

## ▼ Loan Prediction

The aim of this project is to predict real-estate prices using the machine learning algorithms: Logistic Regression, Decision tree, Random Forest. The three of them will show different results for the accuracy.

## Loading the data

In the previous phase - Provisioning, data collection, data cleaning and data preparation were performed on a data scraped from website. In this file, this processed data will be used to train the models.

## Modelling

In this stage, I decided to use several models and eventually I can decide which one performed the best in order to use in the next phase - Deployment. I will explore the machine learning algorithms: Logistic Regression, Decision tree, Random Forest. All three will show different results for the accuracy. I decided to use these four models so as to check more features for comparing and different aspects.

I will compare the models by calculating the MAE, MSE, RMSE and the accuracy.

## Evaluation

After training the models, it is important for the next steps to find out how good the performance of the models is. Based on this, it will be possible to conclude whether Modelling & Evaluation is successful or not.

## ▼ Imports

```
import pandas as pd
import requests
import matplotlib.pyplot as plt
import seaborn as sns
from google.colab import files
from datetime import datetime
import io
import mpl_toolkits
```

```
import numpy as np
%matplotlib inline

# Load the data
local_file = files.upload()
train_data = io.BytesIO(local_file['results.csv'])
df = pd.read_csv(train_data)
```

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Upload widget is only available when the cell has been executed in the current browser session. Please rerun this cell to enable.

Saving results.csv to results (2).csv

## ▼ Preparing the data for training the models

Encoding to numeric data in order to start the training of the models.

```
#drop the uniques loan id
df.drop('Loan_ID', axis = 1, inplace = True)

df.drop('Unnamed: 0', axis = 1, inplace = True)

df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 981 entries, 0 to 980
Data columns (total 14 columns):
#   Column                Non-Null Count  Dtype
---  -
0   Gender                 981 non-null    float64
1   Married                981 non-null    float64
2   Education              981 non-null    int64
3   Self_Employed          981 non-null    float64
4   ApplicantIncome        981 non-null    int64
5   CoapplicantIncome      981 non-null    float64
6   LoanAmount             981 non-null    float64
7   Loan_Amount_Term       981 non-null    float64
8   Credit_History         981 non-null    float64
9   Property_Area          981 non-null    int64
10  Loan_Status            981 non-null    float64
11  LoanAmount_log         981 non-null    float64
12  TotalIncome            981 non-null    float64
13  TotalIncome_log        981 non-null    float64
dtypes: float64(11), int64(3)
memory usage: 107.4 KB
```

### Train-Test Split dataset

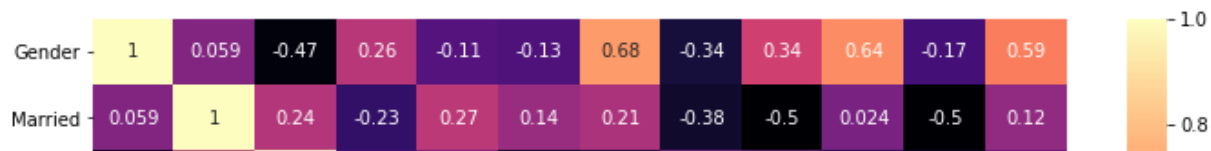
Heatmaps are very useful to find relations between two variables in a dataset and this way the user gets a visualisation of the numeric data. No correlations are extremely high. Each square shows the correlation between the variables on each axis.

- The correlations between the features can be explained:

The close to 1 the correlation is the more positively correlated they are; that is as one increases so does the other and the closer to 1 the stronger this relationship is. It is noticeable that the correlation between the ApplicantIncome and LoanAmount is 0.57, which means that they have a positive correlation, but not strong.

```
from pandas import DataFrame
%matplotlib inline
plt.figure(figsize=(12, 8))
df_temp = df.copy()
Index= ['Gender', 'Married', 'Dependents', 'Education', 'Self_Employed', 'ApplicantIncome']
Cols = ['Gender', 'Married', 'Dependents', 'Education', 'Self_Employed', 'ApplicantIncome']
df_temp = DataFrame(abs(np.random.randn(12, 12)), index=Index, columns=Cols)

sns.heatmap(df_temp.corr(), annot=True, cmap = 'magma')
plt.show()
```



## Importing sklearn libraries

```
from sklearn.model_selection import train_test_split
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.linear_model import LogisticRegression
from sklearn.metrics import f1_score
```



## Splitting into train and test set after choosing the right features X and labels y

```
y = df['Loan_Status']
X = df.drop('Loan_Status', axis = 1)
```



To split the dataset, I will use random sampling with 80/20 train-test split; that is, 80% of the dataset will be used for training and set aside 20% for testing:

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.20, random_state=0)
```

## Analyzing the numeric features.

```
numeric_features = df.select_dtypes(include=[np.number])
```

```
numeric_features.columns
```

```
Index(['Gender', 'Married', 'Education', 'Self_Employed', 'ApplicantIncome',
      'CoapplicantIncome', 'LoanAmount', 'Loan_Amount_Term', 'Credit_History',
      'Property_Area', 'Loan_Status', 'LoanAmount_log', 'TotalIncome',
      'TotalIncome_log'],
      dtype='object')
```

```
# use only those input features with numeric data type
df = df.select_dtypes(include=["int64", "float64"])
```

```
# set the target and predictors
y = df.Loan_Status # target
```

```
# use only those input features with numeric data type
df_temp = df.select_dtypes(include=["int64", "float64"])
```

```
X = df_temp.drop(["Loan_Status"],axis=1) # predictors
```

## ▼ Modeling:

Three models will be built and evaluated by their performances with R-squared metric. Additionally, insights on the features that are strong predictors of house prices, will be analysed .

```
from sklearn.metrics import mean_squared_error
from sklearn.metrics import mean_absolute_error
```

### Logistic Regression:

1. Creating
2. Fitting with train data

```
model = LogisticRegression()
model.fit(X_train, y_train)
```

```
/usr/local/lib/python3.7/dist-packages/sklearn/linear_model/_logistic.py:940: ConvergenceWarning:
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>

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/modules/linear_model.html#logistic-regression)

```
extra_warning_msg=_LOGISTIC_SOLVER_CONVERGENCE_MSG)
LogisticRegression(C=1.0, class_weight=None, dual=False, fit_intercept=True,
                    intercept_scaling=1, l1_ratio=None, max_iter=100,
                    multi_class='auto', n_jobs=None, penalty='l2',
                    random_state=None, solver='lbfgs', tol=0.0001, verbose=0,
                    warm_start=False)
```

Root Mean Square Error (RMSE) is the standard deviation of the residuals (prediction errors). Residuals are a measure of how far from the regression line data points are; RMSE is a measure of how spread out these residuals are. In other words, it tells you how concentrated the data is around the line of best fit.

RMSE values between 0.2 and 0.5 shows that the model can relatively predict the data accurately.

```
# model evaluation for training set
```

```

y_train_r_predict = model.predict(X_train)
rmse = (np.sqrt(mean_squared_error(y_train, y_train_r_predict)))

print("The model performance for training set:")
print('RMSE is {}'.format(rmse))

The model performance for training set:
RMSE is 0.43594841484763225

```

Do predictions on a test set. **Testing** the model by testing the test data.

```

#predict y_values using X_test set
y_reg=model.predict(X_test)

```

Comparing these metrics:

MAE is the easiest to understand because it's the average error. MSE is more popular than MAE because MSE "punishes" larger errors, which tends to be useful in the real world. RMSE is even more popular than MSE because RMSE is interpretable in the "y" units.

```

# model evaluation for testing set
print('MAE:', mean_absolute_error(y_test, y_reg))
print('MSE:', mean_squared_error(y_test, y_reg))
print('RMSE:', np.sqrt(mean_squared_error(y_test, y_reg)))

MAE: 0.19796954314720813
MSE: 0.19796954314720813
RMSE: 0.44493768456628635

```

```

logistic_score =model.score((X_test),y_test)
print("Accuracy: ", logistic_score)

```

```
Accuracy: 0.8020304568527918
```

The F1 score can be interpreted as a weighted average of the precision and recall, where an F1 score reaches its best value at 1 and worst score at 0.

```

evaluation = f1_score(y_test, y_reg)
evaluation

```

```
0.8869565217391304
```

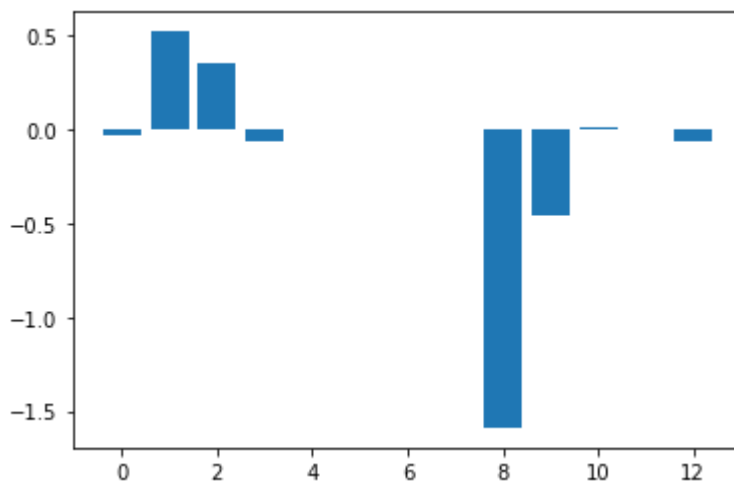
Reporting the coefficient value for each feature. Notice that the coefficients are both positive and negative. The positive scores indicate a feature that predicts class 1, whereas the negative scores

indicate a feature that predicts class 0.

The importance of a feature is measured by calculating the increase in the model's prediction error after permuting the feature. A feature is "important" if shuffling its values increases the model error, because in this case the model relied on the feature for the prediction.

```
# get importance
importance = model.coef_[0]
# summarize feature importance
for i,v in enumerate(importance):
    print('Feature: %0d, Score: %.5f' % (i,v))
# plot feature importance
plt.bar([x for x in range(len(importance))], importance)
plt.show()
```

```
Feature: 0, Score: -0.02992
Feature: 1, Score: 0.52301
Feature: 2, Score: 0.35065
Feature: 3, Score: -0.06703
Feature: 4, Score: -0.00001
Feature: 5, Score: 0.00002
Feature: 6, Score: 0.00293
Feature: 7, Score: -0.00061
Feature: 8, Score: -1.58855
Feature: 9, Score: -0.46054
Feature: 10, Score: 0.00910
Feature: 11, Score: 0.00001
Feature: 12, Score: -0.06359
```



What coefficient of data says:

It shows the relationship between the features. It is either positive or negative, depending on the value shown on the graph. If it is below 0, it is a negative one, which means that whenever there is an increase in the value, there is a decrease in the price. What goes for the positive relationship - there is a parallel movement.

### Decision tree:

1. Creating classifier
2. Fitting classifier with train data

```
dtree = DecisionTreeClassifier()
dtree.fit(X_train, y_train)

DecisionTreeClassifier(ccp_alpha=0.0, class_weight=None, criterion='gini',
                        max_depth=None, max_features=None, max_leaf_nodes=None,
                        min_impurity_decrease=0.0, min_impurity_split=None,
                        min_samples_leaf=1, min_samples_split=2,
                        min_weight_fraction_leaf=0.0, presort='deprecated',
                        random_state=None, splitter='best')

# model evaluation for training set
y_train_r_predict = dtree.predict(X_train)
rmse = (np.sqrt(mean_squared_error(y_train, y_train_r_predict)))

print("The model performance for training set:")
print('RMSE is {}'.format(rmse))

The model performance for training set:
RMSE is 0.0
```

Do predictions on a test set. **Testing** the model by testing the test data.

```
y_tree=dtree.predict(X_test)
```

Comparing these metrics:

MAE is the easiest to understand because it's the average error. MSE is more popular than MAE because MSE "punishes" larger errors, which tends to be useful in the real world. RMSE is even more popular than MSE because RMSE is interpretable in the "y" units.

```
# model evaluation for testing set
print('MAE:', mean_absolute_error(y_test, y_tree))
print('MSE:', mean_squared_error(y_test, y_tree))
print('RMSE:', np.sqrt(mean_squared_error(y_test, y_tree)))
```

```
MAE: 0.2131979695431472
```



```
MSE: 0.2131979695431472
RMSE: 0.4617336564981453
```

There is a 0.21 improvement, determining this from the MAE

```
tree_score =dtree.score((X_test),y_test)
print("Accuracy: ", tree_score)
```

```
Accuracy: 0.7868020304568528
```

The F1 score can be interpreted as a weighted average of the precision and recall, where an F1 score reaches its best value at 1 and worst score at 0.

```
evaluation = f1_score(y_test, y_tree)
evaluation
```

```
0.8653846153846154
```

Evaluate classssifier measures accuracy by using F1 score. The result shows that the model is precise.

## Random forests

```
# Feature Scaling
from sklearn.preprocessing import StandardScaler
from sklearn.ensemble import RandomForestRegressor
from sklearn.model_selection import cross_val_score, train_test_split, GridSearchCV

sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

regressor = RandomForestRegressor(n_estimators=20, random_state=0)
regressor.fit(X_train, y_train)
y_pred = regressor.predict(X_test)
```

Comparing these metrics:

MAE is the easiest to understand because it's the average error. MSE is more popular than MAE because MSE "punishes" larger errors, which tends to be useful in the real world. RMSE is even more popular than MSE because RMSE is interpretable in the "y" units.

```
from sklearn import metrics

print('Mean Absolute Error:', metrics.mean_absolute_error(y_test, y_pred))
print('Mean Squared Error:', metrics.mean_squared_error(y_test, y_pred))
print('Root Mean Squared Error:', np.sqrt(metrics.mean_squared_error(y_test, y_pred)))

Mean Absolute Error: 0.23807106598984776
Mean Squared Error: 0.12558375634517768
Root Mean Squared Error: 0.354377985130535
```

```
# Import the model we are using
from sklearn.ensemble import RandomForestRegressor
# Instantiate model with 1000 decision trees
rf = RandomForestRegressor(n_estimators = 1000, random_state = 42)
# Train the model on training data
rf.fit(X_train, y_train)

RandomForestRegressor(bootstrap=True, ccp_alpha=0.0, criterion='mse',
                       max_depth=None, max_features='auto', max_leaf_nodes=None,
                       max_samples=None, min_impurity_decrease=0.0,
                       min_impurity_split=None, min_samples_leaf=1,
                       min_samples_split=2, min_weight_fraction_leaf=0.0,
                       n_estimators=1000, n_jobs=None, oob_score=False,
                       random_state=42, verbose=0, warm_start=False)
```

```
# Use the forest's predict method on the test data
predictions = rf.predict(X_test)
# Calculate the absolute errors
errors = abs(predictions - y_test)
# Print out the mean absolute error (mae)
print('Mean Absolute Error:', round(np.mean(errors), 2), 'degrees.')
```

Mean Absolute Error: 0.24 degrees.

There is a 0.24 improvement.

```
# Calculate mean absolute percentage error (MAPE)
mape = 100 * (errors / y_test)
# Calculate and display accuracy
accuracy = 100 - np.mean(mape)
print('Accuracy:', round(accuracy, 2), '%.')
```

Accuracy: 81.91 %.

```
#Random forest determined feature importances
rf.feature_importances_
```

array([0.01838005, 0.02412061, 0.02132158, 0.0197354 , 0.17087993,

```
0.10110834, 0.1105108 , 0.04064278, 0.12931742, 0.03864331,
0.11603298, 0.10494534, 0.10436145])
```

```
forest = RandomForestClassifier()
forest.fit(X_train, y_train)
```

```
RandomForestClassifier(bootstrap=True, ccp_alpha=0.0, class_weight=None,
                        criterion='gini', max_depth=None, max_features='auto',
                        max_leaf_nodes=None, max_samples=None,
                        min_impurity_decrease=0.0, min_impurity_split=None,
                        min_samples_leaf=1, min_samples_split=2,
                        min_weight_fraction_leaf=0.0, n_estimators=100,
                        n_jobs=None, oob_score=False, random_state=None,
                        verbose=0, warm_start=False)
```

**Testing** the model by testing the test data.

```
y_forest=forest.predict(X_test)
```

The F1 score can be interpreted as a weighted average of the precision and recall, where an F1 score reaches its best value at 1 and worst score at 0.

Result:

```
evaluation_f= f1_score(y_test, y_forest)
evaluation_f
```

```
0.8975903614457832
```

After using the F1, it is determined that the model is precised to be used in the deployment.

```
importance = rf.feature_importances_

# map feature importance values to the features
feature_importances = zip(importance, X.columns)

#list(feature_importances)
sorted_feature_importances = sorted(feature_importances, reverse = True)

#print(sorted_feature_importances)
top_15_predictors = sorted_feature_importances[0:15]
values = [value for value, predictors in top_15_predictors]
predictors = [predictors for value, predictors in top_15_predictors]
print(predictors)
```

```
['ApplicantIncome', 'Credit_History', 'LoanAmount_log', 'LoanAmount', 'TotalIncome', 'Tc
```

Saving the model that I am going to use in the deployment phase of the project

```
# Saving the model
import pickle

filename = 'classifier.pkl'
pickle.dump(forest, open(filename, 'wb'))
```

## Conclusion

I used three models to determine the accuracy - Logistic Regression, Decision Tree and Random Forest.

From the exploring of the models RMSE:

- Linear Regression score: 0.44
- Decision Tree score: 0.46
- Random forest score: 0.36

RMSE values between 0.2 and 0.5 shows that the model can relatively predict the data accurately. All of the models showed values in this range.

From the exploring of the models accuracy:

- Linear Regression score: 0.73 (73%)
- Decision Tree score: 0.79 (79%)
- Random forest score: 81.91 %

From the exploring of the models after the F1 score validation:

- Linear Regression score: 0.89
- Decision Tree score: 0.6532616143265344
- Random forest: 0.89

Random forest turns out to be the more accurate model for predicting the house price.

All of the models showed RMSE values between 0.2 and 0.5 so that they show relatively accurate predictions of the data.

I evaluated the models performances with F1 score metric and the one that is overfitting the least is the Random forest.

In the end, I tried three different models and evaluated them using Mean Absolute Error. I chose MAE because it is relatively easy to interpret and outliers aren't particularly bad in for this type of model. The one I will be using for the deplyment is the **Random forest**.

