In [1]:	<pre>import numpy as np import pandas as pd</pre>									
In [2]:										
In [3]:	df.head()									
Out[3]:	Pregna 0	ancies Glud	cose BloodPi	ressure SkinThic	kness Insu 35	0 33.6	betes Pedigree	Function 0.627	Age 50	Outcome 1
	2	8	85 183	66 64 66	0	0 26.60 23.394 28.1		0.351	31 32	1
	4	0	137	40		168 43.1		0.167 2.288	33	1
In [4]:	df.info		core.frame.	DataFrame'>						
	RangeInd Data cod # Cod	dex: 768	entries, (otal 9 colu) to 767	l Count	Dtype				
	1 Gli 2 Blo	egnancies ucose oodPressu inThickne	ıre	768 non- 768 non- 768 non- 768 non-	-null -null	<pre>int64 int64 int64 int64</pre>				
	5 BM: 6 Dia 7 Age	abetesPec e	ligreeFunct	768 non-	-null -null -null	int64 float64 float64 int64				
	dtypes:	tcome float64 usage: 54	(2), int64	768 non-	-null	int64				
In [5]: Out[5]:	df.shar									
In [6]:	df.desc	cribe()								
Out[6]:		regnancies 768.000000	Glucose 768.000000	BloodPressure S	kinThicknes		BMI 768.000000	Diabetes		e Functio 768.00000
	mean std	3.845052 3.369578	120.894531 31.972618	69.105469 19.355807	20.53645 15.95221		31.992578 7.884160			0.47187 0.33132
	min 25%	0.000000	0.000000 99.000000	0.000000 62.000000	0.00000	0.000000	0.000000 27.300000			0.07800 0.24375
	50% 75%	3.000000 6.000000	117.000000	72.000000 80.000000	23.00000 32.00000	0 127.250000	32.000000 36.600000			0.37250
In [7]:	df.isn	17.000000	199.000000	122.000000	99.00000	0 846.000000	67.100000			2.42000
Out[7]:	Pregnand	cies	. ()	0						
	BloodPre SkinThio Insulin BMI	ckness	Function	0 0 0 0						
	Age Outcome dtype:	_	or unceron	0						
In [8]:		<pre>clearn.pr = MinMax</pre>		g import MinN	MaxScaler					
In [9]:	df.drop	o(['SkinT	hickness']	,inplace = Tr	rue, axis	= 1)				
In [10]: Out[10]:	df	unancies G	ilucosa Rloor	dPressure Insulir	s RMI Dia	ahatas Padiaraa	Function Ag	e Outco	me	
out[IV].	0	6	148 85	72 () 33.6) 26.6	abetesi edigiree	0.627 5 0.351 3	0	1 0	
	2	8	183 89	64 (23.3		0.672 3 0.167 2	2	1	
	4	0		40 168 	3 43.1		2.288 3		1	
	763 764	10 2	101 122	70 (32.9			7	0	
	765 766	1	121	60 (2 26.2		0.245 3 0.349 4	7	1	
	767 768 rows :	1 × 8 columr	93 ns	70 (30.4		0.315 2	3	0	
In [11]:	<pre>def remove_outliers_iqr(df): # Calculate the first quartile (Q1) and third quartile (Q3) q1 = np.percentile(df, 25)</pre>									
	q3 # (= np.per	centile (df		ge (IQR)					
	# 1 lov	wer_bound	ne lower and = q1 - 1.		ds for ou	ıtliers				
	# I	- Remove ou		f lower_bound	d <= x <=	upper_bour	nd]			
	ret	curn df								
In [12]: In [13]:		seaborn								
Out[13]:			corr(), an	not =True , fmt	c='.2f')					
		_		0.14 -0.07 0.02 0.15 0.33 0.22	_					
			ire - 0.14 0.15 lin0.07 0.33	1.00 0.09 0.28 0.09 1.00 0.20	0.04 0.24	- ().6			
	DiabetesPe	digreeFuncti	on0.03 0.14	0.28 0.20 1.00 0.04 0.19 0.14 0.24 -0.04 0.04	1.00 0.01	3 0.17).4			
		Outcon	ne - 0.22 0.47	0.07 0.13 0.29	0.17 0.24	4 1.00 - 0).0			
			Pregnancies Glucose	BloodPressure Insulin BMI	DiabetesPedigreeFunction Age	Outcome				
					DiabetesPec					
In [14]:	X = df	drop('Ou	tcome', ax	ion import to	rain_test	_split				
In [15]:	y = df	['Outcome	.']							
In [16]:		_		r.transform()	ζ)					
In [17]:	<pre>X = standardised_data y = df['Outcome']</pre>									
In [18]:	<pre>X_train, X_test, y_train, y_test = train_test_split(X,y,test_size=0.2, random_state=0 from sklearn.ensemble import RandomForestClassifier</pre>									
In [19]:	<pre>rf = RandomForestClassifier(n_estimators=20, criterion='entropy', random_state=42) rf.fit(X train, y train)</pre>									
Out[19]:	<pre>rf.fit(X_train, y_train) RandomForestClassifier(criterion='entropy', n_estimators=20, random_state=42)</pre>									
In [20]:	<pre>y_pred = rf.predict(X_test)</pre>									
In [21]:	accurac	cy_score(y_test, y_	rt accuracy_s pred)	score					
Out[21]: In [22]:		584415584 DOM FORES								
	'RANDOM	FOREST.								