Business Understanding

The local police department has required the development of a binary prediction automated system that could determine the sex of individuals from the footprints that have been left at crime scenes, for the automated model, the local police force requires, needs to be able to make reasonably high predictions accuracy, within a limited of time, that will be used on a new device and to help the investigation team to narrow down suspects on the initial stages.

To achieve these targets, we have been given a set of data that contains 18 landmarks in the form of X and y coordinates, the report below will provide a detailed examination of the data and its findings, the decision-making of each process, and recommendations for potential improvements and future work.

step 0: Prepareing

At step zero, we will first be setting up the necessary components for the work to work seamlessly and error free.

local RUN setup

import zipfile

```
pip install kaggle pandas joblib numpy matplotlib seaborn xgboost scipy statsmodels

!pip install kaggle xgboost joblib statsmodels

pip install —upgrade kaggle pandas joblib numpy matplotlib seaborn xgboost scipy st

pip install —upgrade kaggle pandas joblib numpy matplotlib seaborn xgboost scipy st
```

Import List

```
import kaggle
import pandas as pd
from joblib import dump, load
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from xgboost import XGBClassifier, p
from scipy import stats
from scipy.stats import spearmanr
from scipy.stats.mstats import winsor
from sklearn.svm import SVC
from sklearn import metrics
from sklearn.metrics import classific
from sklearn.impute import KNNImputer
from sklearn.experimental import enak
from sklearn.impute import Iterativel
from sklearn.preprocessing import Rok
from sklearn.naive_bayes import Gauss
from sklearn.ensemble import Gradient
from sklearn.ensemble import Isolatic
from sklearn.neighbors import LocalOt
from sklearn. model selection import t
from sklearn.linear model import Logi
from sklearn import linear_model
from sklearn, neighbors import KNeight
from sklearn.tree import DecisionTree
from sklearn.metrics import accuracy
```

funstion list

In function list it will hold all the implement function for features use and robustness.

```
X_train_variants = {}
X_test_variants = {}
y_train_variants = {}
y_test_variants = {}
```

distance clataiton

lengths and widths

```
def lengths_widths_calculation(df):
    df_lengths_widths = df.copy()
    df_lengths_widths['lengths'] =
    df_lengths_widths['widths'] =
    return df_lengths_widths
```

7 foot point

```
def point7_calculation(df):
    df_7_point_footprints = df.cop
    df_7_point_footprints['T1'] =
    df_7_point_footprints['T2'] =
    df_7_point_footprints['T3'] =
    df_7_point_footprints['T4'] =
    df_7_point_footprints['T5'] =
    df_7_point_footprints['BAB'] =
    df_7_point_footprints['BAB'] =
```

```
return \quad df\_7\_point\_footprints
```

IQR missing value function

```
def IQR(df):
    df_outlier_IQR = df.copy()
    for column in df_outlier_IQR.
    Q1 = df_outlier_IQR[cc
    Q3 = df_outlier_IQR[cc
    IQR = Q3 - Q1
    lower_bound = Q1 - 3
    upper_bound = Q3 + 3

    df_outlier_IQR[column]
```

Cap Outliers and Apply Robust and standard Scaling

```
def cap_outliers_and_scale(df):
    df_outlier_capped_scale = df.c

for column in df_outlier_cappe
    Q1 = df_outlier_cappec
    Q3 = df_outlier_cappec
    IQR = Q3 - Q1
    lower_bound = Q1 - 1
    upper_bound = Q3 + 1
    df_outlier_capped_scale[

robust_scaler = RobustScaler()
    df_outlier_capped_scale = robu

standard_scaler = StandardScal
    df_outlier_capped_scale = star

return pd.DataFrame(df_outlier_
```

Winsorization

z score

```
def z_score(df):
    df_z_score = df.copy()
    z_threshold = 4
    for column in df_z_score.columly column = restant to the score of the score
```

isolation_forest

```
def isolation_forest(df, contamination
    df_isolation = df.copy()
    model = IsolationForest(contam

    model.fit(df_isolation)

    outlier_predictions = model.pr

    for column in df_isolation.cc
        median_value = df_isol
        df_isolation[column] =

    return df_isolation
```

step 1: Understanding the data

開始使用 AI 編寫或生成程式碼。

Although all landmarks are provided, it does not necessarily mean all of them will be positive for the model, therefore we will implement features engineering, This involves both adding new features and feature selection to improve model learning, details on feature engineering will be discussed in a later section.

The data has been standardized between 0 and 1, if needed, we can recover to the original values by scaling back to 2240x3200, this will bring us back to its true data form, for more data understanding.

The dataset contains 2,000 entries, which will be used to train the model and between them, x1 to y17 contain 6 to 17 missing values in between that require handling to ensure the data quality, and we will experiment with different imputation methods in step 3.

```
footprints_data = pd.read_csv('SexLand
print(footprints_data.info())
footprints_data.head()

footprints data.isnull().sum()
```

In this step, on "Box Plots for Outliers", outliers are present on the dataset, for early outlier handling, we can scale back the standardized data and calculate basic length and width, as it is difficult to gain meaningful information from the basic box plots, by doing so, we can identify extreme outliers more easily and correct them manually if needed, this method allows us to clean data more consistently, as leaving unreasonable extreme outliers most likely hurt the robustness of the dataset and effectiveness of the deployment.

```
plt.figure(figsize=(15, 10))
sns.boxplot(data=footprints_data)
plt.xticks(rotation=90)
plt.title("Box Plots for Outliers")
plt.show()

width, height = 2240, 3200
```

```
original scaled data = footprints data
for column in original scaled data.cc
       if column.startswith('x'):
               original scaled data[col
       elif column.startswith('y'):
               original scaled data[col
print(original scaled data.head())
plt.figure(figsize=(15, 10))
sns.boxplot(data=original scaled data)
plt. xticks (rotation=90)
plt.title("Box Plots for Outliers")
plt.show()
original scaled data with lengths widths
for name, group in original scaled of
       plt.plot(group.lengths, group.w
plt.legend()
```

The graph below shows the length and width of each footprint, Based on it we can observe extreme outliers, we will check if should we remove or correct these outliers, based on the landmark and dose it relistic.

```
import matplotlib.pyplot as plt

plt.figure(figsize=(10, 6))

plt.scatter(original_scaled_data_with_le

plt.scatter(big_feet['lengths'], big_fe
plt.scatter(small_feet['lengths'], smal

plt.xlabel('Lengths')
plt.ylabel('Widths')
plt.legend()
plt.show()
```

As shown in the graph below, the coordinates of the small feet, has shown a spread that are hardly can be recognized as human, therefore drop these data point from the dataset should improve the dataset.

On the other hand both of the big foot seems to be showing a normal spared therefore they will be kept.

```
plot_footprint(small_foot_1.iloc[0], 'S
```





I have use iqr base dataset for all of them now to just have atry, if need change it back

```
plot_footprint(small_foot_2.iloc[0], 'S

plot_footprint(big_foot_1.iloc[0], 'Big

plot_footprint(big_foot_2.iloc[0], 'Big

indices_to_drop = [small_foot_1.index[

footprints_data = footprints_data.drop

footprints_data.describe().T
```

The graph below shown there is a class imbalance on the dataset, it will be the best practice to implement the Synthetic Minority Over-sampling Technique (SMOTE) to prevent model bias. SMOTE will generate synthetic samples for the minority class, this can help to balance the dataset and improve the model's ability to generalize both classes.

```
barplot=(sns.countplot(data= footprints
plt. title('0 v/s 1 n')
corr = footprints data.corr(method='sr
triangle = np. triu(corr)
plt.figure(figsize=(16, 7))
sns.heatmap(data=corr, annot=True, mas
plt.figure(figsize=(20, 12))
sns.set context('notebook', font scale =
sns.heatmap(footprints data.corr(), annot
plt.tight_layout()
ax = sns.heatmap(
       corr,
       vmin=-1, vmax=1, center=0,
       cmap=sns.diverging_palette(20,
       square=True
ax.set xticklabels(
       ax.get_xticklabels(),
       rotation=45,
```

```
horizontalalignment='right'
);
ax
```

Th is dataset has shown there is no duplicated, therefore no action needed

```
footprints\_data.\,duplicated().\,value\_count
```

step 2: data processing

outliers handling

```
footprints_data_df = footprints_data.c
footprints_data_df.describe().T
```

we will uses 4 method to handle outliers, and we will not be dropping outliers, because as seen there is meaningful data with in the outliers, therefore Dropping them could bring loss of important patterns.

1.Basic IQR Method:

 The Interquartile Range (IQR) is a standard technique used to identify outliers, the outliers will be capped to a bounds, to limit their range.

2.Cap Outliers and Apply Robust Scaling

 Similar to the IQR method but apply robusts and standard scaling to create deviation of the data.

3.Winsorization

 limits exteme values by capping them within specified boundaries.

4.Use Z score

 uses standard deviation to identify outliers, which are then replaced replaced with the median to reduce their effect.

```
footprints_data_df = footprints_data.c
```

use IQR for outliners

```
footprints_data_df = footprints_data.c
footprints_data_df.describe().T
```

```
X = footprints_data_df.drop('sex', ax
y = footprints_data_df['sex']
```

```
X_train, X_test, y_train, y_test =

X_train_iqr = IQR(X_train)
X_test_iqr = IQR(X_test)

y_train_iqr = y_train.loc[X_train_iqr.
y_test_iqr = y_test.loc[X_test_iqr.inc
```

```
X_train_variants['IQR'] = X_train_iqr
X_test_variants['IQR'] = X_test_iqr
y_train_variants['IQR'] = y_train_iqr
y_test_variants['IQR'] = y_test_iqr
```

X_train_iqr.describe()

Cap Outliers and Apply Robust and standard Scaling

```
X_train, X_test, y_train, y_test =

X_train_robust = cap_outliers_and_scal
X_test_robust = cap_outliers_and_scale
y_train = y_train.reset_index(drop=Tru
y_test = y_test.reset_index(drop=True)

y_train_robust = y_train.loc[X_train_r
y_test_robust = y_test.loc[X_test_robust]

X_train_variants['RobustScaling'] = X_t
y_train_variants['RobustScaling'] = y_t
y_test_variants['RobustScaling'] = y_t

lengths_widths_df_robust = pd.concat([
for name, group in lengths_widths_df
plt.plot(group.x1, group.x14,
```

lengths widths df robust.describe()

plt.legend()

for now we have done with the grouping onto lengths and widths which we have mentioned earlier and we have use 2 ways to deal with outliers

Winsorization

```
X_train, X_test, y_train, y_test =

X_train_Winsorization = Winsorization()

X_test_Winsorization = y_train.loc[X_y_train_Winsorization = y_test.loc[X_test_Winsorization = y_test.loc[X_test_Winsorization'] = X_X_test_variants['Winsorization'] = X_ty_train_variants['Winsorization'] = y_y_test_variants['Winsorization'] = y_t

lengths_widths_df_Winsorization = pd.c

for name, group in lengths_widths_df
```

```
plt.plot(group.xl, group.xl4, plt.legend()
```

z_score_df

```
X_train, X_test, y_train, y_test =

X_train_z_score = z_score(X_train)
X_test_z_score = z_score(X_test)
y_train = y_train.reset_index(drop=Tru
y_test = y_test.reset_index(drop=True)

y_train_z_score = y_train.loc[X_train_y_test_z_score = y_test.loc[X_test_rob]

X_train_variants['z_score'] = X_train_X_test_variants['z_score'] = X_test_z_y_train_variants['z_score'] = y_train_y_test_variants['z_score'] = y_test_z_

lengths_widths_df_z_score = pd.concat(
for name, group in lengths_widths_df_plt.plot(group.xl, group.xl4, plt.legend()
```

lengths widths df z score.describe()

step 3: missing value handle

At this step, we group the data that has been processed for outliers handling, assign key values for easier management, then we apply KNN Imputer and Iterative Imputer, this avoids data leakage meanwhile being efficient. These two imputation methods are chosen because:

1. KNN Imputer:

- The KNN fills up missing values by averaging the values from the nearest neighbours, this helps missing values while keeping the patterns related to those neighbours.
- 2. Iterative Imputer:
- The Iterative predicts each missing value by running an iterative regression.

```
datasets = {
       'IQR': (X_train_iqr, X_test_iq
       'RobustScaling': (X train robus
       'Winsorization': (X train Winso
       'Zscore': (X train z score, X
KNNImputer = KNNImputer(n neighbors=4)
IterativeImputer = IterativeImputer(ma
imputed variants = {}
for variant name, (X train, X test,
       X train imputed = pd.DataFrame
       X test imputed = pd.DataFrame(
       key = f"{variant_name}_Iterati
       imputed variants[key] = (X tra
for variant name, (X train, X test,
       X train imputed = pd.DataFrame
       X test imputed = pd.DataFrame(
       key = f"{variant_name}_knn"
       imputed variants[key] = (X tra
print(imputed_variants.keys())
```

step 4: baseline test

before features engineering In this step, we will prepare a baseline test for the performance of the models, as at this point we have already cleaned up our data with the basic method we have covered, and now the data are already for a baseline test and we will choose models that perform well for further development.

```
models = {
    "Logistic Regression": Logisti
    "Gaussian Naïve Bayes": Gauss
    "Support Vector Machine": SVC
    "KNN Classifier": KNeighborsCl
    "Decision Tree": DecisionTree(
    "Random Forest": RandomForestC
    "Gradient Boosting": GradientF
    "AdaBoost": AdaBoostClassifier(
    "XGBoost": XGBClassifier(use_la"
    "SGDOneClassSVM":linear_model.SC
```

The results from the baseline test has shown a acceptable performance concider we have only processed with basic methodology to clean the

The list below has shown the results in order of accuracy score, along with the model, variant of the dataset, outliers method and imputation method. According to the results, the best-performing model so far is XGBoost, which uses IQR_Iterative and it able to achieve 0.8250, This suggests further development of XGBoost will be worthwhile, followed by Random Forest and Gradient Boosting, with a different set of variants, it has also shown there is

not yet have a clear idea of which variants will be the best for us to achieve our goal there for more test will be needed in future steps.

```
results = []
for model name, model in models.item
       for dataset name, (X train, )
               model.fit(X train, y tr
               y pred = model.predict
               accuracy = accuracy so
               results.append({
                       "Model": model
                       "Variant": data
                       "Outlier Handli
                       "Imputation Met
                       "Accuracy": acc
               })
results df = pd. DataFrame (results)
results df = results df. sort values (by
pd.set_option('display.max_rows', 100)
pd. set option ('display. max columns', No
pd. set option ('display. width', 1000)
print(results df)
```

step 5: features engineering

At this step we will implement two different kinds of feature engineering, first, we will simply calculate the fundamental lengths and widths of the footprints, to obtain extra measurements that can help understanding of the size.

Second approach involves using a more unique feature extraction technique based on research of Abledu et al. (2015), published by the NIH (National Library of Medicine), they have implemented an calculation of Seven dimensions—length of each toe to the bottom (t1 to t5), breadth at the ball (BAB) and breadth at heel (BAH), will this approach they have able to achieve a remarkable accuracy in a similar tasks, therefore we will implement this along with the basic lengths and widths calculation.

ref

Abledu, J. K., Abledu, G. K., Offei, E. B., and Antwi, E. M., 2015.

Determination of sex from footprint dimensions in a Ghanaian population [online]. PloS one. Available from: https://pmc.ncbi.nlm.nih.gov/articles/PMC4596846/ [Accessed 5 Nov 2024].

step 6: Testing all the model after features

engineering (baseline)

As we have now implement features engineering, it will be beneficial to did an other baseline test to have a better understanding dose the features we create bring positive or negative impact to the model learning

```
X train point7, X test point7, y trair
engineered results = []
for model name, model in models.item
        for dataset name, (X train, )
                model.fit(X train, y tr
               y pred = model.predict
               accuracy = accuracy so
                engineered results.apper
                        "Model": model
                        "Variant": data
                        "Outlier Handli
                        "Imputation Met
                        "Feature Engine
                        "Accuracy": acc
                })
engineered results df = pd. DataFrame (ε
engineered results df = engineered res
pd. set option ('display. max rows', 1000)
pd. set option ('display. max columns', No
pd. set option ('display. width', 1000)
print (engineered results df)
```

step 7: Hyperparameter Tuning For model

```
print("Imputed Variants:")
print(imputed_variants.keys())

print("Feature-Engineered Variants:")
print(feature_engineered_variants.keys()
```

```
print("Holdout Imputed Variants:")
print(imputed_variants_holdout.keys())
print("Holdout Feature-Engineered Vari
print(holdout_feature_engineered_variant
```

Hyperparameter Tuning for XGBoost (kaggle 0.8334)

```
from sklearn.model selection import F
best variant name = 'Winsorization knr
X train best, X test best, y train bes
param grid = {
       'learning rate': [0.01, 0.05,
       'max depth': [3, 5, 7, 9, 1
       'n_estimators': [50, 100, 200
       'subsample': [0.5, 0.7, 0.9,
       'colsample bytree': [0.5, 0.7,
       'colsample bylevel': [0.5, 0.7
       'colsample bynode': [0.5, 0.7,
       'min child weight': [1, 3, 5,
       'gamma': [0, 0.1, 0.3, 0.5,
       'reg_lambda': [0.5, 1, 1.5],
       'reg alpha': [0, 0.5, 1, 1.5
       'booster': ['gbtree', 'dart'],
       'tree method': ['auto', 'exact
xgb = XGBClassifier(use label encoder=
grid search = RandomizedSearchCV(
       xgb, param_grid, n_iter=50, c
grid_search.fit(X_train_best, y_train_k
print("Best Parameters:", grid search.
print("Best Accuracy from Grid Searc
```

```
best_variant_name = 'Winsorization_Ite

X_train_best, X_test_best, y_train_bes

param_grid = {

    'n_estimators': [50, 100, 200

    'learning_rate': [0.05, 0.1,

    'max_depth': [3, 5, 7],

    'subsample': [0.5, 0.6, 1.0],

    'colsample_bytree': [0.5, 0.7,

    'gamma': [0, 0.1, 0.3, 0.5,

    'min_child_weight': [1, 3, 5,

    'reg_alpha': [0, 0.1, 1],

    'reg_lambda': [0.5, 1, 2, 5]
```

```
xgb = XGBClassifier(use_label_encoder=
grid_search = GridSearchCV(xgb, param
grid_search.fit(X_train_best, y_train_t)
print("Best Parameters:", grid_search.
print("Best Accuracy from Grid Searc
best_model = grid_search.best_estimate
y_pred_best = best_model.predict(X_tes
accuracy_best = accuracy_score(y_test_
print(f"Test Accuracy for Best Model
```

RandomizedSearchCV first than GridSearchCV to safe time as there will be less to try on and close down the candidates

```
best variant name = 'Winsorization knr
X_train, X_test, y_train, y_test =
print(f"Running model for variant:
all accuracies = []
num runs = 10
for i in range (num runs):
       X train best, X test best, y t
               X train, y train, test
       print(f"Features before fitting
       model = XGBClassifier(
               learning rate=0.05,
               max depth=5,
               n estimators=100,
               subsample=0.5,
               eval metric='logloss',
               reg lambda=0.5,
               reg alpha=1,
               min_child_weight=5,
               gamma=0.1,
               colsample bytree=0.9,
               random state=i,
       model.fit(X train best, y train
       y_pred = model.predict(X test
       accuracy = accuracy score(y te
       all accuracies. append (accuracy)
       scores = cross validate(
```

```
model, X_train_best, ,,
)

print(f"Run {i + 1}:")
print(f"Accuracy (Testing): {a
print(f"Accuracy (CV Mean): +

conf_matrix = confusion_matrix(y_test_
sns.heatmap(conf_matrix, annot=True, f
plt.title('Confusion Matrix')
plt.show()

print("\nSummary of accuracies across
print(f"Mean accuracy over {num_runs})

print(classification_report(y_test_best,
```

training to test on the robustness of the process we are getting 85% with almost the same CV mean which means it is generalising well

```
best_xgb = XGBClassifier(
    learning_rate=0.05,
    max_depth=5,
    n_estimators=100,
    subsample=0.5,
    eval_metric='logloss',
    reg_lambda=0.5,
    reg_alpha=1,
    min_child_weight=5,
    gamma=0.1,
    colsample_bytree=0.9,
)
```

model output for more data understanding later

XGBoost play ground

```
def my_plot_importance(booster, figsiz plt.rcParams["figure.figsize"] plot_importance(booster=booster)
my_plot_importance(best_xgb, figsize=(1))
import shap
```

```
\begin{tabular}{lll} explainer &=& shap.TreeExplainer(best\_x \begin{tabular}{lll} explainer.shap\_values(X \begin{tabular}{lll} shap\_values, & X\_test\_t \end{tabular} \end{tabular}
```

```
from sklearn.inspection import permut

result = permutation_importance(best_x)
for i in result.importances_mean.args
    print(f"{X_test_best.columns[i]})
```

up to now we can see which features are more imporatnat which are not, it will allow us to do features selection, base on the infrmation above

```
 barplot=(sns.\,countplot\,(data=\,\,data\_imbal\\ plt.\,title('0\,\,v/s\,\,1\n')
```

as shown above there is heavy data imblance and there is ouliers with in the engineered features, to move forword for better XGBoost performacne we will implnemnt 3 different ways for outliners and ways uses SMOTE for class imblance than perform feature selection and compaire there perfomance together

```
X_train, X_test, y_train, y_test =

X_train_point7_iqr = IQR(X_train)
X_test_point7_iqr = IQR(X_test)

y_train_point7_iqr = y_train.loc[X_train)
y_test_point7_iqr = y_test.loc[X_test_incollabel]
```

```
point7_df_iqr = pd.concat([X_train_poi
for name, group in point7_df_iqr.grc
```

```
plt.plot(group.BAB, group.HB ir
plt.legend()
X train point7 z score = z score(X tra
X test point7 z score = z score(X test
y_train = y_train.reset_index(drop=Tru
y test = y test.reset index(drop=True)
y train point7 z score = y train.loc[}
y_test_point7_z_score = y_test.loc[X_t
point7_df_z_score = pd.concat([X_train
for name, group in point7_df_z_score
       plt.plot(group.BAB, group.HB ir
plt.legend()
X train point7 isolation forest = isol
X_test_point7_isolation_forest = isola
y train = y train.reset index(drop=Tru
y test = y test.reset index(drop=True)
y train point7 isolation forest = y tr
y_test_point7_isolation_forest = y_tes
point7 df isolation forest = pd. concat
for name, group in point7 df isolati
```

plt.plot(group.BAB, group.HB ir plt.legend()

do the same for hold out

now we have done all 3 ways that we have talked about for daeling with the ouliners, and each of them have perfrom abit different which it will be provide a good variants on the outcome

```
from imblearn.over sampling import SN
from imblearn.combine import SMOTETon
from imblearn.over_sampling import SN
from imblearn.combine import SMOTETon
point7 datasets = {
       'point7 IQR': (X train point7 i
```

```
'point7 Zscore': (X train point
       'point7 isolationforest': (X tr
XGBoost outliers variants = {}
for dataset name, (X train, X test,
       smote = SMOTE()
       X_train_resampled, y_train_resa
       X train imputed = X train resε
       X test imputed = X test
       key = f"{dataset name} SMOTE"
       XGBoost outliers variants[key]
for dataset_name, (X_train, X_test,
       smote = BorderlineSMOTE()
       X_train_resampled, y_train_resa
       X train imputed = X train resa
       X test imputed = X test
       key = f"{dataset name} Border1
       XGBoost outliers variants[key]
for dataset_name, (X_train, X_test,
       smote = SVMSMOTE()
       X_train_resampled, y_train_resa
       X train imputed = X train resa
       X test imputed = X test
       key = f"{dataset_name}_SVMSMOT
       XGBoost outliers variants[key]
for dataset_name, (X_train, X_test,
       smote = RandomOverSampler(rand)
       X_train_resampled, y_train_resa
       X train imputed = X train resa
       X_{test_imputed} = X_{test_i}
       key = f"{dataset_name}_RandomS
       XGBoost outliers variants[key]
for dataset_name, (X_train, X_test,
       smote = SMOTETomek(random stat
       X_train_resampled, y_train_resa
       X train imputed = X train resa
       X test imputed = X test
```

```
key = f"{dataset_name}_SMOTETC
XGBoost_outliers_variants[key]

print(XGBoost_outliers_variants.keys())

X_train_point7_IQR_SMOTE, X_test_point7

train_data_point7_IQR_SMOTE = pd.concat
test_data_point7_IQR_SMOTE = pd.concat

barplot=(sns.countplot(data= train_data
barplot=(sns.countplot(data= train_data
plt.title('0 v/s 1\n')
```

as shown the smote is applied only to the test set to avoid data leakage

```
point7_smote_engineered_results df = [
for model name, model in models.item
        for dataset name, (X train, )
               model.fit(X train, y tr
               v pred = model.predict
               accuracy = accuracy_sc
               point7 smote engineered
                       "Model": model_
                       "Variant": data
                       "Feature Engine
                        "Imputation Met
                       "Smote Method":
                       "Accuracy": acc
               })
point7_smote_engineered_df = pd.DataFr
point7 smote engineered results df = r
pd. set option ('display. max rows', 1000)
pd. set option ('display. max columns', No
pd. set option ('display. width', 1000)
print(point7_smote_engineered_results_df
```

for now the Accuracy seems like the same as before but we should try on XGBoost to get more inforemation about its performacne

```
from sklearn.metrics import classific
best variant name = 'point7 Zscore SMC
X_train, X_test, y_train, y_test =
print(f"Running model for variant:
all accuracies = []
num runs = 10
for i in range (num runs):
       X train best, X test best, y t
              X_train, y_train, test
       print(f"Features before fitting
       model = XGBClassifier(
               learning rate=0.05,
               max depth=5,
               n estimators=100,
               subsample=0.5,
               eval metric='logloss',
               reg lambda=0.5,
               reg alpha=1,
               min child weight=5,
               gamma=0.1,
               colsample bytree=0.9,'
               random state=43,
       model.fit(X train best, y train
       y_pred = model.predict(X_test_
       accuracy = accuracy score(y te
       all accuracies. append (accuracy)
        scores = cross_validate(
              model, X train best, y
       )
        print(f"Run {i + 1}:")
        print(f"Accuracy (Testing): {ε
        print(f"Accuracy (CV Mean):
conf matrix = confusion matrix(y test
sns.heatmap(conf matrix, annot=True, f
plt.title('Confusion Matrix')
plt.show()
print("\nSummary of accuracies across
print(f"Mean accuracy over {num_runs}
print(classification_report(y_test_best,
```

we can see our performance have largely increased on different areas

with out features selection, now we will move onto features selection.

```
smote_best_xgb = XGBClassifier(
    learning_rate=0.05,
    max_depth=5,
    n_estimators=100,
    subsample=0.5,
    eval_metric='logloss',
    reg_lambda=0.5,
    reg_alpha=1,
    min_child_weight=5,
    gamma=0.1,
    colsample_bytree=0.9,
)

smote_best_xgb.fit(X_train_best, y_train_best)

explainer = shap.TreeExplainer(smote_t shap_values = explainer.shap_values(X_shap.summary_plot(shap_values, X_test_t)
```

we can see that even after we done isoforest and smotie, it stay the same as before because (look at i can talk about for smote)

```
def my_plot_importance(booster, figsiz
plt.rcParams["figure.figsize"]
plot_importance(booster=booster)
my_plot_importance(smote_best_xgb, figs
```

```
feature indices = [list(model.
       X train selected = X train.ilc
       X test selected = X test.iloc[
       key = f"{dataset_name}_importε
       XGBoost outliers variants featur
for dataset name, (X train, X test,
       model = XGBClassifier()
       model.fit(X train, y train)
       importance scores = model.get
       importance df = pd.DataFrame(1
       selected features = importance
       feature indices = [list(model.
       X train filtered = X train.ilc
       X test filtered = X test.iloc[
       estimator = XGBClassifier()
       rfe = RFE(estimator, n featur
       rfe.fit(X_train_filtered, y_tra
       rfe selected features mask = >
       X train rfe = X train filtered
       X_test_rfe = X_test_filtered.1
       key = f"{dataset name} rfeAftε
       XGBoost_outliers_variants_featur
for dataset name, (X train, X test,
       estimator = XGBClassifier()
       sfs = SequentialFeatureSelecto
       sfs.fit(X train, y train)
       X_train_sfs = X_train.loc[:,
       X test sfs = X test.loc[:, sf
       key = f"{dataset_name}_sfsforw
       XGBoost outliers variants featur
for dataset_name, (X_train, X_test,
       estimator = XGBClassifier(rand
       rfecv = RFECV(
               estimator=estimator,
               step=1,
               cv=cv,
               scoring='accuracy',
               min_features_to_select=1
       )
```

rfecv.fit(X_train, y_train)

```
optimal_feature_count = rfecv.
feature_ranking = rfecv.rankir
total_mean_score = np.mean(rfe

X_train_rfecv = X_train.loc[:,
X_test_rfecv = X_test.loc[:,

key = f"{dataset_name}_rfecv"
XGBoost_outliers_variants_featur

print(f"Dataset: {dataset_name})
print(f"Optimal number of featur)
print(f"Total mean score: {tc

print(XGBoost_outliers_variants_features)
print(timportance_df)
```

print(imputed_variants.keys())
print(feature_engineered_variants.keys())
print(XGBoost_outliers_variants.keys())
print(XGBoost_outliers_variants_features_variants_featur

```
X_train_test, X_test_test, y_train_test
data_check = pd.concat([X_train_test,
data_check.describe().T
```

X_train_test, X_test_test, y_train_tes
X_train_test = pd.DataFrame(X_train_te
y_train_test = y_train_test.reset_inde
data_check = pd.concat([X_train_test,
data_check.describe().T

```
"Imputation Met
"Smote Method":
"Features Selec
"Accuracy": acc
})

model_results_df = pd.DataFrame(model_
model_results_df = model_results_df.sc

pd.set_option('display.max_rows', 10000
pd.set_option('display.max_columns', Nc
pd.set_option('display.width', 10000)
print(model_results_df)
```

point7_Zscore_SMOTETomek_importa nceScore 88%

point7_IQR_SMOTETomek_importanc eScore and 88%

point7_IQR_SMOTETomek_rfecv 89% (have the best for now overall)

point7_IQR_SMOTETomek_rfeAfterIm portanceFiltered 89%

```
best_variant_name = 'point7_Zscore_SMC
X train, X test, y train, y test =
print(f"Running model for variant:
all accuracies = []
num_runs = 10
for i in range (num runs):
       X_train_best, X_test_best, y_t
               X train, y train, test
       print(f"Features before fittir
       model = XGBClassifier(
               learning rate=0.05,
               max_depth=11,
               n estimators=200,
               subsample=0.5,
               eval_metric='logloss',
               reg_lambda=1.5,
               reg_alpha=0.5,
               min_child_weight=10,
               gamma=0.5,
               colsample bytree=1,
               colsample bynode=1,
               colsample bylevel=0.5,
```

```
)
       model.fit(X train best, v train
       y pred = model.predict(X test
       accuracy = accuracy score(v te
       all accuracies. append (accuracy)
       scores = cross validate(
               model, X train best, y
       print(f"Run {i + 1}:")
       print(f"Accuracy (Testing): {ε
       print(f"Accuracy (CV Mean):
conf matrix = confusion matrix(y test
sns.heatmap(conf matrix, annot=True, f
plt.title('Confusion Matrix')
plt. show()
print("\nSummary of accuracies across
print(f"Mean accuracy over {num runs}
print(classification report(y test best,
all runs results = []
num runs = 10
       all accuracies = []
```

booster='gbtree',
random state=42,

```
for dataset name, (X train, X test, y train, y test) in XGBoost outliers variants featur
       for i in range (num runs):
               X_train_best, X_test_best, y_train_best, y_test_best = train_test_split(
                       X train, y train, test size=0.2, random state=i
               print(f"Features before fitting (run {i + 1}): {X train best.columns}")
               print(f"Running model for variant: {dataset name}")
               model = XGBClassifier(
                       learning rate=0.05,
                       max depth=10,
                       n estimators=200,
                       subsample=0.8,
                       objective='reg:squarederror',
                       reg_lambda=1.5,
                       reg alpha=1.5,
                       min child weight=10,
                       gamma=0.5,
                       colsample bytree=1,
                       colsample bynode=1,
                       colsample bylevel=0.5,
                       booster='gbtree'.
```

```
random state=i
               model.fit(X train best, y train best)
               y pred = model.predict(X test best)
               accuracy = accuracy score(y test best, y pred)
               all accuracies, append (accuracy)
               scores = cross validate(
                      model, X_train_best, y_train_best, cv=5, return_train_score=True,
               print(f'' \setminus nRun \{i + 1\}: ")
               print(f"Accuracy (Testing): {accuracy:.2f}")
               print(f"Accuracy (CV Mean): {np.mean(scores['test score']):.2f} (+/- {np.s
       mean accuracy = np. mean(all accuracies)
       std accuracy = np. std(all accuracies)
       all runs results.append({
               "Dataset": dataset name,
               "Mean Accuracy": mean accuracy,
               "Std Accuracy": std accuracy,
               "Details": X train.columns.tolist()
       })
print("\nSummary of accuracies across runs:")
for result in all runs results:
       print(f"Dataset: {result['Dataset']}, Mean Accuracy: {result['Mean Accuracy']:.2f}
results df = pd. DataFrame(all runs results)
sorted results df = results df.sort values(by="Mean Accuracy", ascending=False)
print("\nSorted Results by Accuracy:")
print(sorted results df)
import ace tools as tools
tools.display_dataframe_to_user(name="Sorted Model Results by Accuracy", dataframe=sorted_
```

output model and safe work state

```
best_variant_name = 'point7_Zscore_SM(
XGB_finial_X_train, X_test, XGB_finial

best_XGB_After_proccess = XGBClassifi&
    learning_rate=0.05,
    max_depth=11,
    n_estimators=200,
    subsample=0.5,
    eval_metric='logloss',
    reg_lambda=1.5,
    reg_alpha=0.5,
    min_child_weight=10,
    gamma=0.5,
    colsample_bytree=1,
    colsample_bytroe=1,
```

```
colsample_bylevel=0.5,
booster='gbtree',
random_state=42,
)

best_XGB_After_proccess.fit(XGB_finial_)

import joblib
joblib.dump(best_XGB_After_proccess, 't

best_XGB_After_process = joblib.load('
y_pred = best_XGB_After_process.predic

y_pred = best_XGB_After_process.predic

y_pred

joblib.dump(best_XGB_After_process, 'be

print(imputed_variants.keys())
print(feature_engineered_variants.keys())
print(XGBoost_outliers_variants_features)
```

Hyperparameter Tuning forGradient Boosting (kaggle 0.8505)

```
from skopt import BayesSearchCV
best variant name = 'point7 IQR SMOTE1
X train, X test, y train, y test =
param_space = {
       'n estimators': (50, 100, 200
       'learning_rate': (0.01, 0.05,
       'max depth': (3, 5, 7),
       'min_samples_split': (2, 5, 1
       'min_samples_leaf': (1, 2, 5,
       'max_features': ['sqrt', 'log2
       'subsample': (0.7, 0.8, 1.0),
       'loss': ['log_loss', 'exponent
       'min impurity decrease': (0.001
       'warm start': [True, False],
       'max_leaf_nodes': [None, 10,
       'n iter no change': [None, 5,
       'tol': (0.0001, 0.001, 0.01)
model = GradientBoostingClassifier(rar
```

```
bayes_opt = BayesSearchCV(
    estimator=model,
    search_spaces=param_space,
    n_iter=50,
    scoring='accuracy',
    n_jobs=-1,
    cv=3,
    verbose=1,
    random_state=42
)

bayes_opt.fit(X_train, y_train)

print("Best Parameters: ", bayes_opt.
print("Best Accuracy from Grid Searc
```

```
best_variant_name = 'point7_IQR_SMOTET
X_train, X_test, y_train, y_test =

param_grid = {
        'n_estimators': [50, 100, 200
        'learning_rate': [0.01, 0.05,
        'max_depth': [3, 5, 7],
        'min_samples_split': [2, 5, 1
        'min_samples_leaf': [1, 2, 5,
        'max_features': ['sqrt', 'log2
        'subsample': [0.7, 0.8, 1.0]
}

gb_model = GradientBoostingClassifier(
grid_search = GridSearchCV(estimator=grid_search = GridSearch = Gri
```

```
all_runs_results = []

num_runs = 10

for dataset_name, (X_train, X_test,

all_accuracies = []

for i in range(num_runs):

X_train_best, X_test_be
X_train, y_trai
)
```

```
print(f"Features before
                print(f"Running model
                model = GradientBoosti
                       learning rate=0.
                       max depth=7,
                        n estimators=50,
                        subsample=1.0,
                        max features='1c
                        min samples leaf
                        min samples spli
                        random state=42,
                        warm start=False
                        to1=0.001.
                        min impurity dec
                        max leaf nodes=N
                       loss='exponentia
                        n iter no change
                model.fit(X train best,
                y_pred = model.predict
                accuracy = accuracy sc
                all accuracies. append (ac
                scores = cross validat
                       model, X_train_
                print(f'' \setminus nRun \{i + 1\}
                print(f"Accuracy (Testi
               print(f"Accuracy (CV N
        mean accuracy = np. mean (all ac
        std_accuracy = np. std(all_accu
        all runs results.append({
               "Dataset": dataset name
                "Mean Accuracy": mean_
                "Std Accuracy": std ac
                "Details": X train.colu
       })
print("\nSummary of accuracies across
for result in all runs results:
        print(f"Dataset: {result['Datas
results df = pd. DataFrame (all runs res
sorted results df = results df. sort νε
print("\nSorted Results by Accuracy:"
print(sorted results df)
import ace tools as tools
tools.display dataframe to user(name="Sc
```

best variant name = 'point7 Zscore SMOTETomek rfecv'

```
X_train, X_test, y_train, y_test = XGBoost_outliers_variants features selected[best varian
print(f"Running model for variant: {best variant name}")
all accuracies = []
num runs = 10
for i in range(num_runs):
       X_train_best, X_test_best, y_train_best, y_test_best = train_test_split(
               X train, y train, test size=0.2, random state=i
       print(f"Features before fitting (run {i + 1}): {X train best.columns}")
       model = GradientBoostingClassifier(
              learning rate=0.2,
               max depth=7,
               n estimators=50,
               subsample=1.0,
               max features='log2',
               min samples leaf=1,
               min samples split=10,
               random state=42,
               warm start=False,
               to1=0.001,
               min impurity decrease=0.001,
               max leaf nodes=None,
               loss='exponential',
               n iter no change=None
       )
       model.fit(X_train_best, y_train_best)
       y pred = model.predict(X test best)
       accuracy = accuracy score(y test best, y pred)
       all accuracies. append (accuracy)
       scores = cross validate(
               model, X_train_best, y_train_best, cv=5, return_train_score=True, return_e
       print(f"Run {i + 1}:")
       print(f"Accuracy (Testing): {accuracy:.2f}")
       print(f"Accuracy (CV Mean): {np.mean(scores['test score']):.2f} (+/- {np.std(score
conf_matrix = confusion_matrix(y_test_best, y_pred)
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues')
plt.title('Confusion Matrix')
plt. show()
print("\nSummary of accuracies across runs:")
print(f"Mean accuracy over {num runs} | runs: {np.mean(all accuracies):.2f} (+/- {np.std(a
print(classification_report(y_test_best, y_pred))
best variant name = 'point7 Zscore SMC
GB finial X train, X test, GB finial y
```

```
best GB After process = GradientBoost
               learning rate=0.2,
                max depth=7,
               n estimators=50,
                subsample=1.0,
                max_features='log2',
                min samples leaf=1,
                min samples split=10,
                random state=42,
                warm start=False,
                to1=0.001,
                min impurity decrease=0.
                max leaf nodes=None,
                loss='exponential',
               n iter no change=None
best_GB_After_proccess.fit(GB_finial_X_t
joblib. dump (best GB After proccess, 'bε
best XGB After process = joblib.load('
y_pred = best_XGB_After_process.predic
y_pred = best_XGB_After_process.predic
y_pred
y pred = best gb model.predict(X test)
y_pred
```

Gradient Boosting play ground

```
X = p7_point_footprints_df_iqr_knn_bla
y = p7_point_footprints_df_iqr_knn_bla
x = 0
count = 0
num_runs = 1

for x in range (num_runs):
    count += 1
    model = GradientBoostingClassi
SMOTE_iso_X_train, SMOTE_iso_X_
```

```
model.fit(SMOTE_iso_X_train, SN_SMOTE_iso_y_pred = model.predi

scores = cross_validate(model, X, y, precision = precision_score(SMOTE_iso_recall = recall_score(SMOTE_iso_y_test)

print(metrics.confusion_matrix(SMOTE_iso_print("\nAccuracy (Testing): %0.2f
print("Accuracy (Testing): %0.2f (print("count:", count))
print("Precision: %.2f" % precision)
print("recall: %.2f" % recall)

from sklearn.metrics import confusion print(confusion_matrix(SMOTE_iso_y_test, sns.heatmap(confusion_matrix(SMOTE_iso_y.
```

Hyperparameter Tuning for Support Vector Machines

SVM overall will do better after StandardScaler there for we will use it to improve SVM score

```
from sklearn.preprocessing import Sta
from sklearn.decomposition import PCF
from sklearn.svm import SVC
from skopt import BayesSearchCV
best_variant_name = 'point7_IQR_SMOTE1
X_train, X_test, y_train, y_test =
scaler = StandardScaler()
X_train_scaled = scaler.fit_transform(X_test_scaled = scaler.transform(X_test_pca = PCA(n_components=0.95)
X_train_pca = pca.fit_transform(X_train_pca = pca.fit_transform(X_train
```

```
X test pca = pca.transform(X test scal
param grid = {
       'C': (0.1, 1000, 'log-uniform
        'gamma': (0.001, 10, 'log-uni
       'kernel': ['rbf'],
        'tol': (1e-4, 1e-2, 'log-unif
        'max iter': (1000, 10000),
        'class_weight': [None, 'balanc
mode1 = SVC(random state=42)
bayes opt = BayesSearchCV(
       estimator=model,
       search spaces=param grid,
       n iter=50,
       scoring='accuracy',
       n iobs=-1,
       cv=3,
       verbose=1,
       random state=42
bayes opt. fit (X train pca, y train)
print ("Best Parameters:", bayes opt. be
print ("Best Score:", bayes opt. best so
```

按兩下 (或按 Enter 鍵) 即可編輯

```
X_train_test, X_test_test, y_train_tes

X_train_test = pd.DataFrame(X_train_te

y_train_test = y_train_test.reset_inde
```

```
X_train_test, X_test_test, y_train_tes
X_train_test = pd.DataFrame(X_train_te)
y_train_test = y_train_test.reset_inde
data_check = pd.concat([X_train_test,
data_check.describe().T
```

```
all runs results = []
num runs = 10
best params = {
       'C': 1.1930801848463657,
       'class weight': 'balanced',
       'gamma': 0.120601154417892,
       'kernel': 'rbf',
        'max iter': 1000,
       'tol': 0.0001
for dataset_name, (X_train, X_test,
       all accuracies = []
       for i in range(num_runs):
               X_train_best, X_test_be
                       X_train, y_trai
               print(f"Features before
               print(f"Running model
               mode1 = SVC(
                       C=best_params['(
                       kernel=best para
                       gamma=best param
                       class_weight=bes
                       max iter=best pε
                       tol=best params[
                       random_state=42
               )
               model.fit(X_train_best,
               y_pred = model.predict
               accuracy = accuracy so
               all accuracies. append (ac
```

```
scores = cross validat
                       model, X train
               print(f'' \setminus nRun \{i + 1\}
               print(f"Accuracy (Testi
               print(f"Accuracy (CV N
        mean accuracy = np. mean (all ac
        std accuracy = np. std(all accu
        all runs results, append({
               "Dataset": dataset name
               "Mean Accuracy": mean
               "Std Accuracy": std ac
               "Details": X train.colu
       })
print("\nSummary of accuracies across
for result in all runs results:
       print(f"Dataset: {result['Datas
results df = pd. DataFrame (all runs res
sorted results df = results df.sort va
print("\nSorted Results by Accuracy:"
print(sorted results df)
tools. display dataframe to user (name="Sc
```

```
best variant name = 'point7 Zscore SMC
X train, X test, y train, y test =
print(f"Running model for variant:
all accuracies = []
num runs = 10
for i in range (num runs):
       X_train_best, X_test_best, y_t
              X train, y train, test
       print(f"Features before fittir
       mode1 = SVC(
              C =1.9650743261576813,
              class weight = 'balanc
               gamma = 0.090070545592
              kernel = 'rbf',
               max iter = 4979,
               to1 = 0.01,
               random state=42
       model.fit(X train best, y train
       y pred = model.predict(X test
       accuracy = accuracy_score(y_te
```

Support Vector Machines play ground

```
best_variant_name = 'point7_Zscore_SMC
SCV finial X train, X test, SCV finial
best SVC After process = SVC(
               C =1.9650743261576813,
               class weight = 'balanc
               gamma = 0.090070545592
               kernel = 'rbf',
               max iter = 4979,
               to1 = 0.01,
               random state=42,
               probability=True
best SVC After process.fit(SCV finial )
joblib.dump(best_SVC_After_proccess, 't
best_SVC_After_proccess = joblib.load(
y pred = best SVC After process.predi
print(y pred)
```

Hyperparameter Tuning for Random Forest

```
best_variant_name = 'point7_Zscore_SMC
X_train, X_test, y_train, y_test =
```

```
param grid = {
        'n estimators': [50, 100, 200
        'max depth': [5, 10, 15, 20,
       'min samples split': [2, 5, 1
        'min samples leaf': [1, 2, 5,
        'max features': ['sqrt', 'log2
        'bootstrap': [True],
        'max leaf nodes': [10, 20, 50
        'min impurity_decrease': [0.0,
        'criterion': ['gini', 'entropy
        'class_weight': [None, 'balanc
        'oob score': [True, False]
model = RandomForestClassifier(random
bayes opt = BayesSearchCV(
       estimator=model.
       search spaces=param grid,
       n iter=50,
       scoring='accuracy',
       n jobs=-1,
       cv=3.
       verbose=1,
       random state=42
bayes_opt.fit(X_train_pca, y_train)
print ("Best Parameters:", bayes opt. be
print ("Best Score:", bayes_opt.best_sc
```

```
best variant name = 'point7 Zscore SMC
X_{train}, X_{test}, y_{train}, y_{test} =
print(f"Running model for variant:
all accuracies = []
num runs = 10
for i in range (num runs):
       X train best, X test best, y t
               X train, y train, test
       print(f"Features before fitting
       model = RandomForestClassifier
               class weight =None,
               criterion = 'entropy',
               max depth = None,
               max features = None,
               max leaf nodes = 100,
               min impurity decrease =
               min samples leaf = 2,
               min_samples_split = 2,
               n_estimators = 200,
```

```
oob score = False,
               random state=42
       model.fit(X train best, y train
       y pred = model.predict(X test
       accuracy = accuracy score(y te
       all accuracies. append (accuracy)
       scores = cross validate(
              model, X train best, y
       print(f"Run {i + 1}:")
       print(f"Accuracy (Testing): {ε
       print(f"Accuracy (CV Mean):
conf matrix = confusion matrix(y test
sns.heatmap(conf matrix, annot=True, f
plt.title('Confusion Matrix')
plt. show()
print("\nSummary of accuracies across
print(f"Mean accuracy over {num runs}
print(classification report(y test best,
```

step 8: Ensemble Learning

```
XGB_model = XGBClassifier(best_XGB_Aft
GBC_model = GradientBoostingClassifier
SVM_model = SVC(best_SVC_After_procces
RF_model = RandomForestClassifier(best
```

```
from sklearn.ensemble import VotingCl
from sklearn.metrics import accuracy
voting clf = VotingClassifier(
       estimators=[
               ('xgb', best XGB After
               ('gbc', best GB After p
               ('svm', best_SVC_After_
               ('rf', best RF After pr
       1.
       voting='soft'
voting_clf.fit(X_train, y_train)
y pred = voting clf.predict(X test)
print ("Ensemble Model Accuracy:", acc
print ("Confusion Matrix:\n", confusior
print("Classification Report:\n", clas
print(imputed variants.keys())
print (feature engineered variants. keys ()
print(XGBoost_outliers_variants.keys())
print (XGBoost outliers features selected
print (XGBoost outliers variants features
XGBoost outliers variants = {'point7 1
from sklearn.ensemble import Stacking
from sklearn.linear model import Logi
stacking clf = StackingClassifier(
       estimators=[
               ('xgb', best_XGB_After_
               ('gbc', best GB After p
               ('svm', best_SVC_After_
               ('rf', best_RF_After_pr
       final estimator=LogisticRegressi
stacking clf.fit(X train, y train)
y_pred_stack = stacking_clf.predict(X_
print("Stacking Ensemble Accuracy:",
print ("Confusion Matrix:\n", confusior
print("Classification Report:\n", clas
X train.describe().T
from sklearn.ensemble import VotingCl
ensemble_model = VotingClassifier(
       estimators=[
```

hold_out set change

```
hold out = pd. read csv('SexLandmarks-t
hold out data df = hold out.copy()
hold_out_data_df = IQR(hold_out_data_c
hold out scaled data = hold out.copy()
for column in hold out scaled data.cc
       if column. startswith ('x'):
               hold out scaled data[col
        elif column.startswith('y'):
               hold out scaled data[col
print(hold out scaled data.head())
lengths_upper_threshold = hold_out_sca
lengths lower threshold = hold out sca
widths upper threshold = hold out scal
widths_lower_threshold = hold_out_scal
big feet = hold out scaled data with 1
        (hold_out_scaled_data_with_lengt
        (hold out scaled data with lengt
small feet = hold out scaled data with
        (hold_out_scaled_data_with_lengt
        (hold out scaled data with lengt
print("Big Feet Data Points:")
print(big feet)
print("\nSmall Feet Data Points:")
print(small feet)
```

import matplotlib.pyplot as plt

```
plt.figure(figsize=(10, 6))

plt.scatter(hold_out_scaled_data_with_le

plt.scatter(big_feet['lengths'], big_fe

plt.scatter(small_feet['lengths'], smal

plt.xlabel('Lengths')

plt.ylabel('Widths')

plt.legend()

plt.show()
```

```
holdout_feature_engineered_variants =
lengths_widths_temp_dict = {}
for variant_name, hold_out_data in i
hold_out_lengths = lengths_wic
key = f"{variant_name}_lengths
lengths_widths_temp_dict[key] =
holdout_feature_engineered_variants.upda
```

```
point7 temp dict = {}
for variant name, hold out data in i
       hold out point7 = point7 calcu
       key = f"{variant name} point7"
       point7 temp dict[key] = hold c
holdout feature engineered variants.upda
print (holdout feature engineered variant
print(holdout_feature_engineered_variant
print(imputed variants holdout.keys())
hold out XGBoost outliers variants =
hold_out_data = holdout_feature_engine
hold out point7 datasets = {
        'point7_IQR': (IQR(Winsorization
       'point7_Zscore': (z_score(Winsc
       'point7 isolationforest': (isol
print(hold_out_point7_datasets.keys())
if 'point7 Zscore' in hold out point7
       display(hold out point7 datasets
else:
       print("The key 'point7 Zscore'
if 'point7_IQR' in hold_out_point7_da
       display(hold out point7 datasets
else:
        print ("The key 'point7_IQR' c
if 'point7 isolationforest' in hold of
       display(hold out point7 datasets
else:
       print("The key 'point7 isolati
```

```
selected features = [ #it its from
       'x0', 'y0', 'x1', 'y2', 'x3'
        'x7', 'y7', 'x8', 'y8', 'x10
        'x13', 'y13', 'y14', 'x17',
       'BAH', 'HB index'
]
hold out data filtered unscaled = hold
hold out data filtered unscaled.describe
Used in model = 'point7 Zscore SMOTETo
x train scale, x test scale, y train s
scaler = StandardScaler()
scaler.fit(x test scale)
hold out data filtered unscaled = hold
hold_out_data_filtered_unscaled = hold
try:
       hold out data filtered scaled =
               scaler.transform(hold ou
               columns=hold_out_data_fi
        print("Scaling successful.")
except ValueError as e:
        print("Error during scaling:",
print (hold out data filtered unscaled. hε
print (hold out data filtered scaled head
print("Mean used by scaler: ", scal
print("Scale used by scaler: '
scaler = StandardScaler()
scaler.fit(X train best)
hold_out_data_filtered_scaled = pd.Dat
        scaler.transform(hold out data f
        columns=hold_out_data_filtered_u
print(hold_out_data_filtered_scaled.desc
print (hold out data filtered unscaled. he
print (hold out data filtered scaled head
```

Try to submitting it to kaggle

```
Used in model = 'point7 Zscore SMOTETc
x_train_submit, x_test_submit, y_train
print(x train submit.shape)
scaler = StandardScaler()
scaler.fit(x train submit)
x train submit scaled = pd. DataFrame(s
x train submit_unscaled = x_train_subm
print(x train submit.shape)
print(y train submit.shape)
y_train_submit = pd. Series(y_train_sub
y train submit.reset index(drop=True, i
y train submit = y train submit.values
best_XGB_After_proccess.fit(x_train_subm
best GB After process.fit(x train submi
best_RF_After_proccess.fit(x_train_submi
best_SVC_After_proccess.fit(x_train_subm
```

best_ensemble_Voting_model submit

```
RowID = np.array(hold_out_data_filtere

results = pd.DataFrame({'RowID': RowI

print(results)

results.to_csv('results.csv', index=Fal

'''!kaggle competitions submit -c bu
```

best GB After proccess submit

best_XGB_After_proccess submit

```
best_XGB_After_proccess.fit(x_train_subm

gb_pred = best_XGB_After_proccess.prec

results = pd.DataFrame({
    'RowID': np.array(hold_out_data
    'Sex': gb_pred
})

results.to_csv('best_XGB_After_proccess.
print(results.head)

results.to_csv('best_XGB_After_proccess.
```

best RF After proccess submit

```
best_RF_After_proccess.fit(x_train_submi)

gb_pred = best_RF_After_proccess.predi)

results = pd.DataFrame({
    'RowID': np.array(hold_out_data'
    'Sex': gb_pred
})

results.to_csv('best_RF_After_proccess.comprint(results.head)
```

best SVC After proccess submit

```
best_SVC_After_proccess.fit(x_train_subm
gb_pred = best_SVC_After_proccess.pred
```

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