In [44]:	#Amarendra Shendkar #MIS 111915013										
	<pre>import numpy as np import pandas as pd import matplotlib.pyplot as plt import seaborn as sns</pre>										
In [45]:	<pre>df = pd.read_csv('/input/drug-classification/drug200.csv')</pre>										
In [46]:	df.head()										
Out[46]:	<b>Age So 0</b> 23	ex BP Ch	olesterol Na_	to_K Drug 5.355 DrugY	_						
		M LOW		3.093 drugC ).114 drugC							
	<ul><li>3 28</li><li>4 61</li></ul>	F LOW		7.798 drugX 3.043 DrugY							
In [47]:	df.info()										
	<pre><class 'pandas.core.frame.dataframe'=""> RangeIndex: 200 entries, 0 to 199 Data columns (total 6 columns): # Column Non-Null Count Dtype</class></pre>										
	# Column Non-Null Count Dtype										
In [48]:	df.isna().sum()										
Out[48]:	Age 0 Sex 0 BP 0 Cholesterol 0 Na_to_K 0 Drug 0 dtype: int64										
In [49]:	<pre>df = df.dropna() df.info()</pre>										
	Int64Inde	class 'pandas.core.frame.DataFrame'> nt64Index: 200 entries, 0 to 199 ata columns (total 6 columns): # Column Non-Null Count Dtype									
	0 Age 200 non-null int64 1 Sex 200 non-null object										
	3 Cho. 4 Na_1	lesterol 200 to_K 200	non-null non-null	object float64							
Tn [EQ].	dtypes: float64(1), int64(1), object(4) memory usage: 10.9+ KB										
In [50]: Out[50]:	_	dummies(df)  Na_to_K Sex_F	Sex_M BP_H	IIGH BP_LO	W BP_NORM	AL Cholesterol_HIG	H Cholesterol_NORM	AL Drug_Drug	Y Drug_drug	A Drug_	
	<ul><li>0 23</li><li>1 47</li></ul>	25.355 1 13.093 0	0	1	0		1	-		0	
	<ul><li>2 47</li><li>3 28</li></ul>	10.114 0 7.798 1	1	0	1	0	1			0	
	<b>4</b> 61	18.043 1	0	0	1		1 			0	
	<ul><li>195 56</li><li>196 16</li></ul>	11.567 1 12.006 0	0	0	1	0	1			0	
	<ul><li>197 52</li><li>198 23</li></ul>	9.894 0 14.020 0	1	0	0	1	1 0			0	
	<b>199</b> 40 200 rows ×	11.349 1 14 columns	0	0	1	0	0	1	0	0	
In [51]:	<pre>X = pd.get_dummies(df.drop('Drug',axis=1),drop_first=True) y = df['Drug']</pre>										
In [52]:	<pre>X.head()</pre>										
Out[52]:	y.head()  0 DrugY										
	<pre>1 drugC 2 drugC 3 drugX 4 DrugY</pre>										
In [53]:	<pre>Name: Drug, dtype: object  from sklearn.model_selection import train_test_split X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3, random_state=101)</pre>										
In [54]:	<pre>from sklearn.preprocessing import StandardScaler</pre>										
In [55]:	<pre>scaler = StandardScaler()</pre>										
In [56]:	<pre>scaled_X_train = scaler.fit_transform(X_train) scaled_X_test = scaler.transform(X_test)</pre>										
In [50].	<pre>from sklearn.neighbors import KNeighborsClassifier</pre> knn model = KNeighborsClassifier(n neighbors=10)										
In [58]:	<pre>knn_model = KNeighborsClassifier(n_neighbors=10) knn_model.fit(scaled_X_train,y_train)</pre>										
Out[58]:	KNeighborsClassifier(n_neighbors=10)										
In [59]: Out[59]:	<pre>full_test = pd.concat([X_test,y_test],axis=1) len(full_test)</pre>										
In [60]:	y_pred =	<pre>y_pred = knn_model.predict(scaled_X_test)</pre>									
In [61]:	from skl	<pre>from sklearn.metrics import classification_report,confusion_matrix,accuracy_score</pre>									
In [62]:		accuracy_score(y_test,y_pred) 0.76666666666667									
Out[62]: In [63]:		0.7666666666666666666666666666666666666									
Out[63]:	array([[21, 1, 0, 0, 4],										
Tn [C4]	[ 6, 0, 0, 3, 0], [ 2, 0, 0, 0, 11]])										
In [64]:	print(classification_report(y_test,y_pred))  precision recall f1-score support										
	d: d:	rugY 0. rugA 0. rugB 1.	75 1.0 00 0.8	0. 0.	86 91	6 6 6					
	d: accu:		73 0.8	0.	79 1 77 6	9 3					
	macro weighted					0					