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# Medical Image Analysis Assignment

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## 1)

```
clear all
close all
clc
clf

load MyImages

%a
RGB_im = ind2rgb(Question1Image,Question1cmap); %Generete RGB-image.

%b
Gray_im = rgb2gray(RGB_im); %Convert to gray scale.

%c
seg = Gray_im == 1; %Segment the allocated number

%d
subplot(1,3,1)
imshow(RGB_im);
title('1a - RGB','FontSize',14)

subplot(1,3,2)
imshow(Gray_im);
title('1b - Gray','FontSize',14)

subplot(1,3,3)
imshow(seg);
title('1c - Binary','FontSize',14)
```



2)

```
clc
clf
close all

%a

%[im3,TFORM,overlay] =
  Registratiodemo(Question2Image_B,Question2Image_A); %Select common
  points and

      %generate the affine transfrom object.
%Did the above and saved the data in MyImages.

subplot(1,3,1)
imshow(Question2Image_B);
title('Image B','FontSize',14)

subplot(1,3,2)
imshow(Question2Image_A);
title('Image A','FontSize',14)

subplot(1,3,3)
imshow(im3); %The registrered image B.
```

```

title('Registrered image B','FontSize',14)

%b
A=TFORM.tdata.T(1:2,1:2); %The affine transformation matrix without
    translation data.
[U,S,V]=svd(A); %Break down into three components: rotation U,
    deformation S (diagonal -> scaling), rotation V.
xscale = S(1,1); %Scaling value along x-axis.
yscale = S(2,2); %Scaling value along y-axis.
display('The amount that imageB has been enlarged in x direction after
    the first rotation')
display(xscale)
display('The amount that imageB has been enlarged in y direction after
    the first rotation')
display(yscale)

%c
ROTX = [1 0]*A; %How much imageB was rotated (Just multiply affine
    transform by x base vector).
x_rotation = atan2(ROTX(2),ROTX(1))*360/(2*pi); %The angle in degrees.
display('How much imageB was rotated in degrees')
display(x_rotation)

The amount that imageB has been enlarged in x direction after the
    first rotation

xscale =

    1.0388

The amount that imageB has been enlarged in y direction after the
    first rotation

yscale =

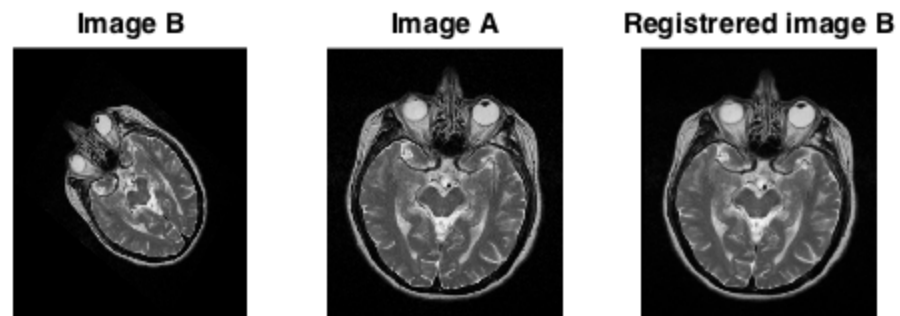
    0.8350

How much imageB was rotated in degrees

x_rotation =

    48.8640

```



3)

```
clc
clf
close all

warning('off','all') %Supress "Image is too big to fit on screen;
    displaying at 67%".

%a
Q3_RGB = ind2rgb(Question3Image,Question3cmap); % Convert to RGB.

%b
figure(1)
subplot(1,3,1)
imshow(Q3_RGB);
title('RGB image','FontSize',14)

subplot(1,3,2)
Q3_gray = rgb2gray(Q3_RGB); %Phase contrast channel.
imshow(Q3_gray)
title('Phase contrast channel','FontSize',14)

subplot(1,3,3)
```

```

Q3_g = Q3_RGB;
Q3_g(:,:, [1, 3]) = 0; %Fluorescence channel.
imshow(Q3_g)
title('Fluorescence channel','FontSize',14)

%c
Segmentnuclei_self = Q3_g(:,:,2) > 0.5; %Manually set threshold.
threshold=graythresh(Q3_g(:,:,2)); %Global threshold through Otsu's
method.
Segmentnuclei_otsu=im2bw(Q3_g(:,:,2),threshold); %Create binary image
from the threshold.

figure(2)
subplot(1,3,1)
imshow(Q3_gray)
title('Grayscale image','FontSize',14)

subplot(1,3,2)
imshow(Segmentnuclei_self)
title('Manual threshold','FontSize',14)

subplot(1,3,3)
imshow(Segmentnuclei_otsu)
title('Otsu method','FontSize',14)

%Otsu's method contains more of the nuclei than the manually set.
%Didn't
%spend much time on the manually set threshold though. If carefully
%setting
%the threshold by some trial and error it would probably be better
%then
%Otsu's. I.e manual will be better then automated tresholds if you
%have
%the time to tune it.

%d
figure(3)
[BW,threshold] = edge(Q3_gray,'log'); %Find edges using LOG edge
detection method.
subplot(2,2,1)
imshow(BW)
title('After LOG-filter','FontSize',14)

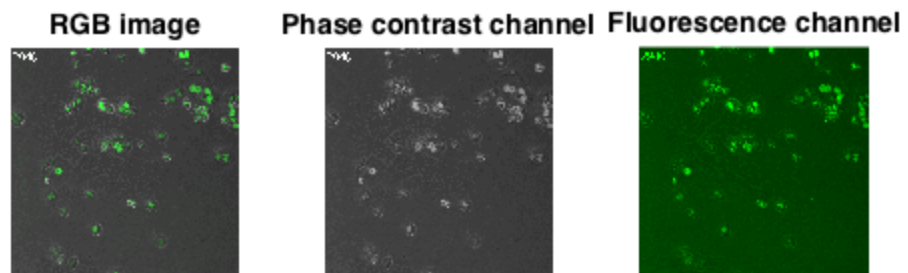
BW2 = imclose(BW,strel('disk',6)); %Close to smooth and connect
irregular features
subplot(2,2,2)
imshow(BW2)
title('After closing','FontSize',14)

BW2 = imfill(BW2,'holes'); %Fill the holes in the cell bodies.
subplot(2,2,3)
imshow(BW2)
title('After filling holes','FontSize',14)

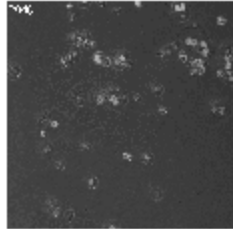
```

```
BW2 = imopen(BW2,strel('disk',10)); %Open to remove small scatter
                                     %while preserving the shape and
                                     size of the cell bodies.
subplot(2,2,4)
imshow(BW2)
title('After opening, done','FontSize',14)

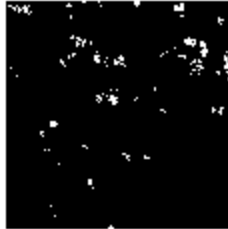
figure;
imshow(Q3_RGB)
hold on
himage = imshow(BW2);
himage.AlphaData = 0.4; %Add transparency.
title('The segmenting on top of the original image','FontSize',14)
```



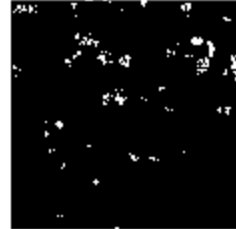
**Grayscale image**



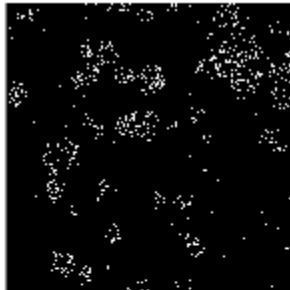
**Manual threshold**



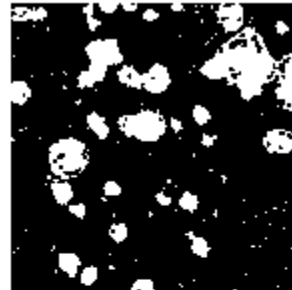
**Otsu method**



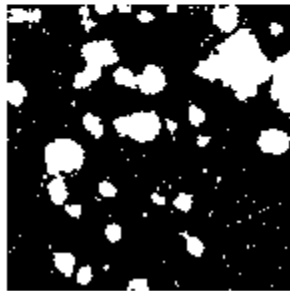
**After LOG-filter**



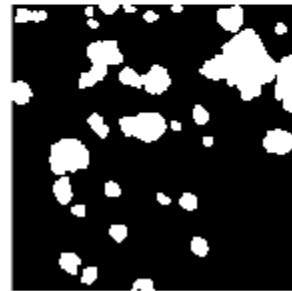
**After closing**

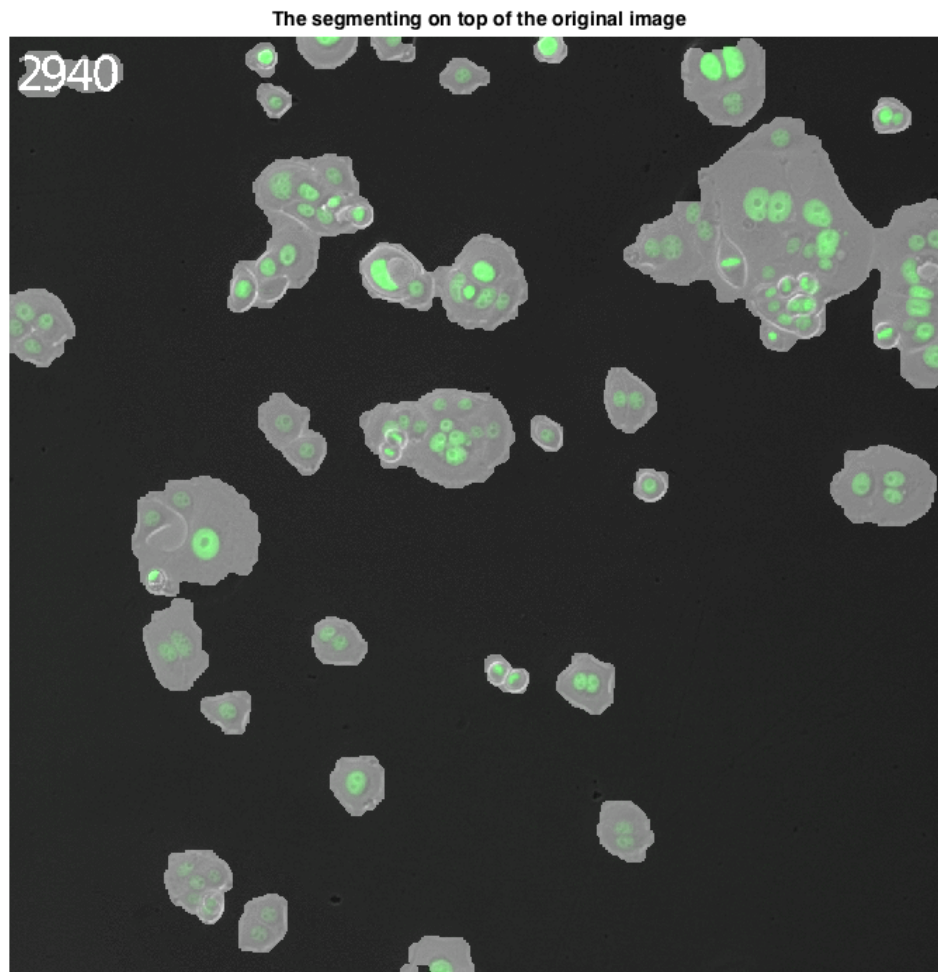


**After filling holes**



**After opening, done**





4)

```
close all
clc
clf

img = Q3_RGB(:,:,2) - Q3_RGB(:,:,1); %Remove the cell bodies but not
nuclei.

subplot(3,3,1)
imshow(img)
title('Original image','FontSize',10)

img = imgaussfilt(img,1.1); %Gaussian filter for noise removal with
edge preservation.
subplot(3,3,2)
imshow(img)
title('After Gaussian filter','FontSize',10)
```



```

img = adapthisteq(img); %Imrove local contrast and enhance the
    definitions of edges
                                %in each region through adaptive histogram
    equalization.
subplot(3,3,3)
imshow(img)
title('After adaptive histogram equalization','FontSize',10)

threshold=graythresh(img); %Create binary image through Otsu's method.
img=im2bw(img,threshold*0.8); %Scale threshold (absence of noise).
subplot(3,3,4)
imshow(img)
title('After (scaled) Otsu's method','FontSize',10)

img = imdilate(img, strel('disk',2)); %Dilate to enlarge nuclei.
subplot(3,3,5)
imshow(img)
title('After dilation','FontSize',10)

img = bwareaopen(img, 75); %Remove scatter.
subplot(3,3,6)
imshow(img)
title('After removing small-area-objects','FontSize',10)

img = imopen(img, strel('disk',5)); %Open to remove small objects but
    preserve nuclei.
subplot(3,3,7)
imshow(img)
title('After opening','FontSize',10)

C = ~img; %Invert binary image.
D = -bwdist(C,'quasi-euclidean'); %Distance transform using quasi-
    euclidean distance.

D(C) = -inf; %Background to -inf.
regmax = imregionalmax(D); %Find regional maxima.
regmax = imdilate(regmax,strel('rectangle',[1 1])); %Dilate to
    coalesce several regional maximas of nuclei.
regmax = imdilate(regmax,strel('disk',1));
D(regmax) = -inf; %To -inf.
% Watershed will now fill from the centre of nuclei to its edge.

subplot(3,3,7)
imshow(D,[],'InitialMagnification','fit');
title('Distance-transform','FontSize',10)

L = watershed(D); %Do the watershed-transform.
rgb = label2rgb(L,'hsv','b','shuffle');
subplot(3,3,8)

```

```

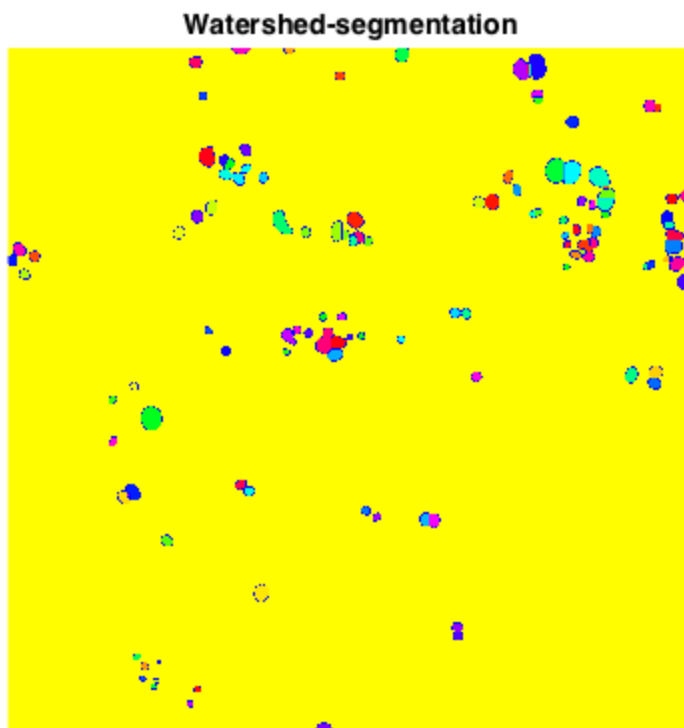
imshow(rgb,'InitialMagnification','fit')
title('Watershed-segmentation','FontSize',10)

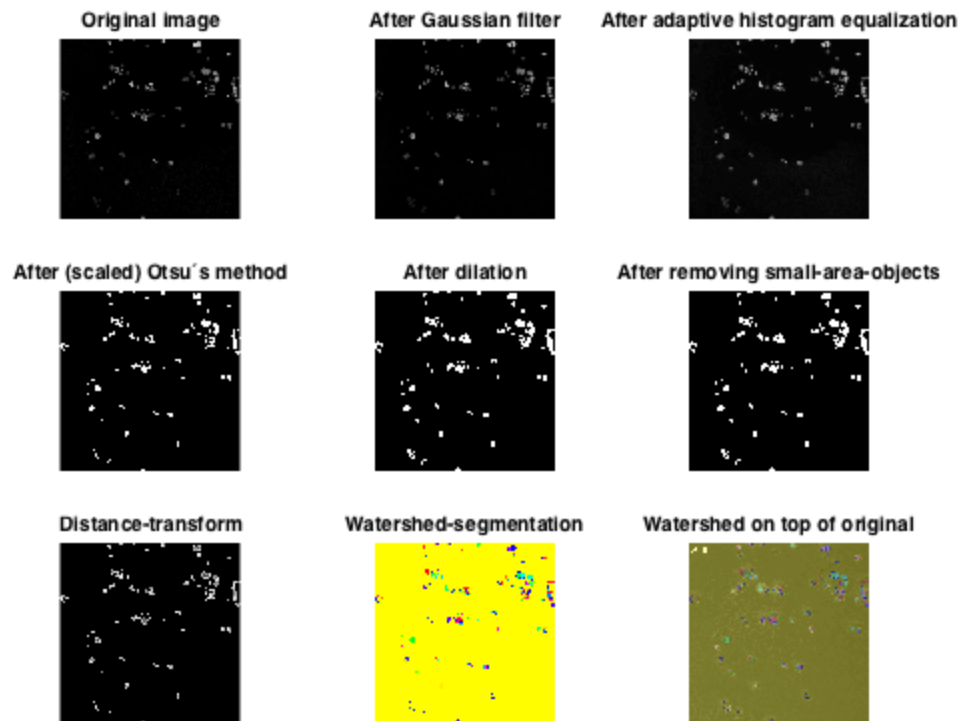
figure;
imshow(rgb,'InitialMagnification','fit') %Show in seperate image for
visibility.
title('Watershed-segmentation','FontSize',14)

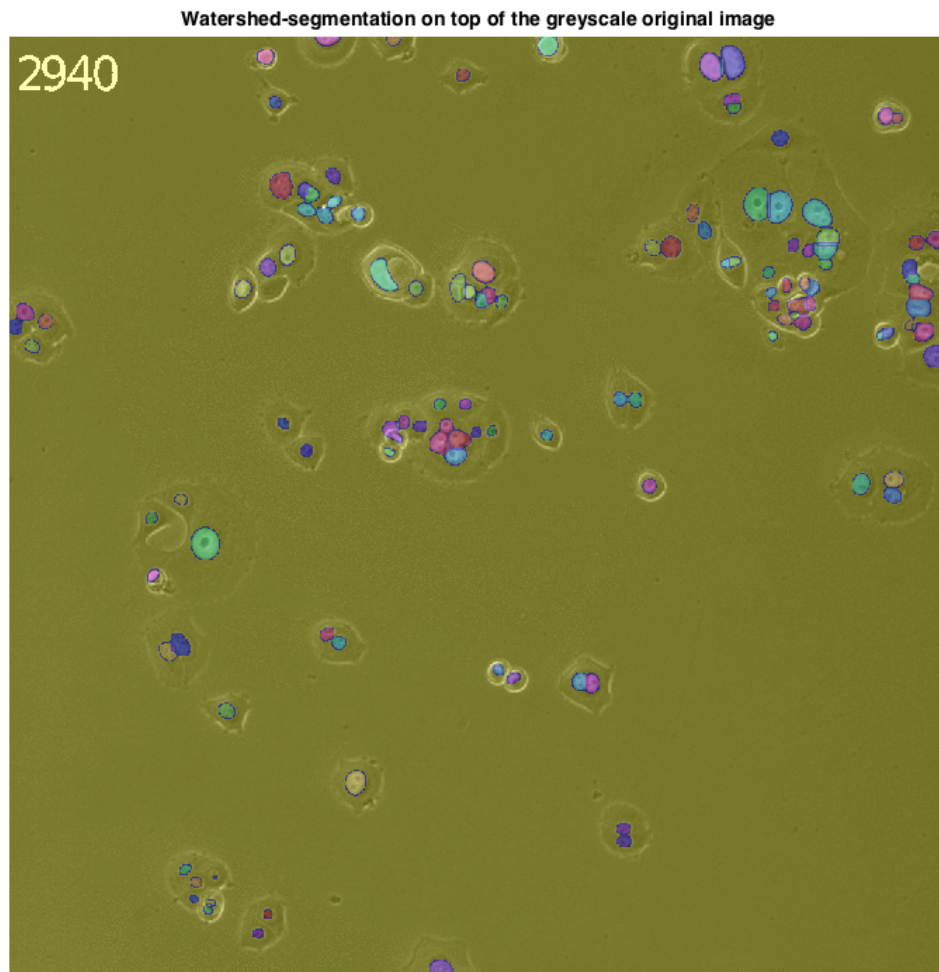
figure(1)
subplot(3,3,9)
imshow(Q3_gray)
hold on
himage = imshow(rgb); %Show segmentation on top of original greyscale
image.
himage.AlphaData = 0.3;
title('Watershed on top of original','FontSize',10)

figure;
imshow(Q3_gray)
hold on
himage = imshow(rgb); %Show in seperate image for visibility.
title('Watershed-segmentation on top of the greyscale original
image','FontSize',14)
himage.AlphaData = 0.3;

```







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