

h5-pca-cs

June 11, 2018

Topics: Sparsity (PCA and Compressive Sensing)

Assigned: Wednesday May 23

Due: Sunday June 10 by midnight

```
In [1]: # -*- coding: utf-8 -*-
import numpy as np
from math import *
import matplotlib.pyplot as plt
from matplotlib import cm
from mpl_toolkits.mplot3d import Axes3D
from scipy.stats import norm

# Params
N = 2000
dims = 100
ulength = 7
```

1 Code & Logs

1.1 Part I PCA

Generate a random vector u in d dimensions as follows: The components of u are i.i.d., with $-P[u[i] = 0] = 2/3$; $P[u[i] = +1] = 1/6$; $P[u[i] = 1] = 1/6$

```
In [2]: # Generate the IID
def generateMultiDimGaussian(d, ulength):
    u = np.ndarray((ulength,d))
    angles = np.zeros((ulength,ulength))
    deviation = 10
    while np.amax(angles+90*np.eye(ulength)) > (90+deviation) or np.amin(angles+90*np.
        for i in range(ulength):
            for j in range(d):
                r = np.random.rand()
                if(r<4/6):
```

```

        u[i,j] = 0
    elif(r<5/6):
        u[i,j] = 1
    else:
        u[i,j] = -1
    if(u.any(axis=1).all()):
        for i in range(ulength):
            for j in range(i):
                angles[i,j] = np.arccos( np.clip( np.dot(u[i,:]/np.linalg.norm(u[i,:]), -1, 1) ) )
                angles[j,i] = angles[i,j]
    #print(angles)
    return u

# Uj be i.i.d
Uj = generateMultiDimGaussian(dims, ulength)
print(Uj.shape)

```

(7, 100)

Generate d-dimensional data samples for a Gaussian mixture distribution with 3 equiprobable components 1. Z_m : Standard Gaussian ($N(0, 1)$) distribution 2. N : noise vector" $N(0, 2Id)$ (default value 2 = 0.01) 3. Component 1: Generate $X = u_1 + Z_1u_2 + Z_2u_3 + N$. 4. Component 2: Generate $X = 2u_4 + \sqrt{2}Z_1u_5 + Z_2u_6 + N$. 5. Component 3: Generate $X = \sqrt{2}u_6 + Z_1(u_1 + u_2) + (1/\sqrt{2})Z_2u_5 + N$

In [3]: """

Generate the higher dimension dataset and sample equiprobable from components
 """

```

def generateDataset(u, num_data = 50, d = 30):
    sigma_sq = 0.01
    #print('\nX(Nxd): ', num_data, "x", d, ', \tUj: ', Uj[0].shape)
    dataset = np.ndarray((num_data, d))
    labels = np.zeros((num_data, 3)) # will be containing [0,1,0] one hot value
    # Assign the values based on the three component function
    for i in range(0, num_data):
        # Random numbers  $Z_m \{Z_1, Z_2\}$  and  $N$  are drawn afresh
        Z1 = np.random.normal()
        Z2 = np.random.normal()
        noise = np.random.multivariate_normal(np.zeros(d), (sigma_sq)*np.eye(d))
        # choose which component to pick from
        idx_comp = np.random.choice([0, 1, 2], 1, p=[0.333, 0.333, 0.334])
        if(idx_comp == 0): # Sample from component 1
            dataset[i,:] = Uj[1,:] + Z1*Uj[2,:] + Z2*Uj[3,:] + noise
        elif(idx_comp == 1): # Sample from component 3
            dataset[i,:] = 2*Uj[4,:] + np.sqrt(2)*Z1*Uj[5,:] + Z2*Uj[6,:] + noise
        elif(idx_comp == 2): # Sample from component 3
            dataset[i,:] = np.sqrt(2)*Uj[6,:] + Z1*(Uj[1,:] + Uj[2,:]) + 1/np.sqrt(2)*

```

```

        # Assign a label
        labels[i,idx_comp]=1
    return dataset, labels

#X_data, Y_hot_labels = generateDataset(Uj, 50, 30)
#print("> X: Data Set:",X_data.shape," Y: One hot :",Y_hot_labels.shape)

```

1.1.1 1. SVD of the A(N CE d) data matrix

$A = U \Sigma V^T$

1. V must diagonalize $A^T A$ and v_i are eigenvectors of $A^T A$.
2. Σ where Σ_{ii} are singular values of A.
3. U must diagonalize $A A^T$ and u_i are eigenvectors of $A A^T$

```

In [4]: # input sample N=? from components
def getSingularValues(N = 50, d=30):
    X_data, Y_hot_labels = generateDataset(Uj, N, d)
    print("> X: Data Set:",X_data.shape," Y: One hot :",Y_hot_labels.shape)
    #Xt = X_data.T
    U, S, V = np.linalg.svd(X_data, full_matrices=False)
    X_a = np.dot(np.dot(U, np.diag(S)), V)
    print("> SD-inp:", np.std(X_data), "\nSD-out:", np.std(X_a), "\nSD-Err:",np.std(X_
    print("Singular values ",N,"E",d," : ",S[0:7])
    return S

plt.figure(figsize=(14,6))
plt.title('Singular Values vs N')
plt.xlabel('Singular value index')
plt.ylabel('Singular Value magnitude')

for Num in [50, 60, 80, 100, 120, 150, 200, 300]:
    S = getSingularValues(Num, dims)
    plt.plot(range(len(S)), S,label = str('N='+str(Num)))
plt.legend()
plt.show()

```

```

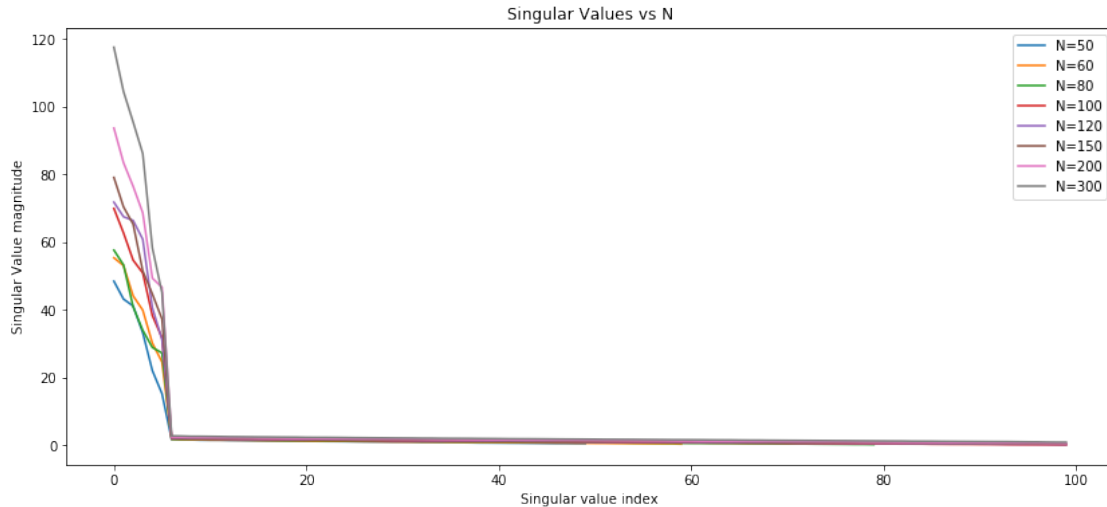
> X: Data Set: (50, 100) , Y: One hot : (50, 3)
> SD-inp: 1.2425699236269871
SD-out: 1.242569923626987
SD-Err: 3.048454173224238e-15
Singular values  50 E 100 : [48.39048927 43.11330642 41.06337937 33.16084801 21.96589821 15.
1.5994595 ]
> X: Data Set: (60, 100) , Y: One hot : (60, 3)
> SD-inp: 1.3498319267986385
SD-out: 1.3498319267986376
SD-Err: 3.4819962065686176e-15

```

```

Singular values  60 @ 100  : [55.31684119 53.0200592  44.0650705  39.84964441 30.09112566 24.1
1.69841237]
> X: Data Set: (80, 100) , Y: One hot : (80, 3)
> SD-inp: 1.1498704791356968
SD-out: 1.1498704791356964
SD-Err: 1.806028648889445e-15
Singular values  80 @ 100  : [57.5893642  53.216893  40.83616548 33.89971405 28.80635373 27.1
1.76064902]
> X: Data Set: (100, 100) , Y: One hot : (100, 3)
> SD-inp: 1.2997668301927499
SD-out: 1.2997668301927496
SD-Err: 4.440394990401163e-15
Singular values  100 @ 100  : [69.90863454 62.67001101 54.6252193  50.84244765 38.22080618 31
1.90438301]
> X: Data Set: (120, 100) , Y: One hot : (120, 3)
> SD-inp: 1.3060641823925117
SD-out: 1.3060641823925123
SD-Err: 1.6985712333628717e-15
Singular values  120 @ 100  : [71.74773111 67.46007854 66.34224228 60.79853815 40.5416491 31
1.98174145]
> X: Data Set: (150, 100) , Y: One hot : (150, 3)
> SD-inp: 1.1982438899074066
SD-out: 1.198243889907406
SD-Err: 2.0649813402642227e-15
Singular values  150 @ 100  : [79.06779613 70.36741234 65.26769035 51.45642124 44.6072953 37
2.11513307]
> X: Data Set: (200, 100) , Y: One hot : (200, 3)
> SD-inp: 1.2448438513004056
SD-out: 1.2448438513004045
SD-Err: 1.9863919347571614e-15
Singular values  200 @ 100  : [93.67871151 83.432265  76.45619056 68.60885691 49.28696661 46
2.31861979]
> X: Data Set: (300, 100) , Y: One hot : (300, 3)
> SD-inp: 1.24972911012126
SD-out: 1.2497291101212598
SD-Err: 1.6247784093964919e-15
Singular values  300 @ 100  : [117.56893054 104.47921109 95.45651027 86.14543701 58.243263
45.01554453  2.66531009]

```



1.1.2 1.(a) $d_0 = 6$ are the dominant singular values

we can see this based on the variation for $N = [50, 60, 80, 100, 120, 150, 200, 300]$

1.1.3 Now, project the data down to the dominant d_0 components to obtain an $N \times d_0$ data matrix.

```
In [5]: X_data, Y_hot_labels = generateDataset(Uj, N, dims)
print("> X: Data Set:", X_data.shape, ", Y: One hot :", Y_hot_labels.shape)
U, S, V = np.linalg.svd(X_data, full_matrices=False)
X_a = np.dot(np.dot(U, np.diag(S)), V)
print(">\nSD-inp:", np.std(X_data), "\nSD-out:", np.std(X_a), "\nSD-Err:", np.std(X_data - X_a))
print("Singular values ", N, "\n", dims, " : ", S[0:7])

d0 = 6 # Dominant vectors d0 = 6
print("Singular: ", np.diag(S[:d0]).shape)
reconst_matrix = np.dot(U[:, :d0], np.dot(np.diag(S[:d0]), V[:, :d0, :]))
print(">\nSD-inp:", np.std(X_data), "\nSD-out:", np.std(reconst_matrix), "\nSD-Err:", np.std(X_data - reconst_matrix))

# Using eigen vector V as the basis for projecting the data
evecs = V[:, :d0]
X_reduced_matrix = np.dot(X_data, evecs)
print("Reduced Matrix[From =", N, "\n", dims, "]:", " To =", X_reduced_matrix.shape)
#print(X_reduced_matrix)

> X: Data Set: (2000, 100) , Y: One hot : (2000, 3)
>
SD-inp: 1.2395057807174528
```

```

SD-out: 1.2395057807174528
SD-Err: 1.7785550318342964e-15
Singular values 2000 @ 100 : [282.76080712 266.71486558 242.47285716 229.01120717 161.325888
133.6358575 5.38798619]
Singular: (6, 6)
>
SD-inp: 1.2395057807174528
SD-out: 1.235722754594632
SD-Err: 0.09679696927444671
Reduced Matrix[From = 2000 @ 100 ]: To = (2000, 6)

```

```

In [47]: def runKmeansCluster(data, labels, d):
    class_means_dict=dict()
    preds_dict=dict()
    fig = plt.figure(figsize=(8,8))
    subplot_id = 221
    print("Data Dims: ",N,"x",d)
    print("Dataset : ", data.shape)
    # Kmeans trial
    for K in range(2,6):
        pred = np.zeros((N, K))
        class_means = np.ndarray((K, d))
        initial_indices = np.random.choice(N,K)

        for k in range(K):
            class_means[k,:] = data[initial_indices[k],:]
            old_class_means = np.zeros((K,d))

        print("\n----- K=",K,"-----")

        while np.linalg.norm(old_class_means-class_means)/np.linalg.norm(class_means)
            norm_mse = np.linalg.norm(old_class_means-class_means)
            norm_mu = np.linalg.norm(class_means)
            #print(">",norm_mse,"/",norm_mu," \t=\t ",norm_mse/norm_mu)
            old_class_means = np.array(class_means)
            for i in range(N):
                distance_to_means = np.zeros(K)
                for k in range(K):
                    distance_to_means[k] = np.linalg.norm(class_means[k] - data[i])
                #print("Distance: ",distance_to_means)
                nearest_mean = np.argmin(distance_to_means)
                #print("Nearest : ",nearest_mean)
                # labels as the min dist
                pred[i,:] = np.zeros(K)
                pred[i, nearest_mean] = 1 # one hot encoding

            # new mean

```

```

        for k in range(K):
            class_means[k] = np.mean(data[np.where(pred[:,k]==1)], axis=0)
            #print("Class MU : ", class_means)
    ax = fig.add_subplot(subplot_id)
    subplot_id = subplot_id + 1

    for k in range(K):
        colors = ('#fc0d1b', '#041ca2', '#162214', '#fd8008', '#c41bb6')
        ax.scatter(data[np.where(pred[:,k] == 1), 0], data[np.where(pred[:,k] == 1), 1],
            ax.scatter(class_means[k,0], class_means[k,1], s=200, marker='+', color = colors[k])
    ax.legend()
    #plt.savefig('dim_%d_q1_%d_means.png' % (d,K), dpi=600)
    #plt.show

    # save means
    class_means_dict[K] = class_means
    preds_dict[K] = pred
    probabilities = np.ndarray((3,K))

    for predicted in range(K):
        print('----- \n')
        for true_label in range(3):
            interssect = np.intersect1d(np.where(labels[:, true_label] == 1), np.where(preds_dict[K] == predicted))
            tots = np.where(labels[:, true_label] == 1)[0]
            #print('K=', predicted, ", idx=", true_label, "\nP -> ", interssect, "\nT -> ", tots)
            probabilities[true_label, predicted] = len(interssect) / len(tots)
            print('K=', predicted, ", idx=", true_label, ' -> Prob=%.2f' % probabilities[true_label, predicted])

    fig.show
    print("Run on the reduced dimensions data with d0 = ", d0)
    runKmeansCluster(X_reduced_matrix, Y_hot_labels, d0)

```

Run on the reduced dimensions data with d0 = 6

Data Dims: 2000 x 6

Dataset : (2000, 6)

----- K= 2 -----

K= 0 , idx= 0 -> Prob=0.00

K= 0 , idx= 1 -> Prob=0.80

K= 0 , idx= 2 -> Prob=0.00

K= 1 , idx= 0 -> Prob=1.00

K= 1 , idx= 1 -> Prob=0.20

K= 1 , idx= 2 -> Prob=1.00

```

----- K= 3 -----
-----

K= 0 , idx= 0   -> Prob=0.00
K= 0 , idx= 1   -> Prob=0.30
K= 0 , idx= 2   -> Prob=0.49
-----

K= 1 , idx= 0   -> Prob=1.00
K= 1 , idx= 1   -> Prob=0.00
K= 1 , idx= 2   -> Prob=0.51
-----

K= 2 , idx= 0   -> Prob=0.00
K= 2 , idx= 1   -> Prob=0.70
K= 2 , idx= 2   -> Prob=0.00

----- K= 4 -----
-----

K= 0 , idx= 0   -> Prob=0.00
K= 0 , idx= 1   -> Prob=0.51
K= 0 , idx= 2   -> Prob=0.00
-----

K= 1 , idx= 0   -> Prob=0.00
K= 1 , idx= 1   -> Prob=0.45
K= 1 , idx= 2   -> Prob=0.00
-----

K= 2 , idx= 0   -> Prob=0.00
K= 2 , idx= 1   -> Prob=0.04
K= 2 , idx= 2   -> Prob=0.56
-----

K= 3 , idx= 0   -> Prob=1.00
K= 3 , idx= 1   -> Prob=0.00
K= 3 , idx= 2   -> Prob=0.44

----- K= 5 -----
-----

K= 0 , idx= 0   -> Prob=0.00
K= 0 , idx= 1   -> Prob=0.30
K= 0 , idx= 2   -> Prob=0.46
-----

K= 1 , idx= 0   -> Prob=0.50

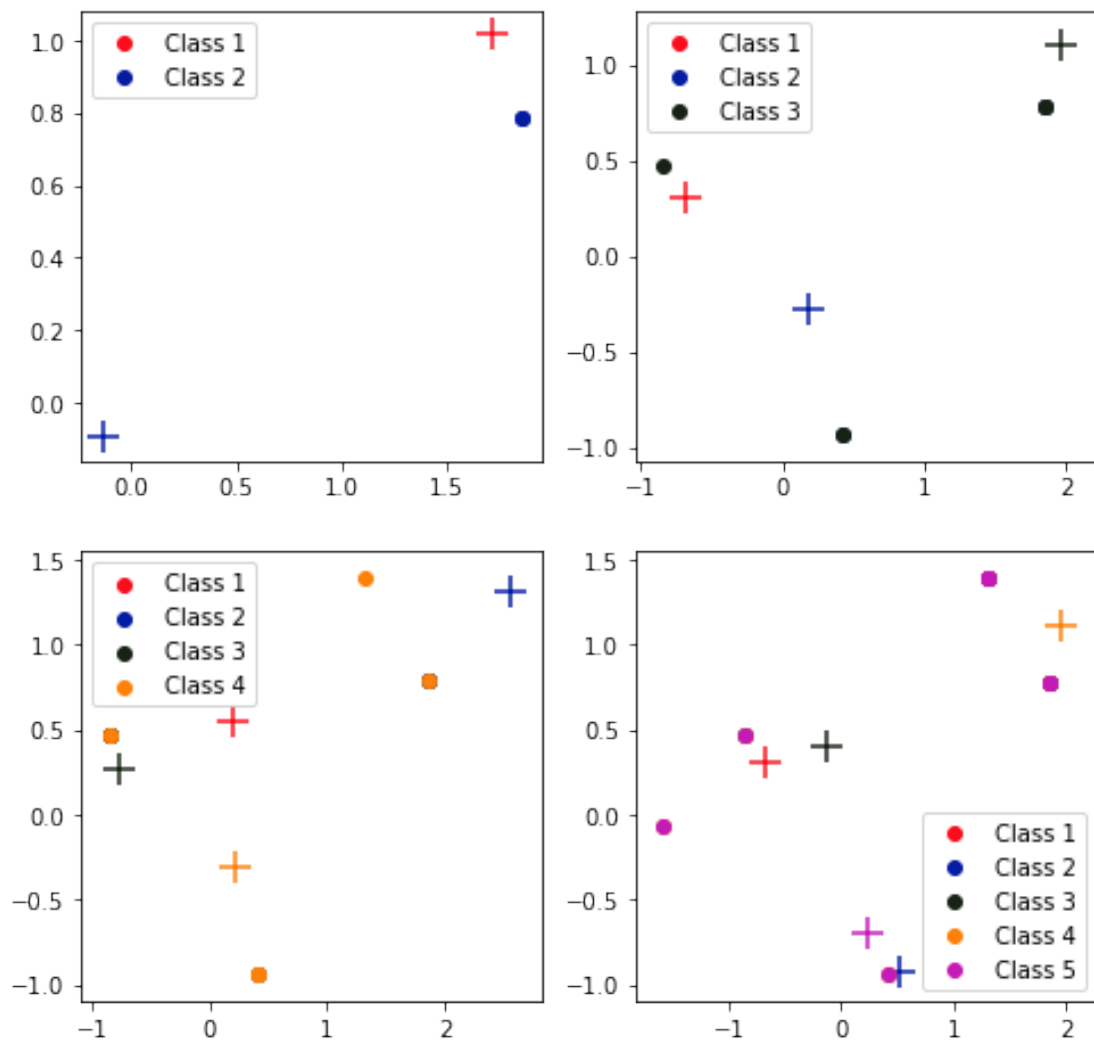
```


K= 1 , idx= 1 -> Prob=0.00
K= 1 , idx= 2 -> Prob=0.00

K= 2 , idx= 0 -> Prob=0.13
K= 2 , idx= 1 -> Prob=0.00
K= 2 , idx= 2 -> Prob=0.54

K= 3 , idx= 0 -> Prob=0.00
K= 3 , idx= 1 -> Prob=0.70
K= 3 , idx= 2 -> Prob=0.00

K= 4 , idx= 0 -> Prob=0.37
K= 4 , idx= 1 -> Prob=0.00
K= 4 , idx= 2 -> Prob=0.00



1.2 Part II : Random Projections and Compressed Sensing

1.3 3.

(a.) Generate $m \times d$ matrix, IID drawn such as $P[ij = +1] = 1/2$; $P[ij = -1] = 1/2$

(b.) Compressive Projection, $y = 1/\sqrt{m} * (\Phi x)$ with sparse reconstruction of s based on y

```
In [7]: def generatePhiMatrix(m, d):
        u = np.ndarray((m,d))
        for i in range(m):
            for j in range(d):
                r = np.random.rand()
                if(r<1/2):
                    u[i,j] = 1
                else:
                    u[i,j] = -1
        #print("(mxd):", u.shape)
        return u

def getCompressedProjection(m, d, log=False):
    # Keeping dimensions as dimensions d = 30 and number of data N = 50
    Xp_data, Zp_hot_labels = generateDataset(Uj, N, d)
    # Generate Phi
    phi = generatePhiMatrix(m,d)
    # Compressive Projection (m-dim projection of the d-dim matrix x)
    Y_xp = (1/np.sqrt(m)) * np.matmul(Xp_data, phi.T)
    B = np.transpose(Uj)
    if(log == True):
        print("> N =",N, ", M =",m, ", Y_xp:", Y_xp.shape, ", Labels:",Zp_hot_labels.shape)
    return Xp_data, Y_xp, Zp_hot_labels, phi, B

M = 20
X, Y_xp, labels, phi, B_evecs = getCompressedProjection(M, dims, True)

> N = 2000 , M = 20 , Y_xp: (2000, 20) , Labels: (2000, 3) , Phi : (20, 100) , B : (100, 7)
```

1.3.1 4. Lasso problem (using sklearn.linear_model.Lasso)

- (Φ) = projection matrix also called A in examples
- x = latent data variables with gaussian noise
- y = observed results (projection matrix is applied on latent data variables and the result has been reduced in dimensions)

```

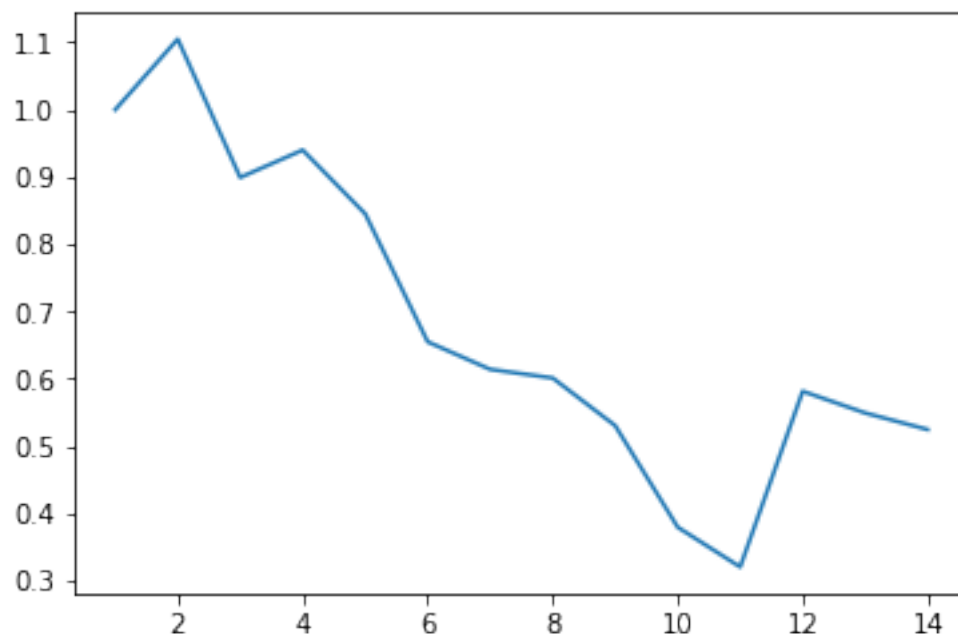
In [11]: from sklearn import linear_model

def normalizedME(s, s_hat):
    return np.linalg.norm(s_hat - s)/np.linalg.norm(s)

def findMinimumM():
    mses = []
    # Keeping dimension fixed and varying M for this
    rng = range(1,15)
    for m in rng:
        s, Y_xp, labels, phi, B = getCompressedProjection(m, dims, False)
        matrix = (1/np.sqrt(m)) * np.matmul(phi, B)
        clf = linear_model.Lasso(alpha = 1.0) # Set lambda ( called alpha here )
        clf.fit(matrix, Y_xp.T) # Fit the reduced Y to the one hot
        a_hat = clf.coef_ # Get a_hat
        s_hat = np.matmul(a_hat, B.T) # this is the output
        #print("Mat:",matrix.shape, ",\ts_hat:",s_hat.shape, ",\ts:",s.shape)
        mses.append(normalizedME(s, s_hat))
    plt.plot(list(rng), mses)
    plt.show()
    return np.argmin(mses)+1

minM = findMinimumM()
print("The minimum M is ", minM)

```



The minimum M is 11

1.3.2 5. Normalized MSE vs Lambda

```
In [37]: def normalize(lst):
    s = sum(lst)
    return map(lambda x: float(x)/s, lst)

def mean(lst):
    return sum(lst)/float(len(lst))

def normalizedMSE(s, s_hat):
    return (np.linalg.norm(s_hat - s)/np.linalg.norm(s))**2

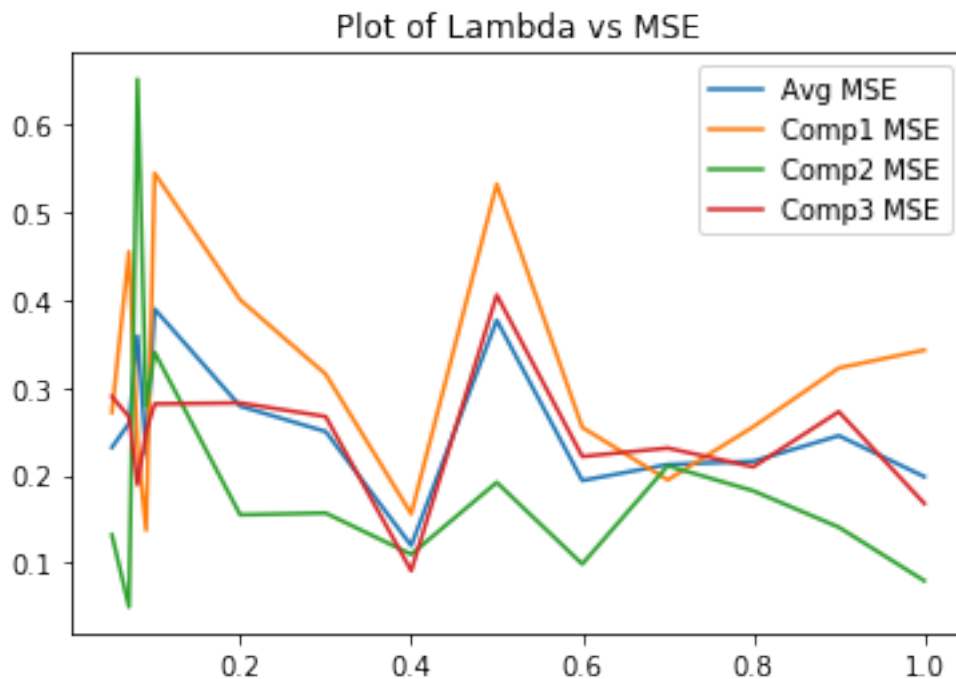
def computeMSELambda(N, m, d):
    mse_ = []
    mse_comp1_ = []
    mse_comp2_ = []
    mse_comp3_ = []
    s, Y_xp, labels, phi, B = getCompressedProjection(m, d, False)
    matrix = (1/np.sqrt(m)) * np.matmul(phi, B)
    clf = linear_model.Lasso(alpha = 1.0) # Set lambda ( called alpha here )
    clf.fit(matrix, Y_xp.T) # Fit the reduced Y to the one hot
    a_hat = clf.coef_ # Get a_hat
    s_hat = np.matmul(a_hat, B.T) # this is the output
    for i in range(N):
        mse = normalizedMSE(s[i],s_hat[i])
        index = np.where(labels[i]==1)[0]
        if index == 0: #component 1
            mse_comp1_.append(mse)
        if index == 1:
            mse_comp2_.append(mse)
        if index == 2:
            mse_comp3_.append(mse)
        mse_.append(mse)
    return mean(mse_), mean(mse_comp1_),mean(mse_comp2_),mean(mse_comp3_)

mse_list=[]
mse_comp1_list = []
mse_comp2_list = []
mse_comp3_list = []

lambda_list = [0.05,0.07,0.08,0.09,0.1,0.2,0.3,0.4,0.5,0.6,0.7,0.8,0.9,1.0]
for l in lambda_list:
    me, me1, me2, me3 = computeMSELambda(N, minM, dims)
    mse_list.append(me)
    mse_comp1_list.append(me1)
```

```
mse_comp2_list.append(me2)
mse_comp3_list.append(me3)
```

```
In [38]: plt.title('Plot of Lambda vs MSE')
plt.plot(lambda_list, mse_list, label="Avg MSE")
plt.plot(lambda_list, mse_comp1_list, label="Comp1 MSE")
plt.plot(lambda_list, mse_comp2_list, label="Comp2 MSE")
plt.plot(lambda_list, mse_comp3_list, label="Comp3 MSE")
plt.legend()
plt.show()
```



1.3.3 6. Compare the Euclidean distances squared vs the corresponding quantity in projected space

```
In [39]: m = minM
J = 6
B = np.transpose(Uj)
projected_u = (1/np.sqrt(m)) * np.matmul(phi, B)
projected_dist_matrix = np.zeros(shape=(J,J))
for i in range(J):
    for j in range(J):
        dist = np.linalg.norm(projected_u[:,i]-projected_u[:,j])**2
        projected_dist_matrix[i][j] = dist
print("projected_dist_matrix\n",projected_dist_matrix)
```

```

dist_matrix = np.zeros(shape=(J,J))
for i in range(J):
    for j in range(J):
        dist = np.linalg.norm(Uj[:,i]-Uj[:,j])**2
        dist_matrix[i][j] = dist
print("dist_matrix\n",dist_matrix)

projected_dist_matrix
[[ 0.    37.45  56.36  73.45  23.64  46.82]
 [ 37.45   0.    66.91  66.55  34.91  87.55]
 [ 56.36  66.91   0.   131.27  59.64 114.45]
 [ 73.45  66.55 131.27   0.    70.18  67.18]
 [ 23.64  34.91  59.64  70.18   0.    43.18]
 [ 46.82  87.55 114.45  67.18  43.18   0.   ]]
dist_matrix
[[ 0.  2.  2.  7.  3.  4.]
 [ 2.  0.  2.  3.  3.  2.]
 [ 2.  2.  0.  7.  3.  4.]
 [ 7.  3.  7.  0. 10.  7.]
 [ 3.  3.  3. 10.  0.  7.]
 [ 4.  2.  4.  7.  7.  0.]]

```

1.3.4 7. K-means algorithm post-projection

```

In [48]: x, labels = generateDataset(Uj, N, dims)
         phi = generatePhiMatrix(minM,dims)
         Y = np.matmul(phi,np.transpose(x))/(np.sqrt(minM))
         Z = np.transpose(Y)
         runKmeansCluster(Z,labels, minM)

```

```

Data Dims:  2000 x 11
Dataset   :  (2000, 11)

```

```

----- K= 2 -----

```

```

-----
K= 0 , idx= 0  -> Prob=0.69
K= 0 , idx= 1  -> Prob=0.63
K= 0 , idx= 2  -> Prob=0.27

```

```

-----
K= 1 , idx= 0  -> Prob=0.31
K= 1 , idx= 1  -> Prob=0.37
K= 1 , idx= 2  -> Prob=0.73

```

```

----- K= 3 -----

```

K= 0 , idx= 0 -> Prob=0.00
K= 0 , idx= 1 -> Prob=0.43
K= 0 , idx= 2 -> Prob=0.08

K= 1 , idx= 0 -> Prob=0.31
K= 1 , idx= 1 -> Prob=0.02
K= 1 , idx= 2 -> Prob=0.66

K= 2 , idx= 0 -> Prob=0.69
K= 2 , idx= 1 -> Prob=0.55
K= 2 , idx= 2 -> Prob=0.26

----- K= 4 -----

K= 0 , idx= 0 -> Prob=0.23
K= 0 , idx= 1 -> Prob=0.01
K= 0 , idx= 2 -> Prob=0.57

K= 1 , idx= 0 -> Prob=0.03
K= 1 , idx= 1 -> Prob=0.30
K= 1 , idx= 2 -> Prob=0.01

K= 2 , idx= 0 -> Prob=0.00
K= 2 , idx= 1 -> Prob=0.39
K= 2 , idx= 2 -> Prob=0.06

K= 3 , idx= 0 -> Prob=0.74
K= 3 , idx= 1 -> Prob=0.30
K= 3 , idx= 2 -> Prob=0.36

----- K= 5 -----

K= 0 , idx= 0 -> Prob=0.00
K= 0 , idx= 1 -> Prob=0.25
K= 0 , idx= 2 -> Prob=0.01

K= 1 , idx= 0 -> Prob=0.00
K= 1 , idx= 1 -> Prob=0.37
K= 1 , idx= 2 -> Prob=0.05

```

-----
K= 2 , idx= 0  -> Prob=0.47
K= 2 , idx= 1  -> Prob=0.28
K= 2 , idx= 2  -> Prob=0.13
-----

```

```

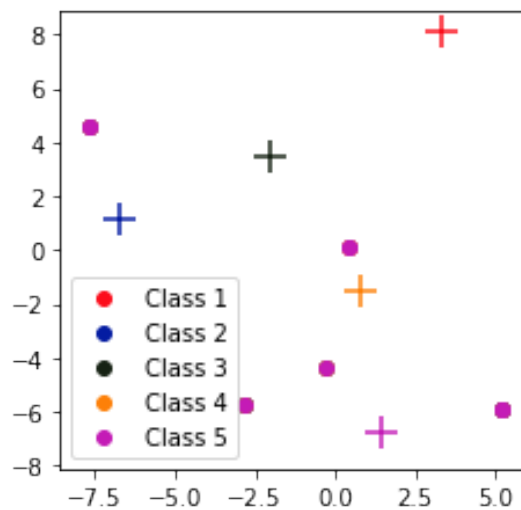
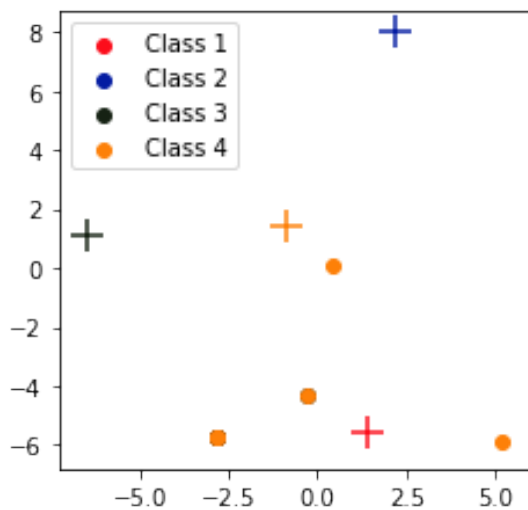
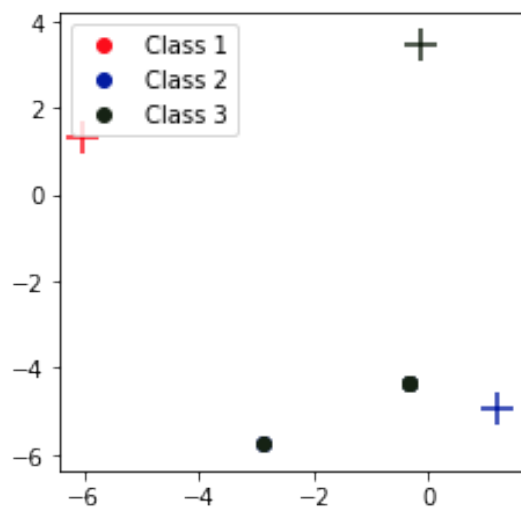
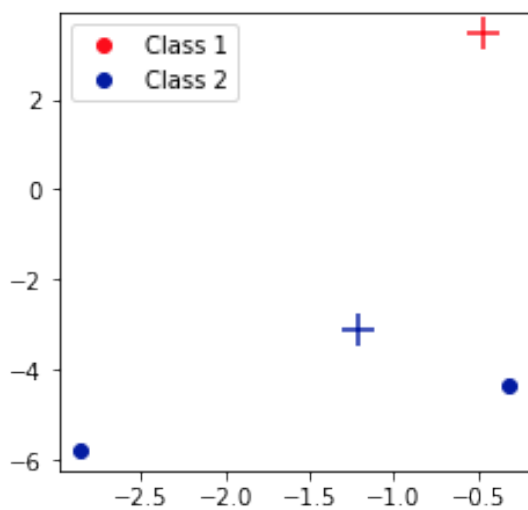
K= 3 , idx= 0  -> Prob=0.43
K= 3 , idx= 1  -> Prob=0.10
K= 3 , idx= 2  -> Prob=0.41
-----

```

```

K= 4 , idx= 0  -> Prob=0.10
K= 4 , idx= 1  -> Prob=0.00
K= 4 , idx= 2  -> Prob=0.40

```



1.3.5 8. Geometric insights

1.3.6 END