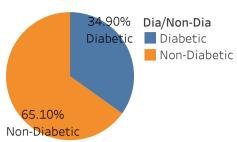
<pre>In [1]: In [2]: Out[2]:</pre>	import numpy as np import pandas as pd import matplotlib.pyplot as plt import seaborn as sns *** **mmatplotlib inline**  data_df=pd.read_csv(r'C:\\Users\Mohd Shadab\\Downloads\\Projects\\CAPSTONE PROJECTS\\Project 2\\Healthcare - Diabetes\\health care diabetes.csv')  data_df*  Pregnancies Glucose BloodPressure SkinThickness Insulin BMI 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 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 1
<pre>In [3]: Out[3]: In [4]:</pre>	(768, 9)  data_df.info()
In [5]:	<pre>cclass 'pandas.core.frame.DataFrame'&gt; RangeIndex: 768 entries, 0 to 767 Data columns (total 9 columns): # Column</pre>
Out[5]: In [6]:	count         768.0000000         768.0000000         768.0000000         768.00000
Out[6]:	Pregnancies         Glucose         Blood/Pressure         SkinThickness         Insulin         BM         Diabetes/PedigreeFunction         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         1  <
In [7]: In [9]:	#Statistical data above shows the presence of nulls in the columns Glucose, BloodPressure, SkinThickness, Insulin, BMI  #Since the value 0 doesn't make any sense.  data_df.hist(figsize=(18,10));  Pregnancies Glucose BloodPressure  250 200 150 150 150 150 150 150 150 150 150 1
	0.0 2.5 5.0 7.5 10.0 12.5 15.0 17.5 0 0 25 50 75 100 12.5 150 175 200 0 20 40 60 80 100 120 SkinThickness Insulin BMI 250 200 150 150 150 150 150 150 150 150 150 1
In [10]: In [11]: In [12]:	#Histogram visualisation confirms the presence of @ values.The rows with @ in such columns shall be treated as null  data_df['Glucose'].replace(@,np.nan,inplace=True) data_df['BloodPressure'].replace(@,np.nan,inplace=True) data_df['SkinThickness'].replace(@,np.nan,inplace=True) data_df['Insulin'].replace(@,np.nan,inplace=True) data_df['BMI'].replace(@,np.nan,inplace=True) data_df['BMI'].replace(@,np.nan,inplace=True)
n [13]: nut[13]:	Pregnancies 0 6 6 6 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
in [14]: in [15]: in [16]:	BMI 1.432292 DiabetesPedigreeFunction 0.000000 Age 0.000000 Outcome 0.000000 dtype: float64
Out[16]: In [17]:	Pregnancies 0
	75 50 40 60 80 100 120 140 160 180 200 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 100 120 140 160 180 180 180 180 180 180 180 180 180 18
n [18]:	300 250 200 150 100 50 0.0 0.5 1.0 1.5 2.0 2.5 020 30 40 50 60 70 80 0.0 0.2 0.4 0.6 0.8 1.0
n [19]:	Q1. Check the balance of the data by plotting the count of outcomes by their value. Describe your findings and plan future course of action.  data_df['Outcome'].value_counts()*100/len(data_df)  0 65.104167
ut[19]: n [20]: n [21]: ut[21]: ut[22]:	1 34.895833 Name: Outcome, dtype: float64  #The above percentage shows that the data is not imbalanced.  data_df.columns  Index(['Pregnancies', 'Glucose', 'BloodPressure', 'SkinThickness', 'Insulin', 'BMI', 'DiabetesPedigreeFunction', 'Age', 'Outcome'], dtype='object')  Q2. Create scatter charts between the pair of variables to understand the relationships. Describe your findings.  sns.pairplot(data_df) <seaborn.axisgrid.pairgrid 0x2095b8b0df0="" at=""></seaborn.axisgrid.pairgrid>
	17.5   12
	120 100 100 100 100 100 100 100 100 100
	5 5 5 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
n [23]: ut[23]:	Q3. Perform correlation analysis. Visually explore it using a heat map.  Plt. figure(figsize=(8,5)) data=data_df.corr() sns.heatmap(data=data_,annot=True) <axessubplot:></axessubplot:>
	Pregnancies - 1 0.13 0.21 0.082 0.025 0.013 0.034 0.54 0.22  Glucose - 0.13 1 0.22 0.19 0.42 0.064 0.14 0.27 0.49  BloodPressure - 0.21 0.22 1 0.19 0.045 0.17 0.0024 0.32 0.17  SkinThickness - 0.082 0.19 0.19 1 0.16 0.27 0.1 0.13 0.097 0.2  Insulin - 0.025 0.42 0.045 0.16 1 0.073 0.13 0.097 0.2  BMI - 0.013 0.064 0.17 0.27 0.073 1 0.069 0.011 0.13 0.097 0.2  DiabetesPedigreeFunction - 0.034 0.14 0.0024 0.1 0.13 0.069 1 0.034 0.17  Age - 0.54 0.27 0.32 0.13 0.097 0.011 0.034 1 0.24  Outcome - 0.22 0.49 0.17 0.21 0.2 0.13 0.17 0.24 1 -0.0
In [27]: In [28]: Out[29]: In [30]:	print(x_train.shape) print(x_train.shape) print(y_train.shape) print(y_t
Out[30]: In [31]: Out[31]: In [32]:	<pre>KNeighborsClassifier()  #Predicting the test set result y_pred= KNN_classifier.predict(x_test) y_pred  array([1, 0, 0, 1, 0, 0, 1, 1, 1, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,</pre>
In [33]:	<pre>for i in range(1,7):     for j in range(1,5):         KNN_classifier= KNeighborsClassifier(n_neighbors=i, metric='minkowski', p=j)         KNN_classifier.fit(x_train, y_train)         y_pred= KNN_classifier.predict(x_test)         print("Accuracy =",accuracy_score(y_test,y_pred),"n_neighbors =",i," p =",j)  Accuracy = 0.6979166666666666 n_neighbors = 1</pre>
n [36]:	Accuracy = 0.713541666666666 n_neighbors = 2
ut[36]: n [37]: ut[37]: n [38]:	<pre>KNN_classifier= KNeighborsClassifier(n_neighbors=6, metric='minkowski', p=2 ) KNN_classifier.fit(x_train, y_train) y_pred= KNN_classifier.predict(x_test) KNN_Accuracy =accuracy_score(y_test,y_pred) KNN_Accuracy  0.7864583333333334  Logistic regression model  from sklearn.linear_model import LogisticRegression LR_classifier= LogisticRegression(random_state=0) LR_classifier.fit(x_train, y_train)  LogisticRegression(random_state=0)</pre>
n [38]: ut[38]: n [39]:	##Healting the test is tresh. y_pred   array([1, 0, 0, 1, 0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 1, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
n [40]: n [41]: ut[41]: n [42]:	[ 25 37]] Accuracy = 0.791666666666666  LR_Accuracy_accuracy_score(y_test,y_pred)  Using Decision tree  #Fitting Decision Tree classifier to the training set from sklearn.tree import DecisionTreeClassifier DT_classifier= DecisionTreeClassifier(criterion='entropy', random_state=0) DT_classifier.fit(x_train, y_train)  DecisionTreeClassifier(criterion='entropy', random_state=0)  #Predicting the test set result y_pred= DT_classifier.predict(x_test)
	<pre>y_pred = DT_classifier.predict(x_test) y_pred  array([1, 1, 1, 1, 0, 0, 1, 0, 0, 1, 1, 0, 0, 1, 1, 0, 0, 0, 0, 1, 0,</pre>
n [44]:	[ 25 37]] Accuracy = 0.713541666666666
n [45]: n [46]:	<pre>#Tuning hyperparameter increases the accuracy by around 5%  Random Forest  #Fitting Decision Tree classifier to the training set from sklearn.ensemble import RandomForestClassifier for i in range(5,15):     RF_classifier= RandomForestClassifier(n_estimators= 10, criterion="entropy")     RF_classifier= Riclassifier.fit(x_train, y_train)     y_pred= RF_classifier.predict(x_test)     cm= confusion_matrix(y_test, y_pred)     print("Accuracy = ",accuracy_score(y_test,y_pred), "n_estimators = ",i)  Accuracy = 0.76520833333333334</pre>
n [47]:	
n [48]: ut[48]: n [49]: ut[49]:	Accuracy   KNN   0.786458   LR   0.791667
n [50]: n [51]: n [52]:	LR 0.791667  DT 0.760417  RF 0.776042  GB 0.776042  #Highest accuracy is given by Logistic Regression  Using cross validation to avoid the data leakage effect on model and checking the accuracy  #KNN  #Testing on train dataset without exposing the test dataset from sklearn.model_selection import cross_val_score scores=cross_val_score(KNN_classifier,x_train,y_train,scoring='accuracy',cv=10)
n [53]: ut[53]: n [54]: ut[54]: n [55]:	scores=cross_val_score(KNN_classifier,x_train,y_train, scoring='accuracy',cv=10)  scores.mean()  0.7045674531155475  #fitting the model KNN_classifier.fit(x_train,y_train)  KNeighborsClassifier(n_neighbors=6)  y_final_pred=KNN_classifier.predict(x_test) accuracy_score(y_test,y_final_pred)  0.7864583333333334
ut[55]: n [56]: n [57]: ut[57]: ut[58]:	#Logistic Regression  from sklearn.model_selection import cross_val_score scores=cross_val_score(LR_classifier,x_train,y_train,scoring='accuracy',cv=10) scores.mean()  0.7447368421052631  LR_classifier.fit(x_train,y_train) y_final_pred=LR_classifier.predict(x_test) accuracy_score(y_test,y_final_pred)  0.791666666666666666666666666666666666666
ut[58]: n [59]: n [60]: ut[60]: ut[61]:	#Decision Tree  from sklearn.model_selection import cross_val_score scores=cross_val_score(DT_classifier,x_train,y_train,scoring='accuracy',cv=10) scores.mean()  DT_classifier.fit(x_train,y_train) y_final_pred=DT_classifier.predict(x_test) accuracy_score(y_test,y_final_pred)  0.76041666666666666
n [62]: n [63]: ut[63]: n [64]:	#Random Forest  from sklearn.model_selection import cross_val_score scores=cross_val_score(RF_classifier,x_train,y_train,scoring='accuracy',cv=10) scores.mean()  0.7290683605565638  RF_classifier.fit(x_train,y_train) y_final_pred=RF_classifier.predict(x_test) accuracy_score(y_test,y_final_pred)  0.7760416666666666
Out[64]: In [65]: In [66]: Out[66]: Out[67]:	<pre>#Gradient Boosting  from sklearn.model_selection import cross_val_score scores=cross_val_score(GB_classifier,x_train,y_train,scoring='accuracy',cv=10) scores.mean()  0.7394736842105264  GB_classifier.fit(x_train,y_train) y_final_pred=GB_classifier.predict(x_test) accuracy_score(y_test,y_final_pred)  0.7864583333333334</pre>
In [68]:	#After cross validation, Logistic regression still gives the highest accuracy.  Prediction  def diabetes_prediction (Pregnancies, Glucose, BloodPressure, SkinThickness, Insulin, BMI, DiabetesPedigreeFunction, Age):     pr=int(Pregnancies)     gluc=float(Glucose)     bp=float(BluoodPressure)     st=float(SkinThickness)     ins=float(Insulin)     bmi=float(BMI)     dpf=float(DiabetesPedigreeFunction)     age=int(Age)
in [71]: in [72]:	<pre>age=int(Age)  x=[[pr,gluc,bp,st,ins,bmi,dpf,age]] x=st_x.transform(x)  return LR_classifier.predict(x)  #Providing random values to the parameters to check our model  #Prediction 1 pred=diabetes_prediction(2,80,73,16,75,31.1,0.555,25)[0] if pred:     print("You have diabetes.") else:     print("You don't have diabetes")</pre>
n [73]:	<pre>You don't have diabetes  #Prediction 2 pred=diabetes_prediction(1,140,92,30,145,35,0.555,70)[0] if pred:     print("You have diabetes.") else:     print("You don't have diabetes")  You have diabetes.</pre>

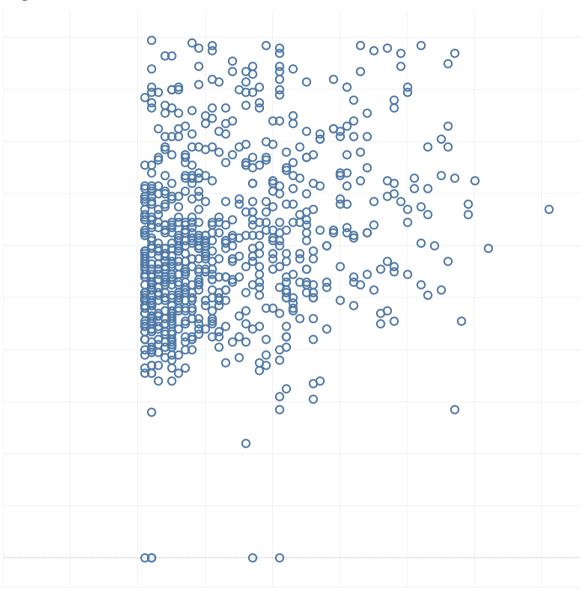
## Diabetic vs Non-Diabet-

ic

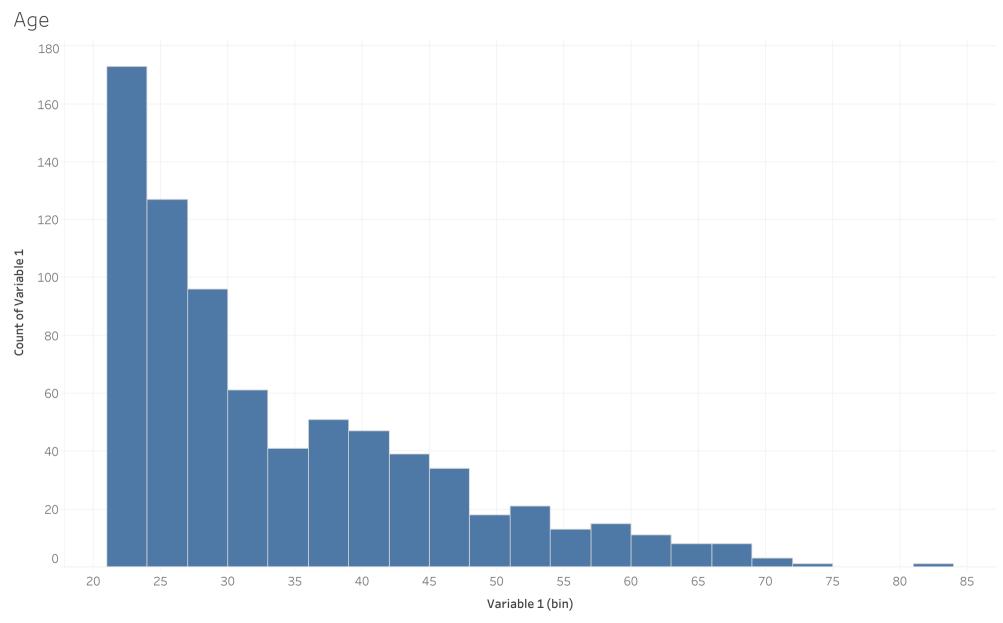


% of Total Count of
Outcome and
Dia/Non-Dia. Color
shows details about
Dia/Non-Dia. The
marks are labeled by
% of Total Count of
Outcome and
Dia/Non-Dia.

Age vs Glucose -Scatter Plots

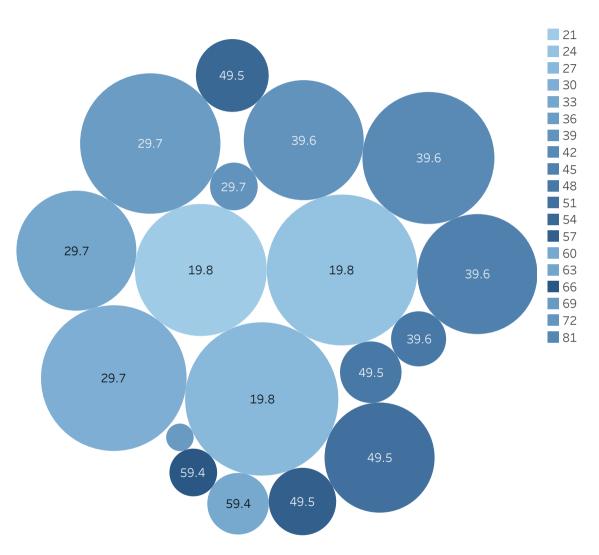


Variable 1 vs. Variable 2.

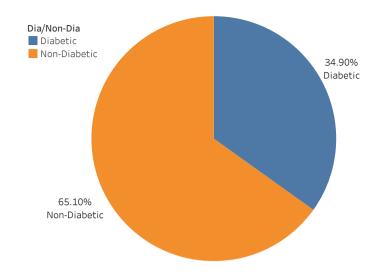


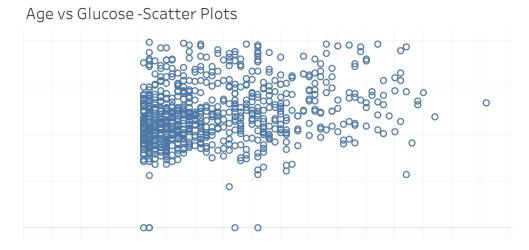
The trend of count of Variable 1 for Variable 1 (bin).

## Age vs Outcome - Bubble Charts



Variable 1 (bin) 2. Color shows details about Variable 1 (bin). Size shows sum of Outcome. The marks are labeled by Variable 1 (bin) 2.



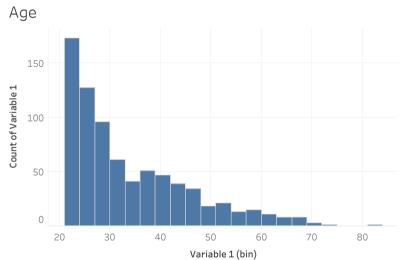


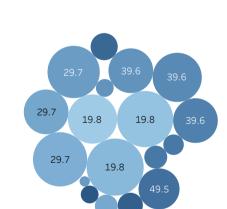
Variable 1

Variable 2

Glucose

Age





Age vs Outcome - Bubble Charts