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NEUROGENESIS RBM
% Execute this file second.
% NOTE:
% epsilon (learning rate) = 0.3
% sparsity cost = 0
% weight decay = 0.005
trialNum = 10;
visible_data = zeros(trialNum,10);
hidden_data = zeros(trialNum,50);
visible recon = zeros(trialNum,10);
hidden_recon = zeros(trialNum,50);
delta_weight = zeros(10,50);
weight = zeros(10,50);
Pmat_h = zeros(1,50);
Pmat v = zeros(1,10);
Pmat_h_r = zeros(1,50);
test_visible_data = zeros(trialNum,10);
test_hidden_data = zeros(trialNum,50);
test_visible_recon = zeros(trialNum,10);
test_hidden_recon = zeros(trialNum,50);
p = 0.05; % target activation
q = zeros(1,trialNum);
weightdecay = 0.005;
% INITIALIZE PATTERN INPUT % WEIGHTS ************************
% length = 10;
% percent = 0.1;
% [orig_data v1 v2 v3 v4 v5 v6 v7 v8 v9 v10] = patternGen(length, percent);
% visible_data = [v1; v2; v3; v4; v5; v6; v7; v8; v9; v10];
%!!! Computations if NOT using normal distribution.
% for i = 1:size(weight,1)
% for j = 1:size(weight,2)
      weight(i,j) = rand(1);
   end
% end
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% weight;
% !!! Computations if randomizing weight matrix using a normal
% distribution.
sigma = 0.0001;
mu = 0.0;
weight = normrnd(mu, sigma, 10, 50); % Random normal distribution fxn.
%weight(weight<0) = 0; % Make negatives into zeros.</pre>
weight;
% COMPUTE LEARNING RATE 1-q(t) AND SPARSITY COST(q(t)) *****************
%!!! This code should only be used when factoring in sigmoidal aging.
epsilon = zeros(1,trialNum); % Learning rate
gt_vect = zeros(1,trialNum); % Sparsity cost
dt = 1/trialNum;
for i = 1:trialNum
    gt_vect(i) = gompertz(-1+(dt*i)); %add to the min. which is -1
    epsilon(i) = 1-gt_vect(i);
end
gt vect;
epsilon;
gt_normalized = normalize_var(gt_vect, 0.1, 0.3);
epsilon normalized = normalize var(epsilon, 0, 0.9);
%!!! Make epsilon and cost constants for trials when not accounting for
% sigmoidal aging.
    % Learning rate 1-q(t) = 0.1 for non neurogen; 0.3 for neurogen.
    % Sparsity cost = 0.9 for non neurogen; 0 for neurogen.
% COMPUTE VISIBLE DATA, HIDDEN DATA, AND THE RECONS ******************
for i = 1:size(visible data, 1) %i is trial number
    % Getting values of Pmat h:
    Pmat_h = 1./(1+exp(-visible_data(i,:) * weight));
    % Set hidden units to 1 depending on the values of Pmat h:
    for j = 1:50;
        if rand(1) < Pmat h(j);
            hidden_data(i,j) = 1;
        end
    end
    q(i) = mean(hidden_data(i,:));
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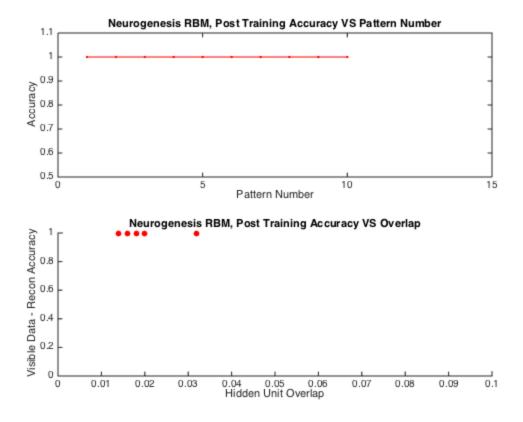
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% Getting values of Pmat v:
    Pmat_v = 1./(1+exp(-hidden_data(i,:) * weight'));
    % Set visible recon units to 1 depending on the values of Pmat_v:
    for j = 1:10;
        if rand(1) < Pmat_v(j);</pre>
            visible_recon(i,j) = 1;
        end
    end
    % Getting values of Pmat h r:
   Pmat_h_r = 1./(1+exp(-visible_recon(i,:) * weight));
    % Set hidden recon units to 1 depending on the vales of Pmat_H_r:
    for j = 1:50
        if rand(1) < Pmat h r(j);</pre>
            hidden_recon(i,j) = 1;
        end
    end
    %!!!!!! Since we are assuming neurons are all new, replaced epsilon and
    %sparsity cost as a constant across all trials:
    for k = 1:size(visible data,2) %1:10
        for m = 1:size(hidden_data,2) %1:50
            % 0.3 replaced epsilon(i)
            delta_weight(k,m) = 0.3*((visible_data(i,k)*hidden_data(i,m))...
                - (visible_recon(i,k)*hidden_recon(i,m))) - ...
                (weightdecay * weight(k,m)) ...
                - (0 * (q(i)-p));% (0 replaced gt_normalized(i));
        end
    end
   weight = weight + delta_weight; % Update weight
end
% COMPUTE VISIBLE DATA, HIDDEN DATA, AND THE RECONS *******************
for i = 1:size(visible_data, 1) %i is trial number
    % Getting values of Pmat h:
   Pmat_h = 1./(1+exp(-visible_data(i,:) * weight));
    % Set hidden units to 1 depending on the values of Pmat_h:
    for j = 1:50;
        if rand(1) < Pmat_h(j);</pre>
            hidden data(i,j) = 1;
        end
    end
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q(i) = mean(hidden data(i,:));
    % Getting values of Pmat_v:
    Pmat_v = 1./(1+exp(-hidden_data(i,:) * weight'));
    % Set visible recon units to 1 depending on the values of Pmat_v:
    for j = 1:10;
        if rand(1) < Pmat_v(j);p</pre>
            visible_recon(i,j) = 1;
        end
    end
    % Getting values of Pmat_h_r:
    Pmat_h_r = 1./(1+exp(-visible_recon(i,:) * weight));
    % Set hidden recon units to 1 depending on the vales of Pmat_H_r:
    for j = 1:50
        if rand(1) < Pmat_h_r(j);</pre>
            hidden_recon(i,j) = 1;
        end
    end
    %!!!!!! Since we are assuming neurons are all new, replaced epsilon and
    %sparsity cost as a constant across all trials:
    for k = 1:size(visible_data,2) %1:10
        for m = 1:size(hidden data,2) %1:50
            % 0.1 replaced epsilon(i)
            delta_weight(k,m) = 0.1*((visible_data(i,k)*hidden_data(i,m))...
                - (visible recon(i,k)*hidden recon(i,m))) - ...
                (weightdecay * weight(k,m)) ...
                - (0.9 * (q(i)-p));% (0.9 replaced gt_normalized(i));
        end
    end
   weight = weight + delta_weight; % Update weight
% COMPUTE VISIBLE DATA, HIDDEN DATA, AND THE RECONS FOR POST TRAINING *****
for i = 1:size(visible_data, 1) %i is trial number
    % Getting values of Pmat_h:
    Pmat_h = 1./(1+exp(-test_visible_data(i,:) * weight));
    % Set hidden units to 1 depending on the values of Pmat h:
    for j = 1:50;
        if rand(1) < Pmat_h(j);</pre>
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end

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test_hidden_data(i,j) = 1;
       end
   end
   q(i) = mean(test_hidden_data(i,:));
   % Getting values of Pmat v:
   Pmat_v = 1./(1+exp(-test_hidden_data(i,:) * weight'));
    % Set visible recon units to 1 depending on the values of Pmat_v:
   for j = 1:10;
       if rand(1) < Pmat_v(j);p</pre>
           test_visible_recon(i,j) = 1;
       end
   end
    % Getting values of Pmat h r:
   Pmat_h_r = 1./(1+exp(-test_visible_recon(i,:) * weight));
   % Set hidden recon units to 1 depending on the vales of Pmat_H_r:
   for j = 1:50
       if rand(1) < Pmat_h_r(j);</pre>
           test_hidden_recon(i,j) = 1;
       end
    end
end
% VDATA - VRECON / ACCURACY FOR POST TRAINING ********************
vDiff_PT_neurogen = zeros(1, trialNum);
for i = 1:trialNum
       vDiff_PT_neurogen(i) = 1-(abs(sum(test_visible_data(i,:) - test_visible_re
end
% VDATA - VRECON / ACCURACY FOR DURING TRAINING ********************
vDiff_DT_neurogen = zeros(1, trialNum);
for i = 1:trialNum
   vDiff_DT_neurogen(i) = 1-(abs(sum(test_visible_data(i,:) - test_visible_recon(
end
hiddenOverlap = zeros(10, trialNum-1);
sizeV = size(visible_data,1) * size(visible_data,2);
for i = 1:trialNum
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for j = 1:trialNum
       if (j ~= i)
          hiddenOverlap(i,j) = abs(sum(test_hidden_data(i,:)) - sum(test_hidden_
   end
end
hiddenOverlapsum_neurogen = zeros(1,trialNum);
for i = 1:trialNum
   hiddenOverlapsum_neurogen(i) = sum(hiddenOverlap(i,:))/trialNum;
end
hiddenOverlapsum_neurogen;
figure(2)
subplot(2,1,1)
plot(1:10, vDiff_DT_neurogen, '.-r')
axis([0 15 0.5 1.1])
title('Neurogenesis RBM, Post Training Accuracy VS Pattern Number')
xlabel('Pattern Number')
ylabel('Accuracy')
subplot(2,1,2)
scatter(hiddenOverlapsum_neurogen, vDiff_PT_neurogen, 'filled', 'r')
axis([0 0.1 0 1]);
title('Neurogenesis RBM, Post Training Accuracy VS Overlap')
xlabel('Hidden Unit Overlap')
ylabel('Visible Data - Recon Accuracy')
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