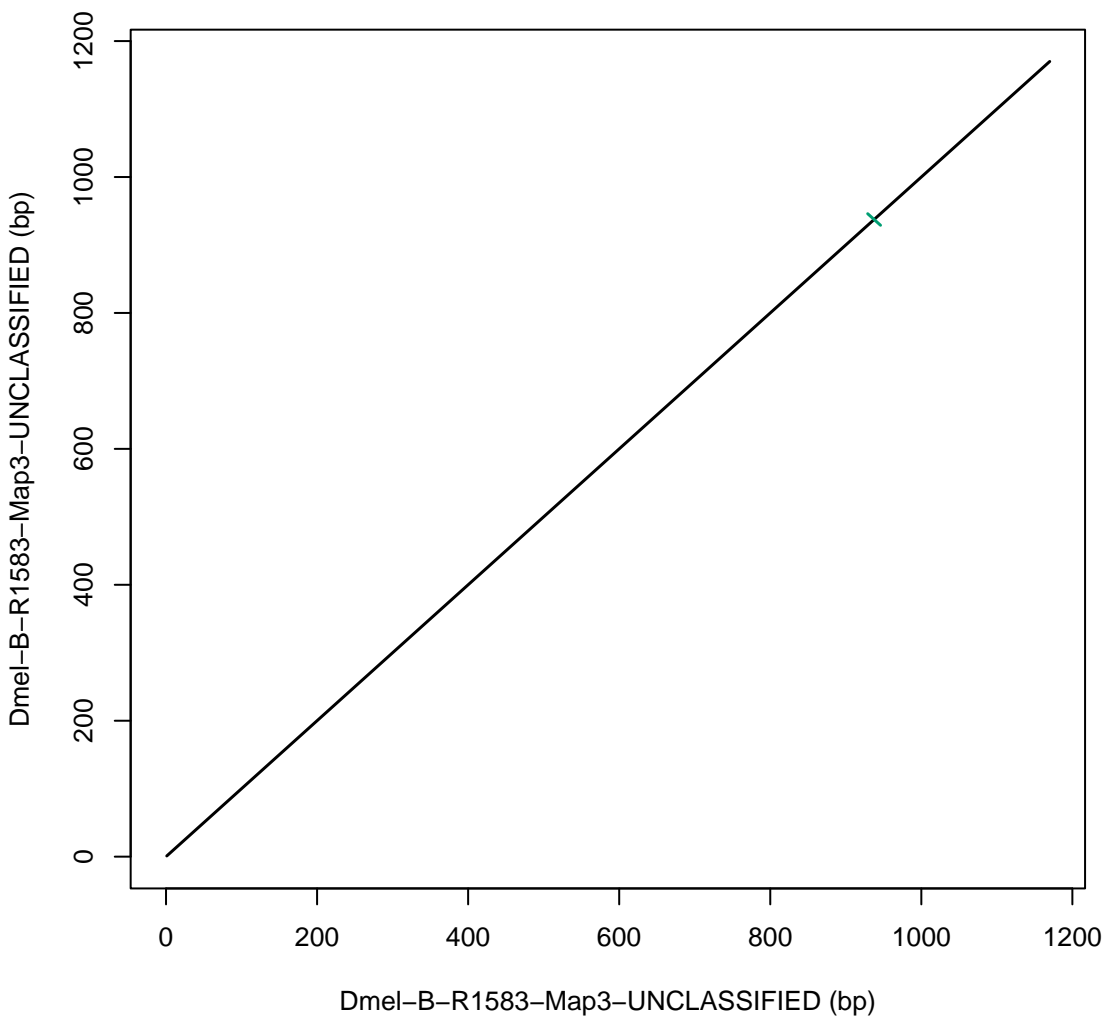
TE: Dmel-B-R1583-Map3-UNCLASSIFIED
consensus size: 1170bp; fragments: 102; full length: 1 (>=1053bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

