

# EEE 178 Test 2

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```
% pre-processing
clear all, close all, clc;
I1=imread('exam2letters.jpg');
I2=I1;
I1=rgb2gray(I1);
figure; imshow(I1);
```

The intrinsic matrix containing 5 intrinsic parameters. These parameters encompass focal length, image format, and principal point. The parameters and represent focal length in terms of pixels, where and are the scale factors relating pixels to distance. represents the skew coefficient between the x and the y axis, and is often 0. and represent the principal point, which would be ideally in the centre of the image. Nonlinear intrinsic parameters such as lens distortion are also important although they cannot be included in the linear camera model described by the intrinsic parameter matrix. This process reduces the dimensions of the data taken in by the camera from three to two (light from a 3D scene is stored on a 2D image). Camera resectioning determines which incoming light is associated with each pixel on the resulting image. In an ideal pinhole camera, a simple projection matrix is enough to do this. With more complex camera systems, errors resulting from misaligned lenses and deformations in their structures can result in more complex distortions in the final image. The camera projection matrix is derived from the intrinsic and extrinsic parameters of the camera, and is often represented by the series of transformations; e.g., a matrix of camera intrinsic parameters, a  $3 \times 3$  rotation matrix, and a translation vector. The camera projection matrix can be used to associate points in a camera's image space with locations in 3D world space.

## I. PROBLEM 1

I was unable to use optimal thresholding due to the fact that the image contains yellow text, which falls below the Otsu's thresholding value. To compensate, a higher threshold was used. Unfortunately, using such a high threshold grabs a lot of noise from the image, and even combines some letter pairs into single objects. I worked around this issue by selecting one of the *m*'s from the yellow text, and evaluating its regionprops. Because the *m* chosen was smaller in size than the majority of the other *m*'s, I set a testing range for the regionprops. The values of the testing regionprops had to be within  $90\% < m < 140\%$ . Satisfying these conditions, there were 51 *m*'s found in the image.

```
clc; close all;
clearvars -except I1 I2 I3; close all
%thresh=graythresh(I1);
thresh=0.83;
I = im2bw(I1,thresh); %convert to bw
I = imcomplement(I); %objects are white in matlab

%I=imerode(I,B);
%I=imclose(I,B);
imshow(I);

%example M
%bigM=imcrop(I, [173,47,(195-173),(63-47)]);
```

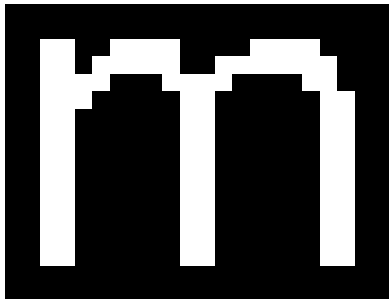
```
smallM=imcrop(I, [525,495,(546-525),(511-495)]);
;
theM=smallM;
figure('name','the m'), imshow(theM), title('the m');
mSTATS=regionprops(theM,'all');

reconstructed = false(size(I));
stats = regionprops(I,'all');
figure('name','ms are highlighted');
imshow(I, title('ms are highlighted')); hold on
;

lowerBound=0.9;
upperBound=1.4;
for i=1:size(stats)
    if ( (stats(i).Solidity < (mSTATS.Solidity*upperBound)) && ...
        && (stats(i).Solidity > (mSTATS.Solidity*lowerBound)) && ...
        (stats(i).FilledArea < (mSTATS.FilledArea*upperBound)) && ...
        (stats(i).FilledArea > (mSTATS.FilledArea*lowerBound)) && ...
        (stats(i).Area < (mSTATS.Area*upperBound)) && ...
        (stats(i).Area > (mSTATS.Area*lowerBound)) && ...
        (stats(i).EulerNumber == 1) && ...
        (stats(i).ConvexArea < (mSTATS.ConvexArea*upperBound)) && ...
        (stats(i).ConvexArea > (mSTATS.ConvexArea*lowerBound)) && ...
        (stats(i).Perimeter > (mSTATS.Perimeter*lowerBound)) && ...
        (stats(i).Perimeter < (mSTATS.Perimeter*upperBound)) && ...
        (stats(i).Extent > (mSTATS.Extent*lowerBound)) && ...
        (stats(i).Extent < (mSTATS.Extent*upperBound)) && ...
        (stats(i).MajorAxisLength < (mSTATS.MajorAxisLength*upperBound)) && ...
        (stats(i).MajorAxisLength > (mSTATS.MajorAxisLength*lowerBound)) && ...
        (stats(i).MinorAxisLength < (mSTATS.MinorAxisLength*upperBound)) && ...
        (stats(i).MinorAxisLength > (mSTATS.MinorAxisLength*lowerBound)) && ...
        (stats(i).Eccentricity < (mSTATS.Eccentricity*upperBound)) && ...
        (stats(i).Eccentricity > (mSTATS.Eccentricity*lowerBound)))
        N= floor(stats(i).Centroid(1));
        M= floor(stats(i).Centroid(2));
        reconstructed(M,N) = true;
        plot(N,M,'s','color','red')
    end
end
hold off;
```

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the m



ms are highlighted

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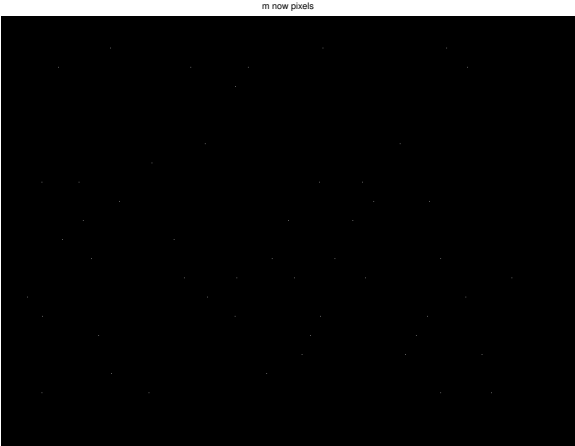
A. Part 2: Transform m's Into Pixels

To transform the m's into single pixels, we can simply run a hit and miss operation using our m mask and the

complement of the m mask. But because I elected to use region properties for this example, it was relatively easy to just include a centroid calculation and toggle the centroid of the detected m into a true pixel value. This code can be seen in the previous section's for loop. The total number of m's in the image was 51.

```
figure('name','m are now pixels'), imshow(
    reconstructed), title('m now pixels')
mCount = sum(sum(reconstructed))
```

mCount =  
  
51



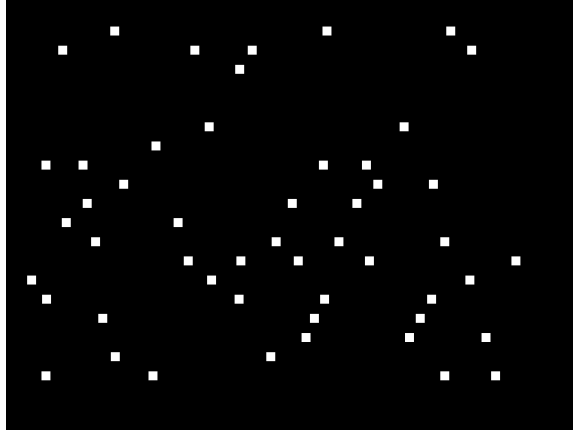
B. Part 3: Transform m's into squares's

I chose to transform the m's in the image into squares. By exploiting the functionality of dilation, I dilate the pixel image with a square, and add the resultant image to the original image.

```
remover=zeros(size(theM));
remover=imcomplement(remover);
Img = imdilate(reconstructed,remover);

square = strel('square',15);
add = imdilate(reconstructed, square);
figure('name','adder'), imshow(adder)
Img2=floor(I+Img); %figure('name','img'),
    imshow(Img2);
Im3=Img2+adder;

figure('name','m are now squares'), imshow(Im3)
    , title('m now different')
```



The intrinsic matrix containing 5 intrinsic parameters. These parameters encode focal length, image format, and principal point. The parameters represent focal length in terms of pixels, where  $f_x$  and  $f_y$  are the scale factors relating pixels to distance.  $s$  represents the skew coefficient between the  $x$  and the  $y$  axis, and is often 0, and represent the principal point, which would be ideally in the centre of the image. Nonlinear intrinsic parameters such as lens distortion are also important although they cannot be included in the linear camera model described by the intrinsic parameter matrix. This process reduces the dimensions of the data taken in by the camera from three to two (light from a 3D scene is stored on a 2D image). Camera resectioning determines which incoming light is associated with each pixel on the resulting image. In an ideal pinhole camera, a simple projection matrix is enough to do this. With more complex camera systems, errors resulting from misaligned lenses and deformations in their structures can result in more complex distortions in the final image. The camera projection matrix is derived from the intrinsic and extrinsic parameters of the camera, and is often represented by the series of transformations; e.g., a matrix of camera intrinsic parameters, a  $3 \times 3$  rotation matrix, and a translation vector. The camera projection matrix can be used to associate points in a camera's image space with locations in 3D world space.

### C. Part 4: Discussion of Results

After counting the  $m$ 's individually, I have verified that they were all found using my method.

## II. PROBLEM 2

In order to classify these two classes, they are first plotted to determine if they are separable. The plot reveals that they are linearly separable by a plane.

```
clear all; clc; close all;
class1=[0.8963, 1.4780, 0.9023;
1.1023, 0.7144, 0.8438;
0.7078, 1.0981, 1.3255;
0.8516, 1.1091, 0.8739;
1.0255, 1.2991, 1.0301;
0.7408, 0.7857, 0.9575;
0.9521, 0.7835, 1.0452;
1.3166, 0.9372, 0.9489;
0.4496, 0.8064, 0.6517;
0.9034, 1.1314, 1.3808;]

class2=[3.7985, 1.8831, -0.1202;
2.8175, 2.4902, 0.9653;
2.7783, 0.5197, 0.7404;
1.9085, 1.2397, -0.4936;
3.2815, 1.1903, -1.0368;
2.2147, 4.0108, 0.2256;
2.3083, 1.0618, 1.8742;
2.1250, 2.5301, -0.7521;
2.8540, 2.3891, -0.9560;
```

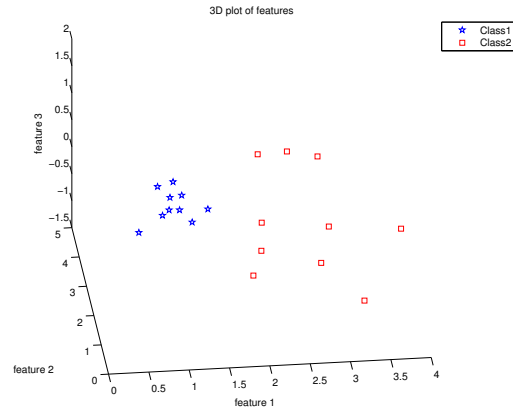
```
2.0397, 1.5494, 0.3092;]
figure('name','3dplot')
plot3(class1(:,1), class1(:,2), class1(:,3), 'p', 'color','blue'), hold on
plot3(class2(:,1), class2(:,2), class2(:,3), 's', 'color','red'), hold on
xlabel('feature 1')
ylabel('feature 2')
zlabel('feature 3')
title('3D plot of features');
legend('Class1','Class2'), hold off
```

```
class1 =

0.8963    1.4780    0.9023
1.1023    0.7144    0.8438
0.7078    1.0981    1.3255
0.8516    1.1091    0.8739
1.0255    1.2991    1.0301
0.7408    0.7857    0.9575
0.9521    0.7835    1.0452
1.3166    0.9372    0.9489
0.4496    0.8064    0.6517
0.9034    1.1314    1.3808
```

```
class2 =

3.7985    1.8831   -0.1202
2.8175    2.4902    0.9653
2.7783    0.5197    0.7404
1.9085    1.2397   -0.4936
3.2815    1.1903   -1.0368
2.2147    4.0108    0.2256
2.3083    1.0618    1.8742
2.1250    2.5301   -0.7521
2.8540    2.3891   -0.9560
2.0397    1.5494    0.3092
```



### A. Part 3: Minimum Distance Classifier

Using the equations defined in class, the following equation was derived in order to find the Minimum Distance Classifier.

$$\%Z = \frac{(-2 * X * \bar{X}_{\{1\}} - (\bar{X}_{\{1\}}^2) + (2 * Y * \bar{Y}_{\{1\}}) - (\bar{Y}_{\{1\}}^2) - (\bar{Z}_{\{1\}}^2))}{2 * (X * \bar{X}_{\{1\}} + Y * \bar{Y}_{\{1\}} + Z * \bar{Z}_{\{1\}})}$$

```

        {1})^2)-(2*X*\bar{X_{2}})+(\bar{X_{2}})^2)
        -(2*Y*\bar{Y_{2}})+(\bar{Y_{2}})^2+(\bar{Z_{2}}
        {2})^2){2*(\bar{Z_{2}}-\bar{Z_{1}}))}$
2  clc; close all;

4  %MDC C:
x1b=mean(class1(1,:));
6  x2b=mean(class2(1,:));

8  y1b=mean(class1(2,:));
y2b=mean(class2(2,:));

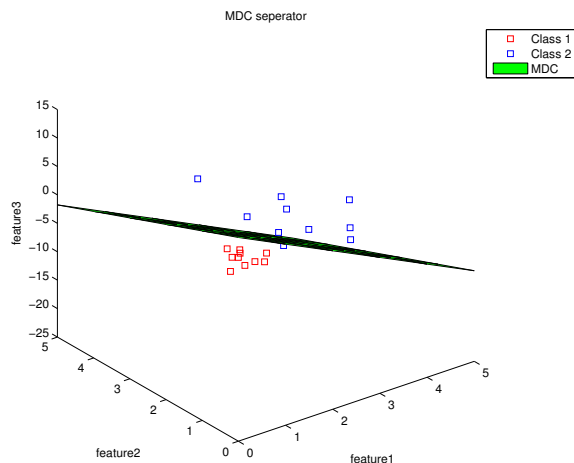
10 z1b=mean(class1(3,:));
12 z2b=mean(class2(3,:));

14 X=0:5;
Y=0:5;

16 Z=zeros(6,6);
for i=0:5
    for j=0:5
18        Z(i+1,j+1)=((2*i*x1b)-(x1b^2)+(2*j*y1b)
20            -(y1b^2)-(z1b^2)-(2*i*x2b)+(x2b^2)-(2*j*y2b
            +(y2b^2)+(z2b^2))/(2*(z2b-z1b));
    end
22 end

24 figure('name','MDC wooooo');
%
26 %surf(X,Y,Z), hold on
28 plot3(class1(:,1), class1(:,2), class1(:,3), 's
    ','color','red'), hold on
30 plot3(class2(:,1), class2(:,2), class2(:,3), 's
    ','color','blue'), hold on
32 surf(X, Y, Z, colormap([0,1,0]));
%axis([0, 2, 0, 2, 0, 2]), title('MDC wooooooo
    ');
34 xlabel('feature1'); ylabel('feature2'); zlabel(
    'feature3');
title('MDC seperator'); legend('Class 1', '
    Class 2', 'MDC'), hold off;

```



### B. Part 4: Perceptron

```

1  clc;
2  clear Z
3  %build training set

```

```

trainingSet = [class1;class2]
[m, n] = size(trainingSet);
resultSet=ones(1,m)';
resultSet(11:20)=0;
weightVector = ones(1,n)/5; %

threshold = 0;%threshold to decide if the
    output is good or bad. usually this is 0
error_count = 1;
bias = 0.1;
iterationNo = 1;
learningRate = 1;
% training phase
while (error_count > 0)
    error_count = 0;
    for i=1:m
        gx=dot(weightVector,trainingSet(i,:))+
            bias;
        if (gx > threshold)
            result = 1;
        else
            result = 0;
        end

        error = resultSet(i)-result;
        if (error ~= 0)
            error_count = error_count + 1;
            weightVector = weightVector + (
                learningRate*(error))*trainingSet(i,1:n);
            bias = bias + learningRate*
                error;
        end
    end

    if (iterationNo >= 100000)
        disp('Neuron input calculation couldn''
            t completed in timely fashion. ');
        break
    end
    iterationNo = iterationNo +1;
end
disp(['answer converged in ' num2str(
    iterationNo) ' iterations']);
disp(['weights: ' num2str(weightVector)]);
disp(['bias: ' num2str(bias)]);

43 X=0:5;
45 Y=0:5;
47 Z=zeros(6,6);
for i=0:5
    for j=0:5
49        Z(i+1,j+1)=(-(weightVector(1)*i)-(
            weightVector(2)*j)-bias)/weightVector(3);
    end
51 end

53 figure('name','Now with NN!');
55 plot3(class1(:,1), class1(:,2), class1(:,3), 's
    ','color','red'), hold on
57 plot3(class2(:,1), class2(:,2), class2(:,3), 's
    ','color','blue'), hold on
59 surf(X,Y,Z, colormap([0,1,0]), hold on
    xlabel('feature1'); ylabel('feature2'); zlabel(
        'feature3');
61 legend('Class 1', 'Class 2', 'Perceptron'),
    title('Perceptron Solution'), hold off;

```

```

trainingSet =

```

```

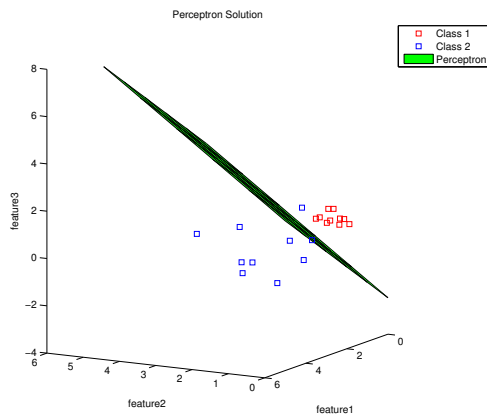
0.8963    1.4780    0.9023
1.1023    0.7144    0.8438
0.7078    1.0981    1.3255
0.8516    1.1091    0.8739
1.0255    1.2991    1.0301
0.7408    0.7857    0.9575
0.9521    0.7835    1.0452
1.3166    0.9372    0.9489
0.4496    0.8064    0.6517
0.9034    1.1314    1.3808
3.7985    1.8831   -0.1202
2.8175    2.4902    0.9653
2.7783    0.5197    0.7404
1.9085    1.2397   -0.4936
3.2815    1.1903   -1.0368
2.2147    4.0108    0.2256
2.3083    1.0618    1.8742
2.1250    2.5301   -0.7521
2.8540    2.3891   -0.9560
2.0397    1.5494    0.3092

```

```

answer converged in 7 iterations
weights: -3.8326    -0.9679    3.5008
bias: 3.1

```



### C. Part 5 Support Vector Machine

The equations derived in class were modified to produce the optimization problem. First, the data points were inspected and the points that were closest to the other groups were chosen. Next, a  $Q$  function was solved in terms of  $\alpha$  and dot product of the closest classes. Setting the differential of this  $Q = 0$ ,  $\alpha$  was solved. This was plugged into the equations to produce a weight matrix and a bias, and the plane was plotted.

```

1 clc; close all;
2 clearvars -except class1 class2
3
4 x1=[1.3166, 0.9372, 0.9489]'
5 x2=[2.0397, 1.5494, 0.3092]'
6
7 x11=dot(x1,x1);
8 x12=dot(x1,x2);
9 x22=dot(x2,x2);
10
11 %syms a X Y Z

```

```

12 %Q=a+a*(1/2)*((a^2)*x11-(2*a*a*x12)+(a^2)*x22);
13 %alpha=double(solve(diff(Q)==0,a))
14
15 alpha = 1.5304 %FOUND BY USING THE ABOVE
16 SYMBOLIC TOOLBOX CODE^^^^
17
18 W=(alpha*(1)*x1) + (alpha*(-1)*x2)
19 b=1-(alpha*(1)*x11+alpha*(-1)*x22)
20
21 %SVM_solX=(-W(2)*Y-W(3)*Z-b)/W(1);
22 X=0:5;
23 Y=0:5;
24
25 Z=zeros(5,5);
26 for i=0:5
27     for j=0:5
28         Z(i+1,j+1)=(-(W(1)*i)-(W(2)*j)-b)/W(3);
29     end
30 end
31
32 figure('name','Now with SVM!');
33 plot3(class1(:,1), class1(:,2), class1(:,3), 's', 'color','green'), hold on
34 plot3(class2(:,1), class2(:,2), class2(:,3), 's', 'color','blue'), hold on
35 surf(X,Y,Z, colormap([1,0,0])); hold on;
36 xlabel('feature1'); ylabel('feature2'); zlabel('feature3');
37 legend('Class 1', 'Class 2', 'SVM'), title('SVM Solution'); hold off;

```

```
x1 =
```

```

1.3166
0.9372
0.9489

```

```
x2 =
```

```

2.0397
1.5494
0.3092

```

```
alpha =
```

```
1.5304
```

```
W =
```

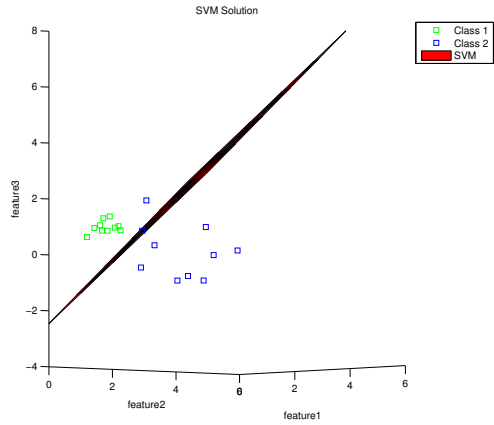
```

-1.1066
-0.9369
0.9790

```

```
b =
```

```
2.4061
```



#### *D. Part 6 Discussion of Results*

This problem was slightly harder to understand because the classes were defined to have three dimensions, so visualizing their distributions was a little more complicated than the two dimensional examples that I was used to. The separators all worked very effectively, but it seemed that the MDC method produced the best separator. This was surprising because the MDC separator worked better than the SVM method, which by theory should be the optimal solution. This is likely because I plotted the flat plane produced by the output of the SVM method, as opposed to the ideal kernel that would be produced through using the quadprog function.