

Experiment 1:

Labelling The Unlabelled Data

```
In [33]: # Importing the General purpose libraries.
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns

# Importing the Scikit-Learn modules for preprocessing and model selection.
from sklearn.preprocessing import StandardScaler, label_binarize
from sklearn.model_selection import train_test_split

# Importing the Scikit-Learn classifiers.
from sklearn.ensemble import IsolationForest
from sklearn.linear_model import LogisticRegression
from sklearn.neural_network import MLPClassifier

# Importing the Scikit-Learn metrics and pipeline.
from sklearn.metrics import classification_report, confusion_matrix, roc_auc_score, accuracy_score
from sklearn.pipeline import Pipeline

# Importing Extra scikit-learn tools
from sklearn.cluster import DBSCAN
from sklearn.neighbors import LocalOutlierFactor
from sklearn.svm import SVC

# Importing the TensorFlow/Keras for deep learning
import tensorflow as tf
from tensorflow.keras.models import Model
from tensorflow.keras.layers import Input, Dense, Dropout
from tensorflow.keras import regularizers
from keras.models import Sequential
from keras.layers import LSTM, Dense, Dropout
from keras import utils as np_utils

# Importing dictionary and counter from python Library.
from collections import defaultdict, Counter
```

```
In [22]: # Reading the original file.
df=pd.read_csv("682200107170331.csv")
# Reading the zero filled value file.
df_zero= pd.read_csv("682200107170331_fill_zero.csv")
# Reading the Knn neighbour 2 imputed value file.
df_knn2= pd.read_csv("682200107170331_fill_knn_2.csv")
# Reading the Knn neighbour 3 imputed value file.
df_knn3= pd.read_csv("682200107170331_fill_knn_3.csv")

# Displaying the first five samples.
df_knn2.head()
```

```
Out[22]:
```

	FPAC	BLAC	CTAC	TH	MH	EGT_1	EGT_2	EGT_3	EGT_4	IVV	...	DWPT	OIPL	EHRS_4	EHRS_3	EHRS_2	TI
0	0.0	-0.011724	0.0	85.690800	98.522450	19.0	21.5	16.0	15.0	6.0	...	32832.0	0.0	8902.0	8498.0	7702.0	
1	0.0	-0.014655	0.0	86.014890	98.846535	19.0	21.5	16.0	15.0	6.0	...	32832.0	0.0	8902.0	8498.0	7702.0	
2	0.0	-0.019540	0.0	86.289540	99.121185	19.0	21.5	16.0	15.0	5.0	...	32832.0	0.0	8902.0	8498.0	7702.0	
3	0.0	-0.022471	0.0	86.580666	99.412315	19.0	21.5	16.0	15.0	3.0	...	32832.0	0.0	8902.0	8498.0	7702.0	
4	0.0	-0.023448	0.0	86.904755	99.736404	19.0	21.5	16.0	15.0	2.0	...	32832.0	0.0	8902.0	8498.0	7702.0	

5 rows × 137 columns

```
In [23]: # Selecting features - excluding 'FRMC' for model training.
# Defining the feature that are not contributing in the anomaly detection.
exclude_columns = ['FRMC', 'DATE_YEAR', 'DATE_MONTH', 'DATE_DAY']
# Collecting all the feature names except excluded columns.
features = [col for col in df_knn2.columns if col not in exclude_columns]
# Printing the final features
print(features)

['FPAC', 'BLAC', 'CTAC', 'TH', 'MH', 'EGT_1', 'EGT_2', 'EGT_3', 'EGT_4', 'IVV', 'GS', 'TRK', 'TRKM', 'DA', 'W
S', 'MW', 'DFGS', 'WD', 'ALT', 'NSQT', 'RALT', 'ALTR', 'FQTY_1', 'OIT_1', 'OIT_2', 'AOA1', 'AOA2', 'PTCH', 'FF_
1', 'PSA', 'FF_2', 'FF_3', 'ROLL', 'FF_4', 'N1_1', 'N1_2', 'MACH', 'CAS', 'APFD', 'PH', 'CASM', 'TAS', 'VRTG',
'LATG', 'PI', 'PS', 'N1_3', 'EVNT', 'MRK', 'VIB_1', 'PT', 'VHF1', 'VHF2', 'LGDN', 'LGUP', 'VIB_2', 'VHF3', 'PUS
H', 'SHKR', 'MSQT_2', 'VIB_3', 'LONG', 'PLA_1', 'N1_4', 'HYDY', 'HYDG', 'VIB_4', 'PLA_2', 'PLA_3', 'PLA_4', 'GM
T_HOUR', 'GMT_MINUTE', 'GMT_SEC', 'ACMT', 'FQTY_2', 'OIT_3', 'OIT_4', 'BLV', 'EAI', 'PACK', 'AOAI', 'AOAC', 'BA
L1', 'BAL2', 'WOW', 'N2_1', 'N2_2', 'N2_3', 'N2_4', 'TAT', 'SAT', 'N1T', 'N1C', 'OIP_1', 'OIP_2', 'FQTY_4', 'CR
SS', 'HDGS', 'ALTS', 'CASS', 'N1CO', 'VSPS', 'MNS', 'MSQT_1', 'VMODE', 'LMOD', 'A_T', 'OIP_3', 'OIP_4', 'LOC',
'GLS', 'LONP', 'ABRK', 'ECYC_1', 'ECYC_2', 'FLAP', 'SPLY', 'SPLG', 'BPGR_2', 'BPYR_1', 'ECYC_3', 'ECYC_4', 'EHR
S_1', 'DWPT', 'OIPL', 'EHRS_4', 'EHRS_3', 'EHRS_2', 'TMODE', 'ATEN', 'LATP', 'FGC3', 'ILSF']
```

1. Anomaly Detection (Label Generation) Using iForest Model

```
In [24]: # Defining the isolation forest method which take data and threshold values which is 9% here.
def apply_isolation_forest(df, threshold_factor=0.09):
    # Exclding the columns from the features that are not contributing for the anomaly detection.
    excluded_columns = ['FRMC', 'DATE_YEAR', 'DATE_MONTH', 'DATE_DAY']
    # Getting all the features except the excluded ones.
    features = [col for col in df.columns if col not in excluded_columns]
    # Defining the isolation forest model with 100 estimators and keeping the contamination rate to 0.01.
    model = IsolationForest(n_estimators=100, contamination=0.01, random_state=42)
    # fitting the data into model.
    model.fit(df[features])

    # Calculating the Anomaly scores.
    anomaly_scores = model.decision_function(df[features])

    # Choosing a threshold for classifying anomalies based on the provided threshold factor.
    threshold = np.percentile(anomaly_scores, threshold_factor * 100)

    # Labelling instances as normal (1) or anomalous (-1) based on the threshold.
    labels = np.where(anomaly_scores < threshold, -1, 1)

    # Creating a column named Labels and passing the labels into it.
    df['labels'] = labels
    # Mapping 1 for Anomaly and 0 for Normal.
    df['labels'] = df['labels'].apply(lambda x: 1 if x == -1 else 0)
    # Storing the Anomaly Score into anomaly_scores column.
    df['anomaly_scores'] = anomaly_scores
    return df
```

```
In [25]: # Identifying Anomaly using Isolation Forest for both knn2 and knn3 file.
df_knn2 = apply_isolation_forest(df_knn2)
df_knn3 = apply_isolation_forest(df_knn3)
df_knn2.head()
```

```
Out[25]:
```

	FPAC	BLAC	CTAC	TH	MH	EGT_1	EGT_2	EGT_3	EGT_4	IVV	...	EHRS_4	EHRS_3	EHRS_2	TMODE	ATEN
0	0.0	-0.011724	0.0	85.690800	98.522450	19.0	21.5	16.0	15.0	6.0	...	8902.0	8498.0	7702.0	2.0	0.0
1	0.0	-0.014655	0.0	86.014890	98.846535	19.0	21.5	16.0	15.0	6.0	...	8902.0	8498.0	7702.0	2.0	0.0
2	0.0	-0.019540	0.0	86.289540	99.121185	19.0	21.5	16.0	15.0	5.0	...	8902.0	8498.0	7702.0	2.0	0.0
3	0.0	-0.022471	0.0	86.580666	99.412315	19.0	21.5	16.0	15.0	3.0	...	8902.0	8498.0	7702.0	2.0	0.0
4	0.0	-0.023448	0.0	86.904755	99.736404	19.0	21.5	16.0	15.0	2.0	...	8902.0	8498.0	7702.0	2.0	0.0

5 rows × 139 columns

2. Anomaly Detection (Label Generation) Using Autoencoder

```

In [26]: # Defining the autoencoder function.
def detect_anomalies_autoencoder(df):
    # Preprocess data: scale features, exclude non-feature columns
    excluded_columns = ['FRMC', 'DATE_YEAR', 'DATE_MONTH', 'DATE_DAY', 'labels']
    features = [col for col in df.columns if col not in excluded_columns]
    # Assigning the features
    df_features = df[features]
    scaler = StandardScaler()
    # performing the scaling operation on the feature values.
    df_scaled = scaler.fit_transform(df_features.values)

    # Splitting the data for testing and training which is 80% and 20% respectively.
    X_train, X_test = train_test_split(df_scaled, test_size=0.2, random_state=42)

    # Defining the autoencoder model.
    input_dim = X_train.shape[1]
    input_layer = Input(shape=(input_dim,))

    # Encoding the data using relu function.
    encoded = Dense(64, activation='relu', activity_regularizer=regularizers.l1(10e-5))(input_layer)
    encoded = Dense(32, activation='relu')(encoded)
    encoded = Dense(16, activation='relu')(encoded)

    # Decoding the data using relu function where using linear activation function.
    decoded = Dense(16, activation='relu')(encoded)
    decoded = Dense(32, activation='relu')(decoded)
    decoded = Dense(64, activation='relu')(decoded)
    decoded = Dense(input_dim, activation='linear')(decoded)

    # Making the autoencoder by defining the layers into it.
    autoencoder = Model(inputs=input_layer, outputs=decoded)

    # Compiling the model using adam optimizer and for loss function using mean squared error for the model
    autoencoder.compile(optimizer='adam', loss='mean_squared_error')

    # Training the model (Having 100 episodes with having batch size of 100)
    autoencoder.fit(X_train, X_train,
                    epochs=100,
                    batch_size=256,
                    shuffle=True,
                    validation_data=(X_test, X_test),
                    verbose=0)

    # Predicting on the full data
    predictions = autoencoder.predict(df_scaled)

    # Computing the reconstruction error
    mse = np.mean(np.power(df_scaled - predictions, 2), axis=1)

    # Determining a threshold for anomaly
    threshold = np.quantile(mse, 0.95)

    # Creating the 'labels_2' column, where 1 indicates an anomaly and 0 indicates normal
    df['labels_2'] = [1 if e > threshold else 0 for e in mse]

    # Comparing the labels generated by both the models.
    matches = (df['labels'] == df['labels_2'])

    # Counting the matching rows and mismatched rows.
    matching_rows = matches.sum()
    mismatching_rows = len(df) - matching_rows

    # Filtering the rows where labels are matching
    matching_df = df[matches]
    # Returns only rows where both the model detects anomaly
    return matching_df.drop(columns=['labels_2']), matching_rows, mismatching_rows

# Detecting Anomaly Using Autoencoder
df_knn2, matches_knn2, mismatches_knn2 = detect_anomalies_autoencoder(df_knn2)

df_knn3, matches_knn3, mismatches_knn3 = detect_anomalies_autoencoder(df_knn3)

```

6708/6708 ————— 10s 1ms/step
 6708/6708 ————— 10s 1ms/step

```
In [20]: # Printing the matching and mismatching rows for the df_knn2
print(f'For df_knn2:')
print(f'Number of matching rows: {matches_knn2}')
print(f'Number of mismatching rows: {mismatches_knn2}')
print(f'Proportion of accurate labeling:{matches_knn2/(matches_knn2+mismatches_knn2)*100:.2f}%')

# Printing the matching and mismatching rows for the df_knn3
print(f'\nFor df_knn3:')
print(f'Number of matching rows: {matches_knn3}')
print(f'Number of mismatching rows: {mismatches_knn3}')
print(f'Proportion of accurate labeling:{matches_knn3/(matches_knn3+mismatches_knn3)*100:.2f}%')

For df_knn2:
Number of matching rows: 192156
Number of mismatching rows: 22500
Proportion of accurate labeling:89.52%

For df_knn3:
Number of matching rows: 191228
Number of mismatching rows: 23428
Proportion of accurate labeling:89.09%
```

Visualize Normal Vs Anomaly for Selected Features.

The visualization is done on the few critical features that are commonly monitored in aviation for safety and performance:

- Engine Gas Temperature (EGT) - represented here as EGT_1, EGT_2, EGT_3, and EGT_4 for multiple engines.
- Mach Number (MH) - indicating the aircraft's speed relative to the speed of sound.
- Total Air Temperature (TAT) - labeled as TH here, which affects engine performance and efficiency.

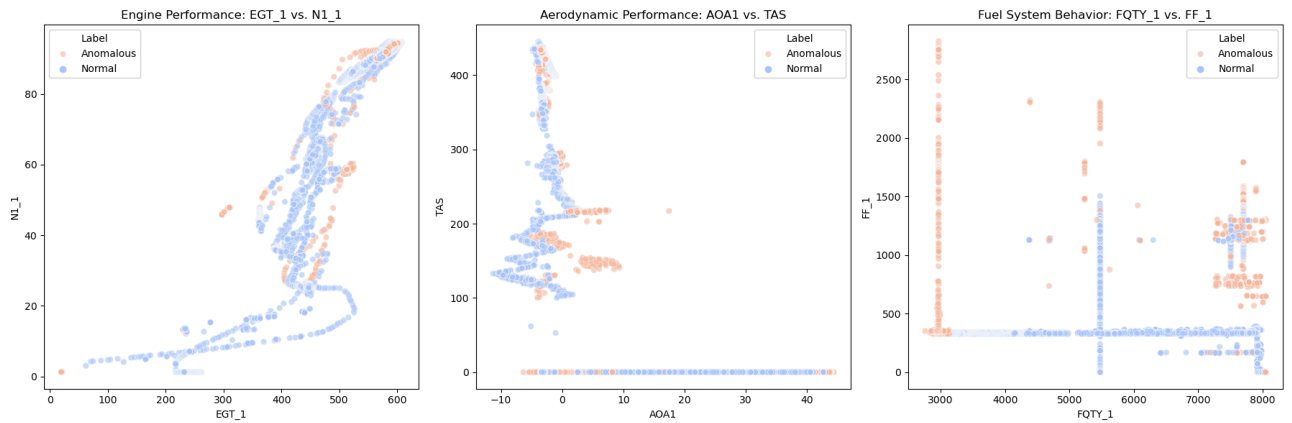
```
In [27]: # Scatter Plot of Engine Performance, aerodynamic performance, fuel system behavior on df_knn2
#       EGT_1 vs. N1_1 to understand engine performance.
#       AOA1 vs. TAS for aerodynamic performance.
#       FQTY_1 vs. FF_1 for insights into fuel system behavior.

# Visualization setup for the selected feature pairs
feature_pairs = [('EGT_1', 'N1_1'), ('AOA1', 'TAS'), ('FQTY_1', 'FF_1')]
# Defining the titles for the graphs
titles = ['Engine Performance: EGT_1 vs. N1_1',
          'Aerodynamic Performance: AOA1 vs. TAS',
          'Fuel System Behavior: FQTY_1 vs. FF_1']

# Setting up the matplotlib figure.
plt.figure(figsize=(18, 6))

# Iterating the data for graphs.
for i, (x, y) in enumerate(feature_pairs, 1):
    # Generating 3 subplots.
    plt.subplot(1, 3, i)
    # Printing the graph for the knn 2 file.
    sns.scatterplot(data=df_knn2, x=x, y=y, hue='labels', palette='coolwarm', alpha=0.6)
    # Plotting the graph with the label and title.
    plt.title(titles[i-1])
    plt.xlabel(x)
    plt.ylabel(y)
    # Defining the legend.
    plt.legend(title='Label', labels=['Anomalous', 'Normal'])

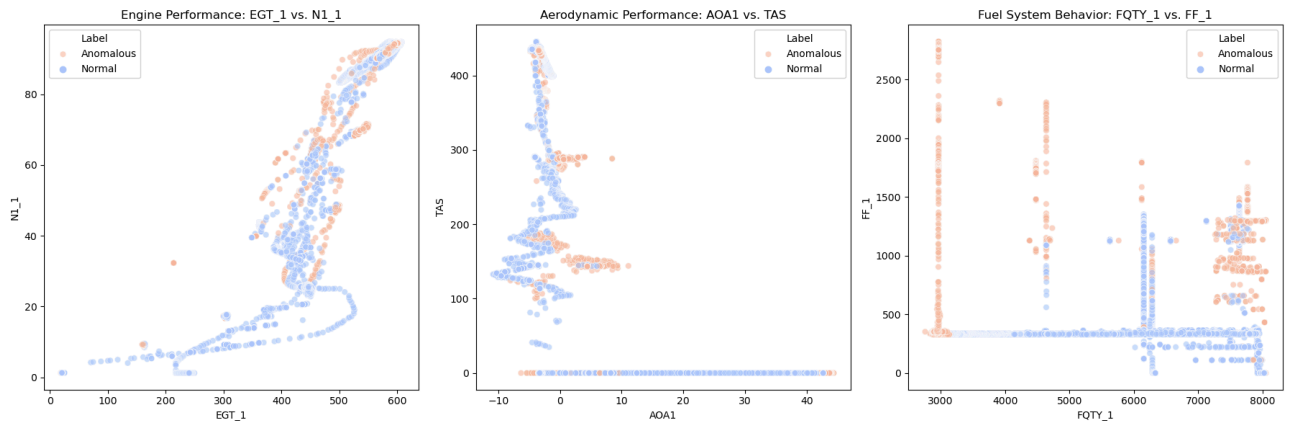
# Plotting the three subplots.
plt.tight_layout()
plt.show()
```



```
In [28]: # Scatter Plot of Engine Performance, aerodynamic performance, fuel system behavior on df_knn3
#         EGT_1 vs. N1_1 to understand engine performance.
#         AOA1 vs. TAS for aerodynamic performance.
#         FQTY_1 vs. FF_1 for insights into fuel system behavior.

# Setting up the matplotlib figure.
plt.figure(figsize=(18, 6))

# Iterating through Loop
for i, (x, y) in enumerate(feature_pairs, 1):
    # Generating 3 subplots.
    plt.subplot(1, 3, i)
    sns.scatterplot(data=df_knn3, x=x, y=y, hue='labels', palette='coolwarm', alpha=0.6)
    plt.title(titles[i-1])
    plt.xlabel(x)
    plt.ylabel(y)
    plt.legend(title='Label', labels=['Anomalous', 'Normal'])
# Displaying the layout
plt.tight_layout()
plt.show()
```



3. Supervised Learning - Classification of Labels (Confidence Checking)

In the below defined method (Process_and_evaluate_models), we are performing the validation of the labels using three models which are Logistic Regression, SVM (Support Vector Machines), Neural Network (MLP Classifier). Using 80% data for training and 20% Data for Testing.

```
In [34]: def process_and_evaluate_models(df, target_column, test_size=0.2, random_state=42):
# Define features excluding the target column and other non-feature columns.
excluded_columns = ['FRMC', 'DATE_YEAR', 'DATE_MONTH', 'DATE_DAY', target_column]
features = [col for col in df.columns if col not in excluded_columns]

# Preprocessing the data: scaling the data
scaler = StandardScaler()
X = scaler.fit_transform(df[features])
```

```

y = df[target_column]

# Splitting the data into training and testing datasets
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=test_size, random_state=random_state)

# Defining the number of models to be evaluated
models = {
    # For Logistic Regression we are using 10000 maximum iteration.
    "Logistic Regression": LogisticRegression(max_iter=10000),
    # Using inbuilt SVM model
    "Support Vector Machine": SVC(probability=True),
    # For Neural Network we are using MLP (Multilayer Preceptron Classifier) Classifier.
    "Neural Network": MLPClassifier(max_iter=1000)
}

# Evaluating each model
evaluation_results = {}
# Iterating through each model.
for name, model in models.items():
    # fitting the data into model
    model.fit(X_train, y_train)
    # Getting the predictions from the model
    y_pred = model.predict(X_test)
    y_pred_proba = model.predict_proba(X_test)[: , 1] if hasattr(model, "predict_proba") else None
    # Getting accuracy from the model
    accuracy = accuracy_score(y_test, y_pred)
    # Getting the details for the AUC score
    auc = roc_auc_score(y_test, y_pred_proba) if y_pred_proba is not None else "N/A"
    # Getting the confusion matrix
    conf_matrix = confusion_matrix(y_test, y_pred)
    report = classification_report(y_test, y_pred, output_dict=True)

    # Defining the structure for evaluation for the models
    evaluation_results[name] = {
        "Classification Report": pd.DataFrame(report).transpose(),
        "Confusion Matrix": conf_matrix,
        "AUC": auc,
        "Accuracy": accuracy
    }

# Displaying the results for the models.
for model_name, results in evaluation_results.items():
    print(f"Model: {model_name}")
    print(f"Accuracy: {results['Accuracy']:.4f}, AUC: {results['AUC']}")
    print("Confusion Matrix:")
    print(results['Confusion Matrix'])
    print("Classification Report:")
    print(results['Classification Report'])
    print("\n" + "-"*60 + "\n")

```

1. Complete Data Based Analysis

```

In [35]: # Performing the Operation for df_knn2
process_and_evaluate_models(df_knn2, 'labels')

```

Model: Logistic Regression
 Accuracy: 0.9996, AUC: 0.9999863765766756
 Confusion Matrix:
 [[37632 3]
 [11 736]]
 Classification Report:

	precision	recall	f1-score	support
0	0.999708	0.999920	0.999814	37635.000000
1	0.995940	0.985274	0.990579	747.000000
accuracy	0.999635	0.999635	0.999635	0.999635
macro avg	0.997824	0.992597	0.995196	38382.000000
weighted avg	0.999634	0.999635	0.999634	38382.000000

Model: Support Vector Machine
 Accuracy: 0.9994, AUC: 0.9998751126911437
 Confusion Matrix:
 [[37628 7]
 [15 732]]
 Classification Report:

	precision	recall	f1-score	support
0	0.999602	0.999814	0.999708	37635.000000
1	0.990528	0.979920	0.985195	747.000000
accuracy	0.999427	0.999427	0.999427	0.999427
macro avg	0.995065	0.989867	0.992451	38382.000000
weighted avg	0.999425	0.999427	0.999425	38382.000000

Model: Neural Network
 Accuracy: 0.9997, AUC: 0.99983502496768
 Confusion Matrix:
 [[37629 6]
 [7 740]]
 Classification Report:

	precision	recall	f1-score	support
0	0.999814	0.999841	0.999827	37635.000000
1	0.991957	0.990629	0.991293	747.000000
accuracy	0.999661	0.999661	0.999661	0.999661
macro avg	0.995886	0.995235	0.995560	38382.000000
weighted avg	0.999661	0.999661	0.999661	38382.000000

```
In [36]: # Performing the operation for the df_knn3
process_and_evaluate_models(df_knn3, 'labels')
```

Model: Logistic Regression
 Accuracy: 0.9997, AUC: 0.9999895192155674
 Confusion Matrix:
 [[37619 0]
 [12 617]]
 Classification Report:

	precision	recall	f1-score	support
0	0.999681	1.000000	0.999841	37619.000000
1	1.000000	0.980922	0.990369	629.000000
accuracy	0.999686	0.999686	0.999686	0.999686
macro avg	0.999841	0.990461	0.995105	38248.000000
weighted avg	0.999686	0.999686	0.999685	38248.000000

Model: Support Vector Machine
 Accuracy: 0.9995, AUC: 0.9999975065875745
 Confusion Matrix:
 [[37618 1]
 [18 611]]
 Classification Report:

	precision	recall	f1-score	support
0	0.999522	0.999973	0.999748	37619.000000
1	0.998366	0.971383	0.984690	629.000000
accuracy	0.999503	0.999503	0.999503	0.999503
macro avg	0.998944	0.985678	0.992219	38248.000000
weighted avg	0.999503	0.999503	0.999500	38248.000000

Model: Neural Network
 Accuracy: 0.9996, AUC: 0.9996806741646255
 Confusion Matrix:
 [[37614 5]
 [10 619]]
 Classification Report:

	precision	recall	f1-score	support
0	0.999734	0.999867	0.999801	37619.000000
1	0.991987	0.984102	0.988029	629.000000
accuracy	0.999608	0.999608	0.999608	0.999608
macro avg	0.995861	0.991984	0.993915	38248.000000
weighted avg	0.999607	0.999608	0.999607	38248.000000

4. Extra models for Evaluation (LSTM, LOF)

```
In [37]: # Defining the method with the same feature as mentioned above.
def process_and_evaluate_models(df, target_column, test_size=0.2, random_state=42, lstm=False):
    # Defining features excluding the target column and non-feature columns
    excluded_columns = ['FRMC', 'DATE_YEAR', 'DATE_MONTH', 'DATE_DAY', target_column]
    features = [col for col in df.columns if col not in excluded_columns]

    # performing scalling operation on features
    scaler = StandardScaler()
    X = scaler.fit_transform(df[features])
    y = df[target_column]

    # Splitting the data
    X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=test_size, random_state=random_state)

    # Defining the models that are being used.
    models = {
        # Using DBSCAN (Density-Based Spatial Clustering of Applications with Noise) model
        "DBSCAN": DBSCAN(),
        # Using Local outlier factor model with 20 neighbours
        "Local Outlier Factor": LocalOutlierFactor(n_neighbors=20)
    }

    # performing the operations for the LSTM model
    if lstm:
        # For LSTM Model - Assuming binary classification out project.
        y_train_enc = np_utils.to_categorical(y_train)
```



```

y_test_enc = np_utils.to_categorical(y_test)

# Reshaping the input to be [samples, time steps, features]
X_train_lstm = X_train.reshape((X_train.shape[0], 1, X_train.shape[1]))
X_test_lstm = X_test.reshape((X_test.shape[0], 1, X_test.shape[1]))

# Defining the LSTM model layers
lstm_model = Sequential()
lstm_model.add(LSTM(50, input_shape=(X_train_lstm.shape[1], X_train_lstm.shape[2])))
# Keeping the dropout rate of 50%
lstm_model.add(Dropout(0.5))
# Using softmax function as the activation function for the model.
lstm_model.add(Dense(2, activation='softmax'))
lstm_model.compile(loss='binary_crossentropy', optimizer='adam', metrics=['accuracy'])
# Assigning the defined model
models["LSTM"] = lstm_model

# Evaluating the models
evaluation_results = {}
# Iterating through models
for name, model in models.items():
    # printing the for DBSCAN and LOF.
    if name in ["DBSCAN", "Local Outlier Factor"]:
        # If the model is DBSCAN then it will perform the operation.
        if name == "DBSCAN":
            model.fit(X)
            labels = model.labels_
        else:
            model.fit(X)
            labels = model.fit_predict(X)

        # Converting the Labels to binary (0: normal, 1: anomaly)
        labels = [1 if x == -1 else 0 for x in labels]
        accuracy = accuracy_score(y, labels)
        evaluation_results[name] = {"Accuracy": accuracy}
        # if the model is LSTM then performing the operation for the LSTM.
    elif name == "LSTM":
        # fitting the data into model
        model.fit(X_train_lstm, y_train_enc, epochs=10, batch_size=64, validation_data=(X_test_lstm, y_test_
_, accuracy = model.evaluate(X_test_lstm, y_test_enc, verbose=0)
        evaluation_results[name] = {"Accuracy": accuracy}

# Printing the results
for model_name, results in evaluation_results.items():
    print(f"Model: {model_name}, Accuracy: {results.get('Accuracy', 'N/A')}")
    if model_name == "LSTM":
        print(f"LSTM Test Accuracy: {results['Accuracy']:.4f}")

```

```

In [38]: # Performing the evaluation for Knn3 for DBSCAN, LOF and LSTM model
process_and_evaluate_models(df_knn3, 'labels')

```

```

Model: DBSCAN, Accuracy: 0.8806657742266101
Model: Local Outlier Factor, Accuracy: 0.8808278775962685

```

```

In [ ]:

```