# MLPR Assignment

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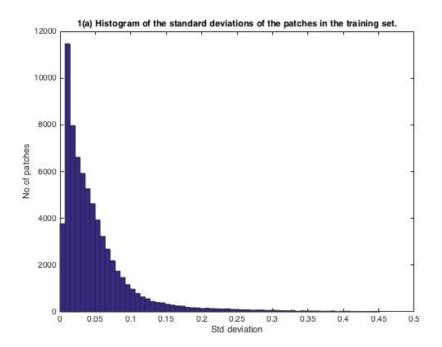
24th Nov 2015

## 1. The Next Pixel Prediction Task

## Q1 Data preprocessing and visualization

(a)

Figure 1: Histogram



#### Observations from histogram:

1. Many patches have low deviations eg. roughly 3900 patches have 0 std deviations whereas roughly 11,000 out of 70,000 patches have 0.01 std deviation, indicating that a lot of these pixels values are similar, (smooth patches).

- 2. Since we assume all patches with std dev < 0.063(4/63) are flat patches, from the plot is can be inferred that more than half of the patches are flat.
- 3. Some patches have very high deviations as well. of the standard deviations of the patches in the training set indicating that they can help in modelling (interesting patches).

Here 64 bins is used because the image has been discretized by 64 grey scales and this way each bin will plot each grey scale value for the image.

## (b)

A simple way is to predict a flat patch is by comparing the y(j) pixel with the mean of all pixels in this same patch. It can also be done by comparing the mean of all pixels of this patch with the mean of x(j,end), x(j,end-34) and y(j). If both values are very similar, the chances of the patch being flat is high.

## (c)

(c) Figures for flat and non-flat image patch is given below. Matlab code at the end of this section.

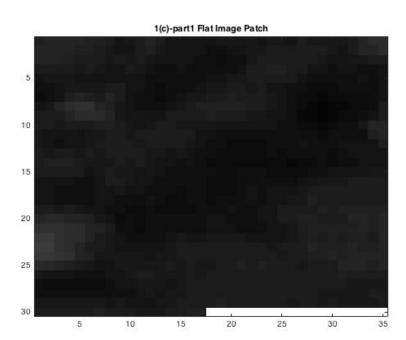
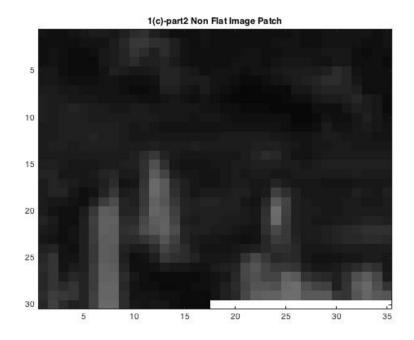


Figure 2

## Matlab Code - Q1.1

Figure 3



```
load('imgregdata.mat');
  % First scale all of the data to [0, 1] by dividing each pixel value by
       63.
  xtr sc=xtr/63;
  ytr_sc=ytr/63;
  xte_sc=xte/63;
  yte_sc=yte/63;
  %compute the standard deviation of each x(j, :) patch
  xtr_sd=std(xtr_sc');
  % 1(a) - Plot a histogram
  figure
  hist(xtr_sd,64)
  xlabel('Std deviation');
  ylabel ('No of patches');
  title ({ '1(a) Histogram of the standard deviations of the patches in the
       training set.'});
  rows = size(xtr_sc, 1);
  % as adviced in the assignment sheet.
  threshold = 4/63;
17
  for i = 1 : rows
       if (xtr sd(i) <= threshold)</pre>
```

```
patch f=xtr sc(i,:);
           break;
21
22
       end
  end
23
  Membed patch with 1's s, increase no of cols from 1032 to 1050 first
  patch embed=horzcat (patch f, ones (1,18));
  % now reshape into (35X30 piel size)
  patch reshaped=reshape(patch embed, 35, 30);
  figure
28
  colormap gray;
  imagesc(patch reshaped',[0,1]);
   title ('1(c)-part1 Flat Image Patch');
   for i = 1 : rows
32
       if (xtr sd(i) > threshold)
           patch nf=xtr sc(i,:);
34
           break;
       end
36
  end
37
  %embed patch with 1's s, increase no of cols from 1032 to 1050 first
  patch embed=horzcat (patch nf, ones (1,18));
  % now reshape into (35X30 piel size)
  patch reshaped=reshape(patch embed, 35, 30);
  figure
  colormap gray;
  imagesc(patch reshaped',[0,1]);
  title ('1(c)-part2 Non Flat Image Patch')
```

#### Q1.2 Linear regression with adjacent pixels

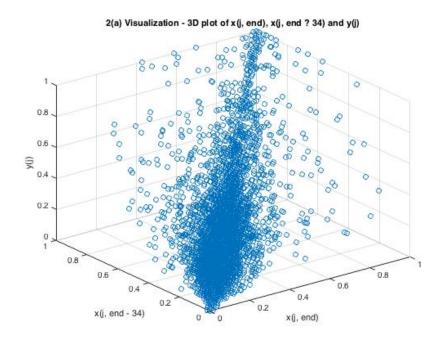
(a)

The 3D plot of x(j, end), x(j, end? 34) and y(j) is presented in Figure 4. For this 3D plot, the data has been subsampled by taking every other row from both datasets Both subsampled sets had roughly 8600 instances after this operation. Strong positive linear correlations between all the 3 pixels are visible in this scatter plot. Both pixels x(j;end) and x(j:end-34) can help in predicting pixel y(j) because of this behaviour. This also indicates that linear regression might be a good technique to explore the interesting statistical trends in this image.

(b)

Bias weight has been added as 3nd column to matrix of x(j, end) and x(j, end - 34), and denoted as X. Let Y represent the vector for y(j) values. MLE solution  $\mathbf{w} = (\mathbf{inv}(\mathbf{X}^{\top} \mathbf{X}))\mathbf{X}^{\top}\mathbf{y}$  The inv() function above is exponent to the power -1 (the inverse function).

Figure 4: .



(c)

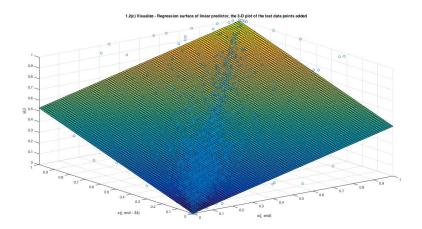
The linear regression predictor is implemented in matlab as shown in code below. The weight vector values are:  $\mathbf{w} = [\ 0.46064\ ;\ 0.52412\ ;\ 0.00256\ ]$ . The root mean squared error on the training and test sets are RMSE training = 0.0506 and RMSE test = 0.0503. Both values are very similar which indicates that the model is a good model and the training data was not over fitted. Had that been the case, RMSE of the test would have been much higher that RMSE of the training set.

The regression surface of this linear predictor in 3-D using Matlab function surf() is in Figure 5 below. From this plot, the data is linearly separable as almost all test points like above this regression surface. which means that the linear classifier is successful in predicting the pixel values(y(j)) given the training set of pixels.

#### Matlab Code - Q1.2- Linear Regression - 2 variables

```
load imgregdata.mat xte_nf yte_nf xtr_nf ytr_nf
% 1.2(a) - Subsampled by taking every other row from both datasets
% Both subsampled sets have roughly 8600 instances
xtr_ss=xtr_nf(1:2:end,:);
ytr_ss=ytr_nf(1:2:end,:);
```

Figure 5



```
figure
   plot (xtr ss (:, end-34), xtr ss (:, end), 'r:+');
   figure
10
   scatter3(xtr_ss(:,end),xtr_ss(:,end-34),ytr_ss);
   title (\{'2(a) \text{ Visualization } -3D \text{ plot of } x(j, \text{ end}), x(j, \text{ end }? 34) \text{ and } y
      (j)'});
   xlabel('x(j, end)');
   ylabel('x(j, end - 34)');
   zlabel('y(j)');
15
   ytr=ytr nf;
17
   yte=yte_nf;
19
  \% 1.2(b) Denote the feature matrix as X, where each row is 3-
      dimensional,
  % the first 2 dimensions are x(j, end) and x(j, end? 34), and the
      third is simply 1.
   xtr = [(xtr_nf(:,end)), (xtr_nf(:,end-34)), ones((size(xtr_nf,1)),1)];
   xte = [(xte_nf(:, end)), (xte_nf(:, end-34)), ones((size(xte_nf, 1)), 1)];
  X=xtr;
  y=ytr;
^{25}
  \% max likelihood weight (w) = inv(X'*X)*X'*y
  w = ((X'*X)^{(-1)})*(X'*y);
  \% \text{ w} = [0.46064 ; 0.52412 ; 0.00256]
  ytr_pr=xtr*w;
  yte pr=xte*w;
31 % Squared errors
```

```
ytr se=(ytr nf-ytr pr).^2;
yte se=(yte nf-yte pr).^2;
% Root Mean of squared errors
rmse tr=sqrt (mean (ytr se));
rmse te=sqrt (mean (yte se));
\% \ RMSE \ training = 0.0506 , RMSE test = 0.0503
% 1.2(c) Visualize the regression surface of this linear predictor in
    3-D using Matlab function surf().
figure
[\dim 1, \dim 2] = \operatorname{meshgrid}(0:0.01:1, 0:0.01:1);
ysurf = [[dim1(:), dim2(:)], ones(numel(dim1),1)]*w;
surf(dim1, dim2, reshape(ysurf, size(dim1)));
scatter3 (xte nf(:, end)), xte nf(:, end-34), yte nf(:, end-34)
xlabel('x(j, end)');
ylabel('x(j, end - 34)');
zlabel('y(j)');
title ('1.2(c) Visualize - Regression surface of linear predictor, the
    3-D plot of the test data points added');
```

#### Q1.3 RBF regression with adjacent pixels

(a)

The RBF model was tried with following number of basis functions 5, 10, 15, 20, 25, 30. using crossval function. The plot for the cross-validation RMSE against the number of RBFs used, is shown in Figure 6 below. Hence best model selected is for nbf = 10. The matlab code which implements this RBF model is shown below.

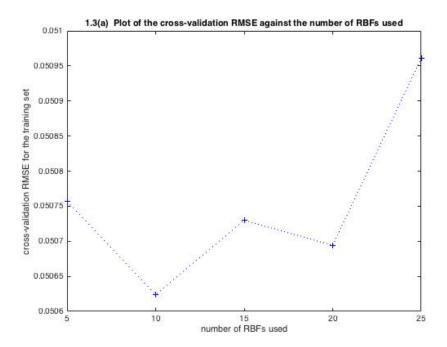
(b)

Choosing nbf = 10 (for which rmse is 0.05063), the RBF network is trained using all the non-flat training data. The values are -RMSE training = 0.0498 and RMSE test = 0.0526. Again, both the values are very close indicated a good fit for this model over given data. There is a slight improvement in RMSE compared to the previous linear regression model.

#### Matlab Code - Q1.3 - RBF

```
load imgregdata.mat xte_nf yte_nf xtr_nf ytr_nf  
 xtr_nf = [xtr_nf(:,end), xtr_nf(:,end-34)]; 
 xte_nf = [xte_nf(:,end), xte_nf(:,end-34)]; 
 options = foptions; % nbf is the number of basis functions  
options(1) = 1; % Display EM training  
options(14) = 5; % number of iterations of EM  
count=1;  
<math display="block"> rmse = [0,0,0,0,0]; 
 n rbf = [5,10,15,20,25];
```

Figure 6



```
while (count \leq 5)% nbf using the candidates \{5, 10, 15, 20, 25, 30\}.
  rbf fn=@(xtr,ytr,xte)(rbffwd(rbftrain((rbf(2, n rbf(count), 1,
      gaussian')), options, xtr, ytr), xte));
  rmse(count) = sqrt(crossval('mse',xtr nf,ytr nf,'Predfun',rbf fn));
  count = count + 1;
13
  end
  figure
15
  plot (n rbf, rmse, 'b+:');
  xlabel('number of RBFs used');
  ylabel ('cross-validation RMSE for the training set');
   title ('1.3(a) Plot of the cross-validation RMSE against the number of
      RBFs used');
  net_tr = rbf(2, 10, 1, 'gaussian'); % 1.3(b) best model nbf = 10
20
  net tr = rbftrain(net tr, options, xtr nf, ytr nf);
  ytr pr = rbffwd(net tr,xtr nf);
  ytr se=(ytr nf-ytr pr).^2;
  rmse\_tr=sqrt(mean(ytr\_se)); % rmse tr = 0.0498
  net_te = rbf(2, 10, 1, 'gaussian');
  net te = rbftrain (net te, options, xte nf, yte nf);
  yte pr = rbffwd (net te, xte nf);
  yte se=(yte nf-yte pr).^2;
```

```
rmse_te=sqrt(mean(yte_se)); % rmse te = 0.0504
```

#### Q1.4 Linear regression with all pixels

Extending the model and implementing the linear regression with all pixels, RMSE training = 0.0456 and RMSE test = 0.0371. Compared to its simpler version of linear model, it performs better on both the training and test sets, but marginally only. Given that so many extra parameters were provided for this model, the performance has not improved significantly. It indicates that the 2 previous pixels alone can predict the target pixel in a satisfactory manner. Compared to the RBF model too, if performs a little better as seen from the RMSE values, but since there is no concrete proof that it is a better model that RBF.

#### Matlab Code - Q1.4 - Linear Regression - all pixels

```
load imgregdata.mat xte_nf yte_nf xtr_nf ytr_nf
max likelihood weight(w) = inv(X'*X)*X'*y
weight = ((xtr_nf'*xtr_nf)^(-1))*(xtr_nf'*ytr_nf);
ytr_pr = xtr_nf*weight;
yte_pr = xte_nf*weight;
% Squared errors
ytr_se=(ytr_nf-ytr_pr).^2;
yte_se=(yte_nf-yte_pr).^2;
% Root mean squared errors
rmse_tr=sqrt(mean(ytr_se)); % RMSE training = 0.0456
rmse te=sqrt(mean(yte_se)); % RMSE test = 0.0371
```

## Q1.5 Neural Network with all pixels

(a)

Using the well trained MLP provided, a neural network was built with 10 hidden units. The RMSE training = 0.0333 and RMSE test = 0.0473.

Comparing its RMSEs with those from linear regression, there is a small difference only. What is noticeable is that the linear regression model seems to be less overfitting that the NN since the difference if test and training values is smaller. At the same time, the RMSE of training of NN is better than linear model which indicates makes it harder to compare both these models.

(b)

Using first 5000 data points to train the MLP and running the training 5 times using different random seeds, the resulting RMSE's are given below. The Matlab code is provided at the end of the section.

All the training and test RMSE values for the small dataset (first 5000 points) are higher that what was observed in 1.5.a. Even though different initial values (random seeds) were used for 5 different times, it did not show improvement This suggests that this subset of data taken might not be very representative of the original dataset. It is also possible that since NNs adapts its network as it learns from data, the lesser data it saw in this part is the reason for higher error values. If so,

then it is expected to perform better with larger datasets and over time.

Training RMSE	Test RMSE
0.0475	0.0499
0.0488	0.0515
0.0485	0.0515
0.0486	0.0527
0.0489	0.0526

## Matlab Code - Q1.5

```
load welltrainedMLP.mat
  load imgregdata.mat xte nf yte nf xtr nf ytr nf
3 % net struture is provided with data
_{4} ytr pr = mlpfwd(net, xtr nf);
  yte pr = mlpfwd(net, xte nf);
  rmse tr = sqrt(mean((ytr nf - ytr pr).^2)); % RMSE on training set
  rmse te = sqrt(mean((yte nf - yte pr).^2)); % RMSE on test set
  %part(b) - first 5000 data points
  arr rmse tr = [0 \ 0 \ 0 \ 0 \ 0];
  arr rmse te=[0 \ 0 \ 0 \ 0 \ 0];
   for i= [2015,2016,2017,2018,2019]
11
     rng(i, 'twister')
12
     nhid = 10; % number of hidden units
13
     net = mlp(size(xtr nf,2), nhid, 1, 'linear'); Set up the network
14
     options = zeros(1,18); % Set up vector of options for the optimiser.
     options(1) = 1; % This provides display of error values.
16
     options (9) = 1; % Check the gradient calculations.
17
     options (14) = 200; % Number of training cycles.
18
    % Train using scaled conjugate gradients.
     [net, options] = netopt(net, options, xtr nf(1:5000,:), ytr nf
20
        (1:5000,:), 'scg');
     ytr_pr = mlpfwd(net, xtr_nf);
21
     yte pr = mlpfwd(net, xte nf);
22
     tr rmse = sqrt(mean(((ytr nf - ytr pr).^2))); % RMSE on training set
23
     te_rmse = sqrt(mean(((yte_nf - yte_pr).^2))); % RMSE on test set
24
     arr rmse tr(i-2014)=tr rmse;
     arr rmse te (i-2014)=te rmse;
26
```

#### Q1.6 Discussion

Above we have considered linear regression and RBF network using 2 neighbouring pixels, and linear regression and a neural network on all pixels.

Compare these methods Especially for all pixels case, NN has the advantage that it can model non linearities in multiple pixels automatically when compared to a linear regression model which implements a statistical model . For example incase the training data has noise, NN is better in

mapping hidden and nonlinear input-output dependencies. But with the freedom of adding multiple hidden layers and nodes, the NN is also more prone to overfitting of the data that linear model.

In this particular task of pixel prediction for image compression, NN model minimizes the training root mean squared error, thus the neural networks prediction, leads to better results.

Briefly, what experiment would you try next if you wanted to improve on these predictors?

As number of pixels grow, I would like to try which of above two models handle large numerical data better. Also, once implemented, which model is better at adjusting with the changes, say for example accommodating the historical data. The next experiment I would try is changing the dataset and running the models on these new datasets since comparing the models on a single dataset can be dangerous.

Due to the complexities which increase as the network and hidden layers increase in an adaptive NN model, I would like to compare the models in terms of execution time. It might be possible that the NN model is unnecessarily slow for this task.

## Q.2. Robust modelling

#### Q2.1 Fitting the baseline model

(a)

The bias feature was added in the logistic regression model by augmenting the data and adding an extra dimension. The bias weight bw (d+1) recorded is

(b)

(b) Maximizing the likelihood

Test set Accuracy = 0.3194 Mean Log Probability = -0.7140 Std error = 0.0116 Training set Accuracy = 0.2977 Mean Log Probability = -0.7311 Std error = 0.0057 Performance of predictions on training and test

We know that a baseline that predicts P(y|x)=0.5 makes no assumptions about the data, so it will have error of 0.5 for every prediction. Mean probability, however, will predict closer to the label that it predicts as more common. So long as the label it predicts as more common IS more common, a baseline that uses mean probability will be better.

(c)

#### (c) Limited training data

Avg log probability = -0.6196 Because a small training set was used, many of the weights in the weight vector quicky minimize to either 0 or 1, indicating that in this small training set, all feature vectors with a certain value for a certain feature had the same label. In our concrete text classification example, this would indicate that for instance that only sports-labelled documents contained the word "pass." When this weight vector is used to classify a document, the linear regressor returns values of zero or one. This is an artifact of weight vectors that over-generalize the likelihood of seeing a given feature in a document of a given label, but the effect is to make the regressor entirely certain of a document's label. If the regressor returns a probability of 0 for any of the documents in the test set, then the mean log probability becomes -Inf +- NaN.

## Matlab Code - Q2.1 - Fitting the baseline model

```
load text data.mat;
2 x train = [x train ones(size(x train,1),1)]; %training and set expanded
  x \text{ test} = [x \text{ test ones}(\text{size}(x \text{ test}, 1), 1)]; \% \text{ and bias feature added}
  %weights = ones(101,1); % initialize using all 1's weight vector
   weights=rand(101,1); % initialize randomly weight vector
  %create a negative-log-likelihood function
   [NLp, dNLp dw] = log like negative(weights, x train, y train);
  % Minimize it given the training data x train and y train
   minimized weights = minimize (weights, @log like negative, 1, x train,
      y train);
   bias_weight=weights(101); %bias_weight
12
  % using fitted weights, find the probability that y=+1 for each of the
       test
 % inputs x test.
prob te = 1./(1 + \exp(-y \text{ test.*}(x \text{ test*minimized weights})));
  % the probability is compared to y test which has values (-1 \text{ or } +1), and
  \% for prediction of label, prob >=0.5, round is used to roundoff the
      prob.
   prob te rd=round(prob te);
   acc te=mean(prob te rd == y test);
   var_te=var(prob_te_rd == y_test);
   std err te = sqrt(var te/size(y test,1));
23
  % find mean log probability that predictions assign to test labels
   mean \log \text{ prob } \text{ te} = \text{mean}(\log (\text{prob te}));
  % performance of predictions on training set
27
   prob tr = 1./(1 + \exp(-y \text{ train.}*(x \text{ train}*minimized weights)));
   prob tr rd=round(prob tr);
   acc tr=mean(prob tr rd == y train);
   var tr=var(prob tr rd == y train);
   std err tr = sqrt(var tr/size(y train,1));
   mean_log_prob_tr = mean(log(prob_tr));
34
   weights ltd = rand(101,1);
36
  % Fit the model with only the first N = 100 training cases
   min weights ltd= minimize(weights ltd, @log like negative, 100, x train
      (1:100,:), y train(1:100);
  \% probability of test set being classified as +1
  prob ltd = 1./(1 + \exp(-y \operatorname{train}(1:100).*(x \operatorname{train}(1:100,:)*)
```

```
\begin{array}{c} \text{minimized\_weights)));} \\ \text{41} \quad \text{avg\_log\_prob} = \left( \frac{\text{sum}(\log(\text{prob\_ltd}))}{100} \right) \end{array}
```

#### Q2.2 Label noise model

(a)

(a) Modifying the likelihood:

Functions created which return log likelihood of this model, given data are:  $\log_l ike_n oise_w() and log_l ike_n oise_e() w.r.twar$ 

(b)

(b) Fitting a constrained parameter:

Create a function that evaluates the negative log-likelihood of the new model and evaluates the derivatives with respect to w and a. Hence fit both w and a. You are advised to wrap the function from the previous part, rather than starting from scratch. Report the fitted noise level e=?(a). Given that the test labels were checked more carefully, predict them using the newly fitted weights but using the original model (1). Report and interpret the new test accuracy and mean log probability.

#### Matlab Code - Q2.2

```
load text data.mat;
2 x train = [x train ones(size(x train,1),1)]; %training and set expanded
x_{\text{test}} = [x_{\text{test}} \text{ ones}(\text{size}(x_{\text{test}}, 1), 1)]; \% \text{ and bias feature added}
4 weights = ones(101,1); % initialize using all 1's weight vector
5 %2.2(a) modifying the likelihood
[lp1, dw] = log like noise w(weights, x train, y train, 0.1);
  grad w = checkgrad(@lr loglike noise w, weights, 1e-1, x train, y train
      , 0.1);
   [lp2, de] = log like noise e(0.1, x train, y train, weights);
   grad e = checkgrad(@lr loglike noise e, 0.1, 1e-1, x train, y train,
      weights);
10 %2.2(b) fitting a constrained param : adding a
  weights = \operatorname{rand}(102,1);
   min weights = minimize(weights, @log like noise a, 100, x train,
      y train);
  e = 1./(1 + \exp(-\min \text{ weights}(end)));
14 % mlp for noisy weights on initial model
nlp noise te = mean(log(1./(1 + exp(-x test * min weights.*y test))));
16 % probability of test set being classified as +1
ytest = x_test * min_weights;
 probability yte = 1./(1 + \exp(-y test));
```

#### Q2.3 Hierarchical model and MCMC

## (a)

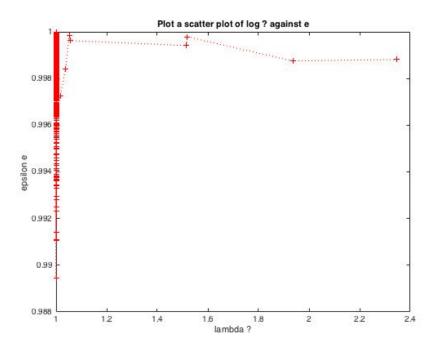
If a model were somehow able to predict every training label correctly, the binary noise variable from Eq 3 will become 0 and the log likelihood will behave as a model with no noise. For w=0, the likelihood will become  $\sup(\log(1/(1+e^0))) = \sup(\log(0.5)) = N*(-0.6)$  Nisthenumber of training/test samples.

## (b)

Putting w, e and log lambda into a single vector, and writing a wrapper to the log posterior function to evaluate Eq(6), slice sample fn was called. Step out values was toggled between false and true, and a number of combinations of width, burn and N were tried. For initialization of the row vector with e,lambda,w, rand fn was called to set the values.

The function returned N samples of e,lambda,w. Plot of lambda against e is shown below. Since the step-out was set to false, the function did not performed as expected.

Figure 7



## (c)

The slice sampler was hung indefinitely with step out = true. All the samples returned with stepout = false are valid, even though the distribution is not as expected. These observations can be utilized as part of test sets of the same model. The values observed are epsilon values are high - between 0.9 and 1.0 and the lambda values are between 1 and 3. Since the function did not complete the execution as expected, the comparison is not possible.

#### Matlab Code - Q2.3 - Hierarchial Model and MCMC

```
load text data.mat;
 x train = [x train ones(size(x train,1),1)]; %training and set expanded
 x \text{ test} = [x \text{ test ones}(size(x \text{ test},1),1)]; \% \text{ and bias feature added}
  %weights = ones(101,1); % initialize using all 1's weight vector
  weights=rand(101,1); % initialize randomly weight vector
  dim=size (weights, 1);
  log lambda=0.1; % initilize with random value btw 0 and 1
  e=0.5; % initilize with random value btw 0 and 1
  % handle of log_posterior passed through slice_sample
 log_p=log_posterior(e,log_lambda,weights,dim,x_train,y_train);
11 % set the values for slice sample
N=10; % no of iterations
  burn=0; % default value
  widths=1; % default value
  step out=false; % value set to true hangs the code.
  rng(0, 'twister');
  init=init params(dim); % row vector initialized
 \log p \ s = @(args) \log posterior(args\{:\}, dim, x train, y train);
  % wrapper created so as to hide extra parameters
 \log s = @(vector) \log p s(split params(vector));
21 % make sure that the initial row vector is inside the prob density area
  assert(~isinf(log s s(init))); % before passing it to slice sample
  % MCMC slice sampling for heirarchial posterior
  result = slice sample(N, burn, log s s, init, widths, step out);
  \% the result contains 3 params – e, lambda and w
  params cells=split params (result);
  e=params cells(1);
  log lambda=params cells(2);
  weights=params cells(3);
  res lambda = exp(log lambda \{1,1\});
  res e=e\{1,1\};
  res weights=weights {1,1};
33 % scatter plot to confirm posterior belie about log lambda and e
plot (res lambda, res e, 'r:+');
xlabel('lambda?');
 ylabel ('epsilon e';');
  title({'Plot a scatter plot of log ? against e'});
```

## Appendix A - Additional Code

```
function log post = log posterior(e, log lambda, ww, D, xx, yy)
   if (e \le 0 \mid \mid e > 1) \% zero prior density
       \log post = -Inf;
       return;
5
   end
6
   if (log lambda <= 0 | log lambda > 1)
       \log post = -Inf;
       return;
9
   end
10
   sigmas = 1./(1 + exp(-yy.*(xx*ww)));
  lambda=exp(log lambda);
  \% Dropped one constant term = -\log(pi)
   \log \text{prior} = -\text{lambda.}*((\text{ww'})*\text{ww}) + (D/2)*\log_{\text{lambda}};
   \log \text{like} = \text{sum}(\log(((1-e)*\text{sigmas})+(e/2)));
   log post = log like + log prior;
17
   end
  function [params] = split params(vector)
  % slice sample returns a matrix, this fn
  % extracts the 3 sampled parameters
  [rows, cols] = size(vector);
  %initial case while starting slice sample
   if rows = 1
       e=vector(1);
       log lambda=vector(2);
       weights=vector (3:end);
       params={e, log lambda, weights'};
10
       return
11
   else
12
       e=vector(1,:);
13
       log lambda=vector (2,:);
14
       weights=vector(3:end,:);
15
       params={e,log lambda, weights}; % the returned result of
16
           slice sample
  end
  function params = init params (dim)
  % initializes the row vector to be passed to slice sample()
   params=ones(1, dim+2);
   params(1)=rand(); % 1st param is epsilon (e) - range btw 0 and 1
  params(2)=rand(); % 2nd param is log(lambda) - range btw 0 and 1
  b=9.376307381550576e+04; % range of weight vector - random value
  a = -7.460274498610420e + 05;
  for i = 3 : dim+2 % 3rd element till last element is weight vector
```

```
params(i)=(b-a).*rand() + a;
10 end
function [NLp, dNLp dw] = log like negative (ww, xx, yy)
2 % this fn returns the negative log likelihood
yy = (yy==1)*2 - 1;
sigmas = 1./(1 + exp(-yy.*(xx*ww)));
_{5} NLp = -sum(log(sigmas));
6 if nargout > 1% additionally returns the derivate w
      dNLp \ dw = (((-1)*(1-sigmas).*yy)' * xx)';
  end
function [Lp, dLp dw] = log like noise w(ww, xx, yy, e)
2 % this fn returns the log likelihood for a model with noise e
3 % and the gradient w
_{4} yy = (yy==1)*2 - 1;
sigmas = 1./(1 + exp(-yy.*(xx*ww))); \% Nx1
6 Lp = sum(log((1-e)*sigmas+(e/2)));
7 if nargout > 1
      dLp dw = (((((1-e) \cdot /((1-e) * sigmas + e/2)) \cdot * sigmas \cdot *(1-sigmas)) \cdot * yy))
          * xx) ';
  end
function [Lp, dLp de] = log like noise e(e, xx, yy, ww)
_{2} vy = (yy==1)*2 - 1;
  % this fn returns the log likelihood for a model with noise e
4 % and the gradient e
sigmas = 1./(1 + exp(-yy.*(xx*ww))); \% Nx1
<sub>6</sub> Lp = sum(log((1-e)*sigmas+(e/2)));
  if nargout > 1
       dLp de = sum((-1.*sigmas + (1/2)).* (1./((1-e)*sigmas+(e/2))));
  end
1 function [NLp, dLp dw, dLp da] = log like noise a(ww, xx, yy)
yy = (yy==1)*2 - 1;
  % this fn returns the -ve log likelihood for a model with noise
4 % unconstrained parameter a and returns derivative wrt w and a
a = ww(end,:);
6 \text{ ww} = \text{ww}(1: \text{end} - 1,:);
  e = 1/(1 + \exp(-a));
 sigmas = 1./(1 + exp(-yy.*(xx*ww)));
  [Lp, dLp dw] = log like noise w(ww, xx, yy, e);
10 NLp=-Lp;
^{11} dLp dw=dLp dw*a;
_{12} dLp da= sum(((log((1-log(a))*sigmas+(log(a)/2))).^(-1))*(((1/a)*sigmas)
      +(0.5*a));
```

# Appendix B - References

1. http://uk.mathworks.com/help/

2. http://mlg.eng.cam.ac.uk/zoubin/tut06/mcmc.pdf

3..http://homepages.inf.ed.ac.uk/imurray2/teaching