

# 1. Definition:

## 1.1. Project Overview:

Supervised learning is the machine learning task of inferring a function from labeled training data. The training data consist of a set of training examples. In supervised learning, each example is a pair consisting of an input object (typically a vector) and a desired output value (also called the supervisory signal). A supervised learning algorithm analyzes the training data and produces an inferred function, which can be used for mapping new examples. An optimal scenario will allow for the algorithm to correctly determine the class labels for unseen instances. This requires the learning algorithm to generalize from the training data to unseen situations in a "reasonable" way.

Nowadays, machine learning has been used vastly in insurance companies in order to recommend decent contracts to their clients based on the past information of them. For example, buying a new car and its insurance need a precise evaluation on both side client and insurance company to make a decent contract. Nothing ruins the thrill of buying a brand new car more quickly than seeing your new insurance bill. The sting's even more painful when you know you're a good driver. It doesn't seem fair that you have to pay so much if you've been cautious on the road for years. Therefore, having a robust predictive model is a key factor for successful companies.

Fig. 1 shows the road map of the project. The project includes four main parts: data analysis, feature engineering, modeling and prediction. In the data analysis part, we applied several statistical techniques to investigate the data in order to get better understanding about our data and prepare a good sample for our analysis. In the feature engineering section, we will deal with missing data, creating interaction variables, making new features by frequency encoding and binary encoding, calculating feature importance and finally select the importance feature and to reduce the size of the features. In the modeling part we will use several machine learning algorithms and stack them in three levels to make a prediction. Finally, by the prediction is made of the mean of the two last prediction models.

The main file that we can run the project is: *project.ipynb*

## 1.2. Problem- Statement

Porto Seguro, one of Brazil's largest auto and homeowner insurance companies. Inaccuracies in car insurance company's claim predictions raise the cost of insurance for good drivers and reduce the price for bad ones. The goal of this project is to build a model that predicts the probability that a driver will initiate an auto insurance claim in the next year.

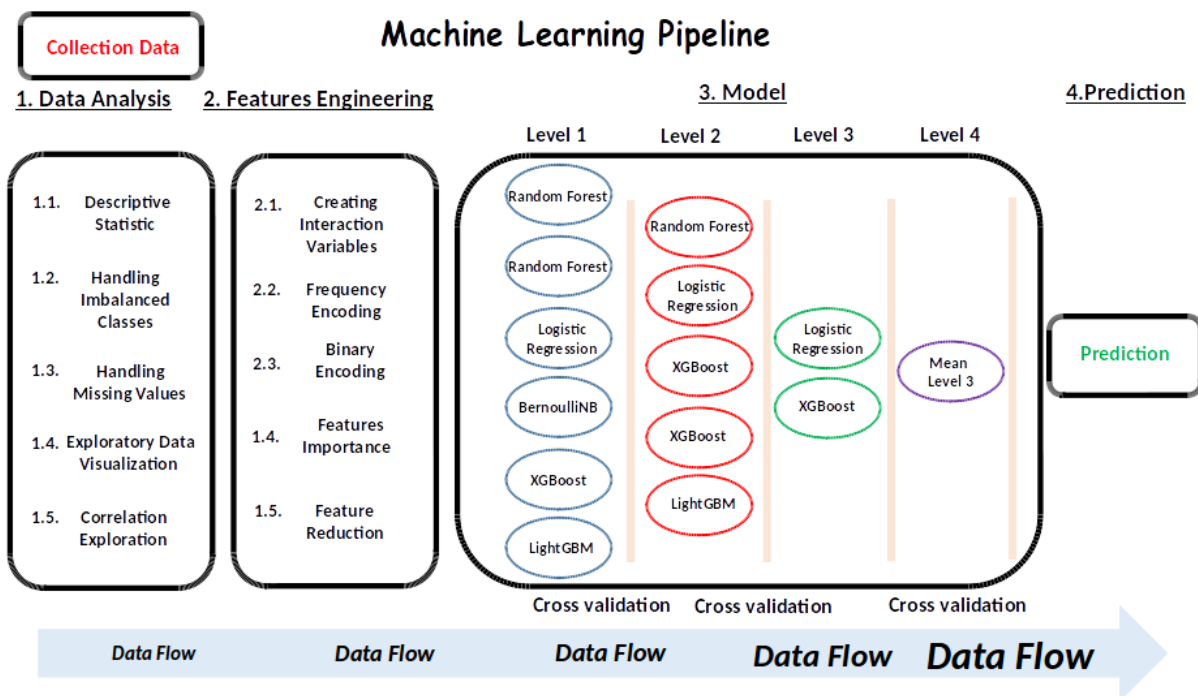


Fig.1. the road map of the project.

An accurate prediction will allow them to further tailor their prices, and hopefully make auto insurance coverage more accessible to more drivers. Thus, I will predict the probability that an auto insurance policy holder files a claim. The dataset is available dataset.

### 1.3. Metrics

code: `crossValidation/gini.py`

Our problem is binary classification. For binary classifiers, this option reports the AUC (area under curve) and Gini coefficient evaluation metrics. Both of these evaluation metrics are calculated together for each binary model. The values of the metrics are reported in a table in the analysis output browser.

The AUC evaluation metric is calculated as the area under an ROC (receiver operator characteristic) curve, and is a scalar representation of the expected performance of a classifier. The AUC is always between 0 and 1, with a higher number representing a better classifier. A

diagonal ROC curve between the coordinates (0,0) and (1,1) represents a random classifier, and has an AUC of 0.5. Therefore, a realistic classifier will not have an AUC of less than 0.5.

The Gini coefficient evaluation metric is sometimes used as an alternative evaluation metric to the AUC, and the two measures are closely related. The Gini coefficient is calculated as twice the area between the ROC curve and the diagonal, or as  $Gini = 2AUC - 1$ . The Gini coefficient is always between 0 and 1, with a higher number representing a better classifier. The Gini coefficient is negative in the unlikely event that the ROC curve is below the diagonal.

The Gini coefficient measures the inequality among values of a frequency distribution (for example, levels of income). A Gini coefficient of zero expresses perfect equality, where all values are the same (for example, where everyone has the same income). A Gini coefficient of 1 (or 100%) expresses maximal inequality among values (e.g., for a large number of people, where only one person has all the income or consumption, and all others have none, the Gini coefficient will be very nearly one). However, a value greater than one may occur if some persons represent negative contribution to the total (for example, having negative income or wealth).

The metric used for this Kaggle competition is the Normalized Gini Coefficient. During scoring, observations are sorted from the largest to the smallest predictions. Predictions are only used for ordering observations; therefore, the relative magnitude of the predictions are not used during scoring. The scoring algorithm then compares the cumulative proportion of positive class observations to a theoretical uniform proportion.

## **2. Analysis:**

### ***2.1. Data Exploration***

The dataset contains 2 .csv files with information necessary to make a prediction. In the train and test data, features that belong to similar groupings are tagged as such in the feature names (e.g., ind, reg, car, calc). In addition, feature names include the postfix bin to indicate binary features and cat to indicate categorical features. Features without these designations are either continuous or ordinal. Values of -1 indicate that the feature was missing from the observation. The shape of train is: (595212, 59) and memory usage: 267.9 MB and test set shape is (892816, 28). The column id shows the 'id' of customers and the target column is 'target'. There is no NaN in the data set.

### **2.1.1 Meta Data:**

*Analysis/featureAnalysis/fetureAnalysis.py*

To facilitate the data management, we'll store meta-information about the variables in a data frame. This will be helpful when we want to select specific variables for analysis, visualization, modeling, etc.. We can also apply the describe method on the data frame. However, it doesn't make much sense to calculate the mean, std, etc. on categorical variables and the id variable. We'll explore the categorical variables visually later. Thanks to our meta file we can easily select the variables on which we want to compute the descriptive statistics. To keep things clear, we'll do this per data type.

The meta data frame includes this information:

Feature, type of feature, number unique, number of duplicate, 1<sup>st</sup> frequency of data, number of 1<sup>st</sup> frequency of data, 2<sup>nd</sup> frequency of data, number of 2<sup>nd</sup> frequency of data, 3<sup>rd</sup> frequency of data, number of 3<sup>rd</sup> frequency of data, mean, standard deviation, min, max, 25%, 50%, 75% of the data information. If the data does not include these info, the NaN is replaced in the data frame cell. Thus, based on the analysis, the 'id' is unique and the target is classified to 0 or 1. Therefore, the problem is binary classification. The brief summary of the this analysis is:

reg variables:

- only ps\_reg\_03 has missing values
- the range (min to max) differs between the variables. We could apply scaling (e.g. StandardScaler), but it depends on the classifier we will want to use.

car variables:

- ps\_car\_12 and ps\_car\_15 have missing values
- again, the range differs and we could apply scaling.

calc variables:

- no missing values
- this seems to be some kind of ratio as the maximum is 0.9
- all three \_calc variables have very similar distributions

Ordinal variables:

- Only one missing variable: ps\_car\_11
- We could apply scaling to deal with the different ranges

Binary variables :

- A priori in the train data is 3.645%, which is strongly imbalanced.
- From the means we can conclude that for most variables the value is zero in most cases.

Target :

- A priori in the train data is 3.645%, which is strongly imbalanced.
- From the means we can conclude that for most variables the value is zero in most cases.

## 2.2. Exploratory Visualization

The dataset includes binary, categorical and numerical features. The plots of all figures are shown in the Jupyter Notebook. The binary and categorical features are plotted by histogram methods. The numerical features are plot by histogram and boxplot methods in order to find out their distributions and the outliers. However, the dataset does not suffer from outliers problem. The correlation of numerical variables are obtained and plot by the heat map plot. Besides, the relation between numerical data is obtain by the seaborn linear regression library package for the target.

The target distribution is shown on Fig. 2. It can be seen that the target has imbalance data.

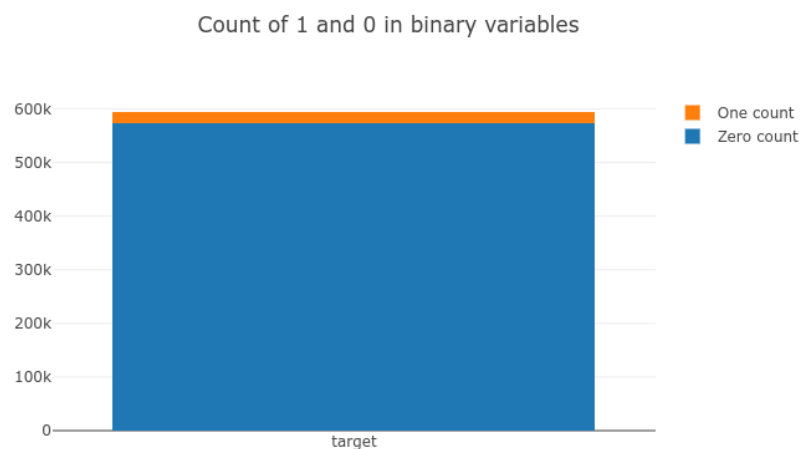


Fig.2: target distribution.

Hence, we need to treat the imbalance target in the project. The distribution of target in binary features is shown Fig.3 after sampling.

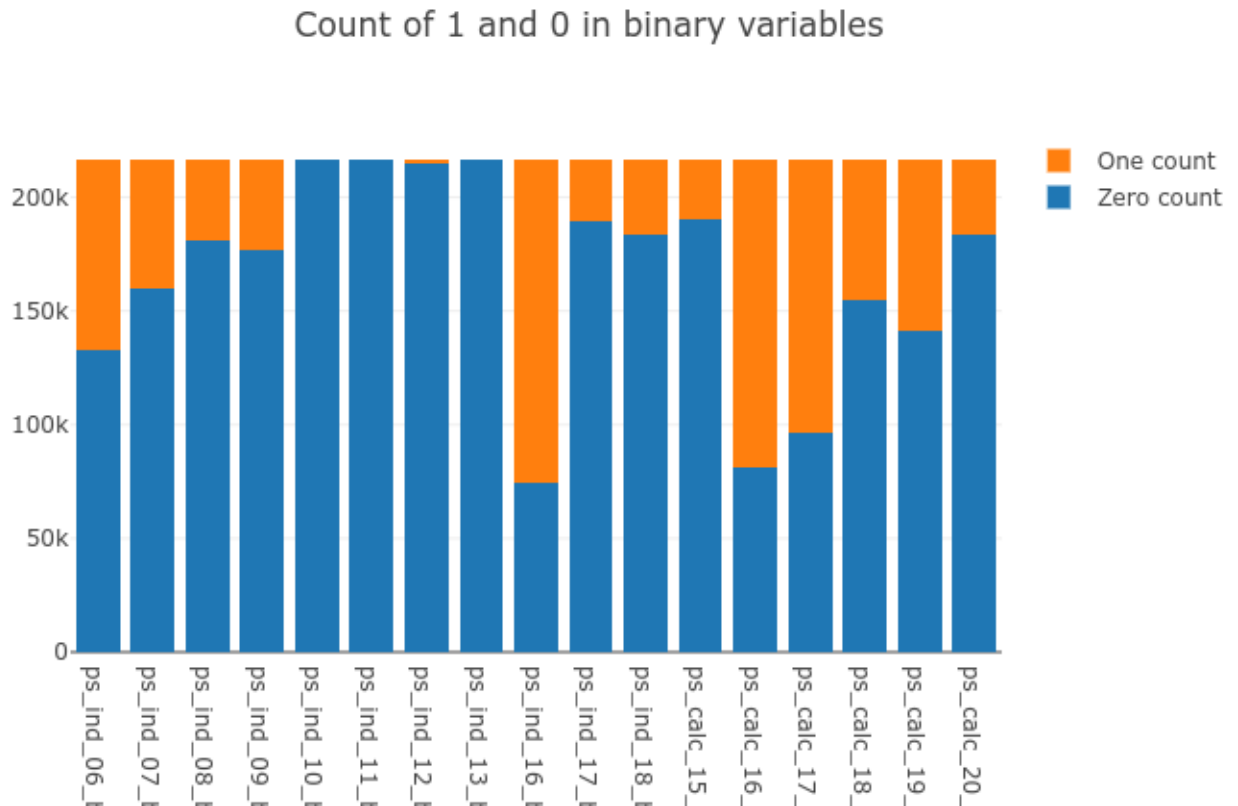


Fig. 3: The distribution of target in binary features.

The correlation among the numerical values is plotted Fig.4. The relation between the numerical values is shown in Fig. 5

Fig.4: Heat Map

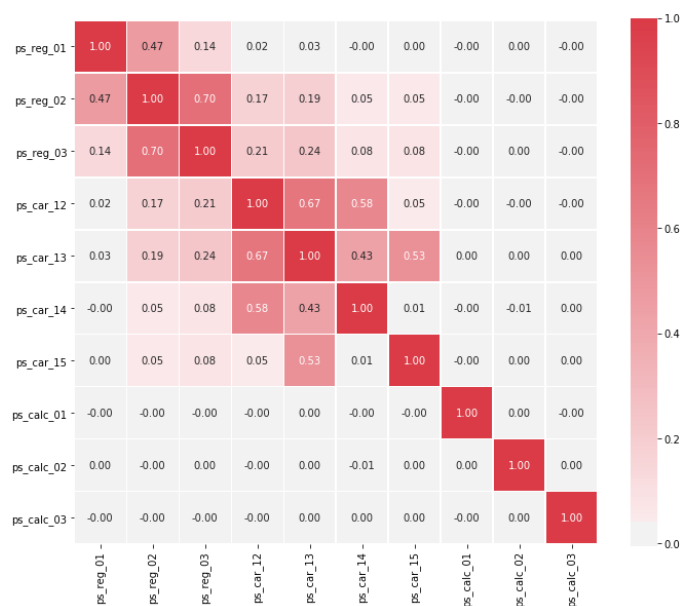
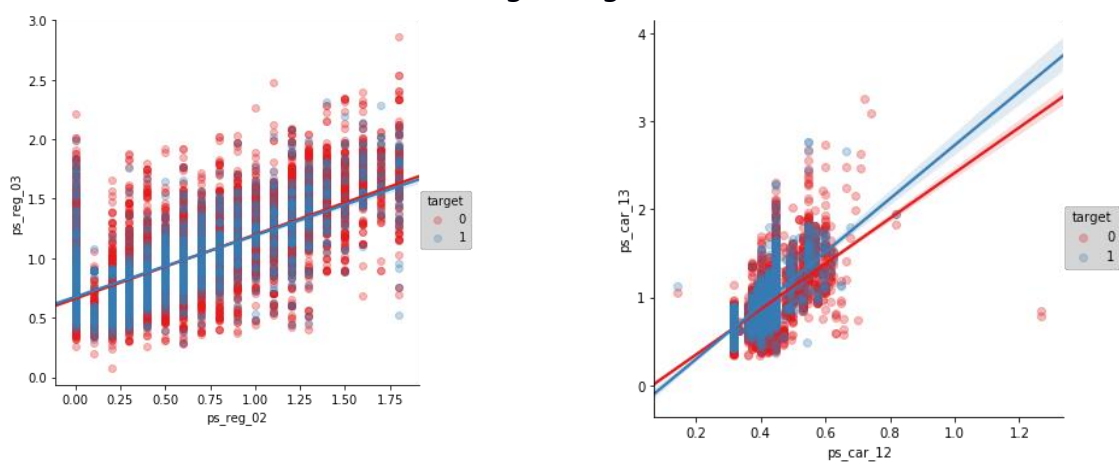
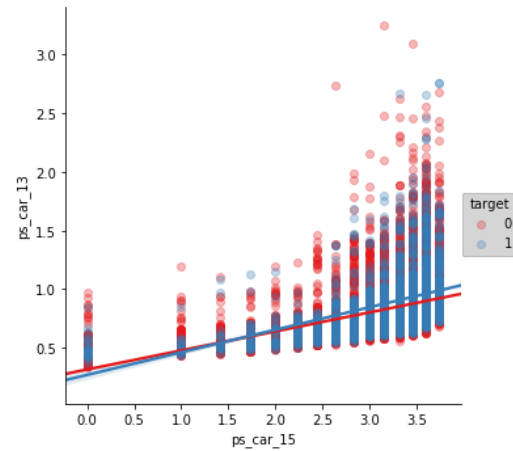
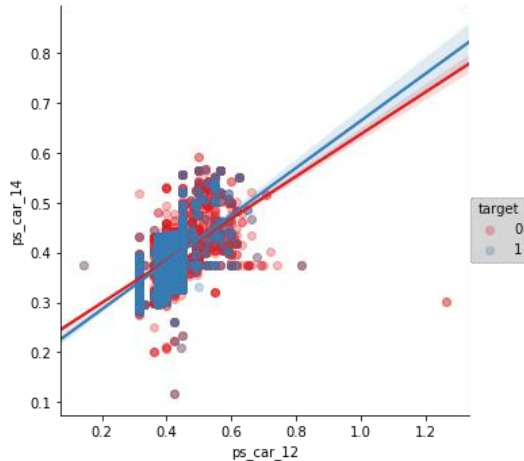


Fig.5: regression relation





### 2.3. Algorithm and Techniques

In this section I will explain the five algorithms that I use in this project: logistic regression, BernoulliNB, Random Forest, XGBoost and Light GBM.

#### logistic regression model:

Logistic regression is a statistical method for analyzing a dataset in which there are one or more independent variables that determine an outcome. The outcome is measured with a dichotomous variable (in which there are only two possible outcomes).

#### BernoulliNB model:

BernoulliNB implements the naive Bayes training and classification algorithms for data that is distributed according to multivariate Bernoulli distributions; i.e., there may be multiple features but each one is assumed to be a binary-valued (Bernoulli, boolean) variable. Therefore, this class requires samples to be represented as binary-valued feature vectors; if handed any other kind of data, a BernoulliNB instance may binarize its input (depending on the binarize parameter).

The decision rule for Bernoulli naive Bayes is based on

$$P(x_i|y) = P(i|y)x_i + (1-P(i|y))(1-x_i)$$

which differs from multinomial NB's rule in that it explicitly penalizes the non-occurrence of a feature  $i$  that is an indicator for class  $y$ , where the multinomial variant would simply ignore a non-occurring feature.



## **Ensemble methods:**

The goal of ensemble methods is to combine the predictions of several base estimators built with a given learning algorithm in order to improve generalizability / robustness over a single estimator.

Two families of ensemble methods are usually distinguished:

In averaging methods, the driving principle is to build several estimators independently and then to average their predictions. On average, the combined estimator is usually better than any of the single base estimator because its variance is reduced.

Example: Bagging methods, Forests of randomized trees, ...

By contrast, in boosting methods, base estimators are built sequentially and one tries to reduce the bias of the combined estimator. The motivation is to combine several weak models to produce a powerful ensemble.

Examples: AdaBoost, Gradient Tree Boosting, ...

The `sklearn.ensemble` module includes two averaging algorithms based on randomized decision trees: the `RandomForest` algorithm and the `Extra-Trees` method.

## **Random Forests:**

In random forests, each tree in the ensemble is built from a sample drawn with replacement (i.e., a bootstrap sample) from the training set. In addition, when splitting a node during the construction of the tree, the split that is chosen is no longer the best split among all features. Instead, the split that is picked is the best split among a random subset of the features. As a result of this randomness, the bias of the forest usually slightly increases (with respect to the bias of a single non-random tree) but, due to averaging, its variance also decreases, usually more than compensating for the increase in bias, hence yielding an overall better model

A random forest is a meta estimator that fits a number of decision tree classifiers on various sub-samples of the dataset and use averaging to improve the predictive accuracy and control over-fitting. The sub-sample size is always the same as the original input sample size but the samples are drawn with replacement if `bootstrap=True` (default).

## **XGBoost**

XGBoost stands for eXtreme Gradient Boosting. The name xgboost, though, actually refers to the engineering goal to push the limit of computations resources for boosted tree algorithms. Which is the reason why many people use xgboost.

### **Algorithm Use by XGBoost**

The XGBoost library implements the gradient boosting decision tree algorithm. This algorithm goes by lots of different names such as gradient boosting, multiple additive regression trees, stochastic gradient boosting or gradient boosting machines. Boosting is an ensemble technique where new models are added to correct the errors made by existing models. Models are added sequentially until no further improvements can be made. A popular example is the AdaBoost algorithm that weights data points that are hard to predict.

Gradient boosting is an approach where new models are created that predict the residuals or errors of prior models and then added together to make the final prediction. It is called gradient boosting because it uses a gradient descent algorithm to minimize the loss when adding new models. This approach supports both regression and classification predictive modeling problems.

### **1. The XGBoost Advantage**

- a. Regularization: Standard GBM implementation has no regularization like XGBoost, therefore it also helps to reduce overfitting. In fact, XGBoost is also known as 'regularized boosting' technique.
- b. Parallel Processing: XGBoost implements parallel processing and is blazingly faster as compared to GBM.
- c. High Flexibility: XGBoost allow users to define custom optimization objectives and evaluation criteria. This adds a whole new dimension to the model and there is no limit to what we can do.
- d. Handling Missing Values: XGBoost has an in-built routine to handle missing values. User is required to supply a different value than other observations and pass that as a parameter. XGBoost tries different things as it encounters a missing value on each node and learns which path to take for missing values in future.
- e. Tree Pruning: A GBM would stop splitting a node when it encounters a negative loss in the split. Thus it is more of a greedy algorithm. XGBoost on the other hand make splits

upto the `max_depth` specified and then start pruning the tree backwards and remove splits beyond which there is no positive gain. Another advantage is that sometimes a split of negative loss say -2 may be followed by a split of positive loss +10. GBM would stop as it encounters -2. But XGBoost will go deeper and it will see a combined effect of +8 of the split and keep both.

- f. Built-in Cross-Validation: XGBoost allows user to run a cross-validation at each iteration of the boosting process and thus it is easy to get the exact optimum number of boosting iterations in a single run. This is unlike GBM where we have to run a grid-search and only a limited values can be tested.
- g. Continue on Existing Model: User can start training an XGBoost model from its last iteration of previous run. This can be of significant advantage in certain specific applications.

## 1. Light GBM

Light GBM is a fast, distributed, high-performance gradient boosting framework based on decision tree algorithm, used for ranking, classification and many other machine learning tasks.

Since it is based on decision tree algorithms, it splits the tree leaf wise with the best fit whereas other boosting algorithms split the tree depth wise or level wise rather than leaf-wise. So when growing on the same leaf in Light GBM, the leaf-wise algorithm can reduce more loss than the level-wise algorithm and hence results in much better accuracy which can rarely be achieved by any of the existing boosting algorithms. Also, it is surprisingly very fast, hence the word 'Light'.

Before is a diagrammatic representation by the makers of the Light GBM to explain the difference clearly.

## 2. Advantages of Light GBM

- a. Faster training speed and higher efficiency: Light GBM use histogram based algorithm i.e it buckets continuous feature values into discrete bins which fasten the training procedure.
- b. Lower memory usage: Replaces continuous values to discrete bins which result in lower memory usage.
- c. Better accuracy than any other boosting algorithm: It produces much more complex trees by following leaf wise split approach rather than a level-wise approach which is the main factor in achieving higher accuracy. However, it can sometimes lead to overfitting which can be avoided by setting the `max_depth` parameter.
- d. Compatibility with Large Datasets: It is capable of performing equally good with large datasets with a significant reduction in training time as compared to XGBOOST.

- e. Parallel learning supported.

## **2.4. Benchmark**

In this section, I show the result of a logistic regression model which is applied on this problem. Our metric is Gini which is related to  $Gini = 2AUC - 1$ . For the simple logistic regression on the sampled data, we obtained  $Gini = 0.19$  score. I am going to improve the result of the prediction. Although this calculation shows an acceptable gini score, the prediction cannot predict the real data because the train data is unbalance, hence, our prediction has overfit problem. Therefore, to establish a robust model we have to overcome the unbalance data problem in our data set.

## **3. Methodology**

### **3.1. Data Preprocessing**

#### **3.1.1 Handling Imbalanced Classes**

Preparing/sampleing.py

What can we do when we have imbalanced data? Mainly three things: 1) Ignoring the problem. 2) Undersampling the majority class. 3) Oversampling the minority class.

IGNORING THE PROBLEM:

Building a classifier using the data as it is, would in most cases give us a prediction model that always returns the majority class. The classifier would be biased. Let's build the models:

UNDERSAMPLING THE MAJORITY CLASS:

One of the most common and simplest strategies to handle imbalanced data is to undersample the majority class. While different techniques have been proposed in the past, typically using more advanced methods did not bring any improvement with respect to simply selecting samples at random. So, for this analysis I will simply select  $n$  samples at random from the majority class, where  $n$  is the number of samples for the minority class, and use them during training phase, after excluding the sample to use for validation.

OVERSAMPLING THE MINORITY CLASS

Oversampling the minority class can result in overfitting problems if we oversample before cross-validating. What is wrong with oversampling before cross-validating? Let's consider the simplest oversampling method ever, as an example that clearly explains this point.

The easiest way to oversample is to re-sample the minority class, i.e. to duplicate the entries, or manufacture data which is exactly the same as what we have already. Now, if we do so before cross-validating, i.e. before we enter the leave one participant out cross-validation loop, we will be training the classifier using N-1 entries, leaving 1 out, but including in the N-1 one or more instances that are exactly the same as the one being validated. Thus, defeating the purpose of cross-validation altogether.

As we have a rather large training set, we can go for undersampling. Therefore, the percentage of 0 in train 0.96, the percentage of 1 in train 0.03, the percentage of 0 in the sample train 0.9, the percentage of 1 in the sample train 0.1

### ***3.1.1 Handling Missing Value in the Sample dataset***

analysis/missingvalue/missingvalue.py, Analysis/traintest/traintest.py

The missing values for each feature are:

Variable ps\_ind\_02\_cat has 216 records (0.04%) with missing values.

Variable ps\_ind\_04\_cat has 83 records (0.01%) with missing values.

Variable ps\_ind\_05\_cat has 5809 records (0.98%) with missing values.

Variable ps\_reg\_03 has 107772 records (18.11%) with missing values.

Variable ps\_car\_01\_cat has 107 records (0.02%) with missing values.

Variable ps\_car\_02\_cat has 5 records (0.00%) with missing values.

Variable ps\_car\_03\_cat has 411231 records (69.09%) with missing values.

Variable ps\_car\_05\_cat has 266551 records (44.78%) with missing values.

Variable ps\_car\_07\_cat has 11489 records (1.93%) with missing values.

Variable ps\_car\_09\_cat has 569 records (0.10%) with missing values.

Variable ps\_car\_11 has 5 records (0.00%) with missing values.

Variable ps\_car\_12 has 1 records (0.00%) with missing values.

Variable `ps_car_14` has 42620 records (7.16%) with missing values.

In total, there are 13 variables with missing values. Thus, we deal with missing values as follows:

`ps_car_03_cat` and `ps_car_05_cat` have a large proportion of records with missing values. Remove these variables.

For the other categorical variables with missing values, we can leave the missing value -1 as such.

`ps_reg_03` (continuous) has missing values for 18% of all records. Replace by the mean.

`ps_car_11` (ordinal) has only 5 records with missing values. Replace by the mode.

`ps_car_12` (continuous) has only 1 records with missing value. Replace by the mean.

`ps_car_14` (continuous) has missing values for 7% of all records. Replace by the mean.

### **3.1.2. Features Engineering**

CREATING INTERACTION VARIABLE:

Preparing/categoricalEncoding.py

The `klearn.preprocessing` library is used for flood data to make new features from interaction between features with `PolynomialFeatures(2)`. Before creating interactions, we have 57 variables. After creating interactions, we have 112 variables. Before creating interactions, we have 56 variables. After creating interactions, we have 111 variable.

FREQUENCY ENCODING:

Preparing/categoricalEncoding.py

For the categorical data, `ps_ind_02_cat`, `ps_car_04_cat`, `ps_car_09_cat`, `ps_ind_05_cat`, `ps_car_01_cat`, `ps_car_11_cat`, the number of repetition data for each feature is counted and the results is saved into a new data frame column.

BINARY ENCODING:

Preparing/categoricalEncoding.py

Additionally, for the categorical data with low number of category, `ps_ind_02_cat`, `ps_car_04_cat`, `ps_car_09_cat`, `ps_ind_05_cat`, `ps_car_01_cat`, `ps_car_11_cat`, I did binary encoding.

Finally, train shape is: (216940, 135) and test shape is: (892816, 134).

### 3.2. Implementation

The first step of the project is data analysis. I determine the type of the problem, features, missing value and imbalance data in this section. Then, since the target is imbalance, I used undersampling model to produce more balance data. After that, in the data engineering section, the new features are added to base dataset. The interaction between numerical features are obtained by polynomial method. The categorical features are treated in two ways: binary encoding and frequency encoding. The important features in the data are extract by random forest feature importance model.

In the modeling part three level of machine learning algorithms are stacked together. In the first level, Logistic Regression, BernouliNB, Random Forest, XGBoost and Light GBM are applied. The result of the five models are stacked and it is used as the input of the level 2. In the level 2, Logistic Regression, Random Forest, XGBoost and Light GBM algorithms are used. Finally, in the last level we applied Logistic Regression, XGBoost models and their result are avarged to make the final prediction.

### 3.3. Refinement

**for the logistic regression tuning parameters:**

Tuning logistic regression penalty = l1

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## Tuning logistic regression $c = 0.1$

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sklearn: logit0

fold cv 0 AUC score is 0.600559, Gini Norm score is 0.201117

fold cv 1 AUC score is 0.596478, Gini Norm score is 0.192955

fold cv 2 AUC score is 0.590134, Gini\_Norm score is 0.180268

fold cv 3 AUC score is 0.597373, Gini Norm score is 0.194746

fold cv 4 AUC score is 0.604743, Gini Norm score is 0.209487

cv AUC score is 0.597826, Gini Norm score is 0.195652

it takes 241.420 seconds to perform cross validation

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Tuning logistic regression  $c = 0.2$

sklearn: logit1

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## Tuning logistic regression $c = 0.5$

sklearn: logit2

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Tuning logistic regression  $c = 0.6$

sklearn: logit3

```
fold cv 0 AUC score is 0.600121, Gini_Norm score is 0.200242
fold cv 1 AUC score is 0.596735, Gini_Norm score is 0.193469
fold cv 2 AUC score is 0.589916, Gini_Norm score is 0.179831
fold cv 3 AUC score is 0.597286, Gini_Norm score is 0.194572
fold cv 4 AUC score is 0.604630, Gini_Norm score is 0.209261
cv AUC score is 0.597702, Gini_Norm score is 0.195404
it takes 464.058 seconds to perform cross validation
```



[illegible]

Tuning logistic regression c = 0.1

```

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sklearn: logit4

```

```
fold cv 0 AUC score is 0.600112, Gini_Norm score is 0.200223
fold cv 1 AUC score is 0.596640, Gini_Norm score is 0.193279
fold cv 2 AUC score is 0.589793, Gini_Norm score is 0.179587
fold cv 3 AUC score is 0.597297, Gini_Norm score is 0.194594
fold cv 4 AUC score is 0.604579, Gini_Norm score is 0.209159
cv AUC score is 0.597649, Gini_Norm score is 0.195299
it takes 25.757 seconds to perform cross validation
```

Tuning logistic regression c = 0.2

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sklearn: logit5
```

```
fold cv 0 AUC score is 0.600042, Gini_Norm score is 0.200083
fold cv 1 AUC score is 0.596696, Gini_Norm score is 0.193391
fold cv 2 AUC score is 0.589805, Gini_Norm score is 0.179610
fold cv 3 AUC score is 0.597295, Gini_Norm score is 0.194589
fold cv 4 AUC score is 0.604579, Gini_Norm score is 0.209158
cv AUC score is 0.597647, Gini_Norm score is 0.195295
it takes 26.582 seconds to perform cross validation
```

Tuning logistic regression c = 0.5

[illegible]

fold cv 0 AUC score is 0.599978, Gini\_Norm score is 0.199956

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Based on the calculation it seems that the first model: penalty = 0.1 and l1 has the best gini = 0.1956 score.

Tuning Random Forest = 80

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Tuning Random Forest = 100

fold cv 0 AUC score is 0.600872, Gini\_Norm score is 0.201744  
fold cv 1 AUC score is 0.599315, Gini\_Norm score is 0.198630  
fold cv 2 AUC score is 0.594168, Gini\_Norm score is 0.188336  
fold cv 3 AUC score is 0.597708, Gini\_Norm score is 0.195416  
fold cv 4 AUC score is 0.607792, Gini\_Norm score is 0.215583  
cv AUC score is 0.599804, Gini\_Norm score is 0.199608  
it takes 74.736 seconds to perform cross validation

fold cv 0 AUC score is 0.603556, Gini\_Norm score is 0.207113  
fold cv 1 AUC score is 0.605790, Gini\_Norm score is 0.211580  
fold cv 2 AUC score is 0.601267, Gini\_Norm score is 0.202535  
fold cv 3 AUC score is 0.605055, Gini\_Norm score is 0.210110  
fold cv 4 AUC score is 0.612224, Gini\_Norm score is 0.224448  
cv AUC score is 0.605451, Gini\_Norm score is 0.210903  
it takes 139.322 seconds to perform cross validation

[illegible]

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Tuning xgb eta = 0.5

fold cv 0 AUC score is 0.610521, Gini\_Norm score is 0.221042  
fold cv 1 AUC score is 0.608286, Gini\_Norm score is 0.216573  
fold cv 2 AUC score is 0.606437, Gini\_Norm score is 0.212874  
fold cv 3 AUC score is 0.609278, Gini\_Norm score is 0.218555  
fold cv 4 AUC score is 0.616090, Gini\_Norm score is 0.232180  
cv AUC score is 0.610019, Gini\_Norm score is 0.220039  
it takes 23.814 seconds to perform cross validation

/////////  
Tuning xgb eta = 0.05

fold cv 0 AUC score is 0.614052, Gini\_Norm score is 0.228104  
fold cv 1 AUC score is 0.614145, Gini\_Norm score is 0.228289  
fold cv 2 AUC score is 0.611498, Gini\_Norm score is 0.222996  
fold cv 3 AUC score is 0.614662, Gini\_Norm score is 0.229324  
fold cv 4 AUC score is 0.621993, Gini\_Norm score is 0.243987  
cv AUC score is 0.615154, Gini\_Norm score is 0.230307  
it takes 69.842 seconds to perform cross validation

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xgb: xgb7





Tuning lgb learning rate= 0.1

lgb: lgb1

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lgb: lgb2

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lgb: lgb3

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it takes 26.670 seconds to perform cross validation

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it takes 13.958 seconds to perform cross validation

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fold cv 4 AUC score is 0.619785, Gini Norm score is 0.239571

Tuning lgb learning rate= 0.1

```
fold cv 0 AUC score is 0.614036, Gini_Norm score is 0.228071
fold cv 1 AUC score is 0.612551, Gini_Norm score is 0.225101
fold cv 2 AUC score is 0.608329, Gini_Norm score is 0.216657
fold cv 3 AUC score is 0.610588, Gini_Norm score is 0.221176
fold cv 4 AUC score is 0.619877, Gini_Norm score is 0.239753
cv AUC score is 0.612752, Gini_Norm score is 0.225504
it takes 30.125 seconds to perform cross validation
```

```

-_-*-*-*-*-*-*-*-*-*-*-*-*-*-*-*-*-*-*-*-*
lgb: lgb8

```

max\_depth = 4 and learning rate = 0.05 has the best result

The stacking technique in three levels is applied to improve the result of the calculation.

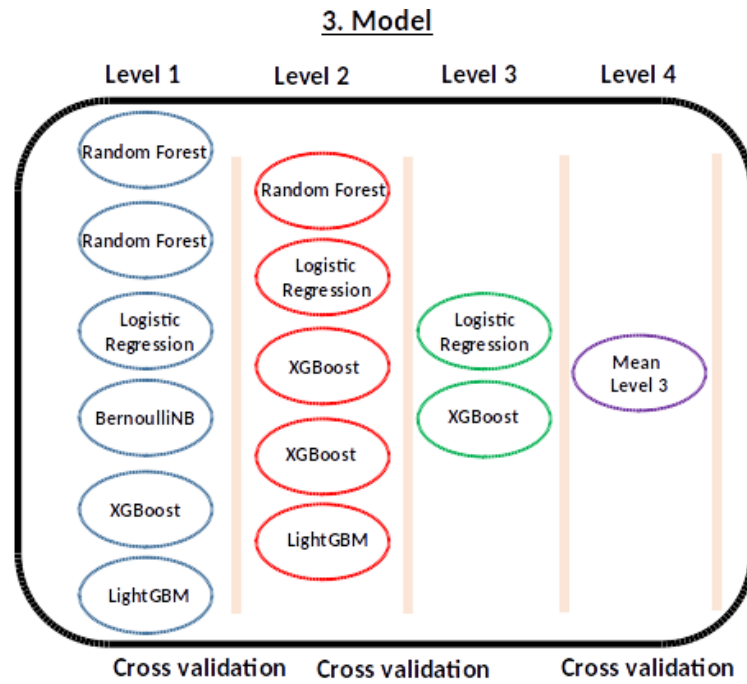


Fig.6: The stacking model to improve the prediction.

The result of the improvement in gibi score shown in the Fig.6 is listed in the Table 1. We keep the same order as Fig.6 to report in table.

Without tuning hyper parameters:

Table 1.

Level 1	Level 2	Level 3	Level 4
0.212			
0.199	0.227		
0.195	0.232	0.230	
0.165	0.105	0.227	0.229
0.232	0.232		
0.230	0.232		

With tuning hyper parameters

Table 2.

Level 1	Level 2	Level 3	Level 4
0.207			
0.199	0.226		
0.195	0.228	0.232	
0.165	0.114	0.244	0.230
0.228	0.228		
0.221	0.229		

## 4. Results

### 4.1. Model Evaluation and Validation

In this project, in order to improve our prediction in comparison a single model which is explained in Benchmark, we tried Four ways:

1) Tuning each model by its hyper-parameter. I tune hyper parameters. Without tuning I obtain gini = 0.228 and with tuning we obtain gini = 0.230

2) In the project2.py, I change the random value from 37 to 2018 in sampling model to obtain new sample as well as changing random in modeling from 2017 to 2018. Then, I tune the hyper parameters for the new model. I obtain a new gini = 0.224 score

3) using different models to predict because usually model such as logistic regression tends to under fitting and modeled such as xgboost and lightbm tend to over fitting. Therefore, we stack these model to overcome the under fitting and over fitting problem. For example in the level two we can see we use different method in xgboost model, with raking without ranking in order to try different model in our prediction and different gini score.

4) The models are stacked together in three deferent levels in order to improve the results. Therefore, we obtain the higher gini score after three levels.

Finally by considering the above model and discussions, at the level four which is mean of level three output, ( logistic regression, xgboost) the gini score = 0.230 is obtained.

## 4.2. Justification

In comparison in our benchmark model, simple logistic regression gini = 0.19 score, we obtained less gini = 0.23 which shows the improvement on our modeling. The submission results in 0.27947 leader board score in competition. Top 40 %.

## 5. Conclusion

### 5.1. Free-From Visualization

one of the main problem of this classification challenge is imbalance target. Fig. 7 shows that the undersampling approach can solve the imbalance sample.

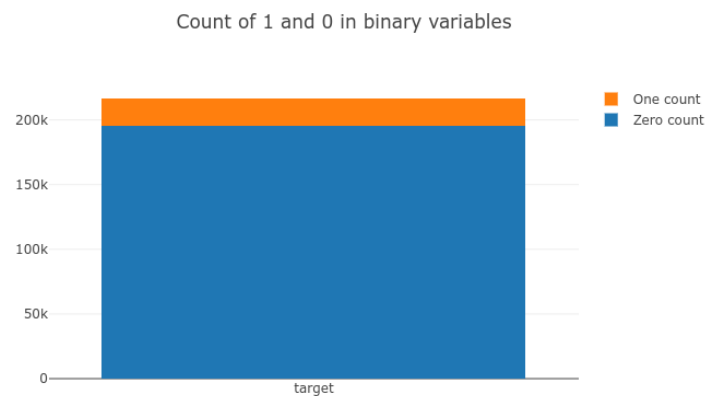
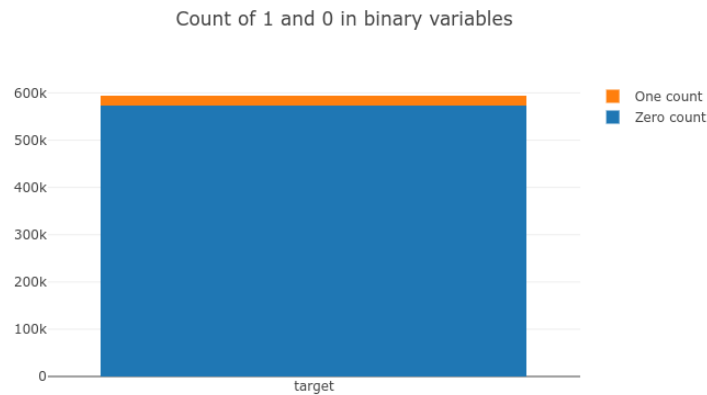


Fig.7: Figures show the target values before and after undersampling respectively.

The main approach to solve this binary classification challenge is stacking of different machine learning algorithms. Therefore, it is necessary to investigate whether the gini score improves through this approach. Therefore, we show how the mean of the gini score changes for each level in Fig. 8. It is seen that the mean of gini score is increasing by increasing the level. It means that the prediction is getting better in level 3.

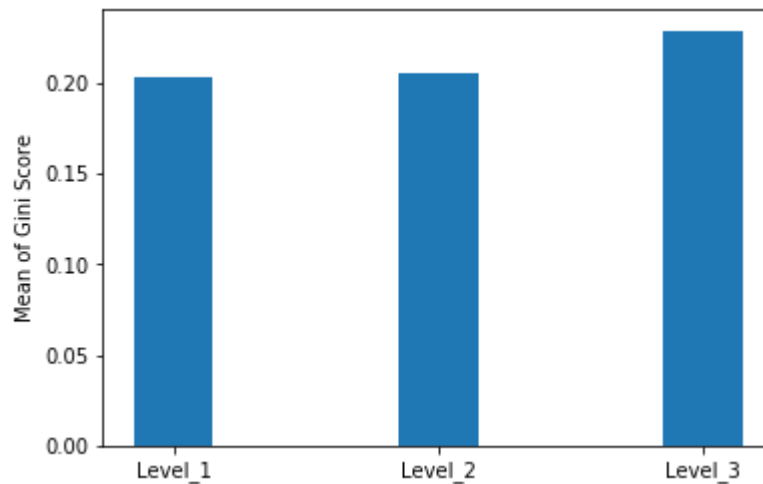


Fig.8: Figures show the target values before and after undersampling respectively.

## 5.2. Reflection

In this project, I work on the dataset from kaggle competition. The dataset is provided by the Porto Seguro company. Porto Seguro, one of Brazil's largest auto and homeowner insurance companies. Inaccuracies in car insurance company's claim predictions raise the cost of insurance for good drivers and reduce the price for bad ones. The goal of this project was to build a model that predicts the probability that a driver will initiate an auto insurance claim in the next year. After the target exploration, I figure out that the dataset target is classified into 0 and 1. Thus, the problem is binary classification. Since the problem is binary classification, the gini metric is



used for the calculation of the cross validation score. The interesting aspect of this binary problem is found after the statical exploration of the target. The exploration shows that the target values are imbalance. Therefore, the undersampling method is applied to over come this problem as well as to obtain the new sampling from the dataset. The data analysis shows that we have binary, categorical, numerical features. We use binary encoding and frequency encoding to extract the new features from the categorical values. The new features from numerical features are obtained by using regression model with polynomial kernels. The data engineer part of the sample was the most difficult part of the project because I had to find new relevant features to improve the prediction score. After data analysis and feature engineering, several machine learning algorithms, Logistic Regression, Random Forest, Brenoulli, Xgboost, Light GBM are stacked together in three levels to make prediction for this classification challenge. Therefore, we reached the gini = 0.23 score for predicting the target value. Additionally, I also apply the whole machine learning pipeline with different random state and I obtained gini = 0.224.

Therefore, the machine learning pipeline is design for solving the binary classification problem which can be used for the similar classification problem.

### ***5.3. Improvement***

The result of the calculation can be improved by using target encoding and stacking more diverse machine learning algorithm. Moreover, the stacking model also can be improved by giving the wight for each model and then tuning that wight. In the other word, every model has wight = 1 now but it is possible to tune this wight too.