Homework 3

Part 1: Imbalanced Dataset

In this homework, you will be working with an imbalanced Dataset. The dataset is Credit Card Fraud Detection dataset which was hosted on Kaggle. The aim is to detect fraudlent transactions.

Instructions

Please push the .ipynb, .py, and .pdf to Github Classroom prior to the deadline. Please include your UNI as well.

Setup

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

In [2]:  # Feel free to import any other packages you'd like to
```

Data Preprocessing and Exploration

Download the Kaggle Credit Card Fraud data set. Features V1, V2, ... V28 are the principal components obtained with PCA, the only features which have not been transformed with PCA are 'Time' and 'Amount'. Feature 'Time' contains the seconds elapsed between each transaction and the first transaction in the dataset. The feature 'Amount' is the transaction Amount, this feature can be used for example-dependant cost-sensitive learning. Feature 'Class' is the response variable and it takes value 1 in case of fraud and 0 otherwise.

```
In [3]:
         raw df = pd.read csv('https://storage.googleapis.com/download.tensorflow.org/dat
         raw df.head()
           Time
                      V1
                               V2
                                        ٧3
                                                 V4
                                                                   V6
                                                                             ۷7
                                                                                      V8
                                                          V5
Out[3]:
        0
                -1.359807 -0.072781 2.536347
                                            1.378155 -0.338321
                                                              0.462388
            0.0
                                                                        0.239599
                                                                                 0.098698
        1
            0.0
                 1.191857
                          0.266151 0.166480
                                            0.448154
                                                    0.060018 -0.082361 -0.078803
                                                                                 0.085102
        2
               -1.358354 -1.340163 1.773209
                                            0.379780 -0.503198 1.800499
                                                                        0.791461 0.247676
        3
            1.0 -0.966272 -0.185226 1.792993 -0.863291 -0.010309
                                                              1.247203
                                                                        0.237609 0.377436
                4
            2.0
                                                              0.095921
                                                                        0.592941 -0.270533
```

5 rows × 31 columns

1.1 Examining the class Imbalance

1.1.1 How many observations are in this dataset? How many are positive and negative?

(Note: Positive labels are labeled as 1)

Ans: there are 284807 observations, while 492 are positive, and 284315 are negative

1.2 Cleaning and normalizing the data

The raw data has a few issues. We are not sure what the time column actually means so drop the Time column. The Amount column also has a wide range of values covered so we take the log of the Amount column to reduce its range.

```
In [6]:
    cleaned_df = raw_df.copy()

# You don't want the `Time` column.
    cleaned_df.pop('Time')

# The `Amount` column covers a huge range. Convert to log-space.
    eps = 0.001 # 0 => 0.1¢
    cleaned_df['Log Ammount'] = np.log(cleaned_df.pop('Amount')+eps)
In [7]:
    cleaned_df
```

Out[7]:		V1	V2	V 3	V4	V 5	V6	V7	V
	0	-1.359807	-0.072781	2.536347	1.378155	-0.338321	0.462388	0.239599	0.09869
	1	1.191857	0.266151	0.166480	0.448154	0.060018	-0.082361	-0.078803	0.08510
	2	-1.358354	-1.340163	1.773209	0.379780	-0.503198	1.800499	0.791461	0.24767
	3	-0.966272	-0.185226	1.792993	-0.863291	-0.010309	1.247203	0.237609	0.37743
	4	-1.158233	0.877737	1.548718	0.403034	-0.407193	0.095921	0.592941	-0.27053
	•••								
2	84802	-11.881118	10.071785	-9.834783	-2.066656	-5.364473	-2.606837	-4.918215	7.30533
2	84803	-0.732789	-0.055080	2.035030	-0.738589	0.868229	1.058415	0.024330	0.29486
2	84804	1.919565	-0.301254	-3.249640	-0.557828	2.630515	3.031260	-0.296827	0.70841
2	84805	-0.240440	0.530483	0.702510	0.689799	-0.377961	0.623708	-0.686180	0.67914

```
284806 -0.533413 -0.189733 0.703337 -0.506271 -0.012546 -0.649617 1.577006 -0.41465
```

284807 rows × 30 columns

```
In [8]: cleaned_df["Class"].value_counts(normalize=True)
Out[8]: 0     0.998273
     1     0.001727
     Name: Class, dtype: float64
```

1.2.1 Split the dataset into development and test sets. Please set test size as 0.2 and random state as 42. Print the shape of your development and test features

```
In [9]: # Your Code Here
   X = cleaned_df.drop("Class", axis = 1)
   y = cleaned_df["Class"]

In [10]: from sklearn.model_selection import train_test_split
   X_dev, X_test, y_dev, y_test = train_test_split(X, y==1, stratify = y, test_size
   print(f"X_dev shape: {X_dev.shape}")
   print(f"X_test shape: {X_test.shape}")
   print(f"y_dev shape: {y_dev.shape}")
   print(f"y_test shape: {y_test.shape}")

   X_dev shape: (227845, 29)
   X_test shape: (56962, 29)
   y_dev shape: (227845,)
   y test shape: (56962,)
```

1.2.2 Normalize the features using Standard Scaler from Sklearn.

```
In [11]: # Your Code Here
    from sklearn.preprocessing import StandardScaler

    scaler1 = StandardScaler()
    X_test = scaler1.fit_transform(X_test)
    X_dev = scaler1.fit_transform(X_dev)
```

1.3 Defining Model and Performance Metrics

1.3.1 First, let us fit a default Decision tree classifier. (use max_depth=10 and random_state=42). Print the AUC and Average Precision values of 5 Fold Cross Validation

```
In [12]: # Your Code here
    from sklearn.tree import DecisionTreeClassifier
    from sklearn.model_selection import cross_validate
    tree1 = DecisionTreeClassifier(max_depth = 10, random_state = 42)
    score1 = cross_validate(tree1, X_dev, y_dev, cv = 5, scoring = ['roc_auc', 'aver

In [13]: print(score1["test_roc_auc"])
```

```
print(score1["test_average_precision"])

[0.88756328 0.88400873 0.81260522 0.77454358 0.82104149]
[0.62653551 0.71014615 0.60399496 0.58692296 0.68112078]

In [14]:
    print(f'mean AUC: {score1["test_roc_auc"].mean()}')
    print(f'mean AP: {score1["test_average_precision"].mean()}')

mean AUC: 0.8359524571279693
mean AP: 0.641744070059615
```

1.3.2 Perform random oversampling on the development dataset.

- What many positive and negative labels do you observe after random oversampling?
- What is the shape of your development dataset?

(Note: Set random state as 42 when performing oversampling)

```
In [15]: !pip install imblearn
```

Requirement already satisfied: imblearn in /Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages (0.0)

Requirement already satisfied: imbalanced-learn in /Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages (from imblearn) (0.9.1)

Requirement already satisfied: numpy>=1.17.3 in /Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages (from imbalanced-learn->imblearn) (1.21.4)

Requirement already satisfied: scipy>=1.3.2 in /Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages (from imbalanced-learn->imblearn) (1.7.3)

Requirement already satisfied: threadpoolctl>=2.0.0 in /Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages (from imbalanced-learn->imblearn) (3.0.0)

Requirement already satisfied: scikit-learn>=1.1.0 in /Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages (from imbalanced-learn->imblearn) (1.1.3)

Requirement already satisfied: joblib>=1.0.0 in /Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages (from imbalanced-learn->imblearn) (1.1.0)

```
In [16]: # Your Code here
    from imblearn.over_sampling import RandomOverSampler

    ros = RandomOverSampler(random_state = 42)
    X_dev_oversample1, y_dev_oversample1 = ros.fit_resample(X_dev, y_dev)
    print(y_dev_oversample1.value_counts())
    print(X_dev_oversample1.shape)
False 227451
```

```
True 227451
Name: Class, dtype: int64
(454902, 29)
```

Ans: Both positive and negative labels have 227451 samples, and the shape is about (454902, 29)

1.3.3 Repeat 1.3.1 using the dataset you created in the above step(1.3.2 Random oversampling). (Make sure you use the same hyperparameters as 1.3.1. i.e., max_depth=10 and random_state=42. This will help us to compare the models)

```
In [17]: tree2 = DecisionTreeClassifier(max_depth = 10, random_state = 42)
```

1.3.4 Perform Random undersampling on the development dataset.

- What many positive and negative labels do you observe after random undersampling?
- What is the shape of your development dataset?

(Note: Set random state as 42 when performing undersampling)

Ans: Both positive and negative labels have 394 samples, and the shape is about (788, 29)

1.3.5 Repeat 1.3.1 using the dataset you created in the above step(1.3.4 Random undersampling). (Make sure you use the same hyperparameters as 1.3.1. i,e., max_depth=10 and random_state=42. This will help us to compare the models)

```
In [25]: print(f'mean AUC: {score33["test_roc_auc"].mean()}')
    print(f'mean AP: {score33["test_average_precision"].mean()}')
mean AUC: 0.9303029182535673
```

mean AUC: 0.9303029182535673 mean AP: 0.9022723811037444

1.3.6 Perform Synthetic Minority Oversampling Technique (SMOTE) on the development dataset

- What many positive and negative labels do you observe after performing SMOTE?
- What is the shape of your development dataset? (Note: Set random state as 42 when performing SMOTE)

Ans: Both positive and negative labels have 227451 samples, and the shape is about (454902, 29)

1.3.7 Repeat 1.3.1 using the dataset you created in the above step(1.3.6 SMOTE). (Make sure you use the same hyperparameters as 1.3.1. i.e., max_depth=10 and random_state=42. This will help us to compare the models)

1.3.8 Make predictions on the test set using the four models that you built and report their AUC values.

```
In [31]: # Your Code here
from sklearn.metrics import roc_auc_score
```

```
tree1.fit(X_dev, y_dev)
score_normal = tree1.predict_proba(X_test)[:, 1]
print(f"The normal model has an AUC of {roc_auc_score(y_test, score_normal)}")

tree2.fit(X_dev_oversample1, y_dev_oversample1)
score_over = tree2.predict_proba(X_test)[:, 1]
print(f"The oversampling model has an AUC of {roc_auc_score(y_test, score_over)}

tree3.fit(X_dev_subsample1, y_dev_subsample1)
score_under = tree3.predict_proba(X_test)[:, 1]
print(f"The undersampling model has an AUC of {roc_auc_score(y_test, score_under)}

tree4.fit(X_dev_smote1, y_dev_smote1)
score_smote = tree4.predict_proba(X_test)[:, 1]
print(f"The smote model has an AUC of {roc_auc_score(y_test, score_smote)}")

The normal model has an AUC of 0.819676270198569
The oversampling model has an AUC of 0.9017043350120015
The undersampling model has an AUC of 0.896389918516647
```

In []:

1.3.9 Plot Confusion Matrices for all the four models on the test set. Comment your results

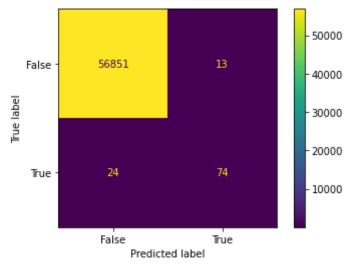
```
# Your Code here
from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(tree1, X_test, y_test)
```

/Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages/sklearn/utils/dep recation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. U se one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from estimator.

warnings.warn(msg, category=FutureWarning)

The smote model has an AUC of 0.8753799792989789

Out[32]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x7fdb6032c250 >

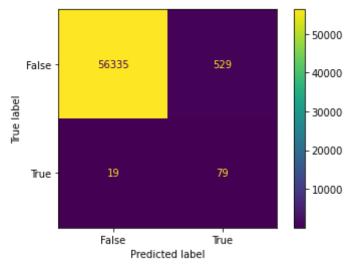


```
In [34]: plot_confusion_matrix(tree2, X_test, y_test)
```

/Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages/sklearn/utils/dep recation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. U se one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from_estimator.

warnings.warn(msg, category=FutureWarning)

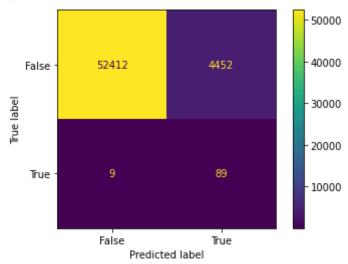
Out[34]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x7fdb8329ccd0 >



/Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages/sklearn/utils/dep recation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. U se one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from estimator.

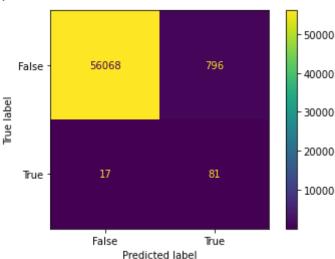
warnings.warn(msg, category=FutureWarning)

Out[35]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x7fdb8329cd30 >



```
In [36]: plot_confusion_matrix(tree4, X_test, y_test)
```

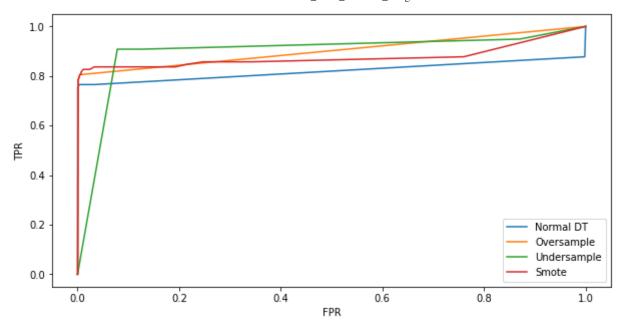
/Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages/sklearn/utils/dep recation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. U se one of the class methods: ConfusionMatrixDisplay.from_predictions or Confusion



Comment: the normal decision tree classifier has the highest accuracy, while the undersampling tree has the lowest accuracy. The undersampling tree has the best recall, while the normal decision tree has lowest recall. The normal decision tree has the highest precision, while the undersampling has the lowest precision.

1.3.10 Plot ROC for all the four models on the test set in a single plot. Make sure you label axes and legend properly. Comment your results

```
In [37]:
          # Your code
          from sklearn.metrics import roc curve
          normal fpr, normal tpr, normal thr = roc curve(y test, score normal, pos label =
          over fpr, over tpr, over thr = roc curve(y test, score over, pos label = 1)
          under_fpr, under_tpr, under_thr = roc_curve(y_test, score_under, pos_label = 1)
          smote_fpr, smote_tpr, smote_thr = roc_curve(y_test, score_smote, pos_label = 1)
          plt.figure(figsize = (10,5))
          plt.plot(normal fpr, normal tpr, label = "Normal DT")
          plt.plot(over fpr, over tpr, label = "Oversample")
          plt.plot(under fpr, under tpr, label = "Undersample")
          plt.plot(smote fpr, smote tpr, label = "Smote")
          plt.legend()
          plt.xlabel("FPR")
          plt.ylabel("TPR")
          plt.show()
```



ANS: The normal decision tree has the worst AUC, and undersampling has the best AUC, which is relatively higher than oversampling and smote. While smote and oversampling have similar AUC

1.3.11 Train a balanced default Decision tree classifier, using max_depth = 10 and random_state = 42 (balance the class weights). Print the AUC and average precision on dev set

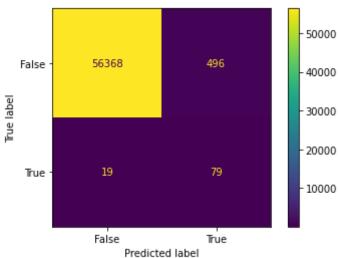
1.3.12 Plot confusion matrix on test set using the above model and comment on your results

```
# Your code here
tree5.fit(X_dev, y_dev)
plot_confusion_matrix(tree5, X_test, y_test)
```

/Users/clarencestudy/opt/anaconda3/lib/python3.8/site-packages/sklearn/utils/dep recation.py:87: FutureWarning: Function plot_confusion_matrix is deprecated; Function `plot_confusion_matrix` is deprecated in 1.0 and will be removed in 1.2. U se one of the class methods: ConfusionMatrixDisplay.from_predictions or ConfusionMatrixDisplay.from estimator.

warnings.warn(msg, category=FutureWarning)

Out[40]: <sklearn.metrics._plot.confusion_matrix.ConfusionMatrixDisplay at 0x7fdb500d75b0 >



ANS:It has an accuracy similar to oversampling decision tree, which is only slightly less than the normal decision tree. Its recall is also similar to the oversampling decision tree, which is relatively less than other models. Its precision is also similar to the oversampling decision tree, which is the only slightly less than the normal DT

In []:		