# Predicting Flu Vaccinations:

Using survey questions to predict which respondents will receive a seasonal flu vaccine or novel flu vaccine

# The Problem

 Hard to predict vaccination status based off demographics alone

- Spread of misinformation/conspiracies
- Not enough clean data on COVID
- Public health efforts not targeted with precision
- Herd immunity has only been a reality for a few viruses
  - Virus eradication even less common







#### The Data

- CDC's National Center for Health Statistics (NCHS) and National Center for Immunization and Respiratory Diseases (NCIRD)
  - O The National 2009 H1N1 Flu Survey (October 2009 and June 2010)
  - O 35 questions:
    - 13 yes/no questions about behaviors, health and demographics
    - 8 opinion questions rated 0-5
    - 14 demographic questions with categorical answers

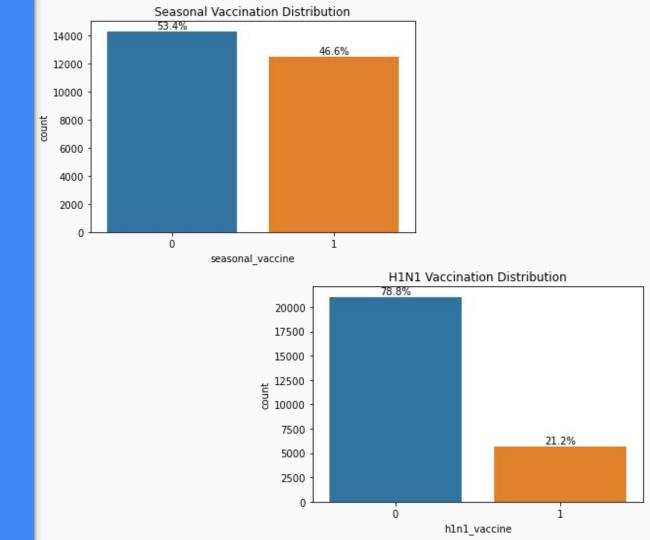
## Data Cleaning

- Relatively clean raw data
- Missing values addressed by using mode of that particular column/feature
  - Non-Normal distributions
  - For all but 3 features
- 3 columns given additional categorical response of "unknown"
  - o Due to large proportion of missing values

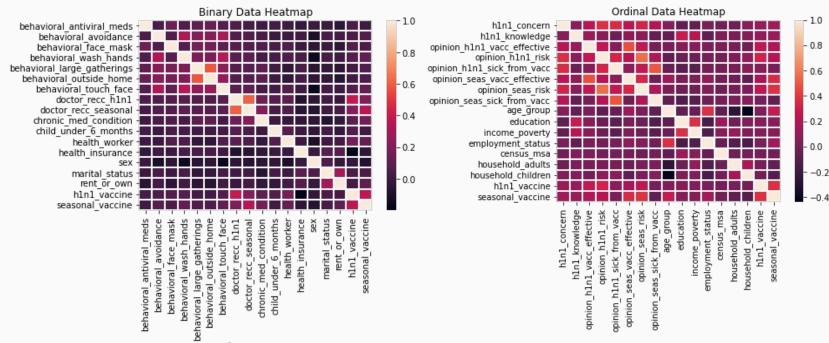


## **Data Exploration**

- Target variables were binary:
  - o 0/1 for no/yes
  - H1N1 vaccine and
  - Seasonal flu vaccine
  - Relatively balanced
    - H1N1 less so



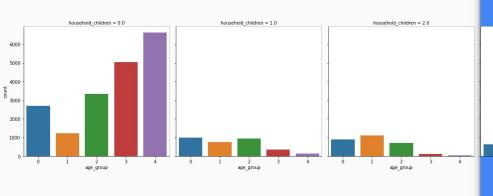
### Data Exploration: Heatmaps

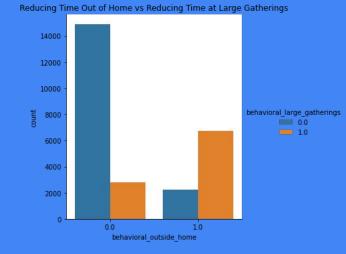


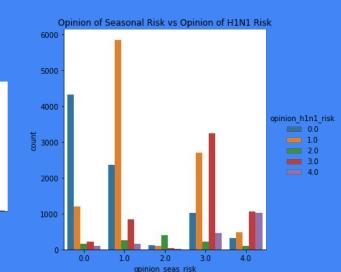
- Heatmaps of binary data and ordinal data used to visualize correlations
- Few strong correlations seen
- Good candidate for machine learning instead of simple correlation study

# Strongest Correlations

- Highest 3 correlation coefficients were 0.58, 0.56 and -0.43
- Above scores for behavioral, opinion and demographic features respectively







household children = 3.0

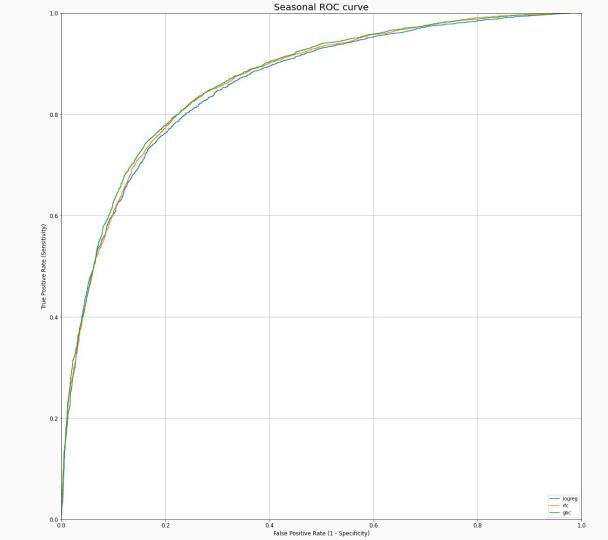
# Modeling Overview/Steps

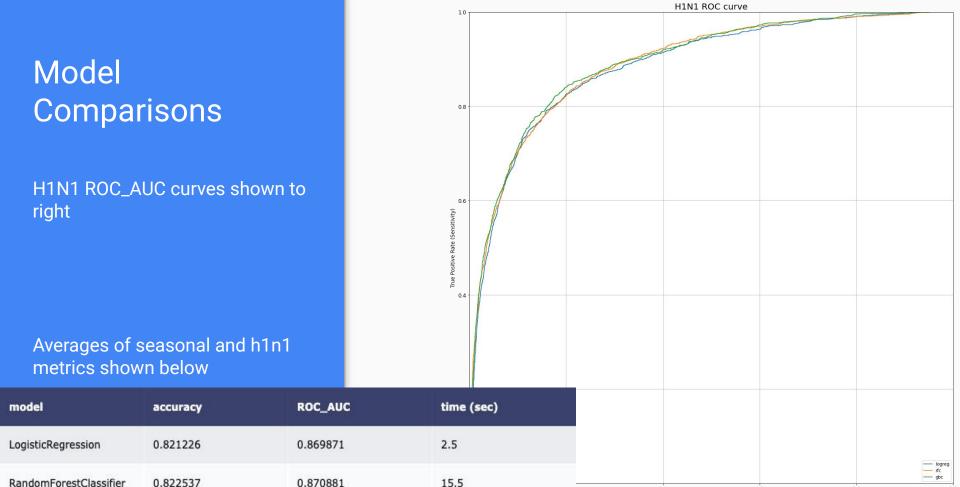
- 1. Data Preprocessing:
  - a. One Hot Encoding of categorical variables
    - i. Results in ~60 additional columns
  - b. Splitting data into train and test sets
    - i. 70/30 split
  - c. Scaling only necessary for certain models
  - d. Data is already relatively well balanced
- 2. Hyperparameter Tuning:
  - a. Used scikit-learn's GridSearchCV
    - i. 5 fold cross validation
    - ii. Evaluation metric = ROC\_AUC
- 3. Train models on 70% of data (train set)
  - a. Evaluate model performance on remaining 30% (test set)

# Model Comparisons

- ROC\_AUC used to choose best of 3 most accurate models
- Cross validated accuracy, roc\_auc and CPU time also compared

Seasonal ROC\_AUC curves shown to right





9.0

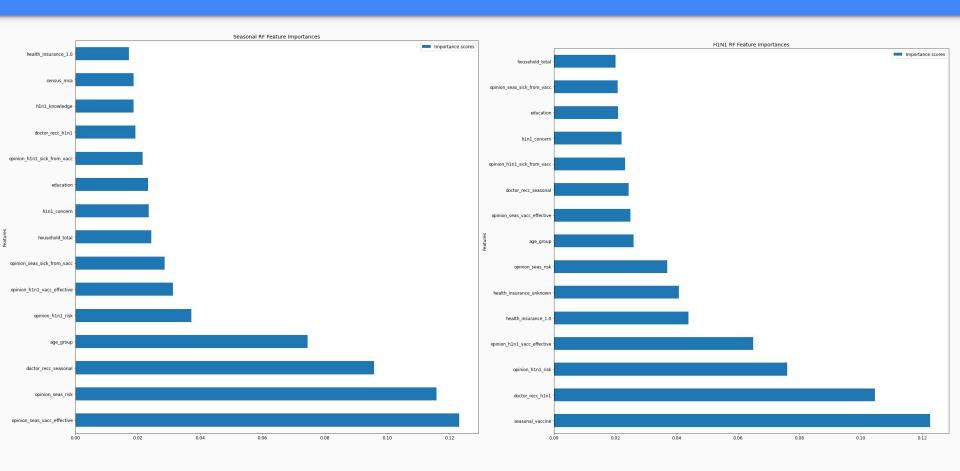
0.87478

GradientBoostingClassifier

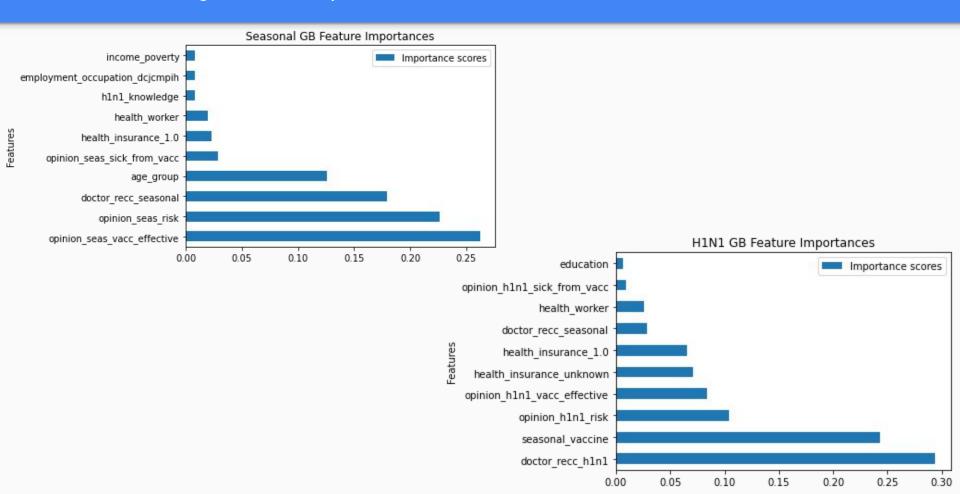
0.825408

False Positive Rate (1 - Specificity)

#### Random Forest Feature Importances



#### **Gradient Boosting Feature Importances**



## Using the GBC Model

- The GradientBoostingClassifier model was shown to be most accurate and displayed the highest ROC\_AUC
- ROC\_AUC was the metric of choice for the contest being entered
- First submission to contest resulted in a ROC\_AUC of 0.8535 which placed 490th among 4000 entries
  - With minor tweaks I was able to increase ROC\_AUC to 0.8572, placing 410th out of 4000
- My model submission scored better than 90% of all other competing user's submissions

# Conclusions and Future

- Any of these 3 models provide reasonable results
  - These 3 were compared after KNN, GNB and others were deemed least accurate
- Choice between these 3 depends on what health officials find most important
  - Lowest CPU time → Logistic Regression
  - Feature reduction → Random Forest
  - Simplest tuning → Random Forest
  - Highest accuracy/roc\_auc → Gradient Boosting
  - Most complex/fine tuned → Gradient Boosting
- Future questionnaires may want to increase number of opinion questions and experience questions as opposed to demographic questions
- My Recommendation: Random Forest Classifier
  - Extremely close to GBC in accuracy but requires less tuning
  - Allows for large reduction in features/questions on questionnaire