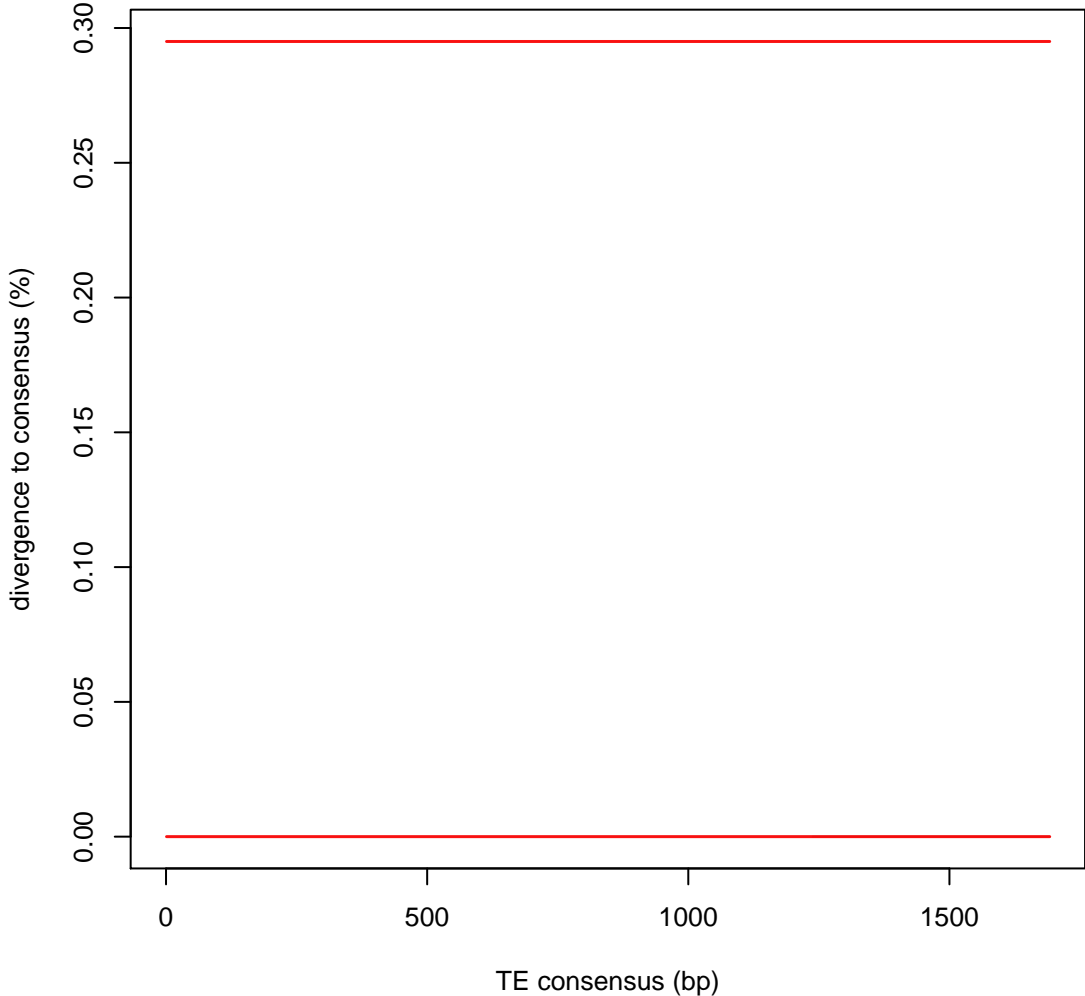
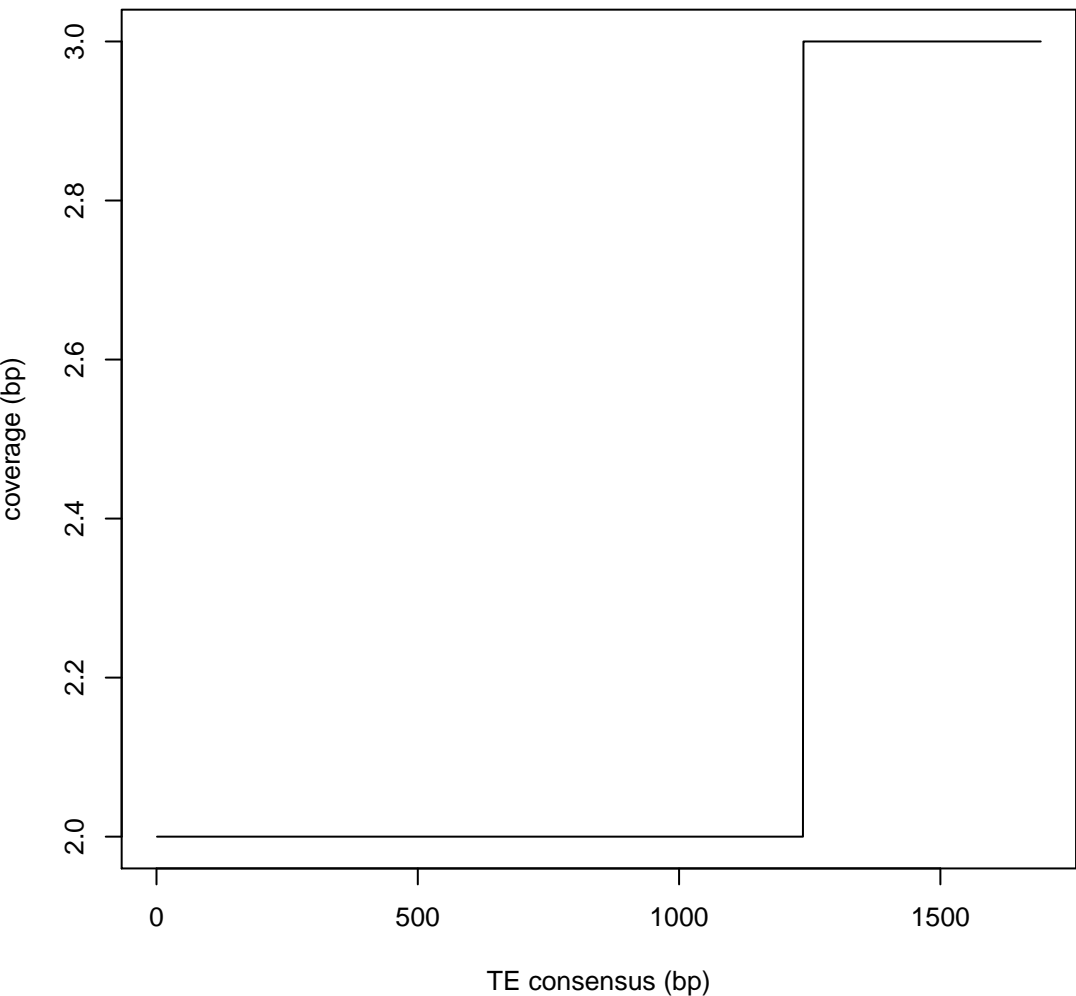


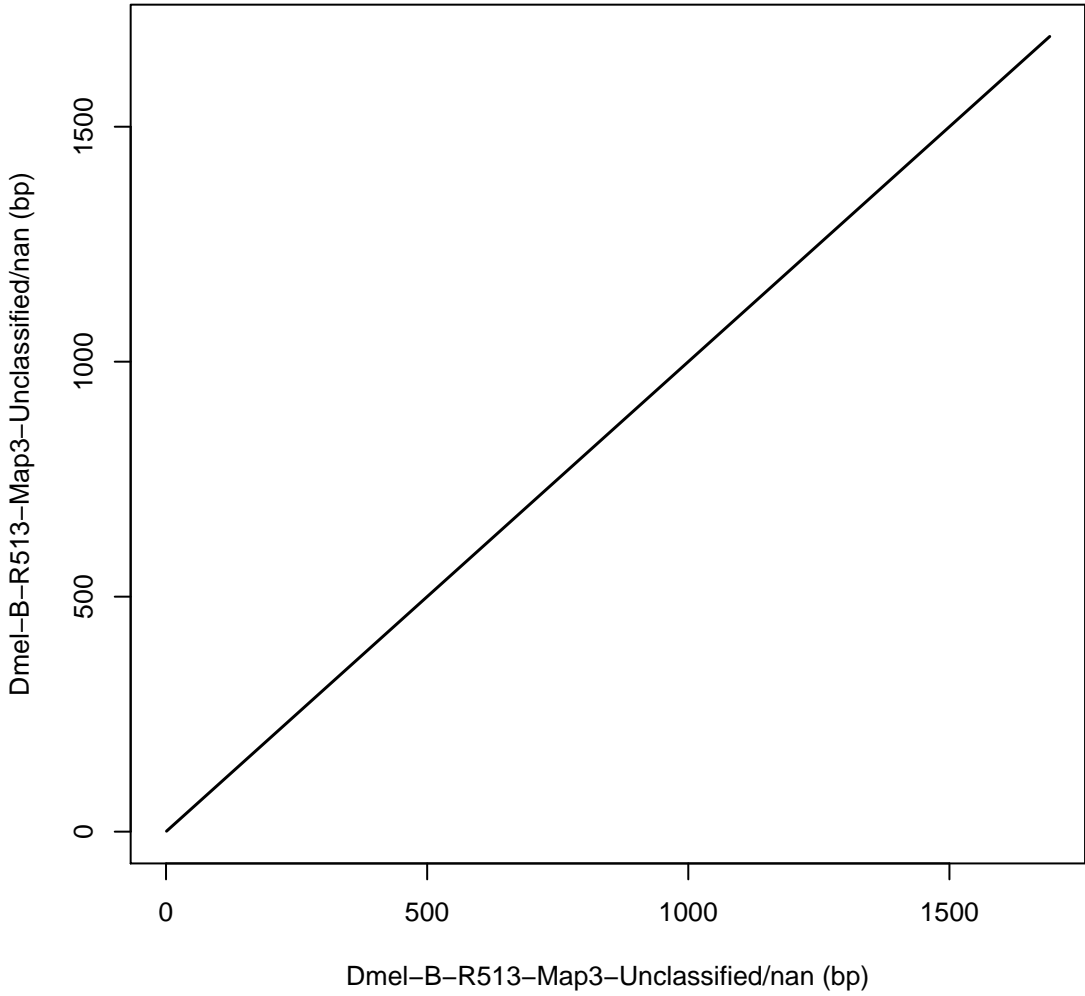
TE: Dmel-B-R513-Map3-Unclassified/nan
consensus size: 1692bp; fragments: 3; full length: 2 (≥ 1522.8 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

