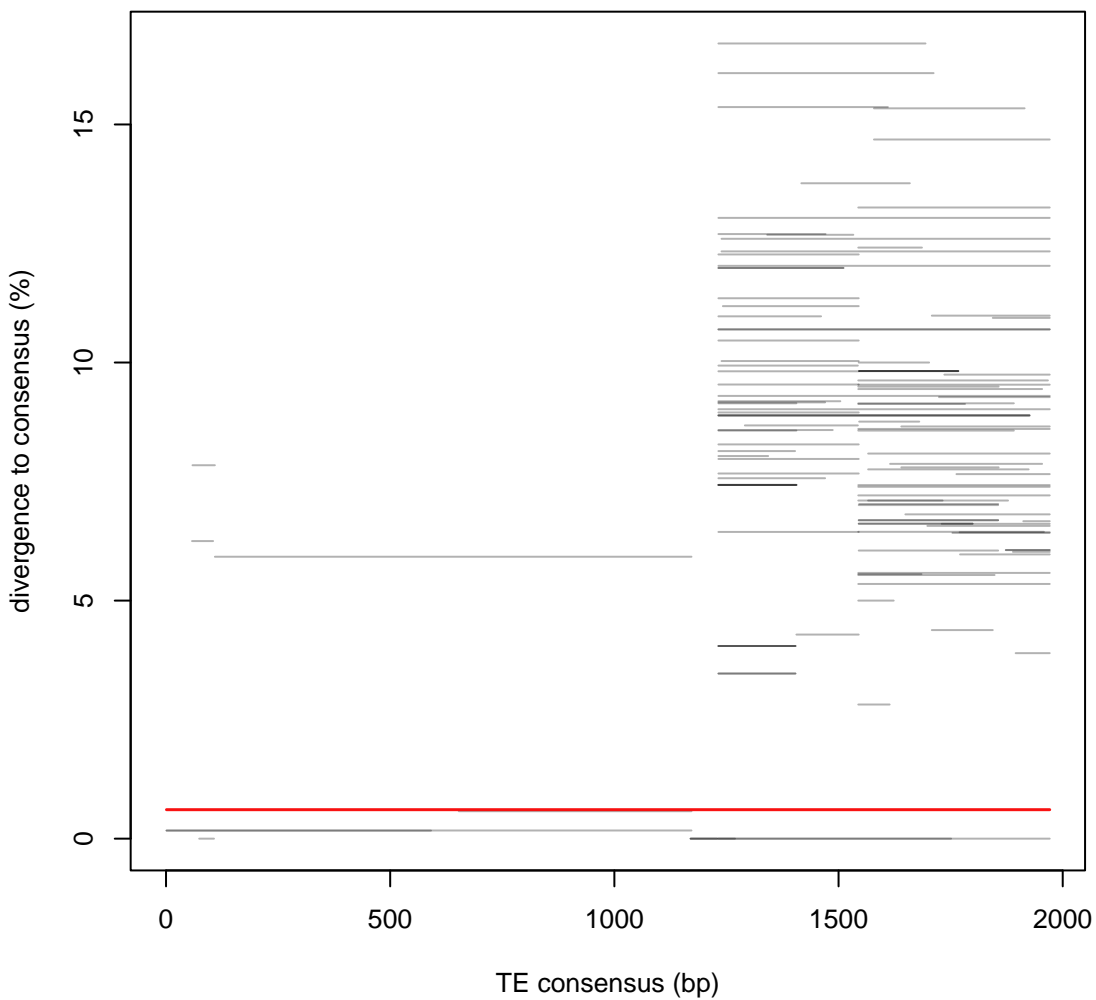
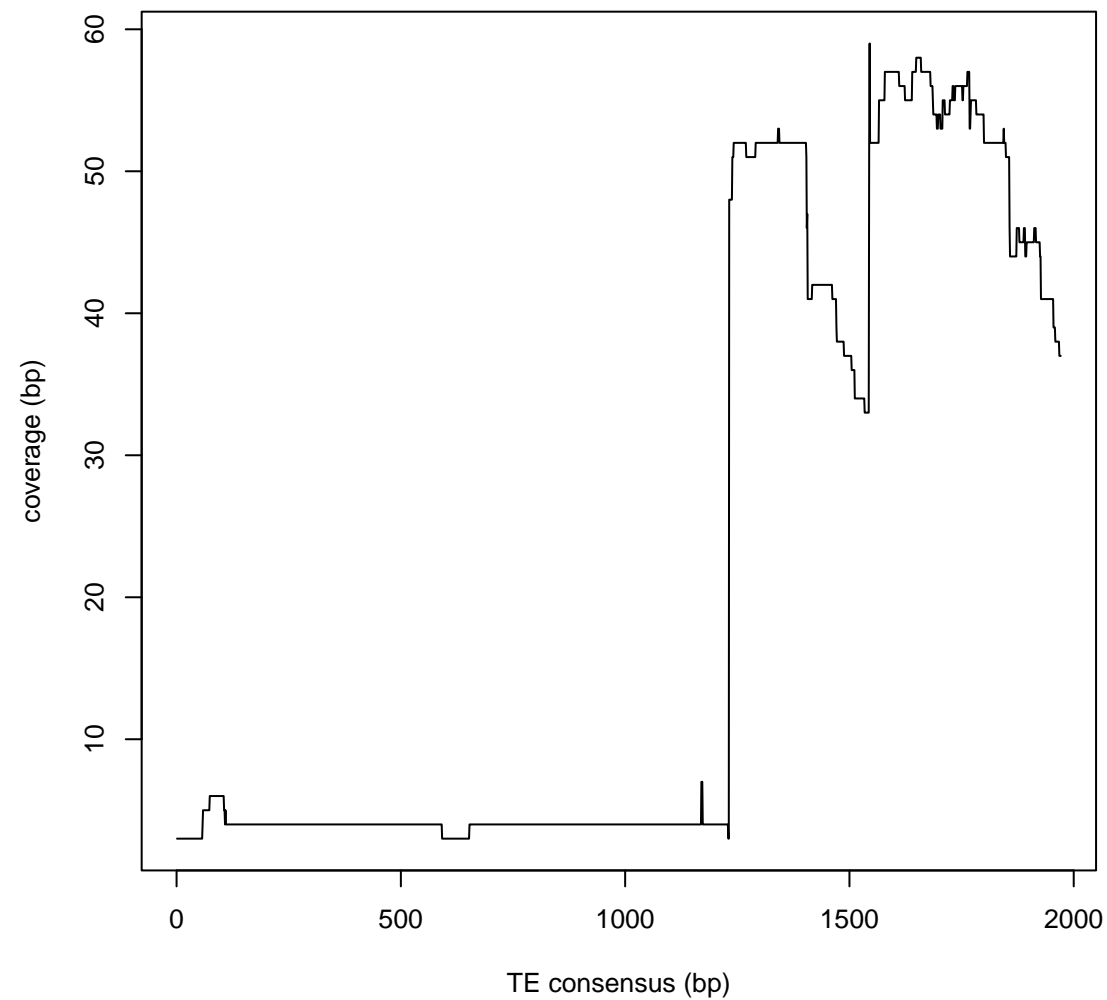


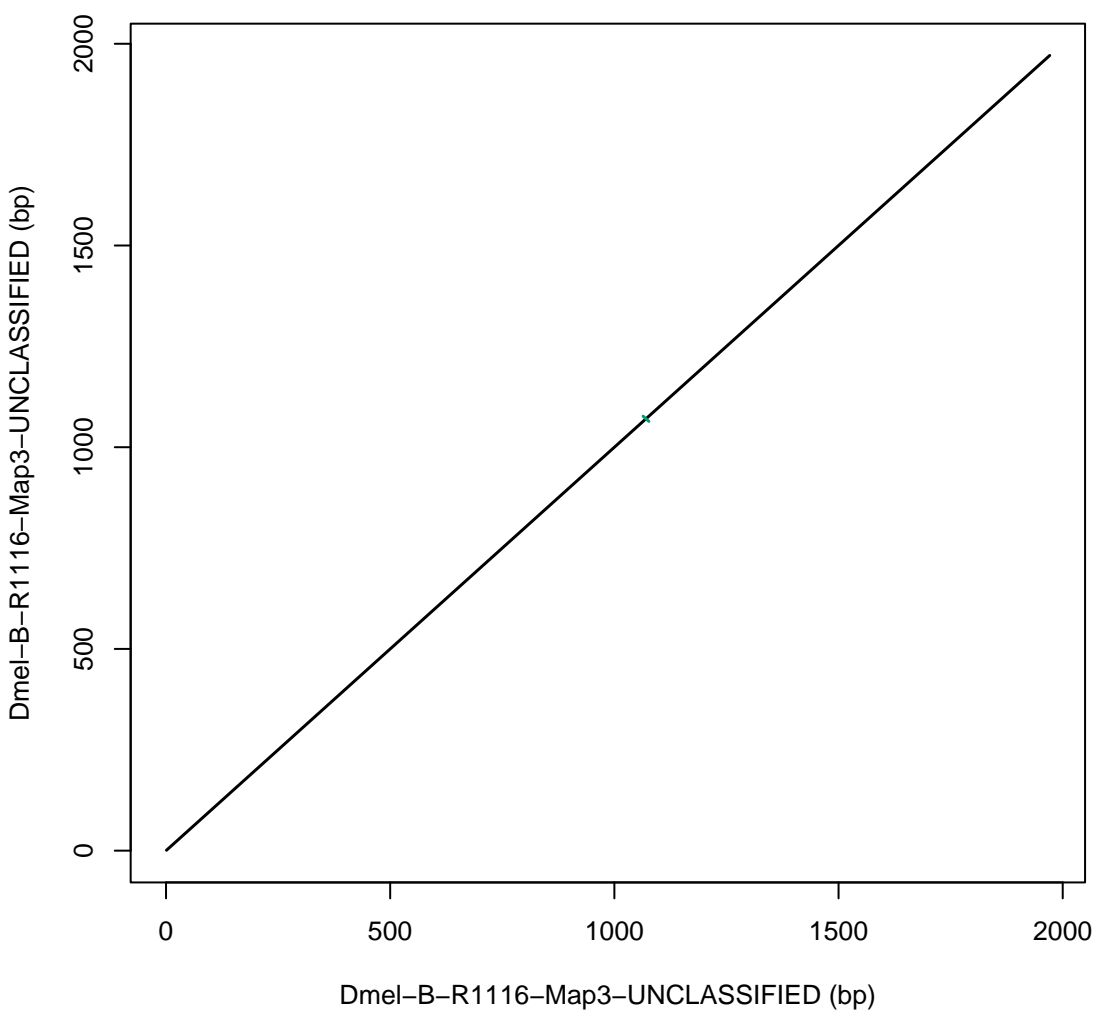
TE: Dmel-B-R1116-Map3-UNCLASSIFIED  
consensus size: 1971bp; fragments: 123; full length: 1 (>=1773.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

