• Group: **14**

Student pace: Part-time

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Instructor name:Blog post URL: N/A

Predicting H1N1 Flu Vaccination Status ¶

Overview

As the world struggles to vaccinate the global population against COVID-19, an understanding of how people's backgrounds, opinions, and health behaviors are related to their personal vaccination patterns can provide guidance for future public health efforts. Such findings can guide policymakers and public health professionals develop public health efforts to improve vaccine uptake to mitigate spread of preventable communicable dieases.

This project utilises data from a United States' conducted National 2009 H1N1 Flu Survey to predict whether someone reviewed H1N1 flu vaccines. Gaining deeper insights into how these attributes correlate with individual vaccination behaviors can offer valuable direction for upcoming public health initiatives.

Business Understanding

The National 2009 H1N1 Flu Survey data was downloaded from DrivenData and the purpose of this project is to use data to forecast whether or not a person received the H1N1 flu vaccination, using features such as social, economic, and demographic background, opinions on risks of illness and vaccine effectiveness, and behaviors towards mitigating transmission, etcetera. The findings would also be applicable for use by the Kenya's Ministry of Health to more effectively target public health initiatives that boost vaccination rates and localise for other communicable dieases like influenza.

Research Questions

1. How may false information regarding vaccines be effectively refuted to encourage factual knowledge and vaccination acceptance of the flu?

- 2. Which major groups are most likely to be deceived about the flu shot, especially those with lower levels of education? How might focused messaging be developed to allay their fears?
- 3. How can resource allocation for flu vaccination programs be optimized by the use of predictive modeling, guaranteeing effective distribution to high-risk populations and regions?
- 4. What programs may be put in place to promote the general adoption of preventative measures like donning face masks, avoiding crowds, and washing your hands properly?
- 5. How can cooperation with medical professionals be improved to encourage routine physical examinations and support physicians' recommendations for flu shots among high-risk individuals?

Objectives

- 1. Analyze how well various communication tactics work to debunk vaccine misconceptions and spread factual information about getting the flu shot.
- 2. Determine which high-risk groups to target with messaging and interventions, especially those who are less educated and show little concern for the H1N1.
- 3. Create a predictive model that makes the best use of behavioral, epidemiological, and demographic data to allocate resources for flu vaccine campaigns.
- 4. Create and put into action policies and initiatives that will increase people's adherence to preventive measures like hand hygiene, avoiding crowds, and wearing face masks.
- 5. Create efficient channels of communication with medical professionals to enable routine physical examinations and improve physician recommendations for influenza shots in high-risk patients. nts.

Data Description

- h1n1 concern Level of concern about the H1N1 flu.
 - 0 = Not at all concerned; 1 = Not very concerned; 2 = Somewhat concerned; 3 = Very concerned.
- h1n1 knowledge Level of knowledge about H1N1 flu.
 - 0 = No knowledge; 1 = A little knowledge; 2 = A lot of knowledge.
- behavioral_antiviral_meds Has taken antiviral medications. (binary)
- behavioral_avoidance Has avoided close contact with others with flu-like symptoms. (binary)
- behavioral face mask Has bought a face mask. (binary)
- behavioral_wash_hands Has frequently washed hands or used hand sanitizer. (binary)
- behavioral large gatherings Has reduced time at large gatherings. (binary)
- behavioral_outside_home Has reduced contact with people outside of own household. (binary)

- behavioral touch face Has avoided touching eyes, nose, or mouth. (binary)
- doctor recc h1n1 H1N1 flu vaccine was recommended by doctor. (binary)
- doctor recc seasonal Seasonal flu vaccine was recommended by doctor. (binary)
- chronic_med_condition Has any of the following chronic medical conditions: asthma or an other lung condition, diabetes, a heart condition, a kidney condition, sickle cell anemia or other anemia, a neurological or neuromuscular condition, a liver condition, or a weakened immune system caused by a chronic illness or by medicines taken for a chronic illness. (binary)
- child_under_6_months Has regular close contact with a child under the age of six months. (binary)
- health worker Is a healthcare worker. (binary)
- health_insurance Has health insurance. (binary)
- opinion h1n1 vacc effective Respondent's opinion about H1N1 vaccine effectiveness.
 - 1 = Not at all effective; 2 = Not very effective; 3 = Don't know; 4 = Somewhat effective; 5 = Very effective.
- opinion_h1n1_risk Respondent's opinion about risk of getting sick with H1N1 flu without vaccine.
 - 1 = Very Low; 2 = Somewhat low; 3 = Don't know; 4 = Somewhat high; 5 = Very high.
- opinion h1n1 sick from vacc Respondent's worry of getting sick from taking H1N1 vaccine.
 - 1 = Not at all worried; 2 = Not very worried; 3 = Don't know; 4 = Somewhat worried; 5 = Very worried.
- opinion seas vacc effective Respondent's opinion about seasonal flu vaccine effectiveness.
 - 1 = Not at all effective; 2 = Not very effective; 3 = Don't know; 4 = Somewhat effective; 5 = Very effective.
- opinion seas risk Respondent's opinion about risk of getting sick with seasonal flu without vaccine.
 - 1 = Very Low; 2 = Somewhat low; 3 = Don't know; 4 = Somewhat high; 5 = Very high.
- opinion_seas_sick_from_vacc Respondent's worry of getting sick from taking seasonal flu vaccine.
 - 1 = Not at all worried; 2 = Not very worried; 3 = Don't know; 4 = Somewhat worried; 5 = Very worried.
- age_group Age group of respondent.
- · education Self-reported education level.
- race Race of respondent.
- · sex Sex of respondent.
- income poverty Household annual income of respondent with respect to 2008 Census poverty thresholds.
- marital_status Marital status of respondent.
- · rent or own Housing situation of respondent.
- employment_status Employment status of respondent.
- hhs_geo_region Respondent's residence using a 10-region geographic classification defined by the U.S. Dept. of Health and Human Services. Values are represented as short random character strings.
- census msa Respondent's residence within metropolitan statistical areas (MSA) as defined by the U.S. Census.
- household adults Number of other adults in household, top-coded to 3.
- household children Number of children in household, top-coded to 3.
- employment_industry Type of industry respondent is employed in. Values are represented as short random character strings.

• employment_occupation - Type of occupation of respondent. Values are represented as short random character strings.resented as short

Import Libraries

```
In [1]: import pandas as pd
        import numpy as np
        from sklearn.experimental import enable iterative imputer
        from sklearn.impute import IterativeImputer
        from sklearn.preprocessing import OneHotEncoder, StandardScaler, MinMaxScaler
        from sklearn.model selection import train test split, cross validate
        from imblearn.over sampling import SMOTE
        from sklearn.neighbors import KNeighborsClassifier
        from sklearn.metrics import accuracy score, precision score, recall score, f1 score, roc curve, roc auc score
        from sklearn.metrics import accuracy score, classification report, confusion matrix
        from sklearn.naive bayes import GaussianNB
        from sklearn.feature selection import RFE, mutual info classif
        from sklearn.linear model import LogisticRegression
        from sklearn.tree import DecisionTreeClassifier
        from sklearn.model selection import GridSearchCV
        from xgboost import XGBClassifier
        from sklearn.ensemble import BaggingClassifier, RandomForestClassifier
        from sklearn.pipeline import Pipeline
        import matplotlib.pyplot as plt
        import seaborn as sns
        plt.style.use('seaborn-v0 8-darkgrid')
```

Load Data

```
In [2]: # Loading data [original data from 'DrivenData' was seperated into training and testing sets]
    train_features_df = pd.read_csv("Data/training_set_features.csv", index_col="respondent_id")

    train_labels_df = pd.read_csv("Data/training_set_labels.csv",index_col="respondent_id")

    test_validation_df = pd.read_csv("Data/test_set_features.csv",index_col="respondent_id")

In [3]: # check first 5 rows of features_df
    train_features_df.head()
```

Out[3]:

	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands bel
respondent_id						
0	1.0	0.0	0.0	0.0	0.0	0.0
1	3.0	2.0	0.0	1.0	0.0	1.0
2	1.0	1.0	0.0	1.0	0.0	0.0
3	1.0	1.0	0.0	1.0	0.0	1.0
4	2.0	1.0	0.0	1.0	0.0	1.0

5 rows × 35 columns

In [4]: # check first 5 rows of labels_df
train_labels_df.head()

Out[4]:

h1n1_vaccine seasonal_vaccine

respondent_id							
	0	0	0				
	1	0	1				
	2	0	0				
	3	0	1				
	4	0	0				

In [5]: # check first 5 rows of test_validation_df
test_validation_df.head()

Out[5]:

	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands bel
respondent_id						
26707	2.0	2.0	0.0	1.0	0.0	1.0
26708	1.0	1.0	0.0	0.0	0.0	0.0
26709	2.0	2.0	0.0	0.0	1.0	1.0
26710	1.0	1.0	0.0	0.0	0.0	0.0
26711	3.0	1.0	1.0	1.0	0.0	1.0

5 rows × 35 columns

```
In [6]: # Double-check that the rows between the features and the labels match up
np.testing.assert_array_equal(train_features_df.index.values, train_labels_df.index.values)
```

In [7]: ## Merge the features and labels data
merged_df = pd.concat([train_features_df, train_labels_df], axis =1)
merged_df

Out[7]:

	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands	bel
respondent_id							
0	1.0	0.0	0.0	0.0	0.0	0.0	
1	3.0	2.0	0.0	1.0	0.0	1.0	
2	1.0	1.0	0.0	1.0	0.0	0.0	
3	1.0	1.0	0.0	1.0	0.0	1.0	
4	2.0	1.0	0.0	1.0	0.0	1.0	
26702	2.0	0.0	0.0	1.0	0.0	0.0	
26703	1.0	2.0	0.0	1.0	0.0	1.0	
26704	2.0	2.0	0.0	1.0	1.0	1.0	
26705	1.0	1.0	0.0	0.0	0.0	0.0	
26706	0.0	0.0	0.0	1.0	0.0	0.0	

26707 rows × 37 columns

In [8]: merged_df.info()

<class 'pandas.core.frame.DataFrame'>
Index: 26707 entries, 0 to 26706
Data columns (total 37 columns):

# 	Column	Non-Null Count	Dtype
0	h1n1_concern	26615 non-null	float64
1	h1n1_knowledge	26591 non-null	float64
2	behavioral_antiviral_meds	26636 non-null	float64
3	behavioral_avoidance	26499 non-null	float64
4	behavioral_face_mask	26688 non-null	float64
5	behavioral_wash_hands	26665 non-null	float64
6	behavioral_large_gatherings	26620 non-null	float64
7	behavioral_outside_home	26625 non-null	float64
8	behavioral_touch_face	26579 non-null	float64
9	doctor_recc_h1n1	24547 non-null	float64
10	doctor_recc_seasonal	24547 non-null	float64
11	<pre>chronic_med_condition</pre>	25736 non-null	float64
12	child_under_6_months	25887 non-null	float64
13	health_worker	25903 non-null	float64
14	health_insurance	14433 non-null	float64
15	<pre>opinion_h1n1_vacc_effective</pre>	26316 non-null	float64
16	opinion_h1n1_risk	26319 non-null	float64
17	opinion_h1n1_sick_from_vacc	26312 non-null	float64
18	opinion_seas_vacc_effective	26245 non-null	float64
19	opinion_seas_risk	26193 non-null	float64
20	opinion_seas_sick_from_vacc	26170 non-null	float64
21	age_group	26707 non-null	object
22	education	25300 non-null	object
23	race	26707 non-null	object
24	sex	26707 non-null	object
25	income_poverty	22284 non-null	object
26	marital_status	25299 non-null	object
27	rent_or_own	24665 non-null	object
28	employment_status	25244 non-null	object
29	hhs_geo_region	26707 non-null	object
30	census_msa	26707 non-null	object
31	household_adults	26458 non-null	float64
32	household_children	26458 non-null	float64
33	employment_industry	13377 non-null	object
34	employment_occupation	13237 non-null	object
35	h1n1_vaccine	26707 non-null	int64

36 seasonal vaccine 26707 non-null int64

dtypes: float64(23), int64(2), object(12)

memory usage: 7.7+ MB

Additionally the datatypes consist of 23 columns of float type, 2 interger(int64) columns and 12 columns of object datatype.

In [9]: merged_df.describe()

Out[9]:

	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands	behavioral_				
count	26615.000000	26591.000000	26636.000000	26499.000000	26688.000000	26665.000000					
mean	1.618486	1.262532	0.048844	0.725612	0.068982	0.825614					
std	0.910311	0.618149	0.215545	0.446214	0.253429	0.379448					
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000					
25%	1.000000	1.000000	0.000000	0.000000	0.000000	1.000000					
50%	2.000000	1.000000	0.000000	1.000000	0.000000	1.000000					
75%	2.000000	2.000000	0.000000	1.000000	0.000000	1.000000					
max	3.000000	2.000000	1.000000	1.000000	1.000000	1.000000					
8 rows	8 rows x 25 columns										

8 rows × 25 columns

Inference

Target Variable

• h1n1_vaccine: The sample popluation show that 21.24% received the vacccine, therefore 78.76% during the survey year (2009)

Data Cleaning

In [10]:	<pre>merged_df.isna().sum()</pre>	
Out[10]:	h1n1_concern	92
	h1n1_knowledge	116
	behavioral_antiviral_meds	71
	behavioral_avoidance	208
	behavioral_face_mask	19
	behavioral_wash_hands	42
	behavioral_large_gatherings	87
	behavioral_outside_home	82
	behavioral_touch_face	128
	doctor_recc_h1n1	2160
	doctor_recc_seasonal	2160
	chronic_med_condition	971
	child_under_6_months	820
	health_worker	804
	health_insurance	12274
	<pre>opinion_h1n1_vacc_effective</pre>	391
	opinion_h1n1_risk	388
	opinion_h1n1_sick_from_vacc	395
	<pre>opinion_seas_vacc_effective</pre>	462
	opinion_seas_risk	514
	opinion_seas_sick_from_vacc	537
	age_group	0
	education	1407
	race	0
	sex	0
	income_poverty	4423
	marital_status	1408
	rent_or_own	2042
	employment_status	1463
	hhs_geo_region	0
	census_msa	0
	household_adults	249
	household_children	249
	employment_industry	13330
	<pre>employment_occupation</pre>	13470
	h1n1_vaccine	0
	seasonal_vaccine	0
	dtype: int64	

In [11]: missing = [[column, round(sum(merged_df[column].isna())/len(merged_df[column]),2)] for column in merged_df.columns]
pd.DataFrame(missing, columns= ['column name', 'missing proportion']).sort_values(by='missing proportion', ascending=

Out[11]:

	column name	missing proportion
34	employment_occupation	0.50
33	employment_industry	0.50
14	health_insurance	0.46
25	income_poverty	0.17
27	rent_or_own	0.08
9	doctor_recc_h1n1	0.08
10	doctor_recc_seasonal	0.08
26	marital_status	0.05
22	education	0.05
28	employment_status	0.05
11	chronic_med_condition	0.04
13	health_worker	0.03
12	child_under_6_months	0.03
19	opinion_seas_risk	0.02
20	opinion_seas_sick_from_vacc	0.02
18	opinion_seas_vacc_effective	0.02
31	household_adults	0.01
15	opinion_h1n1_vacc_effective	0.01
16	opinion_h1n1_risk	0.01
17	opinion_h1n1_sick_from_vacc	0.01
32	household_children	0.01
3	behavioral_avoidance	0.01
30	census_msa	0.00
29	hhs_geo_region	0.00
35	h1n1_vaccine	0.00

	column name	missing proportion
0	h1n1_concern	0.00
24	sex	0.00
23	race	0.00
21	age_group	0.00
1	h1n1_knowledge	0.00
8	behavioral_touch_face	0.00
7	behavioral_outside_home	0.00
6	behavioral_large_gatherings	0.00
5	behavioral_wash_hands	0.00
4	behavioral_face_mask	0.00
2	behavioral_antiviral_meds	0.00
36	seasonal_vaccine	0.00

- We will use 40% as the threshold for null frequency in the columns to decide whether to consider the column for analysis or not.
- Sometimes columns with greater null percentage have more semantic meaning and thus they cannot be ignored.
- Here we can see health_insurance, employment_industry and employment_occupation columns with approximately 46%, 50% and 50% missing values percentages respectively, and thus they cannot be used for analysis as nearly half of the data is missing in those columns.

```
In [14]: X = merged_df.drop(columns= 'h1n1_vaccine')
y = merged_df['h1n1_vaccine']

In [15]: # Split data into train and test sets
X_train, X_test, y_train, y_test = train_test_split(X, y, random_state= 40)
```

Impute missing values

```
In [18]: # Numerical columns of X train
         numerical X train = X train.select dtypes(exclude=['object'])
         numerical X train.columns
Out[18]: Index(['h1n1 concern', 'h1n1 knowledge', 'behavioral antiviral meds',
                 'behavioral avoidance', 'behavioral face mask', 'behavioral wash hands',
                'behavioral large gatherings', 'behavioral outside home',
                'behavioral_touch_face', 'doctor_recc_h1n1', 'chronic med condition',
                'child under 6 months', 'health worker', 'opinion h1n1 vacc effective',
                'opinion h1n1 risk', 'opinion h1n1 sick from vacc', 'household adults',
                'household children'],
               dtvpe='object')
In [19]: # Numerical columns of X test
         numerical X test = X test.select dtypes(exclude=['object'])
         numerical X test.columns
Out[19]: Index(['h1n1 concern', 'h1n1 knowledge', 'behavioral antiviral meds',
                 'behavioral avoidance', 'behavioral face mask', 'behavioral wash hands',
                'behavioral large gatherings', 'behavioral outside home',
                'behavioral touch face', 'doctor recc h1n1', 'chronic med condition',
                'child under 6 months', 'health worker', 'opinion h1n1 vacc effective',
                 'opinion h1n1 risk', 'opinion h1n1 sick from vacc', 'household adults',
                'household children'],
               dtvpe='object')
```

```
In [20]: # Imputation of mising values on train set data [for numerical data]
# Instantiate imputer
imputer = IterativeImputer()

# Fit and tranform X_train
imputed_numerical_X_train = imputer.fit_transform(numerical_X_train)
imputed_numerical_X_train = pd.DataFrame(np.round(numerical_X_train, 0), columns = numerical_X_train.columns)
imputed_numerical_X_train
```

Out[20]:

	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands	bel
respondent_id							
7191	1.0	1.0	0.0	1.0	0.0	1.0	
23784	3.0	1.0	0.0	1.0	0.0	1.0	
316	2.0	1.0	0.0	1.0	0.0	1.0	
521	2.0	1.0	0.0	1.0	0.0	1.0	
21647	2.0	2.0	0.0	1.0	0.0	1.0	
21810	2.0	1.0	0.0	1.0	0.0	0.0	
23992	1.0	0.0	0.0	0.0	0.0	0.0	
14501	2.0	1.0	0.0	1.0	0.0	1.0	
14555	2.0	1.0	0.0	0.0	0.0	1.0	
11590	1.0	2.0	0.0	1.0	0.0	1.0	

20030 rows × 18 columns

localhost:8888/notebooks/DSC-phase-3-project/H1N1_and_seasonal_flu_vaccines_prediction.ipynb

```
In [21]: # Tranform X_test
imputed_numerical_X_test = imputer.transform(numerical_X_test)
imputed_numerical_X_test = pd.DataFrame(np.round(numerical_X_test, 0), columns = numerical_X_test.columns)
imputed_numerical_X_test
```

Out[21]:

	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands b	el
respondent_id							
6458	1.0	0.0	0.0	1.0	0.0	1.0	
12839	2.0	1.0	0.0	1.0	0.0	1.0	
4377	3.0	2.0	0.0	0.0	0.0	1.0	
2731	2.0	1.0	0.0	1.0	0.0	1.0	
4982	3.0	2.0	0.0	1.0	0.0	1.0	
18368	1.0	1.0	0.0	1.0	0.0	1.0	
7212	2.0	0.0	0.0	1.0	0.0	1.0	
13165	2.0	0.0	0.0	0.0	0.0	1.0	
17397	3.0	2.0	1.0	1.0	1.0	1.0	
20852	3.0	2.0	0.0	1.0	0.0	1.0	

6677 rows × 18 columns

Out[22]:

	age_group_35 - 44 Years	age_group_45 - 54 Years	age_group_55 - 64 Years	age_group_65+ Years	education_< 12 Years	education_College Graduate	education_Some College	education_nan	race
respondent_id									
7191	0.0	0.0	1.0	0.0	1.0	0.0	0.0	0.0	
23784	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0	
316	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	
521	0.0	1.0	0.0	0.0	0.0	0.0	1.0	0.0	
21647	0.0	0.0	1.0	0.0	0.0	1.0	0.0	0.0	
21810	0.0	1.0	0.0	0.0	0.0	0.0	1.0	0.0	
23992	0.0	1.0	0.0	0.0	0.0	0.0	0.0	1.0	
14501	0.0	0.0	0.0	0.0	0.0	0.0	1.0	0.0	
14555	0.0	0.0	1.0	0.0	0.0	1.0	0.0	0.0	
11590	0.0	0.0	0.0	0.0	0.0	1.0	0.0	0.0	

20030 rows × 33 columns

localhost:8888/notebooks/DSC-phase-3-project/H1N1_and_seasonal_flu_vaccines_prediction.ipynb

In [23]: # Check for columns with all zeros to drop
pd.DataFrame([[column, encoded_categorical_X_train[column].sum() == 0] for column in encoded_categorical_X_train.column

Out[23]:

	0	1
0	age_group_35 - 44 Years	False
1	age_group_45 - 54 Years	False
2	age_group_55 - 64 Years	False
3	age_group_65+ Years	False
4	education_< 12 Years	False
5	education_College Graduate	False
6	education_Some College	False
7	education_nan	False
8	race_Hispanic	False
9	race_Other or Multiple	False
10	race_White	False
11	sex_Male	False
12	income_poverty_> \$75,000	False
13	income_poverty_Below Poverty	False
14	income_poverty_nan	False
15	marital_status_Not Married	False
16	marital_status_nan	False
17	rent_or_own_Rent	False
18	rent_or_own_nan	False
19	employment_status_Not in Labor Force	False
20	employment_status_Unemployed	False
21	employment_status_nan	False
22	hhs_geo_region_bhuqouqj	False
23	hhs_geo_region_dqpwygqj	False
24	hhs_geo_region_fpwskwrf	False

	0	1
25	hhs_geo_region_kbazzjca	False
26	hhs_geo_region_lrircsnp	False
27	hhs_geo_region_lzgpxyit	False
28	hhs_geo_region_mlyzmhmf	False
29	hhs_geo_region_oxchjgsf	False
30	hhs_geo_region_qufhixun	False
31	census_msa_MSA, Principle City	False
32	census msa Non-MSA	False

Out[24]:

	age_group_35 - 44 Years	age_group_45 - 54 Years	age_group_55 - 64 Years	age_group_65+ Years	education_< 12 Years	education_College Graduate	education_Some College	education_nan	race
respondent_id									
6458	0.0	1.0	0.0	0.0	0.0	0.0	0.0	0.0	
12839	0.0	1.0	0.0	0.0	0.0	0.0	1.0	0.0	
4377	0.0	1.0	0.0	0.0	0.0	0.0	1.0	0.0	
2731	1.0	0.0	0.0	0.0	0.0	1.0	0.0	0.0	
4982	1.0	0.0	0.0	0.0	0.0	1.0	0.0	0.0	
					•••				
18368	1.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
7212	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	
13165	0.0	0.0	0.0	1.0	0.0	0.0	0.0	1.0	
17397	0.0	1.0	0.0	0.0	0.0	1.0	0.0	0.0	
20852	0.0	1.0	0.0	0.0	0.0	0.0	1.0	0.0	

6677 rows × 33 columns

localhost:8888/notebooks/DSC-phase-3-project/H1N1_and_seasonal_flu_vaccines_prediction.ipynb

In [25]: # Check for columns with all zeros to drop
pd.DataFrame([[column, encoded_categorical_X_test[column].sum() == 0] for column in encoded_categorical_X_test.columns

Out[25]:

	0	1
0	age_group_35 - 44 Years	False
1	age_group_45 - 54 Years	False
2	age_group_55 - 64 Years	False
3	age_group_65+ Years	False
4	education_< 12 Years	False
5	education_College Graduate	False
6	education_Some College	False
7	education_nan	False
8	race_Hispanic	False
9	race_Other or Multiple	False
10	race_White	False
11	sex_Male	False
12	income_poverty_> \$75,000	False
13	income_poverty_Below Poverty	False
14	income_poverty_nan	False
15	marital_status_Not Married	False
16	marital_status_nan	False
17	rent_or_own_Rent	False
18	rent_or_own_nan	False
19	employment_status_Not in Labor Force	False
20	employment_status_Unemployed	False
21	employment_status_nan	False
22	hhs_geo_region_bhuqouqj	False
23	hhs_geo_region_dqpwygqj	False
24	hhs_geo_region_fpwskwrf	False

```
0
                                          1
25
               hhs geo region kbazzjca False
                hhs geo region Irircsnp False
26
                hhs geo region Izgpxyit False
27
28
              hhs geo region mlyzmhmf False
               hhs geo region oxchigsf False
29
               hhs geo region qufhixun False
30
31
         census msa MSA, Principle City False
32
                 census msa Non-MSA False
```

```
In [26]: # Back concatenate numerical and categorical columns for both training and test data sets
# Training data set
X_train_clean = pd.concat([imputed_numerical_X_train, encoded_categorical_X_train], axis = 1)
```

```
In [27]: # Test data set
X_test_clean = pd.concat([imputed_numerical_X_test, encoded_categorical_X_test], axis = 1)
```

In [28]: X_train_clean.isna().sum()

Out[28]:	h1n1_concern	68
	h1n1_knowledge	88
	behavioral_antiviral_meds	52
	behavioral_avoidance	159
	behavioral_face_mask	18
	behavioral_wash_hands	35
	behavioral_large_gatherings	60
	behavioral_outside_home	59
	behavioral_touch_face	99
	doctor_recc_h1n1	1627 706
	<pre>chronic_med_condition child_under_6_months</pre>	602
	health worker	587
	opinion_h1n1_vacc_effective	276
	opinion_h1n1_risk	269
	opinion_h1n1_sick_from_vacc	278
	household_adults	171
	household_children	171
	age_group_35 - 44 Years	0
	age_group_45 - 54 Years	0
	age_group_55 - 64 Years	0
	age_group_65+ Years	0
	education_< 12 Years	0
	education_College Graduate	0
	education_Some College	0
	education_nan	0
	race_Hispanic	0
	race_Other or Multiple	0
	race_White	0
	sex_Male	0
	income_poverty_> \$75,000	0
	income_poverty_Below Poverty	0
	income_poverty_nan	0
	marital_status_Not Married	0
	marital_status_nan	0
	rent_or_own_Rent	0
	rent_or_own_nan	0
	employment_status_Not in Labor Force	0
	employment_status_Unemployed	0
	employment_status_nan	0
	hhs_geo_region_bhuqouqj	0

hhs_geo_region_dqpwygqj	0
hhs_geo_region_fpwskwrf	0
hhs_geo_region_kbazzjca	0
hhs_geo_region_lrircsnp	0
hhs_geo_region_lzgpxyit	0
hhs_geo_region_mlyzmhmf	0
hhs_geo_region_oxchjgsf	0
hhs_geo_region_qufhixun	0
census_msa_MSA, Principle City	0
census_msa_Non-MSA	0
dtype: int64	

In [29]: X_test_clean.isna().sum()

0+[20]+	h1n1	24
Out[29].	h1n1_concern	24
	h1n1_knowledge	28
	<pre>behavioral_antiviral_meds behavioral avoidance</pre>	19 49
	behavioral_avoluance behavioral face mask	49
	– –	
	behavioral_wash_hands	7
	behavioral_large_gatherings	27
	behavioral_outside_home	23
	behavioral_touch_face	29
	doctor_recc_h1n1	533
	chronic_med_condition	265
	child_under_6_months	218
	health_worker	217
	opinion_h1n1_vacc_effective	115
	opinion_h1n1_risk	119
	opinion_h1n1_sick_from_vacc	117
	household_adults	78
	household_children	78
	age_group_35 - 44 Years	0
	age_group_45 - 54 Years	0
	age_group_55 - 64 Years	0
	age_group_65+ Years	0
	education_< 12 Years	0
	education_College Graduate	0
	education_Some College	0
	education_nan	0
	race_Hispanic	0
	race_Other or Multiple	0
	race_White	0
	sex_Male	0
	income_poverty_> \$75,000	0
	income_poverty_Below Poverty	0
	income_poverty_nan	0
	marital_status_Not Married	0
	marital_status_nan	0
	rent_or_own_Rent	0
	rent_or_own_nan	0
	<pre>employment_status_Not in Labor Force</pre>	0
	<pre>employment_status_Unemployed</pre>	0
	employment_status_nan	0
	hhs_geo_region_bhuqouqj	0

hhs_geo_region_dqpwygqj	0
hhs_geo_region_fpwskwrf	0
hhs_geo_region_kbazzjca	0
hhs_geo_region_lrircsnp	0
hhs_geo_region_lzgpxyit	0
hhs_geo_region_mlyzmhmf	0
hhs_geo_region_oxchjgsf	0
hhs_geo_region_qufhixun	0
census_msa_MSA, Principle City	0
census_msa_Non-MSA	0
dtype: int64	

```
In [30]: # Impute more missing values on X_train data set
X_train = imputer.fit_transform(X_train_clean)
X_train = pd.DataFrame(np.round(X_train, 0), columns = X_train_clean.columns, index = X_train_clean.index)
X_train
```

Out[30]:

	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands bel
respondent_id						
7191	1.0	1.0	0.0	1.0	0.0	1.0
23784	3.0	1.0	0.0	1.0	0.0	1.0
316	2.0	1.0	0.0	1.0	0.0	1.0
521	2.0	1.0	0.0	1.0	0.0	1.0
21647	2.0	2.0	0.0	1.0	0.0	1.0
21810	2.0	1.0	0.0	1.0	0.0	0.0
23992	1.0	0.0	0.0	0.0	0.0	0.0
14501	2.0	1.0	0.0	1.0	0.0	1.0
14555	2.0	1.0	0.0	0.0	0.0	1.0
11590	1.0	2.0	0.0	1.0	0.0	1.0

20030 rows × 51 columns

localhost:8888/notebooks/DSC-phase-3-project/H1N1_and_seasonal_flu_vaccines_prediction.ipynb

```
In [31]: # Impute more missing values on X_test data set
X_test = imputer.fit_transform(X_test_clean)
X_test = pd.DataFrame(np.round(X_test, 0), columns = X_test_clean.columns, index= X_test_clean.index)
X_test
```

Out[31]:

		h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands	bel
respo	ndent_id							
	6458	1.0	0.0	0.0	1.0	0.0	1.0	
	12839	2.0	1.0	0.0	1.0	0.0	1.0	
	4377	3.0	2.0	0.0	0.0	0.0	1.0	
	2731	2.0	1.0	0.0	1.0	0.0	1.0	
	4982	3.0	2.0	0.0	1.0	0.0	1.0	
	18368	1.0	1.0	0.0	1.0	0.0	1.0	
	7212	2.0	0.0	0.0	1.0	0.0	1.0	
	13165	2.0	0.0	0.0	0.0	0.0	1.0	
	17397	3.0	2.0	1.0	1.0	1.0	1.0	
	20852	3.0	2.0	0.0	1.0	0.0	1.0	

6677 rows × 51 columns

In [32]: X_train.isna().sum()

Out[32]:	h1n1_concern	0
	h1n1_knowledge	0
	behavioral_antiviral_meds	0
	behavioral avoidance	0
	behavioral_face_mask	0
	behavioral_wash_hands	0
	behavioral_large_gatherings	0
	behavioral_outside_home	0
	behavioral_touch_face	0
	doctor_recc_h1n1	0
	chronic_med_condition	0
	child_under_6_months	0
	health_worker	0
	opinion_h1n1_vacc_effective	0
	opinion_h1n1_risk	0
	opinion_h1n1_sick_from_vacc	0
	household_adults	0
	household_children	0
	age_group_35 - 44 Years	0
	age_group_45 - 54 Years	0
	age_group_55 - 64 Years	0
	age_group_65+ Years	0
	education_< 12 Years	0
	education_College Graduate	0
	education_Some College	0
	education_nan	0
	race_Hispanic	0
	race_Other or Multiple	0
	race_White	0
	sex_Male	0
	income_poverty_> \$75,000	0
	income_poverty_Below Poverty	0
	income_poverty_nan	0
	marital_status_Not Married	0
	marital_status_nan	0
	rent_or_own_Rent	0
	rent_or_own_nan	0
	employment_status_Not in Labor Force	0
	employment_status_Unemployed	0
	employment_status_nan	0
	hhs_geo_region_bhuqouqj	0

hhs_geo_region_dqpwygqj	0
hhs_geo_region_fpwskwrf	0
hhs_geo_region_kbazzjca	0
hhs_geo_region_lrircsnp	0
hhs_geo_region_lzgpxyit	0
hhs_geo_region_mlyzmhmf	0
hhs_geo_region_oxchjgsf	0
hhs_geo_region_qufhixun	0
census_msa_MSA, Principle City	0
census_msa_Non-MSA	0
dtype: int64	

In [33]: X_test.isna().sum()

Out[33]:	h1n1_concern	0
	h1n1_knowledge	0
	behavioral_antiviral_meds	0
	behavioral_avoidance	0
	behavioral_face_mask	0
	behavioral_wash_hands	0
	behavioral_large_gatherings	0
	behavioral_outside_home	0
	behavioral_touch_face	0
	doctor_recc_h1n1	0
	<pre>chronic_med_condition</pre>	0
	child_under_6_months	0
	health_worker	0
	<pre>opinion_h1n1_vacc_effective</pre>	0
	opinion_h1n1_risk	0
	opinion_h1n1_sick_from_vacc	0
	household_adults	0
	household_children	0
	age_group_35 - 44 Years	0
	age_group_45 - 54 Years	0
	age_group_55 - 64 Years	0
	age_group_65+ Years	0
	education_< 12 Years	0
	education_College Graduate	0
	education_Some College	0
	education_nan	0
	race_Hispanic	0
	race_Other or Multiple	0
	race_White	0
	sex_Male	0
	income_poverty_> \$75,000	0
	<pre>income_poverty_Below Poverty</pre>	0
	<pre>income_poverty_nan</pre>	0
	marital_status_Not Married	0
	marital_status_nan	0
	rent_or_own_Rent	0
	rent_or_own_nan	0
	<pre>employment_status_Not in Labor Force</pre>	0
	employment_status_Unemployed	0
	employment_status_nan	0
	hhs_geo_region_bhuqouqj	0

```
hhs_geo_region_dqpwygqj 0
hhs_geo_region_fpwskwrf 0
hhs_geo_region_kbazzjca 0
hhs_geo_region_lrircsnp 0
hhs_geo_region_lzgpxyit 0
hhs_geo_region_mlyzmhmf 0
hhs_geo_region_oxchjgsf 0
hhs_geo_region_qufhixun 0
census_msa_MSA, Principle City 0
dtype: int64
```

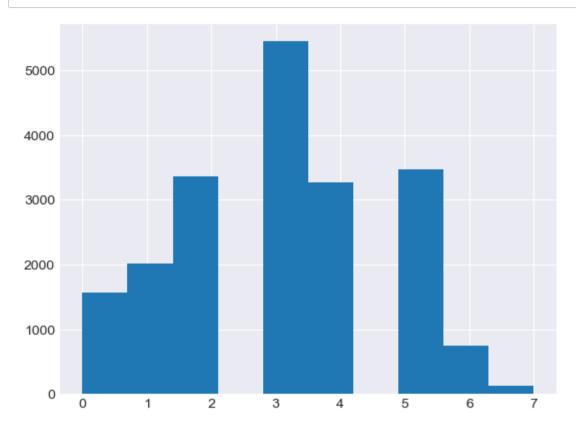
Feature Engineering

i) behavior score

• By adding up all the behavioral variables, create a variable that shows how much a person has done behaviorally to avoid the virus, aside from getting vaccinated. These are all binary columns, where **1** denotes **YES**, indicating that the individual has taken a step to lower their risk of getting the flu. A higher score, calculated by adding the values of these columns, indicates a more circumspect and flu-conscious person.

```
In [35]: # Create the `behavior score` variable for X train data set
         X train['behavior score'] = X train[behavior cols].sum(axis=1)
In [36]: # Create the `behavior score` variable for X test data set
         X test['behavior score'] = X test[behavior cols].sum(axis=1)
In [37]: X test.columns
Out[37]: Index(['h1n1 concern', 'h1n1 knowledge', 'behavioral antiviral meds',
                 'behavioral avoidance', 'behavioral face mask', 'behavioral wash hands',
                 'behavioral large gatherings', 'behavioral outside home',
                 'behavioral touch face', 'doctor recc h1n1', 'chronic med condition',
                 'child under 6 months', 'health worker', 'opinion h1n1 vacc effective',
                 'opinion h1n1 risk', 'opinion h1n1 sick from vacc', 'household adults',
                 'household children', 'age group 35 - 44 Years',
                 'age group 45 - 54 Years', 'age group 55 - 64 Years',
                 'age group 65+ Years', 'education < 12 Years',
                 'education College Graduate', 'education Some College', 'education nan',
                 'race Hispanic', 'race Other or Multiple', 'race White', 'sex Male',
                 'income poverty > $75,000', 'income poverty Below Poverty',
                 'income poverty nan', 'marital status Not Married',
                 'marital status nan', 'rent or own Rent', 'rent or own nan',
                 'employment status Not in Labor Force', 'employment status Unemployed',
                 'employment status nan', 'hhs geo region bhuqouqj',
                 'hhs geo region dqpwygqj', 'hhs geo region fpwskwrf',
                 'hhs geo region kbazzjca', 'hhs geo region lrircsnp',
                 'hhs geo region lzgpxyit', 'hhs geo region mlyzmhmf',
                 'hhs geo region_oxchjgsf', 'hhs_geo_region_qufhixun',
                 'census msa MSA, Principle City', 'census msa Non-MSA',
                 'behavior score'],
               dtype='object')
```

```
In [38]: # Plot the distribution of behavior score variable we created
plt.hist(X_train['behavior_score']);
```



Class Imbalance

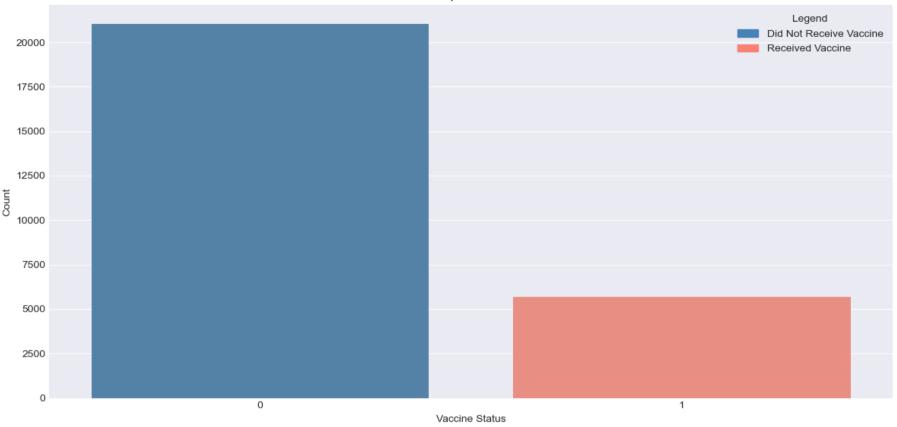
```
In [39]: # Checking for class distribution
fig, ax = plt.subplots(figsize=(12, 6))
sns.countplot(x='hln1_vaccine', data=merged_df, ax=ax, palette=["steelblue", "salmon"])

ax.set_title('HlN1 Vaccine Uptake Class Distribution')
ax.set_xlabel('Vaccine Status')
ax.set_ylabel('Count')

# Add separate Legend Labels for each class
legend_labels = ['Did Not Receive Vaccine', "Received Vaccine"]
colors = ["steelblue", "salmon"]
legend_handles = [plt.Rectangle((0, 0), 1, 1, color=color) for color in colors]
ax.legend(legend_handles, legend_labels, title='Legend')

plt.tight_layout()
plt.savefig('Visualization1')
```

H1N1 Vaccine Uptake Class Distribution



• This plot shows the uptake distribution between those who received the vaccine and those who did not receive the vaccine

```
In [40]: # Checking class imbalance for h1n1_vaccine [target variable]
y.value_counts()
```

Out[40]: h1n1_vaccine 0 21033 1 5674

Name: count, dtype: int64

Presence of class imbalance

Exploratory Data Analysis

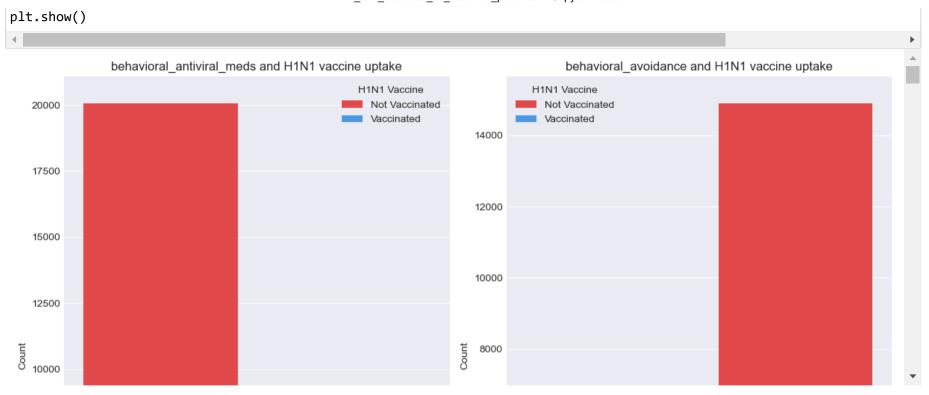
15779

Name: count, dtype: int64

Checking the relationshop between our target variable **h1n1_vaccine** and the features.

```
In [42]: print(merged_df['h1n1_vaccine'].unique())
      [0 1]
```

```
In [43]: # List of columns to plot
         columns = ['behavioral antiviral meds', 'behavioral avoidance',
                   'behavioral face mask', 'behavioral wash hands', 'behavioral large gatherings',
                   'behavioral outside home', 'behavioral touch face',
                    'doctor recc h1n1', 'chronic med condition', 'child under 6 months',
                     'health worker'.
                    'opinion h1n1 vacc effective', 'opinion h1n1 risk', 'opinion h1n1 sick from vacc',
                    'age group', 'education', 'race', 'sex', 'income poverty', 'marital status',
                    'rent or own', 'employment status', 'household adults', 'household children'l
         # Select columns for plotting
         columns to plot = columns[:25]
         # Determine the number of rows and columns for subplots
         num rows = (len(columns to plot) + 1) // 2
         num cols = 2
         # Create subplots
         fig, axes = plt.subplots(num rows, num cols, figsize=(12, 8 * num rows))
         axes = axes.flatten()
         # Iterate over the columns and create count plots
         for i, column in enumerate(columns to plot):
             # Check if the 'h1n1 vaccine' column is present in the DataFrame
             if 'h1n1 vaccine' in merged df.columns:
                 unique values = merged df[column].unique()
                 if len(unique values) > 1: # Proceed only if there are multiple unique values
                     sns.countplot(x=column, data=merged df[merged df['h1n1 vaccine'] == 0], ax=axes[i], color='red', label='Non
                     sns.countplot(x=column, data=merged df[merged df['h1n1 vaccine'] == 1], ax=axes[i], color='dodgerblue', lal
                     axes[i].set title(f"{column} and H1N1 vaccine uptake")
                     axes[i].set xlabel('')
                     axes[i].set ylabel('Count')
                     axes[i].legend(title='H1N1 Vaccine', loc='best')
                 else:
                     axes[i].set title(f"{column} (only one value)")
                     axes[i].set xlabel('')
                     axes[i].set vlabel('Count')
         # Remove any empty subplots
         if len(columns to plot) % 2 == 1:
             fig.delaxes(axes[-1])
         # Adjust spacing between subplots
         plt.tight layout()
         # Display the plot
```



Modeling

Logistic Regression

```
In [44]: # Baseline model
         # Instantiate classifier
         baseline logreg = LogisticRegression(C= 1e12, random state=40, solver='liblinear')
         # Fit model to train set
         baseline logreg.fit(X train resample, y train resample)
Out[44]:
                                      LogisticRegression
         LogisticRegression(C=10000000000000.0, random state=40, solver='liblinear')
In [45]: # Accuracy on training data set
         baseline preds = baseline logreg.predict(X train resample)
         print(f"accuracy score: {accuracy score(y train resample, baseline preds)}")
         print(f"f1 score: {f1 score(y train resample, baseline preds)}")
         print(f"precision score: {precision score(y train resample, baseline preds)}")
         print(f"recall score: {recall score(y train resample, baseline preds)}")
         accuracy score: 0.7676658850370746
         f1 score: 0.7612970438859226
         precision score: 0.7827542344513624
         recall score: 0.740984853286013
```

```
In [46]: # Accuracy on test data set
         baseline preds = baseline logreg.predict(X test)
         print(f"accuracy score: {accuracy score(y test, baseline preds)}")
         print(f"f1 score: {f1 score(y test, baseline preds)}")
         print(f"precision score: {precision score(y test, baseline preds)}")
         print(f"recall score: {recall score(y test, baseline preds)}")
         accuracy score: 0.7753482102740752
         f1 score: 0.5774647887323944
         precision score: 0.48189938881053124
         recall score: 0.7203092059030218
In [47]: # Model2
         # Instantiate classifier
         logreg 2 = LogisticRegression(penalty = 'l1', C= 1e12, random state=40, solver='liblinear')
         # Fit model to train set
         logreg 2.fit(X train resample, y train resample)
Out[47]:
                                    LogisticRegression
         LogisticRegression(C=10000000000000.0, penalty='11', random state=40,
                             solver='liblinear')
In [48]: # Accuracy on training data set
         logreg2 preds = logreg 2.predict(X train resample)
         print(f"accuracy score: {accuracy score(y train resample, logreg2 preds)}")
         print(f"f1 score: {f1 score(y train resample, logreg2 preds)}")
         print(f"precision score: {precision score(y train resample, logreg2 preds)}")
         print(f"recall score: {recall score(y train resample, logreg2 preds)}")
         accuracy score: 0.7676975727232398
         f1 score: 0.76132182972489
         precision score: 0.7828066416711301
         recall score: 0.740984853286013
```

```
In [49]: # Accuracy on test data set
         logreg2 preds = logreg 2.predict(X test)
         print(f"accuracy score: {accuracy score(y test, logreg2 preds)}")
         print(f"f1 score: {f1 score(y test, logreg2 preds)}")
         print(f"precision score: {precision score(y test, logreg2 preds)}")
         print(f"recall score: {recall score(v test, logreg2 preds)}")
         accuracy score: 0.775198442414258
         f1 score: 0.5773021684032668
         precision score: 0.4816729323308271
         recall score: 0.7203092059030218
In [50]: # Model 3
         # Instantiate classifier
         logreg 3 = LogisticRegression(penalty = '12', random state=40, solver='lbfgs')
         # Fit model to train set
         logreg 3.fit(X train resample, y train resample)
         C:\Users\Richard.LAPTOP-2023AAHO\anaconda3\Lib\site-packages\sklearn\linear model\ logistic.py:458: ConvergenceWarnin
         g: lbfgs failed to converge (status=1):
         STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.
         Increase the number of iterations (max iter) or scale the data as shown in:
             https://scikit-learn.org/stable/modules/preprocessing.html (https://scikit-learn.org/stable/modules/preprocessin
         g.html)
         Please also refer to the documentation for alternative solver options:
             https://scikit-learn.org/stable/modules/linear model.html#logistic-regression (https://scikit-learn.org/stable/mo
         dules/linear model.html#logistic-regression)
           n iter i = check optimize result(
Out[50]:
                   LogisticRegression
         LogisticRegression(random state=40)
```

```
In [51]: # Accuracy on training data set
         logreg3 preds = logreg 3.predict(X train resample)
         print(f"accuracy score: {accuracy score(y train resample, logreg3 preds)}")
         print(f"f1 score: {f1 score(v train resample, logreg3 preds)}")
         print(f"precision score: {precision score(y train resample, logreg3 preds)}")
         print(f"recall score: {recall score(y train resample, logreg3 preds)}")
         accuracy score: 0.7677926357817352
         f1 score: 0.7613651165819982
         precision score: 0.7830397213477125
         recall score: 0.7408581025413524
In [52]: # Accuracy on test data set
         logreg3 preds = logreg 3.predict(X test)
         print(f"accuracy score: {accuracy score(y test, logreg3 preds)}")
         print(f"f1 score: {f1 score(y test, logreg3 preds)}")
         print(f"precision score: {precision score(y test, logreg3 preds)}")
         print(f"recall score: {recall score(y test, logreg3 preds)}")
         accuracy score: 0.7754979781338924
         f1 score: 0.5776275007044238
         precision score: 0.4821260583254939
         recall score: 0.7203092059030218
In [53]: # ROC curve for the best Logistic regression model
         fpr logreg, tpr logreg, = roc curve(y test, logreg 3.predict proba(X test)[:,1])
```

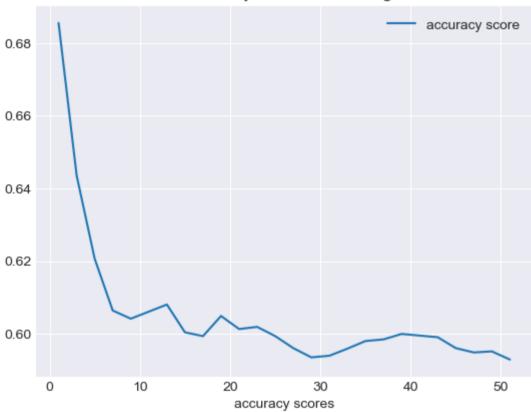
K-Nearest Neighbours

```
In [54]: # Baseline model
         # Instantiate classifier
         baseline KNN clf = KNeighborsClassifier()
         # Fit model to train set
         baseline KNN clf.fit(X train resample, y train resample)
Out[54]:
          ▼ KNeighborsClassifier
         KNeighborsClassifier()
In [55]: # Accuracy on training data set
         baseline preds = baseline KNN clf.predict(X train resample)
         print(f"accuracy score: {accuracy score(y train resample, baseline preds)}")
         print(f"f1 score: {f1 score(y train resample, baseline preds)}")
         print(f"precision score: {precision score(y train resample, baseline preds)}")
         print(f"recall score: {recall score(y train resample, baseline preds)}")
         accuracy score: 0.8445085239875785
         f1 score: 0.8642787996127783
         precision score: 0.7667844522968198
         recall score: 0.9901768172888016
```

```
In [56]: # Accuracy on test data set
         baseline preds = baseline KNN clf.predict(X test)
         print(f"accuracy score: {accuracy score(y test, baseline preds)}")
         print(f"f1 score: {f1 score(y test, baseline preds)}")
         print(f"precision score: {precision score(y test, baseline preds)}")
         print(f"recall score: {recall score(v test, baseline preds)}")
         accuracy score: 0.6206380110828216
         f1 score: 0.4577178334403768
         precision score: 0.3291256157635468
         recall score: 0.7512297962052003
In [57]: # Find best n neighbors parameter
         n neighbors params = []
         accuracy scores = []
         for n in np.arange(1, 52, 2):
             n neighbors params.append(n)
             clf model = KNeighborsClassifier(n neighbors = n)
             clf model.fit(X train resample, y train resample)
             acc = accuracy score(y test, clf model.predict(X test))
             accuracy scores.append(acc)
```

```
In [58]: plt.plot(n_neighbors_params, accuracy_scores, label ='accuracy score')
    plt.xlabel('n_neighbors params values')
    plt.xlabel('accuracy scores')
    plt.title('Model accuracy vs Number of neighbors')
    plt.legend();
```

Model accuracy vs Number of neighbors



```
In [59]: # Scale training and test data sets
         scaler = StandardScaler()
         X train resample scaled = scaler.fit transform(X train resample)
         X test scaled = scaler.transform(X test)
In [60]: # Model Iteration
         # Instantiate classifier
         KNN clf2 = KNeighborsClassifier()
         # Fit model to train set
         KNN clf2.fit(X train resample scaled, y train resample)
         # Accuracy of model on train set
         print(f"accuracy score: {accuracy score(y train resample, KNN clf2.predict(X train resample scaled))}")
         print(f"f1 score: {f1 score(y train resample, KNN clf2.predict(X train resample scaled))}")
         print(f"precision score: {precision score(y train resample, KNN clf2.predict(X train resample scaled))}")
         print(f"recall score: {recall score(v train resample, KNN clf2.predict(X train resample scaled))}")
         accuracy score: 0.8812979276253248
         f1 score: 0.8923129994825505
         precision score: 0.8165412742673752
         recall score: 0.9835857785664491
In [61]: # Accuracy of model on test set
         print(f"accuracy score: {accuracy score(y test, KNN clf2.predict(X test scaled))}")
         print(f"f1 score: {f1 score(y test, KNN clf2.predict(X test scaled))}")
         print(f"precision score: {precision score(y test, KNN clf2.predict(X test scaled))}")
         print(f"recall score: {recall score(y test, KNN clf2.predict(X test scaled))}")
         accuracy score: 0.6601767260745844
         f1 score: 0.43175557225143996
         precision score: 0.335408560311284
         recall score: 0.6057624736472241
```

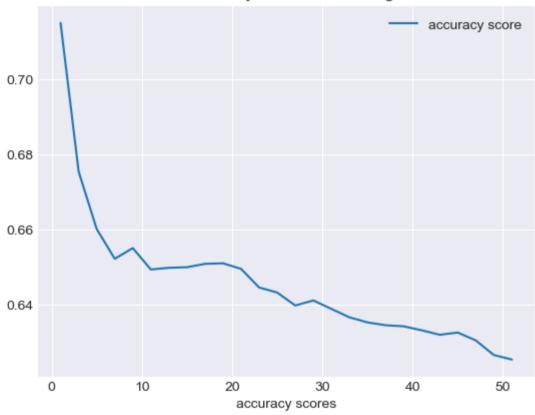
```
In [62]: # Find best n_neighbors parameter
    n_neighbors_params = []
    accuracy_scores = []

for n in np.arange(1, 52, 2):
    n_neighbors_params.append(n)

    clf_model = KNeighborsClassifier(n_neighbors = n)
    clf_model.fit(X_train_resample_scaled, y_train_resample)
    acc = accuracy_score(y_test, clf_model.predict(X_test_scaled))
    accuracy_scores.append(acc)
```

```
In [63]: plt.plot(n_neighbors_params, accuracy_scores, label ='accuracy score')
    plt.xlabel('n_neighbors params values')
    plt.xlabel('accuracy scores')
    plt.title('Model accuracy vs Number of neighbors')
    plt.legend();
```

Model accuracy vs Number of neighbors



Feature selection

```
In [64]: # Define a custom function to compute feature importances
def custom_importance_getter_knn(estimator):
    # Extract the distances to the nearest neighbors
    distances, indices = estimator.kneighbors()
    # Compute the mean distance for each feature
    feature_importances = distances.mean(axis=0)
    return feature_importances

# Initialize RFE with the classifier and desired number of features to select
    rfe = RFE(estimator= KNN_clf2, n_features_to_select= .8, step= 1, importance_getter= custom_importance_getter_knn) # .

# Fit RFE on train data set
    rfe.fit(X_train_resample_scaled, y_train_resample)

print(X_train_resample.columns)
    print(rfe.ranking_)
    print(rfe.support_)
```

```
Index(['h1n1 concern', 'h1n1 knowledge', 'behavioral antiviral meds',
              'behavioral avoidance', 'behavioral face mask', 'behavioral wash hands',
              'behavioral large gatherings', 'behavioral outside home',
              'behavioral touch face', 'doctor recc h1n1', 'chronic med condition',
              'child under 6 months', 'health worker', 'opinion h1n1 vacc effective',
              'opinion h1n1 risk', 'opinion h1n1 sick from vacc', 'household adults',
              'household children', 'age_group_35 - 44 Years',
              'age group 45 - 54 Years', 'age_group_55 - 64 Years',
              'age group 65+ Years', 'education < 12 Years',
              'education College Graduate', 'education Some College', 'education nan',
              'race Hispanic', 'race Other or Multiple', 'race White', 'sex Male',
              'income poverty > $75,000', 'income poverty Below Poverty',
              'income poverty nan', 'marital status Not Married',
              'marital status nan', 'rent or own Rent', 'rent or own nan',
              'employment status Not in Labor Force', 'employment status Unemployed',
              'employment status nan', 'hhs geo region bhuqouqj',
              'hhs geo region dapwygaj', 'hhs geo region fpwskwrf',
              'hhs geo region kbazzjca', 'hhs geo region lrircsnp',
              'hhs geo region lzgpxyit', 'hhs geo region mlyzmhmf',
              'hhs geo region oxchigsf', 'hhs geo region qufhixun',
              'census msa MSA, Principle City', 'census msa Non-MSA',
              'behavior score'],
             dtvpe='object')
        [12 11 10 9 8 7 6 5 4 3 2 1 1 1 1 1 1 1 1 1 1 1 1
         1 1 1 1]
        [False False False False False False False False False False True
         True True True Truel
In [65]: masked columns = X train resample.columns[rfe.get support(indices= True)]
In [66]: X train rfe = X train resample scaled[:,0:11]
       X test rfe = X test scaled[:,0:11]
```

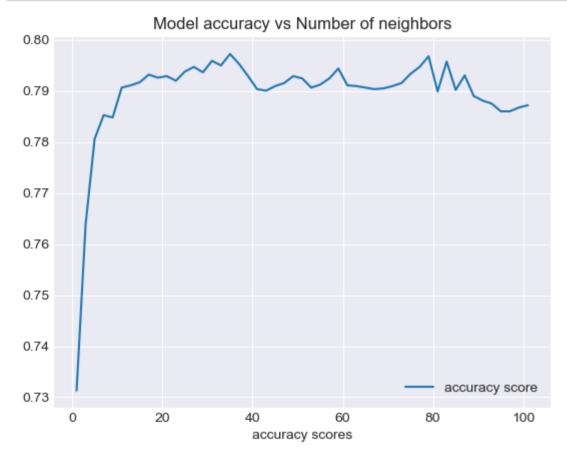
```
In [67]: # Instantiate classifier
         KNN clf3 = KNeighborsClassifier()
         # Fit model to train set
         KNN clf3.fit(X train rfe, y train resample)
Out[67]:
          ▼ KNeighborsClassifier
          KNeighborsClassifier()
In [68]: # Accuracy of model on train set
         print(f"accuracy score: {accuracy score(y train resample, KNN clf3.predict(X train rfe))}")
         print(f"f1 score: {f1 score(y train resample, KNN clf3.predict(X train rfe))}")
         print(f"precision score: {precision_score(y_train_resample, KNN clf3.predict(X train rfe))}")
         print(f"recall score: {recall score(y train resample, KNN clf3.predict(X train rfe))}")
         accuracy score: 0.8234045250015843
         f1 score: 0.8061497791227521
         precision score: 0.8934464148033925
         recall score: 0.7343938145636606
In [69]: # Accuracy of model on test set
         print(f"accuracy score: {accuracy score(y test, KNN clf3.predict(X test rfe))}")
         print(f"f1 score: {f1 score(y test, KNN clf3.predict(X test rfe))}")
         print(f"precision score: {precision score(y test, KNN clf3.predict(X test rfe))}")
         print(f"recall score: {recall score(y test, KNN clf3.predict(X test rfe))}")
         accuracy score: 0.78059008536768
         f1 score: 0.4104627766599598
         precision score: 0.480225988700565
         recall score: 0.3583977512297962
```

```
In [70]: # Find best n_neighbors parameter
    n_neighbors_params = []
    accuracy_scores = []

for n in np.arange(1, 102, 2):
    n_neighbors_params.append(n)

    clf_model = KNeighborsClassifier(n_neighbors = n)
    clf_model.fit(X_train_rfe, y_train_resample)
    acc = accuracy_score(y_test, clf_model.predict(X_test_rfe))
    accuracy_scores.append(acc)
```

```
In [71]: plt.plot(n_neighbors_params, accuracy_scores, label ='accuracy score')
    plt.xlabel('n_neighbors params values')
    plt.xlabel('accuracy scores')
    plt.title('Model accuracy vs Number of neighbors')
    plt.legend();
```



```
In [72]: # Best n_neighbors parameter is 51
index = accuracy_scores.index(max(accuracy_scores))
n_neighbors_params[index]
```

Out[72]: 35

```
In [73]: # Instantiate classifier
         KNN clf final = KNeighborsClassifier(n neighbors = 51)
         # Fit model to train set
         KNN clf final.fit(X train rfe, y train resample)
Out[73]:
                  KNeighbor Classifier
         KNeighborsClassifier(n neighbors=51)
In [74]: # Accuracy of model on train set
         print(f"accuracy score on train set: {accuracy score(y train resample, KNN clf final.predict(X train rfe))}")
         print(f"f1 score on train set: {f1 score(y train resample, KNN clf final.predict(X train rfe))}")
         print(f"precision score on train set: {precision score(y train resample, KNN clf final.predict(X train rfe))}")
         print(f"recall score on train set: {recall score(y train resample, KNN clf final.predict(X train rfe))}")
         print("\n----\\n")
         # Accuracy of model on test set
         print(f"accuracy score on test set: {accuracy score(y test, KNN clf final.predict(X test rfe))}")
         print(f"f1 score on test set: {f1 score(y test, KNN clf final.predict(X test rfe))}")
         print(f"precision score on test set: {precision score(y test, KNN clf final.predict(X test rfe))}")
         print(f"recall score on test set: {recall score(y test, KNN clf final.predict(X test rfe))}")
         accuracy score on train set: 0.7502059699600735
         f1 score on train set: 0.7136890277121999
         precision score on train set: 0.8358856559469117
         recall score on train set: 0.6226630331453197
         accuracy score on test set: 0.7924217462932455
         f1 score on test set: 0.49416058394160584
         precision score on test set: 0.5140470766894457
         recall score on test set: 0.47575544624033733
```

Cross validation

Naive Bayes

```
In [79]: # Accuracy in training data
print(f"accuracy score: {accuracy_score(y_train_resample, baseline_nbayes_clf.predict(X_train_resample_scaled))}")
print(f"f1 score: {f1_score(y_train_resample, baseline_nbayes_clf.predict(X_train_resample_scaled))}")
print(f"precision score: {precision_score(y_train_resample, baseline_nbayes_clf.predict(X_train_resample_scaled))}")
print(f"recall score: {recall_score(y_train_resample, baseline_nbayes_clf.predict(X_train_resample_scaled))}")
```

accuracy score: 0.6946257684263895 f1 score: 0.7131759873805769 precision score: 0.672334455667789 recall score: 0.7593003358894733

In [80]: # Accuracy in testing data print(f"accuracy score: {accuracy_score(y_test, baseline_nbayes_clf.predict(X_test_scaled))}") print(f"f1 score: {f1_score(y_test, baseline_nbayes_clf.predict(X_test_scaled))}") print(f"precision score: {precision_score(y_test, baseline_nbayes_clf.predict(X_test_scaled))}") print(f"recall score: {recall_score(y_test, baseline_nbayes_clf.predict(X_test_scaled))}")

accuracy score: 0.6350157256252809 f1 score: 0.44169530355097364

precision score: 0.32766825288919105
recall score: 0.6774420238931834

Feature Selection

```
In [81]: def custom_importance_getter_gnb(estimator, X_train, y_train):
    # Fit the Gaussian Naive Bayes estimator
    estimator.fit(X_train, y_train)
    # Compute mutual information between each feature and the target variable
    feature_importances = mutual_info_classif(X_train, y_train)
    return feature_importances
```

```
In [83]: # Fit the Gaussian Naive Bayes estimator
    estimator = GaussianNB()

# Compute feature importances using the custom function
    feature_importances = custom_importance_getter_gnb(estimator, X_train_resample_scaled, y_train_resample)

# Get the indices of features sorted by importance
    sorted_indices = feature_importances.argsort()[::-1]

# Get feature names
    feature_names = X_train_resample.columns

# Get feature names and their rankings
    feature_rankings = [[feature_names[i], rank + 1] for rank, i in enumerate(sorted_indices)]

print("Feature rankings:")
    pd.DataFrame(feature_rankings, columns=['Feature', 'Rank'])
```

Feature rankings:

Out[83]:

Feature	Rank			
opinion_h1n1_vacc_effective				
doctor_recc_h1n1				
opinion_h1n1_risk				
h1n1_concern				
behavior_score				
h1n1_knowledge				
opinion_h1n1_sick_from_vacc				
sex_Male				
household_adults				
chronic_med_condition				
census_msa_Non-MSA				
education_Some College				
education_College Graduate				
census_msa_MSA, Principle City				
income_poverty_> \$75,000				
employment_status_Not in Labor Force	16			
household_children	17			
health_worker	18			
marital_status_Not Married				
hhs_geo_region_lzgpxyit				
rent_or_own_Rent				
race_White	22			
age_group_45 - 54 Years	23			
behavioral_touch_face	24			
age_group_55 - 64 Years	25			
	opinion_h1n1_vacc_effective			

	Feature	Rank			
25	behavioral_avoidance	26			
26	hhs_geo_region_qufhixun	27			
27	hhs_geo_region_bhuqouqj	28			
28	behavioral_large_gatherings	29			
29	behavioral_wash_hands	30			
30	hhs_geo_region_fpwskwrf				
31	behavioral_outside_home				
32	child_under_6_months	33			
33	hhs_geo_region_kbazzjca	34			
34	age_group_65+ Years	35			
35	hhs_geo_region_oxchjgsf	36			
36	hhs_geo_region_mlyzmhmf	37			
37	income_poverty_nan	38			
38	income_poverty_Below Poverty	39			
39	age_group_35 - 44 Years	40			
40	hhs_geo_region_lrircsnp	41			
41	education_< 12 Years	42			
42	behavioral_face_mask	43			
43	race_Hispanic	44			
44	race_Other or Multiple	45			
45	employment_status_Unemployed	46			
46	behavioral_antiviral_meds	47			
47	hhs_geo_region_dqpwygqj	48			
48	rent_or_own_nan	49			
49	education_nan	50			
50	employment_status_nan	51			

	Feature	Rank
51	marital_status_nan	52

```
In [84]: feature_rankings = pd.DataFrame(feature_rankings, columns=['Feature', 'Rank'])
    select_columns = feature_rankings['Feature'][0:10]
    X_train_resample[select_columns]
```

Out[84]:

	opinion_h1n1_vacc_effective	doctor_recc_h1n1	opinion_h1n1_risk	h1n1_concern	behavior_score	h1n1_knowledge	opinion_h1n1_sick_from_
0	4.000000	0.000000	2.0	1.000000	5.000000	1.000000	4.00
1	4.000000	1.000000	4.0	3.000000	3.000000	1.000000	4.00
2	3.000000	0.000000	3.0	2.000000	4.000000	1.000000	2.00
3	4.000000	0.000000	2.0	2.000000	4.000000	1.000000	4.00
4	5.000000	0.000000	2.0	2.000000	3.000000	2.000000	2.00
31553	5.000000	0.165731	4.0	2.000000	4.000000	1.834269	1.16
31554	4.000000	0.000000	4.0	2.324656	6.000000	2.000000	5.00
31555	4.640571	0.000000	4.0	2.000000	2.640571	2.000000	2.00
31556	4.562608	0.000000	2.0	2.000000	2.437392	2.000000	1.56
31557	5.000000	0.000000	4.0	2.000000	4.000000	2.000000	1.00

31558 rows × 10 columns

```
In [86]: n_bayes_clf2 = GaussianNB()
# Fit model
n_bayes_clf2.fit(X_train_resample_scaled_select_columns, y_train_resample)
```

Out[86]: ▼ GaussianNB GaussianNB()

```
In [87]: # Accuracy in training data
print(f"accuracy score: {accuracy_score(y_train_resample, n_bayes_clf2.predict(X_train_resample_scaled_select_columns)
print(f"f1 score: {f1_score(y_train_resample, n_bayes_clf2.predict(X_train_resample_scaled_select_columns))}")
print(f"precision score: {precision_score(y_train_resample, n_bayes_clf2.predict(X_train_resample_scaled_select_columns))}")
print(f"recall score: {recall_score(y_train_resample, n_bayes_clf2.predict(X_train_resample_scaled_select_columns))}")
```

accuracy score: 0.7465301983649154 f1 score: 0.7415425377233514

precision score: 0.7564271588661833
recall score: 0.7272323974903353

```
In [88]: # Accuracy in testing data
    print(f"accuracy score: {accuracy_score(y_test, n_bayes_clf2.predict(X_test_scaled_select_columns))}")
    print(f"f1 score: {f1_score(y_test, n_bayes_clf2.predict(X_test_scaled_select_columns))}")
    print(f"precision score: {precision_score(y_test, n_bayes_clf2.predict(X_test_scaled_select_columns))}")
    print(f"recall score: {recall_score(y_test, n_bayes_clf2.predict(X_test_scaled_select_columns))}")
```

accuracy score: 0.7516848884229445

f1 score: 0.553821313240043

precision score: 0.44875708678587006 recall score: 0.7231201686577653

Cross validate

```
In [90]: print(f"test f1 score with cross validation: {np.mean(results['test_accuracy'])}")
    print(f"test accuracy with cross validation: {np.mean(results['test_accuracy'])}")
    print(f"test precision with cross validation: {np.mean(results['test_precision'])}")
    print(f"test recall with cross validation: {np.mean(results['test_recall'])}")

    test f1 score with cross validation: 0.7394549040486306
    test accuracy with cross validation: 0.7445341979391407
    test precision with cross validation: 0.7543987747120069
    test recall with cross validation: 0.72520555016581
In [91]: # ROC curve for the best Gausssian Naive Bayes model
    fpr_gnbayes, tpr_gnbayes, _ = roc_curve(y_test, n_bayes_clf2.predict_proba(X_test_scaled_select_columns)[:,1])
```

Decision Tree

```
In [92]: # Initialize Decision Tree classifier
dt_classifier = DecisionTreeClassifier(random_state=40)

# Train the model
dt_classifier.fit(X_train_resample, y_train_resample)

# Make predictions on the testing set
y_pred = dt_classifier.predict(X_test)

# Evaluate the model
accuracy = accuracy_score(y_test, y_pred)
print("Accuracy:", accuracy)

# Classification report
print("Classification Report:")
print(classification_report(y_test, y_pred))

# Confusion matrix
print("Confusion_matrix:")
print(confusion_matrix(y_test, y_pred))
```

Accuracy: 0.7397034596375618

Classification Report:

	precision	recall	f1-score	support
0	0.85	0.82	0.83	5254
1	0.40	0.45	0.43	1423
accuracy			0.74	6677
macro avg	0.62	0.64	0.63	6677
weighted avg	0.75	0.74	0.75	6677

Confusion Matrix:

[[4295 959]

[779 644]]

```
In [93]: # Define the hyperparameters grid
         param grid = {
             'criterion': ['gini', 'entropy'],
             'max depth': [None, 10, 20, 30, 40, 50],
             'min_samples_split': [2, 5, 10],
             'min samples leaf': [1, 2, 4]
         # Initialize the decision tree classifier
         dt classifier = DecisionTreeClassifier(random state=40)
         # Initialize GridSearchCV
         grid search = GridSearchCV(estimator=dt classifier, param grid=param grid, cv=5, scoring='accuracy')
         # Perform grid search
         grid search.fit(X train resample, y train resample)
         # Get the best parameters and best score
         best params = grid search.best params
         best score = grid search.best score
         print("Best Parameters:", best params)
         print("Best Score:", best score)
```

Best Parameters: {'criterion': 'entropy', 'max_depth': 20, 'min_samples_leaf': 4, 'min_samples_split': 2}
Best Score: 0.8422947921819166

```
In [94]: # Initialize the decision tree classifier with best parameters
         best dt classifier = DecisionTreeClassifier(criterion='entropy', max depth=20, min samples leaf=4, min samples split=2
         # Train the model with best parameters
         best dt classifier.fit(X train resample, y train resample)
         # Make predictions on the testing set
         v pred best = best dt classifier.predict(X test)
         # Evaluate the model with best parameters
         accuracy best = accuracy score(y test, y pred best)
         print("Accuracy with best parameters:", accuracy best)
         # Classification report
         print("Classification Report with best parameters:")
         print(classification report(y test, y pred best))
         # Confusion matrix
         print("Confusion Matrix with best parameters:")
         print(confusion matrix(y test, y pred best))
         Accuracy with best parameters: 0.7795417103489591
         Classification Report with best parameters:
```

	precision	recall	f1-score	support
0	0.85	0.88	0.86	5254
1	0.48	0.42	0.45	1423
accuracy			0.78	6677
macro avg	0.66	0.65	0.65	6677
weighted avg	0.77	0.78	0.77	6677

Confusion Matrix with best parameters: [[4611 643] [829 594]]

Feature selection

```
In [95]: # Initialize RFE with the classifier and desired number of features to select
        rfe = RFE(estimator= best dt classifier, n features to select= .8, step= 1) # Select top 80% features
        # Fit RFE on train data set
        rfe.fit(X train resample, v train resample)
        print(X train resample.columns)
        print(rfe.ranking )
        print(rfe.support )
        Index(['h1n1 concern', 'h1n1 knowledge', 'behavioral antiviral meds',
               'behavioral avoidance', 'behavioral face mask', 'behavioral wash hands',
               'behavioral large gatherings', 'behavioral outside home',
               'behavioral touch face', 'doctor recc h1n1', 'chronic med condition',
               'child under 6 months', 'health worker', 'opinion h1n1 vacc effective',
               'opinion h1n1 risk', 'opinion h1n1 sick from vacc', 'household adults',
               'household children', 'age group 35 - 44 Years',
               'age group 45 - 54 Years', 'age group 55 - 64 Years',
               'age group 65+ Years', 'education < 12 Years',
               'education College Graduate', 'education Some College', 'education nan',
               'race Hispanic', 'race Other or Multiple', 'race White', 'sex Male',
               'income poverty > $75,000', 'income poverty Below Poverty',
               'income poverty nan', 'marital status Not Married',
               'marital status nan', 'rent or own Rent', 'rent or own nan',
               'employment status Not in Labor Force', 'employment status Unemployed',
               'employment status nan', 'hhs geo region bhuqouqj',
               'hhs geo region dapwygaj', 'hhs geo region fpwskwrf',
               'hhs geo region kbazzjca', 'hhs geo region lrircsnp',
               'hhs geo region lzgpxyit', 'hhs geo region mlyzmhmf',
               'hhs geo region oxchjgsf', 'hhs geo region qufhixun',
               'census msa MSA, Principle City', 'census msa Non-MSA',
               'behavior score'],
              dtvpe='object')
        1 9 2 3 1 1 1 5 1 1 11 1 10 1 7 12 1 6 1 1 1 1 1 1
          1 1 1 1]
        [ True True True True False True True True True True True True
          True False False True True True False True False True
         False True False False True False True True True True True True
          True True True True]
```

```
In [96]: masked columns = X train resample.columns[rfe.get support(indices= True)]
         masked columns
Out[96]: Index(['h1n1 concern', 'h1n1 knowledge', 'behavioral antiviral meds',
                 'behavioral avoidance', 'behavioral wash hands',
                 'behavioral large gatherings', 'behavioral outside home',
                 'behavioral touch face', 'doctor recc h1n1', 'chronic med condition',
                 'child under 6 months', 'health_worker', 'opinion_h1n1_vacc_effective',
                 'opinion h1n1 risk', 'opinion h1n1 sick from vacc', 'household adults',
                 'household children', 'age group 35 - 44 Years',
                 'age group 45 - 54 Years', 'age group 55 - 64 Years',
                 'age group 65+ Years', 'education College Graduate',
                 'education Some College', 'race White', 'sex Male',
                 'income poverty > $75,000', 'income poverty nan',
                 'marital status Not Married', 'rent or own Rent',
                 'employment status Not in Labor Force', 'hhs geo region bhuqougi',
                 'hhs geo region fpwskwrf', 'hhs geo region kbazzjca',
                 'hhs geo region lrircsnp', 'hhs geo region lzgpxyit',
                 'hhs geo region mlyzmhmf', 'hhs geo region oxchjgsf',
                 'hhs geo region qufhixun', 'census msa MSA, Principle City',
                 'census_msa_Non-MSA', 'behavior score'l,
               dtvpe='object')
```

```
In [97]: select colums = []
         for column in X train resample.columns:
             if column not in masked columns:
                 select colums.append(column)
         select colums
Out[97]: ['behavioral_face_mask',
           'education < 12 Years',
           'education nan',
           'race Hispanic',
           'race Other or Multiple',
           'income poverty Below Poverty',
           'marital_status_nan',
           'rent_or_own_nan',
           'employment status Unemployed',
           'employment status nan',
           'hhs geo region dapwygaj']
In [98]: X_train_rfe = X_train_resample[select_colums]
         X test rfe = X test[select colums]
```

```
In [99]: # Initialize the decision tree classifier with best parameters
         best dt classifier with rfe = DecisionTreeClassifier(criterion='entropy', max depth=20, min samples leaf=4, min sample
         # Train the model with best parameters
         best dt classifier with rfe.fit(X train rfe, y train resample)
         # Make predictions on the testing set
         v pred best = best dt classifier with rfe.predict(X test rfe)
         # Evaluate the model with best parameters
         accuracy best = accuracy score(y test, y pred best)
         print("Accuracy with best parameters:", accuracy best)
         # Classification report
         print("Classification Report with best parameters:")
         print(classification report(y test, y pred best))
         # Confusion matrix
         print("Confusion Matrix with best parameters:")
         print(confusion matrix(y test, y pred best))
         Accuracy with best parameters: 0.7867305676201887
         Classification Report with best parameters:
```

support	f1-score	recall	precision	
5254	0.88	1.00	0.79	0
1423	0.01	0.00	0.44	1
6677	0.79			accuracy
6677	0.44	0.50	0.62	macro avg
6677	0.69	0.79	0.71	weighted avg

Confusion Matrix with best parameters:

[[5249 5] [1419 4]]

```
In [100]: # ROC curve for the best Decision tree model
fpr_dtree, tpr_dtree, _ = roc_curve(y_test, best_dt_classifier_with_rfe.predict_proba(X_test_rfe)[:,1])
```

Random Forest

```
In [118]: def create_models(seed=40):
    models =[]
    models.append(('random_forest', RandomForestClassifier(random_state=seed)))
    return models
    models= create_models()
```

```
In [102]: # results using default parameters
          results= []
          names=[]
          scoring = 'accuracy'
          for name, model in models:
              # fit model with training data
              model.fit(X train resample, y train resample).predict(X test)
              # make predictions with testing data
              predictions=model.predict(X test)
              # calculating accuracy
              accuracy = accuracy score(y test, predictions)
              # append model name and accuracy to the lists
              results.append(accuracy)
              names.append(name)
              #print classifier accuracy
              print('classifier:{}, Accuracy score: {})'.format(name, accuracy))
              print(classification report(y test, predictions))
          classifier:random forest, Accuracy score: 0.8352553542009885)
                        precision
                                     recall f1-score support
```

```
In [103]: def perform grid search(classifer, param grid):
              pipe = Pipeline([
                  ('scaler', StandardScaler()),
                  ('classifier', RandomForestClassifier())
              1)
              random search = GridSearchCV(estimator = pipe,
                                 param grid = param grid,
                                 scoring = 'accuracy',
                                 cv = 3
                                 verbose =1)
              random search.fit(X train resample, y train resample)
              best params = random search.best params
              print("Best Parameters:", best params)
              # Evaluate the model on the test set
              v pred = random search.predict(X test)
              accuracy = accuracy score(y test, y pred)
              precision = precision score(y test, y pred)
              recall = recall score(y test, y pred)
              f1 = f1 score(y test, y pred)
              print("Test Accuracy:", accuracy)
              print("Precision:", precision)
              print("Recall:", recall)
              print("F1 score:", f1)
              # classifiaction reports
              print("Classification Report:")
              print(classification report(y test, y pred))
              return best params, accuracy
```

```
classifier = RandomForestClassifier()
In [104]:
          param grid = [{'classifier max depth': [5,10],
                  'classifier min samples split': [4,6]}]
          perform grid search(classifier, param grid)
          Fitting 3 folds for each of 4 candidates, totalling 12 fits
          Best Parameters: {'classifier max depth': 10, 'classifier min samples split': 4}
          Test Accuracy: 0.8301632469672008
          Precision: 0.6040316774658028
          Recall: 0.5895994378074491
          F1 score: 0.5967283072546231
          Classification Report:
                                     recall f1-score
                        precision
                                                        support
                             0.89
                                       0.90
                                                 0.89
                                                           5254
                             0.60
                                       0.59
                                                 0.60
                                                           1423
                                                 0.83
                                                           6677
              accuracy
                                                 0.74
                                                           6677
             macro avg
                             0.75
                                       0.74
          weighted avg
                             0.83
                                       0.83
                                                 0.83
                                                           6677
Out[104]: ({'classifier max depth': 10, 'classifier min samples split': 4},
           0.8301632469672008)
In [105]: # Final random forest model
          # Instantiate classifier
          random forest final = RandomForestClassifier(max depth= 10, min samples split= 4)
          # Fit model
          random forest final.fit(X train resample, y train resample)
Out[105]:
                             RandomForestClassifier
          RandomForestClassifier(max depth=10, min samples split=4)
```

```
In [106]: # ROC curve for the best Decision tree model
fpr_rforest, tpr_rforest, _ = roc_curve(y_test, random_forest_final.predict_proba(X_test)[:,1])
```

XGBoost

```
In [107]: # Clean feature names
          clean feature names = [name.replace('[', '').replace(']', '').replace('<', '') for name in X train resample.columns]</pre>
          X train resample.columns = clean feature names
          X test.columns = [name.replace('education < ', 'education ') for name in X test.columns]</pre>
          # Initialize the XGBoost model
          xgboost = XGBClassifier()
          # Train the XGBoost model
          xgboost.fit(X train resample, y train resample)
          # Make predictions on the training set
          v train pred = xgboost.predict(X train resample)
          # Calculate evaluation metrics for training set
          train accuracy = accuracy score(y train resample, y train pred)
          train precision = precision score(y train resample, y train pred)
          train recall = recall score(y train resample, y train pred)
          train f1 score = f1 score(y train resample, y train pred)
          # Make predictions on the test set
          y test pred = xgboost.predict(X test)
          print("confusion matrix for XGBoost\n",confusion matrix(y test,y test pred))
          print(classification report(y test, y test pred))
          print("Test Accuracy Score:", accuracy_score(y_test, y_test_pred))
```

confusion_matrix for XGBoost [[4890 364] [755 668]] precision recall f1-score support 0.87 0.93 0.90 5254 0 0.65 0.47 0.54 1423 1 0.83 6677 accuracy 0.72 macro avg 0.76 0.70 6677 weighted avg 0.82 0.83 0.82 6677

Test Accuracy Score: 0.8324097648644601

```
In [108]: # Define the hyperparameters to tune
          param grid = {
              'max depth': [ 5, 7, 9],
              'learning rate': [0.1, 0.01],
              'n estimators': [100, 300],
              'subsample': [0.8, 1.0],
              'colsample bytree': [0.8, 1.0]
          grid clf = GridSearchCV(xgboost, param_grid, scoring='accuracy', cv=None, n_jobs=1)
          grid clf.fit(X train resample, y train resample)
          best parameters = grid clf.best params
          print('Grid Search found the following optimal parameters: ')
          for param name in sorted(best parameters.keys()):
              print('%s: %r' % (param name, best parameters[param name]))
          training preds = grid clf.predict(X train resample)
          test preds = grid clf.predict(X test)
          training accuracy = accuracy score(y train resample, training preds)
          test accuracy = accuracy score(y test, test preds)
          print('')
          print('Training Accuracy: {:.4}%'.format(training accuracy * 100))
          print('Validation accuracy: {:.4}%'.format(test accuracy * 100))
          Grid Search found the following optimal parameters:
          colsample bytree: 0.8
          learning rate: 0.01
          max depth: 9
          n estimators: 300
          subsample: 0.8
          Training Accuracy: 90.91%
          Validation accuracy: 83.8%
```

enable categorical=False, eval metric=None, feature types=None,

interaction constraints=None, learning rate=0.01, max bin=None,

min child weight=None, missing=nan, monotone constraints=None,

gamma=None, grow policy=None, importance type=None,

multi strategy=None, n estimators=300, n jobs=None,

max_cat_threshold=None, max_cat_to_onehot=None,
max delta step=None, max depth=9, max leaves=None,

In [110]: # ROC curve for the best Decision tree model
fpr_xgboost, tpr_xgboost, _ = roc_curve(y_test, xgboost_final.predict_proba(X_test)[:,1])

Summary of all the tuned models using an Accuracy Scores:

- 1. Logistic Regression 0.7735
- 2. Decision Tree **0.7814**
- 3. KNN **0.7811**
- 4. Naive Bayes 0.7528
- 5. Random Forest **0.8288**
- 6. XGBoost 0.8405

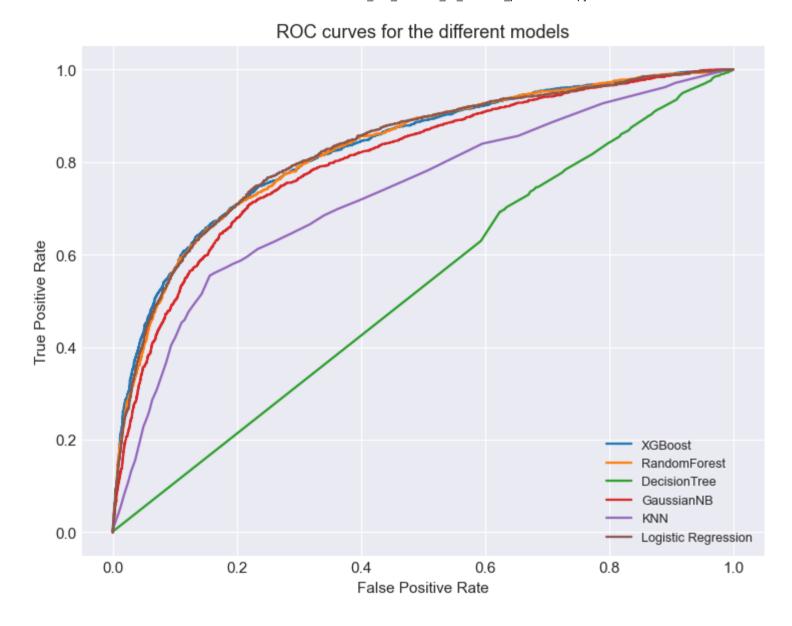
The best model for predicting vaccine uptake:

• XGBoost is therefore our **best and final model** with the best Accuracy of **0.8405**, as compared to all other models

ROC Curve

Summary of all the tuned models using an ROC curve:

```
In [111]: plt.figure(figsize=(8,6), dpi=110)
    plt.plot(fpr_xgboost, tpr_xgboost, label='XGBoost')
    plt.plot(fpr_forest, tpr_forest, label='RandomForest')
    plt.plot(fpr_dtree, tpr_dtree, label='DecisionTree')
    plt.plot(fpr_gnbayes, tpr_gnbayes, label='GaussianNB')
    plt.plot(fpr_knn, tpr_knn, label='KNN')
    plt.plot(fpr_logreg, tpr_logreg, label='Logistic Regression')
    plt.title('ROC curves for the different models')
    plt.xlabel('False Positive Rate')
    plt.ylabel('True Positive Rate')
    plt.legend(fontsize= 'small', loc='lower right')
    plt.show()
```



Conclusion

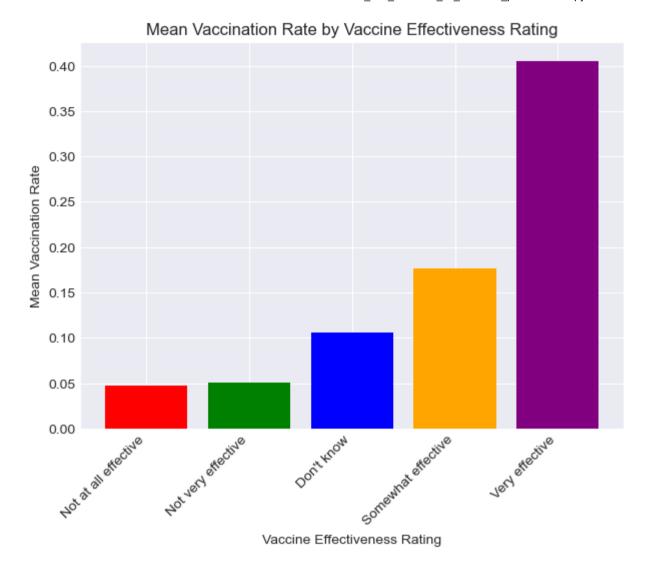
Features that were most important in predicting H1N1 vaccine uptake:

- 1. Opinion on H1N1 vaccine effectiveness.
- 2. Doctor's recommendation.
- 3. H1N1 concern.
- 4. Gender.
- 5. H1N1 Knowledge.
- 6. Perceived side-effects from H1N1 vaccine.
- 7. Chronic medical condition.
- 8 Number of adults in an household

```
In [112]: # Define a function to generate the plots for the important features
          def create vaccination rate plot(feature, target, data, x label, y label, plot title,
                                           x tick labels=None, plot width=7, plot height=5):
              mean vaccination rate = data.groupby(feature)[target].mean().reset index()
              # Define a list of colors to assign to each category
              colors = ['red', 'green', 'blue', 'orange', 'purple', 'black']
              fig, ax = plt.subplots(figsize=(plot_width, plot_height))
              ax.bar(mean vaccination rate[feature], mean vaccination rate[target],
                     color=colors[:len(mean vaccination rate[feature])])
              ax.set title(plot title)
              ax.set xlabel(x label)
              ax.set ylabel(y label)
              if x tick labels:
                  ax.set xticks(mean vaccination rate[feature])
                  ax.set xticklabels(x tick labels, rotation=45, ha='right')
              plt.show()
```

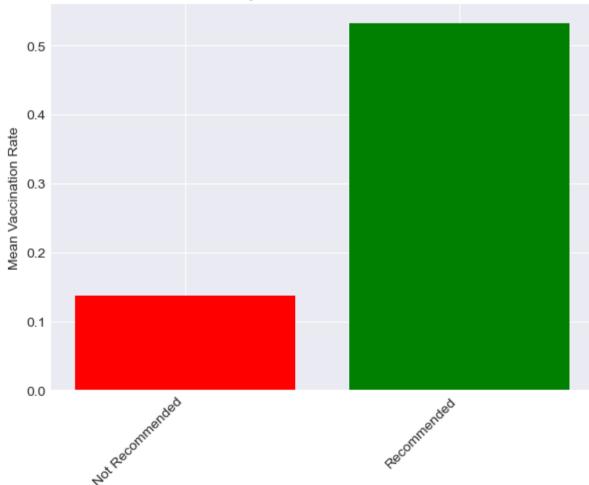
Plot of the top 5 most important features in predicting H1N1 vaccine uptake

1. Respondent's opinion about H1N1 flu vaccine effectiveness.



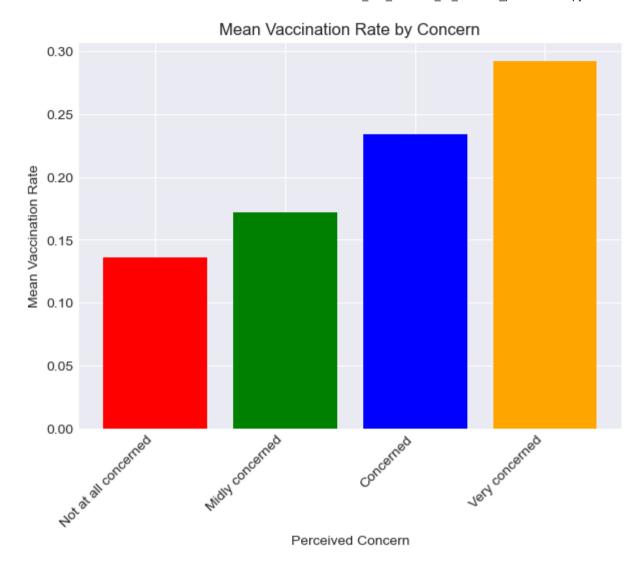
2. Doctor's recommendation.





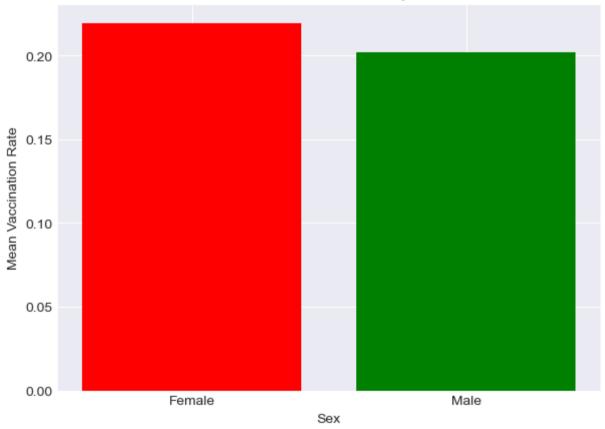
Doctor Recommendation for H1N1 Vaccine

3. H1N1 concern



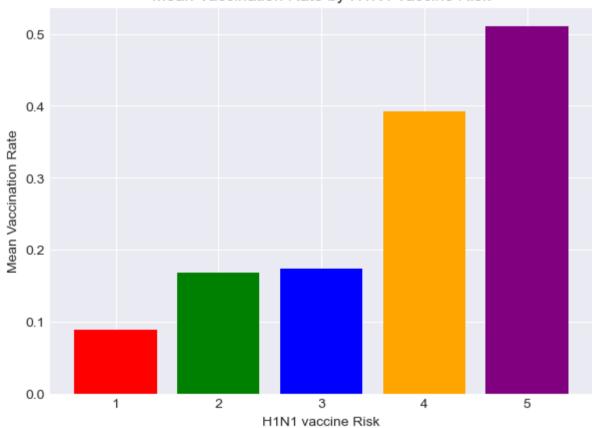
4. Gender





5. Perceived side effects from H1N1 vaccine





Recommendations

- 1. Dispel vaccine myths and promote preventative measures against the flu.
- 2. Identify high risk groups partcularly those with lower levels of education and indviduals expressing low concern for H1N1, to implement targeted messaging to them.
- 3. Utilize predictive model for efficent resource allocation on the vaccination campaign.
- 4. Implement policies that emphasize on face masks, minimizing gatherings and hand washing.
- 5. Collaboration with health experts to encouraage medical checkups and doctor's recommendation. tion

Next Steps

- 1. Monitoring and Assessment: Keep a close eye on the results of the interventions and communication tactics used. Compare the vaccination uptake rates before and after the interventions to assess the efficacy of various strategies.
- 2. Update the predictive model often with fresh information to enhance future interventions and targeting tactics.