

# Report to Project-4

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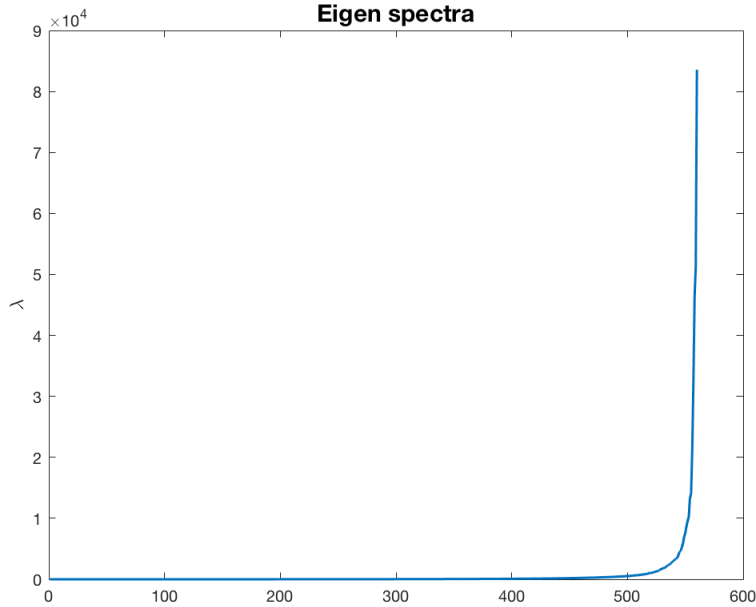


Figure 1: Eigen Spectra

# 1 Dimensionality Reduction

## 1.1 PCA

### 1.1.1 Choice for $k$

I write the script `select_k.m` to read data, convert to double, plot the eigen spectra, and select  $k$ . Figure 1 shows the eigen spectra of  $XX^T/N$  with first removing the mean.

We can observe that only a few eigen values are big, whereas the vast majority are small enough to be omitted. Such being the case, we conclude the main information of pictures are stored in those big eigen values. And therefore, discarding the small ones not only reduces the storage required for saving pictures, but also does good to denoising.

The retention rate of PCA can be defined as

$$rate = \frac{\sum_{i=560-k+1}^{560} \lambda_i}{\sum_{j=1}^{560} \lambda_j}$$

Thus we select  $k$  based on our expected retention rate. For example,  $k = 80$  at the retention rate of 95%, and  $k = 20$  at 80%. As my own preference,  $k = 80$  is a reasonable choice.

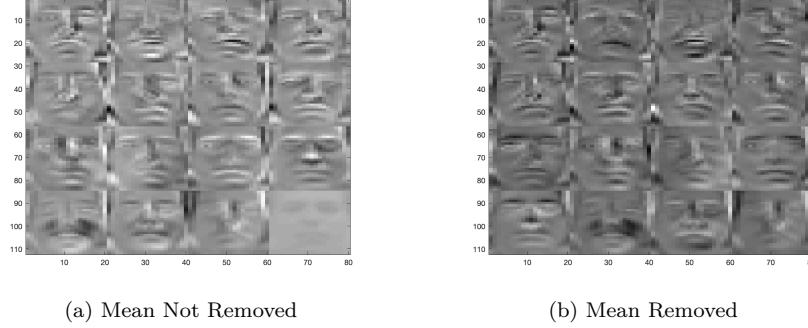


Figure 2: The Top 16 Eigenvectors

### 1.1.2 Visualize Eigenvectors

Figure 2, plotted by the script *showEigenvectors.m*, shows the top 16 eigenvectors with and without first removing the mean. As we can see, each picture characterizes certain features of the original photo, such as the eyes, brow, nose, mouth, moustache and so on. The features showed in each of group are in general corresponding to one of the other group. But the difference is, pictures in Figure 2(b) (with first removing the mean) specifies the details more distinctly than those in Figure 2(a). In conclusion, it is effective and efficient to first remove the mean.

### 1.1.3 Project onto 2D

The script *projectOnto2D.m* projects data onto the top two eigenvectors, and plots the resulting 2D points (Figure 3). As a consequence, the projection makes little sense since the outcome points comes too close to each other. The reason is evident—setting  $k = 2$  loses too much information of the original pictures. Recall that in Section 1.1.1, we have discussed how to select  $k$ .

### 1.1.4 Reconstruction

My function *PCAreconstruct* compose the corresponding projected vector of  $Y$ , where  $V$  is the eigenvectors used in PCA.

As discussed before, we select  $k = 80$  at the retention rate of 95%, and use first remove the mean. For the sake of comparison, my script *showReconstruct.m* plots both the original picture to be processed, and the reconstructed one. Take an informative picture from the data and a randomly generated data as examples, the results are in Figure 4 and Figure 5.

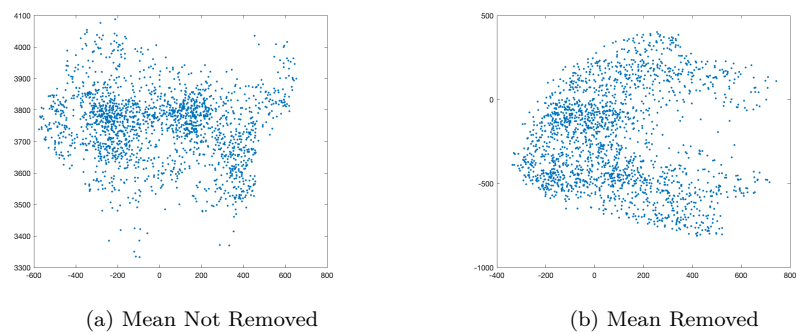


Figure 3: Project onto 2D

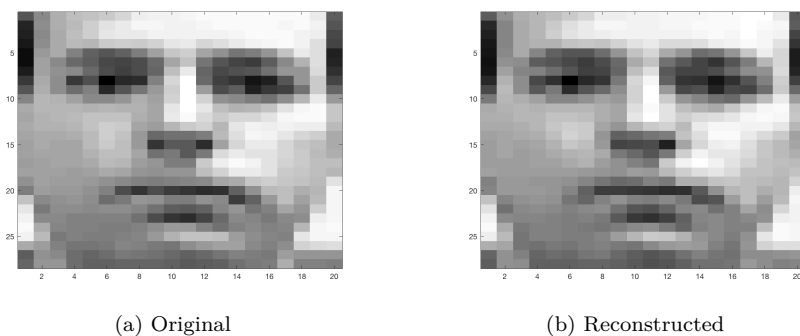


Figure 4: An Informative Example

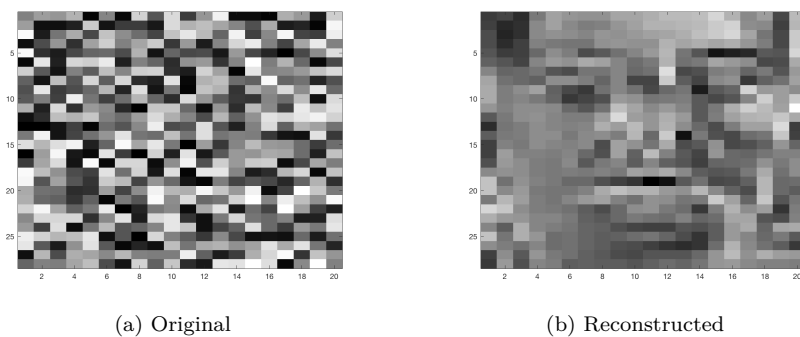


Figure 5: A Random Example

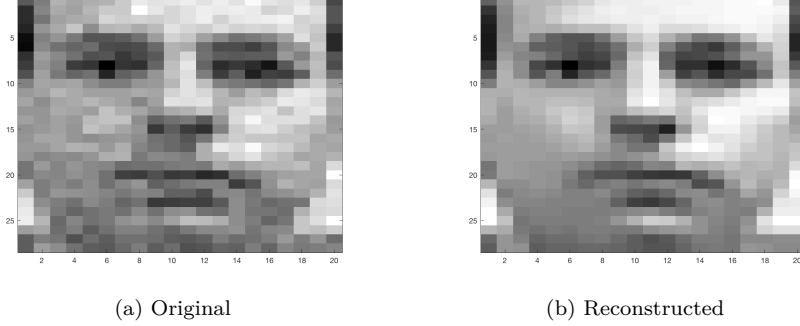


Figure 6: An Informative Example with Noise

### 1.1.5 Test with Noise

Add extra noise to the informative example in Section 1.1.4. See Figure 6, and we will observe the effect of denoising by PCA.

## 1.2 Isomap on Swiss Roll

The code *swissroll.m* provided by Roweis and Saul generates “swiss roll” data and runs LLE (Locally Linear Embedding) algorithm. I extract the data generation part of it and runs the isomap algorithm on the same data set. My script is called *swissroll\_isomap.m*. And the results are in Figure 7.

## 2 Gaussian Mixture Model

### 2.1 EM for Mixture of Gaussians

When  $\Sigma_k = \Sigma$  for all  $k$ , the expected complete-data log-likelihood function becomes

$$\mathbb{E}_z[\ln p(x, z | \mu, \Sigma, \pi)] = \sum_{n=1}^N \sum_{k=1}^K \gamma(z_{nk}) \{\ln \pi_k + \ln \mathcal{N}(x_n | \mu_k, \Sigma)\}$$

where

$$\gamma(z_{nk}) = \frac{\pi_k \mathcal{N}(x_n | \mu_k, \Sigma)}{\sum_{j=1}^K \pi_j \mathcal{N}(x_n | \mu_j, \Sigma)}$$

Differentiating it w.r.t  $\Sigma^{-1}$ , we obtain

$$\frac{N}{2} \Sigma - \frac{1}{2} \sum_{n=1}^N \sum_{k=1}^K \gamma(z_{nk}) (x_n - \mu_k)(x_n - \mu_k)^T$$

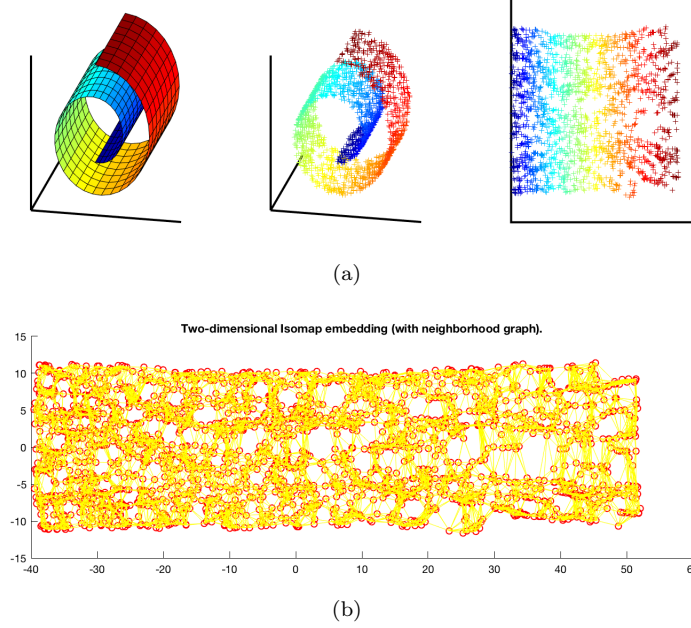


Figure 7: Swiss Roll

Note that we have used  $\sum_{k=1}^K \gamma(z_{nk}) = 1$  for all  $n$ . Setting this to zero, we finally derive

$$\Sigma = \frac{1}{N} \sum_{n=1}^N \sum_{k=1}^K \gamma(z_{nk}) (x_n - \mu_k)(x_n - \mu_k)^T$$

## 2.2 Training

We first experiment with *randConst* in the function *mogEM*. I write script *setRandConst.m* to plot  $\log P(\text{TrainingData})$  against *randConst* (Figure 8). As a result, *randConst* = 1 is a reasonable choice.

Run several times the below codes to train good models.

```

1 [p2,mu2,vary2,logProbX3] = mogEM(train2,2,10,0.01,1);
2 [p3,mu3,vary3,logProbX3] = mogEM(train3,2,10,0.01,1);

```

The final model parameters are separately stored in *model2.mat* and *model3.mat*.

Both the mean and variance vectors are shown as images in Figure 9 and Figure 10 using function *visualize\_digits*. The mixing proportions for the clusters within *model2* are

p =

0.4900

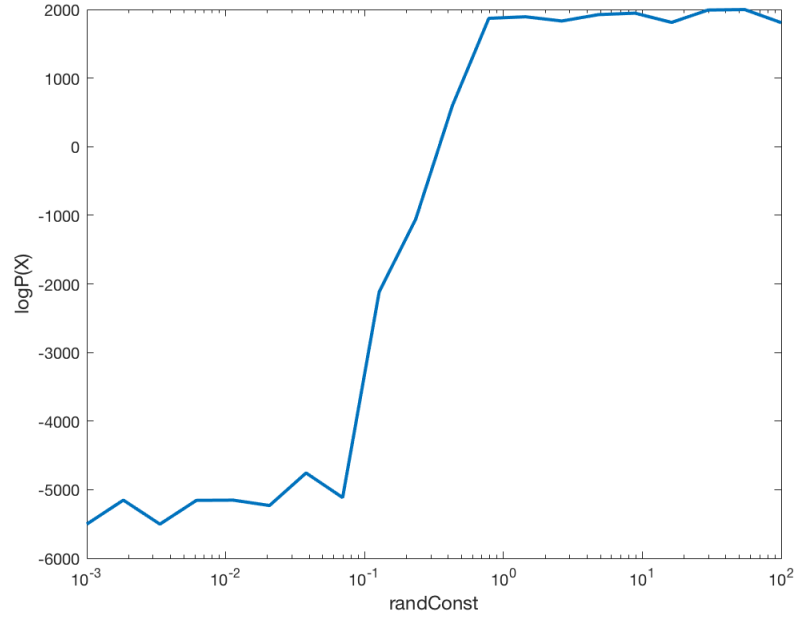


Figure 8:  $\log P(X)$  Against  $randConst$

0.5100

The mixing proportions for the clusters within *model3* are

p =

0.4769

0.5231

Finally,  $\log P(TrainingData)$  is computed by

1 `sum(mogLogProb(p,mu,vary,x));`

The results of two models are respectively

-3.8202e+03

and

2.2582e+03

## 2.3 Initialize with K-Means

Function *mogEM\_kmeans* modified by me does the same thing as *mogEM* does, except that *mu* is initialized with k-means algorithm instead of in random.



(a) Mean

(b) Variance

Figure 9: Mean and Variance of Model 2



(a) Mean

(b) Variance

Figure 10: Mean and Variance of Model 3



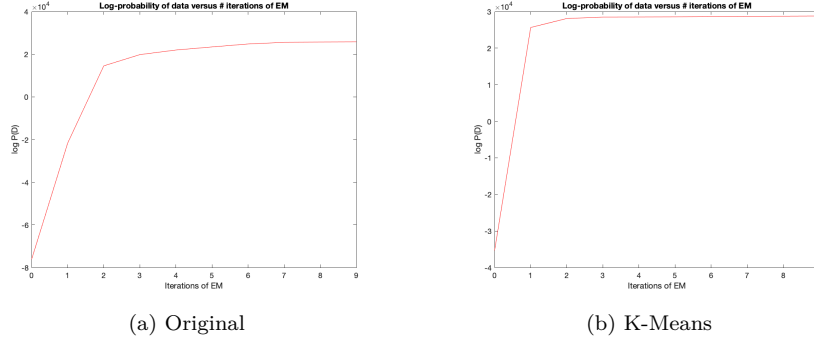


Figure 11: Converging Process

Run `run_q4.m`, and the results are stored in `q4.m`. The two converging processes with and without using k-means are shown in Figure 11. Note that Figure 11(a) converges at 5th iteration, whereas Figure 11(b) only needs 2 iterations.

The final *log – prob*'s are separately

2.4673e+04

and

2.7755e+04

As a consequence, initializing with k-means algorithm provides much more efficiency, and is less likely to be trapped in local optimization.

## 2.4 Classification Using MoGs

`run_q5.m` plots the error rate against the number of clusters (Figure 12). The model with each number of clusters has been trained for 20 times, and the error rate is computed averagely.

Answers:

1. The error rates on the training sets generally decrease as the number of clusters increases, because more patterns are recognized. The more clusters, the more fitting for training data.
2. The error rate curve for the test set decreases rapidly at the beginning, but slowly when the number of clusters is big. These phenomena—fitting and overfitting appear just like in any other machine learning models. A small number of clusters may not include enough necessary patterns, whereas too many clusters causes overfitting.

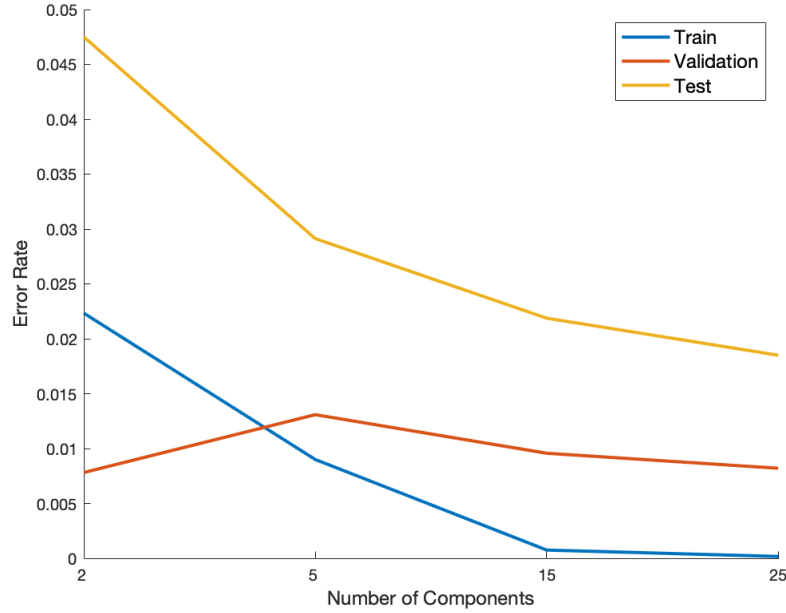


Figure 12: Error Rate

3. If I wanted to choose a particular model as the best, I would certainly choose the 25-cluster model according to my above experiments. But if my aim is to achieve the lowest error rate possible on the new images my system will receive, which is obviously more meaningful, I will select the 15-cluster model as my final model. Because as we see in Figure 12, there is no much difference between 15 and 20 clusters. In this way, we shall use 15 in order to avoid overfitting.

## A MATLAB Codes

### A.1 Dimensionality Reduction

#### A.1.1 *select\_k.m*

```

1 % Edited by Yuanbo Han, Dec. 22, 2017.
2
3 load freyface.mat
4 X = double(X); % convert to double
5
6 % Compute like SVD
7 N = size(X, 2);

```

```

8 [Vun, Dun] = eig(X*X'/N);
9 [lambda_un, order] = sort(diag(Dun));
10 Vun = Vun(:, order);
11 Xctr = X - repmat(mean(X, 2), 1, N);
12 [Vctr, Dctr] = eig(Xctr*Xctr'/N);
13 [lambda_ctr, order] = sort(diag(Dctr));
14 Vctr = Vctr(:, order);
15
16 % Eigen spectra (plot lambda_ctr)
17 figure;
18 plot(lambda_ctr, 'LineWidth', 1.5);
19 ylabel('\lambda', 'FontSize', 12);
20 title('Eigen spectra', 'FontSize', 15);
21
22 % Select k based on retention rate
23 original = sum(lambda_ctr);
24 new = 0;
25 r = 0.95; % retention rate
26 for k=1:560
27     new = new + lambda_ctr(end-k+1);
28     if new / original >= r
29         break;
30     end
31 end
32 fprintf('k = %d, at the retention rate of %.2f%%\n', k, r*100);

```

### A.1.2 *showEigenvectors.m*

```

1 % Show the top 16 eigenvectors with and without first removing the mean.
2 figure;
3 showfreyface(Vun(:, end-15:end));
4 figure;
5 showfreyface(Vctr(:, end-15:end));

```

### A.1.3 *projectOnto2D.m*

```

1 % Project the data onto the top two eigenvectors,
2 % and plot the resulting 2D points.
3 Yctr = Vctr(:, end-1:end)' * X;
4 plot(Yctr(1,:), Yctr(2,:), '.');
5 explorefreymanifold(Yctr, X);
6
7 Yun = Vun(:, end-1:end)' * X;
8 plot(Yun(1,:), Yun(2,:), '.');
9 explorefreymanifold(Yun, X);

```

#### A.1.4 PCAreconstruct.m

```
1 function [X_re] = PCAreconstruct(Y,V)
2 % PCARECONSTRUCT(Y,V) compose the corresponding projected vector of Y,
3 % where V is the eigenvectors used in PCA.
4 %
5 % Edited by Yuanbo Han, Dec. 25, 2017.
6 X_re = V * Y;
7 end
```

#### A.1.5 showReconstruct.m

```
1 % Edited by Yuanbo Han, Dec. 25, 2017.
2
3 k = 80;
4 V = Vctr(:,end-k+1:end);
5
6 % Example for an original picture
7 figure;
8 showfreyface(X(:,50)); % the 50th sample in the data
9 figure;
10 showfreyface(PCAreconstruct(V'*(X(:,50)-mean(X,2)),V)+mean(X,2));
11
12 % Example for a random picture
13 R = randi([1,255], 560, 1);
14 figure;
15 showfreyface(R);
16 figure;
17 showfreyface(PCAreconstruct(V'*(R-mean(X,2)),V)+mean(X,2));
18
19 % Add noise for an original picture
20 X_noise = X(:,50) + 10*randn(560,1); % the 50th sample in the data
21 figure;
22 showfreyface(X_noise);
23 figure;
24 showfreyface(PCAreconstruct(V'*(X_noise-mean(X,2)),V)+mean(X,2));
```

#### A.1.6 distmat.m

```
1 function [dist] = distmat(p, q)
2 % function [dist] = distmat(v1, v2)
3 %
4 % Calculates pairwise distances between vectors. If v1 and v2 are both
5 % provided, a size(v1, 1) by size(v2, 1) matrix is returned, where the
```

```

6 % entry at (i,j) contains the Euclidean distance from v1(i,:) to v2(j,:).
7 % If only v1 is provided, squareform(pdist(v1)) is returned.
8
9 p = p';
10 if nargin == 1
11     q = p;
12 else
13     q = q';
14 end
15
16 [~, pn] = size(p);
17 [~, qn] = size(q);
18
19 pmag = sum(p .* p, 1);
20 qmag = sum(q .* q, 1);
21 dist = repmat(qmag, pn, 1) + repmat(pmag', 1, qn) - 2*p'*q;
22 dist = sqrt(dist);
23 end

```

#### A.1.7 *swissroll\_isomap.m*

```

1 % SWISS ROLL DATASET
2
3 N=2000;
4 K=12;
5 d=2;
6
7 clf; colordef none; colormap jet; set(gcf, 'Position', [200,400,620,200]);
8
9 % PLOT TRUE MANIFOLD
10 tt0 = (3*pi/2)*(1+2*[0:0.02:1]); hh = [0:0.125:1]*30;
11 xx = (tt0.*cos(tt0))'*ones(size(hh));
12 yy = ones(size(tt0))'*hh;
13 zz = (tt0.*sin(tt0))'*ones(size(hh));
14 cc = tt0'*ones(size(hh));
15
16 subplot(1,3,1); cla;
17 surf(xx,yy,zz,cc);
18 view([12 20]); grid off; axis off; hold on;
19 lnx=-5*[3,3,3;3,-4,3]; lny=[0,0,0;32,0,0]; lnz=-5*[3,3,3;3,3,-3];
20 lnh=line(lnx,lny,lnz);
21 set(lnh, 'Color',[1,1,1], 'LineWidth',2, 'LineStyle','-', 'Clipping','off');
22 axis([-15,20,0,32,-15,15]);
23
24 % GENERATE SAMPLED DATA

```

```

25 tt = (3*pi/2)*(1+2*rand(1,N)); height = 21*rand(1,N);
26 X = [tt.*cos(tt); height; tt.*sin(tt)];
27
28 % SCATTERPLOT OF SAMPLED DATA
29 subplot(1,3,2); cla;
30 scatter3(X(1,:),X(2,:),X(3,:),12,tt,'+');
31 view([12 20]); grid off; axis off; hold on;
32 lnh=line(lnx,lny,lnz);
33 set(lnh,'Color',[1,1,1],'LineWidth',2,'LineStyle','-','Clipping','off');
34 axis([-15,20,0,32,-15,15]); drawnow;
35
36 % RUN ISOMAP ALGORITHM
37 options.dims = 1:2;
38 Y = isomap(distmat(X'),'k',K,options);
39
40 % SCATTERPLOT OF EMBEDDING
41 figure(1);
42 subplot(1,3,3); cla;
43 Y_2 = Y.coords{2};
44 scatter(Y_2(1,:),Y_2(2,:),12,tt,'+');
45 grid off;
46 set(gca,'XTick',[]); set(gca,'YTick',[]);

```

## A.2 Gaussian Mixture Model

### A.2.1 mogEM\_test.m

```

1 function [p,mu,vary,logProbX] = ...
2     mogEM_test(x,K,itors,minVary,plotFlag,randConst)
3 % MOGEM_TEST does the same thing as MOGEM does, except that randConst can
4 % be initialized outside.
5 %
6 % Modified by Yuanbo Han, Dec. 26, 2017.
7
8 if nargin==3 minVary = 0; plotFlag = 0; end
9 N = size(x,1); T = size(x,2);
10
11 plotString = cellstr(['bo'; 'gx'; 'r+'; 'c*'; 'ms'; 'yd'; 'kd']);
12 ellColor = ['b'; 'g'; 'r'; 'c'; 'm'; 'y'; 'k'];
13
14 % Initialize the parameters
15 %randConst = 1;
16 p = randConst+rand(K,1); p = p/sum(p);
17 mn = mean(x,2);
18 vr = std(x,[],2).^2;

```

```

19
20 %----- Modify the code here -----
21 % Change the random initialization with k-means algorithm, use 5
22 % iterations.
23 mu = mn*ones(1,K)+randn(N,K).*(sqrt(vr)/randConst*ones(1,K));
24 %mu = kmeans(x,K,5);
25 %-----
26 vary = vr*ones(1,K)*2;
27 vary = (vary>=minVary).*vary + (vary<minVary)*minVary;
28
29 % Do iters iterations of EM
30 logProbX = zeros(iters,1);
31
32 for i=1:iters
33     % Do the E step
34     respTot = zeros(K,1); respX = zeros(N,K);
35     respDist = zeros(N,K); logProb = zeros(1,T);
36     ivary = 1./vary;
37     logNorm = log(p)-0.5*N*log(2*pi)-0.5*sum(log(vary'),2);
38     logPcAndx = zeros(K,T);
39     for k=1:K
40         logPcAndx(k,:) = logNorm(k) - 0.5*...
41             sum((ivary(:,k)*ones(1,T)).*(x-mu(:,k)*ones(1,T)).^2,1);
42     end
43     [mx, mxi] = max(logPcAndx,[],1);
44     PcAndx = exp(logPcAndx-ones(K,1)*mx); Px = sum(PcAndx,1);
45     PcGivenx = PcAndx./(ones(K,1)*Px); logProb = log(Px) + mx;
46     logProbX(i) = sum(logProb);
47
48     % Plot log prob of data
49     %f
50     figure(1);
51     set(gcf,'DoubleBuffer','on')
52     clf;
53     plot([0:i-1],logProbX(1:i),'r-');
54     title('Log-probability of data versus # iterations of EM');
55     xlabel('Iterations of EM');
56     ylabel('log P(D)');
57     drawnow;
58
59     if plotFlag % Plot the data and Gaussians
60         figure(2);
61         set(gcf,'DoubleBuffer','on')
62         clf;
63         hold on;
64         for k=1:K

```

```

65         plotEllipse(mu(1,k),mu(2,k),vary(1,k),vary(2,k),0,...
66                     ellColor(mod(k, length(ellColor))+1));
67     end;
68     for t=1:T plot(x(1,t),x(2,t),...
69                 char(plotString(mod(mx(t), length(plotString))+1))); end
70
71     axis equal;
72 end;
73 %}
74
75 respTot = mean(PcGivenx,2);
76 respX = zeros(N,K); respDist = zeros(N,K);
77 for k=1:K
78     respX(:,k) = mean(x.*(ones(N,1)*PcGivenx(k,:)),2);
79     respDist(:,k) = ...
80         mean((x-mu(:,k)*ones(1,T)).^2.*(ones(N,1)*PcGivenx(k,:)),2);
81 end
82
83 % Do the M step
84 p = respTot;
85 mu = respX./(ones(N,1)*respTot'+eps);
86 vary = respDist./(ones(N,1)*respTot'+eps);
87 vary = (vary>=minVary).*vary + (vary<minVary)*minVary;
88
89 end
90 end

```

### A.2.2 *setRandConst.m*

```

1  % Edited by Yuanbo Han, Dec. 26, 2017.
2  load digits.mat;
3
4  randConst = logspace(-3,2,20);
5  iters = 20;
6  repeat = 20;
7  logProbX_temp = zeros(iters,repeat);
8  logProbX = zeros(1,length(randConst));
9  for i = 1:length(randConst)
10     fprintf('randConst = %f\n', randConst(i));
11     for j = 1:repeat
12         [~,~,~,logProbX_temp(:,j)] = ...
13             mogEM_test(train3,2,iters,0.01,1,randConst(i));
14     logProbX(i) = mean(logProbX_temp(end,:));
15 end
16 fprintf('logProbX = %f\n', logProbX(i));

```



```

17 end
18
19 figure;
20 semilogx(randConst, logProbX, 'LineWidth', 2);
21 xlabel('randConst', 'FontSize', 12);
22 ylabel('logP(X)', 'FontSize', 12);

```

### A.2.3 *mogEM\_kmeans.m*

```

1 function [p,mu,vary,logProbX] = mogEM_kmeans(x,K,itors,minVary,plotFlag)
2 % MOGEM_KMEANS does the same thing as MOGEM does, except that mu is
3 % initialized with k-means algorithm instead of in random.
4 %
5 % Modifiedy by Yuanbo Han, Dec. 26, 2017.
6
7 if nargin==3
8     minVary = 0;
9     plotFlag = 0;
10 end
11
12 N = size(x,1); T = size(x,2);
13
14 plotString = cellstr(['bo'; 'gx'; 'r+'; 'c*'; 'ms'; 'yd'; 'kd']);
15 ellColor = ['b'; 'g'; 'r'; 'c'; 'm'; 'y'; 'k'];
16
17 % Initialize the parameters
18 randConst = 1;
19 p = randConst+rand(K,1); p = p/sum(p);
20 mn = mean(x,2);
21 vr = std(x,[],2).^2;
22
23 %----- Modify the code here -----
24 % Change the random initialization with k-means algorithm, use 5
25 % iterations.
26 %mu = mn*ones(1,K)+randn(N,K).*(sqrt(vr)/randConst*ones(1,K));
27 mu = kmeans(x,K,5);
28 %-----
29 vary = vr*ones(1,K)*2;
30 vary = (vary>=minVary).*vary + (vary<minVary)*minVary;
31
32 % Do iters iterations of EM
33 logProbX = zeros(itors,1);
34
35 for i=1:itors
36     % Do the E step

```

```

37     respTot = zeros(K,1); respX = zeros(N,K);
38     respDist = zeros(N,K); logProb = zeros(1,T);
39     ivary = 1./vary;
40     logNorm = log(p)-0.5*N*log(2*pi)-0.5*sum(log(vary'),2);
41     logPcAndx = zeros(K,T);
42     for k=1:K
43         logPcAndx(k,:) = logNorm(k)- 0.5*...
44             sum((ivary(:,k)*ones(1,T)).*(x-mu(:,k)*ones(1,T)).^2,1);
45     end
46     [mx, mxi] = max(logPcAndx,[],1);
47     PcAndx = exp(logPcAndx-ones(K,1)*mx); Px = sum(PcAndx,1);
48     PcGivenx = PcAndx./(ones(K,1)*Px); logProb = log(Px) + mx;
49     logProbX(i) = sum(logProb);
50
51     % Plot log prob of data
52     %f
53     figure(1);
54     set(gcf,'DoubleBuffer','on')
55     clf;
56     plot([0:i-1],logProbX(1:i),'r-');
57     title('Log-probability of data versus # iterations of EM');
58     xlabel('Iterations of EM');
59     ylabel('log P(D)');
60     drawnow;
61
62     if plotFlag % Plot the data and Gaussians
63         figure(2);
64         set(gcf,'DoubleBuffer','on')
65         clf;
66         hold on;
67         for k=1:K
68             plotEllipse(mu(1,k),mu(2,k),vary(1,k),vary(2,k),0,...
69                 ellColor(mod(k, length(ellColor))+1));
70         end;
71         for t=1:T plot(x(1,t),x(2,t),...
72             char(plotString(mod(mxi(t), length(plotString))+1))); end
73
74         axis equal;
75     end;
76     %f
77
78     respTot = mean(PcGivenx,2);
79     respX = zeros(N,K); respDist = zeros(N,K);
80     for k=1:K
81         respX(:,k) = mean(x.*(ones(N,1)*PcGivenx(k,:)),2);
82         respDist(:,k) = ...

```

```

83         mean((x-mu(:,k)*ones(1,T)).^2.*(ones(N,1)*PcGivenx(k,:)),2);
84     end
85
86     % Do the M step
87     p = respTot;
88     mu = respX./(ones(N,1)*respTot'+eps);
89     vary = respDist./(ones(N,1)*respTot'+eps);
90     vary = (vary>=minVary).*vary + (vary<minVary)*minVary;
91
92 end
93 end

```

#### A.2.4 run\_q4.m

```

1  % Completed by Yuanbo Han, Dec. 26, 2017.
2  load digits;
3  x = [train2, train3];
4
5  % Train a MoG model with 20 components on all 600 training vectors
6  % with both original initialization and your kmeans initialization.
7
8  % Original initialization
9  [p1,mu1,vary1,logProbX1] = mogEM(x,20,10,0.01,0);
10 logProb1 = sum(mogLogProb(p1,mu1,vary1,x))
11
12 % K-means initialization
13 [p2,mu2,vary2,logProbX2] = mogEM_kmeans(x,20,10,0.01,0);
14 logProb2 = sum(mogLogProb(p2,mu2,vary2,x))

```

#### A.2.5 run\_q5.m

```

1  % Completed by Yuanbo Han, Dec. 26, 2017.
2  load digits;
3
4  errorTrain = zeros(1, 4);
5  errorValidation = zeros(1, 4);
6  errorTest = zeros(1, 4);
7  numComponent = [2, 5, 15, 25];
8  maxIter = 20; repeat = 20;
9
10 for i = 1 : 4
11     K = numComponent(i);
12     for r = 1:repeat
13         % Train a MoG model with K components for digit 2

```

```

14     [p2,mu2,vary2,~] = mogEM_kmeans(train2 , K, maxIter , 0.01, 0);
15
16     % Train a MoG model with K components for digit 3
17     [p3,mu3,vary3,~] = mogEM_kmeans(train3 , K, maxIter , 0.01, 0);
18
19     % Caculate the probability  $P(d=1/x)$  and  $P(d=2/x)$ ,
20     % classify examples, and compute the error rate
21     % Hints: you may want to use mogLogProb function
22     [inputs_train , inputs_valid , inputs_test , ...
23         target_train , target_valid , target_test] = load_data();
24
25     P2GivenTrain = mogLogProb(p2,mu2,vary2,inputs_train);
26     P3GivenTrain = mogLogProb(p3,mu3,vary3,inputs_train);
27     train_label = (P3GivenTrain > P2GivenTrain);
28     errorTrain(i) = errorTrain(i) + 1/repeat * ...
29         sum(train_label ~= target_train) / length(inputs_train);
30
31     P2GivenValid = mogLogProb(p2,mu2,vary2,inputs_valid);
32     P3GivenValid = mogLogProb(p3,mu3,vary3,inputs_valid);
33     valid_label = (P3GivenValid > P2GivenValid);
34     errorValidation(i) = errorValidation(i) + 1/repeat * ...
35         sum(valid_label ~= target_valid) / length(inputs_valid);
36
37     P2GivenTest = mogLogProb(p2,mu2,vary2,inputs_test);
38     P3GivenTest = mogLogProb(p3,mu3,vary3,inputs_test);
39     test_label = (P3GivenTest > P2GivenTest);
40     errorTest(i) = errorTest(i) + 1/repeat * ...
41         sum(test_label ~= target_test) / length(inputs_test);
42     end
43 end
44
45 % Plot the error rate
46 figure;
47 hold on;
48 plot(1:4, errorTrain , 'LineWidth', 2);
49 plot(1:4, errorValidation , 'LineWidth', 2);
50 plot(1:4, errorTest , 'LineWidth', 2);
51 set(gca, 'XTick', 1:4);
52 set(gca, 'XTickLabel', numComponent);
53 lgd = legend({'Train', 'Validation', 'Test'}, 'Location', 'NorthEast');
54 set(lgd, 'FontSize', 12);
55 xlabel('Number of Components', 'FontSize', 12);
56 ylabel('Error Rate', 'FontSize', 12);

```