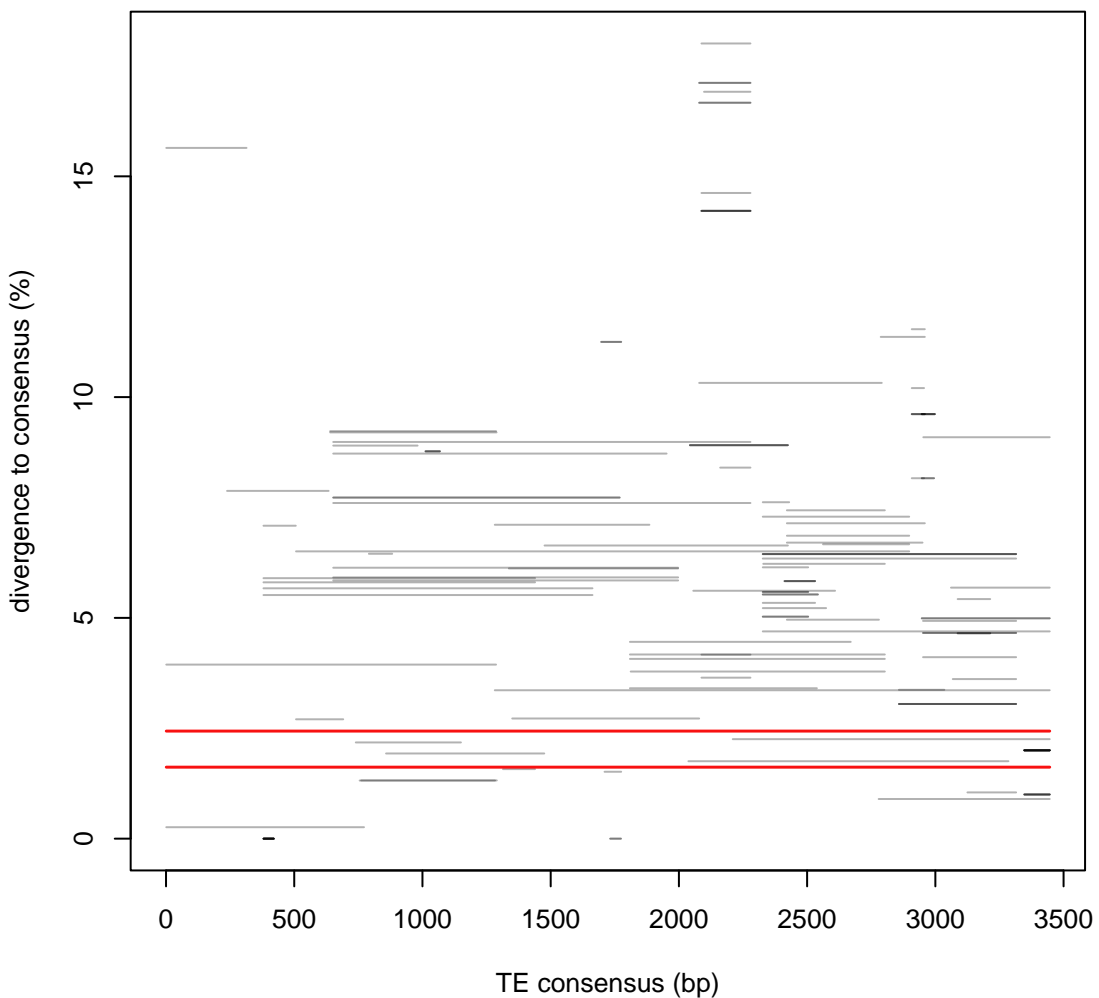
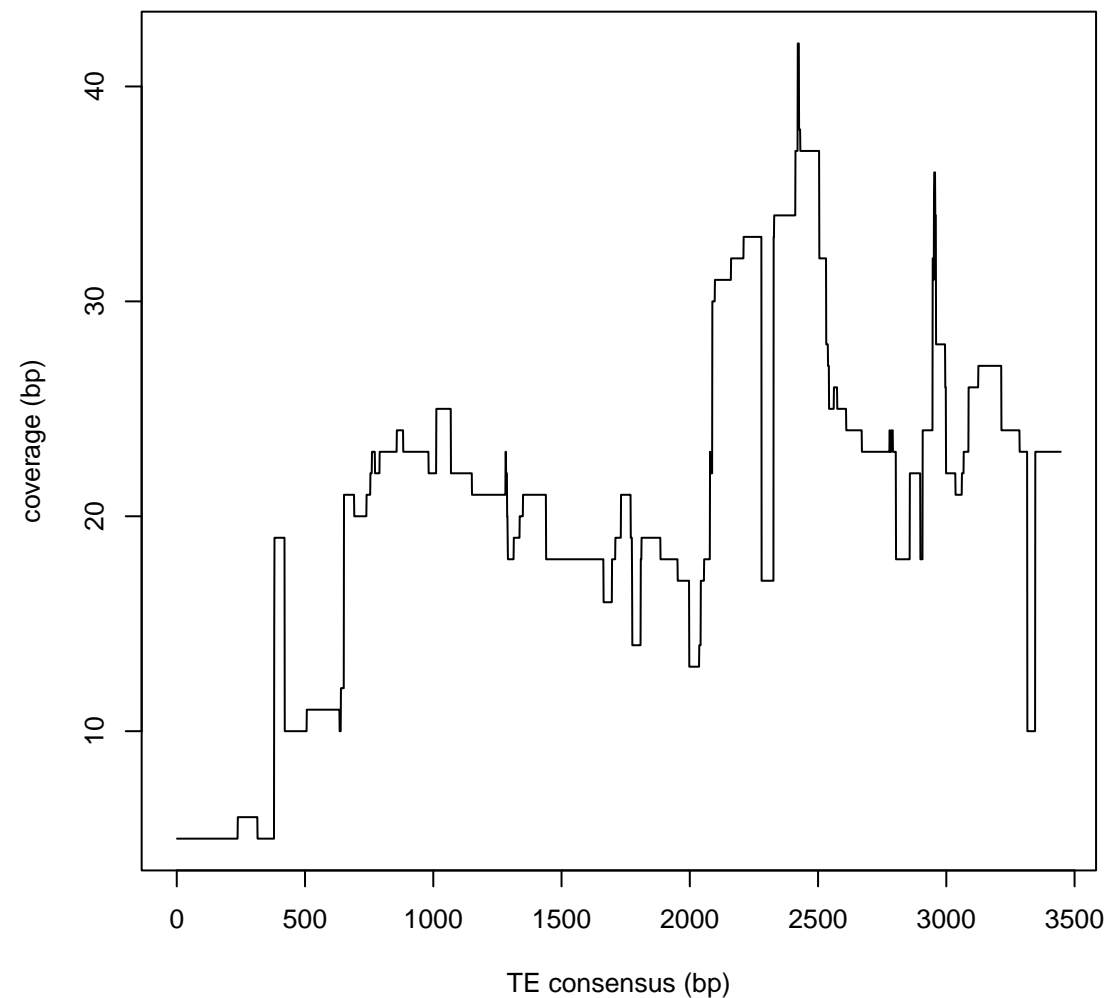


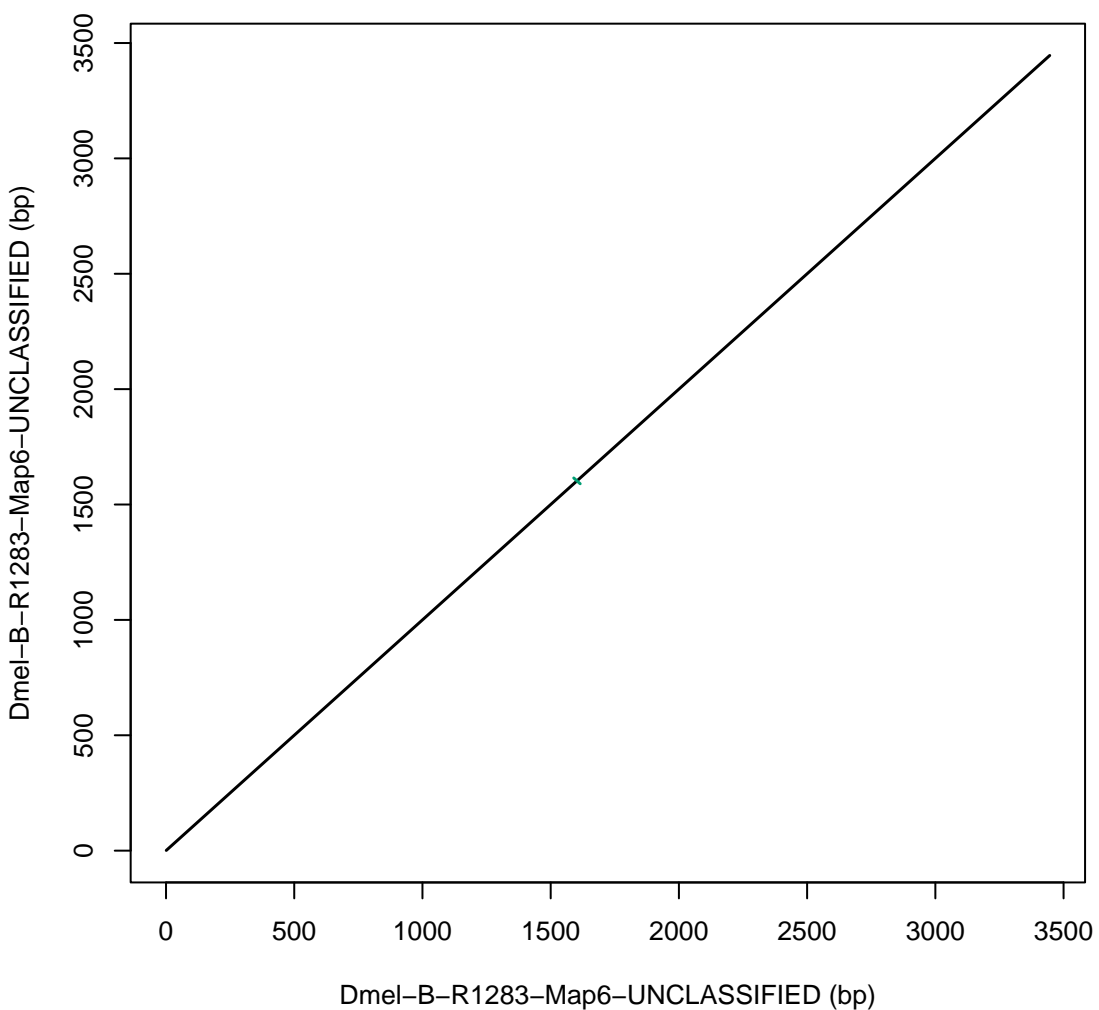
TE: Dmel-B-R1283-Map6-UNCLASSIFIED  
consensus size: 3446bp; fragments: 148; full length: 2 (>=3101.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

