PR Assignment-2

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18MCMB15
M.TECH (IT)

1st question

```
def standerdize (data):
    mean = np.mean(data,axis=0)
    std = np.std(data,axis=0)
    return (data - mean) / std

def getpca(k,data,get_eig = False):
    data = standerdize(data)
    cov = np.cov(data.T)
    eig_vals , eig_vecs = np.linalg.eig(cov)
    eig_pairs = [(np.abs(eig_vals[i]),eig_vecs[:,i]) for i in range(len(eig_vals))]
    eig_pairs.sort(key = lambda k:k[0] , reverse = True)
    mat = np.hstack([eig_pairs[i][1][:,np.newaxis] for i in range(k)])
    if get_eig == True:
    return eigen_vals , data.dot(mat)
    return data.dot(mat)
```

def getIda(k,data,datalabels):

```
labels = np.unique(datalabels)
data = standerdize(data)
mean_vecs = []
mean_labels = []
for label in labels:
mean_vecs.append(np.mean(data[datalabels==label], axis=0))
mean_labels.append(label)
d = data.shape[1]
S_W = np.zeros((d,d))
for label,mv in zip(labels,mean_vecs):
class_scatter = np.zeros((d,d))
for row in data[datalabels==label]:
row, mv = row.reshape(d, 1), mv.reshape(d, 1)
class_scatter += (row-mv).dot((row-mv).T)
S_W += class_scatter
"""for label,mv in zip(range(1, 4), mean_vecs):
class_scatter = np.cov(X_train_std[y_train==label].T)
S W += class scatter"""
```

```
mean_overall = np.mean(data, axis=0)

mean_overall = mean_overall.reshape(d,1)

S_B = np.zeros((d, d))

for label, mean_vec in zip(mean_labels,mean_vecs):

n = data[datalabels == label].shape[0]

mean_vec = mean_vec.reshape(d,1)

S_B += n * (mean_vec - mean_overall).dot((mean_vec - mean_overall).T)

eigen_vals, eigen_vecs = np.linalg.eig(np.linalg.inv(S_W) @ (S_B))

eigen_pairs = [(np.abs(eigen_vals[i]), eigen_vecs[:,i]) for i in range(len(eigen_vals))]
eigen_pairs = sorted(eigen_pairs,key=lambda k: k[0], reverse=True) mat = np.hstack([eigen_pairs[i][1][:,np.newaxis] for i in range(k)])
```

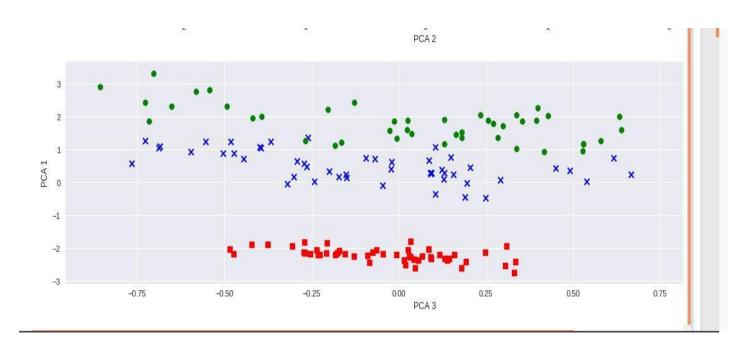
return data.dot(mat)

2nd Question:

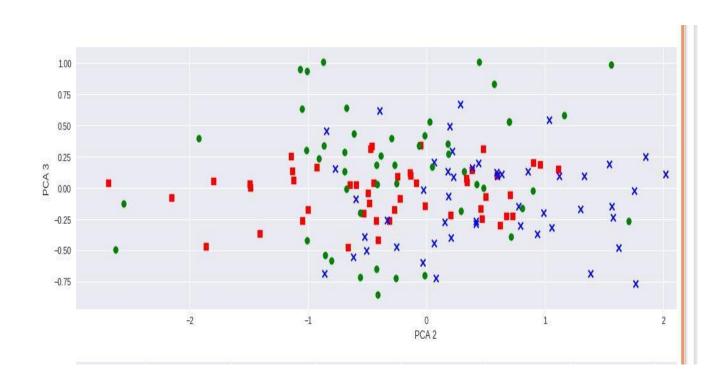
Results:

(Snapshots of the Results TO view them completely Open Notebook)

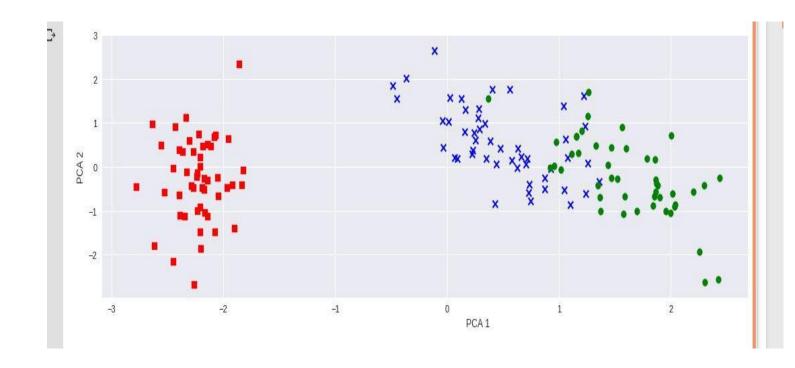
PCA1 vs PCA3



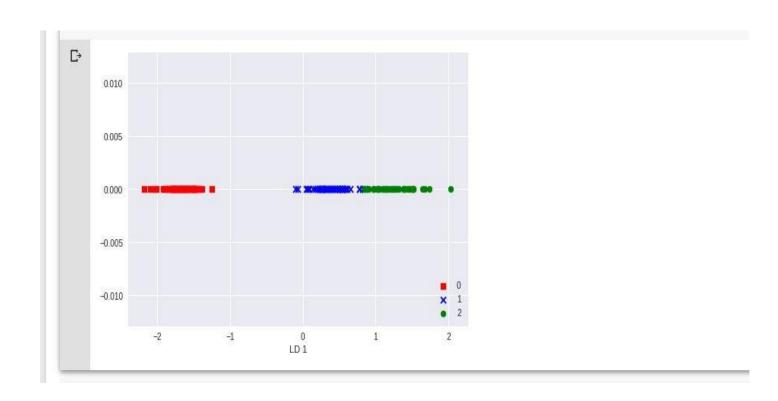
PCA2 vs PCA3



PCA1 vs PCA2

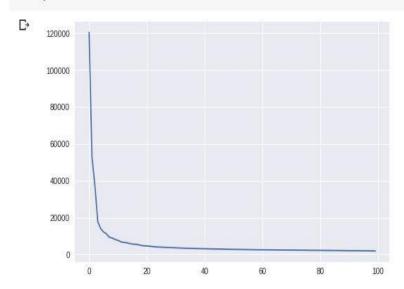


LDA:



Arcene:





3rd question

Libraries Used:

import numpy as np

import pandas as pd

import matplotlib.pyplot as plt

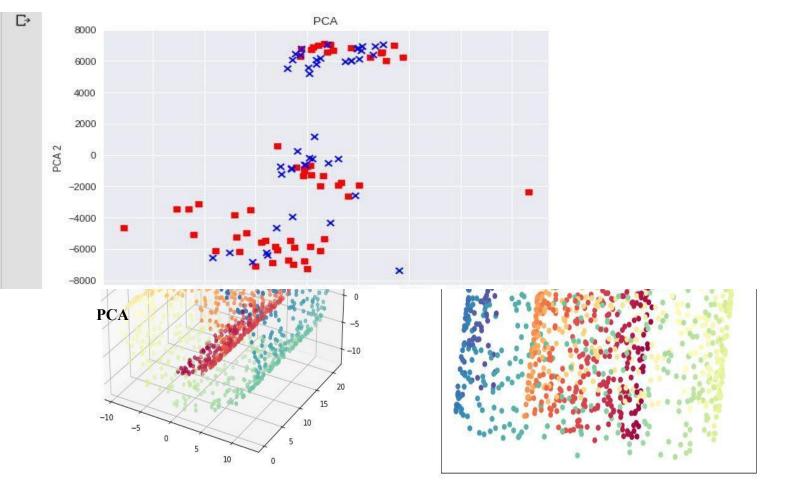
Dataset Used:

UCI Arcene cancer dataset

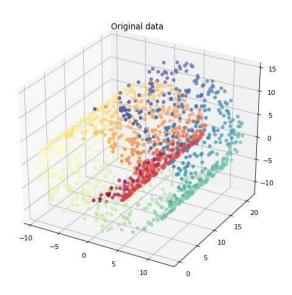
Analysis of components with respect to variance: No of Components required for getting 0.85 variances are: 61 No of Components required for getting 0.90 variances are: 73 No of Components required for getting 0.95 variances are: 86 No of Components required for getting 0.99 variances are: 97

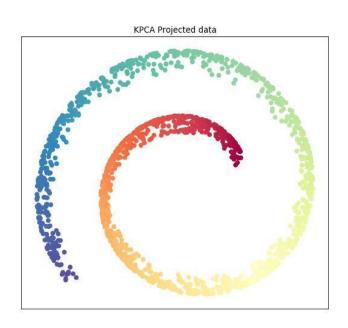
Results:

Pca two principal component

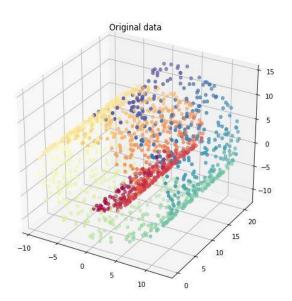


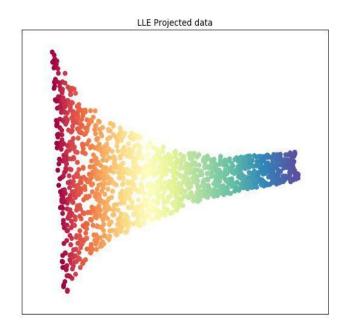
KPCA





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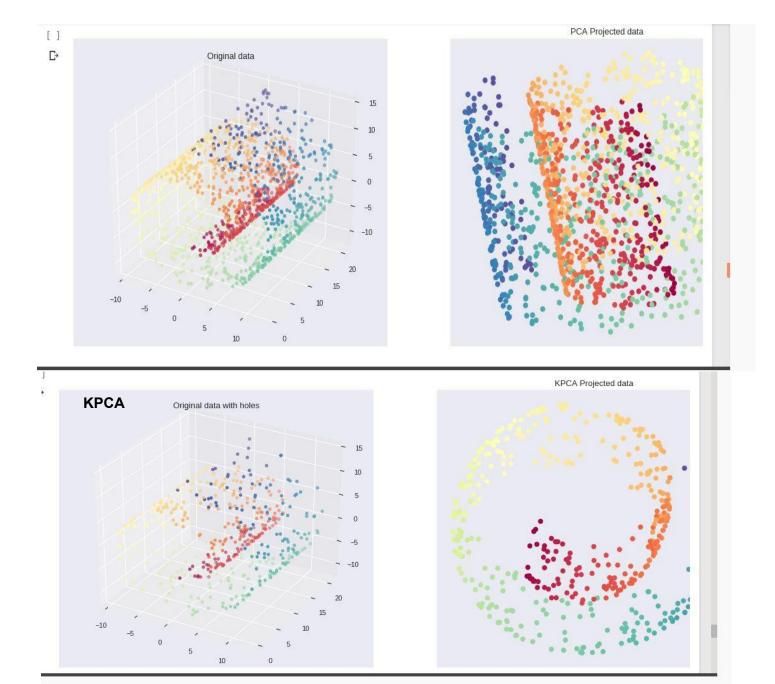
4th Question

Dataset Used:

3D swissroll dataset

Results:

PCA



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