

#### T.C.

# ESKİŞEHİR TECHNICAL UNIVERSITY ENGINEERING FACULTY

#### INDUSTRIAL ENGINEERING PROJECT

# PREDICTION OF H1N1 AND SEASONAL VACCINATION STATUS AND GETTING INDIVIDUALS INFORMED BY MACHINE LEARNING

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#### **Dataset**

#### 1.1 Goal and about the dataset

Each row in the dataset represents one person who responded to the National 2009 H1N1 Flu Survey. The aim is to predict how likely individuals are to receive their H1N1 and seasonal flu vaccines. Specifically, the model will be predicting two probabilities: one for 'h1n1\_vaccine' and one for 'seasonal\_vaccine'. In a possible epidemic, people who have not been vaccinated should be informed by a SMS message.

#### 1.1.1 Meanings of features

For all binary variables: 0 = No; 1 = Yes.

- h1n1\_vaccine Whether respondent received H1N1 flu vaccine.
- seasonal\_vaccine Whether respondent received seasonal flu vaccine.
- 1. h1n1 concern Level of concern about the H1N1 flu.
  - a. 0 = Not at all concerned; 1 = Not very concerned; 2 = Somewhat concerned; 3 = Very concerned.
- 2. h1n1\_knowledge Level of knowledge about H1N1 flu.
  - a. 0 = No knowledge; 1 = A little knowledge; 2 = A lot of knowledge.
- 3. <u>behavioral\_antiviral\_meds</u> Has taken antiviral medications. (binary)
- 4. <u>behavioral avoidance</u> Has avoided close contact with others with flu-like symptoms. (binary)
- 5. <u>behavioral\_face\_mask</u> Has bought a face mask. (binary)
- 6. <u>behavioral\_wash\_hands</u> Has frequently washed hands or used hand sanitizer. (binary)
- 7. <u>behavioral\_large\_gatherings</u> Has reduced time at large gatherings. (binary)

- 8. <u>behavioral outside home</u> Has reduced contact with people outside of own household. (binary)
- 9. <u>behavioral\_touch\_face</u> Has avoided touching eyes, nose, or mouth. (binary)
- 10. doctor\_recc\_h1n1 H1N1 flu vaccine was recommended by doctor. (binary)
- 11. <u>doctor\_recc\_seasonal</u> Seasonal flu vaccine was recommended by doctor. (binary)
- 12. <a href="mailto:chronic\_med\_condition">chronic\_med\_condition</a> 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)
- 13. <a href="mailto:child\_under\_6\_months">child\_under\_6\_months</a> Has regular close contact with a child under the age of six months. (binary)
- 14. health\_worker Is a healthcare worker. (binary)
- 15. health\_insurance Has health insurance. (binary)
- 16. <u>opinion h1n1 vacc effective</u> Respondent's opinion about H1N1 vaccine effectiveness.
  - a. 1 = Not at all effective; 2 = Not very effective; 3 = Don't know; 4 = Somewhat effective; 5 = Very effective.
- 17. <u>opinion\_h1n1\_risk</u> Respondent's opinion about risk of getting sick with H1N1 flu without vaccine.
  - a. 1 = Very Low; 2 = Somewhat low; 3 = Don't know; 4 = Somewhat high; 5 = Very high.
- 18. <u>opinion\_h1n1\_sick\_from\_vacc</u> Respondent's worry of getting sick from taking H1N1 vaccine.
  - a. 1 = Not at all worried; 2 = Not very worried; 3 = Don't know; 4 = Somewhat worried; 5 = Very worried.
- 19. <u>opinion seas vacc effective</u> Respondent's opinion about seasonal flu vaccine effectiveness.
  - a. 1 = Not at all effective; 2 = Not very effective; 3 = Don't know; 4 = Somewhat effective; 5 = Very effective.
- 20. <u>opinion\_seas\_risk</u> Respondent's opinion about risk of getting sick with seasonal flu without vaccine.
  - a. 1 = Very Low; 2 = Somewhat low; 3 = Don't know; 4 = Somewhat high; 5 = Very high.

- 21. <u>opinion\_seas\_sick\_from\_vacc</u> Respondent's worry of getting sick from taking seasonal flu vaccine.
  - a. 1 = Not at all worried; 2 = Not very worried; 3 = Don't know; 4 = Somewhat worried; 5 = Very worried.
- 22. age\_group Age group of respondent.
- 23. <u>education</u> Self-reported education level.
- 24. race Race of respondent.
- 25. <u>sex</u> Sex of respondent.
- 26. <u>income\_poverty</u> Household annual income of respondent with respect to 2008 Census poverty thresholds.
- 27. marital\_status Marital status of respondent.
- 28. <u>rent\_or\_own</u> Housing situation of respondent.
- 29. <a href="mailto:employment\_status">employment\_status</a> Employment status of respondent.
- 30. <u>census msa</u> Respondent's residence within metropolitan statistical areas (MSA) as defined by the U.S. Census.
- 31. household\_adults Number of **other** adults in household, top-coded to 3.
- 32. <a href="household\_children">household\_children</a> Number of children in household, top-coded to 3.
- 33. <u>phone</u> Phone number of observation

# **Data Preprocessing**

#### 2.1 Dataframe preprocessing

The training dataset as two pieces looks like below in its original form;

	respondent_id	h1n1_vaccine	seasonal_vaccine
0	0	0	0
1	1	0	1
2	2	0	0
3	3	0	1
4	4	0	0

Figure 2.1 : Dataframe including target variables

	respondent_id	h1n1_concern	h1n1_knowledge	$behavioral\_antiviral\_meds$	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hands
0	0	1.0	0.0	0.0	0.0	0.0	0.0
1	1	3.0	2.0	0.0	1.0	0.0	1.0
2	2	1.0	1.0	0.0	1.0	0.0	0.0
3	3	1.0	1.0	0.0	1.0	0.0	1.0
4	4	2.0	1.0	0.0	1.0	0.0	1.0

Figure 2.2 : Dataframe including independent variables (with unseen)

To avoid a dangerous index problem while dealing with missing values, they should be combined. After that, because it will never be used, the column 'respondent\_id' should be dropped.

h1n1_	_vaccine	seasonal_vaccine	h1n1_concern	h1n1_knowledge	behavioral_antiviral_meds	behavioral_avoidance	behavioral_face_mask	behavioral_wash_hand
	0	1	3.0	2.0	0.0	1.0	0.0	1.
	1	1	1.0	0.0	0.0	1.0	0.0	1.
	1	1	2.0	1.0	0.0	1.0	0.0	1.
	1	1	1.0	2.0	0.0	1.0	0.0	1.:
	0	0	1.0	1.0	0.0	1.0	0.0	0.

Figure 2.3: The final version of training dataframe (with unseen)

#### 2.2 Data types preprocessing

Because almost all the variables are categorical, it should be better to drop the rows which include missing value. Nevertheless, We still have 13506 observation. It looks enough to train a model which includes 32 independent variables.

As can be seen although some variables originally mean 'categorical', they are saved as 'int64' or 'float64' on the dataframe.

Figure 2.4: Primitive types of variables

The variables which mean categorical should be coordinated, and verbal categorical variables (kind of 'sex', 'age group', 'education' etc.) should be transformed to dummy and integer variables to work well.

Figure 2.5: After transformation and get the verbal variables to dummy

After all the preprocessing operations, to make sure I have saved the last form of dataframe as 'mukemmel.csv'\*. The dataframe that the model will train on is below.

#### 2.3 Determining training and test dataframes

By using 'model\_selection' from Scikit-Learn, the final form of dataframe will be divided into two parts as training and testing. %30 of observations will be used for testing data. Observations will be chosen randomly.

<sup>\*</sup> https://github.com/sametsoekel/flushotlearning/blob/master/mukemmel.csv

## **Model**

#### 3.1 Methodology

The optimal model should classify the target variable by using categorical variables. That is why CatBoost algorithm that is based on decision trees (developed to work on categorical datasets by Yandex in 2017) will be used to build a model.

If we try to explain how CatBoost algorithm works step by step;

- 1. In one randomly chosen row (k-th row in the training data set), we exchange one random level of this categorical feature (i-th level of x) with a number.
- 2. This number is usually based on the target variable (the one we want to predict) conditional on the category level. In other words, the target number is based on the expected outcome variable.
- 3. A splitting attribute is used to create two sets of the training data: One set that has all categories who will have greater target variable than the one computed in step 2, and the other set with smaller target variables.

We will need have 2 different models to predict two target variables (h1n1\_vaccine and seasonal\_vaccine). Here our target variables, I defined them as 'y1' and 'y2'. Everything else is independent variables;

h1n1	_vaccine	season	al_vaccine
0	0	0	0
1	0	1	1
2	1	2	1
3	0	3	0
4	1	4	1
	***	***	
13501	0	13501	0
13502	0	13502	0
13503	0	13503	0
13504	0	13504	0
13505	0	13505	0
13506 rows × 1 columns 13506 rows × 1 columns			

Figure 3.1: Target variables

#### **3.2 Primitive Models**

If we build two models in primitive without any attempt to optimize their outputs will be like;

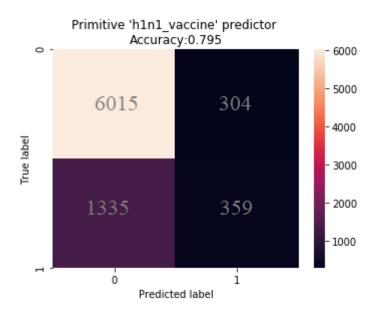


Figure 3.2: Confusion matrix of primitve 'y1' predictor.

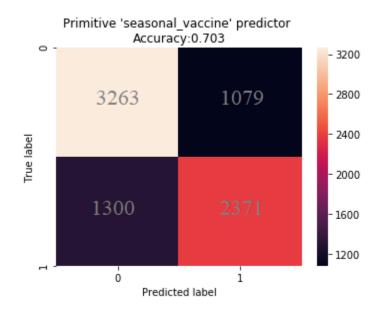


Figure 3.4: Confusion matrix of primitive 'y2' predictor.

Then according to primitive models, features importance be like respectively;

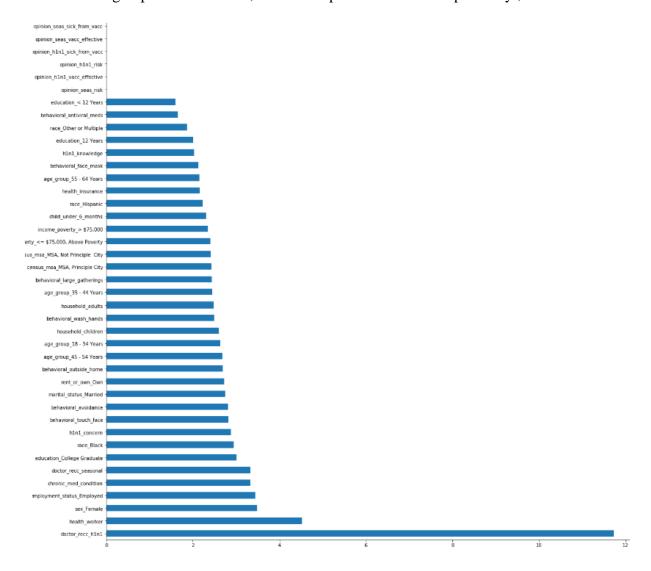


Figure 3.5: The most important feature is 'doctor\_recc\_hlnl'

The predictor of 'seasonal\_vaccine' says almost the same information. It says the most important features is 'doctor\_recc\_seasonal' and 'age\_group\_18 – 34 Years' respectively. According to these results, we can say that the doctor's recommendation is definitely affects people to get vaccinated or not.

#### 3.3 Model Tuning

GridSearchCV published by Scikit-Learn is going to help us to find the optimal hyperparameters of each model. It works on what you give as hyperparameters and how many fold of cross-validation. We can say that the library finds the optimal hyperparameters by brute-force.

I preffered 10 folds of cross-validation to validate my models well. Then to find optimal hyperparameters, I made a list of hyperparameters that are commonly used and highly recommended for CatBoost algorithm. They are like;

```
catb_params={
    'iterations':[100,200,300,500],
    'learning_rate':[0.01,0.05,0.1,0.2],
    'depth':[1,3,5,8]
}
```

Figure 3.6: Hyperparameters to be tested

I used the hyperparameters for each model, and after that different hyperparameters are determined for each.

```
{'iterations': 200, 'learning rate': 0.05, 'depth': 8}
```

Figure 3.7: Optimal hyperparameters for 'h1n1 vaccine' predictor

```
{'depth': 3, 'iterations': 500, 'learning_rate': 0.05}
```

Figure 3.8: Optimal hyperparameters for 'seasonal\_vaccine' predictor

#### 3.4 Model Testing

After validation, the model is ready to be tested. Re-modelling required with found hyperparameters.

#### Their scores will be like;

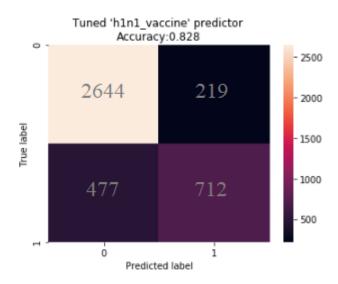


Figure 3.9: Confusion matrix of tuned 'y1' predictor

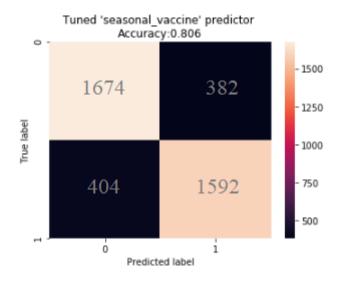


Figure 3.10: Confusion matrix of tuned 'y2' predictor

I saved both of models as 'h1n1_predictor.cbm' and 'seasonal_predictor.cbm' to use somewhere else. They can be downloaded from the link below.
https://github.com/sametsoekel/flushotlearning/blob/master/h1n1_predictor.cbm
https://github.com/sametsoekel/flushotlearning/blob/master/seasonal_predictor.cbm

# Designing interfaces to visualize model

QtDesigner allows very easy and successful interface without coding. These windows are created by QtDesigner.

The first interface offers file upload as \*.csv and choosing what kind of observation. Then shows contact numbers of individuals using tuned models.



Figure 4.1: Both of interfaces

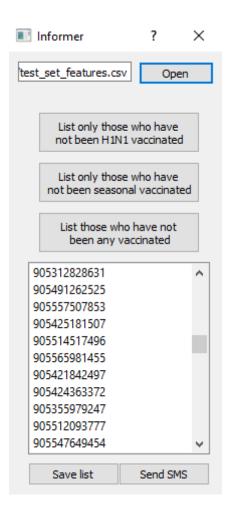


Figure 4.2: Informer interface

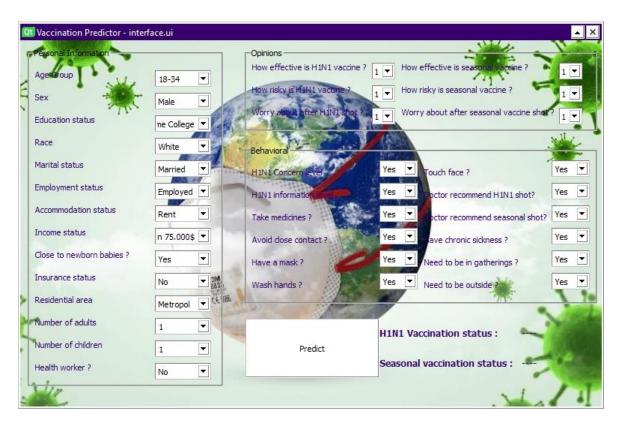


Figure 4.3: The singular prediction interface

After all arrangements the interface is ready to predict. For example;

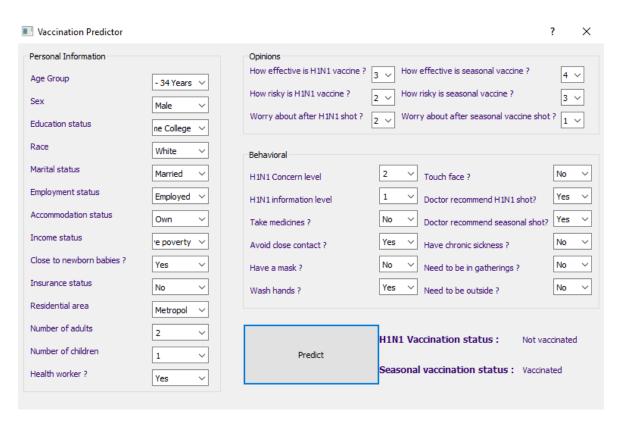


Figure 4.4: A prediction

## **Conclusion**

After validation and optimization, both of models can make predictions with %80 accuracy. If desired, predictions can be made from dataframes or singularly.

One of the biggest advantages of CatBoost algorithm is that datasets which consisting of so many categorical variables can be interpreted very well. As because my dataset is consisting of categorical variables as well, the algorithm work with perfect accuracy.

Almost all model operations were done on JupyterNotebook, that is why file extensions might be '\*.ipynb'. Data preprocessing operations are on 'work\_on\_dataset.ipynb', building tuned model and predicting operations are on 'final\_models.ipynb' and the predictor programs are on 'program.py' and 'second\_program.py'.

All the source codes, interfaces, model files are on the URL.

https://github.com/sametsoekel/flushotlearning