CREDIT CARD FRAUD ANALYSIS

PART 2: Handling Class Imbalance

Imports

```
In [1]: # Imported Libraries
        import numpy as np # linear algebra
        import pandas as pd # data processing, CSV file I/O (e.g. pd.read csv)
        #import tensorflow as tf
        import matplotlib.pyplot as plt
        import seaborn as sns
        from sklearn.manifold import TSNE
        from sklearn.decomposition import PCA, TruncatedSVD
        import matplotlib.patches as mpatches
        import time
        # Classifier Libraries
        from sklearn.linear model import LogisticRegression
        from sklearn.svm import SVC
        from sklearn.neighbors import KNeighborsClassifier
        from sklearn.tree import DecisionTreeClassifier
        from sklearn.ensemble import RandomForestClassifier
        import collections
        # Other Libraries
        import imblearn
        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 score, accuracy score, classification report
        from collections import Counter
        from sklearn.model selection import KFold, StratifiedKFold
        import warnings
        warnings.filterwarnings("ignore")
```

Data Reads

```
In [2]: df = pd.read_csv('creditcard_data.csv')
```

PART 2: Handling Class Imbalance

Check for class imbalance

```
In [3]: # Check for class Imbalance

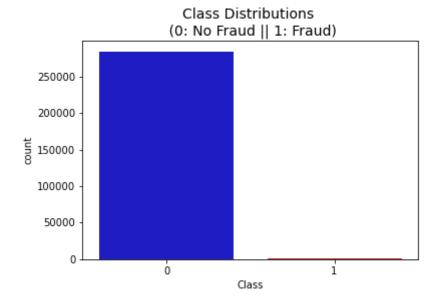
# The classes are heavily skewed we need to solve this issue later.
print('No Frauds', round(df['Class'].value_counts()[0]/len(df) * 100,2),
    '% of the dataset')
print('Frauds', round(df['Class'].value_counts()[1]/len(df) * 100,2), '%
of the dataset')

No Frauds 99.83 % of the dataset
Frauds 0.17 % of the dataset
In [4]: # Check Visual for the class distribution
```

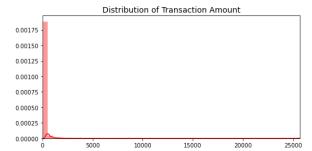
```
In [4]: # Check Visual for the class distribution

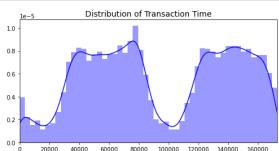
colors = ["#0101DF", "#DF0101"]
    sns.countplot('Class', data=df, palette=colors)
    plt.title('Class Distributions \n (0: No Fraud || 1: Fraud)', fontsize=1
    4)
```





Check distribution for Amount and Time field





Observation: Most of the Transactions are smaller amount: The ferquency of Transactions have a clear peak pattern depending upon the time of the day

Scale Variables Time and Amount

```
In [6]: # Scale using Robust scaler as it is less prone to outliers

# Scale the columns that are left to scale (Amount and Time)
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.resha
pe(-1,1))
df['scaled_time'] = rob_scaler.fit_transform(df['Time'].values.reshape(-
1,1))

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

Out[6]:

	V1	V 2	V 3	V 4	V 5	V 6	V 7	V 8	V!
0	-1.359807	-0.072781	2.536347	1.378155	-0.338321	0.462388	0.239599	0.098698	0.36378
1	1.191857	0.266151	0.166480	0.448154	0.060018	-0.082361	-0.078803	0.085102	-0.25542
2	-1.358354	-1.340163	1.773209	0.379780	-0.503198	1.800499	0.791461	0.247676	-1.51465
3	-0.966272	-0.185226	1.792993	-0.863291	-0.010309	1.247203	0.237609	0.377436	-1.38702
4	-1.158233	0.877737	1.548718	0.403034	-0.407193	0.095921	0.592941	-0.270533	0.817739

5 rows × 31 columns

Split original data set for testing

```
In [7]: from sklearn.model selection import train test split
        from sklearn.model selection import StratifiedShuffleSplit
        # Create X an y Data
        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 i
            original_ytrain, original_ytest = y.iloc[train_index], y.iloc[test_i
        ndex]
        # We already have X train and y train for undersample data thats why I a
        m using original to distinguish and to not overwrite these variables.
        # original Xtrain, original Xtest, original ytrain, original ytest = tra
        in test split(X, y, test size=0.2, random state=42)
        # 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 distri
        buted
        train unique label, train counts label = np.unique(original ytrain, retu
        rn counts=True)
        test unique label, test counts label = np.unique(original ytest, return
        counts=True)
        print('-' * 100)
        print('Label Distributions: \n')
        print(train counts label/ len(original ytrain))
        print(test counts label/ len(original ytest))
```

```
Train: [ 30473 30496 31002 ... 284803 284804 284805] Test: [
      2 ... 57017 57018 570191
                           2 ... 284803 284804 284805] Test: [ 30473
                    1
0496 31002 ... 113964 113965 113966]
                           2 ... 284803 284804 284805] Test: [ 81186
                    1
1609 82400 ... 170946 170947 170948]
                           2 ... 284803 284804 284805] Test: [150647 15
Train: [
                    1
0654 150660 ... 227866 227867 227868]
             0
                    1
                           2 ... 227866 227867 227868] Test: [208651 21
2516 212644 ... 284803 284804 284805]
Label Distributions:
[0.99827514 0.00172486]
[0.99826197 0.00173803]
```

CLASS IMBALANCE HANDLING APPROACHES

1. Random Undersampling

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 [8]: # Create Undersample and Equally represented dataset
# Since our classes are highly skewed we should make them equivalent in
    order to have a normal distribution of the classes.

# 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]

# combine the fraud and non fraud rows
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()
```

Out[8]:

	V1	V2	V 3	V 4	V 5	V 6	V 7	1
260781	-1.164409	-1.279381	-1.304220	-1.200687	-0.779834	0.229462	0.985915	0.0476
154720	-5.552122	5.678134	-9.775528	8.416295	-4.409844	-1.506235	-6.899839	3.7504
108353	0.826988	-0.317576	0.261861	1.125783	-0.209647	0.191533	0.100275	0.1447
151006	-26.457745	16.497472	-30.177317	8.904157	-17.892600	-1.227904	-31.197329	-11.4389
163586	0.949241	1.333519	-4.855402	1.835006	-1.053245	-2.562826	-2.286986	0.2609

5 rows × 31 columns

Undersampling using imblearn

TBD

Check for class distribution after Under Sampling

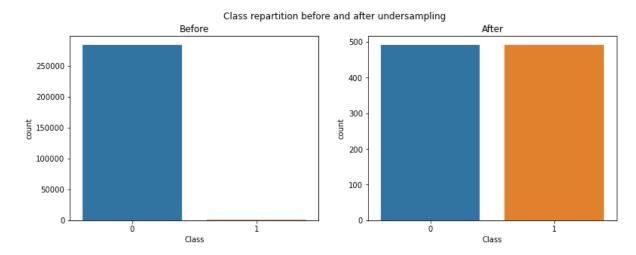
```
In [9]: # Check class distribution after undersampling
    print('Distribution of the Classes in the subsample dataset')
    print(new_df['Class'].value_counts()/len(new_df))

#visualizing undersampling results
    fig, axs = plt.subplots(ncols=2, figsize=(13,4.5))
    sns.countplot(x="Class", data=df, ax=axs[0])
    sns.countplot(x="Class", data=new_df, ax=axs[1])

fig.suptitle("Class repartition before and after undersampling")
    al=fig.axes[0]
    al.set_title("Before")
    a2=fig.axes[1]
    a2.set_title("After")
```

```
Distribution of the Classes in the subsample dataset 1 0.5 0 0.5 Name: Class, dtype: float64
```

Out[9]: Text(0.5, 1.0, 'After')



Check correlation matrix with balanced data set

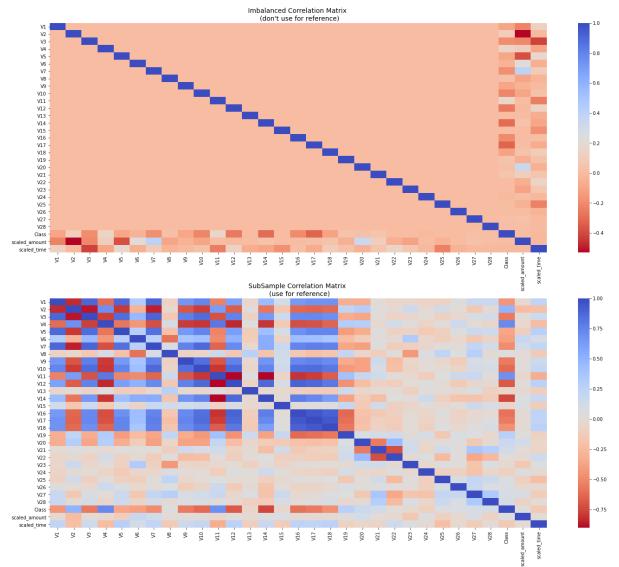
https://towardsdatascience.com/having-an-imbalanced-dataset-here-is-how-you-can-solve-it-1640568947eb (https://towardsdatascience.com/having-an-imbalanced-dataset-here-is-how-you-can-solve-it-1640568947eb)

```
In [10]: # 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)", fontsize=14)

#. Balanced Sub Set
sub_sample_corr = new_df.corr()
sns.heatmap(sub_sample_corr, cmap='coolwarm_r', annot_kws={'size':20}, a
x=ax2)
ax2.set_title('SubSample Correlation Matrix \n (use for reference)', fon
tsize=14)
plt.show()
```



Observations from the Reference Heatmap

Positive correlation with Y Variable V2; V4; V11; V19

Negative correlation with Y Variable V17, V16, V14, V12 and V10

Box plots to Study Distributions by class for Highly correlated variables

```
In [11]: ## Positively correlated variables
    f, axes = plt.subplots(ncols=4, figsize=(20,4))

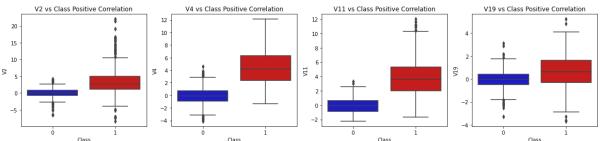
# Positive correlations (The higher the feature the probability increase
    s that it will be a fraud transaction)
    sns.boxplot(x="Class", y="V2", data=new_df, palette=colors, ax=axes[0])
    axes[0].set_title('V2 vs Class Positive Correlation')

sns.boxplot(x="Class", y="V4", data=new_df, palette=colors, ax=axes[1])
    axes[1].set_title('V4 vs Class Positive Correlation')

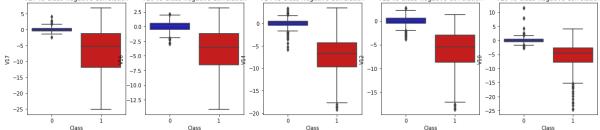
sns.boxplot(x="Class", y="V11", data=new_df, palette=colors, ax=axes[2])
    axes[2].set_title('V11 vs Class Positive Correlation')

sns.boxplot(x="Class", y="V19", data=new_df, palette=colors, ax=axes[3])
    axes[3].set_title('V19 vs Class Positive Correlation')

plt.show()
```



```
In [12]:
         ## Negatively correlated variables
          # Negative Correlations with our Class (The lower our feature value the
           more likely it will be a fraud transaction)
          f, axes = plt.subplots(ncols=5, figsize=(20,4))
          sns.boxplot(x="Class", y="V17", data=new df, palette=colors, ax=axes[0])
          axes[0].set_title('V17 vs Class Negative Correlation')
          sns.boxplot(x="Class", y="V16", data=new df, palette=colors, ax=axes[1])
          axes[1].set_title('V16 vs Class Negative Correlation')
          sns.boxplot(x="Class", y="V14", data=new_df, palette=colors, ax=axes[2])
          axes[2].set_title('V14 vs Class Negative Correlation')
          sns.boxplot(x="Class", y="V12", data=new df, palette=colors, ax=axes[3])
          axes[3].set_title('V12 vs Class Negative Correlation')
          sns.boxplot(x="Class", y="V10", data=new_df, palette=colors, ax=axes[4])
          axes[4].set_title('V10 vs Class Negative Correlation')
          plt.show()
             V17 vs Class Negative Correlation
                            V16 vs Class Negative Correlation
                                            V14 vs Class Negative Correlation
                                                           V12 vs Class Negative Correlation
```



Observation: For the variable studied the low values clearly result in fraud

Visualize and remove extreme outliers for high correlated variables

Steps:

- **Visualize Distributions:** We first start by visualizing the distribution of the feature we are going to use to eliminate some of the outliers.
- **Determining the threshold:** After we decide which number we will use to multiply with the iqr (the lower more outliers removed), we will proceed in determining the upper and lower thresholds by substrating q25 threshold (lower extreme threshold) and adding q75 + threshold (upper extreme threshold).
- Conditional Dropping: Lastly, we create a conditional dropping stating that if the "threshold" is exceeded in both extremes, the instances will be removed.
- **Boxplot Representation:** Visualize through the boxplot that the number of "extreme outliers" have been reduced to a considerable amount.

Visualize the Top Negative correlated Variables with Y

V10, V12, V14

```
In [13]: from scipy.stats import norm

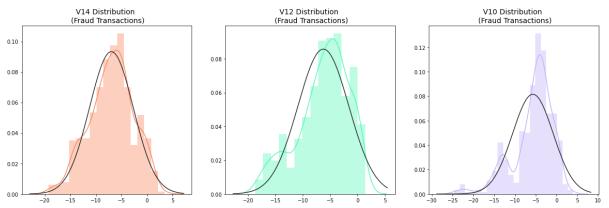
f, (ax1, ax2, ax3) = plt.subplots(1,3, figsize=(20, 6))

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 [14]:  #### Observations :
```

Visualize the Top positive correlated variables with Y

V2,V4,V11

```
In [15]: from scipy.stats import norm
           f, (ax1, ax2, ax3) = plt.subplots(1,3, figsize=(20, 6))
          v14 fraud dist = new_df['V2'].loc[new_df['Class'] == 1].values
           sns.distplot(v14_fraud_dist,ax=ax1, fit=norm, color='#FB8861')
           ax1.set title('V2 Distribution \n (Fraud Transactions)', fontsize=14)
          v12 fraud dist = new_df['V4'].loc[new_df['Class'] == 1].values
           sns.distplot(v12_fraud_dist,ax=ax2, fit=norm, color='#56F9BB')
           ax2.set title('V4 Distribution \n (Fraud Transactions)', fontsize=14)
          v10 fraud dist = new df['V11'].loc[new df['Class'] == 1].values
           sns.distplot(v10_fraud_dist,ax=ax3, fit=norm, color='#C5B3F9')
           ax3.set_title('V11 Distribution \n (Fraud Transactions)', fontsize=14)
          plt.show()
                     V2 Distribution
                                                                             V11 Distribution
                                                 V4 Distribution
                    (Fraud Transactions)
                                                (Fraud Transactions)
                                                                            (Fraud Transactions)
                                       0.14
                                                                   0.175
                                       0.12
                                                                   0.150
           0.150
                                       0.10
                                                                   0.125
           0.125
                                       0.08
                                                                   0.100
           0.100
                                       0.06
           0.075
                                                                   0.075
```

0.050

7.5 10.0

Remove Extreme Outliers From Negative correlated variables

0.04

0.02

Remove outliers from V14

0.050

```
In [16]:
          # ----> V14 Removing Outliers (Highest Negative Correlated with Label
         s)
         # Isolate the fraud rows
         v14_fraud = new_df['V14'].loc[new_df['Class'] == 1].values
         # Get the lower quartile and upper quartile
         q25, q75 = np.percentile(v14_fraud, 25), np.percentile(v14_fraud, 75)
         print('Quartile 25: {} | Quartile 75: {}'.format(q25, q75))
         # Get the interquartile range
         v14 iqr = q75 - q25
         print('iqr: {}'.format(v14_iqr))
         # define cut offs
         v14 cut off = v14 iqr * 1.5
         # define lower and upper cut offs
         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))
         # Get the outliers for Fraud Rows
         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('V14 outliers:{}'.format(outliers))
         # Drop outliers from Fraud rows
         print("row count before outlier drop : {}".format(new_df.shape[0]))
         new_df = new_df.drop(new_df[((new_df['V14'] > v14_upper) | (new_df['V14'])
         ] < v14 lower))&(new df['Class'] ==1)].index)
         print("row count after outlier drop : {}".format(new df.shape[0]))
         print('---' * 44)
         Quartile 25: -9.69272296475 | Quartile 75: -4.2828208495
         igr: 5.40990211525
         Cut Off: 8.114853172875002
         V14 Lower: -17.807576137625002
         V14 Upper: 3.8320323233750013
         Feature V14 Outliers for Fraud Cases: 4
         V14 outliers:[-18.49377336, -18.04999769, -18.82208674, -19.21432549]
         row count before outlier drop: 984
         row count after outlier drop: 980
```

Remove outliers from V12

```
In [17]: # ----> V12 Removing Outliers (Highest Negative Correlated with Labels)
         # Isolate the fraud rows
         v12_fraud = new_df['V12'].loc[new_df['Class'] == 1].values
         # Get the lower quartile and upper quartile
         q25, q75 = np.percentile(v12 fraud, 25), np.percentile(v12 fraud, 75)
         print('Quartile 25: {} | Quartile 75: {}'.format(q25, q75))
         # Get the interquartile range
         v12_{iqr} = q75 - q25
         print('iqr: {}'.format(v12 iqr))
         # define cut offs
         v12 cut off = v12 iqr * 1.5
         # define lower and upper cut offs
         v12 lower, v12 upper = q25 - v12 cut off, q75 + v12 cut off
         print('Cut Off: {}'.format(v12_cut_off))
         print('V12 Lower: {}'.format(v12_lower))
         print('V12 Upper: {}'.format(v12 upper))
         # Get the outliers for Fraud Rows
         outliers = [x for x in v12_fraud if x < v12_lower or x > v12_upper]
         print('Feature V12 Outliers for Fraud Cases: {}'.format(len(outliers)))
         print('V12 outliers:{}'.format(outliers))
         # Drop outliers from Fraud rows
         print("row count before outlier drop : {}".format(new df.shape[0]))
         new df = new df.drop(new df[((new df['V12'] > v12 upper) | (new df['V12'
         ] < v12_lower))&(new_df['Class'] ==1 )].index)</pre>
         print("row count after outlier drop : {}".format(new df.shape[0]))
         print('----' * 44)
         Quartile 25: -8.6730332045 | Quartile 75: -2.8930305682500004
         igr: 5.780002636249999
         Cut Off: 8.670003954374998
         V12 Lower: -17.343037158875
         V12 Upper: 5.776973386124998
         Feature V12 Outliers for Fraud Cases: 4
         V12 outliers:[-18.43113103, -18.55369701, -18.68371463, -18.04759657]
         row count before outlier drop: 980
         row count after outlier drop: 976
```

```
In [18]: #### Remove Outliers from V10
         # ----> V10 Removing Outliers (Highest Negative Correlated with Labels)
         # Isolate the fraud rows
         v10_fraud = new_df['V10'].loc[new_df['Class'] == 1].values
         # Get the lower quartile and upper quartile
         q25, q75 = np.percentile(v10 fraud, 25), np.percentile(v10 fraud, 75)
         print('Quartile 25: {} | Quartile 75: {}'.format(q25, q75))
         # Get the interguartile range
         v10_{iqr} = q75 - q25
         print('iqr: {}'.format(v10_iqr))
         # define cut offs
         v10_cut_off = v10_iqr * 1.5
         # define lower and upper cut offs
         v10_lower, v10_upper = q25 - v10_cut_off, q75 + v10_cut_off
         print('Cut Off: {}'.format(v10 cut off))
         print('V10 Lower: {}'.format(v10 lower))
         print('V10 Upper: {}'.format(v10_upper))
         # Get the outliers for Fraud Rows
         outliers = [x for x in v10 fraud if x < v10 lower or x > v10 upper]
         print('Feature V10 Outliers for Fraud Cases: {}'.format(len(outliers)))
         print('V10 outliers:{}'.format(outliers))
         # Drop outliers from Fraud rows
         print("row count before outlier drop : {}".format(new_df.shape[0]))
         new df = new df.drop(new df[((new df['V10'] > v10 upper) | (new df['V10'
         ] < v10 lower))&(new df['Class'] ==1)].index)
         print("row count after outlier drop : {}".format(new df.shape[0]))
         print('----' * 44)
         Quartile 25: -7.466658536000001 | Quartile 75: -2.51186113825
         igr: 4.954797397750001
         Cut Off: 7.432196096625002
         V10 Lower: -14.898854632625003
         V10 Upper: 4.920334958375001
         Feature V10 Outliers for Fraud Cases: 27
         V10 outliers:[-22.18708856, -15.23996196, -16.60119697, -15.12416281, -
         19.83614885, -24.58826244, -15.34609885, -14.92465477, -16.64962816, -1
         5.56379134, -16.30353766, -22.18708856, -22.18708856, -15.23183337, -1
         6.74604411, -20.94919155, -24.40318497, -16.25561175, -23.22825484, -1
         8.91324333, -15.23996196, -22.18708856, -17.14151364, -15.56379134, -1
         4.92465477, -15.12375218, -18.27116817]
         row count before outlier drop: 976
         row count after outlier drop: 949
```

file:///Users/anishroychowdhury/Desktop/Credit_Card_Fraud_Class_Imbalance_2.html

Removing Extreme Outliers from Highly Positive correlated variables

Remove Outliers from V2

```
In [19]: #### Remove Outliers from V2
         # ----> V2 Removing Outliers (Highest Negative Correlated with Labels)
         # Isolate the fraud rows
         v2 fraud = new df['V2'].loc[new df['Class'] == 1].values
         # Get the lower quartile and upper quartile
         q25, q75 = np.percentile(v2 fraud, 25), np.percentile(v2 fraud, 75)
         print('Quartile 25: {} | Quartile 75: {}'.format(q25, q75))
         # Get the interquartile range
         v2 iqr = q75 - q25
         print('iqr: {}'.format(v2_iqr))
         # define cut offs
         v2_cut_off = v2_iqr * 1.5
         # define lower and upper cut offs
         v2 lower, v2_upper = q25 - v2_cut_off, q75 + v2_cut_off
         print('Cut Off: {}'.format(v2 cut off))
         print('V2 Lower: {}'.format(v2_lower))
         print('V2 Upper: {}'.format(v2_upper))
         # Get the outliers for Fraud Rows
         outliers = [x for x in v2 fraud if x < v2 lower or x > v2 upper]
         print('Feature V2 Outliers for Fraud Cases: {}'.format(len(outliers)))
         print('V2 outliers:{}'.format(outliers))
         # Drop outliers from Fraud rows
         print("row count before outlier drop : {}".format(new_df.shape[0]))
         new df = new df.drop(new df['(new df['V2'] > v2 upper) | (new df['V2'] <</pre>
         v2 lower))&(new df['Class'] ==1 )].index)
         print("row count after outlier drop : {}".format(new df.shape[0]))
         print('---' * 44)
```

```
Ouartile 25: 1.133138588 | Ouartile 75: 4.141986232
igr: 3.008847644
Cut Off: 4.513271466
V2 Lower: -3.380132878
V2 Upper: 8.655257698
Feature V2 Outliers for Fraud Cases: 46
V2 outliers: [12.78597064, 13.76594216, 14.60199804, 10.5417508, -3.9359
189239999997, 16.15570143, 10.39391714, 12.65219683, 12.37398914, 12.78
597064, 15.59819266, 12.78597064, 12.78597064, 9.067613427000001, -4.81
446074, 12.78597064, -7.159041717000001, -6.976420008, 12.78597064, 11.
81792199, 11.58638052, 12.09589323, 14.32325381, -3.4204679839999996, 1
2.93050512, -7.1969796310000005, -8.402153678, 14.04456678, 8.775997152
999999, 10.11481572, -7.449015159, 10.81966537, -5.198360199, 13.208904
28, 16.43452455, 16.71338924, 8.713250171, 9.669900173, -3.488130181000
0003, 13.48738579, 15.87692299, 10.55860019, 9.223691949, 15.36580438,
-3.9523200860000003, -3.930731396]
row count before outlier drop: 949
row count after outlier drop: 903
```

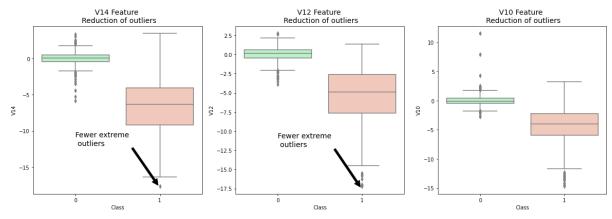
Remove Outliers for V11

```
In [20]: # ----> V11 Removing Outliers (Highest Negative Correlated with Labels)
         # Isolate the fraud rows
         v11_fraud = new_df['V11'].loc[new_df['Class'] == 1].values
         # Get the lower quartile and upper quartile
         q25, q75 = np.percentile(v11 fraud, 25), np.percentile(v11 fraud, 75)
         print('Quartile 25: {} | Quartile 75: {}'.format(q25, q75))
         # Get the interquartile range
         v11_iqr = q75 - q25
         print('iqr: {}'.format(v11 iqr))
         # define cut offs
         v11 cut off = v11 iqr * 1.5
         # define lower and upper cut offs
         v11 lower, v11 upper = q25 - v11 cut off, q75 + v11 cut off
         print('Cut Off: {}'.format(v11 cut off))
         print('V11 Lower: {}'.format(v11_lower))
         print('V11 Upper: {}'.format(v11 upper))
         # Get the outliers for Fraud Rows
         outliers = [x for x in v11_fraud if x < v11_lower or x > v11_upper]
         print('Feature V11 Outliers for Fraud Cases: {}'.format(len(outliers)))
         print('V11 outliers:{}'.format(outliers))
         # Drop outliers from Fraud rows
         print("row count before outlier drop : {}".format(new df.shape[0]))
         new df = new df.drop(new df[((new df['V11'] > v11 upper) | (new df['V11'])
         ] < v11_lower))&(new_df['Class'] ==1 )].index)</pre>
         print("row count after outlier drop : {}".format(new df.shape[0]))
         print('----' * 44)
         Quartile 25: 1.8450858435 | Quartile 75: 4.775434035
         igr: 2.9303481915000003
         Cut Off: 4.39552228725
         V11 Lower: -2.5504364437500007
         V11 Upper: 9.17095632225
         Feature V11 Outliers for Fraud Cases: 11
         V11 outliers:[9.369079057999999, 11.27792073, 10.27776886, 10.18758732,
         9.939819742000001, 10.54526295, 11.1524906, 10.06378975, 9.328799257, 1
         0.85301165, 10.44684681]
         row count before outlier drop: 903
         row count after outlier drop: 892
```

Check outlier removal with Box Plots

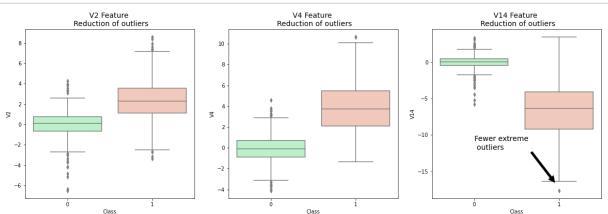
Box plots for highly correlated -ve features: V10, V12, V14

```
In [21]:
         f_{\star}(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 \setminus n outliers', xy=(0.98, -17.5), xytext=(0, -
          12),
                      arrowprops=dict(facecolor='black'),
                      fontsize=14)
          # Feature 12
          sns.boxplot(x="Class", y="V12", data=new df, ax=ax2, palette=colors)
          ax2.set_title("V12 Feature \n Reduction of outliers", fontsize=14)
          ax2.annotate('Fewer extreme n outliers', xy=(0.98, -17.3), xytext=(0, -
          12),
                      arrowprops=dict(facecolor='black'),
                      fontsize=14)
          # Feature V10
          sns.boxplot(x="Class", y="V10", data=new df, ax=ax3, palette=colors)
          ax3.set title("V10 Feature \n Reduction of outliers", fontsize=14)
          ax3.annotate('Fewer extreme \setminus n outliers', xy=(0.95, -16.5), xytext=(0, -
          12),
                      arrowprops=dict(facecolor='black'),
                      fontsize=14)
          plt.show()
```



Box plots for highly correlated +ve features: V2. V4. V11

```
In [22]: f_1(ax1, ax2, ax3) = plt.subplots(1, 3, figsize=(20,6))
         colors = ['#B3F9C5', '#f9c5b3']
         # Boxplots with outliers removed
         # Feature V2
         sns.boxplot(x="Class", y="V2", data=new df,ax=ax1, palette=colors)
         ax1.set_title("V2 Feature \n Reduction of outliers", fontsize=14)
         ax1.annotate('Fewer extreme \setminus n outliers', xy=(0.98, -17.5), xytext=(0, -
         12),
                      arrowprops=dict(facecolor='black'),
                      fontsize=14)
         # Feature 12
         sns.boxplot(x="Class", y="V4", data=new df, ax=ax2, palette=colors)
         ax2.set_title("V4 Feature \n Reduction of outliers", fontsize=14)
         ax2.annotate('Fewer extreme \n outliers', xy=(0.98, -17.3), xytext=(0, -
         12),
                      arrowprops=dict(facecolor='black'),
                      fontsize=14)
         # Feature V10
         sns.boxplot(x="Class", y="V14", data=new_df, ax=ax3, palette=colors)
         ax3.set_title("V14 Feature \n Reduction of outliers", fontsize=14)
         ax3.annotate('Fewer extreme \n outliers', xy=(0.95, -16.5), xytext=(0, -
         12),
                      arrowprops=dict(facecolor='black'),
                      fontsize=14)
         plt.show()
```



Dimensionality Reduction and Clustering Visualization using t-SNE

Summary:

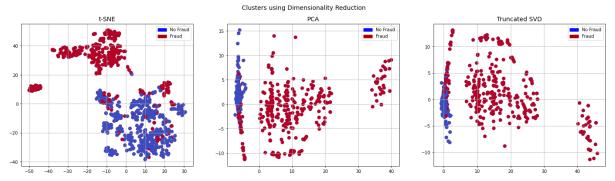
- t-SNE algorithm can pretty accurately cluster the cases that were fraud and non-fraud in our dataset.
- Although the subsample is pretty small, the t-SNE algorithm is able to detect clusters pretty accurately in every scenario (I shuffle the dataset before running t-SNE)
- This gives us an indication that further predictive models will perform pretty well in separating fraud cases from non-fraud cases.

```
In [23]: # New df is from the random undersample data (fewer instances)
         X = new df.drop('Class', axis=1)
         y = new df['Class']
         # T-SNE Implementation
         t0 = time.time()
         X_reduced_tsne = TSNE(n_components=2, random_state=42).fit_transform(X.v
         alues)
         t1 = time.time()
         print("T-SNE took {:.2} s".format(t1 - t0))
         # PCA Implementation
         t0 = time.time()
         X reduced pca = PCA(n components=2, random state=42).fit transform(X.val
         ues)
         t1 = time.time()
         print("PCA took {:.2} s".format(t1 - t0))
         # TruncatedSVD
         t0 = time.time()
         X reduced svd = TruncatedSVD(n components=2, algorithm='randomized', ran
         dom state=42).fit transform(X.values)
         t1 = time.time()
         print("Truncated SVD took {:.2} s".format(t1 - t0))
         T-SNE took 2.2 s
         PCA took 0.018 s
```

Truncated SVD took 0.0034 s

Plot Scatter from the various reduced techniques

```
In [24]: f, (ax1, ax2, ax3) = plt.subplots(1, 3, figsize=(24,6))
         # labels = ['No Fraud', 'Fraud']
         f.suptitle('Clusters using Dimensionality Reduction', fontsize=14)
         blue_patch = mpatches.Patch(color='#0A0AFF', label='No Fraud')
         red_patch = mpatches.Patch(color='#AF0000', label='Fraud')
         # t-SNE scatter plot
         ax1.scatter(X_reduced_tsne[:,0], X_reduced_tsne[:,1], c=(y == 0), cmap=
         'coolwarm', label='No Fraud', linewidths=2)
         ax1.scatter(X_reduced_tsne[:,0], X_reduced_tsne[:,1], c=(y == 1), cmap=
         'coolwarm', label='Fraud', linewidths=2)
         ax1.set_title('t-SNE', fontsize=14)
         ax1.grid(True)
         ax1.legend(handles=[blue patch, red patch])
         # PCA scatter plot
         ax2.scatter(X reduced pca[:,0], X reduced pca[:,1], c=(y == 0), cmap='co
         olwarm', label='No Fraud', linewidths=2)
         ax2.scatter(X reduced pca[:,0], X reduced pca[:,1], c=(y == 1), cmap='co
         olwarm', label='Fraud', linewidths=2)
         ax2.set title('PCA', fontsize=14)
         ax2.grid(True)
         ax2.legend(handles=[blue patch, red patch])
         # TruncatedSVD scatter plot
         ax3.scatter(X reduced svd[:,0], X reduced svd[:,1], c=(y == 0), cmap='co
         olwarm', label='No Fraud', linewidths=2)
         ax3.scatter(X reduced svd[:,0], X reduced svd[:,1], c=(y == 1), cmap='co
         olwarm', label='Fraud', linewidths=2)
         ax3.set title('Truncated SVD', fontsize=14)
         ax3.grid(True)
         ax3.legend(handles=[blue patch, red patch])
         plt.show()
```



Observations: We find that t-SNE offers a good separation - Thus the classes are separable

Classification with - Under Sampling

```
In [25]: # Preparing data for modelling
         # Undersampling before cross validating (prone to overfit)
         X = new df.drop('Class', axis=1)
         y = new_df['Class']
         # 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,
         random_state=42)
In [26]: X_train1 = X_train.values
         X_train1.shape
Out[26]: (713, 30)
In [27]: # Let's implement simple classifiers
         classifiers = {
             "LogisticRegression": LogisticRegression(),
             "KNN": KNeighborsClassifier(),
             "SupportVectorClassifier": SVC(),
             "DecisionTreeClassifier": DecisionTreeClassifier()
         }
```

Cross Validation employed

```
In [28]: 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 trainin
         g score of", round(training score.mean(), 2) * 100, "% accuracy score")
         Classifiers: LogisticRegression Has a training score of 93.0 % accurac
         y score
         Classifiers:
                       KNeighborsClassifier Has a training score of 93.0 % accur
         acy score
         Classifiers:
                       SVC Has a training score of 93.0 % accuracy score
                       DecisionTreeClassifier Has a training score of 89.0 % acc
         Classifiers:
         uracy score
```

Observation

LR has the best training accuracy with CV

Grid Search Employed - Hyperparameter tuned individually for each classifiers

```
In [29]: # Use GridSearchCV to find the best parameters.
        from sklearn.model selection import GridSearchCV
        # Logistic Regression
        #-----
        # penalty is regularization L1 or L2
        # C parameter inverse of Reg strength
        log_reg_params = {"penalty": ['l1', 'l2'], 'C': [0.001, 0.01, 0.1, 1, 10
        , 100, 1000]}
        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
        #_____
        # K nearest neighbours
        #----
        # Number of nearest neighbours
        # Algo to find distance
        # auto
        # ball tree
        # kd tree
        # brute
        knears_params = {"n_neighbors": list(range(2,5,1)), 'algorithm': ['auto'
        , 'ball_tree', 'kd_tree', 'brute']}
        grid knears = GridSearchCV(KNeighborsClassifier(), knears params)
        grid_knears.fit(X_train, y_train)
        # KNears best estimator
        knears neighbors = grid knears.best estimator
        #-----
        # Support Vector Classifier
        #_____
        # Small value of C will cause the optimizer to look for alarger margin h
        ence more misclassification
        svc params = {'C': [0.5, 0.7, 0.9, 1], 'kernel': ['rbf', 'poly', 'sigmoi
        d', 'linear']}
        grid svc = GridSearchCV(SVC(), svc params)
        grid svc.fit(X train, y train)
        # SVC best estimator
        svc = grid svc.best estimator
        #_____
        # DecisionTree Classifier
        #-----
        tree params = {"criterion": ["gini", "entropy"], "max depth": list(range
        (2,4,1)),
                    "min samples leaf": list(range(5,7,1))}
        grid tree = GridSearchCV(DecisionTreeClassifier(), tree params)
        grid tree.fit(X train, y train)
        # tree best estimator
        tree clf = grid tree.best estimator
        #-----
```

Get the Cross Validated Score for each of the best models chosen

```
In [30]: # 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() * 100, 2).astype(str) + '%')

knears_score = cross_val_score(knears_neighbors, X_train, y_train, cv=5)
print('Knears Neighbors Cross Validation Score', round(knears_score.mean() * 100, 2).astype(str) + '%')

svc_score = cross_val_score(svc, X_train, y_train, cv=5)
print('Support Vector Classifier Cross Validation Score', round(svc_score.mean() * 100, 2).astype(str) + '%')

tree_score = cross_val_score(tree_clf, X_train, y_train, cv=5)
print('DecisionTree Classifier Cross Validation Score', round(tree_score.mean() * 100, 2).astype(str) + '%')
```

Logistic Regression Cross Validation Score: 93.41% Knears Neighbors Cross Validation Score 93.69% Support Vector Classifier Cross Validation Score 93.27% DecisionTree Classifier Cross Validation Score 92.56%

Get the ROC Score for All the models

```
In [32]: # Get the ROC Score for all the models

from sklearn.metrics import roc_auc_score

print('Logistic Regression: ', roc_auc_score(y_train, log_reg_pred))
print('KNears Neighbors: ', roc_auc_score(y_train, knears_pred))
print('Support Vector Classifier: ', roc_auc_score(y_train, svc_pred))
print('Decision Tree Classifier: ', roc_auc_score(y_train, tree_pred))

Logistic Regression: 0.9801644202275117
KNears Neighbors: 0.9331126724659847
Support Vector Classifier: 0.9799095051460982
```

Observation Logistics Regression shows the best ROC Score

Decision Tree Classifier: 0.9220756460504095

Plot ROC Curve

```
In [33]: # Get ROC curve parms for Logit
log_fpr, log_tpr, log_threshold = roc_curve(y_train, log_reg_pred)

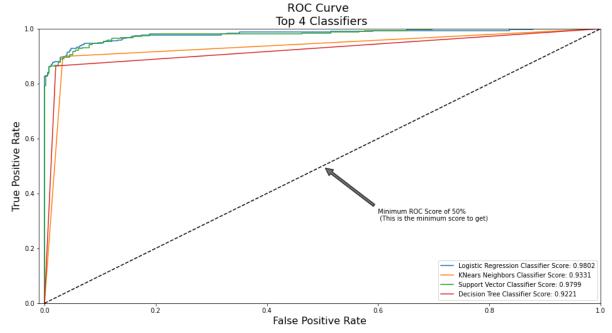
# Get ROC curve parms for Logit
knear_fpr, knear_tpr, knear_threshold = roc_curve(y_train, knears_pred)

# Get ROC curve parms for Logit
svc_fpr, svc_tpr, svc_threshold = roc_curve(y_train, svc_pred)

# Get ROC curve parms for Logit
tree_fpr, tree_tpr, tree_threshold = roc_curve(y_train, tree_pred)
```

```
In [ ]:
```

```
In [34]:
        # plot curve
         def graph roc curve multiple(log fpr, log tpr, knear fpr, knear tpr, svc
         _fpr, svc_tpr, tree_fpr, tree_tpr):
             plt.figure(figsize=(16,8))
             plt.title('ROC Curve \n Top 4 Classifiers', fontsize=18)
             plt.plot(log fpr, log tpr, label='Logistic Regression Classifier Sco
         re: {:.4f}'.format(roc auc score(y train, log reg pred)))
             plt.plot(knear_fpr, knear_tpr, label='KNears Neighbors Classifier Sc
         ore: {:.4f}'.format(roc_auc_score(y_train, knears_pred)))
             plt.plot(svc_fpr, svc_tpr, label='Support Vector Classifier Score:
         {:.4f}'.format(roc_auc_score(y_train, svc_pred)))
             plt.plot(tree fpr, tree tpr, label='Decision Tree Classifier Score:
         {:.4f}'.format(roc_auc_score(y_train, tree_pred)))
             plt.plot([0, 1], [0, 1], 'k--')
             plt.axis([-0.01, 1, 0, 1])
             plt.xlabel('False Positive Rate', fontsize=16)
             plt.ylabel('True Positive Rate', fontsize=16)
             plt.annotate('Minimum ROC Score of 50% \n (This is the minimum score
         to get)', xy=(0.5, 0.5), xytext=(0.6, 0.3),
                         arrowprops=dict(facecolor='#6E726D', shrink=0.05),
             plt.legend()
         graph roc curve multiple(log fpr, log tpr, knear fpr, knear tpr, svc fpr
         , svc tpr, tree fpr, tree tpr)
         plt.show()
```

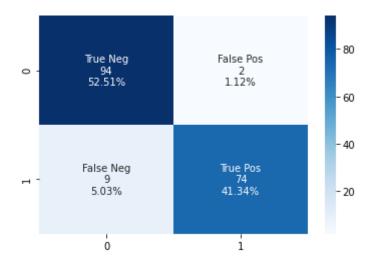


```
In [59]: labels = ['No Fraud', 'Fraud']
    undersample_pred = log_reg.predict(original_Xtest)
    print(classification_report(original_ytest, undersample_pred, target_nam
    es=labels))
```

	precision	recall	f1-score	support
No Fraud Fraud	1.00 0.15	0.99 0.88	1.00 0.26	56862 99
accuracy			0.99	56961
macro avg	0.57	0.94	0.63	56961
weighted avg	1.00	0.99	0.99	56961

```
In [60]: from sklearn.metrics import confusion matrix
         from sklearn.metrics import classification report
         import seaborn as sns
         # Logistic Regression fitted using SMOTE technique
         y_pred_log_reg = log_reg.predict(X_test)
         # Confusion Matrix
         log reg cf = confusion_matrix(y test, y pred_log_reg)
         group_names = ["True Neg", "False Pos", "False Neg", "True Pos"]
         group_counts = ["{0:0.0f}".format(value) for value in log_reg_cf.flatten
         ()]
         group_percentages = ["{0:.2%}".format(value) for value in
                               log reg cf.flatten()/np.sum(log reg cf)]
         labels = [f''(v1)\n(v2)\n(v3)'' for v1, v2, v3 in
                   zip(group names, group counts, group percentages)]
         labels = np.asarray(labels).reshape(2,2)
         sns.heatmap(log_reg_cf, annot=labels, fmt='', cmap='Blues')
```

Out[60]: <matplotlib.axes. subplots.AxesSubplot at 0x7f87655a3670>



Classification with Over Sampling

SMOTE Technique (Over-Sampling):

<img

src="https://raw.githubusercontent.com/rikunert/SMOTE visualisation/master/SMOTE R visualisation 3.png (https://raw.githubusercontent.com/rikunert/SMOTE visualisation/master/SMOTE R visualisation 3.png)", width=800> **SMOTE** stands for Synthetic Minority Over-sampling Technique. Unlike Random UnderSampling, SMOTE creates new synthetic points in order to have an equal balance of the classes. This is another alternative for solving the "class imbalance problems".

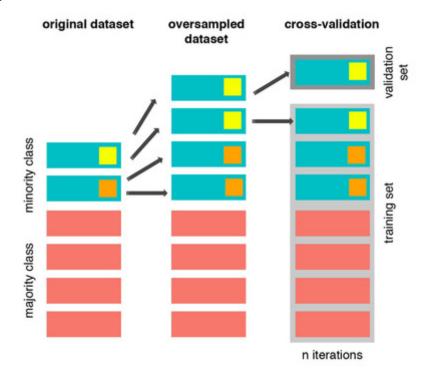
Understanding SMOTE:

- Solving the Class Imbalance: SMOTE creates synthetic points from the minority class in order to reach an equal balance between the minority and majority class.
- Location of the synthetic points: SMOTE picks the distance between the closest neighbors of the minority class, in between these distances it creates synthetic points.
- **Final Effect:** More information is retained since we didn't have to delete any rows unlike in random undersampling.
- Accuracy | Time Tradeoff: Although it is likely that SMOTE will be more accurate than random undersampling, it will take more time to train since no rows are eliminated as previously stated.

Cautionary Note on Cross Validation for SMOTE

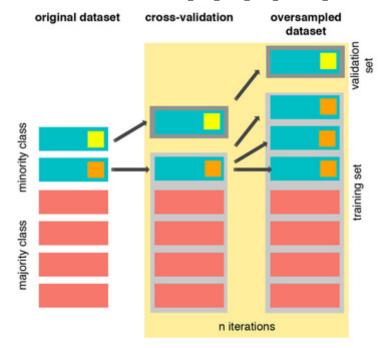
Overfitting during Cross Validation:

The Wrong Way:



As mentioned previously, if we get the minority class ("Fraud) in our case, and create the synthetic points before cross validating we have a certain influence on the "validation set" of the cross validation process. Remember how cross validation works, let's assume we are splitting the data into 5 batches, 4/5 of the dataset will be the training set while 1/5 will be the validation set. The test set should not be touched! For that reason, we have to do the creation of synthetic datapoints "during" cross-validation and not before, just like below:

The Right Way:



As you see above, SMOTE occurs "during" cross validation and not "prior" to the cross validation process. Synthetic data are created only for the training set without affecting the validation set.

References:

- DEALING WITH IMBALANCED DATA: UNDERSAMPLING, OVERSAMPLING AND PROPER CROSS-VALIDATION
- SMOTE explained for noobs
- Machine Learning Over-& Undersampling Python/ Scikit/ Scikit-Imblearn

SMOTE Implementation for OverSampling with CV for Logistics Regression

MODEL TRAIN

```
In [35]: from imblearn.over sampling import SMOTE
         from sklearn.model selection import train test split, RandomizedSearchCV
         # List to append the score and then find the average
         accuracy_lst = []
         precision lst = []
         recall_lst = []
         f1 lst = []
         auc_lst = []
         # Instantiate LR
         log reg sm = LogisticRegression()
         # LR parameters
         log_reg_params = {"penalty": ['l1', 'l2'], 'C': [0.001, 0.01, 0.1, 1, 10
         , 100, 1000]}
         # Instantiate Random Search CV
         rand log reg = RandomizedSearchCV(LogisticRegression(), log reg params,
         n iter=4)
         # Implementing SMOTE Technique
         # Cross Validating the right way
         for train, test in sss.split(original Xtrain, original ytrain):
             pipeline = imbalanced make pipeline(SMOTE(sampling strategy='minorit
         y'), rand log reg) # SMOTE happens during Cross Validation not before..
             model = pipeline.fit(original Xtrain[train], original ytrain[train])
             best est = rand log reg.best estimator
             prediction = best est.predict(original Xtrain[test])
             accuracy lst.append(pipeline.score(original Xtrain[test], original y
         train[test]))
             precision lst.append(precision score(original ytrain[test], predicti
         on))
             recall lst.append(recall score(original ytrain[test], prediction))
             f1 lst.append(f1 score(original ytrain[test], prediction))
             auc lst.append(roc auc score(original ytrain[test], prediction))
         # Get the MEAN Values of the CV accuracies
         print('---' * 45)
         print('')
         print("accuracy: {}".format(np.mean(accuracy lst)))
         print("precision: {}".format(np.mean(precision lst)))
         print("recall: {}".format(np.mean(recall lst)))
         print("f1: {}".format(np.mean(f1_lst)))
         print('---' * 45)
```

accuracy: 0.9425706072110426 precision: 0.0622266391310668 recall: 0.9137617656604998 f1: 0.11454090998118364

MODEL TEST

Classification report on test data

	precision	recall	f1-score	support
No Fraud Fraud	1.00	0.99	0.99 0.18	56862 99
rraud	0.10	0.00	0.10	99
accuracy			0.99	56961
macro avg	0.55	0.92	0.58	56961
weighted avg	1.00	0.99	0.99	56961

Precision and Recall Studies

```
In [37]: from sklearn.metrics import average_precision_score
    y_score = best_est.decision_function(original_Xtest)
    average_precision = average_precision_score(original_ytest, y_score)

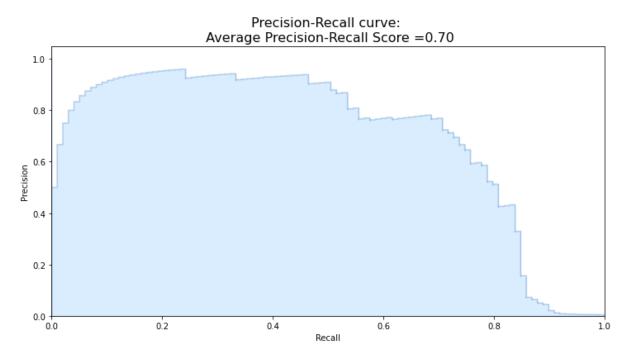
print('Average precision-recall score: {0:0.2f}'.format(
    average_precision))
```

Average precision-recall score: 0.70

Plot Precision Recall Curve

```
from sklearn.metrics import precision_recall_curve
In [38]:
         import matplotlib.pyplot as plt
         fig = plt.figure(figsize=(12,6))
         precision, recall, _ = precision_recall_curve(original_ytest, y_score)
         plt.step(recall, precision, color='#004a93', alpha=0.2,
                  where='post')
         plt.fill_between(recall, precision, step='post', alpha=0.2,
                          color='#48a6ff')
         plt.xlabel('Recall')
         plt.ylabel('Precision')
         plt.ylim([0.0, 1.05])
         plt.xlim([0.0, 1.0])
         plt.title('Precision-Recall curve: \n Average Precision-Recall Score =
         {0:0.2f}'.format(
                   average_precision), fontsize=16)
```

Out[38]: Text(0.5, 1.0, 'Precision-Recall curve: \n Average Precision-Recall Sco
 re =0.70')



Confusion Matrix

```
In [57]: from sklearn.metrics import confusion matrix
         from sklearn.metrics import classification report
         import seaborn as sns
         # Logistic Regression fitted using SMOTE technique
         y pred log reg = best est.predict(X test)
         # Confusion Matrix
         log reg cf = confusion_matrix(y test, y pred log reg)
         group_names = ["True Neg", "False Pos", "False Neg", "True Pos"]
         group_counts = ["{0:0.0f}".format(value) for value in log_reg_cf.flatten
         ()]
         group_percentages = ["{0:.2%}".format(value) for value in
                               log reg cf.flatten()/np.sum(log reg cf)]
         labels = [f''(v1)\n(v2)\n(v3)'' for v1, v2, v3 in
                   zip(group names, group counts, group percentages)]
         labels = np.asarray(labels).reshape(2,2)
         sns.heatmap(log_reg_cf, annot=labels, fmt='', cmap='Blues')
```

Out[57]: <matplotlib.axes. subplots.AxesSubplot at 0x7f8764678460>



```
In [ ]:
```