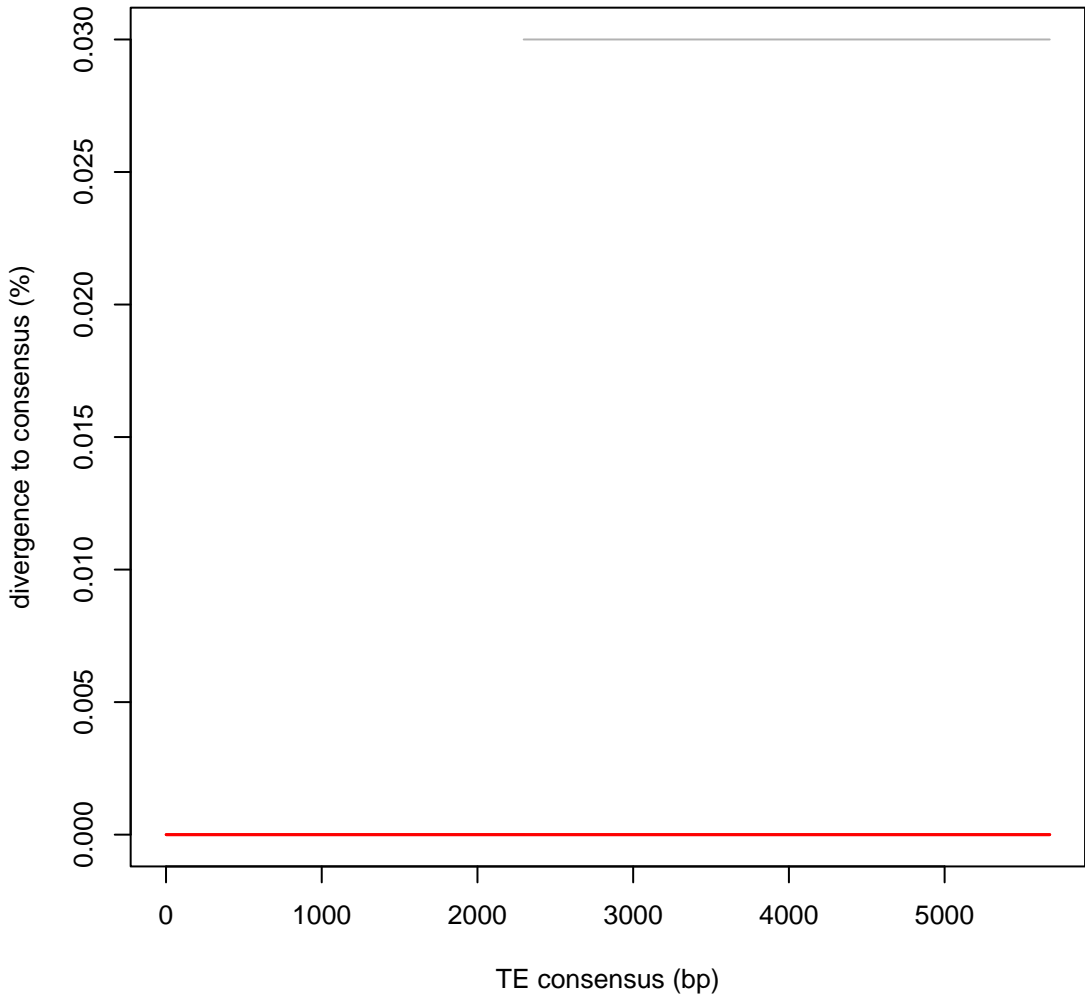
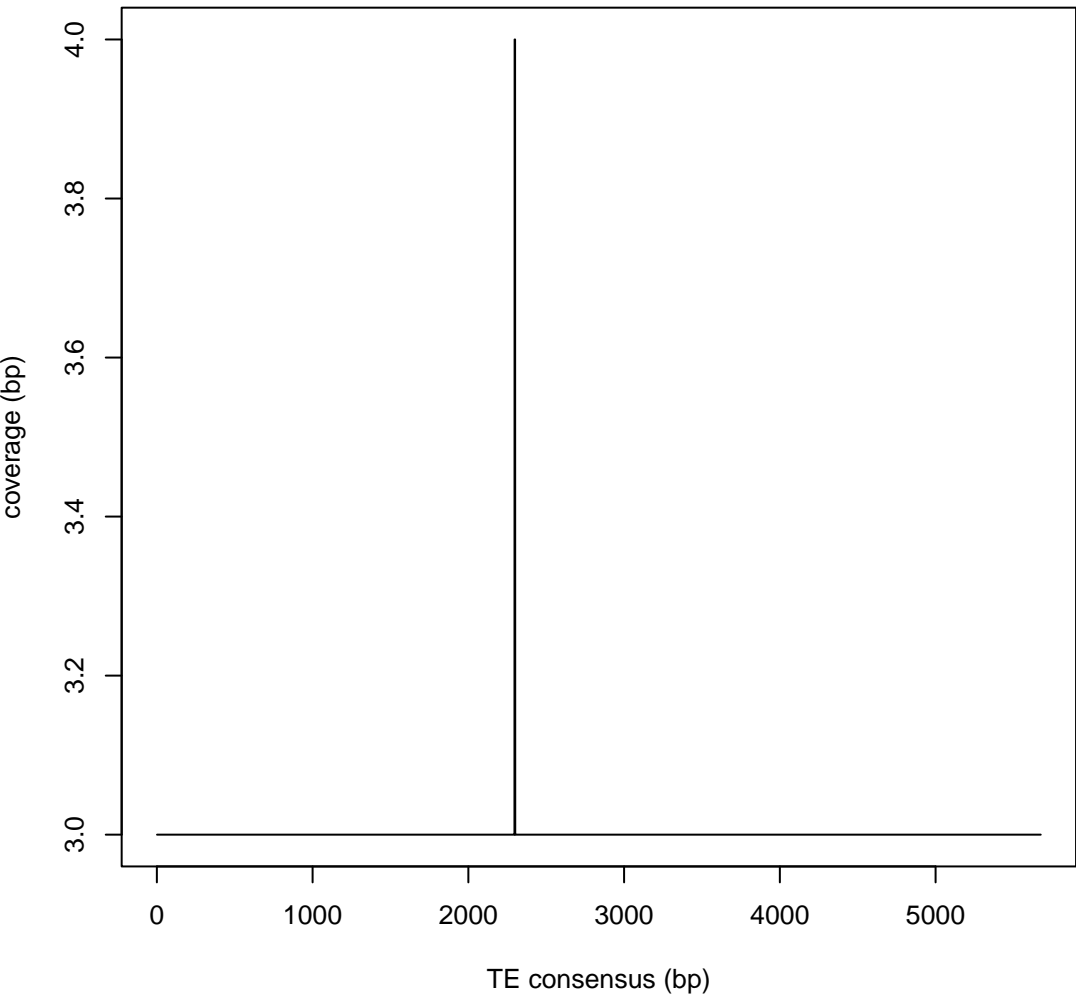


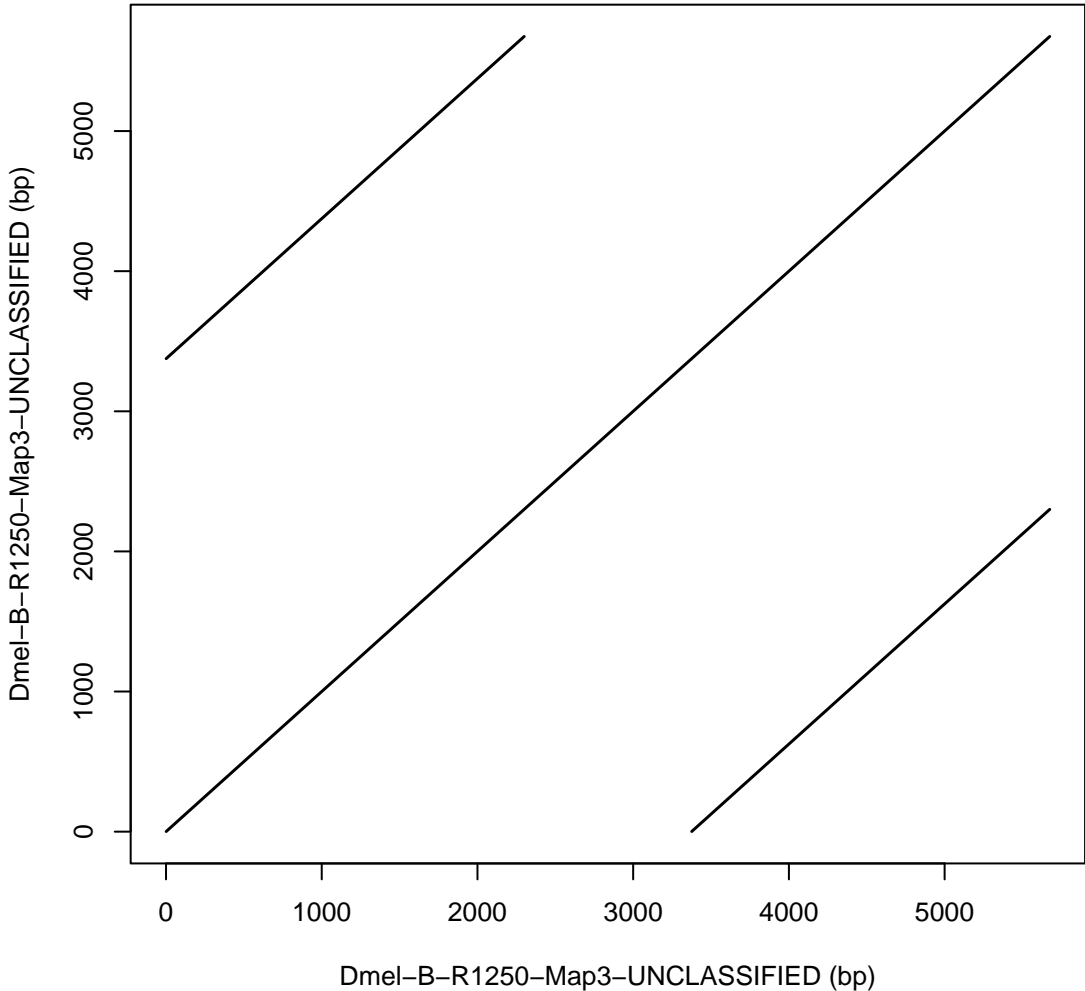
TE: Dmel-B-R1250-Map3-UNCLASSIFIED  
consensus size: 5675bp; fragments: 4; full length: 2 (>=5107.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

