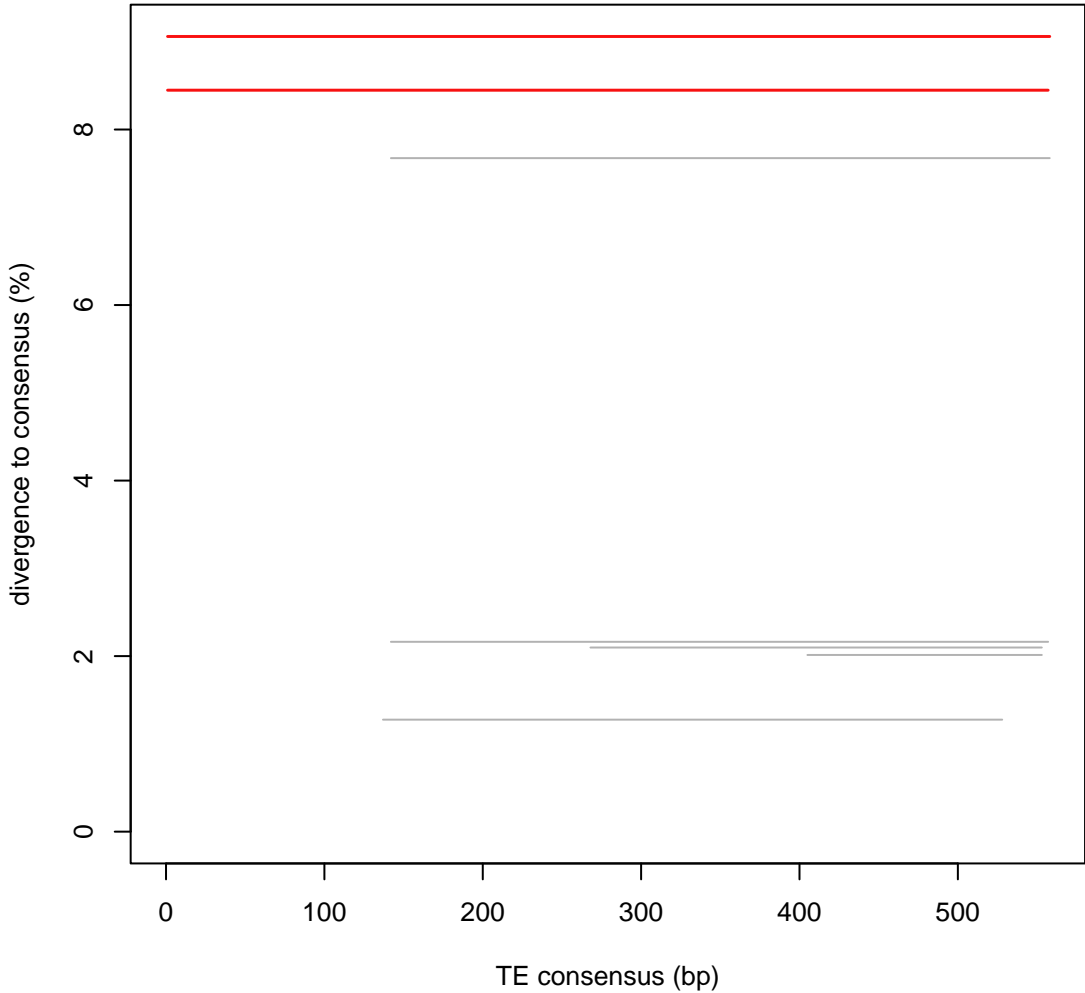
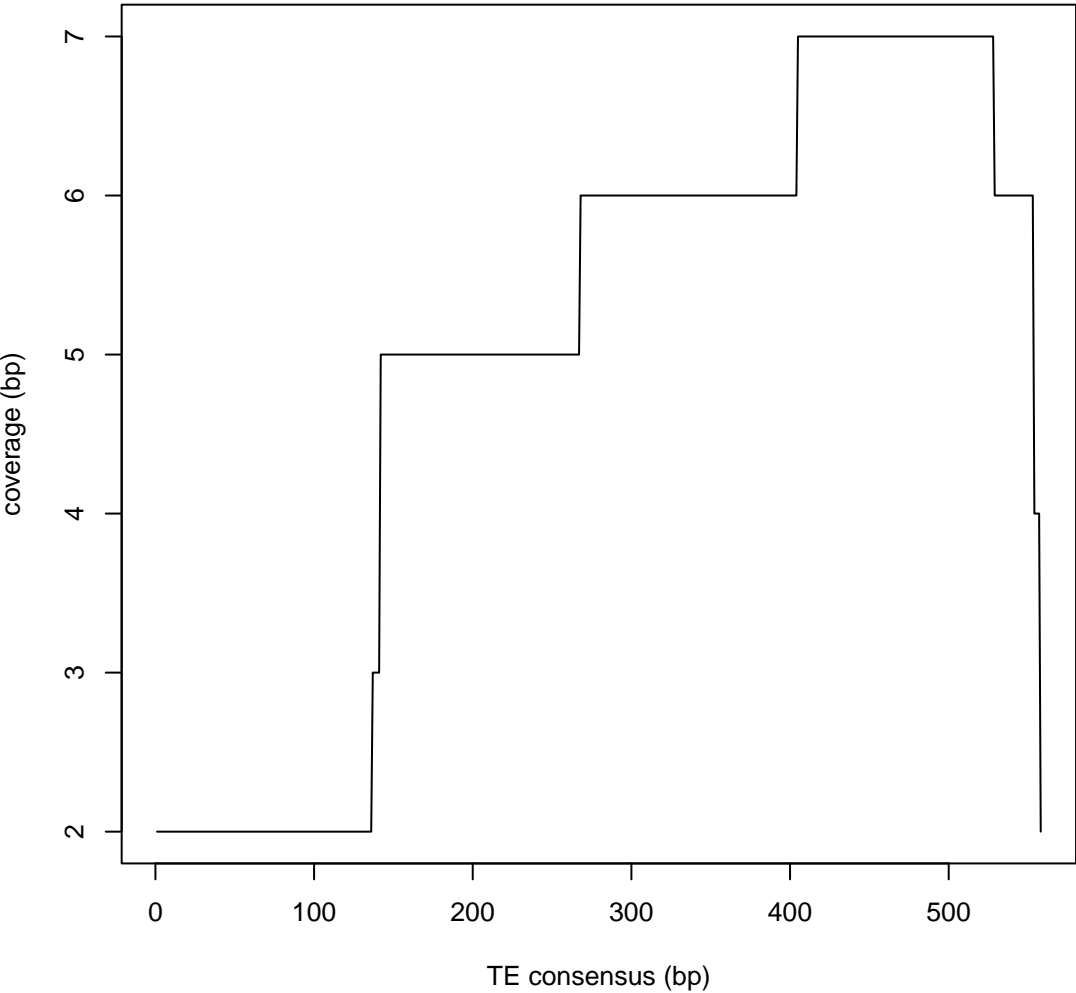


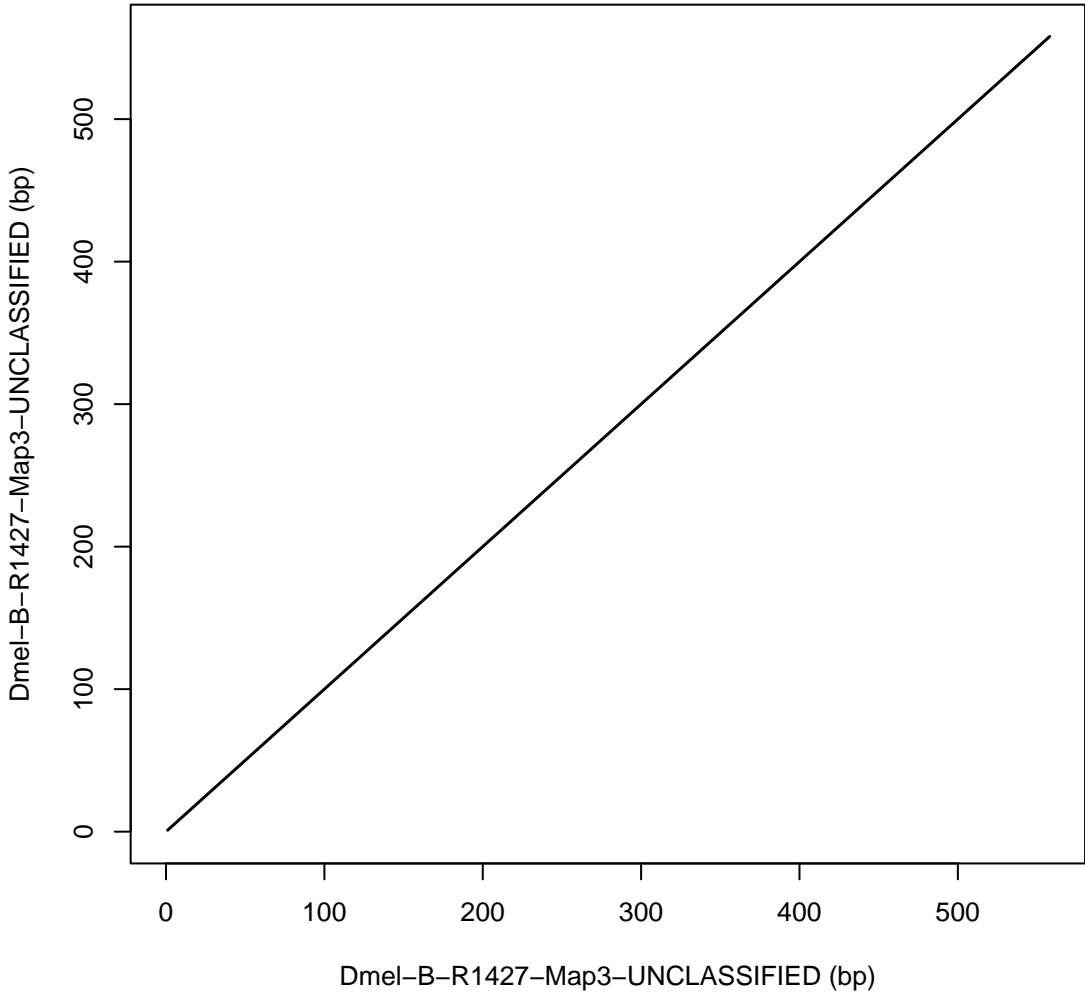
TE: Dmel-B-R1427-Map3-UNCLASSIFIED  
consensus size: 558bp; fragments: 7; full length: 2 (>=502.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

