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OVERVIEW

- 1. Project description
- 2. Data preprocessing
- 3. Data visualization
- 4. Building models
- 5. performance comparison
- 6. further iprovement.





- Objective: To leverage Classification Algorithm to predict Diabetes in female patients.
- Datasource: Prima Indians Database by National Institute of Diabetes and Digestive and Kidney Diseases
- **Tools:** Five classification algorithms including Random ForestClassifier, AdaBoostClassifier, XGBClassifier, SVM, LightBGM; one regression algorithm logistic regression.

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	DiabetesPedigreeFunction	Age	Outcome
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1
3	1	89	66	23	94	28.1	0.167	21	0
4	0	137	40	35	168	43.1	2.288	33	
5	5	116	74	0	0	25.6	0.201	30	
6	3	78	50	32	88	31.0	0.248	26	1,
7	10	115	0	0	0	35.3	0.134	29	0
8	2	197	70	45	543	30.5	0.158	53	1
9	8	125	96	0	0	0.0	0.232	54	1

Source of the data: The National Institute
 of Diabetes and Digestive and Kidney
 Diseases

```
Pregnancies
                             0
Glucose
BloodPressure
SkinThickness
Insulin
BMI
                             Ø
DiabetesPedigreeFunction
Age
                             0
Outcome
dtype: int64
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
```

```
Data columns (total 9 columns):
    Column
                              Non-Null Count
    Pregnancies
                              768 non-nul]
                                                 cleaning
     Glucose
                              768 non-null
    BloodPressure
                              768 non-null
    SkinThickness
                              768 non-null
    Insulin
                              768 non-null
                                              int64
                                              float64
    BMI
                              768 non-null
    DiabetesPedigreeFunction 768 non-null
                                              float64
                              768 non-null
                                              int64
     Age
                                              int64
    Outcome
                              768 non-null
dtypes: float64(2), int64(7)
memory usage: 54.1 KB
None
```

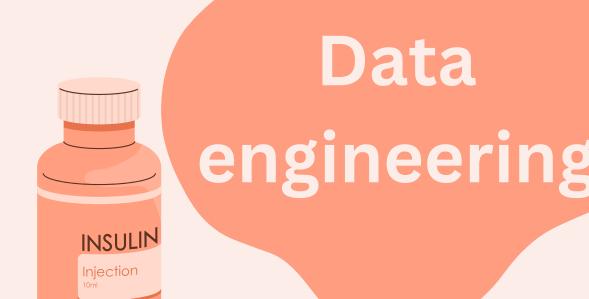
Data

```
# Replace missing values with column medians
diabetes df copy = diabetes df.copy(deep = True)
columns median=np.array([np.median(diabetes df copy["Glucose"]),
                         np.median(diabetes df copy["BloodPressure"]),
                         np.median(diabetes df copy["SkinThickness"]),
                        np.median(diabetes df copy["Insulin"]),
                         np.median(diabetes df copy["BMI"])])
diabetes_df_copy.loc[diabetes_df_copy["Glucose"]==0,"Glucose"]= columns_median[0]
diabetes df copy.loc[diabetes df copy["BloodPressure"]==0,"BloodPressure"]= columns median[1]
diabetes df copy.loc[diabetes df copy["SkinThickness"]==0,"SkinThickness"]= columns median[2]
diabetes_df_copy.loc[diabetes_df_copy["Insulin"]==0,"Insulin"]= columns_median[3]
```

Data Preprocessing

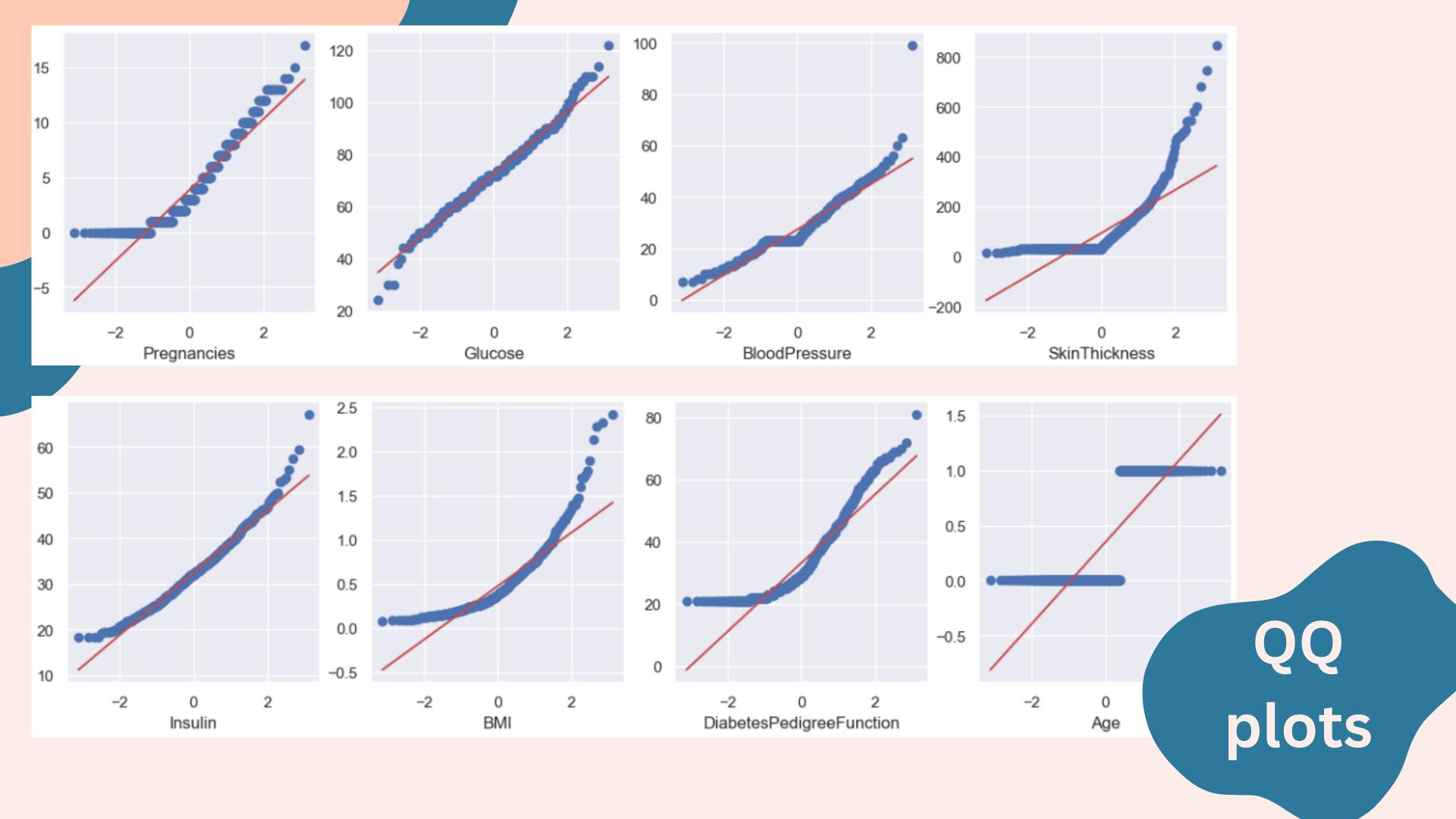
Newly created features:

- BMI_Category (categorical)
- Age_Group (categorical)
- Glucose_Insulin_Interact (numerical)



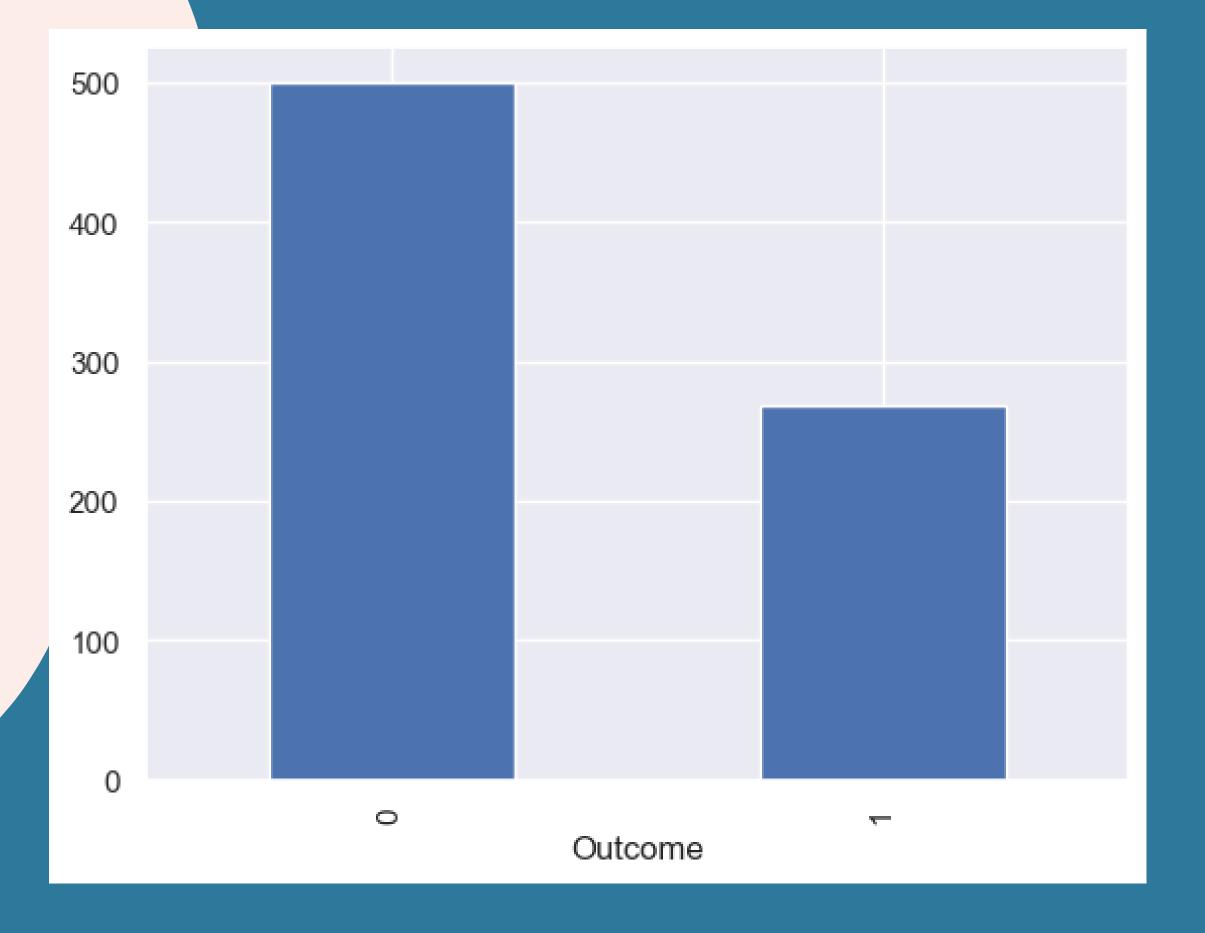
3. Data Visualization

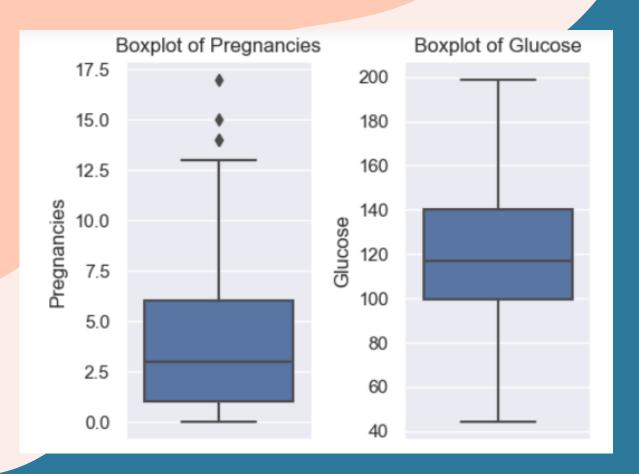


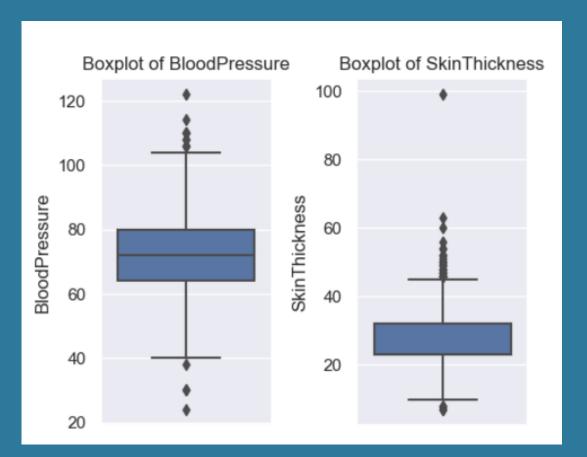


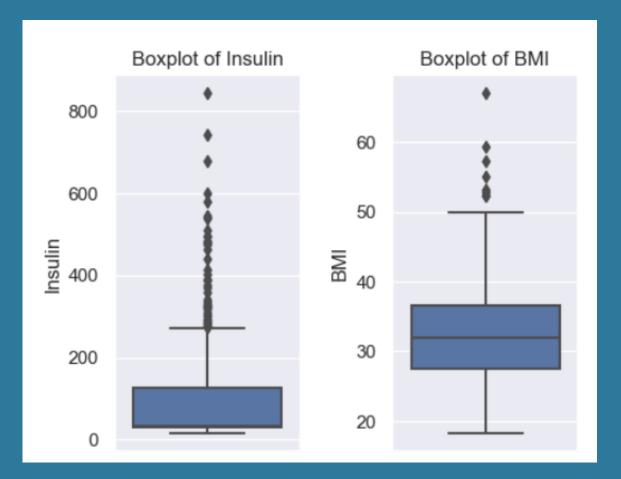
	1	0.13	0.21	0.033	-0.056	0.022	-0.034	0.54	0.22	1.0
Heatma	3	1	0.22	0.17	0.36	0.23	0.14	0.27	0.49	- 0.8
ricatina	J.21	0.22	1	0.15	-0.029	0.28	0.0024	0.32	0.17	
	0.033	0.17	0.15	1	0.24	0.55	0.14	0.055	0.19	- 0.6
	-0.056	0.36	-0.029	0.24	1	0.19	0.18	-0.015	0.15	
	0.022	0.23	0.28	0.55	0.19	1	0.15	0.026	0.31	- 0.4
	-0.034	0.14 -	0.0024	0.14	0.18	0.15	1	0.034	0.17	- 0.2
	0.54	0.27	0.32	0.055	-0.015	0.026	0.034	1	0.24	
	0.22	0.49	0.17	0.19	0.15	0.31	0.17	0.24	1	- 0.0

Checking the Balance of our data

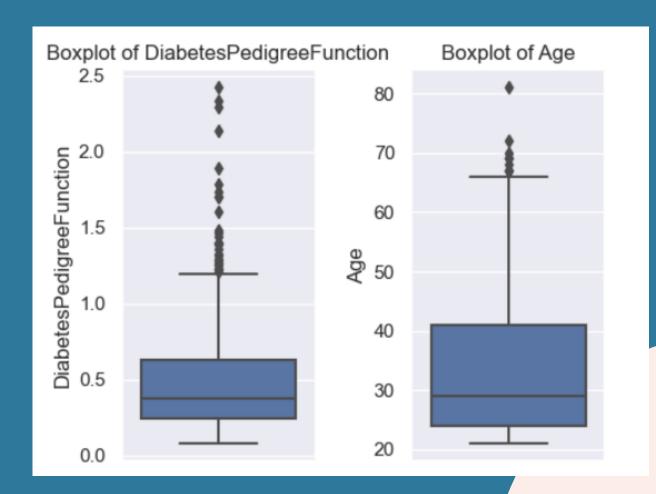








Checking outliers:
Glucose is the only
feature without any
outliers





4. Building models:

1. Logistic regression

```
# One-hot encode categorical variables
diabetes scaled = pd.get dummies(diabetes scaled, columns=['BMI Category', 'Age Group'], drop first=True)
# Logistic Regression
X lr = diabetes scaled[['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI',
                        'DiabetesPedigreeFunction', 'Age', 'Glucose_Insulin_Interact',
                        'BMI Category Normal Weight', 'BMI Category Overweight', 'BMI Category Obese I',
                        'BMI Category Obese II', 'BMI Category Obese III',
                        'Age_Group_30-40', 'Age_Group_40-50', 'Age_Group_50-60', 'Age_Group_60-70', 'Age_Group_70+']]
y lr = diabetes scaled[['Outcome']]
X_train_lr, X_test_lr, y_train_lr, y_test_lr = train_test_split(X_lr, y_lr, test_size=0.2, random_state=42)
model lr = LogisticRegression(penalty="12", max iter=100, solver="lbfgs", random state=42)
model_lr.fit(X_train_lr, y_train_lr.values.ravel())
y estimated lr = model lr.predict(X test lr)
print("Logistic Regression:")
print("Accuracy: ", accuracy_score(y_test_lr, y_estimated_lr))
print("\nConfusion Matrix: ", confusion matrix(y test lr, y estimated lr))
print("\nClassification report: ", classification_report(y_test_lr, y_estimated_lr))
```



accuracy

score:

0.77

2. Random forest

```
n estimators rf = [20, 30, 40, 50, 70, 80, 90, 100, 120]
max depth rf = np.arange(10, 30)
accuracy_matrix_rf = np.zeros([len(n_estimators_rf), len(max_depth_rf)])
for i, estimator in enumerate(n estimators rf):
   for j, depth in enumerate(max depth rf):
        rf model = RandomForestClassifier(n estimators=estimator, max depth=depth, random state=42)
       rf_model.fit(X_train_lr, y_train_lr.values.ravel())
       y estimated_rf = rf_model.predict(X_test_lr)
        accuracy matrix rf[i, j] = accuracy score(y test lr, y estimated rf)
best accuracy rf = np.max(accuracy matrix rf)
best_params_rf = np.unravel_index(np.argmax(accuracy_matrix_rf), accuracy_matrix_rf.shape)
best_n_estimators_rf = n_estimators_rf[best_params_rf[0]]
best_max_depth_rf = max_depth_rf[best_params_rf[1]]
```

<mark>0.75</mark>0.760.77<mark>0.750.75</mark>0.760.76<mark>0.79</mark>0.770.780.780.78<mark>0.79</mark>0.770.780.780.780.780.780.78

0.750.740.770.760.780.770.77<mark>0.79</mark>0.770.780.780.790.790.790.780.790.790.790.790.79

Best accuracy score: 0.79



Best accuracy score: 0.78



Heatmap of accuracy score for A	AdaBoost
---------------------------------	----------

		<u> </u>	,		
20	0.77	0.77	0.78	0.77	0.75
8	0.77	0.77	0.77	0.75	0.73
8	0.77	0.77	0.77	0.73	0.7
22	0.77	0.77	0.77	0.75	0.69
20	0.77	0.77	0.75	0.73	0.68
8	0.77	0.76	0.75	0.73	0.67
8	0.77	0.76	0.75	0.73	0.67
100	0.78	0.77	0.75	0.73	0.69
120	0.78	0.77	0.75	0.75	0.68
	0.01	0.02	0.05	0.1	1

- 0.76

- 0.74

- 0.72

- 0.70

- 0.68

3. AdaBoost

```
# AdaBoost Grid Search
base classifier = DecisionTreeClassifier(max depth=2)
n estimators ab = [20, 30, 40, 50, 70, 80, 90, 100, 120]
learning rate ab = [0.01, 0.02, 0.05, 0.1, 1]
accuracy_matrix_ab = np.zeros([len(n_estimators_ab), len(learning_rate_ab)])
for i, estimator in enumerate(n estimators ab):
    for j, rate in enumerate(learning_rate_ab):
        adaboost class = AdaBoostClassifier(base estimator=base classifier,
                                           n estimators=estimator,
                                           learning rate=rate,
                                           random state=42)
        adaboost_class.fit(X_train_lr, y_train_lr.values.ravel())
        y_estimated_ab = adaboost_class.predict(X_test_lr)
        accuracy_matrix_ab[i, j] = accuracy_score(y test lr, y estimated ab)
```

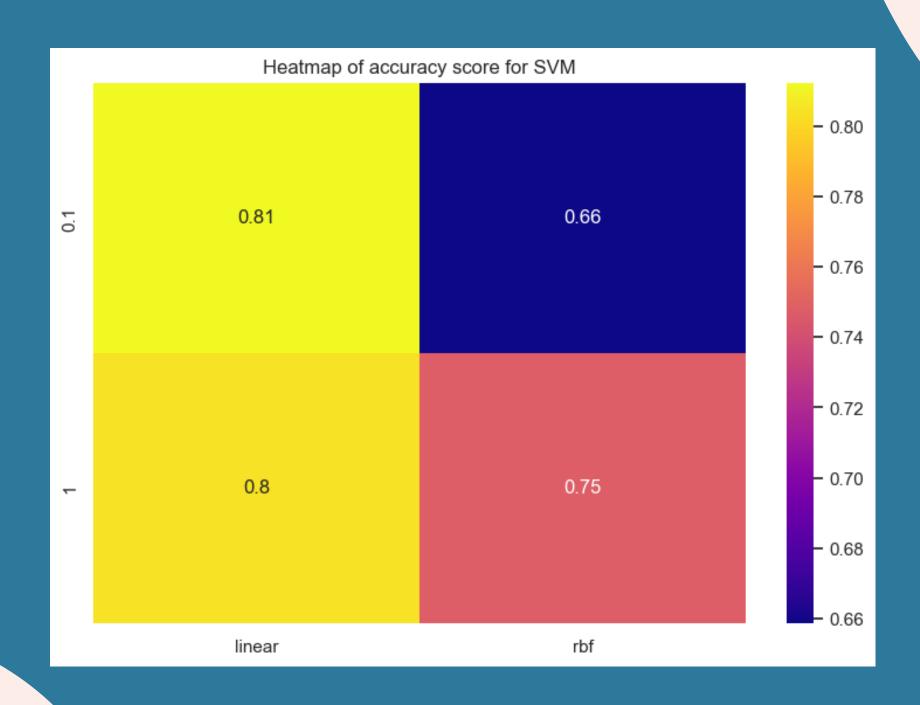
4. LightBGM

```
train_data=lgb.Dataset(X_train_lr,label=y_train_lr)
test_data=lgb.Dataset(X_test_lr,label=y_test_lr,reference=train_data)
params={"objective":"binary",
        "metric": "mse",
        "boosting type": "gbdt",
        "num boost round":100,
        "learning rate":0.05,
        "max depth":6,
        "num leaves":30,
        "feature fraction":0.9}
# train model
lgb_model = lgb.train(params,train_data,valid_sets=[test_data])
# make predictions
predictions=lgb_model.predict(X_test_lr,num_iteration=lgb_model.best_iteration)
binary predictions = [1 if pred>=0.5 else 0 for pred in predictions]
# evalutae the model
lbg_accuracy=accuracy_score(y_test_lr,binary_predictions)
print("Accuracy of LightBMG model is ",lbg accuracy)
```

accuracy score: 0.76

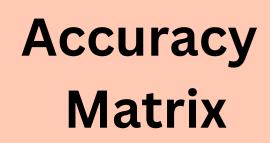
5. SVM

```
Best SVM model parameters:
{'C': 0.1, 'kernel': 'linear'}
The best SVM model accuracy is 0.8123333333333333
```



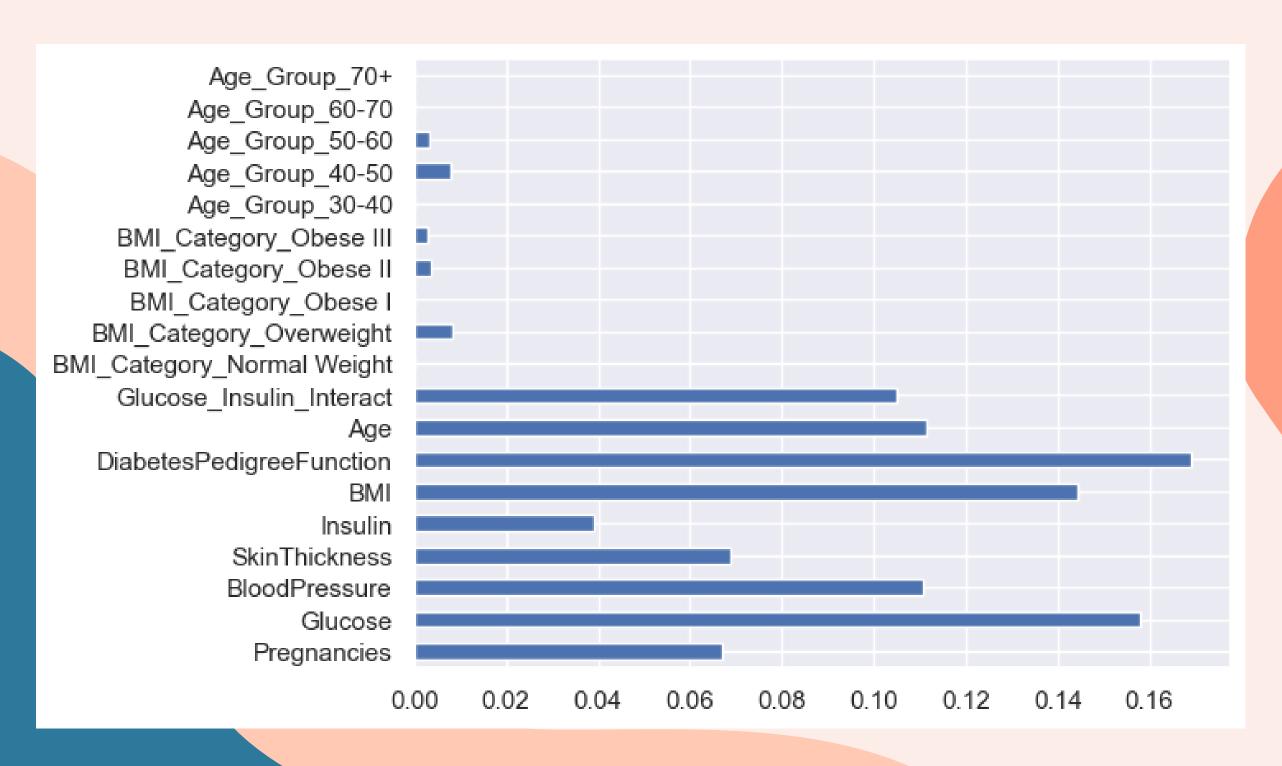
6. XG Boost Classifier

```
Best XGBoost model parameters:
{'colsample_bytree': 1.0, 'learning_rate': 0.1, 'max_depth': 3, 'n_estimators': 50, 'subsample': 0.8}
The best XGBoost model accuracy is 0.783379981340797
```



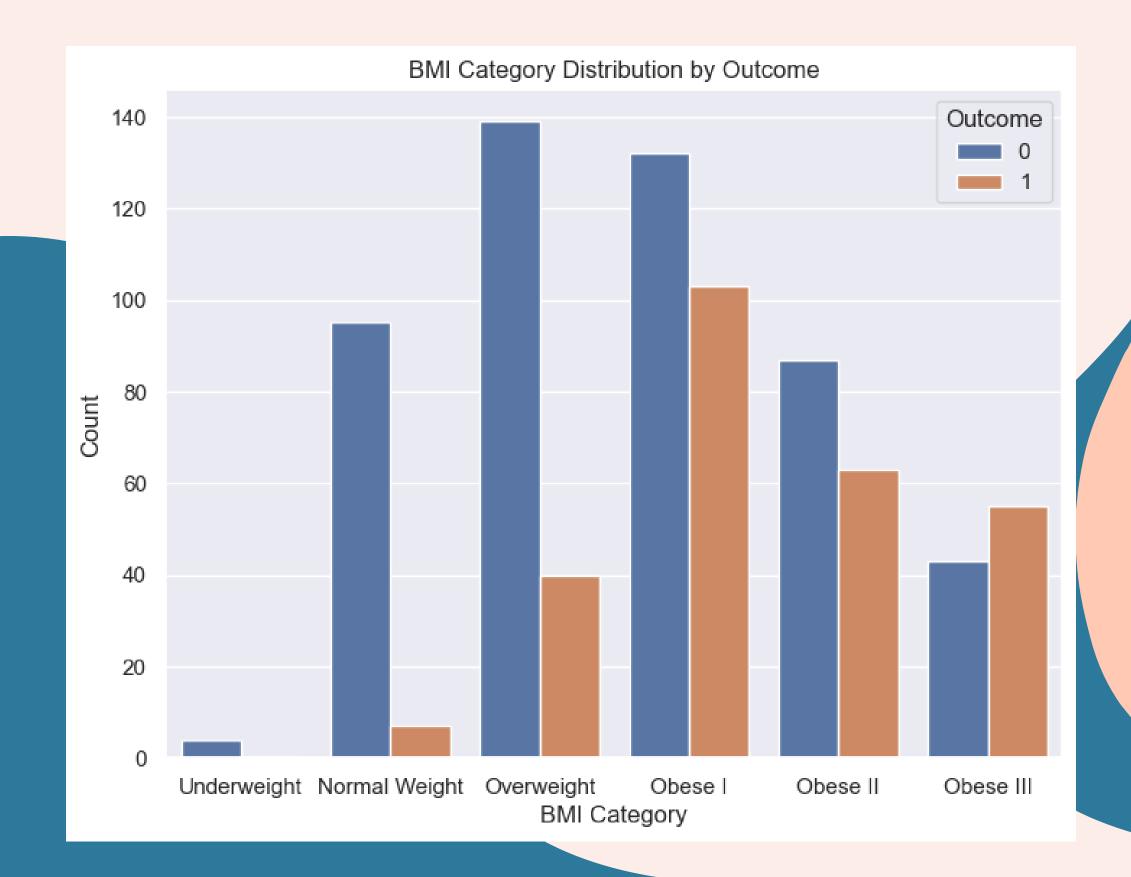
```
[[0.70030654 0.70683727 0.75737705 0.73779821 0.76871918 0.7622551
  0.70686392 0.7100893 0.76060243 0.74435559 0.76222844 0.7605891
 0.71338131 0.7198454 0.75409836 0.73775823 0.76544049 0.74919366
 0.78175397 0.78010129 0.76871918 0.76220179 0.75731041 0.7638278
 0.75731041 0.7703452 0.77361056 0.76378782 0.75893643 0.77031854
 0.75572438 0.75888311 0.75732374 0.76867919 0.75408503 0.76705318]
 [0.75893643 0.76707983 0.76545382 0.76871918 0.76218846 0.75565774
 0.76222844 0.75729708 0.75896308 0.76381447 0.75241903 0.74260962
 0.75728375 0.76057577 0.75403172 0.76546715 0.75404505 0.75893643
 0.70848994 0.72311076 0.76222844 0.75087298 0.77359723 0.7654938
 0.7117553  0.71824603  0.75733707  0.74270292  0.76869252  0.75572438
 0.71339464 0.72152472 0.7605891 0.73451953 0.76869252 0.74594162]
 [0.78337998 0.77527656 0.76548047 0.77198454 0.76221511 0.76380115
 0.76709316 0.76709316 0.76057577 0.77031854 0.76545382 0.76705318
 0.7621618  0.75728375  0.76381447  0.75732374  0.76545382  0.76709316
 0.7621618 0.76875916 0.76214847 0.7621618 0.76218846 0.76706651
 0.77029188 0.76871918 0.76381447 0.7638278 0.75563108 0.76221511
 0.7589231 0.7638278 0.75405838 0.76054911 0.7589231 0.75565774]]
```

Feature Importance



Features that make most contributions are DiabetesPedigreeFunct ion, Glucose, BMI, Age, BloodPressure.

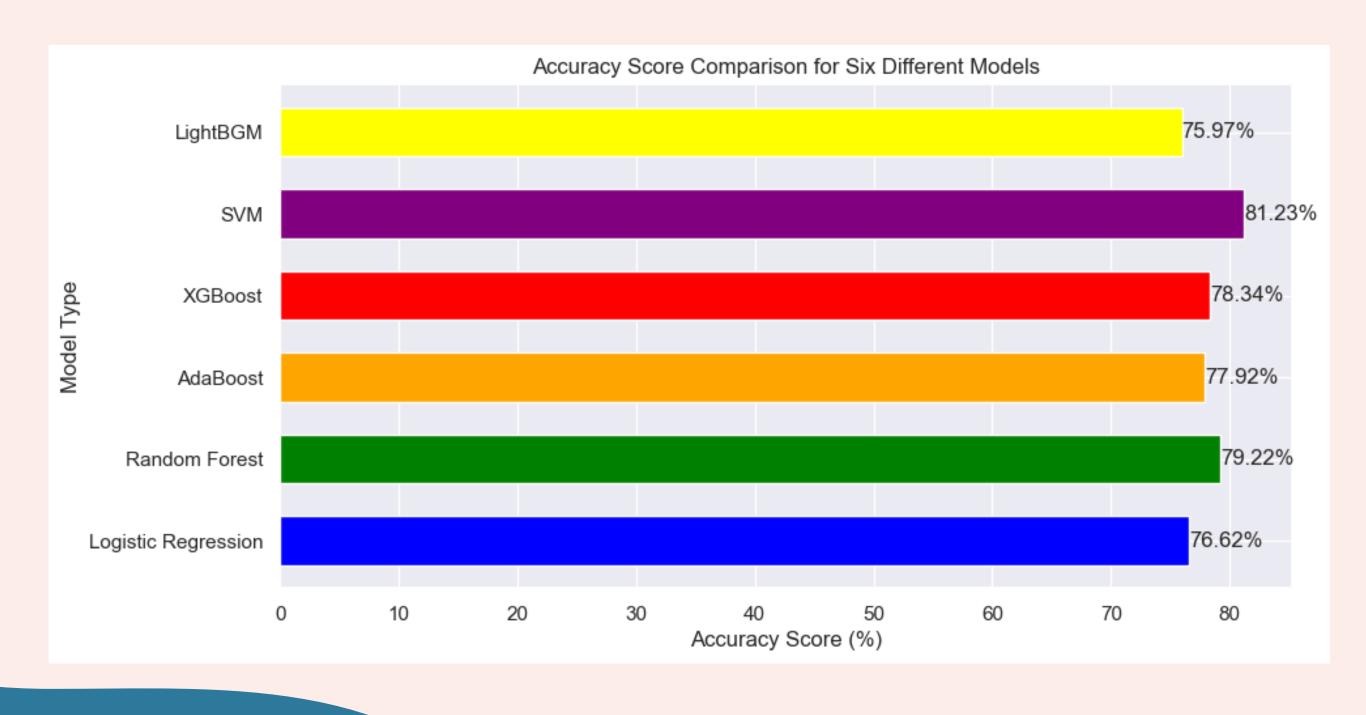
Feature Engineering



The following new features were added:

- BMI_Cateogy
- Glucose_Insulin_Int eract
- Age_Group

Performance Comparison And the winner is!!!



Further Improvements

- Better Feature Engineering
- Regularization
- Better Scaling of Data



THANK YOU FOR YOUR ATTENTION! WE ARE NOW OPEN FOR

ANY QUESTIONS!