



+ Code + Text

Connect ▾

Colab AI



Importing required Libraries



```
# import some libraries
import numpy as np;
import pandas as pd;
import seaborn as sns;
import matplotlib.pyplot as plt;
import os;
```

Importing important libraries for predictions



```
[ ] # importing important libraries for prediction
from sklearn.model_selection import train_test_split;
from sklearn.linear_model import LogisticRegression;
from sklearn.metrics import accuracy_score;
from sklearn.tree import DecisionTreeClassifier;
from sklearn.ensemble import RandomForestClassifier;
```



+ Code + Text

Connect

Colab AI



Loading the dataset

```
[ ] data=pd.read_csv("/diabetes.csv");  
print(data);
```

| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | BMI | \ |
|-----|-------------|---------|---------------|---------------|---------|------|---|
| 0 | 6 | 148 | 72 | 35 | 0 | 33.6 | |
| 1 | 1 | 85 | 66 | 29 | 0 | 26.6 | |
| 2 | 8 | 183 | 64 | 0 | 0 | 23.3 | |
| 3 | 1 | 89 | 66 | 23 | 94 | 28.1 | |
| 4 | 0 | 137 | 40 | 35 | 168 | 43.1 | |
| .. | ... | ... | ... | ... | ... | ... | |
| 763 | 10 | 101 | 76 | 48 | 180 | 32.9 | |
| 764 | 2 | 122 | 70 | 27 | 0 | 36.8 | |
| 765 | 5 | 121 | 72 | 23 | 112 | 26.2 | |
| 766 | 1 | 126 | 60 | 0 | 0 | 30.1 | |
| 767 | 1 | 93 | 70 | 31 | 0 | 30.4 | |

| | DiabetesPedigreeFunction | Age | Outcome |
|-----|--------------------------|-----|---------|
| 0 | 0.627 | 50 | 1 |
| 1 | 0.351 | 31 | 0 |
| 2 | 0.672 | 32 | 1 |
| 3 | 0.167 | 21 | 0 |
| 4 | 2.288 | 33 | 1 |
| .. | ... | ... | ... |
| 763 | 0.171 | 63 | 0 |



+ Code + Text

Connect

Colab AI



```
[ ] ..
763 0.171 63 0
764 0.340 27 0
765 0.245 30 0
766 0.349 47 1
767 0.315 23 0
```

[768 rows x 9 columns]

Explore the data

```
[ ] print(data.info);
```

```
<bound method DataFrame.info of
0      6    148    72     35     0  33.6
1      1     85    66     29     0  26.6
2      8    183    64      0     0  23.3
3      1     89    66     23    94  28.1
4      0    137    40     35   168  43.1
..     ..     ..     ..     ..     ..
763    10    101    76     48   180  32.9
764     2    122    70     27     0  36.8
765     5    121    72     23   112  26.2
766     1    126    60      0     0  30.1
767     1     92    70     21     0  20.1
```



+ Code + Text

Connect ▼

Colab AI



{x}



```
[ ] 767      1      93      70      31      0 30.4
      DiabetesPedigreeFunction Age Outcome
0      0.627 50      1
1      0.351 31      0
2      0.672 32      1
3      0.167 21      0
4      2.288 33      1
..      ...  ...  ...
763     0.171 63      0
764     0.340 27      0
765     0.245 30      0
766     0.349 47      1
767     0.315 23      0
```

[768 rows x 9 columns]>

```
[ ] print(data.describe);
```



```
<bound method NDFrame.describe of
0      6      148      72      35      0 33.6
1      1      85      66      29      0 26.6
2      8      183      64      0      0 23.3
3      1      89      66      23      94 28.1
4      0      137      40      35     168 43.1
..      ...  ...  ...  ...  ...  ...
763     10     101      76      48     190 22.0
```



+ Code + Text

Connect ▼

Colab AI



```
[ ] 4          0      137          40          35      168  43.1
    ..      ...      ...      ...      ...      ...      ...
    763      10      101          76          48      180  32.9
    764          2      122          70          27          0  36.8
    765          5      121          72          23      112  26.2
    766          1      126          60          0          0  30.1
    767          1      93          70          31          0  30.4
```

DiabetesPedigreeFunction Age Outcome

```
0          0.627  50      1
1          0.351  31      0
2          0.672  32      1
3          0.167  21      0
4          2.288  33      1
..      ...      ...      ...
763      0.171  63      0
764      0.340  27      0
765      0.245  30      0
766      0.349  47      1
767      0.315  23      0
```

[768 rows x 9 columns]>

Data cleaning



+ Code + Text

Connect

Colab AI



Data cleaning

{x}

```
[ ] data_cleaning=data.dropna();
```



```
[ ] print("Clean Data:");  
print(data_cleaning.head());
```



Clean Data:

| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | BMI | \ |
|---|-------------|---------|---------------|---------------|---------|------|---|
| 0 | 6 | 148 | 72 | 35 | 0 | 33.6 | |
| 1 | 1 | 85 | 66 | 29 | 0 | 26.6 | |
| 2 | 8 | 183 | 64 | 0 | 0 | 23.3 | |
| 3 | 1 | 89 | 66 | 23 | 94 | 28.1 | |
| 4 | 0 | 137 | 40 | 35 | 168 | 43.1 | |

| | DiabetesPedigreeFunction | Age | Outcome |
|---|--------------------------|-----|---------|
| 0 | 0.627 | 50 | 1 |
| 1 | 0.351 | 31 | 0 |
| 2 | 0.672 | 32 | 1 |
| 3 | 0.167 | 21 | 0 |
| 4 | 2.288 | 33 | 1 |

<>



Checking for missing values

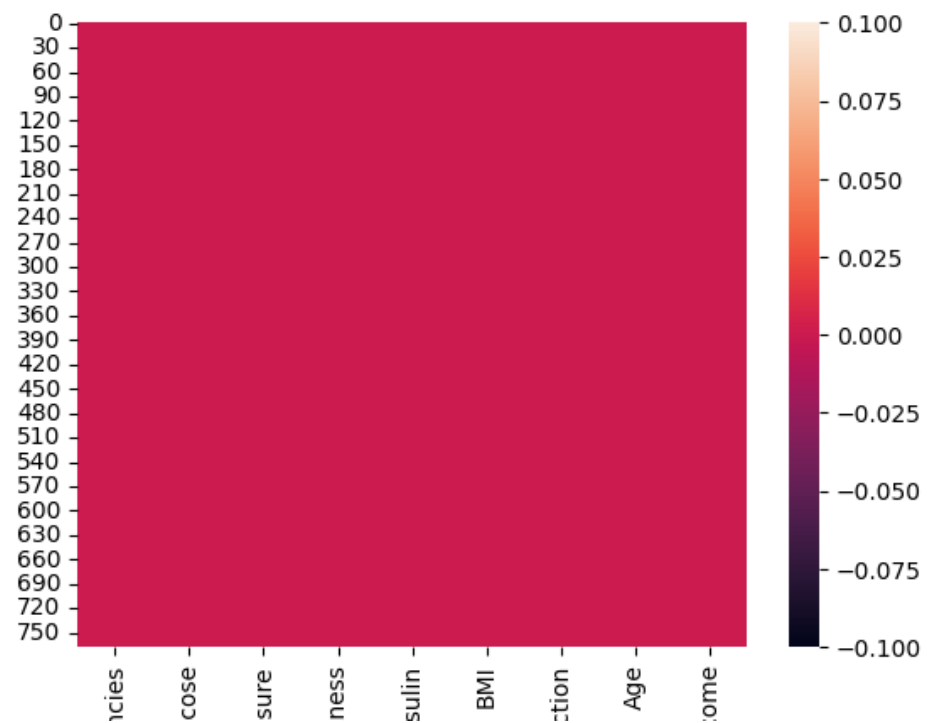
+ Code + Text

Connect ▼

Colab AI

^

```
[ ] sns.heatmap(data.isnull());
```



+ Code + Text

Connect

Colab AI

Co Relation Matrix

```
[ ] Correlation=data.corr();  
    print(Correlation);
```

| | Pregnancies | Glucose | BloodPressure | SkinThickness | \ |
|--------------------------|-------------|----------|---------------|---------------|---|
| Pregnancies | 1.000000 | 0.129459 | 0.141282 | -0.081672 | |
| Glucose | 0.129459 | 1.000000 | 0.152590 | 0.057328 | |
| BloodPressure | 0.141282 | 0.152590 | 1.000000 | 0.207371 | |
| SkinThickness | -0.081672 | 0.057328 | 0.207371 | 1.000000 | |
| Insulin | -0.073535 | 0.331357 | 0.088933 | 0.436783 | |
| BMI | 0.017683 | 0.221071 | 0.281805 | 0.392573 | |
| DiabetesPedigreeFunction | -0.033523 | 0.137337 | 0.041265 | 0.183928 | |
| Age | 0.544341 | 0.263514 | 0.239528 | -0.113970 | |
| Outcome | 0.221898 | 0.466581 | 0.065068 | 0.074752 | |

| | Insulin | BMI | DiabetesPedigreeFunction | \ |
|--------------------------|-----------|----------|--------------------------|---|
| Pregnancies | -0.073535 | 0.017683 | -0.033523 | |
| Glucose | 0.331357 | 0.221071 | 0.137337 | |
| BloodPressure | 0.088933 | 0.281805 | 0.041265 | |
| SkinThickness | 0.436783 | 0.392573 | 0.183928 | |
| Insulin | 1.000000 | 0.197859 | 0.185071 | |
| BMI | 0.197859 | 1.000000 | 0.140647 | |
| DiabetesPedigreeFunction | 0.185071 | 0.140647 | 1.000000 | |
| Age | -0.042163 | 0.036242 | 0.033561 | |
| Outcome | 0.130548 | 0.292695 | 0.173844 | |



+ Code + Text

Connect ▼

Colab AI



```
[ ] Age -0.042163 0.036242 0.033561
Outcome 0.130548 0.292695 0.173844
```

| | Age | Outcome |
|--------------------------|-----------|----------|
| Pregnancies | 0.544341 | 0.221898 |
| Glucose | 0.263514 | 0.466581 |
| BloodPressure | 0.239528 | 0.065068 |
| SkinThickness | -0.113970 | 0.074752 |
| Insulin | -0.042163 | 0.130548 |
| BMI | 0.036242 | 0.292695 |
| DiabetesPedigreeFunction | 0.033561 | 0.173844 |
| Age | 1.000000 | 0.238356 |
| Outcome | 0.238356 | 1.000000 |

Visualizing the Correlation

```
[ ] sns.heatmap(Correlation);
```



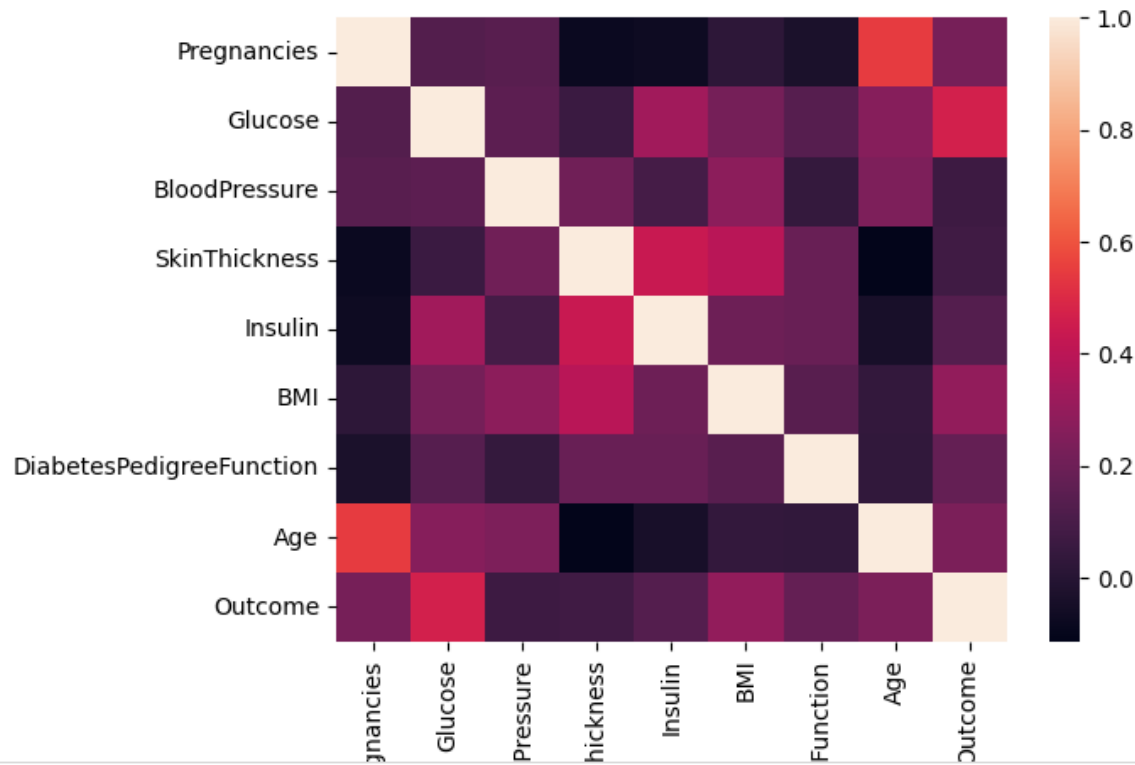
+ Code + Text

Connect ▼

Colab AI



[]





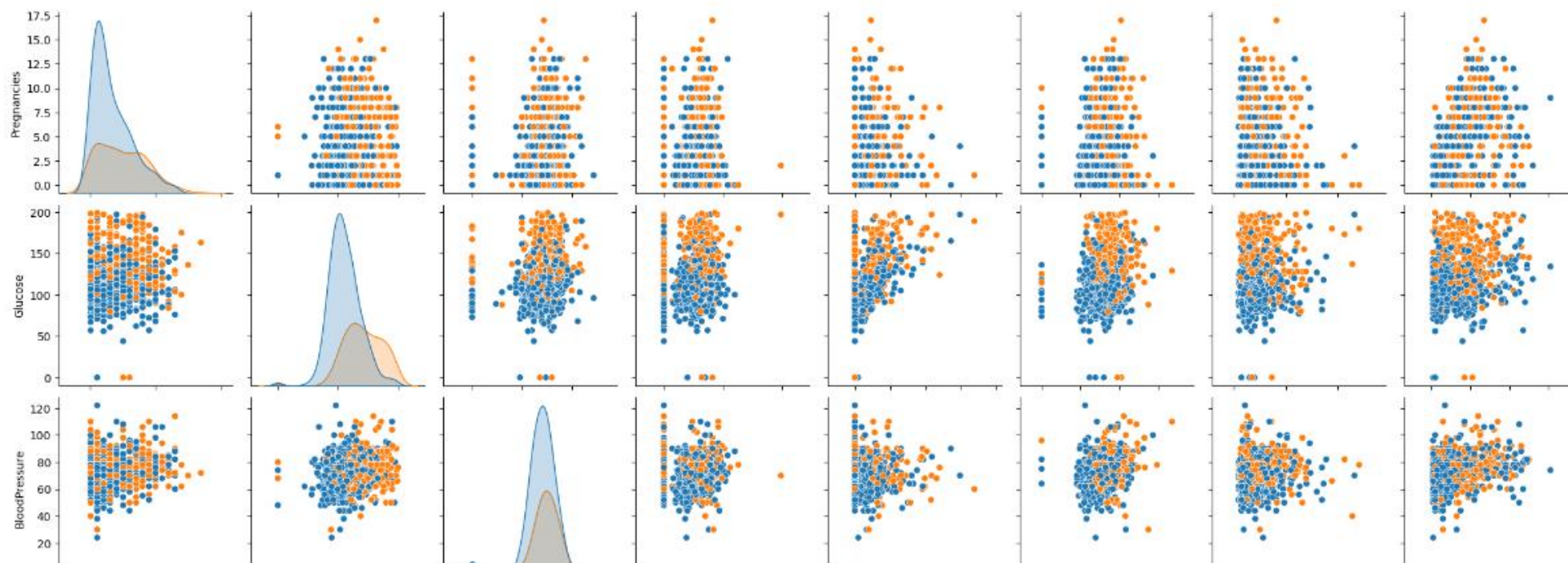
+ Code + Text

Connect

Colab AI



```
[ ] sns.pairplot(data,hue='Outcome');  
plt.show();
```





+ Code + Text

Connect

Colab AI



Training the model with the help of train test split



```
[ ] X = data.drop("Outcome", axis=1)
    Y = data['Outcome'] # Assuming 'Outcome' is the column name containing labels
    X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.2)
```



Training the model

```
[ ] model=LogisticRegression();
    model.fit(X_train,Y_train)
```

/usr/local/lib/python3.10/dist-packages/sklearn/linear_model/_logistic.py:458: ConvergenceWarning: lbfgs failed to converge (status=1):
STOP: TOTAL NO. of ITERATIONS REACHED LIMIT.

Increase the number of iterations (max_iter) or scale the data as shown in:

<https://scikit-learn.org/stable/modules/preprocessing.html>

Please also refer to the documentation for alternative solver options:

https://scikit-learn.org/stable/modules/linear_model.html#logistic-regression

```
n_iter_i = _check_optimize_result(
```

```
    LogisticRegression
```

```
LogisticRegression())
```





+ Code + Text

Connect ▼

Colab AI



Decision Tree



```
[ ] dt_model=DecisionTreeClassifier();  
    dt_model.fit(X_train,Y_train);  
    dt_pred=dt_model.predict(X_test);  
    print("Decision Tree Accuracy:", accuracy_score(Y_test,dt_pred));
```



Decision Tree Accuracy: 0.6103896103896104



Random Forest



```
[ ] rf_model=RandomForestClassifier();  
    rf_model.fit(X_train,Y_train);  
    rf_pred=rf_model.predict(X_test);  
    print("Random Forest Accuracy:", accuracy_score(Y_test,rf_pred));
```

Random Forest Accuracy: 0.7272727272727273



Making prediction



+ Code + Text

Connect ▼

Colab AI



Random Forest Accuracy: 0.7272727272727273

Making prediction

```
[ ] Prediction=model.predict(X_test);  
    print(Prediction);
```

```
[0 0 1 0 0 0 0 0 0 1 0 0 0 0 0 1 1 0 1 1 1 1 0 0 1 0 0 0 0 0 0 0 0 0  
 1 0 0 0 0 0 0 0 1 0 0 0 0 0 1 0 0 0 1 1 0 1 0 0 0 0 0 0 0 0 1 0 1 0 1 0  
 0 1 0 0 0 1 0 0 0 0 0 1 0 0 1 0 0 1 0 0 0 1 1 0 0 1 1 1 0 0 1 0 0 0 0 1 0  
 0 0 0 0 1 0 0 0 1 0 0 0 0 1 0 0 1 0 0 0 1 0 1 0 0 1 0 1 0 0 0 0 0 0 0 0 0  
 0 0 0 0 0 0]
```

```
[ ] Accuracy=accuracy_score(Prediction,Y_test);  
    print(Accuracy);
```

```
0.7337662337662337
```



+ Code + Text

Connect ▼

Colab AI



Total no of prediction



```
[ ] Total_Diabetes=sum(Prediction);  
    print("Total number of people with diabetes:",Total_Diabetes);
```

Total number of people with diabetes: 39



```
[ ] Total_non_Diabetes=len(Prediction);  
    print("Total number of people with no Diabetes:", Total_non_Diabetes);
```

Total number of people with no Diabetes: 154

**Prediction Visualization **



```
[ ] Total=[Total_Diabetes,Total_non_Diabetes];  
    Labels=['Diabetes', 'Non_Diabetes'];  
    plt.bar(Labels,Total,color=['red','blue']);  
    plt.xlabel('Diabetes Status');  
    plt.ylabel('Total number of people');  
    plt.title('Total number of people with and without Diabetes');  
    plt.show();
```





+ Code + Text

Connect ▼

Colab AI



[]

