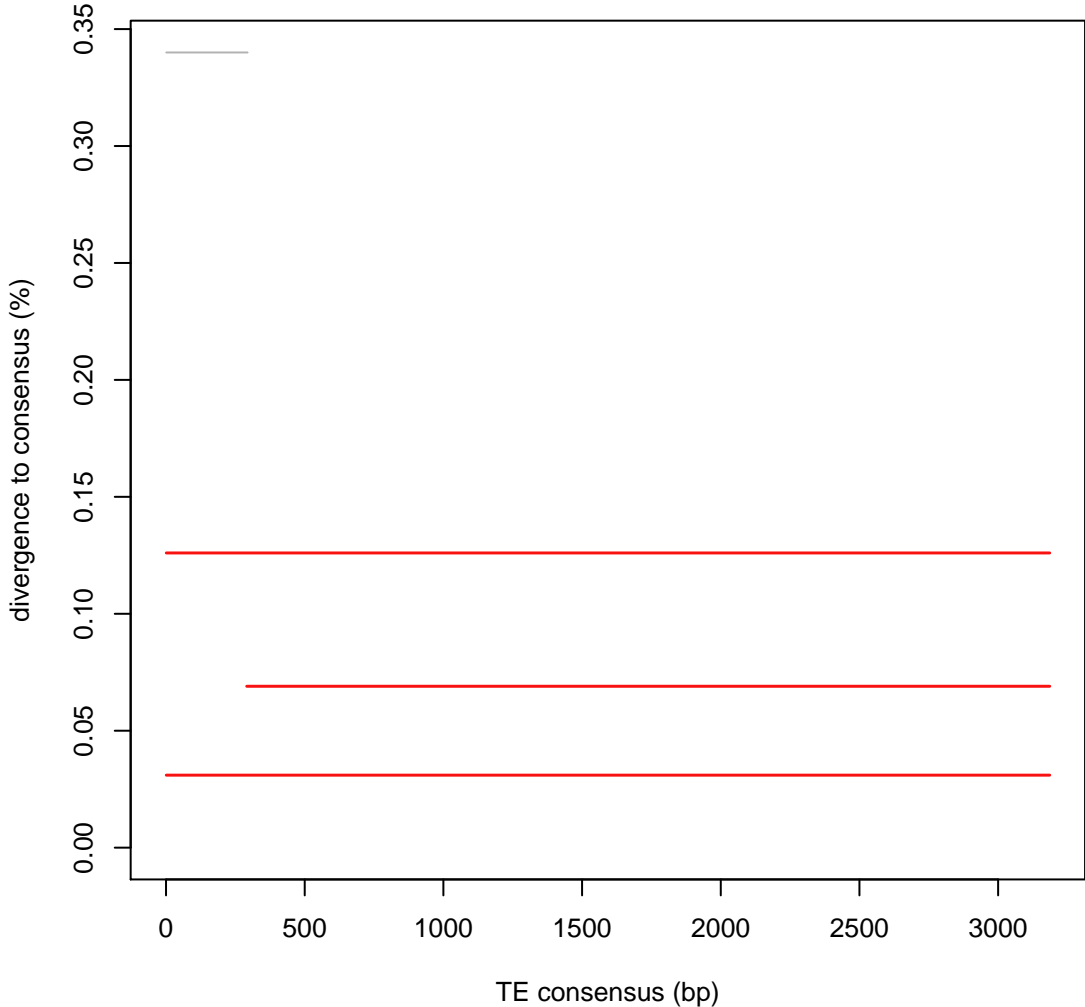
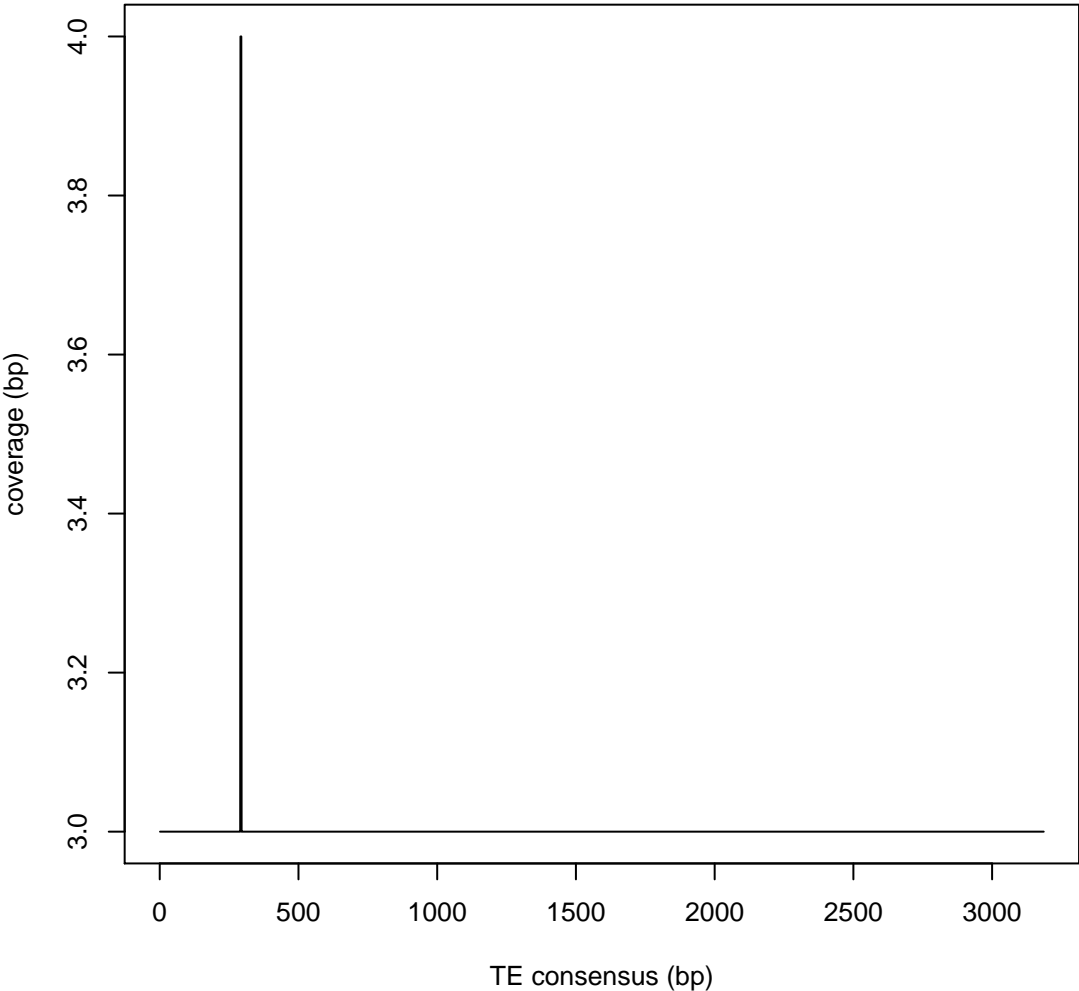


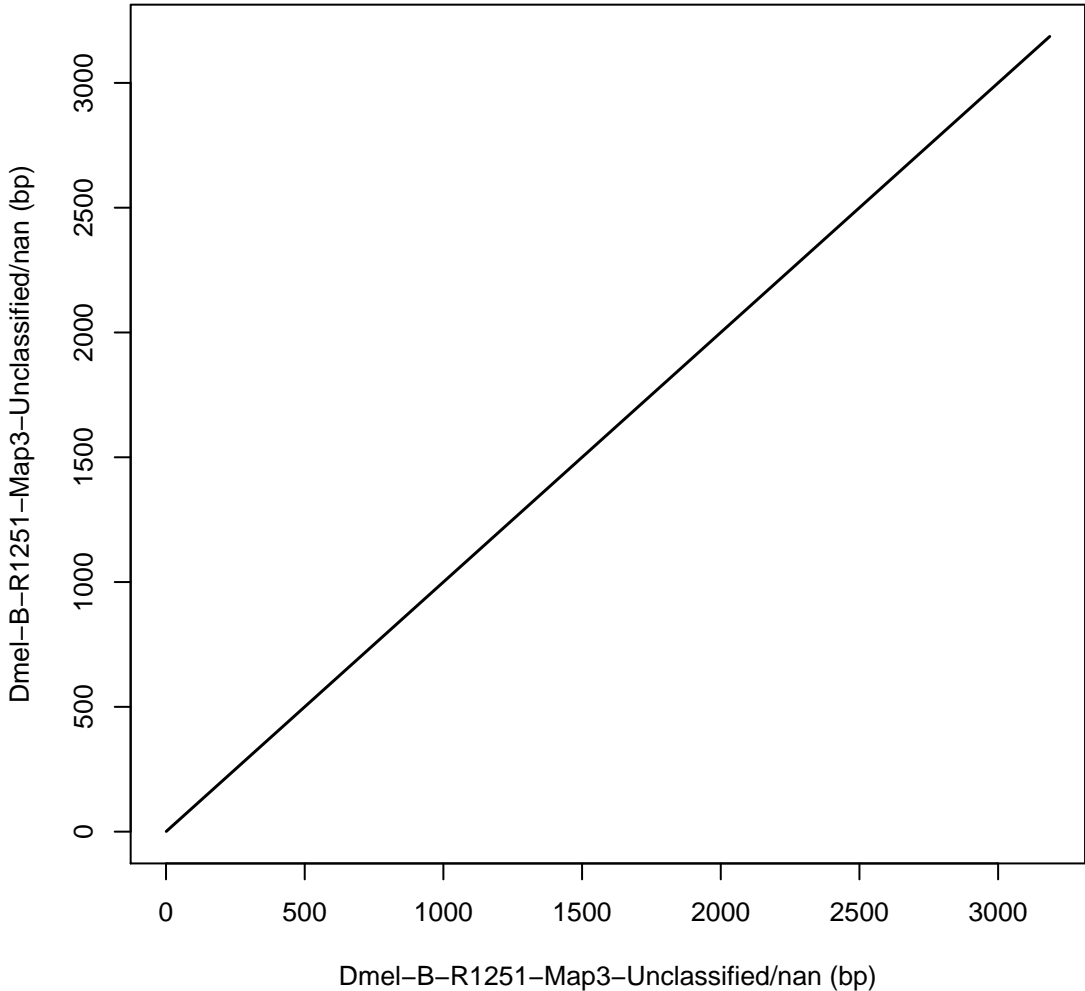
TE: Dmel-B-R1251-Map3-Unclassified/nan  
consensus size: 3186bp; fragments: 4; full length: 3 (>=2867.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

