



**Figure 7– Supplement 1. Individual mRNAs do not show evidence of preferential translation.** Cumulative read counts across the length of individual transcripts were calculated and normalized to the sum of ribosome footprints of mitochondrial transcripts in each respective library. Red = BMMs treated with harringtonine. Black = BMMs untreated. Numbers in parentheses = total read counts. Rows = individual transcript. Columns = infection condition. Data are representative of two independent experiments.