

# Predicting seasonal flu vaccine uptake

Machine Learning Individual Project 2020-21

Camilla Callierotti

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## Outline

- 1 Dataset
- 2 Data preprocessing
- 3 Exploratory analysis
- 4 Train and test splitting
- 5 Logistic regression
- 6 Random forest
- 7 Neural network
- 8 Results
- 9 Discussion
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# Dataset

```
Dimensions of X: (26707, 36)
Dimensions of Y: (26707, 3)
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 26707 entries, 0 to 26706
Data columns (total 36 columns):
```

#	Column	Non-Null Count	Dtype
0	respondent_id	26707 non-null	int64
1	h1n1_concern	26615 non-null	float64
2	h1n1_knowledge	26591 non-null	float64
3	behavioral_antiviral_meds	26636 non-null	float64
4	behavioral_avoidance	26499 non-null	float64
5	behavioral_face_mask	26688 non-null	float64
6	behavioral_wash_hands	26665 non-null	float64
7	behavioral_large_gatherings	26620 non-null	float64
8	behavioral_outside_home	26625 non-null	float64
9	behavioral_touch_face	26579 non-null	float64
10	doctor_recc_h1n1	24547 non-null	float64
11	doctor_recc_seasonal	24547 non-null	float64
12	chronic_med_condition	25736 non-null	float64
13	child_under_6_months	25987 non-null	float64
14	health_worker	25903 non-null	float64
15	health_insurance	14433 non-null	float64
16	opinion_h1n1_vacc_effective	26316 non-null	float64
17	opinion_h1n1_risk	26319 non-null	float64
18	opinion_h1n1_sick_from_vacc	26312 non-null	float64
19	opinion_seas_vacc_effective	26245 non-null	float64
20	opinion_seas_risk	26193 non-null	float64
21	opinion_seas_sick_from_vacc	26170 non-null	float64
22	age_group	26707 non-null	object
23	education	25300 non-null	object
24	race	26707 non-null	object
25	sex	26707 non-null	object
26	income_poverty	22284 non-null	object
27	marital_status	25299 non-null	object
28	rent_or_own	24665 non-null	object
29	employment_status	25244 non-null	object
30	hhs_geo_region	26707 non-null	object
31	census_msa	26707 non-null	object
32	household_adults	26458 non-null	float64
33	household_children	26458 non-null	float64
34	employment_industry	13377 non-null	object
35	employment_occupation	13237 non-null	object

```
dtypes: float64(23), int64(1), object(12)
```

- 26'707 observations
- 36 features
  - Vaccine opinions
  - Behaviours
  - Sociodemographic factors

⇒ Dataset also contained information on H1N1 vaccine but the focus of this project is seasonal vaccine

Figure 1: Original dataset features and structure

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## Missingness

- $>50\%$  observations missing: drop feature
- $<50\%$  observations missing: imputation
  - Mode imputation (categorical features)
  - Mean imputation (numerical features)
- Column containing indecipherable observations: drop feature

## Categorical features

- Label encoding (hierarchical features)
- One-hot encoding (non-hierarchical features)

	age_group	education	race	sex	income_poverty	marital_status	rent_or_own	employment_status	census_msa
0	55 - 64 Years	< 12 Years	White	Female	Below Poverty	Not Married	Own	Not in Labor Force	Non-MSA
1	35 - 44 Years	12 Years	White	Male	Below Poverty	Not Married	Rent	Employed	MSA, Not Principle City
2	18 - 34 Years	College Graduate	White	Male	<= \$75,000, Above Poverty	Not Married	Own	Employed	MSA, Not Principle City
3	65+ Years	12 Years	White	Female	Below Poverty	Not Married	Rent	Not in Labor Force	MSA, Principle City
4	45 - 54 Years	Some College	White	Female	<= \$75,000, Above Poverty	Married	Own	Employed	MSA, Not Principle City

## Collinear features (1/2)

Feature engineering to merge collinear features into one normally distributed feature:

- Cleanliness: Antiviral meds, avoidance, face mask, hand washing, large gatherings, outside home, face touching
- Opinion on seasonal flu vaccine: Vaccine effectiveness, seasonal flu risk, getting seasonal flu from vaccine

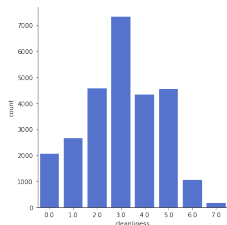


Figure 2: Cleanliness variable distribution

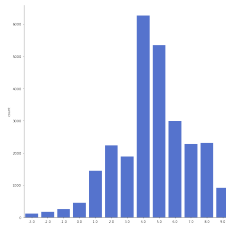


Figure 3: Opinion variable distribution



## Collinear features (2/2)

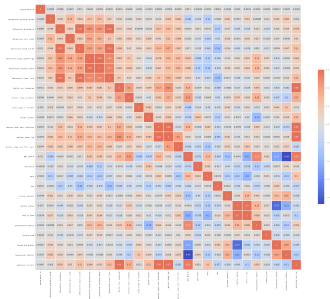


Figure 4: Heatmap before  
feature engineering

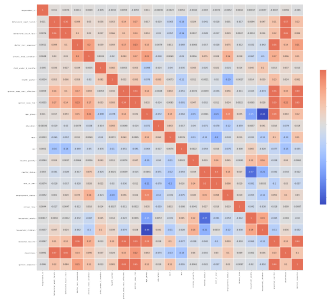


Figure 5: Heatmap after  
feature engineering

## Standardisation

Standardisation to standard normal random variables with mean 0 and standard deviation 1:

$$z = \frac{x - \mu}{\sigma}$$

Motivation: features are in different scales, so standardisation brings them to comparable scales

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## Variable distributions

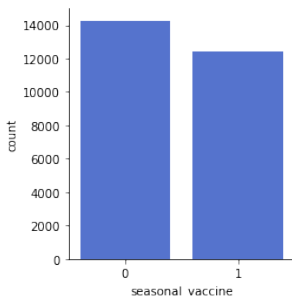


Figure 6: Outcome variable distribution

Approximately equal case-control distribution, meaning precision metrics (specificity and accuracy) won't be affected by balance

# Variable distributions

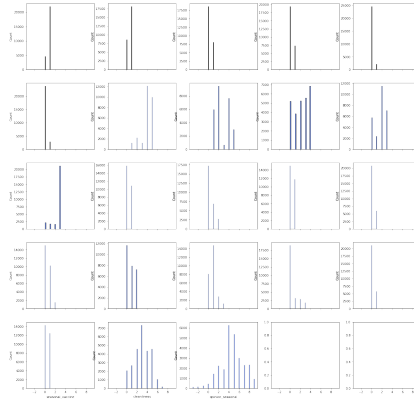


Figure 7: Predictor variable distributions

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## Train and test splitting

80-20 train-test split

Motivation: 26'707 observations so a test set of 20% (small) will still give robust error metrics

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# Logistic regression

Motivation:

- Benchmark model
- Understand the predictive power of a simple model

## Logistic regression: performance

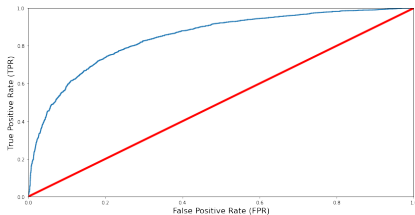


Figure 8: ROC curve  
AUC= 0.84

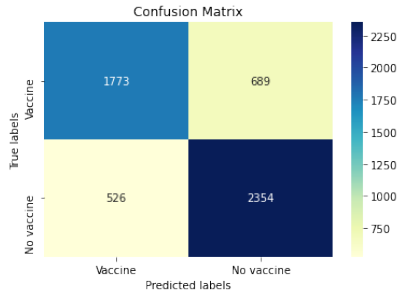


Figure 9: Confusion matrix  
Sensitivity= 0.77  
Specificity= 0.77  
Precision= 0.72  
Accuracy= 0.77  
F1 score= 0.74

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# Random forest

## Motivation:

- Understand the added accuracy of an ensemble algorithm
- Performs very well for classification problems compared to other improvements of decision tree algorithms
- Obtain a variable importance plot

## Random forest: hyperparameter tuning

Hyperparameter	Base model	Tuned model
Estimators	100	1600
Min samples per leaf	2	4
Min samples per split	1	2

Table 1: Hyperparameters of base model vs new model tuned by cross-validated grid search

⇒ The new model was computationally intensive (80 min) but only improved AUC by 2 ⇒ Keep base model

## Random forest: one tree

At the root node there are less samples than training data points highlighting the random forest's training with bagging (on a random subset with replacement)

## Random forest: variable importance plot

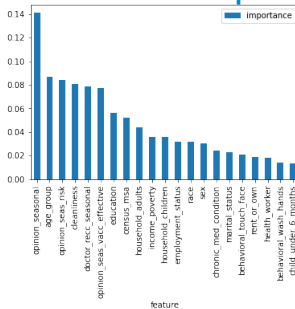


Figure 10: Variable importance plot

Relevance:

- Interpretability: view features most associated with outcome
- Reduce overfitting: remove non-contributing features

## Random forest: performance

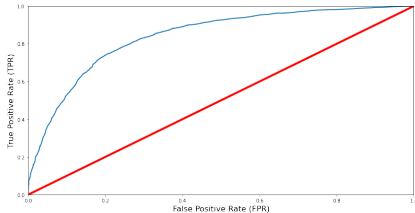


Figure 11: ROC curve  
AUC= 0.84

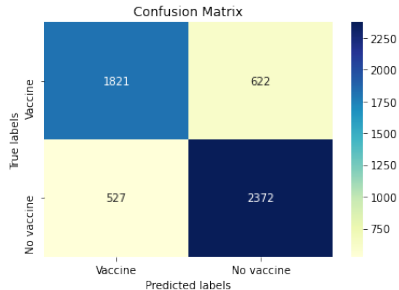


Figure 12: Confusion matrix  
Sensitivity= 0.78  
Specificity= 0.79  
Precision= 0.75  
Accuracy= 0.78  
F1 score= 0.76



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## Neural network: motivation

Motivation:

- Unsupervised learning algorithm
- Can handle complex data structure (21 features)

## Neural network: input and output layers

Input layer depth: 21  
(number of unique features)

Output layer depth: 1 (binary classification with sigmoid activation function)

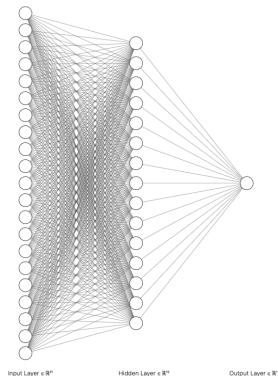


Figure 13: Neural network architecture

## Neural network: hidden layer

Number of layers (depth):

- One hidden layer is sufficient for the majority of problems

Number of neurons in hidden layer (width):

- $<$  input layer width,  
   $>$  output layer width
- $< 2$  times input layer width
- $2/3$  input layer width  
   $+$  output layer width

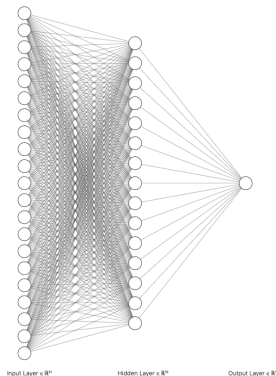


Figure 14: Neural network architecture

## Neural network: hyperparameter tuning

Method: Cross-validated grid search

Hyperparameters:

- Batch size: 100
- Epochs: 50
- Optimisation algorithm: Adam

## Neural network: performance

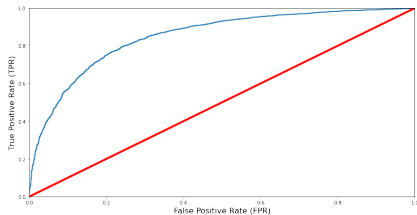


Figure 15: ROC curve  
AUC= 0.85

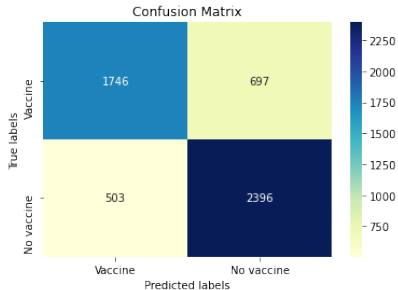


Figure 16: Confusion matrix  
Sensitivity= 0.78  
Specificity= 0.77  
Precision= 0.71  
Accuracy= 0.78  
F1 score= 0.74

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## Results

Model	ROC-AUC	Accuracy	F1 score
Logistic regression	0.845	0.77	0.74
Random forest	0.830	0.78	0.76
Neural network	0.85	0.78	0.74

Table 2: Performance metrics of three classification algorithms

- ⇒ All models have comparable performance metrics
- ⇒ Prediction performance by models is better than random chance, but is far from perfect
- ⇒ Measures are coherent, there is no over/under-representation of either class



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## Discussion

- ⇒ Predicting vaccine uptake does not require complex transformations of data (linearly separable data)
- ⇒ However, perfect prediction is unlikely due to an emotive component to the nature of the decision which cannot be fully captured in the 21 features (in fact the biggest contributor to prediction is the opinion variable)

## Discussion

Study relevance:

- Projecting vaccination rates
  - Informing decision of number of doses to purchase from manufacturers
  - Informing decision of how to allocated health resources
- Predicting whether immunisation programme will reach protective efficacy
- Leveraging factors most associated with vaccine compliancy

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## Evaluation

### Strengths:

- Research question could be answered with a simple model (logistic regression) that did not use too much computational power - a complex model is not always required
- Was able to choose a well-performing random forest over a very slightly better performing one to save computational power

### Limitations:

- Could not tune all hyperparameters due to computational resources needed
- Data available cannot fully capture determinants of outcome due to the nature of the research question