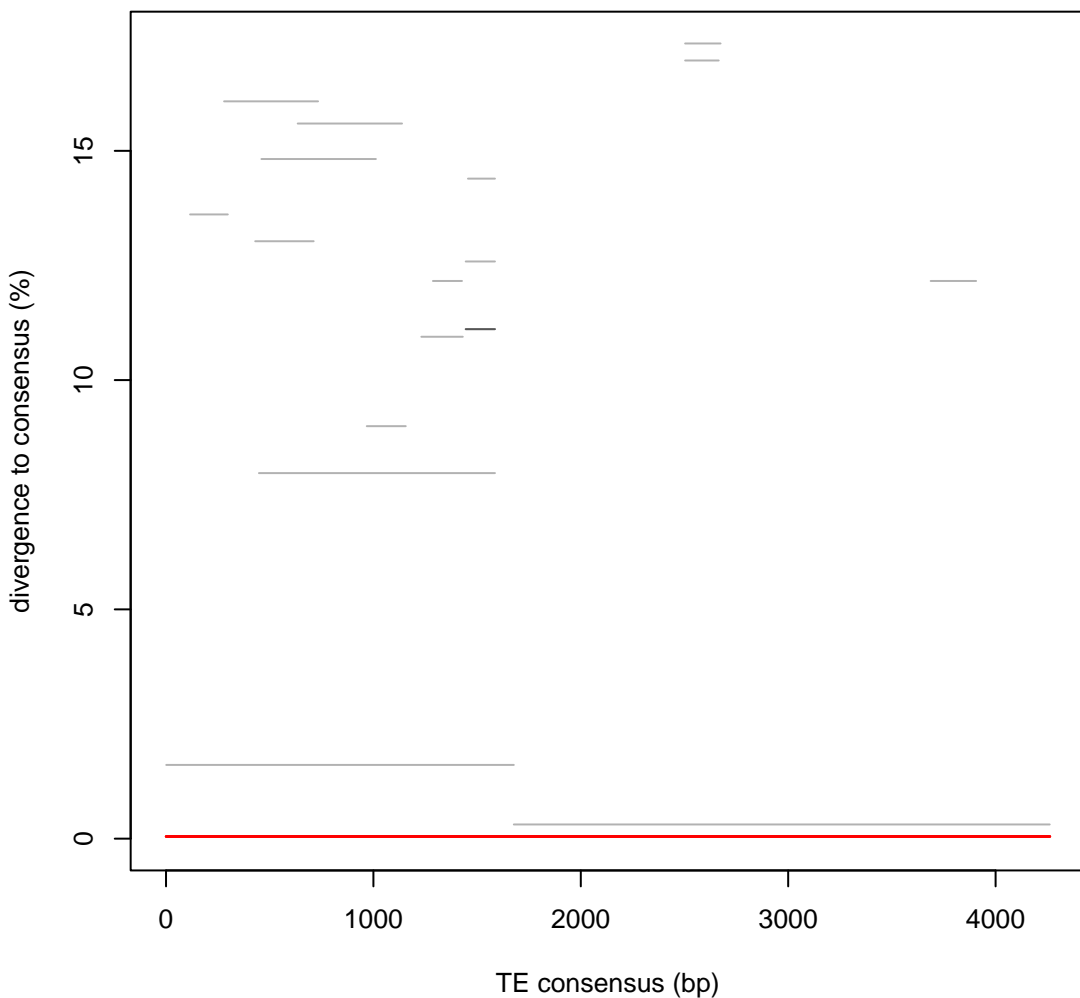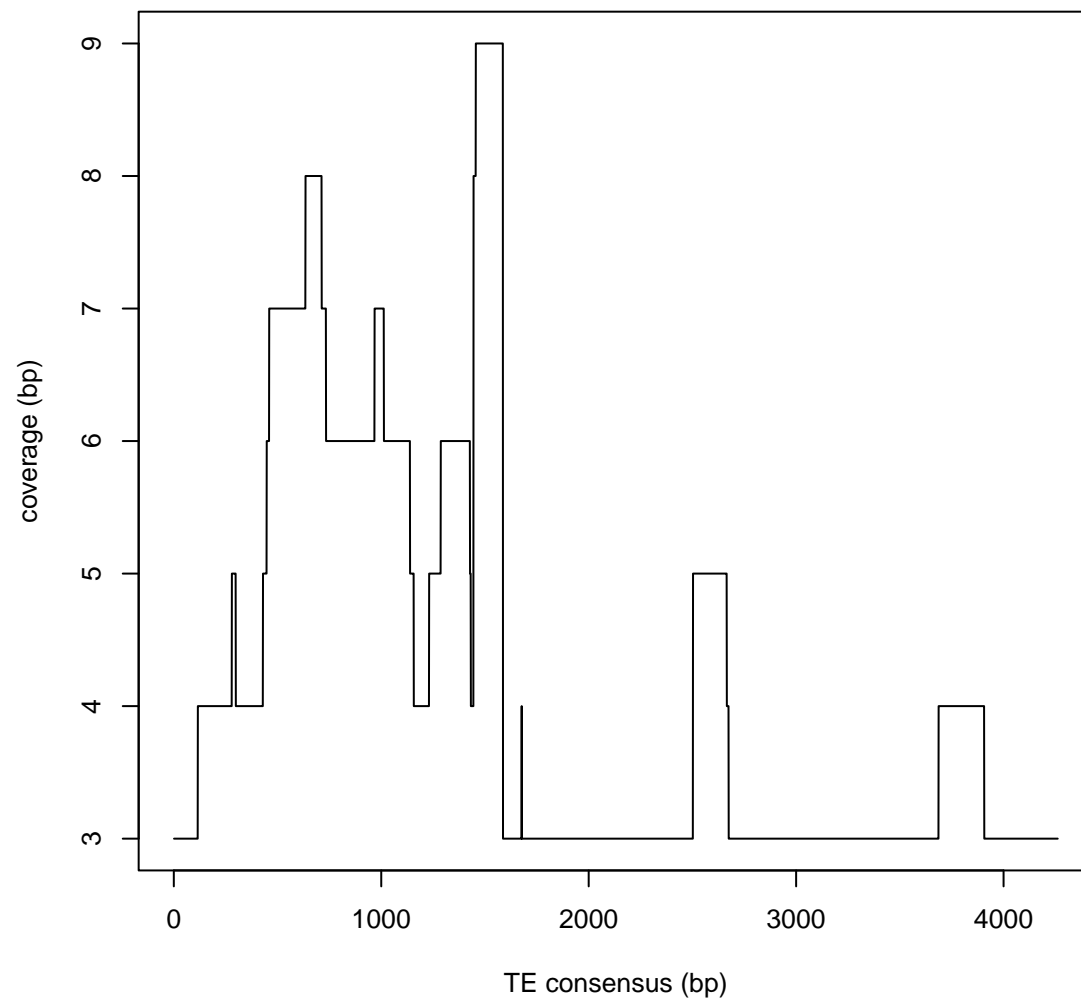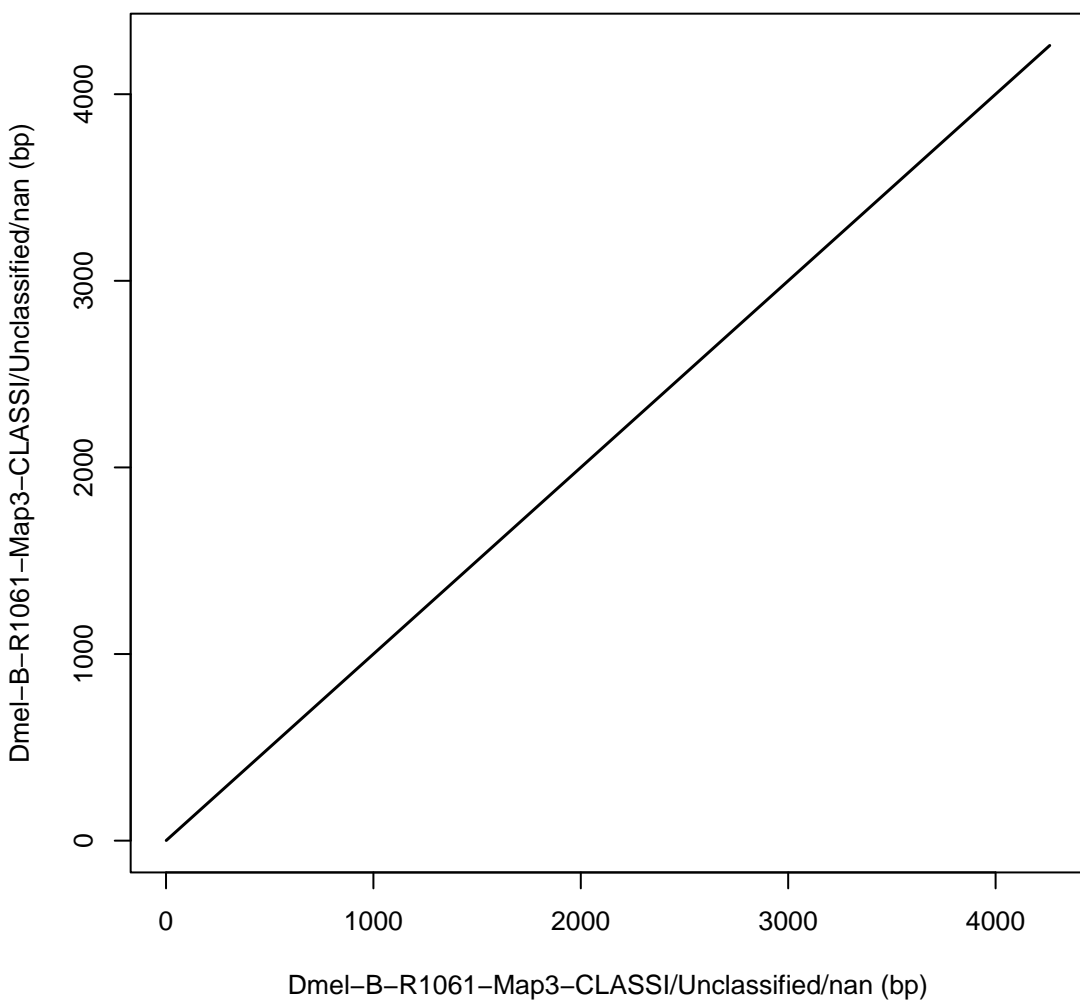TE: Dmel−B−R1061−Map3−CLASSI/Unclassified/nan
consensus size: 4261bp; fragments: 21; full length: 2 (>=3834.9bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

no orf >400 bp detected