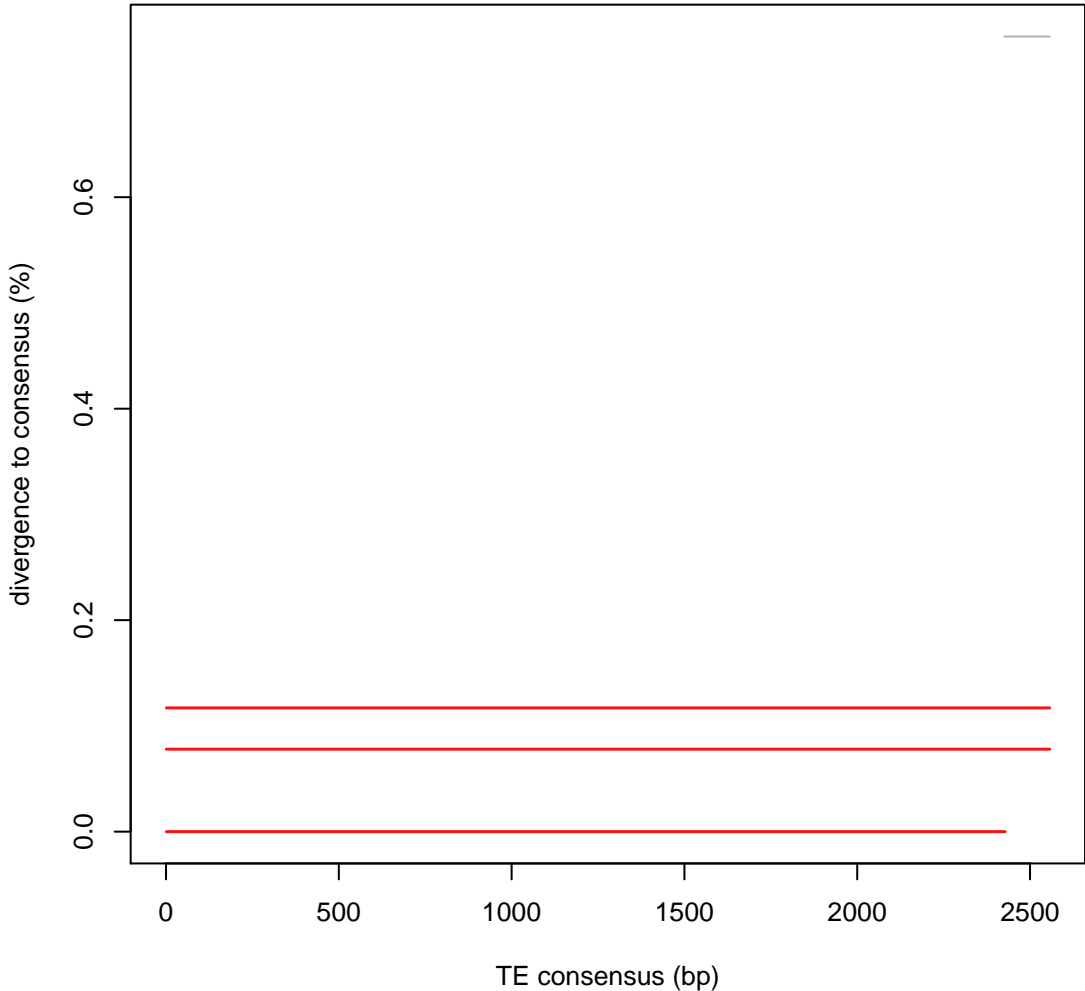
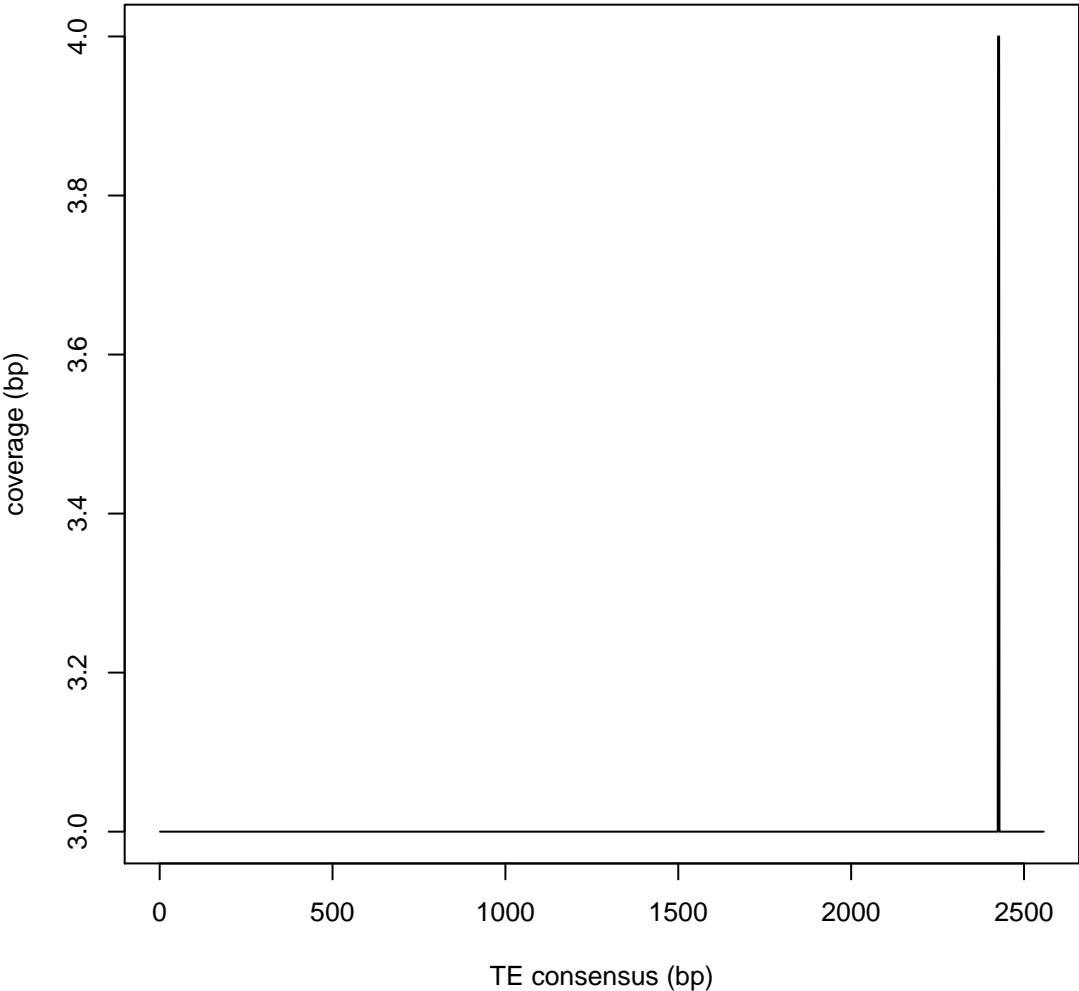


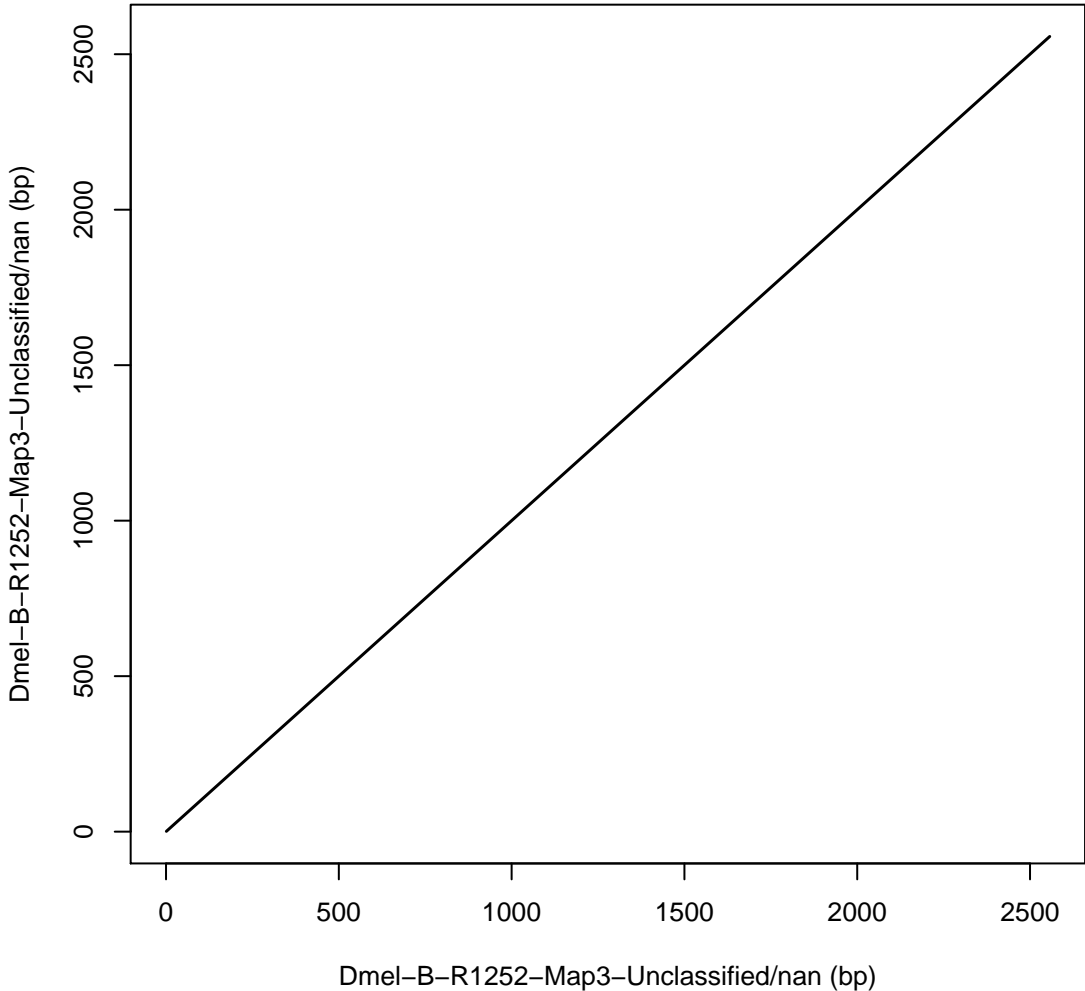
TE: Dmel-B-R1252-Map3-Unclassified/nan  
consensus size: 2557bp; fragments: 4; full length: 3 (>=2301.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

