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INTRODUCTION

- Objective: To predict credit card approval status (Approved/Not Approved) using classification techniques.
- Overview: Our Datasets involve the data of many clients like their Income, Date of Birth, Occupation, Education level, Relationship Status, Housing Status, etc.

Credit Scores to Predict

Approved Status

- 'C': Likely denotes a "Credit Approved" status.
- 'X': Possibly stands for "Accepted" or "Approved." These statuses indicate that the application was successful, so they are grouped under the same category.

Not Approved Status

- '0': Application not approved, but no specific issue provided.
- '1': Denial due to minor reasons, such as insufficient credit history.
- '2': Denial due to more severe issues, such as a high debt-to-income ratio.
- '3', '4', '5': Represent escalating levels of rejection or issues, such as bankruptcy, delinquency, or fraud.

1. DATA EXPLORATION AND STATISTICAL ANALYSIS

Dataset Description:

- Provide details about the credit_record and application_record datasets.
 - credit_record dataset contained the following attributes:



Figure 1: credit_record dataset

application_record dataset contained the following attributes:

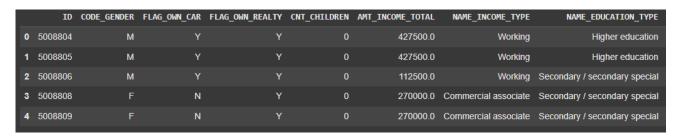


Figure 2: application_record dataset

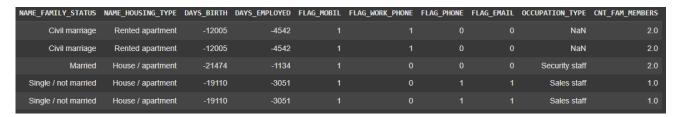


Figure 3: application_record dataset continued

EDA (Exploratory Data Analysis):

We found that there were missing values in the coloum "Occupation Title" and there were duplicates in both datasets which needed to be dealt with.

2. DATA PREPROCESSING

Steps Taken:

We Addressed the null and duplicate values by firstly identifying them using isnull() function.

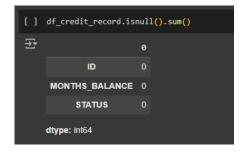


Figure 4: Identifying Null Values (credit_record)

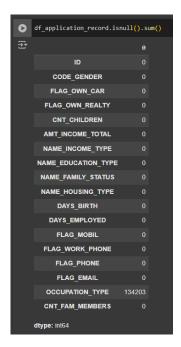


Figure 5: Identifying Null Values (application_record) (30.6%)

The Merge was done based on ID

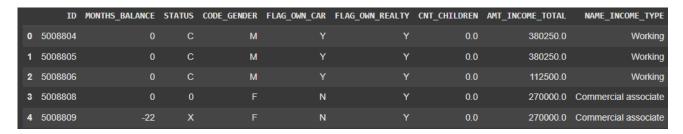


Figure 6: Merged Dataset Based On ID



Figure 7: Merged Dataset Based On ID Continued

- Regarding Scaling, we have done it to fix continuous values like the month balance for example.
- Regarding label encoding, we converted categorical columns to numerical values to help the models and feature scaling function better.
- Regarding SMOTE and SMOTEENN, this after implementation, helped dramatically to fix class imbalance.
- Feature Engineering helped us simplify our data for further optimization to help achieve better accuracy.

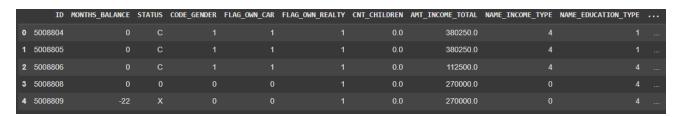


Figure 8: Feature Engineering Results

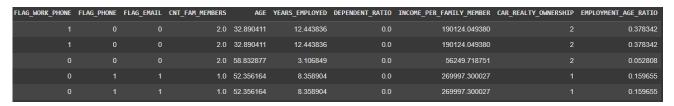


Figure 9: Feature Engineering Results Continued

• The Heatmap allowed us to visualize the correlation between every column and status column to see the effect of each column on the final classification/output.

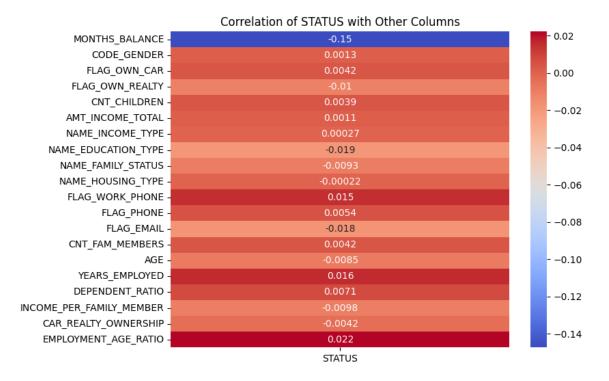


Figure 10: Heatmap

3. FEATURE SELECTION

Genetic Algorithm Approach:

In this project, GAs are applied to feature selection, aiming to identify the most important features that maximize the model's predictive performance. This helps reduce dimensionality, improve model interpretability, and enhance computational efficiency.

Process of Feature Selection Using Genetic Algorithms

• Initialization of Population:

- A population of chromosomes is generated, where each chromosome represents a subset of features.
- A chromosome is a binary array (True for selected features, False otherwise),
 with a mix of included and excluded features initialized randomly.

Fitness Evaluation:

- The fitness of each chromosome is measured by training the model (e.g., DecisionTreeClassifier) on the selected features and evaluating its accuracy on the validation set.
- The fitness_score function ranks chromosomes based on their validation accuracy.

Selection of Elites and Parents:

- Elite Selection: Top-performing chromosomes are retained to ensure the best solutions persist across generations.
- Parent Selection: A subset of top chromosomes is chosen for producing offspring via crossover.

Crossover:

 Chromosomes are combined to create new offspring by swapping segments between two parents at a crossover point. This introduces diversity while preserving strong traits.

Mutation:

 Adaptive mutation modifies certain genes in chromosomes based on the mutation rate and generation. This prevents premature convergence and explores alternative solutions.

Generations:

The GA iteratively evolves the population for a fixed number of generations.
 At each step, the best-performing chromosomes and scores are tracked.

Results of Feature Selection

Selected Features:

After running the GA, the features selected by the best-performing chromosome are used for model training. These features are deemed most significant for predicting credit card approval status.

• Importance to Models:

The selected features improve model accuracy and reduce overfitting by eliminating irrelevant or redundant data. Additionally, they speed up training and evaluation processes.

4. DATA SPLITTING

Procedure:

• We split the dataset into training (70%), validation (15%), and testing (15%).

```
[ ] def split_data(df, label):
    X_train, X_temp, y_train, y_temp = train_test_split(df, label, test_size=0.3, random_state=42)
    X_val, X_test, y_val, y_test = train_test_split(X_temp, y_temp, test_size=0.5, random_state=42)
    return X_train, X_val, X_test, y_train, y_val, y_test
```

Figure 11: Data Splitting

5. MODEL SELECTION AND TRAINING

Algorithms Used:

KNN, Decision Trees, Random Forest, and MLP were used.

Figure 12: Model Selection

• Each Model was trained using the genetic algorithm explained above.

```
[ ] def evaluate_model(model, X_test, y_test):
    predictions = model.predict(X_test)
             accuracy = accuracy_score(y_test, predictions)
             print("Accuracy:", accuracy)
print("Confusion Matrix:")
             print(confusion_matrix(y_test, predictions))
             print(classification_report(y_test, predictions))
             return accuracy
       final_data = standardized_df.copy()
final_data["STATUS"] = y_resampled.values
      label = final_data["STATUS"]
final_data = final_data.drop(columns=["STATUS"], axis=1)
n_features = final_data.shape[1]
       faint_results = {|
for model_name, (model, params) in models_and_params.items():
    print(f"\nRunning Genetic Algorithm with {model_name}...")
    best_chromosomes, best_scores, X_train, y_train, X_test, y_test = generations(final_data,
    label, size=50, n_feat=n_features, n_parents=20, mutation_rate=0.2, n_gen=5,
    model=model, elite_size=2)
             best_chromosome = best_chromosomes[-1]
             selected_features = final_data.columns[best_chromosome]
             print(f"\nHyper-parameter tuning for {model_name}...")
tuned_model = hyperparameter_tuning(model, params, X_train.iloc[:, best_chromosome], y_train)
             print(f"\nEvaluating {model_name}...")
accuracy = evaluate_model(tuned_model, X_test.iloc[:, best_chromosome], y_test)
             final_results.append((model_name, accuracy))
       final_results.sort(key=lambda x: x[1], reverse=True)
       for model_name, accuracy in final_results:
    print(f"{model_name}: {accuracy:.4f}")
       best_model_name, best_model_accuracy = final_results[0]
                   "\nBest Model: {best_model_name} with Accuracy: {best_model_accuracy:.4f}")
```

Figure 13: Model Training

6. HYPERPARAMETER TUNING

- Approach:
 - We used grid-search for each model.
- Results:

Figure 14: Hyperparameter Tuning

```
Hyper-parameter tuning for KNN...
Best Parameters: {'n_neighbors': 3, 'weights': 'distance'}
```

Figure 15: Hyperparameter Tuning Output

7. MODEL EVALUATION

Metrics:

KNN:

```
Evaluating KNN...
Accuracy: 0.9852468690577667
Confusion Matrix:
[[ 808 15
                       0
                               20
                                    33]
    3 2077
                                     1]
       0 2586 0 0
                                     2]
            0 2773
        0
                     0
                           0
                                0
                                     0]
    0
             0 0 2607
    0
                                     0]
                0 0 2405
                                     0]
                2
0
                                   44]
   20
             0
                      0
                           1 692
                              24 1078]]
             a
   28
                      0
Classification Report:
             precision
                        recall f1-score
                                           support
                           0.92
                 0.94
                                    0.93
                                              880
                 0.98
                           1.00
                                    0.99
                                              2085
                                    1.00
                                             2588
                 1.00
                          1.00
                 1.00
                          1.00
                                    1.00
                                             2773
                 1.00
                           1.00
                                    1.00
                                              2607
                 1.00
                           1.00
                                    1.00
                                             2405
                 0.94
                           0.90
                                    0.92
                                              767
                 0.93
                           0.94
                                    0.94
                                             1146
          Х
                                    0.99
                                            15251
   accuracy
  macro avg
                 0.97
                           0.97
                                    0.97
weighted avg
                 0.99
                           0.99
                                    0.99
```

Figure 16: KNN

MLP:

```
Evaluating MLP...
Accuracy: 0.9410530457019212
Confusion Matrix:
[[ 616 39
                              0 120
                                      101]
    8 2058
              a
                  0
                         a
                                        5]
    0 0 2577 0 0 11
0 0 0 2773 0 0
                                   0
                                         0]
              0 2773 0 0
0 0 2607 0
                                        0]
    0
         0
                                   0
                                        0]
              1 0 0 2403 0
5 1 0 7 411
0 0 0 7 118
                                        0]
    0
       26
12
                                      187]
                              7 118 907]]
 T 102
Classification Report:
                           recall f1-score
           0
                   0.72
                             0.70
                                       0.71
                                                   880
                   0.96
                             0.99
                                       0.98
                                                  2085
                   1.00
                             1.00
                                       1.00
                                                  2588
                             1.00
                                       1.00
                                                  2773
                   1.00
                   1.00
                             1.00
                                        1.00
                                                  2607
                   0.99
                             1.00
                                        0.99
                                                  2405
                   0.63
                             0.54
                                        0.58
                                                  767
                                                  1146
                   0.76
                             0.79
                                        0.77
                                        0.94
                                                 15251
    accuracy
                   0.88
                             0.88
  macro avg
                                        0.88
                                                 15251
weighted avg
                   0.94
                             0.94
                                        0.94
                                                 15251
```

Figure 17: MLP

Random Forest:

```
Evaluating RandomForest...
Accuracy: 0.9885253426004852
Confusion Matrix:
[[ 812 24
[ 0 2085
                                           26]
                                      16
                   0
                                            0]
                                            øj
          0 2588
             0 2773
                                            01
                     0 2607
                                            øj
     0
                      0 0 2405
                                 0 699
                                           40]
                                     17 1107]]
Classification Report:
               precision
                              recall f1-score
                                                   support
                     0.95
                                0.92
                                           0.94
                                                       880
                     0.98
                                1.00
                                           0.99
                                                       2085
                     1.00
                                                       2588
                                1.00
                                           1.00
                     1.00
                                1.00
                                           1.00
                     1.00
                                1.00
                                           1.00
                     1.00
                                1.00
                                           1.00
                                                       2405
                     0.95
                                           0.93
                     0.94
                                           0.95
                                                       1146
    accuracy
   macro avg
                     0.98
                                0.98
                                           0.98
                     0.99
                                           0.99
weighted avg
```

Figure 18: Random Forest

Decision Tree:

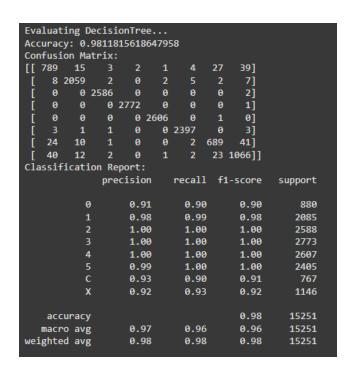


Figure 19: Decision Tree

8. CONCLUSION

Conclusion:

• We Found that the Random Forest Model was best at predicting credit card scores approval. Followed by, KNN and the rest of the models used.

```
Final Accuracy Results:
RandomForest: 0.9885
KNN: 0.9852
DecisionTree: 0.9812
MLP: 0.9411
Best Model: RandomForest with Accuracy: 0.9885
```

Figure 20: Models Comparison

• Following the steps that were given to us and understanding them and implementing them in python and seeing results helped us understand these concepts and allowed us to take a step forward in our learning path.