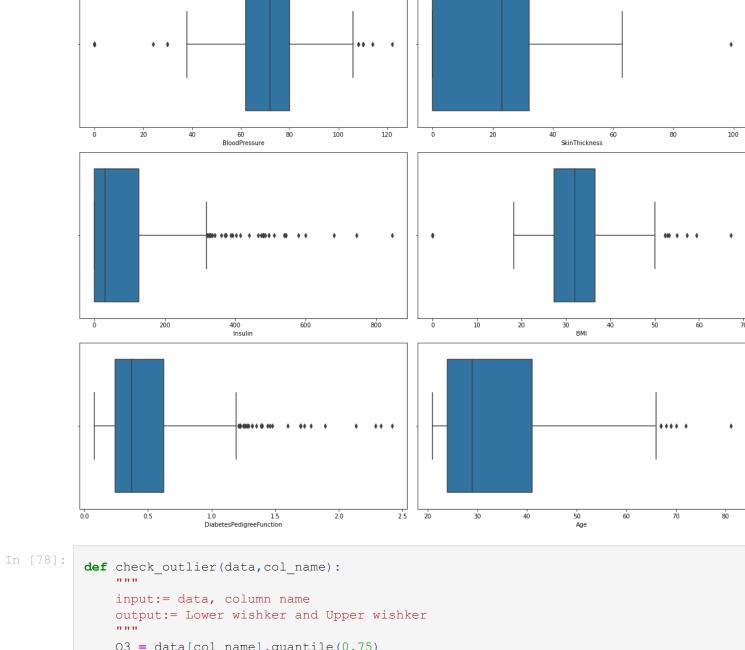
Dataset Description - Pregnancies: Number of times pregnant - Glucose: Plasma glucose concentration a 2 hours in an oral glucose tolerance - BloodPressure: Diastolic blood pressure (mm Hg) - SkinThickness: Triceps skin fold thickness (mm) - Insulin: 2-Hour serum insulin (mu U/ml) BMI: Body mass index (weight in kg/(height in m)^2) - DiabetesPedigreeFunction: Diabetes pedigree function - Age: Age (years) - Outcome: Class variable (0 or 1) - Dataset Link : https://www.kaggle.com/datasets/mathchi/diabetes-dataset/code #Import Libraries import warnings warnings.filterwarnings("ignore") import pandas as pd import seaborn as sns import matplotlib.pyplot as plt import numpy as np # Import Data df = pd.read_csv('diabetes.csv') df.head() Pregnancies Glucose BloodPressure SkinThickness Insulin BMI DiabetesPedigreeFunction Age Outcome 0 6 148 72 35 0 33.6 0.627 50 1 85 29 26.6 0.351 66 2 8 183 64 0 0 23.3 0.672 32 3 89 66 94 28.1 0.167 4 0 137 40 35 168 43.1 2.288 33 1 df.shape Out[71]: (768, 9) df.info() <class 'pandas.core.frame.DataFrame'> RangeIndex: 768 entries, 0 to 767 Data columns (total 9 columns): Non-Null Count Dtype Column ---------0 Pregnancies 768 non-null int64 Glucose 1 768 non-null int64 int64 int64 BloodPressure 768 non-null SkinThickness 768 non-null 3 int64 Insulin 768 non-null 768 non-null float64 5 DiabetesPedigreeFunction 768 non-null 6 float64 768 non-null 7 Age int64 8 768 non-null int64 Outcome dtypes: float64(2), int64(7) memory usage: 54.1 KB df.describe() Glucose BloodPressure SkinThickness BMI DiabetesPedigreeFunction **Pregnancies** Insulin 768.000000 768.000000 768.000000 768.000000 768.000000 768.000000 768.000000 count mean 3.845052 120.894531 69.105469 20.536458 79.799479 31.992578 0.471876 19.355807 3.369578 31.972618 15.952218 115.244002 7.884160 0.331329 std 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 min 0.078000 0.000000 25% 1.000000 99.000000 62.000000 0.000000 27.300000 0.243750 72.000000 32.000000 **50**% 3.000000 117.000000 23.000000 30.500000 0.372500 140.250000 80.000000 32.000000 127.250000 36.600000 **75**% 6.000000 0.626250 17.000000 199.000000 122.000000 99.000000 846.000000 67.100000 2.420000 max In [74]: df.columns Index(['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', Out[74]: 'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome'], dtype='object') df.dtypes Out[75]: Pregnancies int64 Glucose int64 BloodPressure int64 SkinThickness int64 Insulin int64 BMI float64 DiabetesPedigreeFunction float64 int64 Age int64 Outcome dtype: object Applying Some EDA df.isnull().sum() 0 Out[76]: Pregnancies 0 Glucose BloodPressure SkinThickness 0 Insulin 0 BMI 0 DiabetesPedigreeFunction 0 Outcome 0 dtype: int64 fig, ax = plt.subplots(4, 2, figsize=(16,18))axes list = [axes for axes_row in ax for axes in axes_row] column list = list(df.columns) for i, col in enumerate(column list[:-1]): sns.boxplot(data=df, x=col, ax=axes list[i]) plt.tight layout() plt.show() 7.5 Pregnancies 12.5 15.0 125 10.0 Glucose



```
Q3 = data[col_name].quantile(0.75)
    Q1 = data[col name].quantile(0.25)
    IQR = Q3-Q1
    print("75%:", Q3)
    print("25%",Q1)
    print("IQR:",IQR)
    LW = Q1 - 1.5*IQR
    UW = Q3 + 1.5*IQR
    print("Lower and Upper Wishker: ",LW, UW)
    print("Min and Max value: ", np.min(data[col_name]),np.max(data[col_name]))
    print("Full data:", data.shape)
    print("No of outliers: ",data[(data[col_name]<LW) | (data[col_name]>UW)].shape)
    sns.boxplot(x=data[col name])
    sns.stripplot(x=data[col name], color="0.5")
    plt.show()
    return LW, UW
# for c in df.columns:
# print("Column: ",c)
c = 'Age'
LW, UW= check_outlier(df,c)
# print("After removing outliers")
df = df[(df[c] >= LW) & (df[c] <= UW)]
# sns.boxplot(x=df[c])
# plt.show()
# print("="*40)
75%: 38.0
25% 23.5
IQR: 14.5
Lower and Upper Wishker: 1.75 59.75
```

