Homework 4: Morphology Segmentation

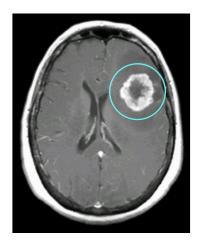
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BMI/CS 567: Medical Image Analysis

4.1 Segmentation

Use any combination of the segmentation techniques we have studied in class (region growing, erosion, dilation, thresholding, etc.) to segment the metastatic tumor highlighted in the following brain MRI:



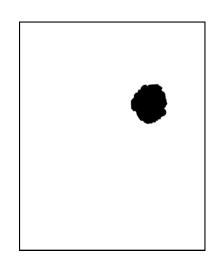
You can find this MRI (metastatic.jpg) on the course website. Aside from the techniques we have studied in class, you may also use any other technique you want (e.g., something you find on a book, research papers, the internet, or something you come up with), but you must explain such technique clearly, where you got it from, and why you chose to use it. Hand in your code and your segmentation.

```
brain = imread('metastatic.jpg');
brain = double(rgb2gray(brain));

seedPoint = [161,236]; % Initial seedpoint (selected near bottom edge of tumor)
threshold = 8; % Region growing criteria
segmented = regionGrow(brain, seedPoint, threshold); % Region grow in tumor
segmented = dilation(segmented); % Dilate to connect inside
```

4.2 Evaluation

On the course website you can also find the file ground_truth.png, which contains the handsegmented tumor:



(a) Compute the Hausdorff distance between your segmentation and the ground truth.

```
truth = imread('ground_truth.png');
truth = rgb2gray(truth)<255; % Binarize and inverse image</pre>
truth = imresize(truth, size(brain)); % Make segmented and ground truth images same size
segmented = imbinarize(segmented); % Binarize segmented image
%Create indices of boundary points
contourSegmented = cell2mat(bwboundaries(segmented)); %Border points in segmented (Nx2)
contourTruth = cell2mat(bwboundaries(truth)); %Get all border points in segmented (Mx2)
% Calculate Hausdorff
for i=1:length(contourSegmented)
    for j=1:length(contourTruth)
        % Find euclidean distance between points
        distances1(i,j) = norm(contourSegmented(i,:)-contourTruth(j,:));
    end
end
firstTerm = max(min(distances1,[],2)); % Get maximum of all mins
for i=1:length(contourTruth)
    for j=1:length(contourSegmented)
        distances2(i,j) = norm(contourTruth(i,:)-contourSegmented(j,:));
    end
end
secondTerm = max(min(distances2,[],2)); % Get maximum of all mins
hausdorff = max(firstTerm, secondTerm); % Find max of either term
disp(['Hausdorff Distance = ' num2str(hausdorff)]);
```

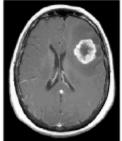
Hausdorff Distance = 3.6056

(b) Compute the Dice coefficient between your segmentation and the ground truth.

```
% Compute dice coefficient
union = segmented.*truth; % Find shared active pixels
dice = 2*( sum(union(:)) )./( sum(truth(:)) + sum(segmented(:)) );
disp(['Dice Coefficient = ' num2str(dice)]);
```

```
figure; subplot(2,2,1); imshow(brain,[]); title('Original Image');
subplot(2,2,2); imshow(truth); title('Ground Truth Image');
subplot(2,2,3); imshow(brain.*segmented,[]); title('My Segmentation');
subplot(2,2,4); imshow(brain.*truth,[]); title('True Segmentation');
```

Original Image



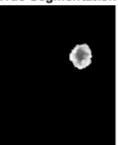
Ground Truth Image



My Segmentation



True Segmentation



Custom Functions

Region Growing

```
function segmented = regionGrow(image, seedPoint, threshold)
    segmented = zeros(size(image)); % Initialize segmented matrix
    segmented(seedPoint(1), seedPoint(2)) = 1; % Starting seed point
    tried = zeros(size(image)); % Initialize already tried points matrix
    image = padarray(image,[1 1]); % Pad array with zeros so there are no edge issues
    while sum( tried(:) ~= segmented(:) )>0 % While there are still points to try ...
    [tryX,tryY] = find(segmented - tried); % Get all coordinates to try
    for i = 1:length(tryX)
        tried(tryX(i),tryY(i)) = 1; % Add test coordinates to 'tried' array
        value = image(tryX(i),tryY(i)); % Get value of current pixel

    % If pixel to the top left of the current pixel is within threshold
```

```
if abs( image(tryX(i)-1,tryY(i)-1)-value ) < threshold</pre>
                 % Then add this point to the segmented image
                 segmented(tryX(i)-1,tryY(i)-1) = 1;
            end
            if abs( image(tryX(i)-1,tryY(i))-value ) < threshold %top pixel</pre>
                 segmented(tryX(i)-1,tryY(i)) = 1;
            end
            if abs( image(tryX(i)-1,tryY(i)+1)-value ) < threshold %top right pixel</pre>
                 segmented(tryX(i)-1,tryY(i)+1) = 1;
            end
            if abs(image(tryX(i),tryY(i)-1)-value ) < threshold %left pixel</pre>
                 segmented(tryX(i),tryY(i)-1) = 1;
            end
            if abs( image(tryX(i),tryY(i)+1)-value ) < threshold %right pixel</pre>
                 segmented(tryX(i),tryY(i)+1) = 1;
            end
            if abs( image(tryX(i)+1,tryY(i)-1)-value ) < threshold %bot left pixel</pre>
                 segmented(tryX(i)+1,tryY(i)-1) = 1;
            end
            if abs( image(tryX(i)+1,tryY(i))-value ) < threshold %bottom pixel</pre>
                 segmented(tryX(i)+1,tryY(i)) = 1;
            end
            if abs( image(tryX(i)+1,tryY(i)+1)-value ) < threshold %bot right pixel</pre>
                 segmented(tryX(i)+1,tryY(i)+1) = 1;
            end
        end
    end
end
```

Dilation

Erosion

```
function eroded = erosion(image)
  eroded = image;
  for i=2:size(image,1)-1
    for j=2:size(image,2)-1
       block = image(i-1:i+1,j-1:j+1);
       if sum(block(:))<9</pre>
```

```
eroded(i-1:i+1,j-1:j+1) = 0;
end
end
end
end
end
```