

# Credit Fraud Detector

#### Introduction

In this kernel we will use various predictive models to see how accurate they are in detecting whether a transaction is a normal payment or a fraud. As described in the dataset, the features are scaled and the names of the features are not shown due to privacy reasons. Nevertheless, we can still analyze some important aspects of the dataset. Let's start!

#### Our Goals:

- Understand the little distribution of the "little" data that was provided to us.
- Create a 50/50 sub-dataframe ratio of "Fraud" and "Non-Fraud" transactions. (NearMiss Algorithm)
- Determine the Classifiers we are going to use and decide which one has a higher accuracy.
- Create a Neural Network and compare the accuracy to our best classifier.
- Understand common mistaked made with imbalanced datasets.

## Outline:

#### 1. Understanding our data

a) [Gather Sense of our data](#gather)

#### **II. Preprocessing**

- a) Scaling and Distributing
- b) Splitting the Data

#### III. Testing

- a) Testing with Logistic Regression
- b) Neural Networks Testing (Undersampling vs Oversampling)

## References:

• Hands on Machine Learning with Scikit-Learn & TensorFlow by Aurélien

Géron (O'Reilly). CopyRight 2017 Aurélien Géron

- Machine Learning Over-& Undersampling Python/ Scikit/ Scikit-Imblearn by Coding-Maniac
- auprc, 5-fold c-v, and resampling methods by Jeremy Lane (Kaggle Notebook)

```
In [1]: # This Python 3 environment comes with many helpful analytics libraries instal
        # It is defined by the kaggle/python docker image: https://github.com/kaggle/c
        # For example, here's several helpful packages to load in
        # Imported Libraries
        import numpy as np # linear algebra
        import pandas as pd # data processing, CSV file I/O (e.g. pd.read csv)
        import matplotlib.pyplot as plt
        import seaborn as sns
        import time
        # Classifier Libraries
        from sklearn.linear model import LogisticRegression
        import collections
        # Other Libraries
        from sklearn.model selection import train test split
        from sklearn.pipeline import make pipeline
        from imblearn.pipeline import make pipeline as imbalanced make pipeline
        from imblearn.over sampling import SMOTE
        from imblearn.under sampling import NearMiss
        from imblearn.metrics import classification report imbalanced
        from sklearn.metrics import precision score, recall score, f1 score, roc auc s
        from collections import Counter
        from sklearn.model selection import KFold, StratifiedKFold
        import warnings
        warnings.filterwarnings("ignore")
        df = pd.read csv('/Users/avijit/Desktop/Himanshu Project/creditcard.csv')
        df.head()
```

| Out[1]: |   | Time | V1        | V2        | V3       | V4        | V5        | V6        | <b>V</b> 7 |
|---------|---|------|-----------|-----------|----------|-----------|-----------|-----------|------------|
|         | 0 | 0.0  | -1.359807 | -0.072781 | 2.536347 | 1.378155  | -0.338321 | 0.462388  | 0.239599   |
|         | 1 | 0.0  | 1.191857  | 0.266151  | 0.166480 | 0.448154  | 0.060018  | -0.082361 | -0.078803  |
|         | 2 | 1.0  | -1.358354 | -1.340163 | 1.773209 | 0.379780  | -0.503198 | 1.800499  | 0.791461   |
|         | 3 | 1.0  | -0.966272 | -0.185226 | 1.792993 | -0.863291 | -0.010309 | 1.247203  | 0.237609   |
|         | 4 | 2.0  | -1.158233 | 0.877737  | 1.548718 | 0.403034  | -0.407193 | 0.095921  | 0.592941   |

 $5 \text{ rows} \times 31 \text{ columns}$ 

In [2]: df.describe()

| Out[2]: |             | Time          | V1            | V2            | V3            | V            |
|---------|-------------|---------------|---------------|---------------|---------------|--------------|
|         | count       | 284807.000000 | 2.848070e+05  | 2.848070e+05  | 2.848070e+05  | 2.848070e+C  |
|         | mean        | 94813.859575  | 1.168375e-15  | 3.416908e-16  | -1.379537e-15 | 2.074095e-1  |
|         | std         | 47488.145955  | 1.958696e+00  | 1.651309e+00  | 1.516255e+00  | 1.415869e+C  |
|         | min         | 0.000000      | -5.640751e+01 | -7.271573e+01 | -4.832559e+01 | -5.683171e+C |
|         | 25%         | 54201.500000  | -9.203734e-01 | -5.985499e-01 | -8.903648e-01 | -8.486401e-C |
|         | 50%         | 84692.000000  | 1.810880e-02  | 6.548556e-02  | 1.798463e-01  | -1.984653e-C |
|         | <b>75</b> % | 139320.500000 | 1.315642e+00  | 8.037239e-01  | 1.027196e+00  | 7.433413e-C  |
|         | max         | 172792.000000 | 2.454930e+00  | 2.205773e+01  | 9.382558e+00  | 1.687534e+C  |

8 rows × 31 columns

```
In [3]: # Good No Null Values!
df.isnull().sum().max()
```

Out[3]: 0

```
In [4]: df.columns
```

In [5]: # The classes are heavily skewed we need to solve this issue later.
print('No Frauds', round(df['Class'].value\_counts()[0]/len(df) \* 100,2), '% of
print('Frauds', round(df['Class'].value\_counts()[1]/len(df) \* 100,2), '% of th

No Frauds 99.83 % of the dataset Frauds 0.17 % of the dataset

**Note:** Notice how imbalanced is our original dataset! Most of the transactions are non-fraud. If we use this dataframe as the base for our predictive models and analysis we might get a lot of errors and our algorithms will probably overfit since it will "assume" that most transactions are not fraud. But we don't want our model to assume, we want our model to detect patterns that give signs of fraud!

**Distributions:** By seeing the distributions we can have an idea how skewed are these features, we can also see further distributions of the other features. There are techniques that can help the distributions be less skewed which will be implemented in this notebook in the future.

```
fig, ax = plt.subplots(1, 2, figsize=(18,4))
In [7]:
          amount val = df['Amount'].values
          time_val = df['Time'].values
          sns.distplot(amount val, ax=ax[0], color='r')
          ax[0].set title('Distribution of Transaction Amount', fontsize=14)
          ax[0].set_xlim([min(amount_val), max(amount_val)])
          sns.distplot(time val, ax=ax[1], color='b')
          ax[1].set_title('Distribution of Transaction Time', fontsize=14)
          ax[1].set xlim([min(time val), max(time val)])
          plt.show()
                       Distribution of Transaction Amount
                                                                          Distribution of Transaction Time
                                                             1.0
                                                             0.8
         0.0025
         0.0020
                                                            <u>₹</u> 0.6
         0.0015
                                                             0.4
         0.0010
                                                             0.2
         0.0005
         0.0000
                            10000
                                            20000
                                                    25000
                    5000
                                    15000
                                                                   20000
                                                                        40000
                                                                            60000
                                                                                 80000 100000 120000 140000 160000
```

# Scaling and Distributing

In this phase of our kernel, we will first scale the columns comprise of **Time** and **Amount**. Time and amount should be scaled as the other columns. On the other hand, we need to also create a sub sample of the dataframe in order to have an equal amount of Fraud and Non-Fraud cases, helping our algorithms better understand patterns that determines whether a transaction is a fraud or not.

## What is a sub-Sample?

In this scenario, our subsample will be a dataframe with a 50/50 ratio of fraud and non-fraud transactions. Meaning our sub-sample will have the same amount of fraud and non fraud transactions.

## Why do we create a sub-Sample?

In the beginning of this notebook we saw that the original dataframe was heavily imbalanced! Using the original dataframe will cause the following issues:

- **Overfitting:** Our classification models will assume that in most cases there are no frauds! What we want for our model is to be certain when a fraud occurs.
- Wrong Correlations: Although we don't know what the "V" features stand for, it will be useful to understand how each of this features influence the result (Fraud or No Fraud) by having an imbalance dataframe we are not able to see the true correlations between the class and features.

## Summary:

- Scaled amount and scaled time are the columns with scaled values.
- There are **492 cases** of fraud in our dataset so we can randomly get 492 cases of non-fraud to create our new sub dataframe.
- We concat the 492 cases of fraud and non fraud, creating a new subsample.

```
In [8]: # Since most of our data has already been scaled we should scale the columns t
    from sklearn.preprocessing import StandardScaler, RobustScaler

# RobustScaler is less prone to outliers.

std_scaler = StandardScaler()

rob_scaler = RobustScaler()

df['scaled_amount'] = rob_scaler.fit_transform(df['Amount'].values.reshape(-1,
    df['scaled_time'] = rob_scaler.fit_transform(df['Time'].values.reshape(-1,1))

df.drop(['Time', 'Amount'], axis=1, inplace=True)

In [9]: scaled_amount = df['scaled_amount']
    scaled_time = df['scaled_time']

df.drop(['scaled_amount', 'scaled_time'], axis=1, inplace=True)
```

```
df.insert(0, 'scaled_amount', scaled_amount)
df.insert(1, 'scaled_time', scaled_time)

# Amount and Time are Scaled!

df.head()
```

| Out[9]: |   | scaled_amount | scaled_time | V1        | V2        | V3       | V4        | 1       |
|---------|---|---------------|-------------|-----------|-----------|----------|-----------|---------|
|         | 0 | 1.783274      | -0.994983   | -1.359807 | -0.072781 | 2.536347 | 1.378155  | -0.3383 |
|         | 1 | -0.269825     | -0.994983   | 1.191857  | 0.266151  | 0.166480 | 0.448154  | 0.0600  |
|         | 2 | 4.983721      | -0.994972   | -1.358354 | -1.340163 | 1.773209 | 0.379780  | -0.5031 |
|         | 3 | 1.418291      | -0.994972   | -0.966272 | -0.185226 | 1.792993 | -0.863291 | -0.0103 |
|         | 4 | 0.670579      | -0.994960   | -1.158233 | 0.877737  | 1.548718 | 0.403034  | -0.4071 |

 $5 \text{ rows} \times 31 \text{ columns}$ 

## Splitting the Data (Original DataFrame)

Before proceeding with the Random UnderSampling technique we have to separate the original dataframe. Why? for testing purposes, remember although we are splitting the data when implementing Random UnderSampling or OverSampling techniques, we want to test our models on the original testing set not on the testing set created by either of these techniques. The main goal is to fit the model either with the dataframes that were undersample and oversample (in order for our models to detect the patterns), and test it on the original testing set.

```
In [10]: from sklearn.model_selection import train_test_split
    from sklearn.model_selection import StratifiedShuffleSplit

print('No Frauds', round(df['Class'].value_counts()[0]/len(df) * 100,2), '% of print('Frauds', round(df['Class'].value_counts()[1]/len(df) * 100,2), '% of th

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

sss = StratifiedKFold(n_splits=5, random_state=None, shuffle=False)

for train_index, test_index in sss.split(X, y):
    print("Train:", train_index, "Test:", test_index)
    original_Xtrain, original_Xtest = X.iloc[train_index], X.iloc[test_index]
    original_ytrain, original_ytest = y.iloc[train_index], y.iloc[test_index]

# We already have X_train and y_train for undersample data thats why I am usin # original_Xtrain, original_Xtest, original_ytrain, original_ytest = train_tes
```

```
# Check the Distribution of the labels
 # Turn into an array
 original Xtrain = original Xtrain.values
 original Xtest = original Xtest.values
 original ytrain = original ytrain.values
 original ytest = original ytest.values
 # See if both the train and test label distribution are similarly distributed
 train unique label, train counts label = np.unique(original ytrain, return cou
 test unique label, test counts label = np.unique(original ytest, return counts
 print('-' * 100)
 print('Label Distributions: \n')
 print(train counts label/ len(original ytrain))
 print(test counts label/ len(original ytest))
No Frauds 99.83 % of the dataset
Frauds 0.17 % of the dataset
                                                                           2
Train: [ 30473 30496 31002 ... 284804 284805 284806] Test: [ 0
... 57017 57018 57019]
Train: [ 0
                 1
                          2 ... 284804 284805 284806] Test: [ 30473 30496 31
002 ... 113964 113965 113966]
Train: [ 0 1 2 ... 284804 284805 284806] Test: [ 81609 82400 83
053 ... 170946 170947 170948]
Train: [ 0 1
                         2 ... 284804 284805 284806] Test: [150654 150660 150
661 ... 227866 227867 227868]
Train: [ 0 1 2 ... 227866 227867 227868] Test: [212516 212644 213
092 ... 284804 284805 284806]
Label Distributions:
[0.99827076 0.00172924]
[0.99827952 0.00172048]
```

# Random Under-Sampling:

In this phase of the project we will implement "Random Under Sampling" which basically consists of removing data in order to have a more **balanced dataset** and thus avoiding our models to overfitting.

#### Steps:

- The first thing we have to do is determine how imbalanced is our class (use "value\_counts()" on the class column to determine the amount for each label)
- Once we determine how many instances are considered fraud transactions (Fraud = "1"), we should bring the non-fraud

**transactions** to the same amount as fraud transactions (assuming we want a 50/50 ratio), this will be equivalent to 492 cases of fraud and 492 cases of non-fraud transactions.

 After implementing this technique, we have a sub-sample of our dataframe with a 50/50 ratio with regards to our classes. Then the next step we will implement is to **shuffle the data** to see if our models can maintain a certain accuracy everytime we run this script.

**Note:** The main issue with "Random Under-Sampling" is that we run the risk that our classification models will not perform as accurate as we would like to since there is a great deal of **information loss** (bringing 492 non-fraud transaction from 284,315 non-fraud transaction)

```
In [11]: # Since our classes are highly skewed we should make them equivalent in order
# Lets shuffle the data before creating the subsamples

df = df.sample(frac=1)

# amount of fraud classes 492 rows.
fraud_df = df.loc[df['Class'] == 1]
non_fraud_df = df.loc[df['Class'] == 0][:492]

normal_distributed_df = pd.concat([fraud_df, non_fraud_df]))

# Shuffle dataframe rows
new_df = normal_distributed_df.sample(frac=1, random_state=42)
new_df.head()
```

| ut[11]: |        | scaled_amount | scaled_time | V1        | V2        | V3         | V4        |
|---------|--------|---------------|-------------|-----------|-----------|------------|-----------|
|         | 100193 | -0.000279     | -0.202610   | 1.317389  | -0.702860 | 0.728638   | -0.734294 |
|         | 215132 | 9.798225      | 0.649197    | -2.921944 | -0.228062 | -5.877289  | 2.201884  |
|         | 226772 | -0.281981     | 0.706129    | -0.395931 | 1.335044  | -0.186761  | -0.049380 |
|         | 189587 | 0.641375      | 0.514327    | 0.909124  | 1.337658  | -4.484728  | 3.245358  |
|         | 150660 | 0.319989      | 0.107626    | -6.185857 | 7.102985  | -13.030455 | 8.010823  |

 $5 \text{ rows} \times 31 \text{ columns}$ 

#### **Correlation Matrices**

Correlation matrices are the essence of understanding our data. We want to know if there are features that influence heavily in whether a specific transaction is a fraud. However, it is important that we use the correct dataframe (subsample) in order for us to see which features have a high positive or negative correlation with regards to fraud transactions.

#### Summary and Explanation:

- **Negative Correlations:** V17, V14, V12 and V10 are negatively correlated. Notice how the lower these values are, the more likely the end result will be a fraud transaction.
- **Positive Correlations:** V2, V4, V11, and V19 are positively correlated. Notice how the higher these values are, the more likely the end result will be a fraud transaction.
- BoxPlots: We will use boxplots to have a better understanding of the distribution of these features in fradulent and non fradulent transactions.

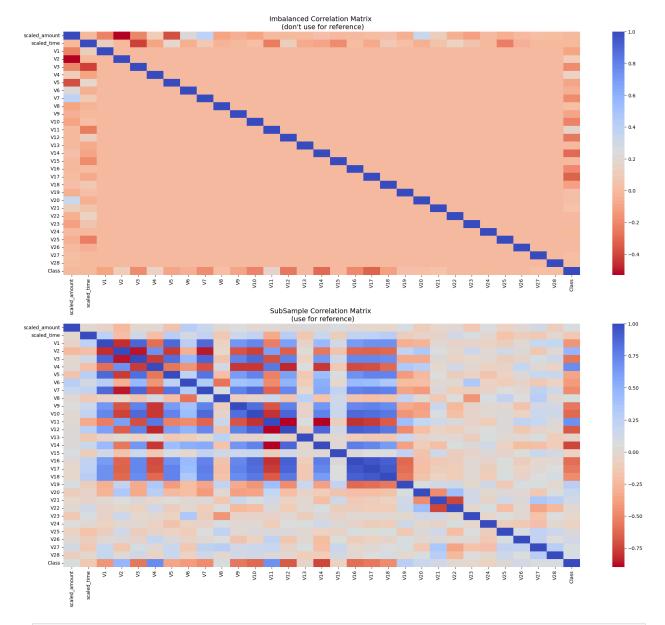
\*\*Note: \*\* We have to make sure we use the subsample in our correlation matrix or else our correlation matrix will be affected by the high imbalance between our classes. This occurs due to the high class imbalance in the original dataframe.

```
In [13]: # Make sure we use the subsample in our correlation

f, (ax1, ax2) = plt.subplots(2, 1, figsize=(24,20))

# Entire DataFrame
corr = df.corr()
sns.heatmap(corr, cmap='coolwarm_r', annot_kws={'size':20}, ax=ax1)
ax1.set_title("Imbalanced Correlation Matrix \n (don't use for reference)", fc

sub_sample_corr = new_df.corr()
sns.heatmap(sub_sample_corr, cmap='coolwarm_r', annot_kws={'size':20}, ax=ax2)
ax2.set_title('SubSample Correlation Matrix \n (use for reference)', fontsize=plt.show()
```



```
In [14]: from scipy.stats import norm

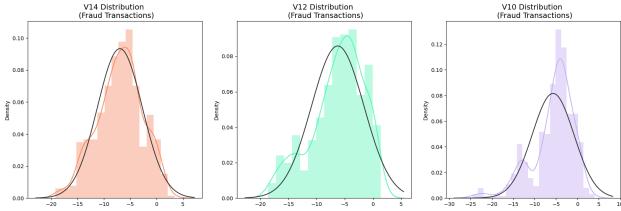
f, (ax1, ax2, ax3) = plt.subplots(1,3, figsize=(20, 6))

v14_fraud_dist = new_df['V14'].loc[new_df['Class'] == 1].values
    sns.distplot(v14_fraud_dist,ax=ax1, fit=norm, color='#FB8861')
    ax1.set_title('V14 Distribution \n (Fraud Transactions)', fontsize=14)

v12_fraud_dist = new_df['V12'].loc[new_df['Class'] == 1].values
    sns.distplot(v12_fraud_dist,ax=ax2, fit=norm, color='#56F9BB')
    ax2.set_title('V12 Distribution \n (Fraud Transactions)', fontsize=14)

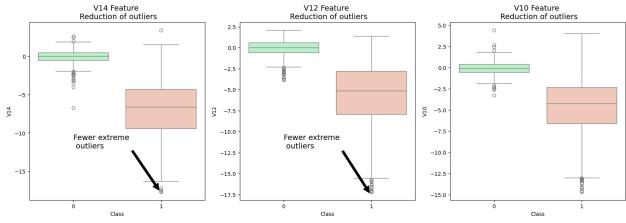
v10_fraud_dist = new_df['V10'].loc[new_df['Class'] == 1].values
    sns.distplot(v10_fraud_dist,ax=ax3, fit=norm, color='#C5B3F9')
    ax3.set_title('V10 Distribution \n (Fraud Transactions)', fontsize=14)
```

#### plt.show()



```
In [15]: # # ----> V14 Removing Outliers (Highest Negative Correlated with Labels)
         v14 fraud = new df['V14'].loc[new df['Class'] == 1].values
         q25, q75 = np.percentile(v14 fraud, 25), <math>np.percentile(v14 fraud, 75)
         print('Quartile 25: {} | Quartile 75: {}'.format(q25, q75))
         v14 iqr = q75 - q25
         print('iqr: {}'.format(v14 iqr))
         v14 cut off = v14 iqr * 1.5
         v14_lower, v14_upper = q25 - v14_cut_off, q75 + v14_cut_off
         print('Cut Off: {}'.format(v14 cut off))
         print('V14 Lower: {}'.format(v14 lower))
         print('V14 Upper: {}'.format(v14 upper))
         outliers = [x for x in v14_fraud if x < v14_lower or x > v14_upper]
         print('Feature V14 Outliers for Fraud Cases: {}'.format(len(outliers)))
         print('V10 outliers:{}'.format(outliers))
         new df = new df.drop(new df['V14'] > v14 upper) | (new df['V14'] < v14
         print('---' * 44)
         # ----> V12 removing outliers from fraud transactions
         v12 fraud = new df['V12'].loc[new df['Class'] == 1].values
         q25, q75 = np.percentile(v12 fraud, 25), np.percentile(v12 fraud, 75)
         v12 iqr = q75 - q25
         v12 cut off = v12 iqr * 1.5
         v12 lower, v12 upper = q25 - v12 cut off, q75 + v12 cut off
         print('V12 Lower: {}'.format(v12 lower))
         print('V12 Upper: {}'.format(v12 upper))
         outliers = [x \text{ for } x \text{ in } v12 \text{ fraud if } x < v12 \text{ lower or } x > v12 \text{ upper}]
         print('V12 outliers: {}'.format(outliers))
         print('Feature V12 Outliers for Fraud Cases: {}'.format(len(outliers)))
         new df = new df.drop(new df['V12'] > V12 upper) | (new df['V12'] < V12
         print('Number of Instances after outliers removal: {}'.format(len(new df)))
         print('---' * 44)
         # Removing outliers V10 Feature
```

```
v10 fraud = new df['V10'].loc[new df['Class'] == 1].values
         q25, q75 = np.percentile(v10 fraud, 25), np.percentile(v10 fraud, 75)
         v10 igr = q75 - q25
         v10 cut off = v10 igr * 1.5
         v10 lower, v10 upper = q25 - v10 cut off, q75 + v10 cut off
         print('V10 Lower: {}'.format(v10 lower))
         print('V10 Upper: {}'.format(v10_upper))
         outliers = [x \text{ for } x \text{ in } v10 \text{ fraud if } x < v10 \text{ lower or } x > v10 \text{ upper}]
         print('V10 outliers: {}'.format(outliers))
         print('Feature V10 Outliers for Fraud Cases: {}'.format(len(outliers)))
         new df = new df.drop(new df['^{10}'] > v10 upper) | (new df['^{10}'] < v16
         print('Number of Instances after outliers removal: {}'.format(len(new df)))
        Quartile 25: -9.692722964972386 | Quartile 75: -4.282820849486865
        igr: 5.409902115485521
        Cut Off: 8.114853173228282
        V14 Lower: -17.807576138200666
        V14 Upper: 3.8320323237414167
        Feature V14 Outliers for Fraud Cases: 4
        V10 outliers:[-18.8220867423816, -19.2143254902614, -18.4937733551053, -18.0499
        976898594]
        V12 Lower: -17.3430371579634
        V12 Upper: 5.776973384895937
        V12 outliers: [-18.0475965708216, -18.5536970096458, -18.6837146333443, -18.431
        13102799931
        Feature V12 Outliers for Fraud Cases: 4
        Number of Instances after outliers removal: 975
        V10 Lower: -14.89885463232024
        V10 Upper: 4.92033495834214
        V10 outliers: [-20.9491915543611, -15.5637913387301, -19.836148851696, -18.2711
        681738888, -15.3460988468775, -23.2282548357516, -24.5882624372475, -18.9132433
        348732, -16.6011969664137, -14.9246547735487, -16.3035376590131, -24.4031849699
        728, -22.1870885620007, -14.9246547735487, -17.1415136412892, -15.123752180345
        5, -15.2318333653018, -22.1870885620007, -22.1870885620007, -15.5637913387301,
        -16.6496281595399, -16.2556117491401, -15.1241628144947, -15.2399619587112, -2
        2.1870885620007, -16.7460441053944, -15.2399619587112]
        Feature V10 Outliers for Fraud Cases: 27
        Number of Instances after outliers removal: 943
In [16]: f_{(ax1, ax2, ax3)} = plt.subplots(1, 3, figsize=(20,6))
         colors = ['#B3F9C5', '#f9c5b3']
         # Boxplots with outliers removed
         # Feature V14
         sns.boxplot(x="Class", y="V14", data=new df,ax=ax1, palette=colors)
         ax1.set title("V14 Feature \n Reduction of outliers", fontsize=14)
         ax1.annotate('Fewer extreme \n outliers', xy=(0.98, -17.5), xytext=(0, -12),
```



```
In [17]: # Undersampling before cross validating (prone to overfit)
X = new_df.drop('Class', axis=1)
y = new_df['Class']
```

In [18]: # Our data is already scaled we should split our training and test sets
 from sklearn.model\_selection import train\_test\_split

# This is explicitly used for undersampling.
X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.2, randometric randomet

```
In [19]: # Turn the values into an array for feeding the classification algorithms.
X_train = X_train.values
X_test = X_test.values
y_train = y_train.values
y_test = y_test.values
```

```
In [20]: # Let's implement simple classifiers
classifiers = {
```

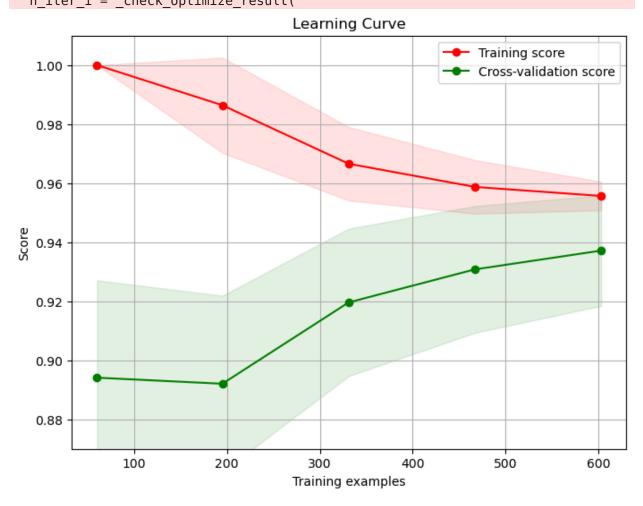
```
"LogisiticRegression": LogisticRegression()
         }
In [21]: # Wow our scores are getting even high scores even when applying cross validat
         from sklearn.model selection import cross val score
         for key, classifier in classifiers.items():
             classifier.fit(X train, y train)
             training score = cross val score(classifier, X train, y train, cv=5)
             print("Classifiers: ", classifier.__class__.__name__, "Has a training scor
       Classifiers: LogisticRegression Has a training score of 94.0 % accuracy score
In [22]: # Use GridSearchCV to find the best parameters.
         from sklearn.model selection import GridSearchCV
         # Logistic Regression
         log reg params = {"penalty": ['l1', 'l2'], 'C': [0.001, 0.01, 0.1, 1, 10, 100,
         grid log reg = GridSearchCV(LogisticRegression(), log reg params)
         grid log reg.fit(X train, y train)
         # We automatically get the logistic regression with the best parameters.
         log reg = grid log reg.best estimator
In [23]: # Overfitting Case
         log reg score = cross val score(log reg, X train, y train, cv=5)
         print('Logistic Regression Cross Validation Score: ', round(log reg score.mean
       Logistic Regression Cross Validation Score: 94.56%
In [24]: # Let's Plot LogisticRegression Learning Curve
         from sklearn.model selection import ShuffleSplit
         from sklearn.model selection import learning curve
         def plot learning curve(estimator1, estimator2, estimator3, estimator4, X, y,
                                 n jobs=1, train sizes=np.linspace(.1, 1.0, 5)):
             f, ((ax1, ax2), (ax3, ax4)) = plt.subplots(2,2, figsize=(20,14), sharey=Tr
             if ylim is not None:
                 plt.ylim(*ylim)
             # First Estimator
             train sizes, train scores, test scores = learning curve(
                 estimator1, X, y, cv=cv, n jobs=n jobs, train sizes=train sizes)
             train_scores_mean = np.mean(train_scores, axis=1)
             train scores std = np.std(train scores, axis=1)
             test scores mean = np.mean(test scores, axis=1)
             test scores std = np.std(test scores, axis=1)
             ax1.fill between(train sizes, train scores mean - train scores std,
                              train_scores_mean + train_scores std, alpha=0.1,
                              color="#ff9124")
```

```
ax1.fill between(train sizes, test scores mean - test scores std,
                 test scores mean + test scores std, alpha=0.1, color="#24
ax1.plot(train sizes, train_scores_mean, 'o-', color="#ff9124",
         label="Training score")
ax1.plot(train sizes, test scores mean, 'o-', color="#2492ff",
         label="Cross-validation score")
ax1.set title("Logistic Regression Learning Curve", fontsize=14)
ax1.set xlabel('Training size (m)')
ax1.set ylabel('Score')
ax1.grid(True)
ax1.legend(loc="best")
# Second Estimator
train sizes, train scores, test scores = learning curve(
    estimator2, X, y, cv=cv, n jobs=n jobs, train sizes=train sizes)
train scores mean = np.mean(train scores, axis=1)
train scores std = np.std(train scores, axis=1)
test_scores_mean = np.mean(test_scores, axis=1)
test scores std = np.std(test scores, axis=1)
ax2.fill_between(train_sizes, train_scores_mean - train_scores_std,
                 train scores mean + train scores std, alpha=0.1,
                 color="#ff9124")
ax2.fill between(train sizes, test scores mean - test scores std,
                 test_scores_mean + test_scores_std, alpha=0.1, color="#24
ax2.plot(train sizes, train scores mean, 'o-', color="#ff9124",
         label="Training score")
ax2.plot(train sizes, test scores mean, 'o-', color="#2492ff",
         label="Cross-validation score")
ax2.set title("Knears Neighbors Learning Curve", fontsize=14)
ax2.set xlabel('Training size (m)')
ax2.set ylabel('Score')
ax2.grid(True)
ax2.legend(loc="best")
# Third Estimator
train_sizes, train_scores, test_scores = learning_curve(
    estimator3, X, y, cv=cv, n jobs=n jobs, train sizes=train sizes)
train scores mean = np.mean(train scores, axis=1)
train scores std = np.std(train scores, axis=1)
test scores mean = np.mean(test scores, axis=1)
test scores std = np.std(test scores, axis=1)
ax3.fill_between(train_sizes, train_scores_mean - train_scores_std,
                 train scores mean + train scores std, alpha=0.1,
                 color="#ff9124")
ax3.fill between(train sizes, test scores mean - test scores std,
                 test scores mean + test scores std, alpha=0.1, color="#24
ax3.plot(train sizes, train scores mean, 'o-', color="#ff9124",
         label="Training score")
ax3.plot(train_sizes, test_scores_mean, 'o-', color="#2492ff",
         label="Cross-validation score")
ax3.set title("Support Vector Classifier \n Learning Curve", fontsize=14)
ax3.set xlabel('Training size (m)')
ax3.set ylabel('Score')
```

```
ax3.legend(loc="best")
             # Fourth Estimator
             train_sizes, train_scores, test_scores = learning_curve(
                 estimator4, X, y, cv=cv, n jobs=n jobs, train sizes=train sizes)
             train scores mean = np.mean(train scores, axis=1)
             train scores std = np.std(train scores, axis=1)
             test scores mean = np.mean(test scores, axis=1)
             test scores std = np.std(test scores, axis=1)
             ax4.fill between(train sizes, train scores mean - train scores std,
                              train scores mean + train scores std, alpha=0.1,
                              color="#ff9124")
             ax4.fill between(train sizes, test scores mean - test scores std,
                              test scores mean + test scores std, alpha=0.1, color="#24
             ax4.plot(train sizes, train scores mean, 'o-', color="#ff9124",
                      label="Training score")
             ax4.plot(train sizes, test scores mean, 'o-', color="#2492ff",
                      label="Cross-validation score")
             ax4.set title("Decision Tree Classifier \n Learning Curve", fontsize=14)
             ax4.set xlabel('Training size (m)')
             ax4.set ylabel('Score')
             ax4.grid(True)
             ax4.legend(loc="best")
             return plt
In [26]: import numpy as np
         import matplotlib.pyplot as plt
         from sklearn.model selection import ShuffleSplit, learning curve
         # Create ShuffleSplit cross-validator
         cv = ShuffleSplit(n splits=100, test size=0.2, random state=42)
         # Plot learning curve function
         def plot learning curve(estimator, X, y, ylim=None, cv=None, n jobs=None, trai
             plt.figure(figsize=(8,6))
             if ylim is not None:
                 plt.ylim(*ylim)
             plt.xlabel("Training examples")
             plt.ylabel("Score")
             train sizes, train scores, test scores = learning curve(
                 estimator, X, y, cv=cv, n jobs=n jobs, train sizes=train sizes
             )
             train scores mean = np.mean(train scores, axis=1)
             train scores std = np.std(train scores, axis=1)
             test scores mean = np.mean(test scores, axis=1)
             test scores std = np.std(test scores, axis=1)
             plt.grid()
             plt.fill between(train sizes, train scores mean - train scores std,
                              train scores mean + train scores std, alpha=0.1, color="r
```

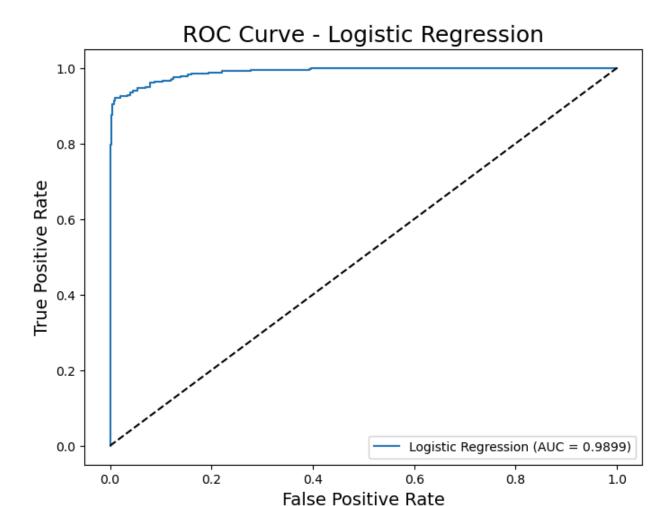
ax3.grid(**True**)

```
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-regressi
on
  n iter i = check optimize result(
/opt/anaconda3/lib/python3.12/site-packages/sklearn/linear model/ logistic.py:4
69: ConvergenceWarning: 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
Please also refer to the documentation for alternative solver options:
    https://scikit-learn.org/stable/modules/linear model.html#logistic-regressi
on
  n iter i = check optimize result(
```



In [27]: from sklearn.metrics import roc\_curve
 from sklearn.model\_selection import cross\_val\_predict
 # Create a DataFrame with all the scores and the classifiers names.
log\_reg\_pred = cross\_val\_predict(log\_reg, X\_train, y\_train, cv=5,

```
method="decision function")
In [28]: from sklearn.metrics import roc auc score
         print('Logistic Regression: ', roc auc score(y train, log reg pred))
       Logistic Regression: 0.9669784309982821
In [30]: from sklearn.metrics import roc curve, roc auc score
         log reg pred prob = log reg.predict proba(X train)[:,1] # probabilities for R
         log fpr, log tpr, log threshold = roc curve(y train, log reg pred prob)
         def graph roc curve single(log fpr, log tpr):
             plt.figure(figsize=(8,6))
             plt.title('ROC Curve - Logistic Regression', fontsize=18)
             plt.plot(log fpr, log tpr, label='Logistic Regression (AUC = {:.4f})'.form
                 roc auc score(y train, log reg pred prob)))
             plt.plot([0, 1], [0, 1], 'k--')
             plt.xlabel('False Positive Rate', fontsize=14)
             plt.ylabel('True Positive Rate', fontsize=14)
             plt.legend(loc="lower right")
             plt.show()
         graph_roc_curve_single(log_fpr, log_tpr)
```



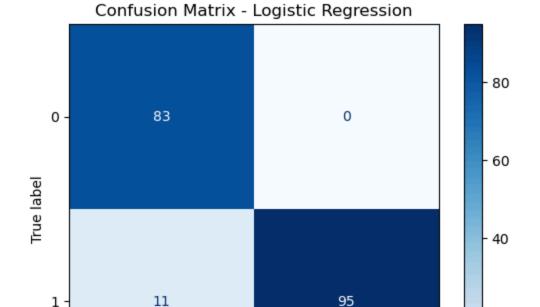
```
In [32]: # Train the Logistic Regression model
log_reg = LogisticRegression()
log_reg.fit(X_train, y_train)

# Predict probabilities and classes
y_pred_proba = log_reg.predict_proba(X_test)[:, 1] # probabilities for class
y_pred = log_reg.predict(X_test) # predicted classes

In [33]: from sklearn.metrics import confusion_matrix, ConfusionMatrixDisplay
import matplotlib.pyplot as plt

y_pred_classes = (y_pred > 0.5).astype(int) # if probabilities
cm = confusion_matrix(y_test, y_pred_classes)

disp = ConfusionMatrixDisplay(confusion_matrix=cm, display_labels=[0, 1])
disp.plot(cmap=plt.cm.Blues, values_format='d')
plt.title("Confusion Matrix - Logistic Regression")
plt.show()
```



Predicted label

1 -

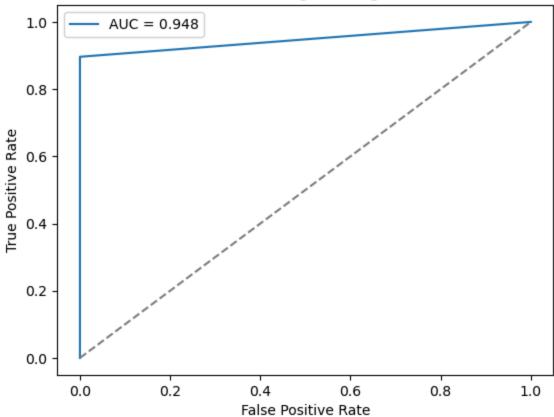
0

```
In [34]:
         from sklearn.metrics import roc_curve, roc_auc_score
         fpr, tpr, thresholds = roc_curve(y_test, y_pred)
         auc_score = roc_auc_score(y_test, y_pred)
         plt.plot(fpr, tpr, label=f"AUC = {auc_score:.3f}")
         plt.plot([0,1],[0,1],'--', color='gray')
         plt.xlabel("False Positive Rate")
         plt.ylabel("True Positive Rate")
         plt.title("ROC Curve - Logistic Regression")
         plt.legend()
         plt.show()
```

1

- 20

#### **ROC Curve - Logistic Regression**



# Conclusion:

Dataset is highly imbalanced → applied SMOTE/undersampling.

Logistic Regression was trained and evaluated.

Confusion matrix + ROC-AUC showed [insert recall/precision tradeoff].

For fraud detection, recall is more important (catch frauds).