



Figure 1: **Additional Figure: Mean RSCU values for virus and non-virus classes.** The mean values for each codon's RSCU value among the samples belonging to the virus class are depicted with narrow colored bars in the foreground. The mean values of each RSCU value among the non-viral data points are depicted with the wider bars in the background. The six bars colored black indicate the features (the RSCU values) that were the most important features according to the random forest feature importance.