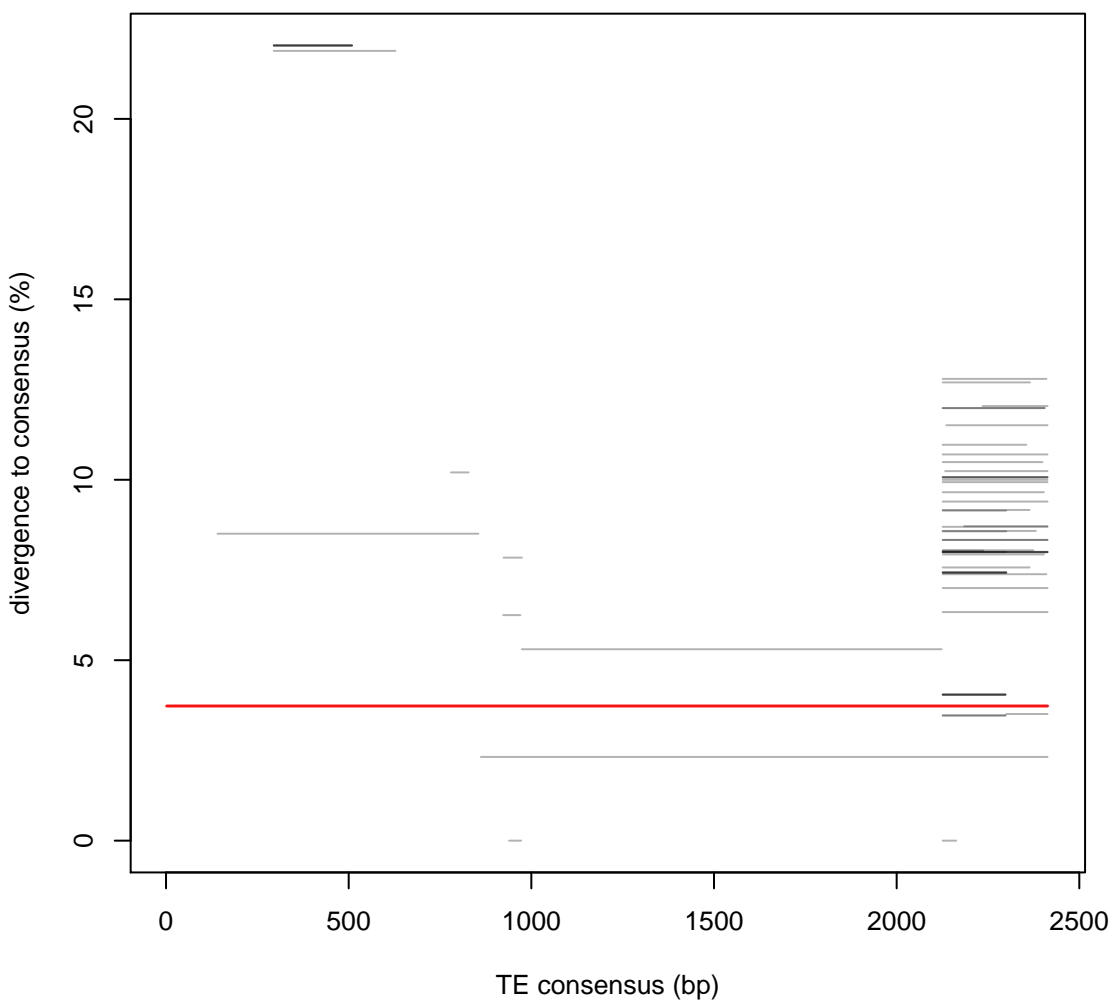
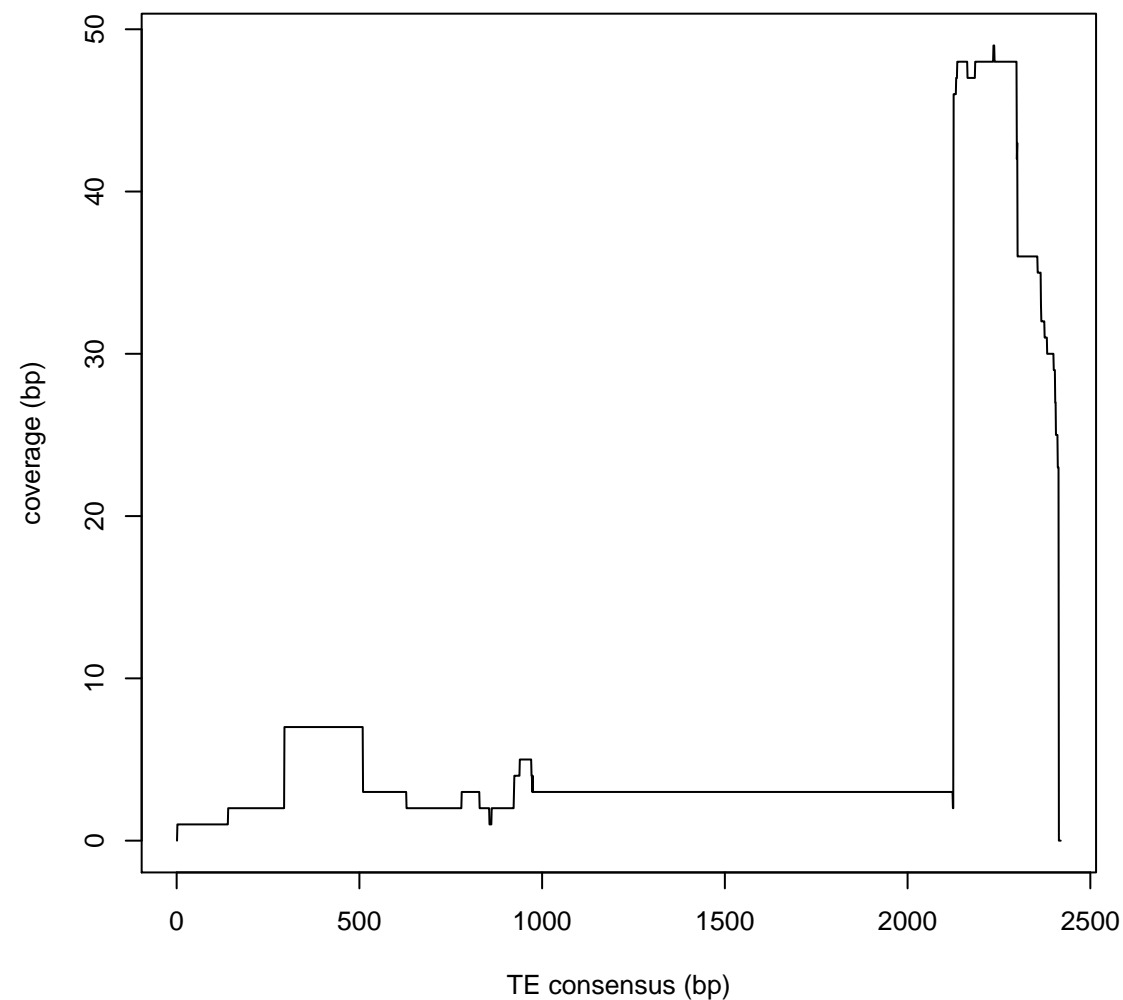


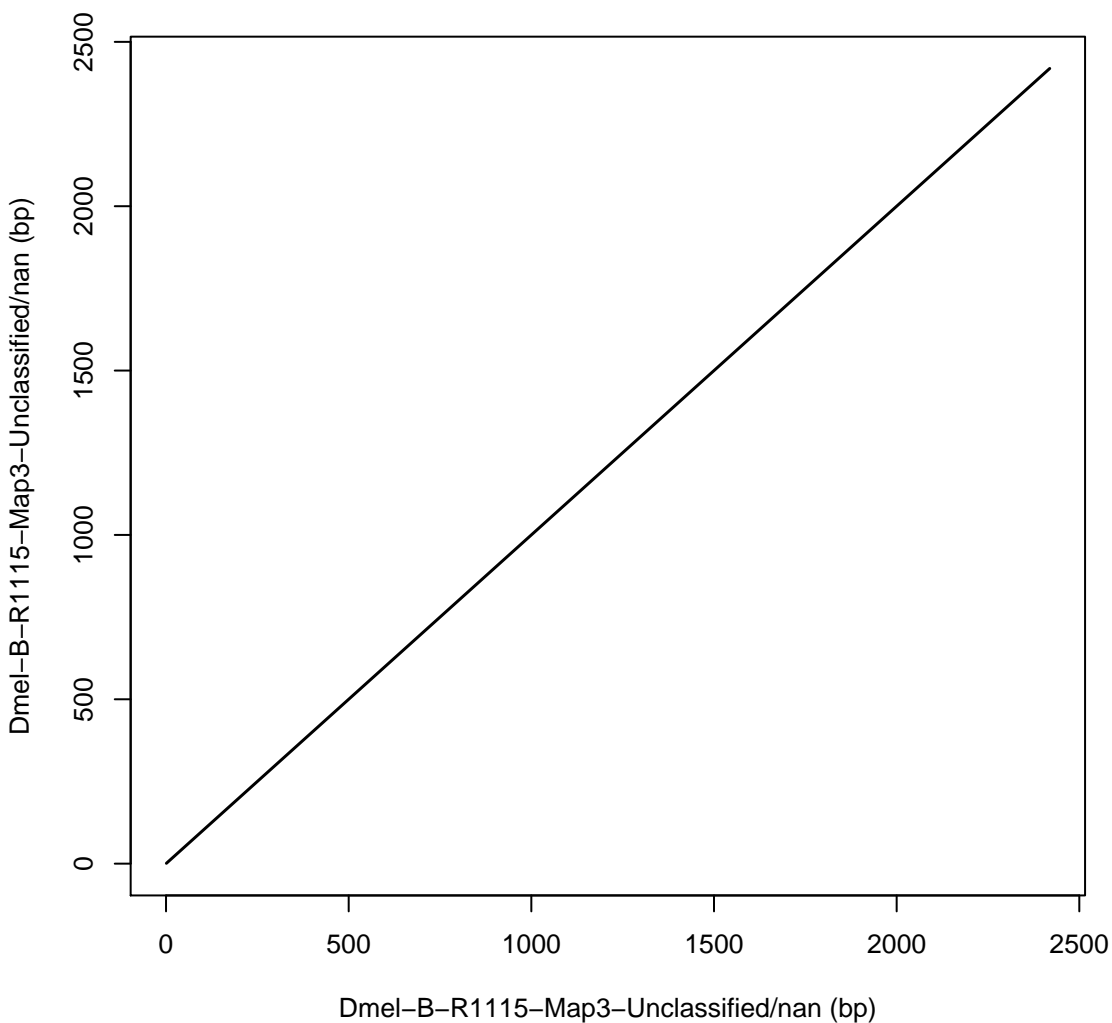
TE: Dmel-B-R1115-Map3-Unclassified/nan
consensus size: 2419bp; fragments: 62; full length: 1 (≥ 2177.1 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

