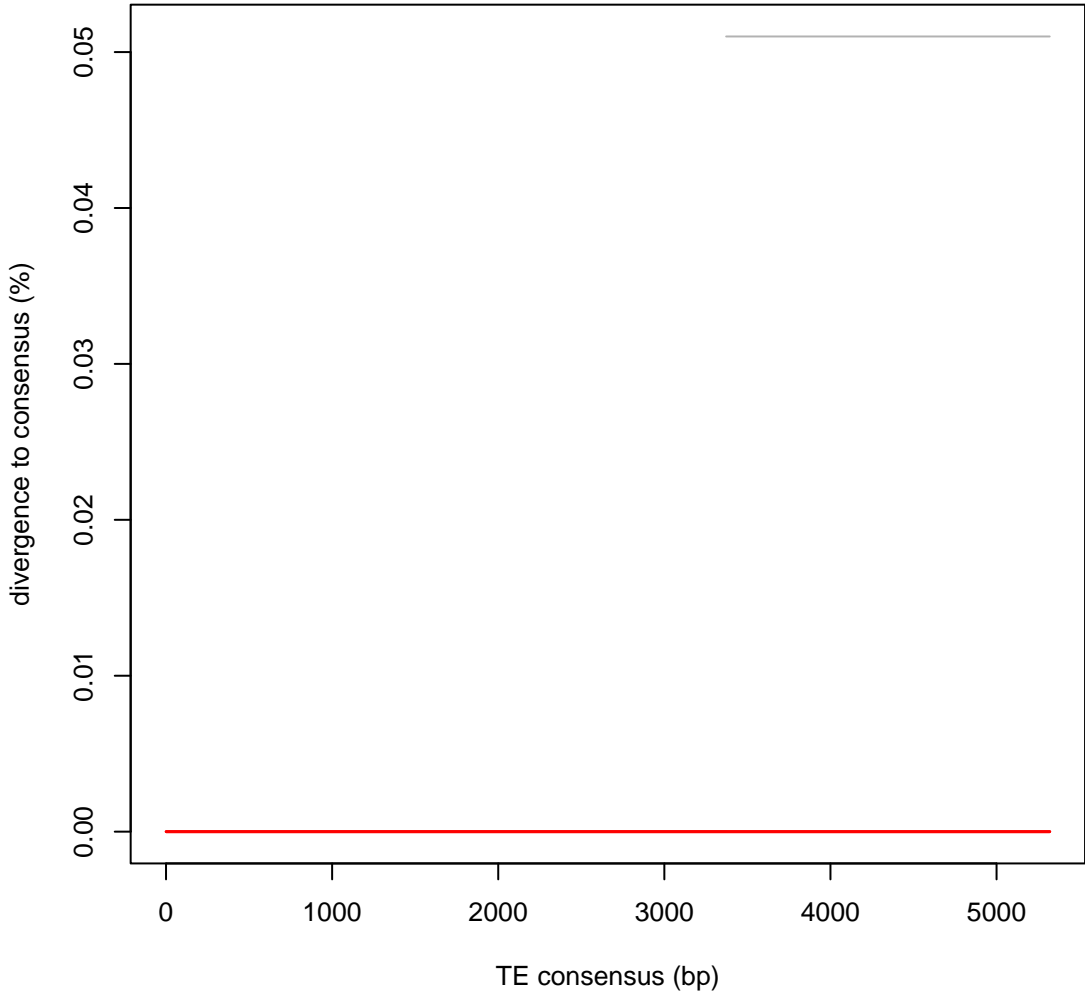
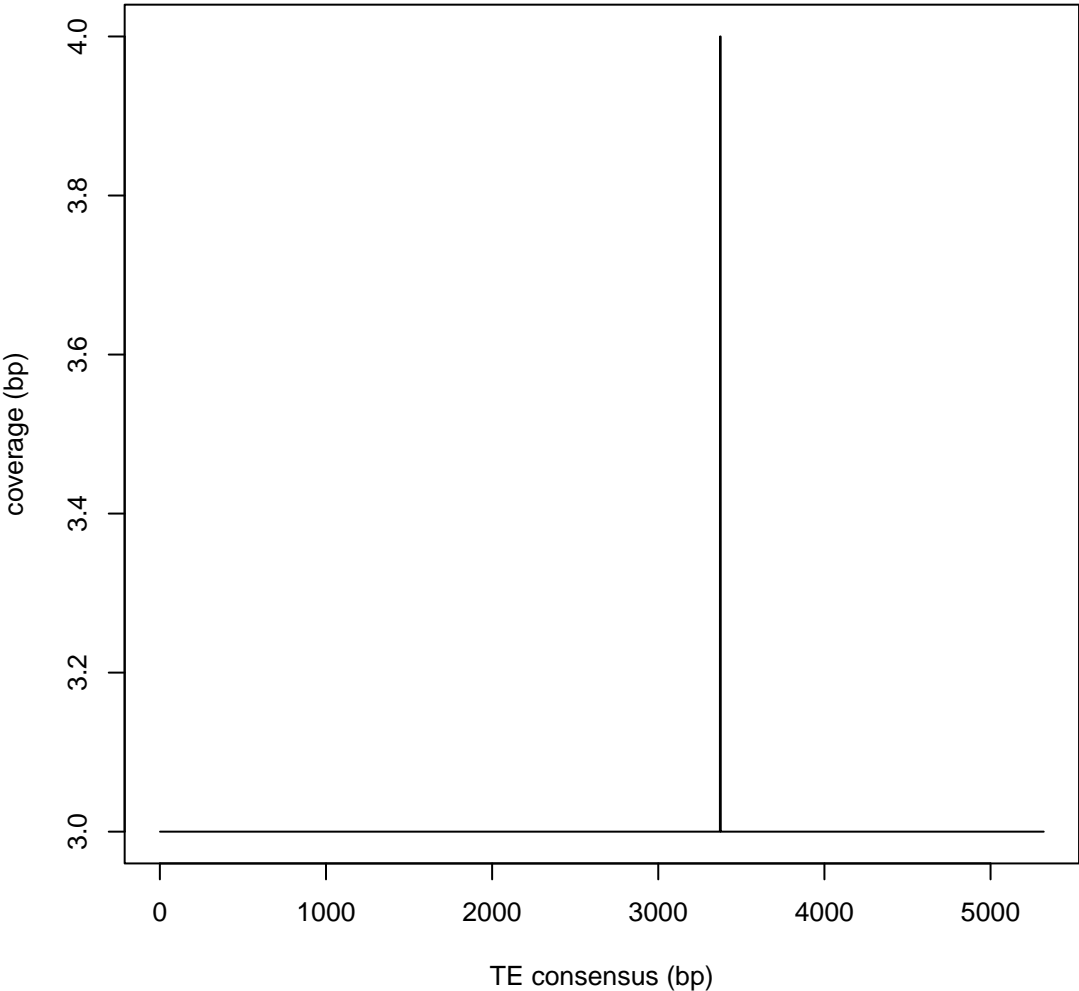


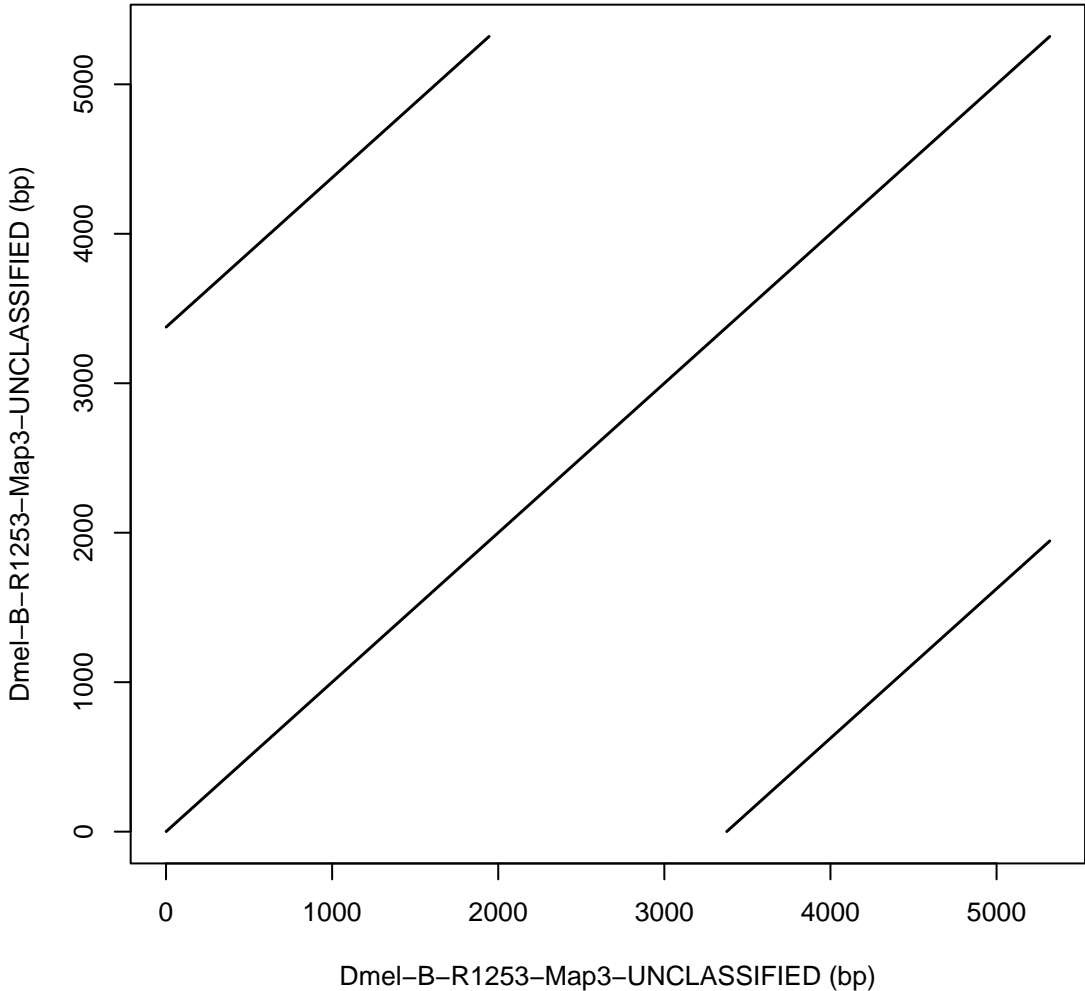
TE: Dmel-B-R1253-Map3-UNCLASSIFIED  
consensus size: 5320bp; fragments: 4; full length: 2 (>=4788bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

