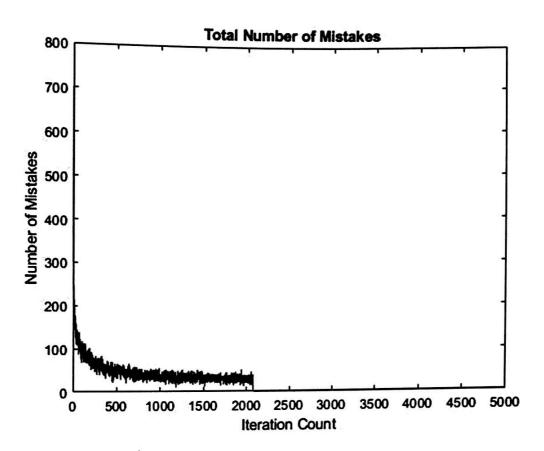
EEE - 485

Statistical Learning and Data Analytics

Problem Set #3

Kerem Ayöz 21501569



Number of mistakes become 0 after ~ 2100 iterations, since perceptron finds the separating hyperplane.

C.) Yes, doto is linearly separable since poceptron gives O error after contain interations.

$$y = 57 \times 1 + 29.221.x_1 + 231.6887.x_3 - 201.7355 \times 4 - 115.8198 \times_5$$

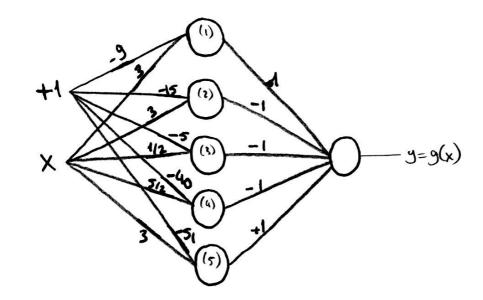
-174.0012 x6 -57.7855 x7 +231.6201 x8 +87.1270 x9 - 115.821 x10

Yes, those might be altiferent separating hyperplanes which could be found with different learning rate and weight initializations.

d.) We can use a method similar to SVM (Support vector machines). We can find the distances of closest points in each closses to each separating hypotheme. We can select the hypotheme with the equal distance to closest points in each class.

QUESTION 1 Code

```
load("perceptrontest.mat");
x = D(:,1:11);
y = D(:,12);
[weights, mistakes] = train_perceptron(x,y,1,5000);
error = calculate_error(x, weights, y);
plot(mistakes);
title("Total Number of Mistakes");
ylabel("Number of Mistakes");
xlabel("Iteration Count");
function error = calculate_error(x,w,y)
    error = 0;
    for i = (1:length(x))
        est = perceptron(x(i,:),w);
        error = error + (y(i,:) - est);
    end
end
function [weig, mistakes] =
train_perceptron(x,y,learning_rate,iteration_count)
    w = zeros(1,11);
    mistakes = zeros(1,5000);
    for i = (1:iteration_count)
       [w, mistakes(i)] = single_iteration(x,y,learning_rate,w);
    weig = w;
end
function [weigh, mistake] = single_iteration(x, y, learning_rate, w)
    mistake = 0;
    for i = (1:length(x))
        est = perceptron(x(i,:),w);
        w = w + learning_rate * (y(i,:) - est) * x(i,:);
        mistake = mistake + abs(y(i,:) - est);
    end
    weigh = w;
end
function estimate = perceptron(x_i,w)
   res = x i*transpose(w);
    if res > 0
        estimate = 1;
    else
        estimate = 0;
   end
end
```



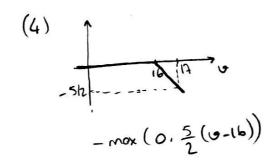
(1)

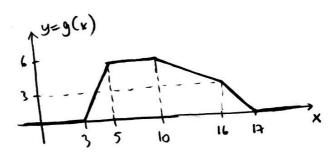
max (0,3(v-3))

- max (0,3(10-5))

(3)

- max(0,1(0-10))





- Output of neural network-

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a.) Because we did not standardize the samples. Third feature for each element in sample is much larger than other features. So not-standardizing the data causes one eight value to be much bigger than others.

$$u_{1} = \begin{cases} 0.8278 \\ 0.5605 \\ -0.0227 \end{cases} u_{2} = \begin{cases} 0.5608 \\ -0.8279 \\ 0.015 \end{cases} u_{3} = \begin{cases} 0.0128 \\ 0.0214 \\ 0.9397 \end{cases}$$

$$e_{1} = 0.2 \qquad e_{2} = 1.9 \qquad e_{3} = 1400.9$$

b.)
$$u_1' = \begin{pmatrix} 0.5245 \\ 0.5650 \\ -0.6370 \end{pmatrix}$$
 $u_2' = \begin{pmatrix} 0.3725 \\ -0.6303 \\ 0.0770 \end{pmatrix}$ $u_3' = \begin{pmatrix} 0.3580 \\ 0.7670 \\ 0.7670 \end{pmatrix}$

$$e_1' = \begin{pmatrix} 0.0742 \\ e_2' = 1.0276 \end{pmatrix}$$

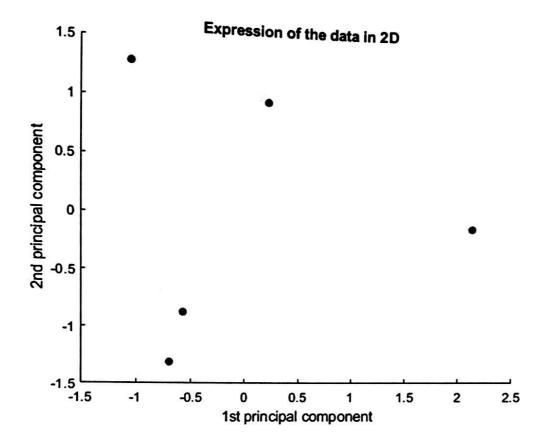
$$e_3' = 1.2981$$

$$x' = \begin{pmatrix} -0.6972 \\ -0.6972 \\ 1.1268 \\ 0.1937 \\ 0.937 \\ 0.9304 \\ -1.0538 \\ 1.12370 \\ -0.5755 \\ -0.8646 \end{pmatrix}$$

$$x_1' = \begin{pmatrix} 0.3580 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.7670 \\ 0.76$$

Yes it is a good representation since it really explains all of the variace in data. (1.96.9)

C.) It means the non-zero dimensions are enough to represent all the varionce in obta. Zero eigen-value alimension also not have any variance, it is some for all the instances. We can remove that feat-ne and represent the obta with other atimansions.



QUESTION 3 Code

```
x1 = [1; 4; 400];
x2 = [3; 5; 500];
x3 = [3; 2; 450];
x4 = [3; 1; 400];
x5 = [1; 3; 425];
x = [x1, x2, x3, x4, x5];
x_centered = zeros(size(x));
x_standardized = zeros(size(x));
% Centralize
for i = (1:3)
    x_{centered(i,:)} = x(i,:) - mean(x(i,:));
cov = x_centered * transpose(x_centered)./5;
[v1, e1] = eig(cov);
% Standardize
for i = (1:3)
    x_standardized(i,:) = x_centered(i,:) / std(x_centered(i,:));
cov = x_standardized * transpose(x_standardized)./5;
[v2, e2] = eig(cov);
x_standardized = transpose(x_standardized);
z\overline{1} = x_{standardized} * v2(:,3);
z2 = x_standardized * v2(:,2);
figure;
scatter(z1,z2, 'filled');
title("Expression of the data in 2D");
ylabel("2nd Principal Component");
xlabel("1st Principal Component");
totalV = sum(sum(x_standardized.*x_standardized))/5
pvm1 = sum(z1.^2) / 5
pvm2 = sum(z2.^2) / 5;
ratio = (pvm1+pvm2)/totalV*100
```

a.) Let's consider $X=A\cdot s$. If we multiply right side by $\vec{PP}=T$ where P is a permetation matrix, we get $X=AP^{-1}Ps$. If we coll AP^{-1} as \vec{A} then $X=\vec{A}Ps$ which indicates that result may be a different permetation than the original s.

For sign antigrity if we both altiply A and s with negotive one we get $X = As = -A \cdot -s = \ddot{A}\ddot{s}$ where \ddot{s} has apposite sign of s, which means recovered signal night have apposite sign than the argumal.

b.) Let's take $S \sim W(0,1)$ which is standart normalizationable. X = Acs is the equation for recovering signal. Then x is also gaussian random variable and its variate C can be additionally from covariance matrix. E(xxT) = E(AsSTAT) = E(AAT) = AAT $\sum x \sim W(0, AAT)$ E(x) = E(As) = 0

Let's define A' = AK where K is orthogonal which means $KK^T = I_P$ If we again apply recovery formulo with A' we get $X' = A' \cdot s$.

Calculate $E[X'X'^T]$ and E[X'];

E(x'x'T) = E(A'SSTA'T) = E(A'A'T) = E(AKKTAT) = E(AAT) = AAT $E(x') = E(A'S) = 0 Then > x' \sim W(O, AAT)$

This means that with some distributed ru's x and x', we cannot decide mixing matrix since with 2 altiferent mixing matrix we reach the same gaussian distribution for dosonators. This, sources need to be non-Gaussian