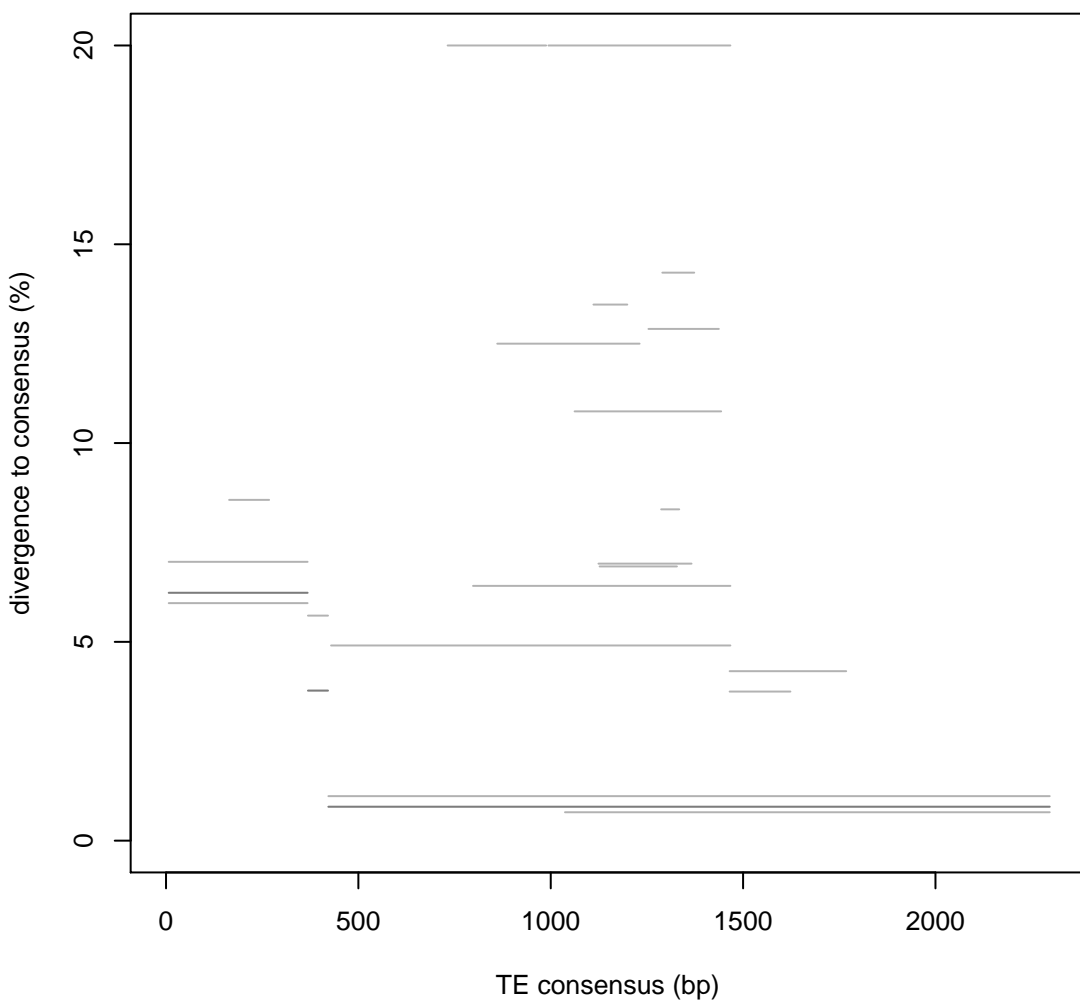
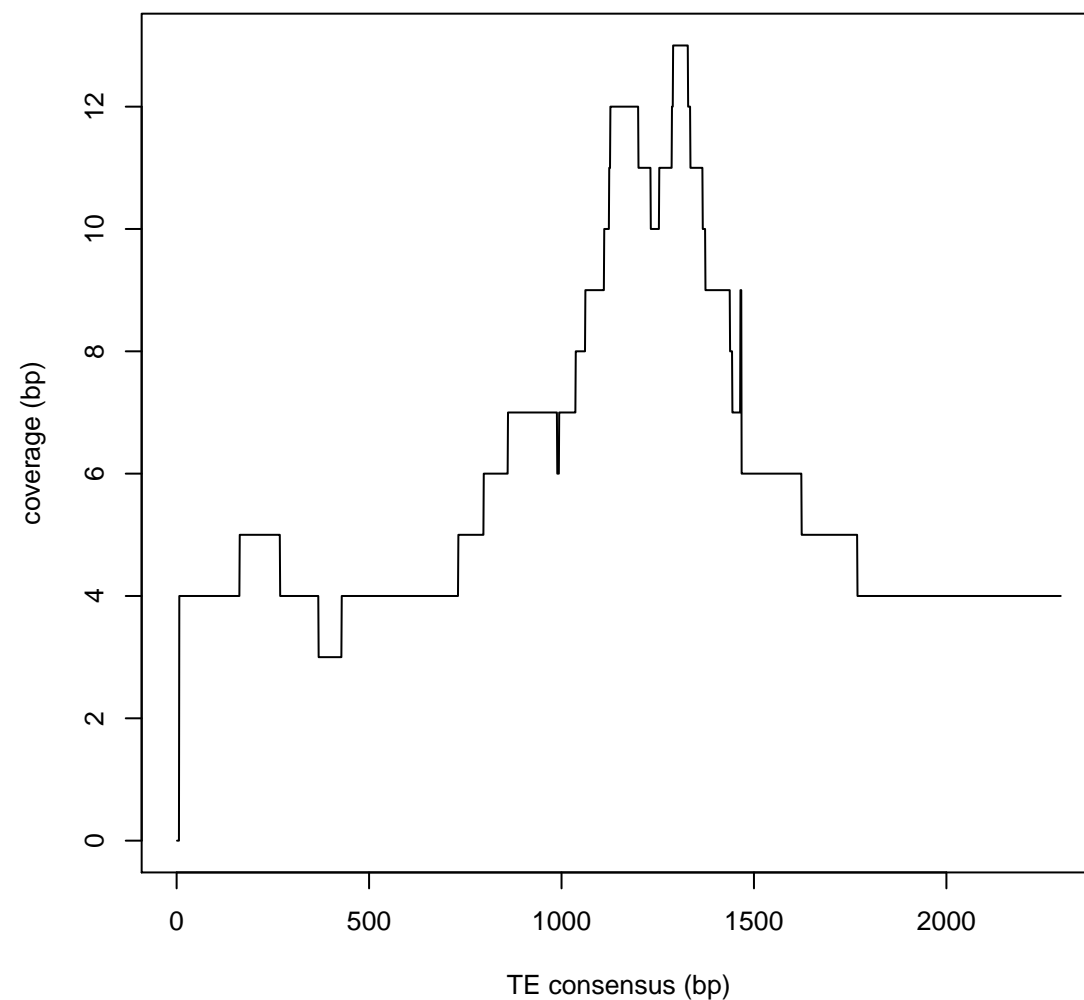


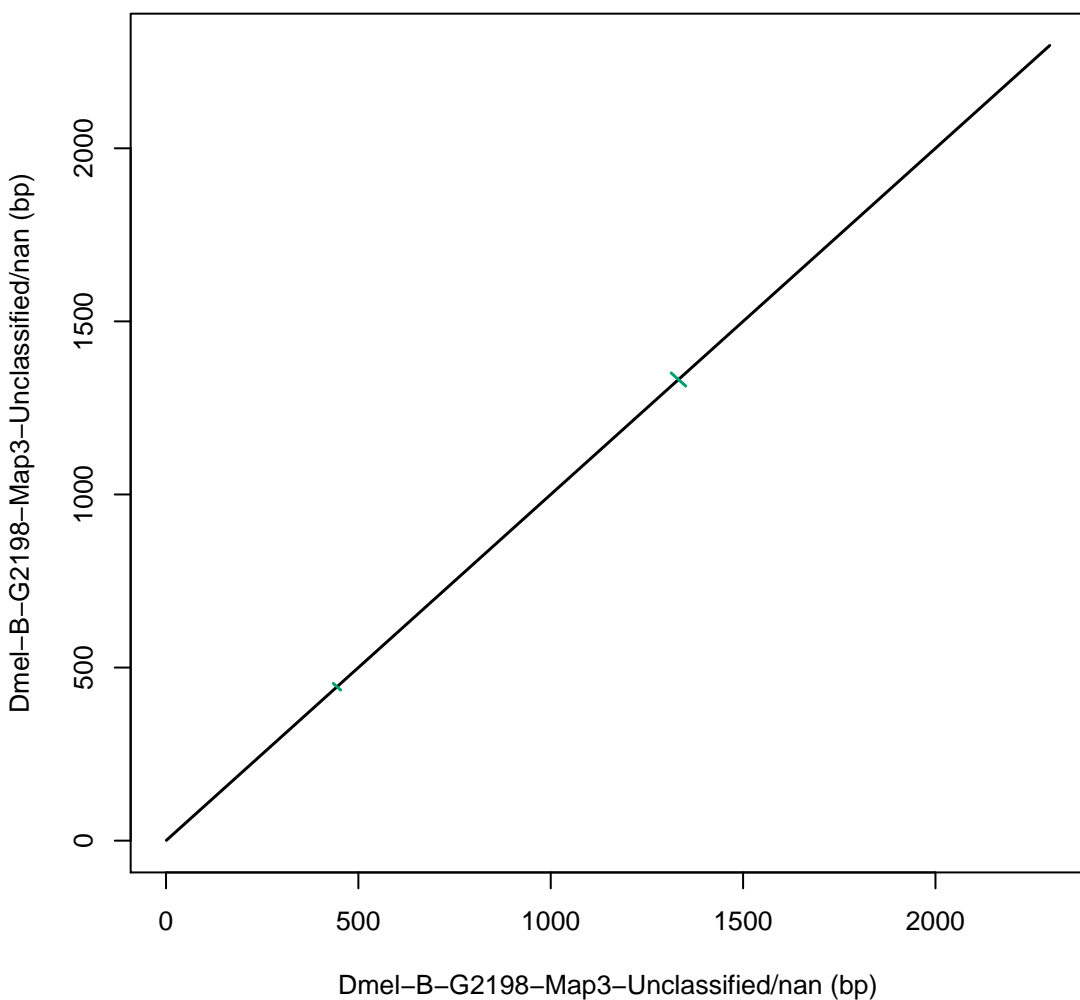
TE: Dmel-B-G2198-Map3-Unclassified/nan
consensus size: 2297bp; fragments: 26; full length: 0 (≥ 2067.3 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

