S5 Fig. Prediction model with the variable ‘total recovered RNA’ (total RNA) included. 
(a) Model 1 which includes miRNAs (classified by MIMAT#########) from mid-(MP) and 
late-(LP) pregnancy as separate variables. Overall misclassification rate was 17.78%.
(b) Model 2 includes the difference in miRNA expression between mid- and late-
pregnancy. Overall misclassification rate was 24.44%. (c) Using classification model #1 
to classify the HEua group, results in a majority of the HEua group predicted to be HEa-
like throughout pregnancy.