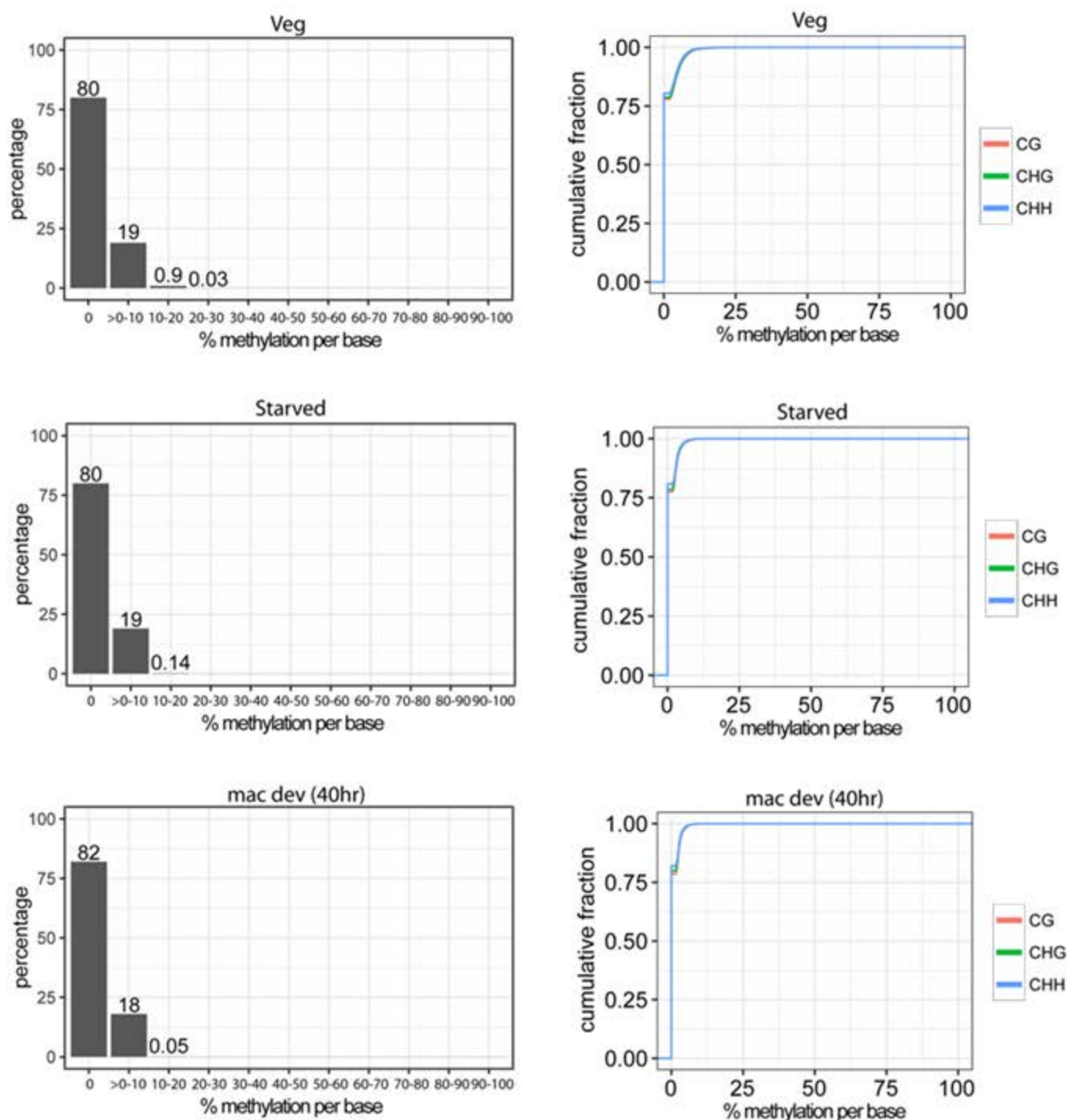


S6 Fig

S6 Fig: Absence of evidence of C methylation in *Paramecium* mac genome. Left panel, percent calculated using $C/(C + T)$ for each C in *Paramecium* genome. There was no bias in C conversion among the three different nucleotide contexts (right panel).