



Supplementary Figure 4. Accuracy and precision of the median coverage approach versus the modal coverage approach. Each cell represents the deviation of the estimated mean rDNA copy number, calculated from the 100 technical replicates for each coverage level, from the “true” rDNA copy number (20), using a sliding window size of 600 bp. The heatmap scales used are indicated, and rDNA copy number was rounded to the nearest integer.