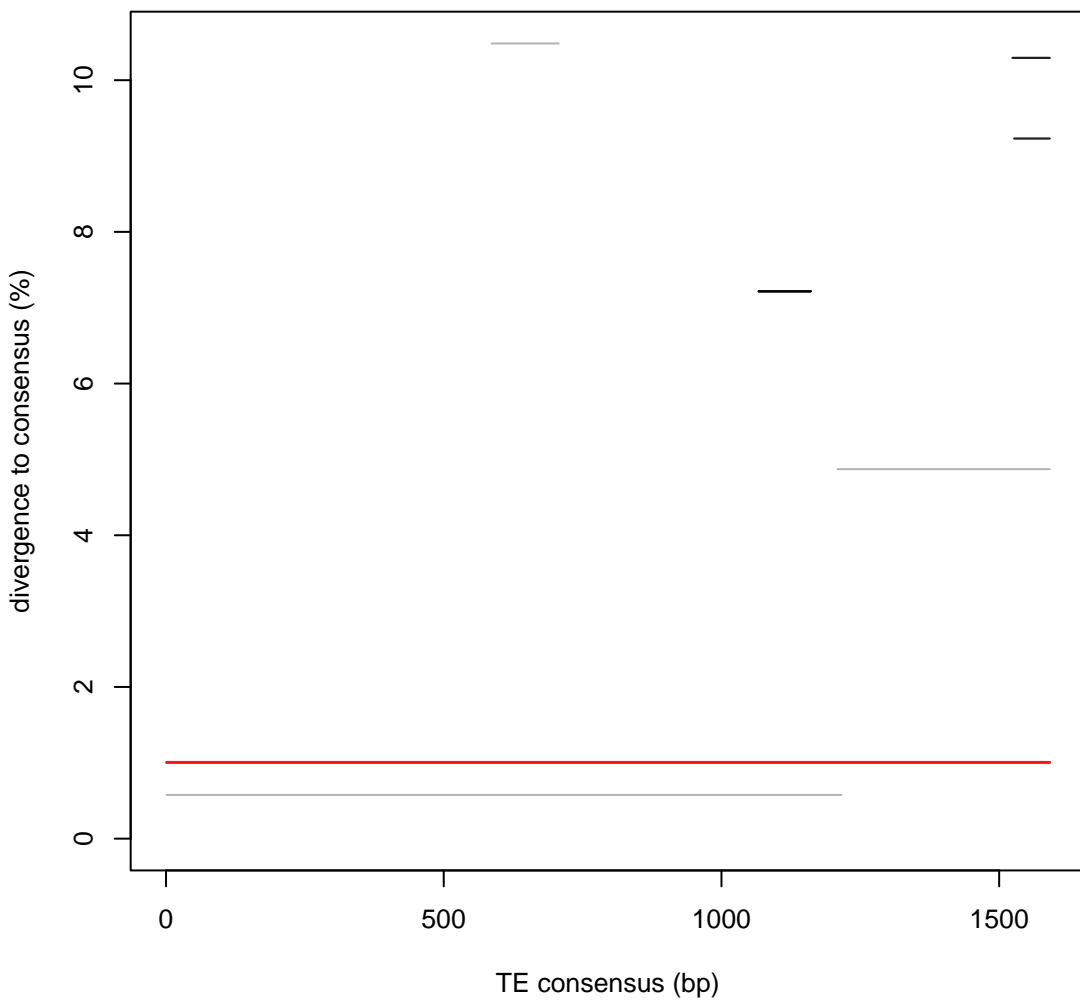
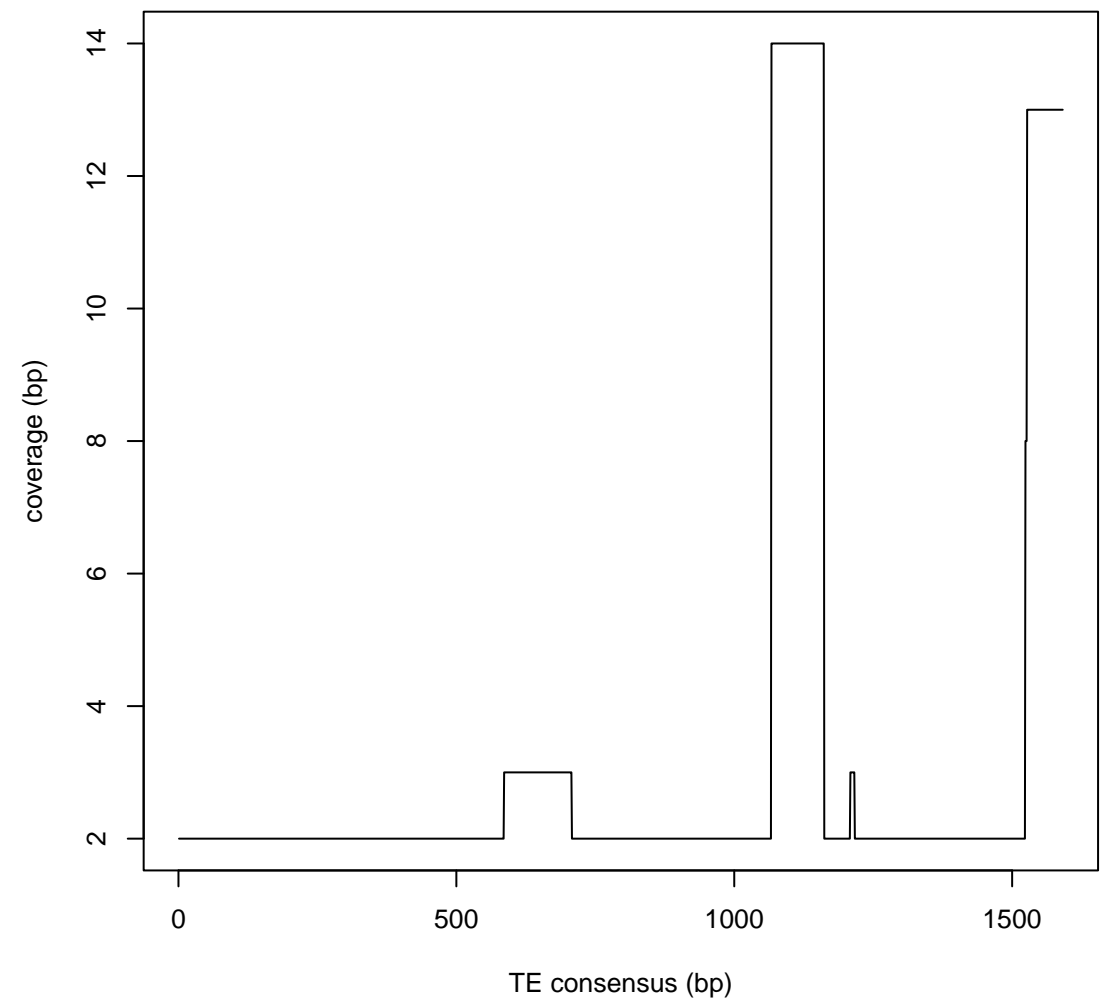


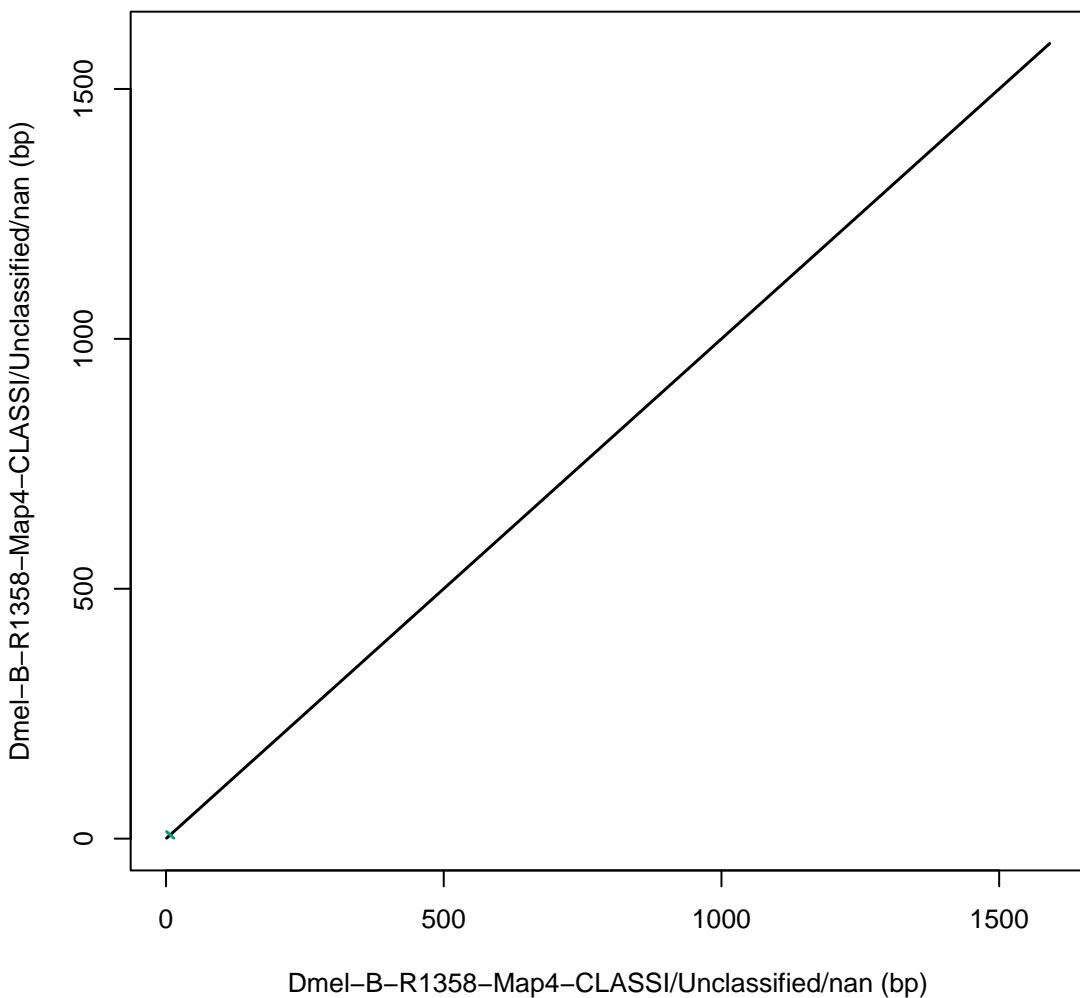
TE: Dmel-B-R1358-Map4-CLASSI/Unclassified/nan
consensus size: 1591bp; fragments: 27; full length: 1 (≥ 1431.9 bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

