

Applied machine learning Group assignment 3

Team members-Group 8:

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Part 1: calculation

Solution:

a) d(a,b) denotes the Euclidean distance between a and b.

It is obtained directly from the distance matrix or calculated as follows:

$$d(a,b) = \sqrt{(xb - xa)^2 + (yb - ya)^2}$$

The initial centroids \rightarrow A2, A4

For A1

$$d(A1, A2) = \sqrt{(5-2)^2 + (8-5)^2} = 3\sqrt{2}$$

$$d(A1, A4) = \sqrt{(1-2)^2 + (2-5)^2} = \sqrt{10}$$
 -> Smaller

 $A1 \in A4$, $A1 \in Cluster2$

For A2

$$d(A1, A2) = \sqrt{(5-5)^2 + (8-8)^2} = Zero$$

$$d(A1, A4) = \sqrt{(1-5)^2 + (2-8)^2} = 2\sqrt{13}$$

 $A2 \in A2$, $A2 \in Cluster1$

For A3

$$d(A3, A2) = \sqrt{(5-7)^2 + (8-5)^2} = \sqrt{13}$$
 -> Smaller

$$d(A3, A4) = \sqrt{(1-7)^2 + (2-5)^2} = 3\sqrt{5}$$

 $A3 \in A2$, $A3 \in Cluster1$

For A4

$$d(A4, A2) = \sqrt{(5-1)^2 + (8-2)^2} = 2\sqrt{13}$$

$$d(A4, A4) = \sqrt{(1-1)^2 + (2-2)^2} = Zero$$

 $A4 \in A4$, $A4 \in Cluster2$

For A5

$$d(A5,A2) = \sqrt{(5-4)^2 + (8-9)^2} = \sqrt{2} - \text{Smaller}$$

$$d(A5,A4) = \sqrt{(1-4)^2 + (2-9)^2} = \sqrt{58}$$

 $A5 \in A2$, $A5 \in Cluster1$

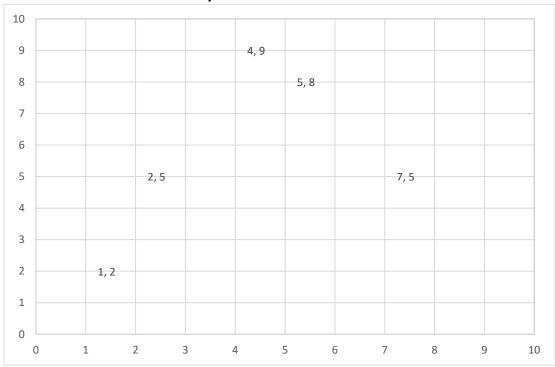
New clusters:-

The Mean for each clusters :-

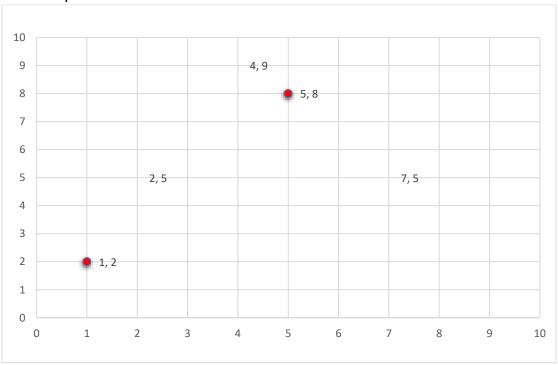
Cluster 1 =
$$\left(\frac{5+7+4}{3}, \frac{8+5+9}{3}\right) = \left(\frac{16}{3}, \frac{22}{3}\right)$$

Cluster 2 =
$$\left(\frac{1+2}{2}, \frac{5+2}{2}\right) = \left(\frac{3}{2}, \frac{7}{2}\right)$$

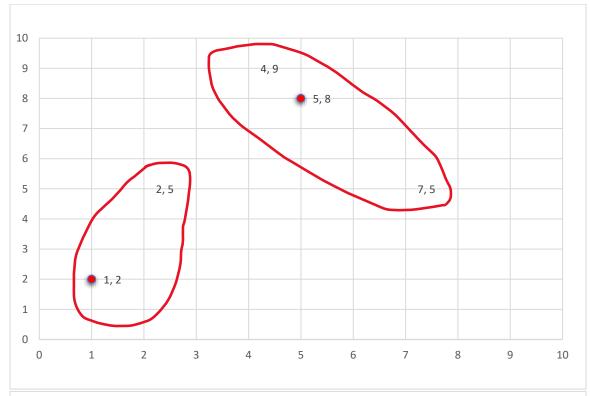
b)
The locations for the data points:



The red points are the initial centroids.



The clusters of cluster 1 and cluster 2:





The X points are the centroid for each clusters.

c)

Calculate the silhouette score and WSS score.

1. Calculate WSS score.

For calculating WSS the equation equal:

$$WSS = \sum_{i=1}^{m} (x_i - c_i)$$

WSS =
$$(5-5.3)^2 + (8-7.3)^2 + (7-5.3)^2 + (5-7.3)^2 + (4-5.3)^2 + (9-7.3)^2 + (2-1.5)^2 + (5-3.5)^2 + (1-1.5)^2 + (2-3.5)^2$$

= 18.34

2. Calculate silhouette score.

For calculating silhouette the equation equal:

$$S(i) = \frac{b(i) - a(i)}{\max\{a(i), b(i)\}}$$

According to cluster 1:

<mark>A2 :</mark>

•
$$d(A2,A3) = \sqrt{(5-7)^2 + (8-5)^2} = 3.6$$

•
$$d(A2,A5) = \sqrt{(5-4)^2 + (8-9)^2} = 1.4$$

$$\bullet \quad a(A2) = \frac{d(A2,A3) + d(A2,A5)}{2} = \frac{3.6 + 1.4}{2} = 2.5$$

• d(A2,A1)=
$$\sqrt{(5-2)^2 + (8-5)^2}$$
 = 4.2

$$d(A2,A4) = \sqrt{(5-1)^2 + (8-2)^2} = 7.2$$

•
$$b(A2) = \frac{4.2 + 7.2}{2} = 5.7$$

$$S(A2) = \frac{b(A2) - a(A2)}{\max\{a(A2), b(A2)\}} = \frac{5.7 - 2.5}{5.7} = 0.56$$

<mark>A3 :</mark>

•
$$d(A3,A2) = \sqrt{(5-7)^2 + (8-5)^2} = 3.6$$

• d(A3,A5)=
$$\sqrt{(7-4)^2 + (5-9)^2} = 5$$

$$\bullet \quad a(A3) = \frac{d(A3,A2) + (A3,A5)}{2} = \frac{3.6 + 5}{2} = 4.3$$

•
$$d(A3,A1) = \sqrt{(7-2)^2 + (5-5)^2} = 5$$

•
$$d(A3,A4) = \sqrt{(7-1)^2 + (5-2)^2} = 6.7$$

• b(A3) =
$$\frac{5+6.7}{2}$$
 = 5.85

$$> S(A3) = \frac{b(A3) - a(A3)}{\max\{a(A3), b(A3)\}} = \frac{5.85 - 4.3}{5.85} = 0.26$$

<mark>A5:</mark>

•
$$d(A5,A2) = \sqrt{(4-5)^2 + (9-8)^2} = 1.4$$

•
$$d(A5,A3) = \sqrt{(4-7)^2 + (9-5)^2} = 5$$

■
$$a(A5) = \frac{d(A5,A2) + (A5,A3)}{2} = \frac{1.4+5}{2} = 3.2$$

•
$$d(A5,A1) = \sqrt{(4-2)^2 + (9-5)^2} = 4.4$$

•
$$d(A5,A4) = \sqrt{(4-1)^2 + (9-2)^2} = 7.6$$

■ b(A5) =
$$\frac{4.4 + 7.6}{2}$$
 = 6

$$> S(A5) = \frac{b(A5) - a(A5)}{\max\{a(A5), b(A5)\}} = \frac{6 - 3.2}{6} = 0.46$$

According to cluster 2:

<mark>A1:</mark>

•
$$d(A1,A4) = \sqrt{(2-1)^2 + (5-2)^2} = 3.16$$

$$\bullet \quad a(A1) = \frac{d(A1,A4) + d(A1,A4)}{1} = \frac{3.16}{1} = 3.16$$

•
$$d(A1,A2) = \sqrt{(5-2)^2 + (8-5)^2} = 4.2$$

•
$$d(A1,A3) = \sqrt{(7-2)^2 + (5-5)^2} = 5$$

• d(A5,A1)=
$$\sqrt{(4-2)^2 + (9-5)^2}$$
 = 4.4

•
$$b(A1) = \frac{4.2 + 5 + 4.4}{3} = 4.53$$

$$> S(A1) = \frac{b(A1) - a(A1)}{\max\{a(A1), b(A1)\}} = \frac{4.53 - 3.16}{4.53} = 0.302$$

<mark>A4 :</mark>

•
$$d(A1,A4) = \sqrt{(2-1)^2 + (5-2)^2} = 3.16$$

$$\bullet \quad a(A4) = \frac{d(A1,A4) + d(A1,A4)}{1} = \frac{3.16}{1} = 3.16$$

•
$$d(A4,A2) = \sqrt{(5-1)^2 + (8-2)^2} = 7.2$$

$$d(A4,A3) = \sqrt{(7-1)^2 + (5-2)^2} = 6.7$$

$$d(A4,A5) = \sqrt{(4-1)^2 + (9-2)^2} = 7.6$$

b
$$(A4) = \frac{7.2 + 6.7 + 7.6}{3} = 7.166$$

$$> S(A4) = \frac{b(A4) - a(A4)}{\max\{a(A4), b(A4)\}} = \frac{7.166 - 3.16}{7.166} = 0.55$$

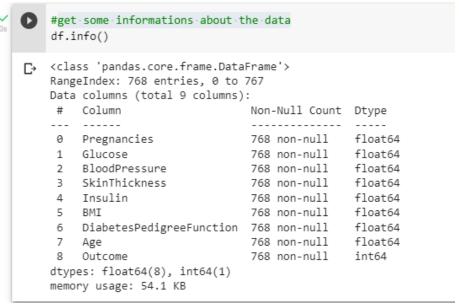
AVG S(i) =
$$\frac{0.302 + 0.55 + 0.46 + 0.26 + 0.56}{5}$$
 = 0.43

Part 2: programming

Import important Libraries and load the dataset

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Outcome	0
0	0.411765	0.623116	0.573770	0.333333	0.254137	0.380030	0.035440	0.266667	0	
1	0.294118	0.542714	0.590164	0.434343	0.088652	0.538003	0.078992	0.200000	0	
2	0.058824	0.437186	0.491803	0.373737	0.088652	0.554396	0.184031	0.016667	0	
3	0.058824	0.723618	0.672131	0.464646	0.212766	0.687034	0.109735	0.416667	1	
4	0.058824	0.557789	0.508197	0.131313	0.215130	0.357675	0.025619	0.033333	0	

get some information about the data and describe it:

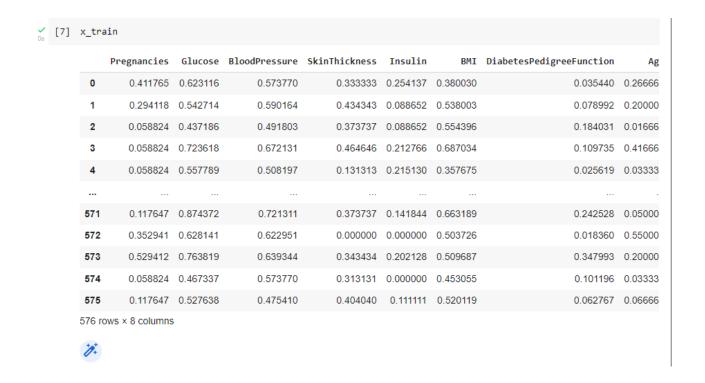


	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	${\tt DiabetesPedigreeFunction}$	Age	Outcome
count	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000
mean	0.226180	0.607510	0.566438	0.207439	0.094326	0.476790	0.168179	0.204015	0.348958
std	0.198210	0.160666	0.158654	0.161134	0.136222	0.117499	0.141473	0.196004	0.476951
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000
25%	0.058824	0.497487	0.508197	0.000000	0.000000	0.406855	0.070773	0.050000	0.000000
50%	0.176471	0.587940	0.590164	0.232323	0.036052	0.476900	0.125747	0.133333	0.000000
75%	0.352941	0.704774	0.655738	0.323232	0.150414	0.545455	0.234095	0.333333	1.000000
max	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000	1.000000

1- Split the dataset to train and test with random_state = 0

```
from sklearn.model_selection import train_test_split
x_train , x_test ,y_train ,y_test = train_test_split(X,y,test_size = 0.25 ,random_state=0,shuffle=False)
```

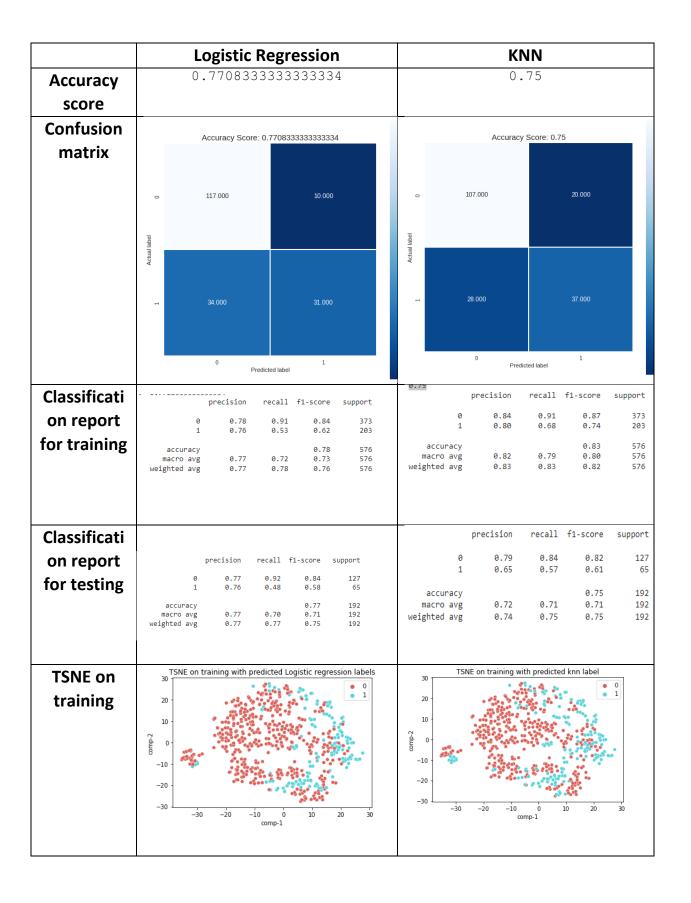
X train will be:

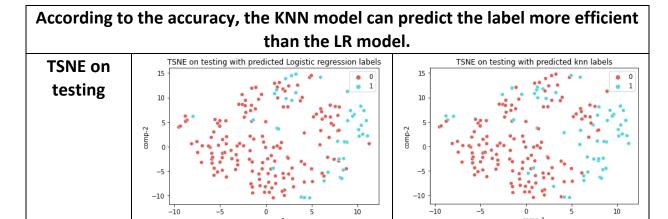


2- Train Logistic regression model and KNN model:

Logistic regression estimates the probability of an event occurring, such as voted or didn't vote, based on a given dataset of independent variables. Since the outcome is a probability, the dependent variable is bounded between 0 and 1.

K Nearest Neighbor algorithm falls under the Supervised Learning category and is used for classification (most commonly) and regression. It is a versatile algorithm also used for imputing missing values and resampling datasets. As the name (K Nearest Neighbor) suggests it considers K Nearest Neighbors (Data points) to predict the class or continuous value for the new Datapoint.

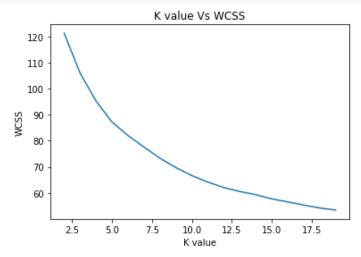




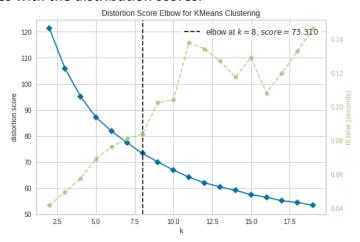
Accosrding to the testing accuracy, the LR model can predict the actual labels slightly than the KNN model.

1- compare between the values of K and silhouette scores:

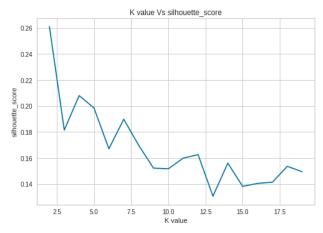
```
[31] from sklearn.cluster import KMeans
    from sklearn.metrics import silhouette_score
    wcss=[]
    scores =[]
    for i in range(2,20):
        kmean = KMeans(n_clusters=i)
        kmean.fit(X)
        wcss.append(kmean.inertia_)
        y_pred= kmean.predict(X)
        scores.append(silhouette_score(X,y_pred))
    plt.xlabel('K value')
    plt.ylabel('WCSS')
    sns.lineplot(x=range(2,20),y=wcss,).set(title="K value Vs WCSS")
```



Visualize the k values with the distribution scores:

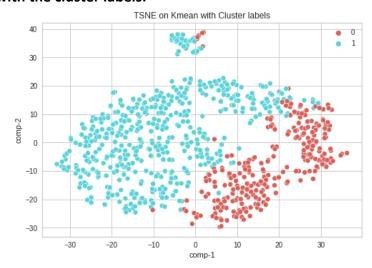


Visualize the k values with the silhouette scores:



So, the optimal number of k based on the silhouette score is when k=2

TSNE on K-means with the cluster labels:



As we can see from the TSNE plot, the K-Means algorithm on can separate between the two cluster labels well.

3- Apply dimensionality reduction methods:

1- Apply the principle component analysis(PCA):

Principal component analysis, or PCA, is a statistical procedure that allows you to summarize the information content in large data tables by means of a smaller set of "summary indices" that can be more easily visualized and analyzed.

```
from sklearn.decomposition import PCA
scores_lr = []
scores_knn = []
for i in range(2,8):
    pca= PCA(n_components=i)
    x_train_pca = pca.fit_transform(x_train)
    x_test_pca = pca.transform(x_test)
    lr = LogisticRegression()
    lr.fit(x_train_pca,y_train)

    y_pred_lr = lr.predict(x_test_pca)
    scores_lr.append(accuracy_score(y_pred_lr,y_test))
    knn = KNeighborsClassifier()
    knn.fit(x_train_pca,y_train)
    y_pred_knn = knn.predict(x_test_pca)
    scores_knn.append(accuracy_score(y_pred_knn,y_test))
```

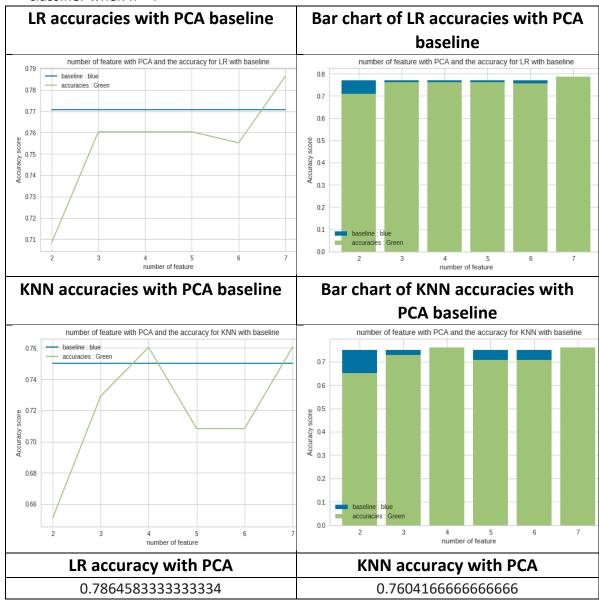
Scores of logistic regression with PCA	Scores of KNN with PCA
[0.70833333333334,	[0.6510416666666666666666666666666666666666
0.7604166666666666666666666666666666666666	0.7291666666666666666666666666666666666666
0.7604166666666666666666666666666666666666	0.7604166666666666666666666666666666666666
0.7604166666666666666666	0.708333333333334,
0.755208333333334,	0.708333333333334,
0.7864583333333333]	0.760416666666666]

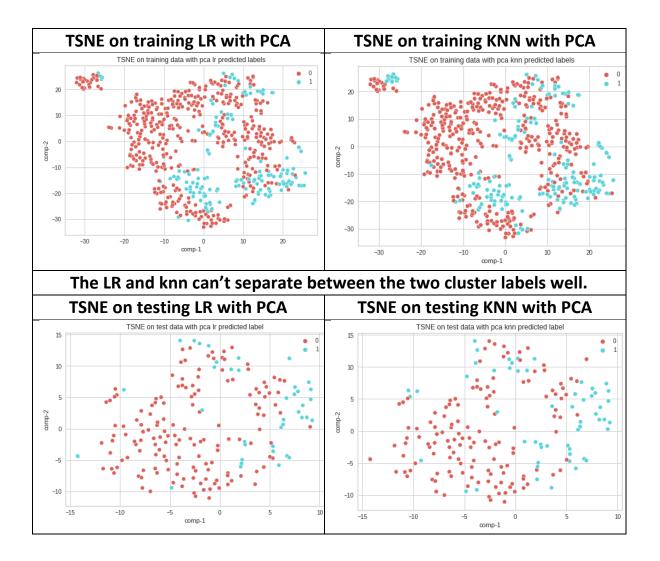
So, n =7 is the best number number of coponebt that achieve high accuarcy on LR and KNN

```
(2, 0.7083333333333334, 0.6510416666666666)
(3, 0.7604166666666666, 0.729166666666666)
(4, 0.7604166666666666, 0.7604166666666666)
(5, 0.7604166666666666, 0.7083333333333333)
```

```
(6, 0.7552083333333334, 0.70833333333333334)
(7, 0.78645833333333334, 0.760416666666666)
```

1- Plot the Number of Components-Accuracy graph with baseline performances for each classifier when n = 7



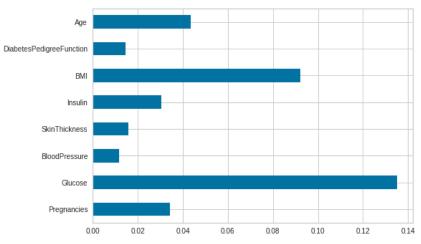


4- Apply feature selection methods:

1- **Apply filter method:** Filter methods pick up the intrinsic properties of the features measured via univariate statistics instead of cross-validation performance. These methods are faster and less computationally expensive than wrapper methods. When dealing with high-dimensional data, it is computationally cheaper to use filter methods.

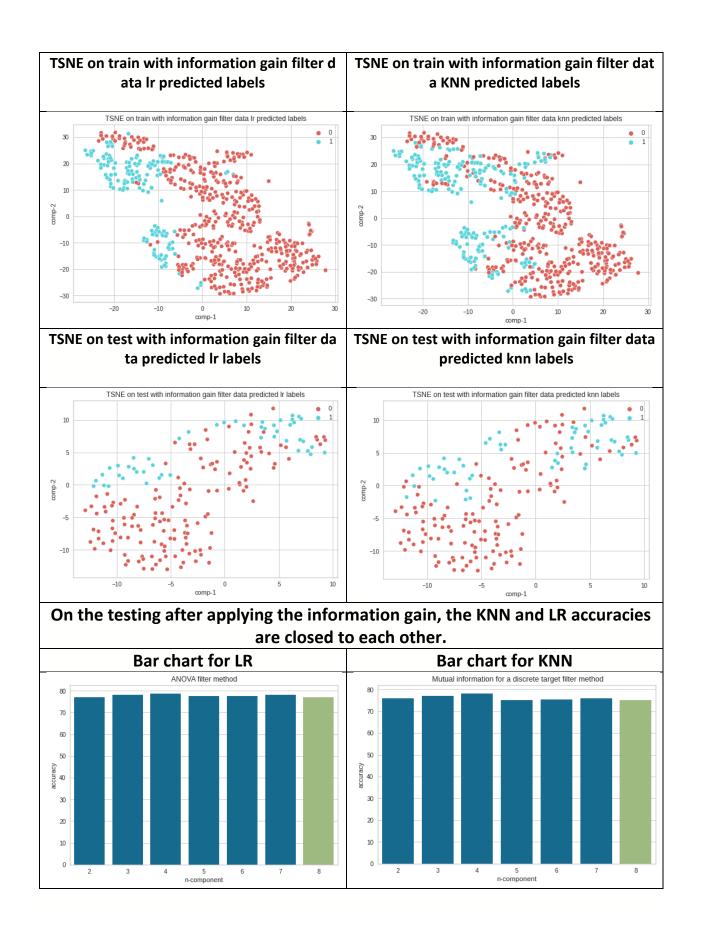
Apply information gain: Information gain calculates the reduction in entropy from the transformation of a dataset. It can be used for feature selection by evaluating the Information gain of each variable in the context of the target variable.

```
# information Gain
from traitlets.traitlets import ForwardDeclaredInstance
from sklearn.feature_selection import mutual_info_classif
importance = mutual_info_classif(X,y)
feat_importance = pd.Series(importance,df.columns[0:len(df.columns)-1])
feat_importance.plot(kind='barh')
```



based on information gain we will select the highest four feature that has information gain.

Accuracy	of LR wi	th info	rmation	gain	Accuracy of KNN with information gain				
0.7864583333333334					0.765625				
Classificat	-		•	Rwith	Classification report on training KNN with				
	informa	ation ga	nin			inform	ation gai	in	
	precision	recall	f1-score	support		precision	recall	f1-score	support
0	0.77	0.90	0.83	373	6	0.85	0.85	0.85	373
1	0.73	0.50	0.59	203	1	0.72	0.71	0.72	203
accuracy			0.76	576	accuracy	,		0.80	576
macro avg	0.75	0.70	0.71	576	macro avg		0.78	0.78	576
weighted avg	0.75	0.76	0.75	576	weighted avg	,	0.80	0.80	576
Classifica	tion repo	rt on te	esting LR	with	Classifica	ition repoi	rt on test	ting KNN	with
	informa	ation ga	nin			inform	ation gai	in	
	precision	recall	f1-score	support		precision	recall f	f1-score	support
0	0.80	0.91	0.85	127	0	0.80	0.86	0.83	127
1	0.75	0.55	0.64	65	1	0.68	0.58	0.63	65
accuracy			0.79	192	accuracy			0.77	192
macro avg	0.77	0.73	0.74	192	macro avg	0.74	0.72	0.73	192
weighted avg	0.78	0.79	0.78	192	weighted avg	0.76	0.77	0.76	192



According to the accuracies, the LR model is better than KNN model with information gain.

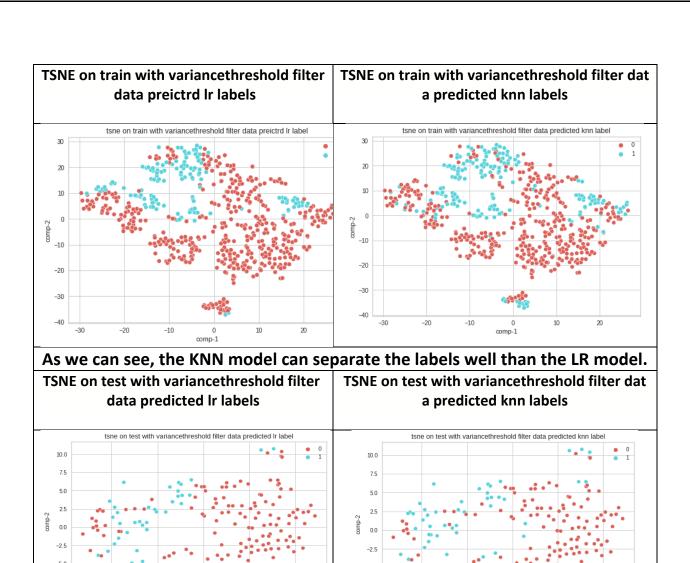
Apply variance threshold:

The variance threshold is a simple baseline approach to feature selection. It removes all features which variance doesn't meet some threshold. By default, it removes all zero-variance features, i.e., features that have the same value in all samples. We assume that features with a higher variance may contain more useful information, but note that we are not taking the relationship between feature variables or feature and target variables into account, which is one of the drawbacks of filter methods.

```
# Variance threshold
from sklearn.feature_selection import VarianceThreshold
v = VarianceThreshold(threshold=0.02)
v.fit(X)
v.get_support()
```

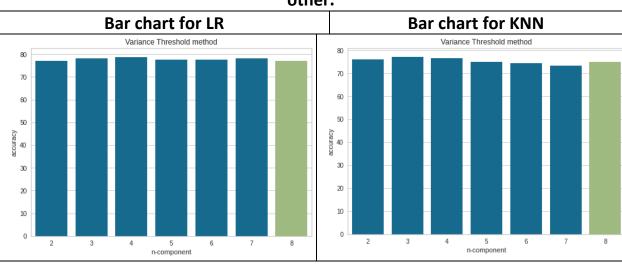
array([True, True, True, True, False, False, False, True])

Accur	acy of L	R with	varian	ce	Accur	acy of KN	NN with	n varian	ce
	thre	shold			threshold				
0	0.770833333333334					0.78125			
Classifica	ation rep	ort fo	r traini	ng LR	Classification report for training KNN				
_ 017700333333	precision	recall	f1-score	support		precision	recall	f1-score	support
0	0.75	0.90	0.82	373	0	0.82	0.88	0.85	373
1	0.71	0.46	0.56	203	1	0.75	0.66	0.70	203
accuracy			0.74	576	accuracy			0.80	576
macro avg	0.73	0.68	0.69	576	macro avg	0.79	0.77	0.77	576
weighted avg	0.74	0.74	0.73	576	weighted avg	0.80	0.80	0.80	576
Classific	ation re	port fo	r testir	ng LR	Classific	ation rep	ort for	testing	KNN
	precision	recall	f1-score	support		precision	recall	f1-score	support
0	0.77	0.92	0.84	127	0	0.83	0.84	0.84	127
1	0.76	0.48	0.58	65	1	0.68	0.66	0.67	65
accuracy			0.77	192					
macro avg	0.77	0.70	0.71	192	accuracy			0.78	192
weighted avg	0.77	0.77	0.75	192	macro avg	0.76	0.75	0.75	192
					weighted avg	0.78	0.78	0.78	192



When testing the two models, we can see that they are too close from each other.

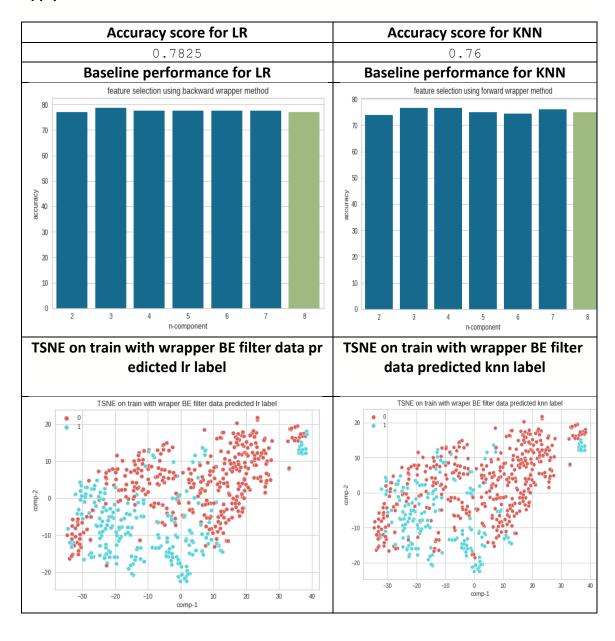
-7.5 -10.0 -5.0

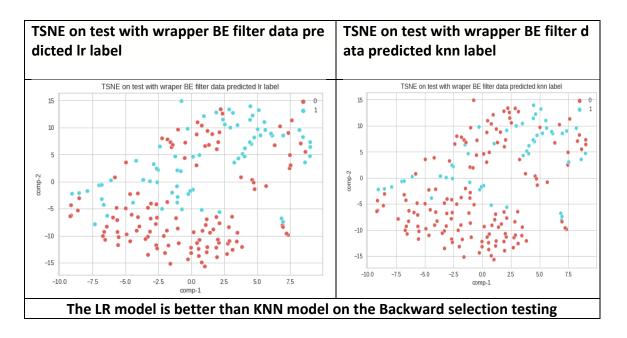


Apply wrapper methods:

Wrappers require some method to search the space of all possible subsets of features, assessing their quality by learning and evaluating a classifier with that feature subset. The feature selection process is based on a specific machine learning algorithm that we are trying to fit on a given dataset. It follows a greedy search approach by evaluating all the possible combinations of features against the evaluation criterion. The wrapper methods usually result in better predictive accuracy than filter methods.

1- Apply backward feature selection:





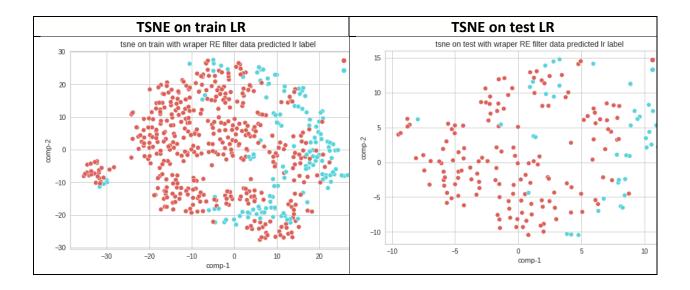
Apply Recursive feature elimination on LR:

the goal of recursive feature elimination (RFE) is to select features by recursively considering smaller and smaller sets of features. First, the estimator is trained on the initial set of features and the importance of each feature is obtained either through a coef_ attribute or through a feature_importances_ attribute.

Accuracy is: 0.7760416666666666

	precision	recall	f1-score	support
0	0.78	0.91	0.84	373
1	0.76	0.53	0.62	203
accuracy			0.78	576
macro avg	0.77	0.72	0.73	576
weighted avg	0.77	0.78	0.76	576
	precision	recall	f1-score	support
0	precision 0.78	recall 0.92	f1-score 0.84	support
0 1				
_	0.78	0.92	0.84	127

can't apply RFE with KNN because doesn't have a feature importance attribute

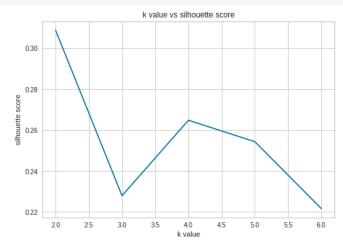


5- Choose the best number of cluster for k-means clustering algorithm on the processed data, using the best features:

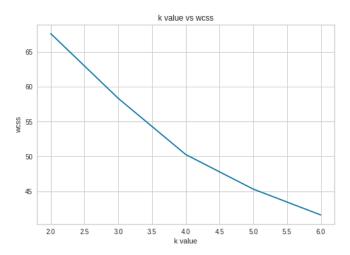
Silhouette score vs number of clusters:

```
scores =[]
for i in range(2,7):
    kmean = KMeans(n_clusters=i)
    y_pred = kmean.fit_predict(x_train_b_knn)
    wcss.append(kmean.inertia_)
    scores.append(silhouette_score(x_train_b_knn,y_pred))

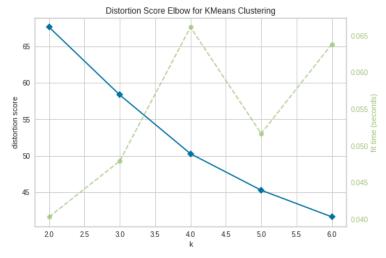
plt.plot(range(2,7),scores)
plt.title('k value vs silhouette score')
plt.xlabel('k value')
plt.ylabel('silhouette score')
```



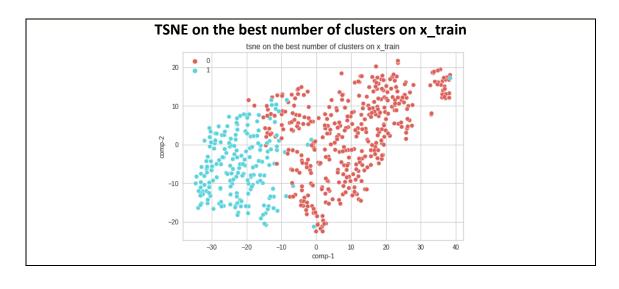
k value vs wcss:

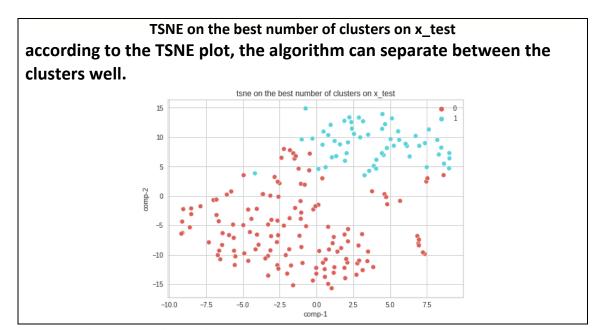


Distribution score elbow for k-means clustering:



So the best number of k is k = 2

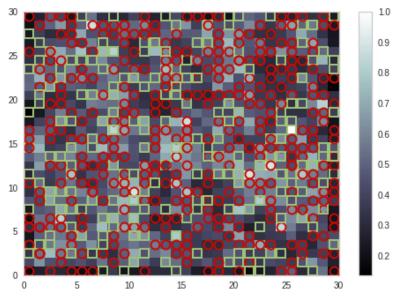




6- Train the SOM algorithm:

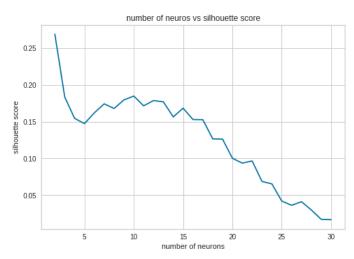
An SOM is mainly used for data visualization and provides a quick visual summary of the training instances. In a 2D rectangular grid, each cell is represented by a weight vector. For a trained SOM, each cell weight represents a summary of a few training examples. Cells in the close vicinity of each other have similar weights, and like examples can be mapped to cells in a small neighborhood of each other.

The final result on training SOM model, the 30*30 neuron:



So the best number of neuron is 2 based on the silhouette score

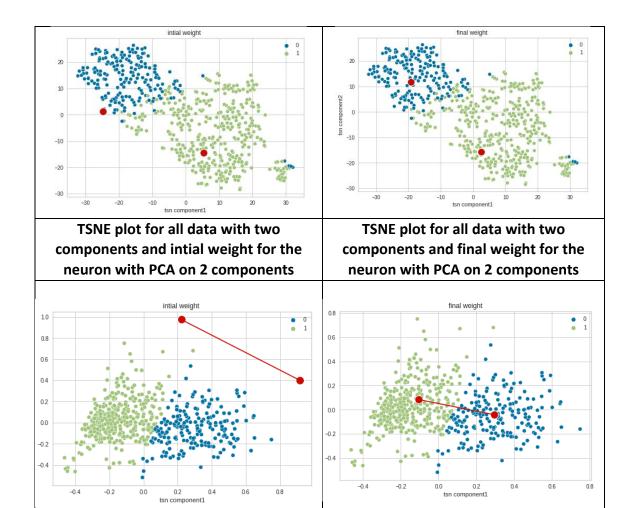
```
# the best number of neuron is 2 based on the highest silhouette score
plt.plot(range(2,31),scores)
plt.xlabel('number of neurons')
plt.ylabel('silhouette score')
plt.title('number of neuros vs silhouette score')
```



Apply PCA on the SOM Model:

```
138] som = MiniSom(2,1, 6, sigma=0.3, learning_rate=0.5,random_seed=0) # initialization of 2*1 SOM
   intial_weight = som.get_weights().copy()
   som.train_batch(np.array(x_train_b_knn), 1000) # trains the SOM with 1000 iterations
   final_weight = som.get_weights()
   w = np.array([som.winner(x) for x in np.array(x_train_b_knn)]).T
   index = np.ravel_multi_index(w,(2,1))
```

intial_weighT	final_weight
array([[[0.16419828, 0.72385141,	array([[[0.47710267, 0.71042386,
0.34567421, 0.15097751, -0.25680907,	0.59288791, 0.08145516, 0.46140247,
0.49075687]], [[-0.09085602,	0.36316582]], [[0.13668112, 0.6537922 ,
0.57031478, 0.67496669, -0.1696774 ,	0.54311031, 0.13120688, 0.47819656,
0.4246722 , 0.04206313]]])	0.12776533]]])
TSNE plot for all data with two	TSNE plot for all data with two
components and intial weight for the	components and final weight for the
neuron	neuron



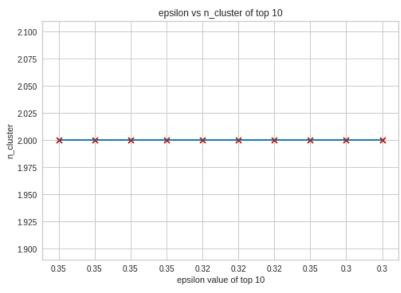
Part 7: Tune the epsilon (0.3-0.7) and minpoints (2-15) values to obtain the same number of clusters in Q6 by using DBSCAN:

The **DBSCAN algorithm** is based on this intuitive notion of "clusters" and "noise". The key idea is that for each point of a cluster, the neighborhood of a given radius has to contain at least a minimum number of points.

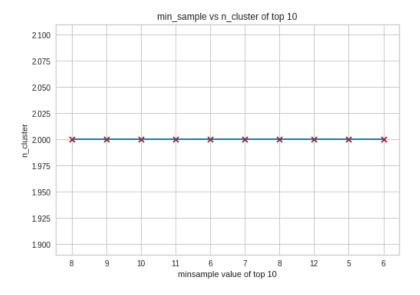
Top 10 cluster	S:
----------------	----

	eps	minsample	silhouette	cluster
34	0.35	8	0.264563	2
35	0.35	9	0.263837	2
36	0.35	10	0.259480	2
37	0.35	11	0.258518	2
18	0.32	6	0.254182	2
19	0.32	7	0.250709	2
20	0.32	8	0.250598	2
38	0.35	12	0.248660	2
3	0.30	5	0.246779	2
4	0.30	6	0.244219	2

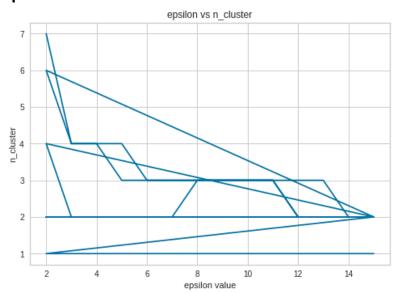
Plot the number of neurons vs number of clusters(top 10):



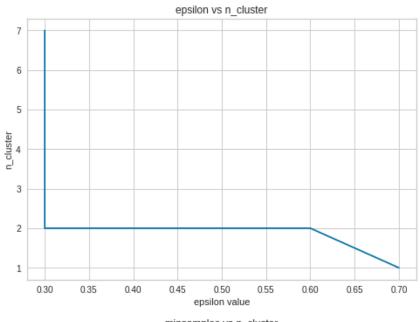
plotting the minpoints vs number of clusters(top 10):

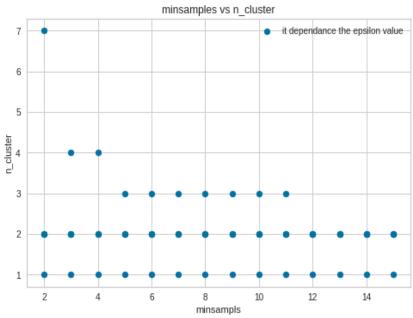


Number of epsilon vs number of clusters:



The best epsilon and min samples is 0.5 as epsilon and samples[2:15] this values achieve the high siluoette score on the best feature

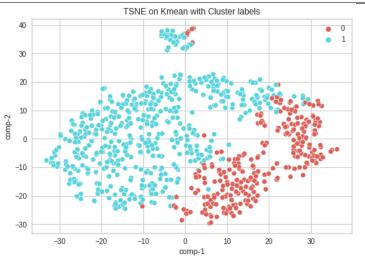




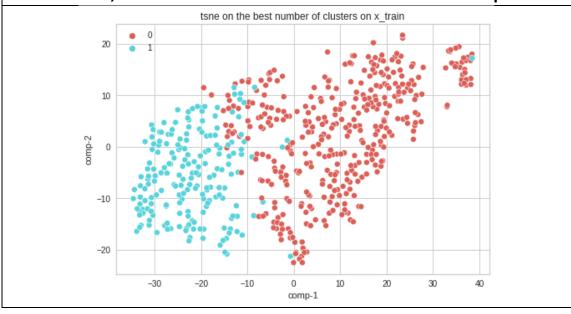
Conclusion:

In this assignment we learned how to apply the logistic regression model and the KNN model, applying the different feature selection methods like the filter selection and wrapper methods, applying the principle component analysis, and applying SOM algorithm and DBSCAN algorithm. According to our results of K-means algorithm we can say that:

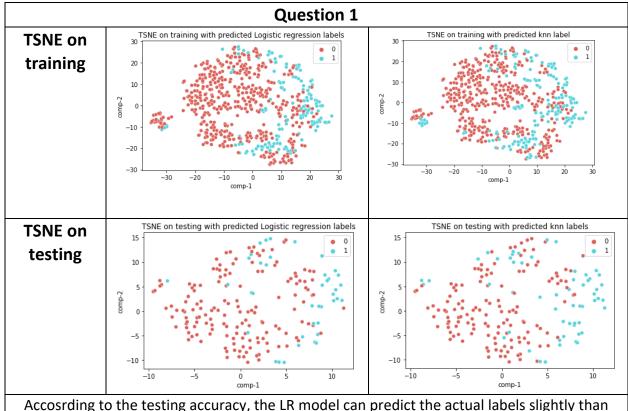




The silhouette score is increased by 0.31 after dimensionality reduction and feature selection, and the highest silhouette score means that we can separate between the clusters and the distance between them is increased, and we can see the difference between the two plots.



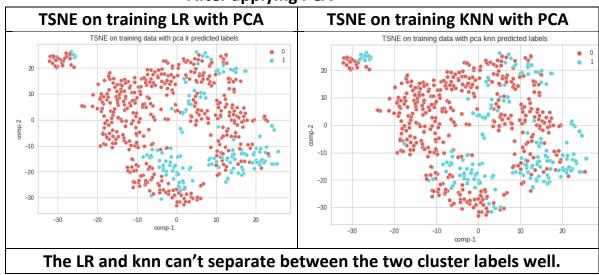
From question 1, 3 and 4 we can compare between the results as:

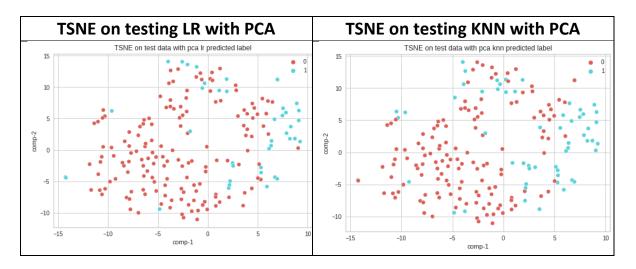


Accosrding to the testing accuracy, the LR model can predict the actual labels slightly than the KNN model.

Accuracy of LR is 77%, and KNN is 75%

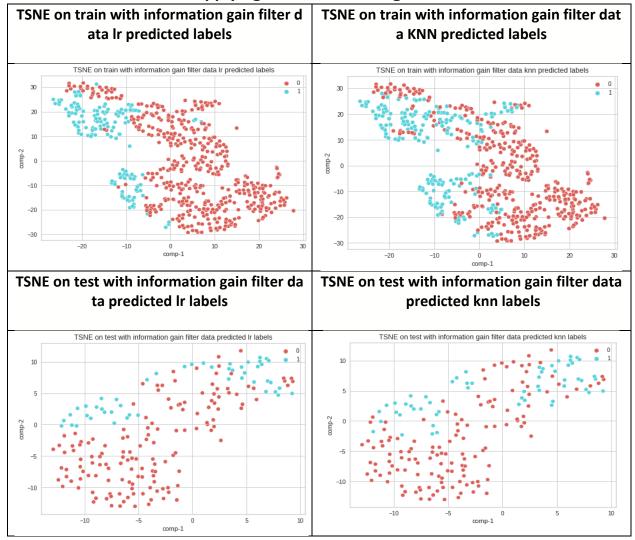
After applying PCA





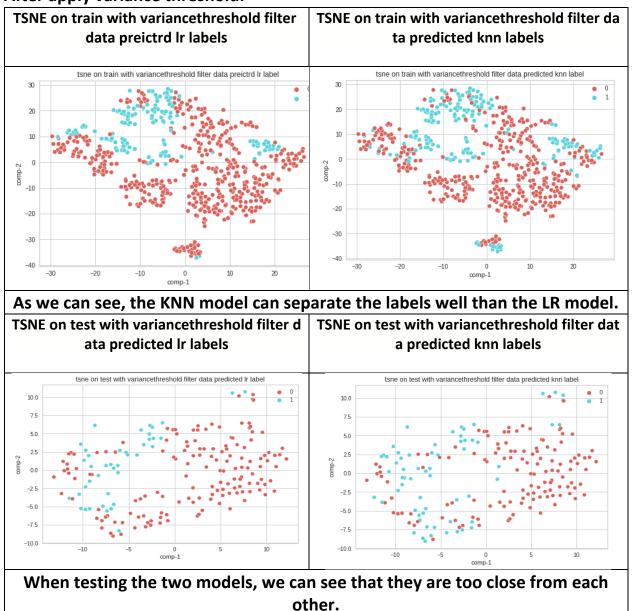
accuracy for LR with PCA: 78% accuracy for KNN with PCA: 76%

after applying the information gain:



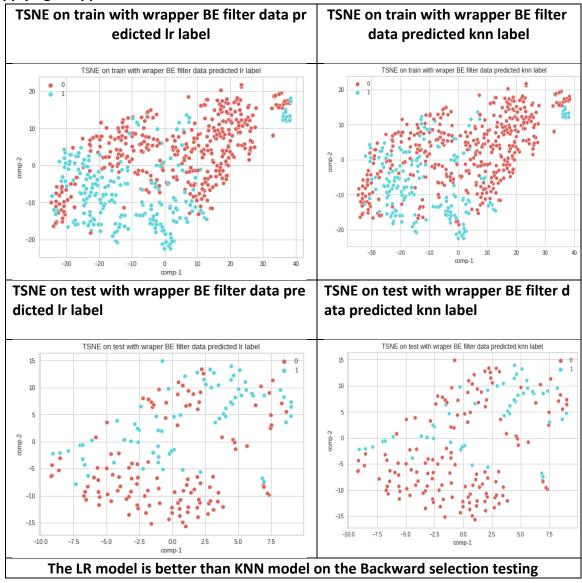
Accuracy of LR is 79% Accuracy of KNN is 76%

After apply variance threshold:



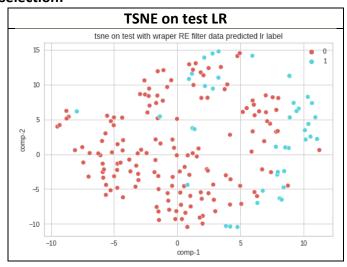
Accuracy of LR is 77% Accuracy of KNN is 78%

After applying wrapper method:



Accuracy of LR is 78% Accuracy of KNN is 76%

Recursive elimination selection:



Accuracy of LR is 78%

So, after comparing between the TSNE plots for all models, we can say that the LR model accuracies are closed to each other from 76: 79%, and the KNN model accuracies are between 75: 78%

References:

https://scikit-

<u>learn.org/stable/modules/generated/sklearn.linear_model.LogisticRegress_ion.html</u>

https://www.datacamp.com/tutorial/k-nearest-neighbor-classification-scikit-learn

https://www.kdnuggets.com/2020/04/dbscan-clustering-algorithm-machine-learning.html

https://stackabuse.com/self-organizing-maps-theory-and-implementation-/in-python-with-numpy

colab code link: https://colab.research.google.com/drive/18YEqf-coNeP721YTjcpqt9DSHEmu4gHJ?usp=sharing