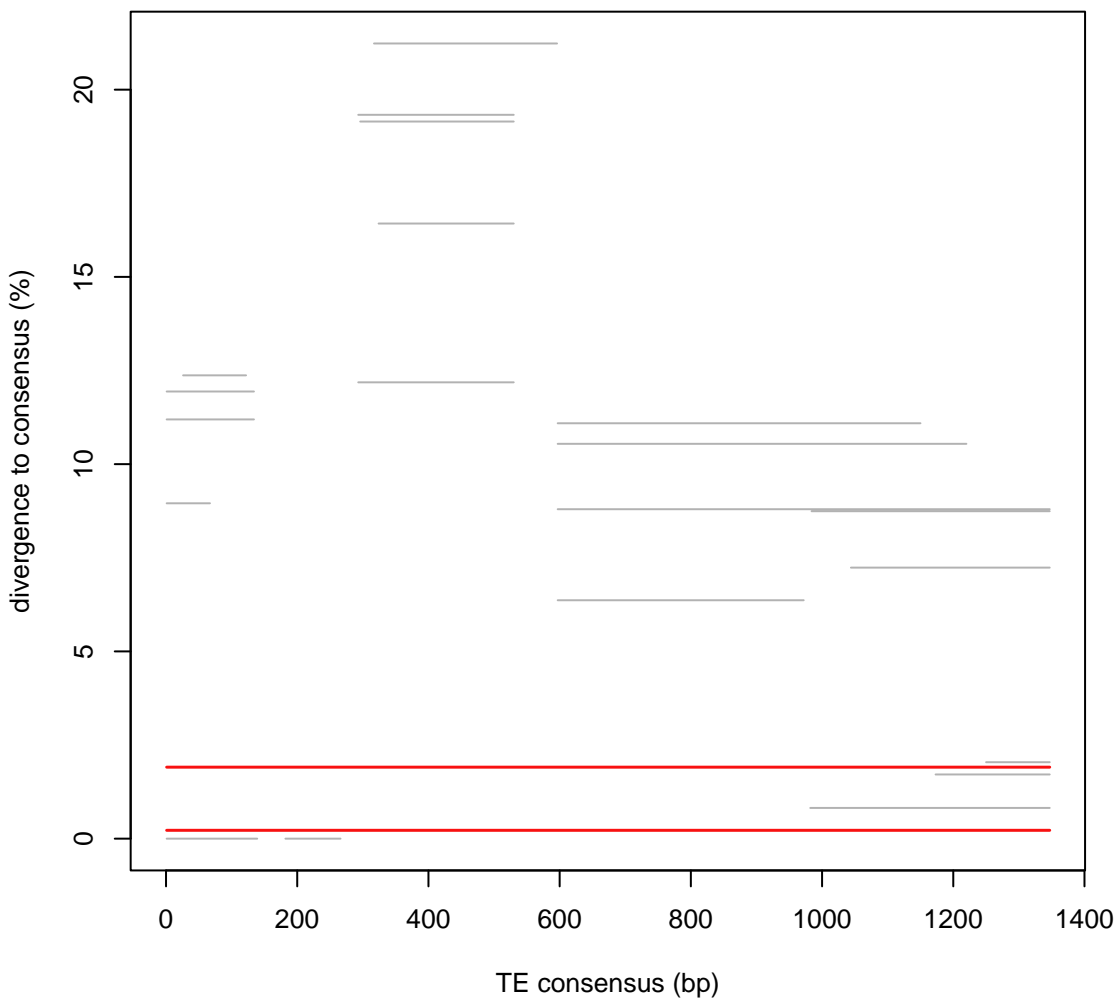
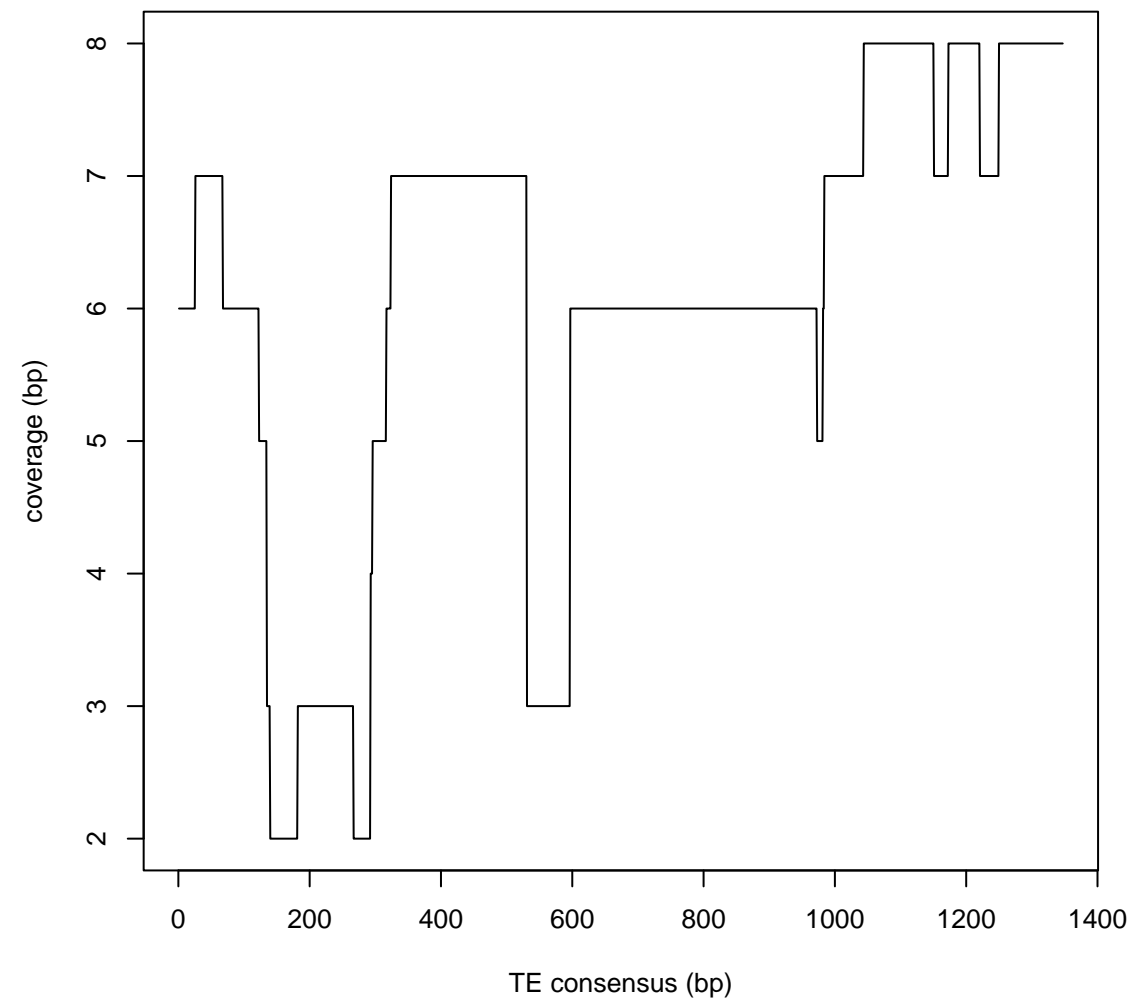


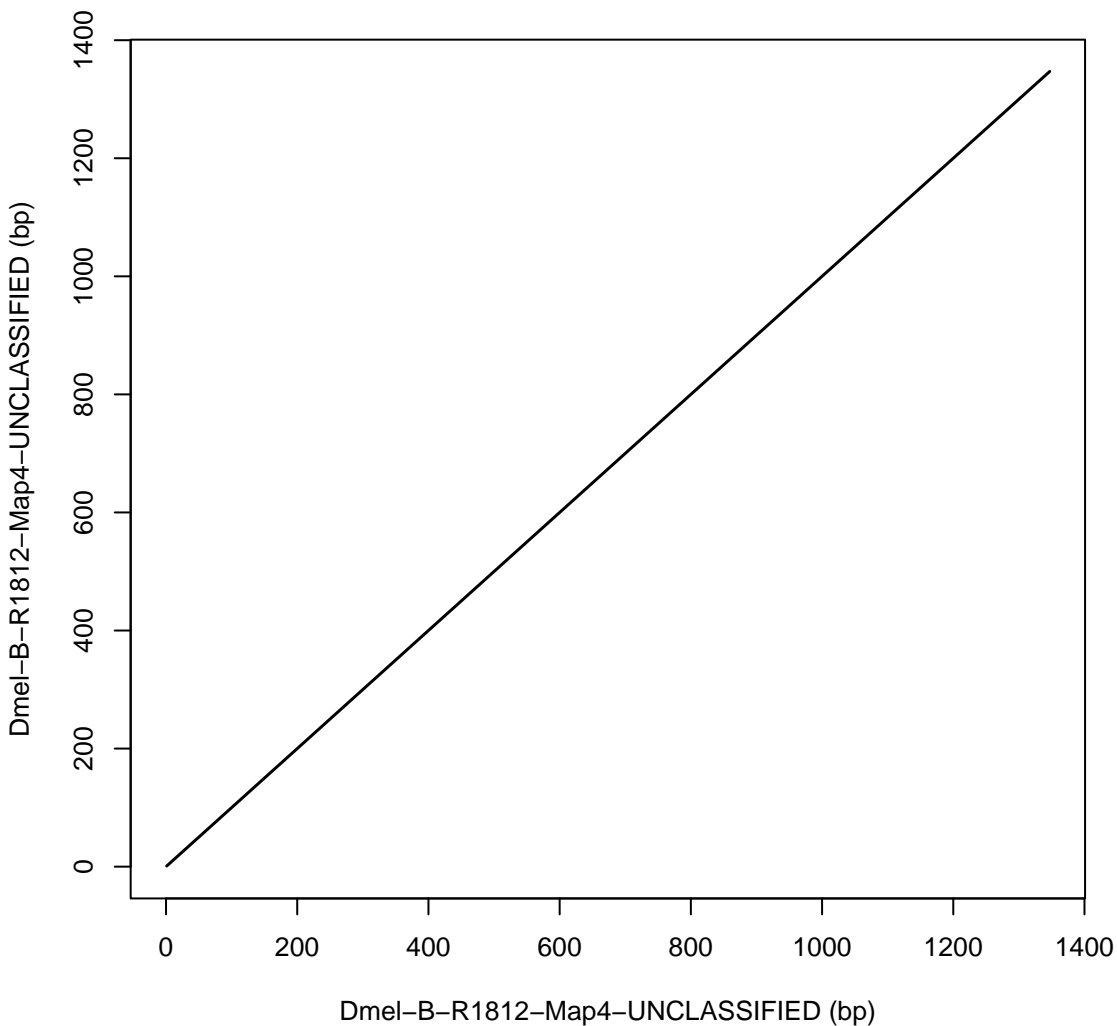
TE: Dmel-B-R1812-Map4-UNCLASSIFIED
consensus size: 1347bp; fragments: 22; full length: 2 ($\geq 1212.3\text{bp}$)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

