Ayin CMNDLINE

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%GENERATES A SIMPLE DATASET
N = 150;
dataset1 = rand(300,N)*5;
dataset1(:,N+1) = 1;
dataset2 = rand(300,N)*5 + 5;
dataset2(:,N+1) = 2;
dataset4 = rand(300,N)*5 + 10;
dataset4(:,N+1) = 4;
observations = [dataset1;dataset2;dataset4];
num_items = size(observations,1);
scramble = randperm(num_items);
observations = observations(scramble,:); %permutes the dataset
%ADJUSTMENTS FOR UCI DATASETS
%Adjustments for the Wine database to: (1) adjust the scale of the last column (it's
too large for the rest of the data); and
%(2) swap the first and last columns after that, to put the labels in the last
column.
clear temp
temp = observations(:,1);
observations(:,14) = observations(:,14)/100;
observations(:,1) = [];
observations(:,14) = temp;
%Adjustments to the Parkinson's database to move the labels to the last column
%and delete non-data entries from the first row and first column
observations(1,:) = [ ];
observations(:,1) = [ ];
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clear temp
temp = observations(:,17);
observations(:, 17) = [ ];
observations(:,23) = temp;
%LOADS CSV DATASET
observations = [ ];
filename = "C:/Users/erdosfan/Desktop/Image Research/External Training
Sets/ionosphere-data.txt";
Note that for the ionosphere dataset, you'll need to do a manual replacement of the
g's and b's with some kind of numerical category label.
%This can be done easily in a text editor using find / replace.
%loads the file
observations = csvread(filename);
%RUN THE ALGORITHM
threshold = .93; %threshold accuracy above which learning stops
N = size(observations,2) - 1; %dimensions of dataset
tic;
[output_matrix dataset accuracy_vector] = Ayin_RTE(observations, threshold, N);
toc
%measures accuracy
num_predictions = size(output_matrix,1);
num_learning_iterations = size(dataset,1) %number of training iterations
learning_ratio = num_learning_iterations/num_predictions %percentage of training
iterations to total iterations
overall_accuracy = sum(output_matrix(:,N+1) == observations(:,N+1))/num_predictions
start_val = 1;
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Ayin_CMNDLINE

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end_val = num_predictions;
figure, plot(start_val:end_val,accuracy_vector(start_val:end_val))
```