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EE 5322 Intelligent Control Systems Assignment no 4

Neural Networks

Ouestion 1:

1- Consider the following training data set

```
x = [5 4 6 5 2 1 2 1
     0 -1 0 -1 -1 -2 -2 -3]; % inputs
y = [0 0 0 0 1 1 1 1 ]; % Target values
```

a- Train a perceptron to classify the data set into two classes. Plot points and decision boundaries.

b- Use a single neuron with sigmoid activation function to do this classification problem. Plot the points and decision boundary and compare the results to part a.

Solution 1 a:

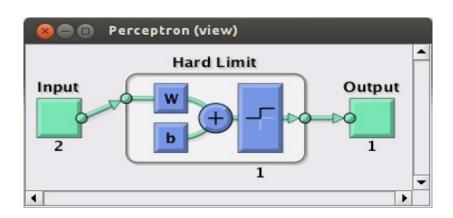
```
Code:
```

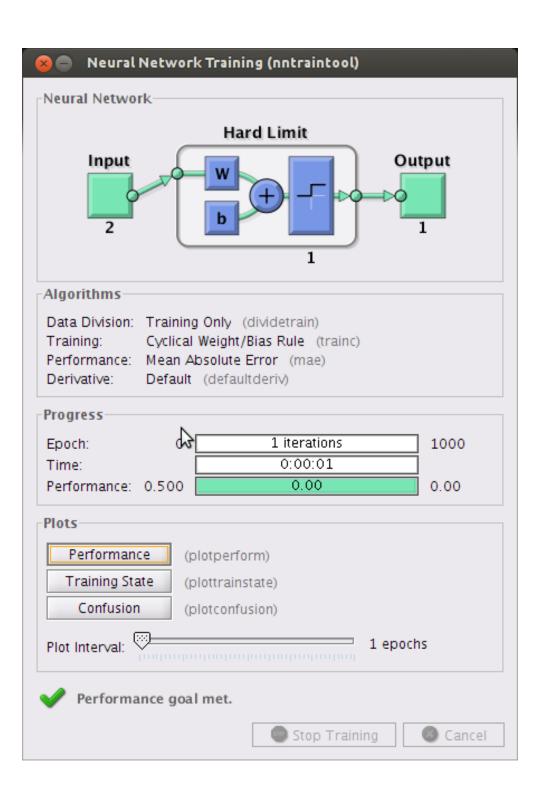
```
close all,
clear all,
clc,
format compact
% number of samples of each class
% define inputs and outputs
x = [5 4 6 5 2 1 2 1]
    0 -1 0 -1 -1 -2 -2 -3]; % inputs
y = [0 \ 0 \ 0 \ 0 \ 1 \ 1 \ 1 \ 1];
% outputs
% Plot input samples with PLOTPV (Plot perceptron input/target vectors)
figure(1)
plotpv(x,y);
net = perceptron;
net = train(net,x,y);
view(net);
figure(1)
plotpc(net.IW{1},net.b{1});
y = net(x)
```

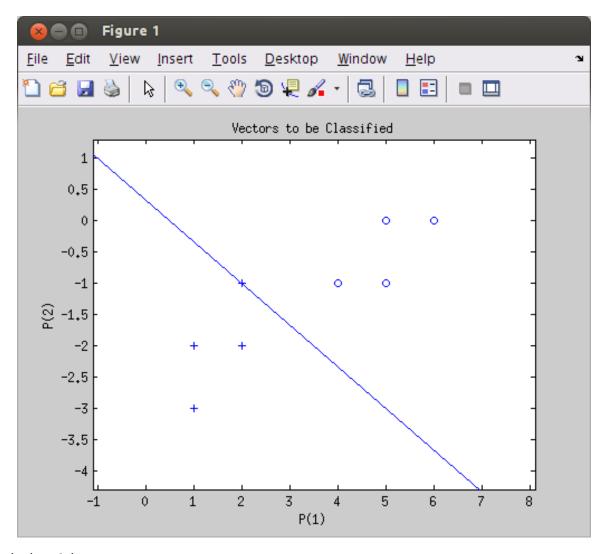
OutPut :

y = 0 0 0 0 1 1 1 1

Plots:





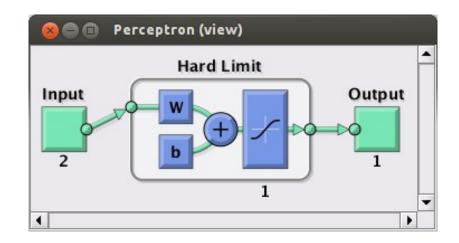


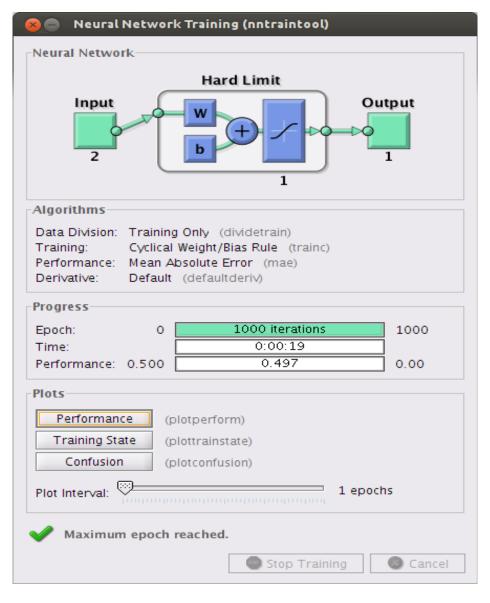
Solution 1 b:

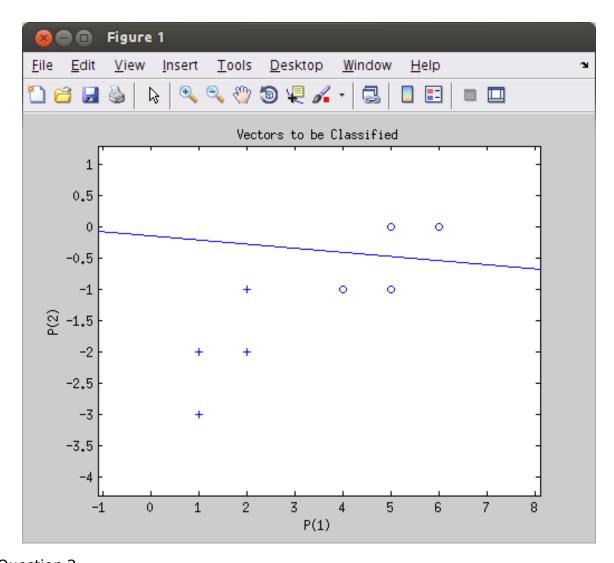
```
Code:
close all,
clear all,
clc,
format compact
% number of samples of each class
% define inputs and outputs
x = [5 \ 4 \ 6 \ 5 \ 2 \ 1 \ 2 \ 1]
    0 -1 0 -1 -1 -2 -2 -3]; % inputs
y = [0 \ 0 \ 0 \ 0 \ 1 \ 1 \ 1 \ 1];
% outputs
% Plot input samples with PLOTPV (Plot perceptron input/target vectors)
figure(1)
plotpv(x,y);
net = perceptron;
net.layers{1}.transferFcn = 'tansig';
net = train(net,x,y);
view(net);
figure(1)
plotpc(net.IW{1},net.b{1});
y = net(x)
```

Output :y =
Columns 1 through 6
-0.9889 0.9969 -0.9946 0.9936 0.9993 1.0000
Columns 7 through 8
1.0000 1.0000

Plots:



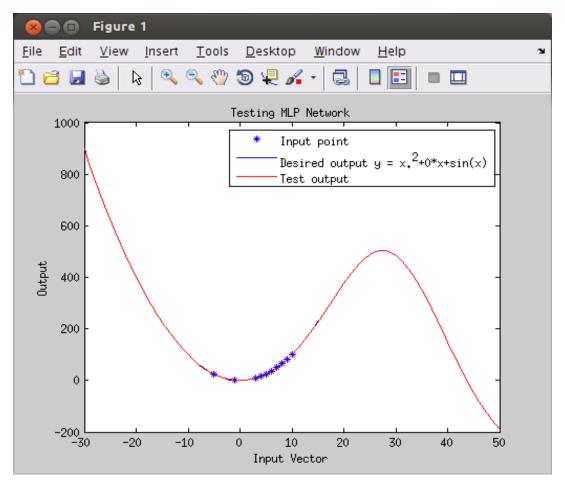


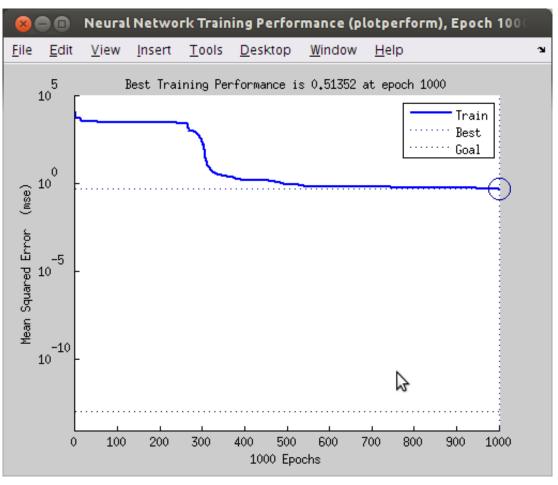


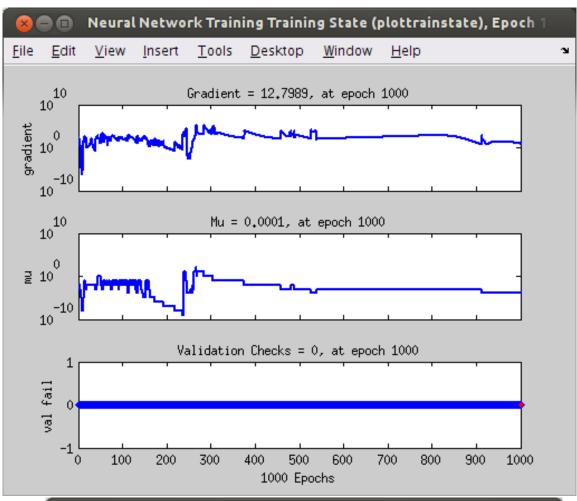
Question 2:
Consider the dynamical system

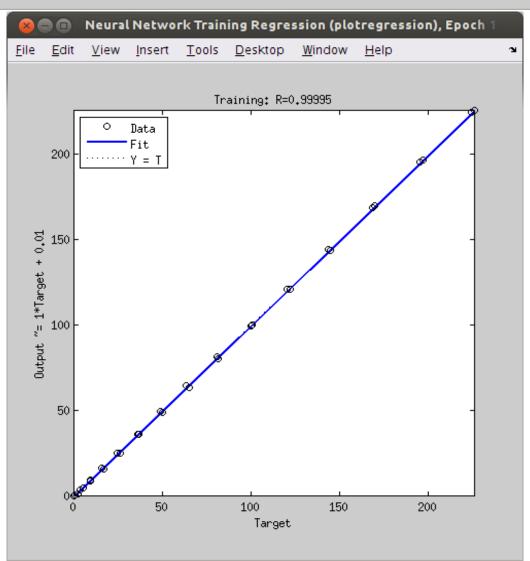
a) Approximate the function f(x) using an MLP neural network and plot the function and the estimation on the same graph.

b) Simulate the system response for exact f(x) and the approximation. Use different initial conditions. Compare the results.









```
Output:
ans = Neural Network
        name: 'Custom Neural Network'
    efficiency: .cacheDelayedInputs, .flattenTime,
            .memoryReduction
      userdata: (your custom info)
  dimensions:
     numInputs: 1
     numLayers: 4
     numOutputs: 1
  numInputDelays: 0
  numLayerDelays: 0
numFeedbackDelays: 0
numWeightElements: 76
    sampleTime: 1
  connections:
    biasConnect: [1: 1: 1: 1]
   inputConnect: [1; 0; 0; 0]
   layerConnect: [4x4 boolean]
   outputConnect: [0 0 0 1]
  subobjects:
       inputs: {1x1 cell array of 1 input}
       layers: {4x1 cell array of 4 layers}
      outputs: {1x4 cell array of 1 output}
       biases: {4x1 cell array of 4 biases}
   inputWeights: {4x1 cell array of 1 weight}
   layerWeights: {4x4 cell array of 3 weights}
```

Use a multilayer perceptron to classify the data set into two classes and Compute the false-positive rate and false-negative rate of your classification results.

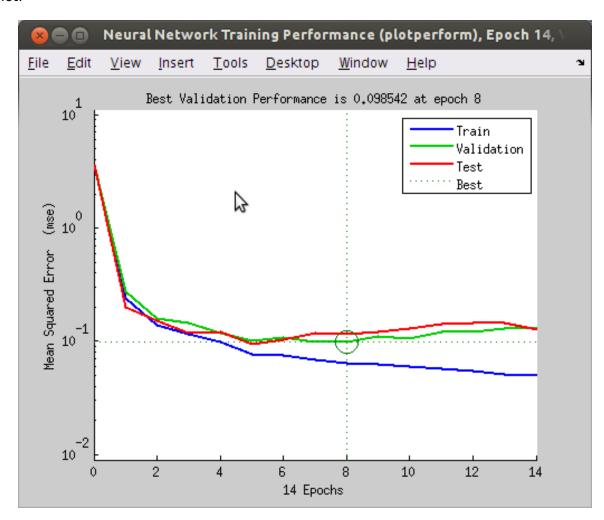
Code:

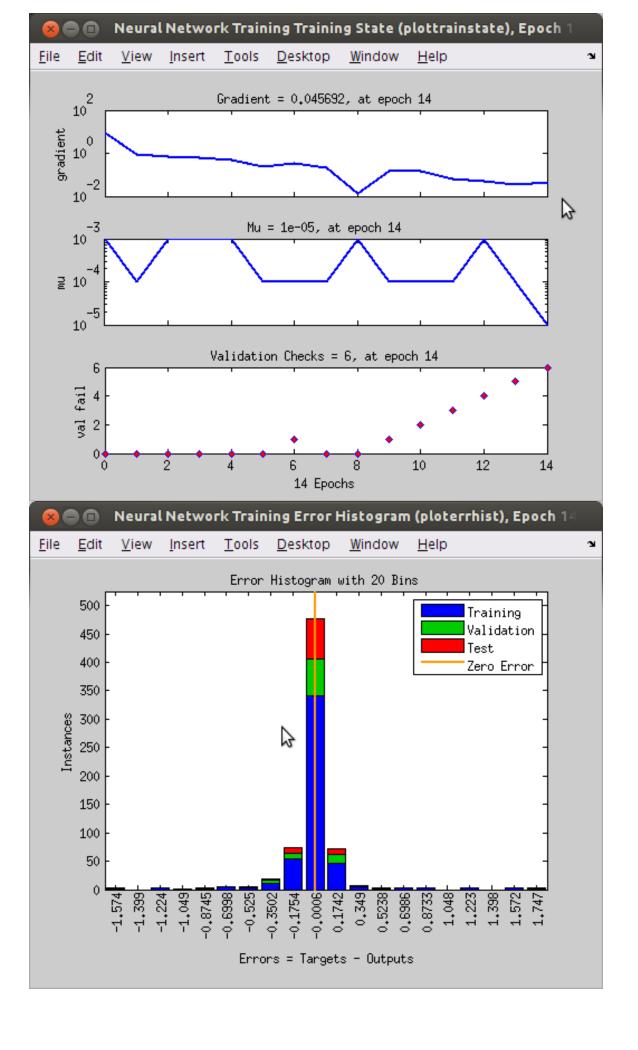
Question 3:

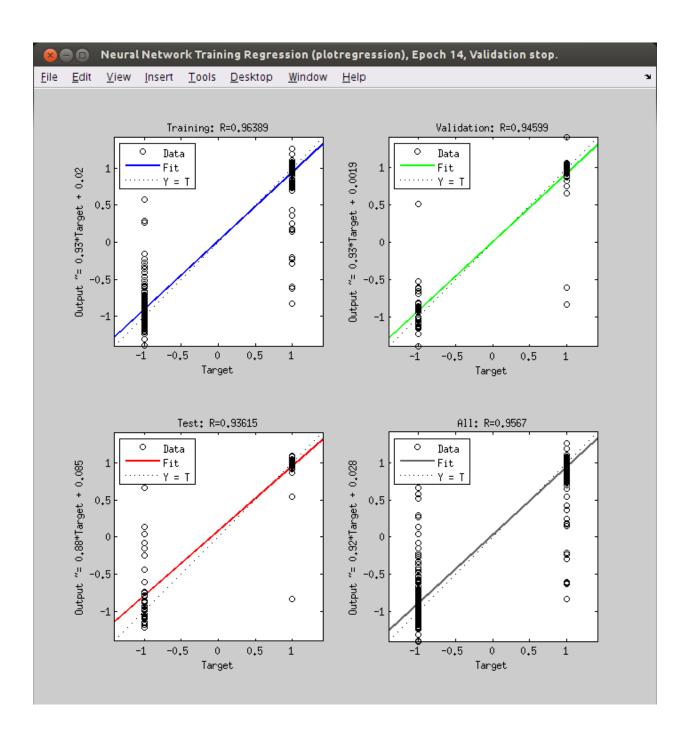
```
clc:
clear all;
close all;
%data classification using MLP
data=open('breastcancer.mat');
Input=data.data(:,2:10)'; %Input matrix
Target=data.data(:,11)'; %Target values
benign=0;
mali=0;
for i=1:length(Target)
    if Target(i)==2
        Target(i)=1; %Benign cell
        benign=benign+1;
    else
        Target(i)=-1; %Malignant cell
        malignant=malignant+1;
    end
end
nnet=feedforwardnet(10);
nnet.divideParam.trainRatio = 70/100;
nnet.divideParam.valRatio = 15/100;
nnet.divideParam.testRatio = 15/100;
```

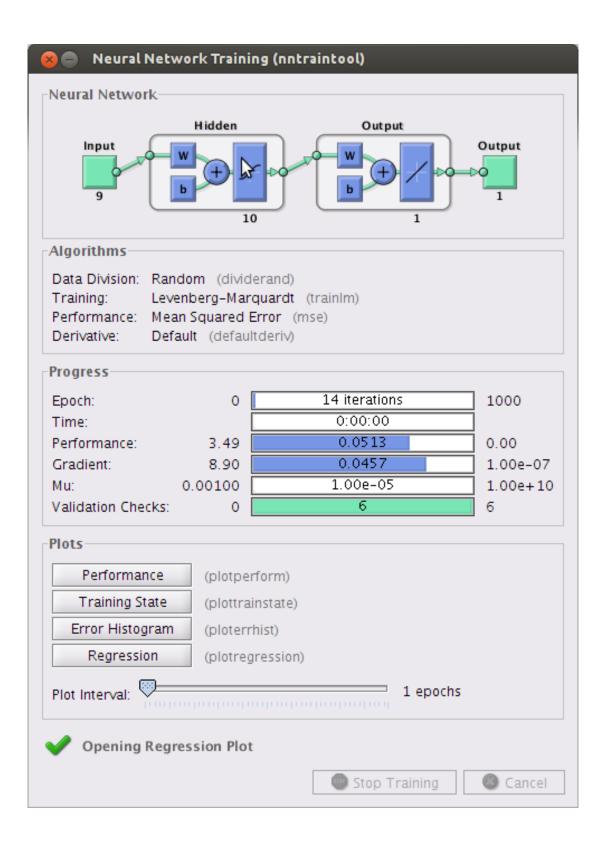
```
nnet=train(nnet,Input,Target);
output=hardlims(sim(nnet,Input));
wrongMalign=0;
wrongBenign=0;
for i=1:length(Target)
    if Target(i)==1 && output(i)==-1
        wrongMalign=wrongMalign+1;
    end
    if Target(i)==-1 && output(i)==1
        wrongBenign=wrongBenign+1;
    end
end
FalsePositive=wrongMalign/benign
FalseNegative=wrongBenign/malignant
Output:
FalsePositive =
  0.0203
FalseNegative =
  0.0293
```

Plot:









```
Solution 3b:
Code:
clc;
clear all;
close all;
%Classification using Self-Organizing Map
data=open('breastcancer.mat');
P=data.data(:,2:10)'; %Input matrix
T=data.data(:,11)'; %Target values
benign=0;
malignant=0;
for i=1:length(T)
    if T(i)==2
        %T(i)=1; %Benign cell
        benign=benign+1;
    else
        %T(i)=0; %Malignant cell
        malignant=malignant+1;
    end
end
SOM=selforgmap([8 8],478);
SOM=train(SOM,P);
output=SOM(P);
classes=vec2ind(output);
```

Output:

