0.

Means:

Glucose: 121.6867627785059

Blood Pressure: 72.40518417462484 Skin Thickness: 29.153419593345657

Insulin: 155.5482233502538 BMI: 32.45746367239099

Standard Deviations:

Glucose: 30.53564107280403

Blood Pressure: 12.382158210105263 Skin Thickness: 10.476982369987212

Insulin: 118.77585518724514 BMI: 6.924988332105907

1.

Covariance matrix:

	Glucose	BloodPressure	SkinThickness	Insulin	BMI
Glucose	932.425376	84.811985	74.025750	2127.014566	49.355133
BloodPressure	84.811985	153.317842	29.240422	145.553584	24.644988
SkinThickness	74.025750	29.240422	109.767160	230.676780	46.725661
Insulin	2127.014566	145.553584	230.676780	14107.703775	190.422831
BMT	49.355133	24.644988	46.725661	190.422831	47.955463

Correlations:

Glucose, Blood Pressure: 0.22319177824954217 Glucose, Skin Thickness: 0.22804322678186384

Glucose, Insulin: 0.581186208912165 Glucose, BMI: 0.23277051103551327

Blood Pressure, Skin Thickness: 0.22683906740782173

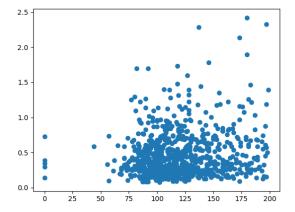
Blood Pressure, Insulin: 0.09827229945465545 Blood Pressure, BMI: 0.28923034040466644 Skin Thickness, Insulin: 0.18488842018975898

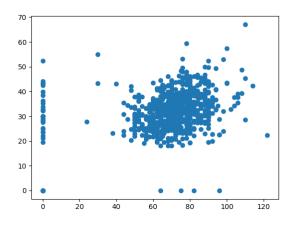
Skin Thickness, BMI: 0.6482139430923206

Insulin, BMI: 0.228050157416571

Because covariance matrices are not standardized, we can only generalize the relationships between the attributes. Since the main diagonal of the matrix is large for Insulin (when compared to its' standard deviation squared), we can say the standard deviation around the mean is largely spread. We can say this for other variables by comparing the standard deviation squared of each variable with the same variable in the matrix. The values here for correlation are generally not very high, representing far linear relationships between each other. Some strongly correlated relationships, compared to others, to note are glucose and insulin, as well as skin thickness and BMI. Again, these are the most correlated attributes, and have the best linear relationships.

2.



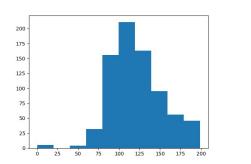


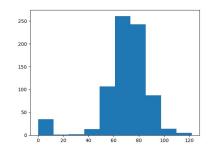
Left Attributes: 2, 7; Right Attributes: 3, 6

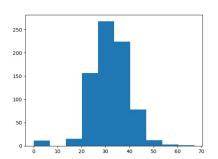
For attributes 2 and 7, the points appear to be most dense around value 100 to 125 on the x axis, with the values ranging from around 60 to 200. On the y axis, values range from around 0 to at most 1.5. For this scatter plot, there is no apparent trend or relationship between the two attributes.

For the scatter plot containing the attributes 3 and 6, the points are heavily centered around 60-80 on the x and 20-50 on the y axis. The trend here appears to be a linearly positive relationship.

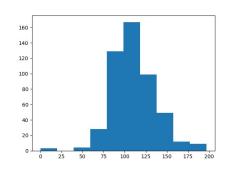
3.

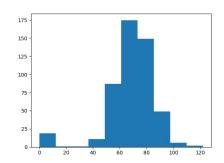


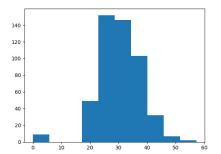




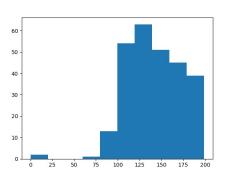
Histograms for attributes: 2, 3, 6, respectively, for the whole data set

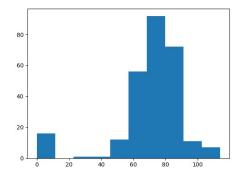


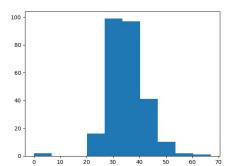




Histograms for attributes: 2, 3, 6, respectively, for outcome 0



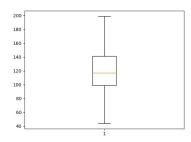


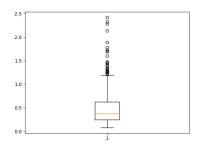


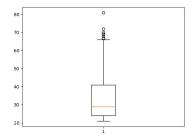
Histograms for attributes: 2, 3, 6, respectively, for outcome 1

Each histogram is generally Gaussian shaped, with sharper peaks for attributes 3 and 6 when the outcome was 0. All of the histograms displayed are unimodal. In this case, the differences are not very apparent. The only histogram that is skewed would be attribute 2 for outcome 0. I believe a larger bin size would allow for more information to be shown, while also revealing more differences between the classes.

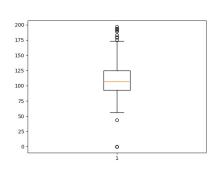
4.

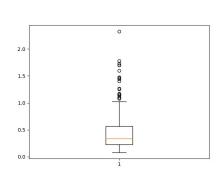


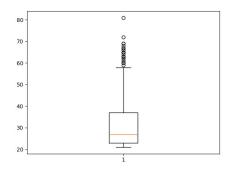




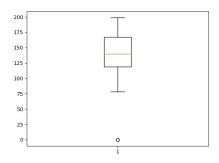
Boxplots for attributes: 2, 7, 8, respectively, whole data set

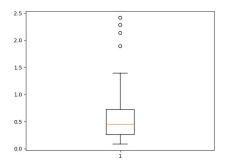


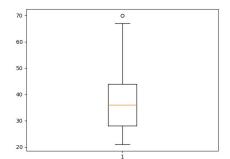




Boxplots for attributes: 2, 7, 8, respectively, for outcome 0

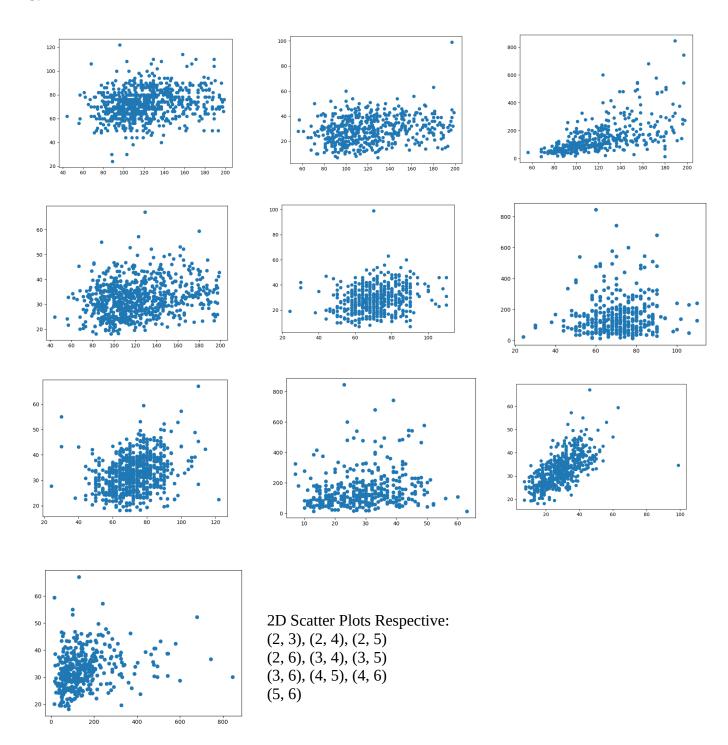




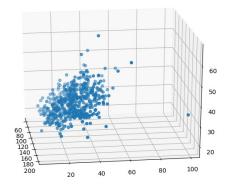


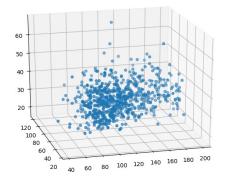
Boxplots for attributes: 2, 7, 8, respectively, for outcome 1 The data for each boxplot is centered at the median, and for the whole data set, the outliers are above the upper quartile. Also for the whole data set, the mean is generally skewed to the right. We can also note that these features also appear when the outcome is 0, as well as outcome 1. When looking at attribute 8, or age, we see that for those with diabetes have a higher median value, without much skew. On the other hand, those without had a lower median, with it being skewed to the right. There were also much more outliers present in those with an outcome of 0. Another attribute to observe between classes is the Glucose levels. Those with diabetes have a higher median, as well as higher 2nd and 3rd quartiles.

5.



The three plots that appear to have a linear and positive relationship are with attributes (2, 5), (4, 6), (5, 6). Each graph has obvious outliers, and the 3 graphs that are particularly centered are with attributes (2, 6), (3, 4), (3, 5), (3, 6). Although debatable, these graphs have less variance at the mean when compared to plots (2, 4), and (4, 5). Because these plots have greater variance around the mean, the two attributes has a weaker relationship with each other, and may be a bad predictor of the outcome. The amount of attributes present does make determining if the patient has diabetes difficult, and for other projects with many more attributes, there will also be an increased amount of graphs to observe.





3D Scatter Plots: Attributes respective (2, 4, 6), (2, 3, 6)

The difficulty with 3D plots is from multiple variables being displayed, cause issues observing the relationships in the 3D space. It can be seen that while both have some outliers, the plot on the left seems to have a positive relationship between the three attributes. On the other graph, the values are somewhat centered near the center of the space. They are generally spread out, due to high variance in these attributes. In this case, it was more difficult to describe the features of the 3 dimensional plot. On a positive note, a 3D space allows for stronger visualization of multivariate data, and may be helpful in determining whether relationships are present in this data.

Dep. Variable:	Outcome	R-square	d:		0.320		
Model:	OLS			0.313			
Method:	Least Squares		F-statistic:		44.61		
Date: T	ue, 18 Sep 2018	Prob (F-statistic): Log-Likelihood: AIC:		9.60e-59 -371.59 761.2			
Time:	10:31:38						
No. Observations:	767						
Df Residuals:	758	BIC:			803.0		
Df Model:	8						
Covariance Type:	nonrobust						
	coef	std err	t	P> t	[0.025	0.975]	
Intercept	-1.0215	0.104	-9.818	0.000	-1.226	-0.817	
Pregnancies	0.0207	0.005	4.093	0.000	0.011	0.031	
Glucose	0.0065	0.001	11.832	0.000	0.005	0.008	
BloodPressure	-0.0011	0.001	-0.861	0.389	-0.004	0.003	
SkinThickness	0.0001	0.002	0.066	0.947	-0.004	0.004	
Insulin	-8.58e-05	0.000	-0.461	0.645	-0.000	0.000	
BMI	0.0144	0.003	5.547	0.000	0.009	0.020	
DiabetesPedigreeFuncti	on 0.1288	0.044	2.924	0.004	0.042	0.21	
Age	0.0020	0.002	1.319	0.187	-0.001	0.005	
======================================	======================================	Durbin-Watson:		=======	1.964		
Prob(Omnibus):	0.000	Jarque-Bera (JB):		23.785			
Skew:	0.324	Prob(JB)	: ' '	6	.84e-06		
Kurtosis:	2.430	Cond. No.		1	.67e+03		
Warnings:							

	OLS Regres	sion Resul	ts			
Dep. Variable: Model: Method: Date: Time: No. Observations: Df Residuals: Df Model: Covariance Type:	Outcome OLS Least Squares ue, 18 Sep 2018 10:31:38 767 760 6 nonrobust	R-squared: Adj. R-squared: F-statistic: Prob (F-statistic): Log-Likelihood: AIC: BIC:		0.320 0.315 59.59 1.78e-60 -371.70 757.4 789.9		
	coef	std err	t	P> t	[0.025	0.975]
Intercept Pregnancies Glucose BloodPressure BMI DiabetesPedigreeFuncti Age	-1.0210 0.0208 0.0064 -0.0011 0.0144 on 0.1282 0.0020	0.104 0.005 0.001 0.001 0.002 0.044 0.002	-9.841 4.105 12.610 -0.839 6.457 2.915 1.311	0.000 0.000 0.000 0.401 0.000 0.004 0.190	-1.225 0.011 0 .005 -0.004 0.010 0.042 -0.001	-0.817 0.031 0.007 0.001 0.019 0.214 0.005
Omnibus: 31.727 Prob(Omnibus): 0.000 Skew: 0.320 Kurtosis: 2.441		Durbin-Watson: Jarque-Bera (JB): Prob(JB): Cond. No.		9 1		
<pre>[1] Standard Errors as [2] The condition numb strong multicollineari</pre>	er is large, 1.1	1e+03. Thi	s might indic			ecitied.

OLS Regression: Top picture is 1st run, bottom picture is 2nd run

Both models return similar results, even after removing blood pressure and skin thickness. The adjusted R-squared values only differed by 0.002, while other values also demonstrated very similar values. With regards to the coefficients, the DiabetesPedigreeFunction has the largest coefficients. It appears that when the coefficients are smaller or negative, they provide a larger p value, also providing less evidence against the null hypothesis. This can be seen for blood pressure, skin thickness, and insulin. On the other hand, when the coefficients are larger, the p value is generally smaller, equating to strong evidence against the null hypothesis. Because two values were removed due to high p values, and little changed in the R-squared values, it is obvious that their removal did not affect the result, and it may be safe to say that those attributes were not strongly related to the remaining attributes.

8.

Note: While trying to use Anaconda to install the graphviz package to show the decision trees, all of the dependencies became undetectable to my IDE, and thus I could not run any of the libraries afterwards like numpy/pandas/matplotlib/etc. Hours of troubleshooting were spent to no avail.

9.

In the linear regression, we observed that the removal of two attributes did not affect the results, thus we can conclude that the removed attributes were not strongly related to determining whether if the patient has diabetes. In the scatter plots, observing strong positive linear relationships between glucose and insulin, insulin and BMI, or skin thickness and BMI, could result in the patient having a higher chance of diabetes. The box plots determining age and diabetes were contradicting, but they did indicate higher glucose along with higher insulin levels, both pertaining to greater chances of diabetes. Overall, only a few attributes, such as the aforementioned insulin, glucose provides more indication of diabetes. It could also be seen that in the regression, that those with lower and negative correlations equated to less evidence against diabetes, while the opposite values in correlations provided greater evidence of diabetes.