

# Problem Statement

In this assignment students have to transform iris data into 3 dimensions and plot a 3d chart with transformed dimensions and color each data point with specific class.

## Import libraries into working environment

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D
from sklearn import decomposition
from sklearn import datasets
import seaborn as sns
from sklearn.decomposition import PCA
```

## Load iris data set

```
In [2]: iris = datasets.load_iris()
X = iris.data
y = iris.target
print("Number of samples:")

print(X.shape[0])
print('-----')
print('Number of features :')
print(X.shape[1])
print('-----')
print("Feature names:")
print('-----')
print(iris.feature_names)
```

Number of samples:

150

-----  
Number of features :

4

-----  
Feature names:

-----  
['sepal length (cm)', 'sepal width (cm)', 'petal length (cm)', 'petal width (cm)']

## Feature scaling prior to applying PCA

```
In [3]: # Feature Scaling
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
X_scaled = sc.fit_transform(X)
print('shape of scaled data points:')
print('-----')
print(X_scaled.shape)
print('first 5 rows of scaled data points :')
print('-----')
print(X_scaled[:5,:])
```

shape of scaled data points:

-----

(150, 4)

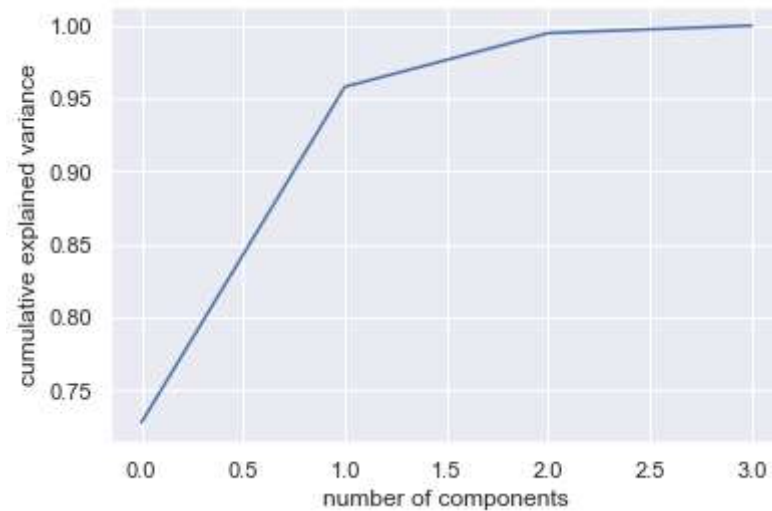
first 5 rows of scaled data points :

-----

```
[[-0.90068117  1.03205722 -1.3412724  -1.31297673]
 [-1.14301691 -0.1249576  -1.3412724  -1.31297673]
 [-1.38535265  0.33784833 -1.39813811 -1.31297673]
 [-1.50652052  0.10644536 -1.2844067  -1.31297673]
 [-1.02184904  1.26346019 -1.3412724  -1.31297673]]
```

**looking at the explained variance as a function of the components**

```
In [4]: sns.set()  
pca = PCA().fit(X_scaled)  
plt.plot(np.cumsum(pca.explained_variance_ratio_))  
plt.xlabel('number of components')  
plt.ylabel('cumulative explained variance')  
plt.show()
```



### Note

Here we see that we'd need about 3 components to retain 100% of the variance. Looking at this plot for a high-dimensional dataset can help us understand the level of redundancy present in multiple observations.

### PCA using Eigen-decomposition: 5-step process

```

In [5]: # 1. Normalize columns of A so that each feature has zero mean
A0 = iris.data
mu = np.mean(A0,axis=0)
A = A0 - mu
print("Does A have zero mean across rows?")
print(np.mean(A,axis=0))
print('-----')
print('Mean value : ')
print('-----')
print(mu)
print('Standardized Feature value first 5 rows: ')
print('-----')
print(A[:5,:])

# 2. Compute sample covariance matrix Sigma = {A^TA}/{(m-1)}
#covariance matrix can also be computed using np.cov(A.T)
m,n = A.shape
Sigma = (A.T @ A)/(m-1)
print("-----")
print("Sigma:")
print(Sigma)

# 3. Perform eigen-decomposition of Sigma using `np.linalg.eig(Sigma)`
W,V = np.linalg.eig(Sigma)
print("-----")
print("Eigen values:")
print(W)
print("-----")
print("Eigen vectors:")
print(V)

# 4. Compress by ordering 3 eigen vectors according to largest eigen values and compute AX_k
print("-----")
print("Compressed - 4D to 3D:")
print("-----")
print('First 3 eigen vectors :')
print(V[:, :3] )
print("-----")
Acomp = A @ V[:, :3]
print('First first five rows of transformed features :')
print("-----")
print(Acomp[:5,:])

```

```
# 5. Reconstruct from compressed version by computing  $A V_k V_k^T$ 
print("-----")
print("Reconstructed version - 3D to 4D:")
print("-----")
Arec = A @ V[:, :3] @ V[:, :3].T # first 3 e vectors
print(Arec[:5, :] + mu) # first 5 obs, adding mu to compare to original
```

Does A have zero mean across rows?

```
[-1.12502600e-15 -6.75015599e-16 -3.23889064e-15 -6.06921920e-16]
```

-----  
Mean value :

```
[5.84333333 3.054      3.75866667 1.19866667]
```

Standardized Feature value first 5 rows:

```
-----  
[[-0.74333333  0.446      -2.35866667 -0.99866667]  
 [-0.94333333 -0.054      -2.35866667 -0.99866667]  
 [-1.14333333  0.146      -2.45866667 -0.99866667]  
 [-1.24333333  0.046      -2.25866667 -0.99866667]  
 [-0.84333333  0.546      -2.35866667 -0.99866667]]  
-----
```

Sigma:

```
[ [ 0.68569351 -0.03926846  1.27368233  0.5169038 ]  
 [-0.03926846  0.18800403 -0.32171275 -0.11798121]  
 [ 1.27368233 -0.32171275  3.11317942  1.29638747]  
 [ 0.5169038  -0.11798121  1.29638747  0.58241432]]  
-----
```

Eigen values:

```
[4.22484077 0.24224357 0.07852391 0.02368303]  
-----
```

Eigen vectors:

```
[ [ 0.36158968 -0.65653988 -0.58099728  0.31725455]  
 [-0.08226889 -0.72971237  0.59641809 -0.32409435]  
 [ 0.85657211  0.1757674  0.07252408 -0.47971899]  
 [ 0.35884393  0.07470647  0.54906091  0.75112056]]  
-----
```

Compressed - 4D to 3D:

-----  
First 3 eigen vectors :

```
[ [ 0.36158968 -0.65653988 -0.58099728]  
 [-0.08226889 -0.72971237  0.59641809]  
 [ 0.85657211  0.1757674  0.07252408]  
 [ 0.35884393  0.07470647  0.54906091]]  
-----
```

First first five rows of transformed features :

```
-----  
[[-2.68420713 -0.32660731 -0.02151184]  
 [-2.71539062  0.16955685 -0.20352143]  
 [-2.88981954  0.13734561  0.02470924]  
 [-2.7464372  0.31112432  0.03767198]  
-----
```

```
[-2.72859298 -0.33392456  0.0962297  ]]
```

-----  
Reconstructed version - 3D to 4D:  
-----

```
[[5.09968079 3.50032609 1.40048267 0.19924425]  
 [4.86840068 3.03228058 1.44778117 0.12518657]  
 [4.69387555 3.20625649 1.30926076 0.18549996]  
 [4.62409716 3.07538332 1.46356281 0.25705157]  
 [5.02002788 3.57954033 1.36971595 0.24741729]]
```

### Original iris feature values

```
In [6]: iris.data[:5,:]
```

```
Out[6]: array([[5.1, 3.5, 1.4, 0.2],  
               [4.9, 3. , 1.4, 0.2],  
               [4.7, 3.2, 1.3, 0.2],  
               [4.6, 3.1, 1.5, 0.2],  
               [5. , 3.6, 1.4, 0.2]])
```

### 3D Visualization

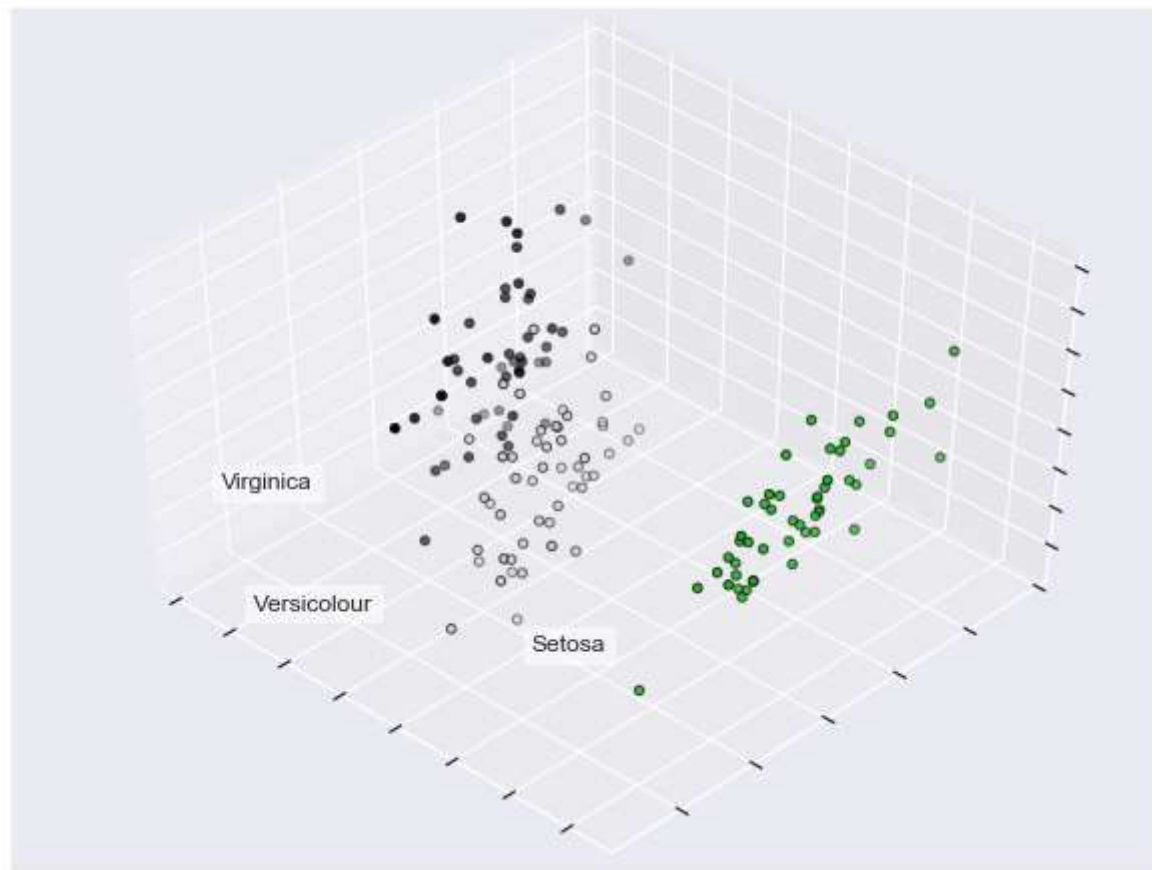


```
In [7]: np.random.seed(5)

centers = [[1, 1], [-1, -1], [1, -1]]
fig = plt.figure(1, figsize=(8, 6))
plt.clf()
ax = Axes3D(fig, rect=[0, 0, 1, 1], elev=48, azimuth=134)
y = iris.target
plt.cla()
for name, label in [('Setosa', 0), ('Versicolour', 1), ('Virginica', 2)]:
    ax.text3D(Acomp[y == label, 0].mean(),
              Acomp[y == label, 1].mean() + 1.5,
              Acomp[y == label, 2].mean(), name,
              horizontalalignment='center',
              bbox=dict(alpha=.5, edgecolor='w', facecolor='w'))
# Reorder the labels to have colors matching the cluster results
y = np.choose(y, [1, 2, 0]).astype(np.float)
ax.scatter(Acomp[:, 0], Acomp[:, 1], Acomp[:, 2], c=y, cmap=plt.cm.nipy_spectral,
           edgecolor='k')

ax.w_xaxis.set_ticklabels([])
ax.w_yaxis.set_ticklabels([])
ax.w_zaxis.set_ticklabels([])

plt.show()
```



**Applying PCA for number of compents = 3 using sklearn**

```
In [8]: pca = PCA(n_components=3)
pca.fit(X_scaled)
print('explained variance :')
print('-----')
print(pca.explained_variance_)
print('-----')
print('PCA Components : ')
print('-----')
print(pca.components_)
print('-----')
X_transformed = pca.transform(X)
print('Transformed Feature values first five rows :')
print('-----')
print(X_transformed[:5,:])
print('-----')
print('Transformed Feature shape :')
print('-----')
print(X_transformed.shape)
print('-----')
print('Original Feature shape :')
print('-----')
print(X.shape)
print('-----')
print('Retransformed Feature :')
print('-----')
X_retransformed = pca.inverse_transform(X_transformed)
print('Retransformed Feature values first five rows :')
print('-----')
print(X_retransformed[:5,:])
```

explained variance :

```
-----  
[2.93035378 0.92740362 0.14834223]  
-----
```

PCA Components :

```
-----  
[[ 0.52237162 -0.26335492  0.58125401  0.56561105]  
 [ 0.37231836  0.92555649  0.02109478  0.06541577]  
 [-0.72101681  0.24203288  0.14089226  0.6338014  ]]  
-----
```

Transformed Feature values first five rows :

```
-----  
[[ 2.66923088  5.18088722 -2.50606121]  
 [ 2.69643401  4.6436453  -2.48287429]  
 [ 2.4811633   4.75218345 -2.30435358]  
 [ 2.57151243  4.62661492 -2.22827673]  
 [ 2.59065822  5.23621104 -2.40975624]]  
-----
```

Transformed Feature shape :

```
-----  
(150, 3)  
-----
```

Original Feature shape :

```
-----  
(150, 4)  
-----
```

Retransformed Feature :

Retransformed Feature values first five rows :

```
-----  
[[5.13018217 3.48569954 1.30770618 0.26031309]  
 [4.92764912 2.98689971 1.31545197 0.25525129]  
 [4.72689213 3.18725838 1.21776678 0.25373858]  
 [4.67248379 3.06565682 1.27835237 0.3448445  ]  
 [5.04029862 3.58090632 1.27677117 0.2805288  ]]  
-----
```

### Note :

Transformed from 4D to 3D using PCA

```
In [9]: print('First Principal Component PC1: ',pca.components_[0])  
print('\nSecond Principal Component PC2: ',pca.components_[1])  
print('\nThird Principal Component PC3: ',pca.components_[2])
```

First Principal Component PC1: [ 0.52237162 -0.26335492 0.58125401 0.56561105]

Second Principal Component PC2: [0.37231836 0.92555649 0.02109478 0.06541577]

Third Principal Component PC3: [-0.72101681 0.24203288 0.14089226 0.6338014 ]

### Note:

Transforming from 3D to 4D

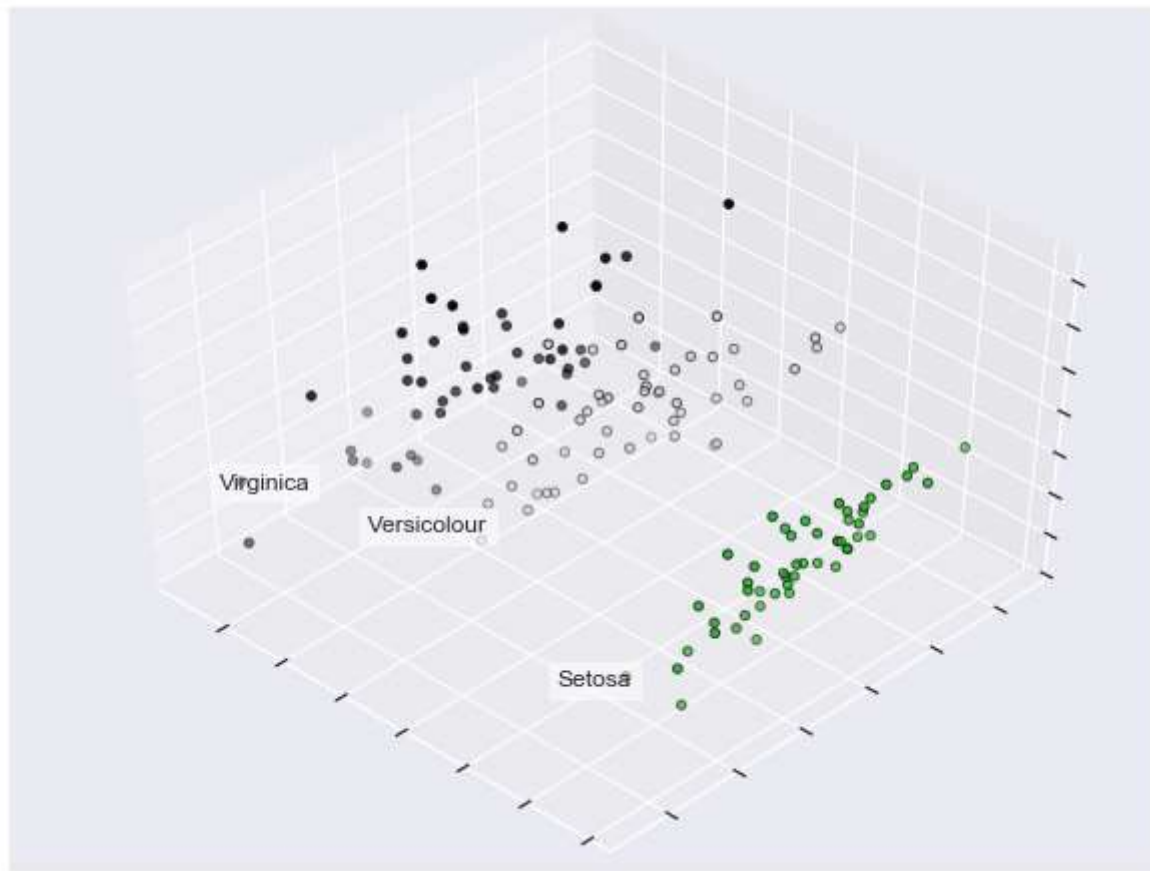
### 3D visualization

```
In [10]: np.random.seed(5)

centers = [[1, 1], [-1, -1], [1, -1]]
fig = plt.figure(1, figsize=(8, 6))
plt.clf()
ax = Axes3D(fig, rect=[0, 0, 1, 1], elev=48, azimuth=134)
y = iris.target
plt.cla()
for name, label in [('Setosa', 0), ('Versicolour', 1), ('Virginica', 2)]:
    ax.text3D(X_transformed[y == label, 0].mean(),
              X_transformed[y == label, 1].mean() + 1.5,
              X_transformed[y == label, 2].mean(), name,
              horizontalalignment='center',
              bbox=dict(alpha=.5, edgecolor='w', facecolor='w'))
# Reorder the labels to have colors matching the cluster results
y = np.choose(y, [1, 2, 0]).astype(np.float)
ax.scatter(X_transformed[:, 0], X_transformed[:, 1], X_transformed[:, 2], c=y, cmap=plt.cm.nipy_spectral,
           edgecolor='k')

ax.w_xaxis.set_ticklabels([])
ax.w_yaxis.set_ticklabels([])
ax.w_zaxis.set_ticklabels([])

plt.show()
```



In [ ]: