

# Dimensionality Reduction Assignment 3 Applied Machine Learning ELG5255[EG]

Group name: Group 6

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#### 1. Load Iris Dataset

```
[ ] pokemon_train = pd.read_csv("Pokemon_train.csv")
    pokemon_test= pd.read_csv("Pokemon_test.csv")

[ ] x_pokemon_train=pokemon_train.iloc[:,:-1].to_numpy()
    y_pokemon_train=pokemon_train.iloc[:,-1]

[ ] x_pokemon_test=pokemon_test.iloc[:,:-1].to_numpy()
    y_pokemon_test=pokemon_test.iloc[:,-1]
```

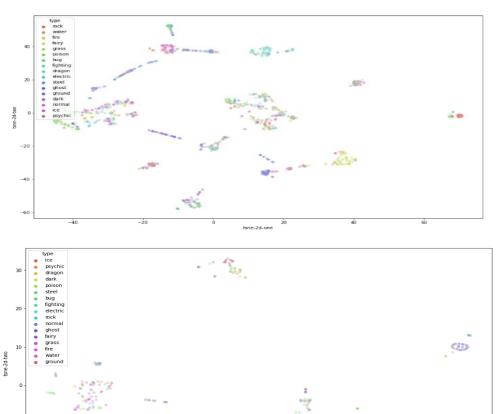
Figure 1: loading the dataset

## 2. Apply Gaussian Naïve Bayes classifier (GNB) and Support Vector Machine (SVM) to Pokémon dataset

Confusion Matrix Gaussian Naïve Bayes: naive clf = GaussianNB() naive\_clf.fit(x\_pokemon\_train,y\_pokemon\_train) y\_pred\_naive=naive\_clf.predict(x\_pokemon\_test) #Making the Confusion Matrix and Classification Report acc\_GNB1=naive\_clf.score(x\_pokemon\_test,y\_pokemon\_test)\*100 0 0 0 0 0 0 0 0 1 0 0 3 0 18 0 0 2 9 1 5 1 print("Accuracy for Gaussian Naïve Bayes in testing data: ",acc\_GNB1) print('\nClassification Report Gaussian Naïve Bayes:\n') print(classification report(y pokemon test,y pred naive)) 6 15 14 13 12 11 10 9 print('Confusion Matrix\_Gaussian Naïve Bayes:\n') hm=sn.heatmap(confusion\_matrix(y\_pokemon\_test, y\_pred\_naive), annot=True) 0 0 0 plt.show() Accuracy for Gaussian Naïve Bayes in testing data: 51.43769968051119 0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16

```
svm_clf= SVC(kernel='rbf',random_state=0)
svm_clf.fit(x_pokemon_train,y_pokemon_train)
# Predicting the Test set results
y_pred_svm=svm_clf.predict(x_pokemon_test)
#Making the Confusion Matrix and Classification
acc_SVC1=svm_clf.score(x_pokemon_test,y_pokemon_print("Accuracy for model_SVM: ",acc_SVC1)
print('\nClassification Report_SVM:\n')
print(classification_report(y_pokemon_test,y_print('Confusion MatrixSVM:\n')
# plot_confusion_matrix(svm_clf,x_pokemon_test,y_pnt)
hm=sn.heatmap(confusion_matrix(y_pokemon_test,y_plt.show())
```

```
x_tsne_train=TSNE(n_components=2, random_state=0).fit_transform(x_pokemon_train)
x_tsne_train
pokemon_train_ts=pd.DataFrame()
pokemon_train_ts['tsne-2d-one'] = x_tsne_train[:,0]
pokemon_train_ts['tsne-2d-two'] = x_tsne_train[:,1]
pokemon_train_ts['type']=y_pokemon_train
plt.figure(figsize=(16,10))
sns.scatterplot(
    x="tsne-2d-one", y="tsne-2d-two",
    hue="type",
    palette=sns.color_palette("hls", 17),
    data=pokemon_train_ts,
    legend="full",
    alpha=0.3
)
```



After applying the Gaussian Naïve Bayes classifier and Support Vector Machine that we know the Gaussian Naïve Bayes (51.44%) is better than Support Vector Machine (12.14%) and applying TSNE on training and testing data to visualize data with number of components= 2.

-10

-20

### 3. Choose the best number of cluster for k-means clustering algorithm

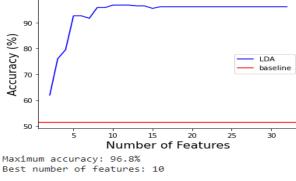
```
inertia1 = []
for i in range(1, 11):
     kmeans = KMeans(n_clusters = i, init = 'k-means++', random_state = 0)
     kmeans.fit(x_train_tsne)
     inertia1.append(kmeans.inertia )
kn1 = KneeLocator(range(1,11), inertia1, curve='convex', direction='decreasing'
plt.plot(range(1, 11), inertia1)
plt.title('The Elbow Method')
plt.xlabel('Number of clusters')
plt.ylabel('WCSS')
plt.vlines(kn1.knee, plt.ylim()[0], plt.ylim()[1], linestyles='dashed')
plt.show()
                 The Elbow Method
                                                                  KMeans
 1.75
                                                  40
 1.50
 1.25
                                                  20
                                                                              ★ Centroid
SS 100
 0.75
 0.50
                                                 -20
 0.25
                                                 -40
 0.00
                        6
                                8
                 Number of clusters
                                                 -60
```

After applying k-means with elbow method on the dataset to Determine the optimal number of clusters. Optimal number of clusters =4. Then plotting data with the optimal number of clusters (4).

4. Apply one of the following Dimensionality Reduction (DR) methods to data. Find the best value for n\_components based on the GNB and SVM classifiers test accuracies.

```
# Models
SVC_model=SVC(kernel='rbf',random_state=0)
GNB_model=GaussianNB()
```

```
[ ] def apply LDA(model,baseModel acc):
        model scores=[]
        for i in range(1,33):
           LDA = LinearDiscriminantAnalysis(n_components=i)
          x train lda = LDA.fit transform(x pokemon train,y pokemon train)
          x_test_lda = LDA.transform(x_pokemon_test)
          model.fit(x train lda,y pokemon train)
          model_scores.append(model.score(x_test_lda,y_pokemon_test)*100)
         max accuracy=max(model scores)
         index best com = model scores.index(max accuracy)+1
         # Plot a simple line chart
         plt.plot(range(1,33), model_scores, 'b', label='LDA')
         # Plot another line on the same chart/graph
         plt.axhline( y=baseModel_acc,c='r', label='baseline')
         plt.xlabel("Number of Features", fontsize=16)
         plt.ylabel("Accuracy (%)", fontsize=16)
         plt.legend()
         plt.show()
         return index best com, max accuracy
```





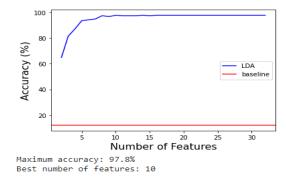
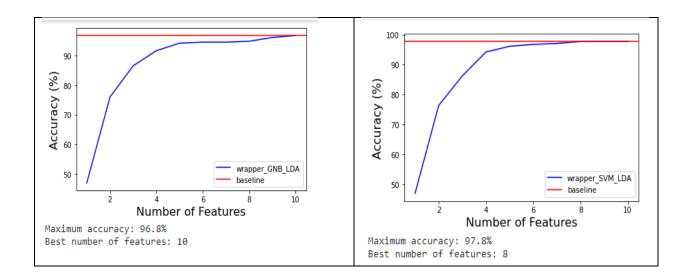


Figure 3: LDA with SVM

We use these parameters with the models because that affect to our accuracy better and after applying LDA on the Pokémon dataset with GNB and SVM classifiers. We get the best accuracy = 97.8% and number of features = 10 in LDA with SVM.

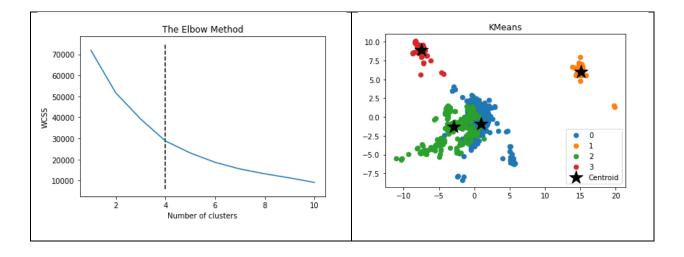
5. Use the following Feature Selection methods (one for each method). Find the best number of features based on the GNB and SVM classifiers' test accuracies on the data that is obtained after Q4. Plot the number of features versus accuracy graph for each case with the improved baseline performance as shown in Q4.

```
# Data after transformation with 10 features
       LDA = LinearDiscriminantAnalysis(n components=10)
       x_train lda = LDA.fit transform(x pokemon train,y pokemon train)
       x test lda = LDA.transform(x pokemon test)
       def select feature(fs, model):
            fs.fit(x train lda, y pokemon train)
            train new = fs.transform(x train lda)
            test new = fs.transform(x test lda)
            model.fit(train new, y pokemon train)
            yPred = model.predict(test new)
            acc = accuracy_score(y_pokemon_test, yPred) * 100
            return acc
                                                   100
   90
                                                    90
Accuracy (%) 8 8 8
                                                 Accuracy (%)
                                 filter GNB LDA
                                                    50
                                                                              filter SVM LDA
   50
                                 baseline
                                                                              baseline
                                                                                    10
              Number of Features
                                                              Number of Features
Maximum accuracy: 96.8%
                                                Maximum accuracy: 97.8%
Best number of features: 10
                                                Best number of features: 10
```



Applying Filter Method (Information Gain) and Wrapper Method (Forward) with the data from LDA, GNB and SVM classifiers. We get the best accuracy = 97.8% and the number of features = 8 when using wrapper method with LDA and SVM.

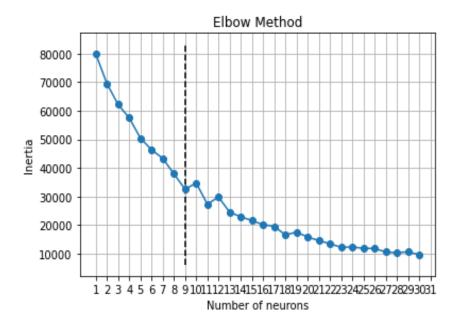
6. Choose the best number of clusters for k-means clustering algorithm on the processed data (using the best features or dimensionality from Q4 and Q5)



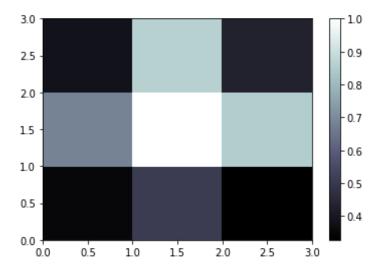
After applying LDA in Q4 and filter and wrapper method in Q5 that we get the best accuracy (97.8%) with the least number of feature (8) in wrapper method with LDA and SVM. After applying k-means on the dataset from wrapper method with LDA and SVM and using elbow method to determine the best number of cluster (4).

7. Choose the best number of neurons for SOM algorithm using the best features or dimensionality from Q4 and Q5

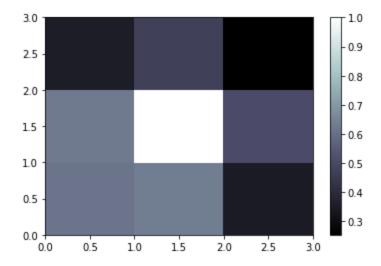
```
score = []
df som = pd.DataFrame(test sfs SVM)
for i in range(1,31):
  SOM_model = SOM(m=i, n=1, dim=train_sfs_SVM.shape[1],max_iter=3000,random_sta
 SOM_model.fit(train_sfs_SVM)
 SOM lables=SOM model.predict(test sfs SVM)
  score.append((SOM model.inertia ,i))
df som['SOM lables'] = SOM lables
df score = pd.DataFrame(score, columns=['score', 'neurons'])
kn3 = KneeLocator(range(1,31), df_score['score'], curve='convex', direction='de
plt.plot(df_score['neurons'], df_score['score'], marker = 'o')
plt.title("Elbow Method")
plt.xlabel("Number of neurons")
plt.ylabel("Inertia")
plt.xticks([i for i in range(1,33)])
plt.grid(True)
plt.vlines(kn3.knee, plt.ylim()[0], plt.ylim()[1], linestyles='dashed')
plt.show()
```



```
np.random.seed(43)
som_rows=3
som_columns=3
som_iterations=3000
sigma=1
learning_rate=0.5
np.random.seed(42)
som_init = MiniSom(x = som_rows, y = som_columns, input_len=8, sigma=sigma, lea som_init.random_weights_init(train_sfs_SVM)
bone()
pcolor(som_init.distance_map().T)
colorbar()
show()
```



```
np.random.seed(42)
som_final=MiniSom(x = som_rows, y = som_columns, input_len=8, sigma=sigma, learn
som_final.train_random(train_sfs_SVM,som_iterations)
som_final.distance_map()
bone()
pcolor(som_final.distance_map().T)
colorbar()
show()
```

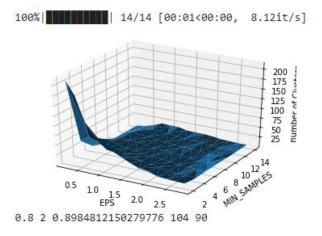


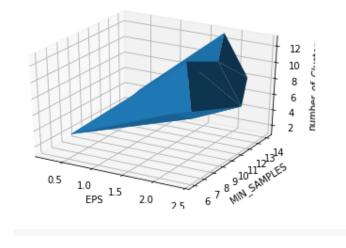
After applying LDA in Q4 and filter and wrapper method in Q5 that we get the best accuracy (97.8%) with the least number of feature (8) in wrapper method with LDA and SVM. After applying the SOM on the dataset from wrapper method with LDA and SVM and using elbow rule by inertia to determine the best number of neurons (9) then plot neurons in the initial (before training) and final positions (after training) on the shape gird 3\*3 because number of neurons = 9.

8. Tune the epsilon (0.2-3) and minpoints (2-15) values in the given intervals to obtain same number of clusters in Q6 by using DBSCAN

```
def unsupervisedLabelMap(labels, y):
     labelDict = dict()
     for label in unique labels(labels):
        tmpY = y[labels == label]
        unique, count = np.unique(tmpY, return_counts=True)
        trueLabel = unique[np.argmax(count)]
        labelDict[label] = trueLabel
     return labelDict
def usLabels2sLabels(labels, y):
     sLabels = np.empty(labels.shape, labels.dtype)
    labelDict = unsupervisedLabelMap(labels, y)
    for usl, tl in labelDict.items():
        sLabels[labels == usl] = tl
     return sLabels
encoder = LabelEncoder()
encoder train y = encoder.fit transform(y pokemon train)
```

```
epsList, mpList, clustersList, accuracyList, noiselist = list(), list(), list(), list(), list()
for eps in tqdm(np.arange(0.2, 3,0.2)):
        for mp in range(2, 15,2):
            model = DBSCAN(eps=eps, min samples=mp)
            y_pred_db = model.fit_predict(train_sfs_SVM)
            labels = model.labels
            n_clusters_ = len(set(labels)) - (1 if -1 in labels else 0)
            n noise = list(labels).count(-1)
            predY = usLabels2sLabels(y_pred_db, encoder_train_y)
            accuracy = accuracy_score(encoder_train_y, predY)
            epsList.append(eps)
            mpList.append(mp)
            clustersList.append(n_clusters_)
            accuracyList.append(accuracy)
            noiselist.append(n noise )
epsList, mpList, accuracyList, clustersList, noiselist = np.array(epsList), np.array(mpList),
   np.array(accuracyList), np.array(clustersList),np.array(noiselist)
ax = plt.axes(projection='3d')
ax.plot trisurf(epsList, mpList, clustersList)
ax.set_xlabel('EPS')
ax.set_ylabel('MIN_SAMPLES')
ax.set_zlabel('number of Clusters')
plt.show()
x = accuracyList.argmax()
print(epsList[x], mpList[x], accuracyList[x],noiselist[x],clustersList[x])
```





	срэттоп	minpoint	number of	CIUSCO
6	0.2	14		1
3	0.2	8		2
13	0.4	14		4
95	2.8	10		6
81	2.4	10		7
12	0.4	12		8
71	2.2	4		9
69	2.0	14		11
11	0.4	10		12
54	1.6	12		13

epsilon minpoint number of cluster

After applying tune, the epsilon (0.2-3) and minpoints (2-15) values in DBSCAN, plotting the epsilon, minpoints and the number of clusters in 3D figure, we know the best epsilon (0.4) and minpoints (14) to obtain the same number of clusters = 4 in (Q6). Showing the 10 combinations from epsilon and minpoints with number of clusters in dataframe to be organized and plotting the 10 combinations from epsilon and minpoints in 3D figure.

### 9. Conclusion

In conclusion, this report could be summarized by loading the data from Pokémon dataset, after we applied the GNB and SVM classifiers and k-means, then applied TSNE to visualize data and LDA with the GNB and SVM classifiers. After applied LDA then applied filter and wrapper methods based on data from LDA. Then we applied k-means and SOM with LDA and wrapper method because the accuracy of wrapper with LDA is the best = 98.1% and the least of number of features = 5 and we determined the optimal number of clusters in k-means and number of neurons in SOM by elbow. We applied tuning the epsilon and minpoints in DBSCAN to obtain the same number of clusters in k-means (Q6) and plotting the epsilon, minpoints and number of clusters in 3D figure.