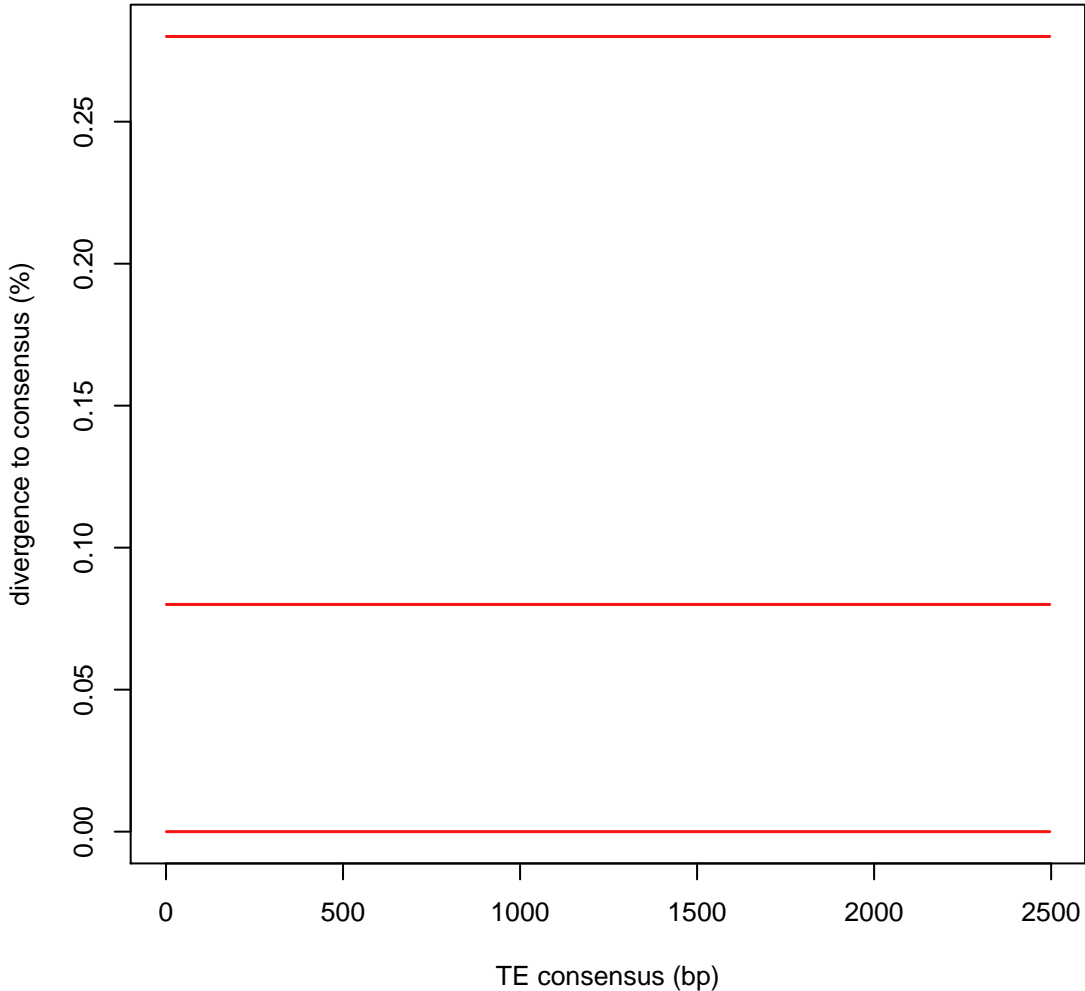
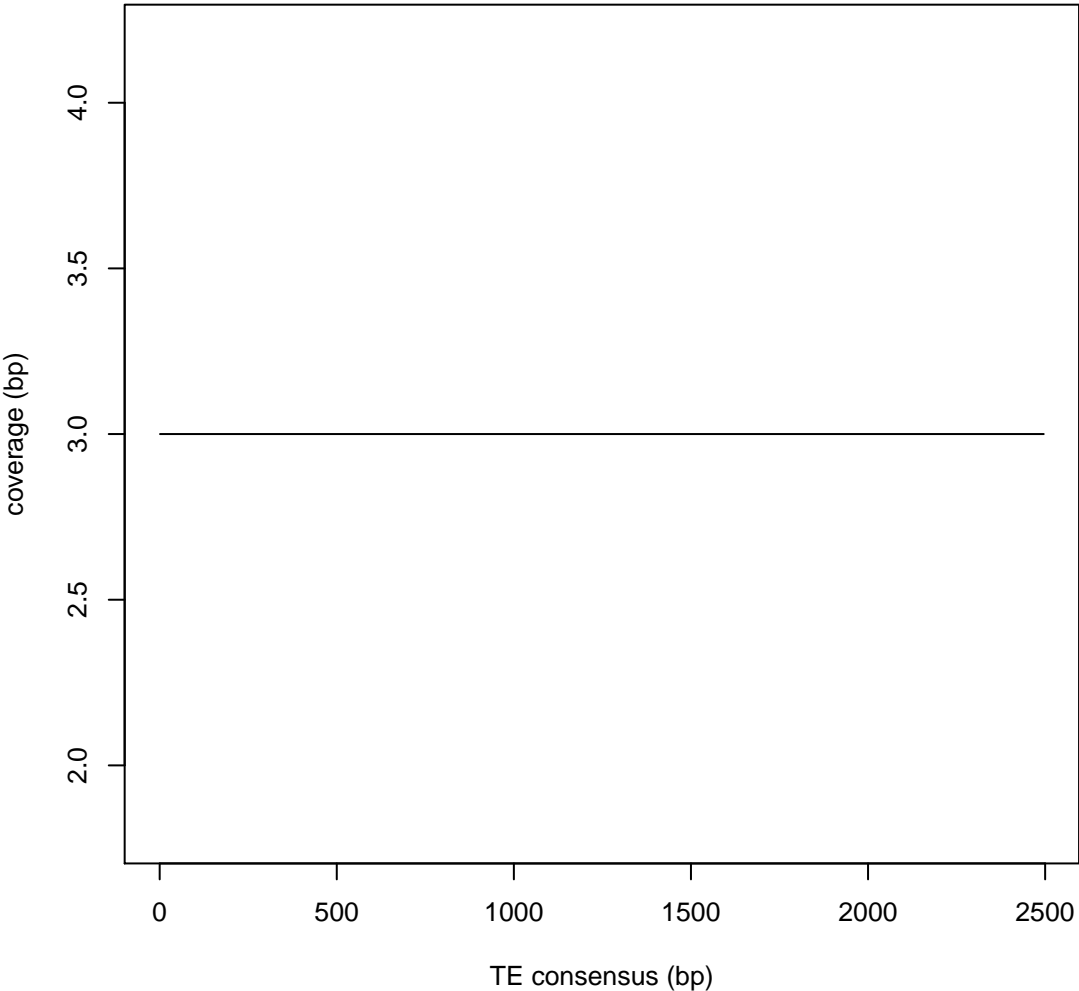


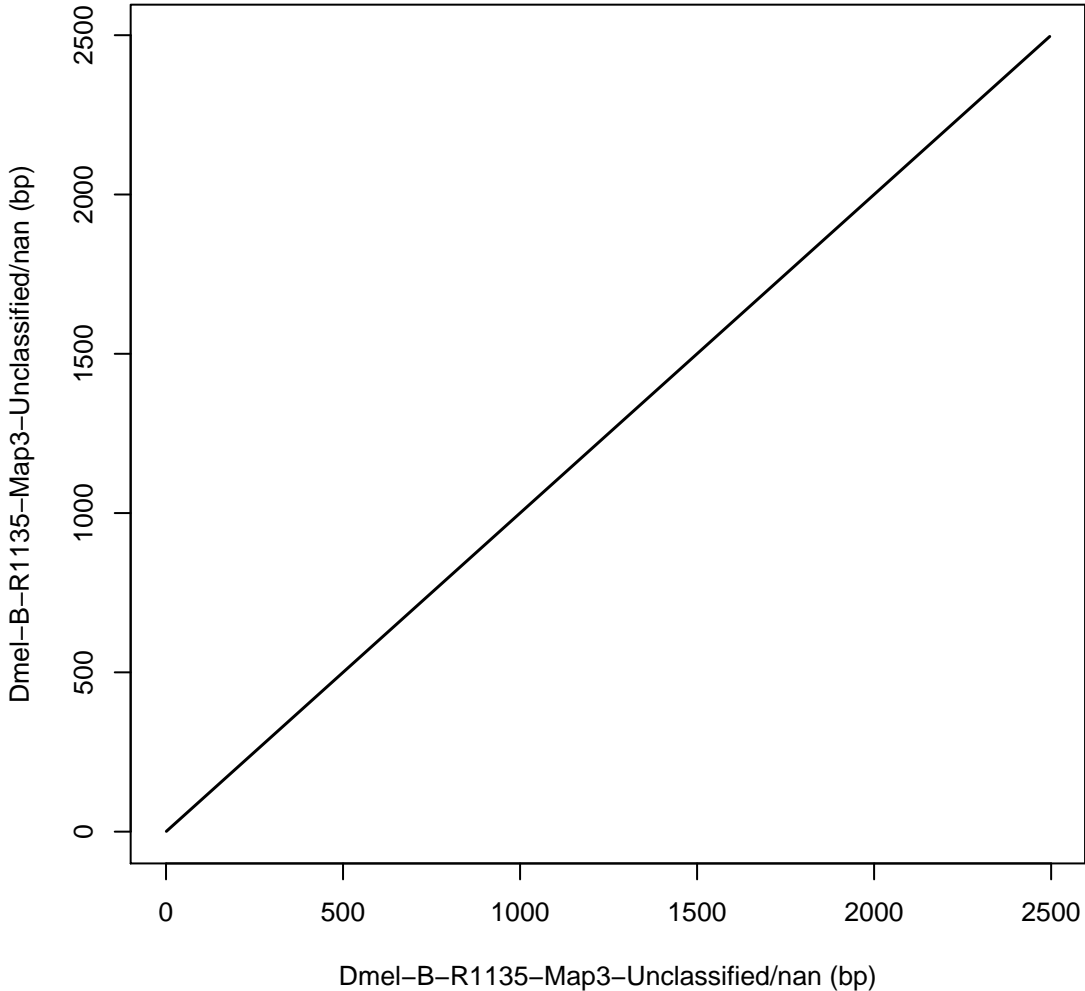
TE: Dmel-B-R1135-Map3-Unclassified/nan
consensus size: 2496bp; fragments: 3; full length: 3 (>=2246.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

