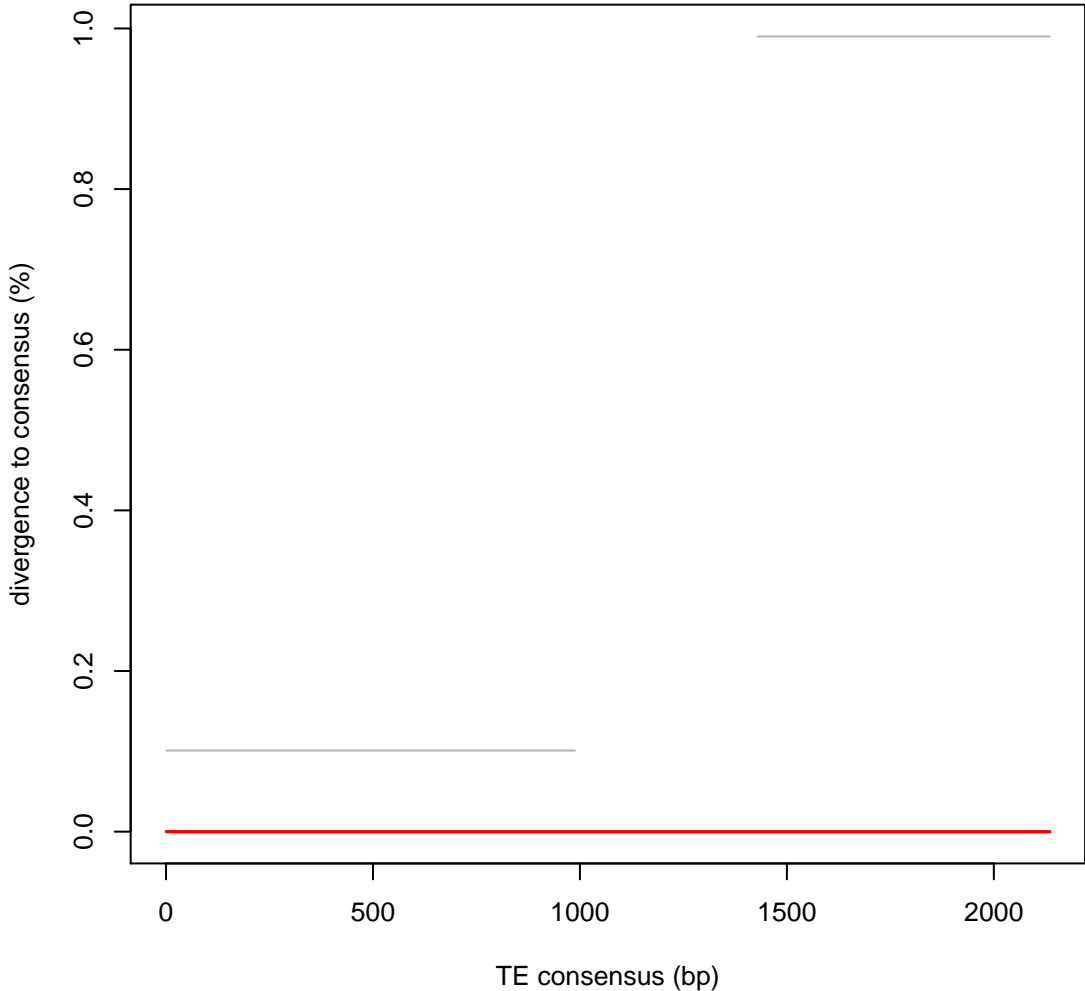
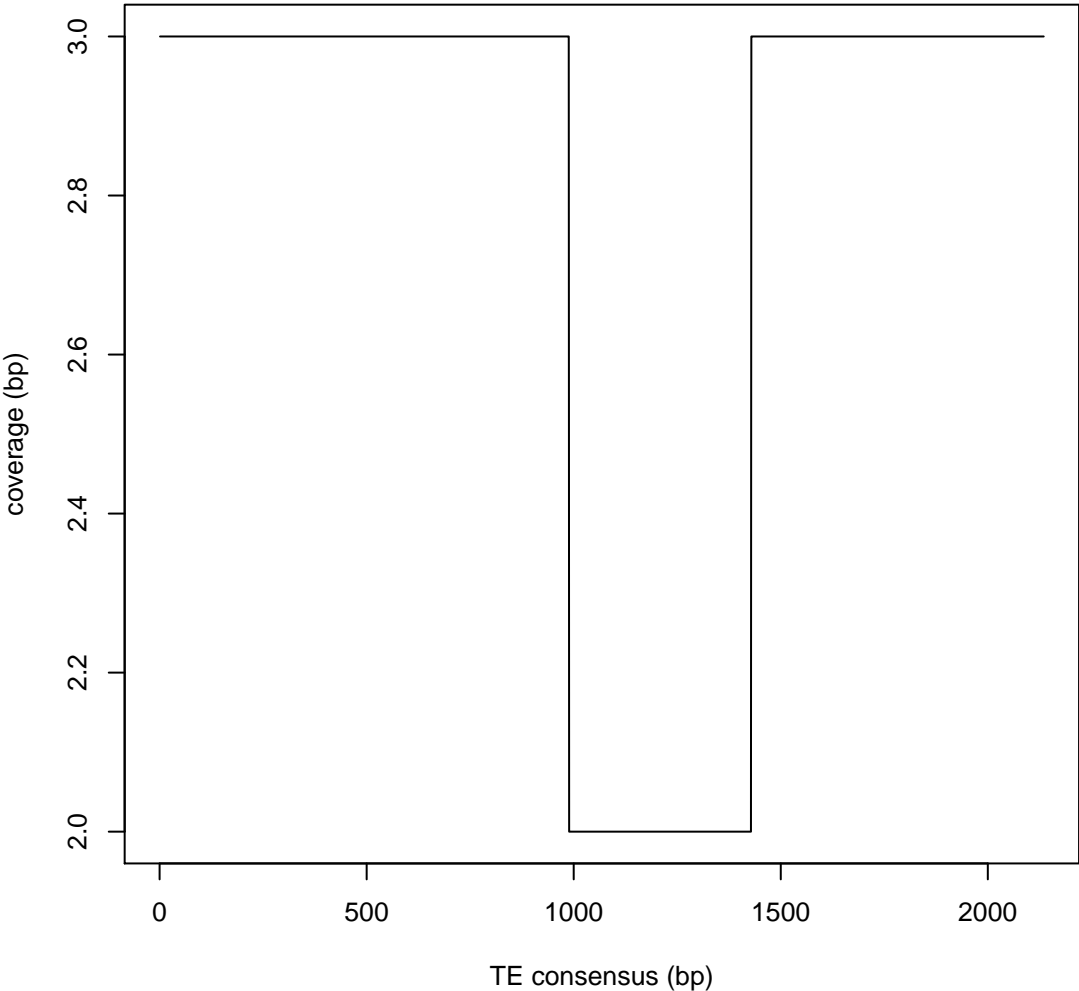


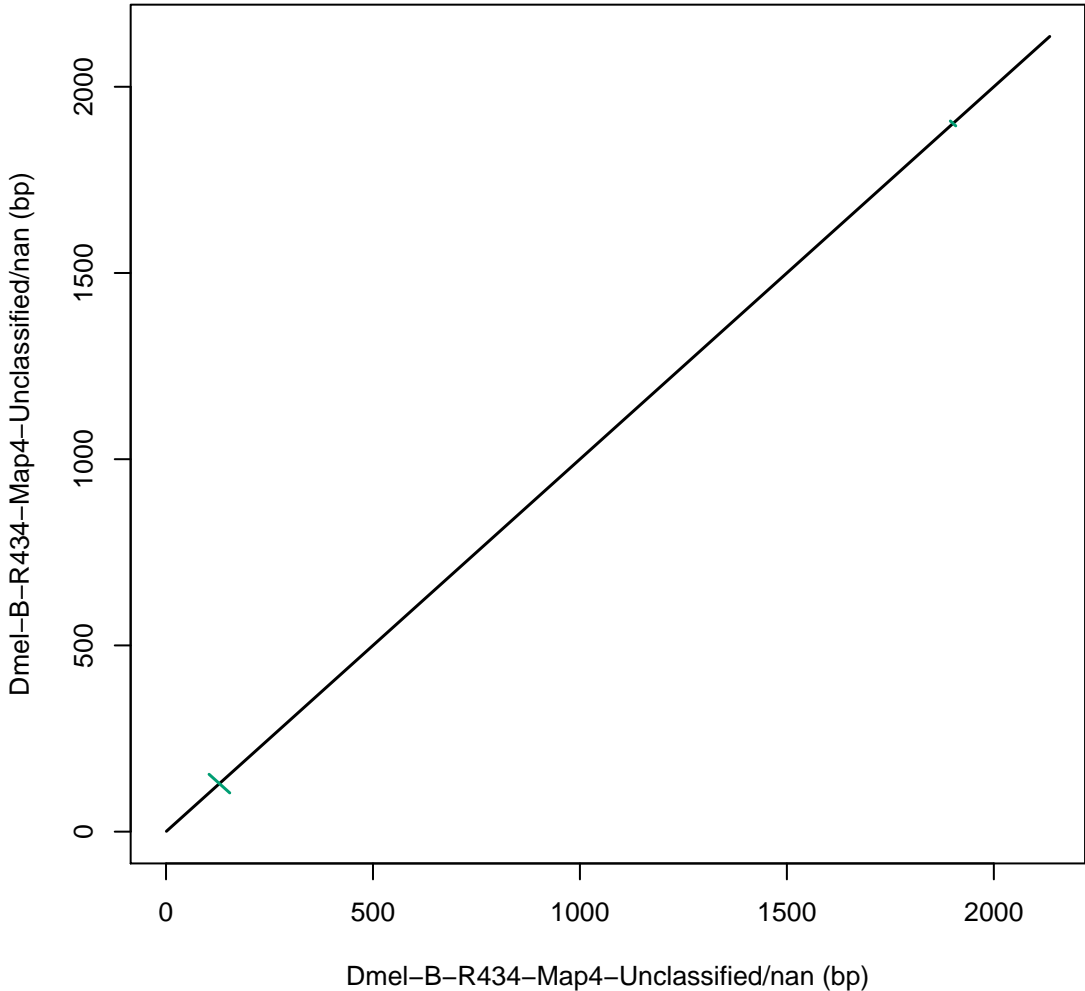
TE: Dmel-B-R434-Map4-Unclassified/nan
consensus size: 2135bp; fragments: 4; full length: 2 (≥ 1921.5 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

