

IVA SEMINAR

LEVEL SET SEGMENTATION OF TUMOR DETECTION

MAHALAKSHMI S

205229118

%% Input

```
s=imread('br.PNG');  
figure;  
imshow(s);  
title('Input image','FontSize',20);
```

%% Filter or preprocessing image

```
num_iter = 10;  
delta_t = 1/7;  
kappa = 15;  
option = 2;  
disp('Preprocessing image please wait . . .');  
inp = anisodiff(s,num_iter,delta_t,kappa,option);  
inp = uint8(inp);
```

```
inp=imresize(inp,[256,256]);  
if size(inp,3)>1  
    inp=rgb2gray(inp);  
end
```

%% Binarization

```
sout=imresize(inp,[256,256]);
t0=60;
th=t0+((max(inp(:))+min(inp(:)))./2);
for i=1:1:size(inp,1)
    for j=1:1:size(inp,2)
        if inp(i,j)>th
            sout(i,j)=1;
        else
            sout(i,j)=0;
        end
    end
end
```

%% Morphological Operation

```
label=bwlabel(sout);
stats=regionprops(logical(sout),'Solidity','Area','BoundingBox');
density=[stats.Solidity];
area=[stats.Area];
high_dense_area=density>0.6;
max_area=max(area(high_dense_area));
tumor_label=find(area==max_area);
tumor=ismember(label,tumor_label);
```

%% Bounding box

```
box = stats(tumor_label);
wantedBox = box.BoundingBox;
```

%% Getting Tumor Outline - image filling, eroding, subtracting
% erosion the walls by a few pixels

```
dilationAmount = 5;
```

```

rad = floor(dilationAmount);
[r,c] = size(tumor);
filledImage = imfill(tumor, 'holes');
for i=1:r
    for j=1:c
        x1=i-rad;
        x2=i+rad;
        y1=j-rad;
        y2=j+rad;
        if x1<1
            x1=1;
        end
        if x2>r
            x2=r;
        end
        if y1<1
            y1=1;
        end
        if y2>c
            y2=c;
        end
        erodedImage(i,j) = min(min(filledImage(x1:x2,y1:y2)));
    end
end

```

%% subtracting eroded image from original BW image

```

tumorOutline=tumor;
tumorOutline(erodedImage)=0;

```

%% Level Set Segmentation

```

switch flag

```

```

    case 1

```

```

    Img=imread('b.jpg');
    Img=double(Img(:,:,2));
    title('100 iterations')

```

```
nu=0.01*255*255;  
sigma = 6;  
iter_outer=200;  
iter_inner=150;  
n =[50;90]; m=[30;90];
```

case 2

```
Img=double(Img(:,:,2));  
title('200 iterations')  
nu=0.01*255*255;  
sigma = 35; % or 20  
iter_outer=150;  
iter_inner=120;  
n =[400;200]; m=[142;247];
```

case 3

```
Img=double(Img(:,:,2));  
title('300 iterations')  
nu=0.1*255*255;  
sigma = 12; % or 6  
iter_outer=100;  
iter_inner=100;  
n =[176;245]; m=[59;114];
```

case 4

```
Img=double(Img(:,:,2));  
title('400 iterations')  
nu=0.1*255*255;  
sigma = 35;  
iter_outer=100;  
iter_inner=70;  
n =[204;70]; m=[25;207];
```

case 5

```

Img=double(Img(:,:,2));
title('500 iterations')
nu=0.1*255*255;
sigma = 12;
iter_outer=100;
iter_inner=150;
n =[190;117]; m=[71;135];

```

case 6

```

Img=double(Img(:,:,1));
title('Final contour, 617 iterations')
nu=0.01*255*255;
sigma = 12;
iter_outer=100;
iter_inner=150;
n =[130;209]; m=[140;49];

```

%% Inserting the outline in filtered image in green color

```

rgb = inp(:,:, [1 1 1]);
red = rgb(:,:,1);
red(tumorOutline)=255;
green = rgb(:,:,2);
green(tumorOutline)=0;
blue = rgb(:,:,3);
blue(tumorOutline)=0;
tumorOutlineInserted(:,:,1) = red;
tumorOutlineInserted(:,:,2) = green;
tumorOutlineInserted(:,:,3) = blue;

```

%% Display Together

```

set(0,'defaultfigureposition',[20 60 1500 700])
figure
subplot(2,3,1);imshow(s);title('Input image','FontSize',20);
subplot(2,3,1);title('Input image','FontSize',20);

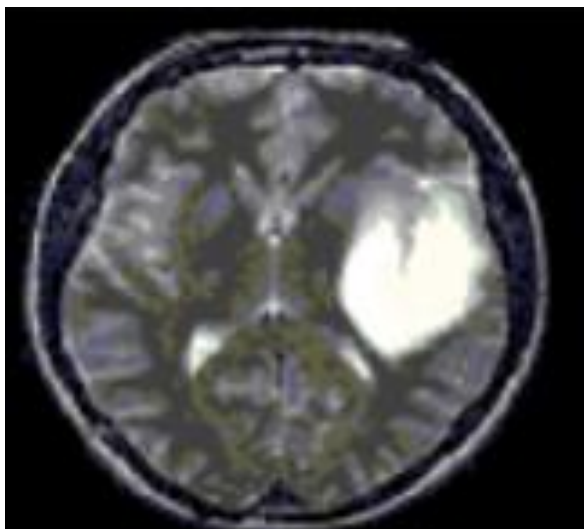
```

```

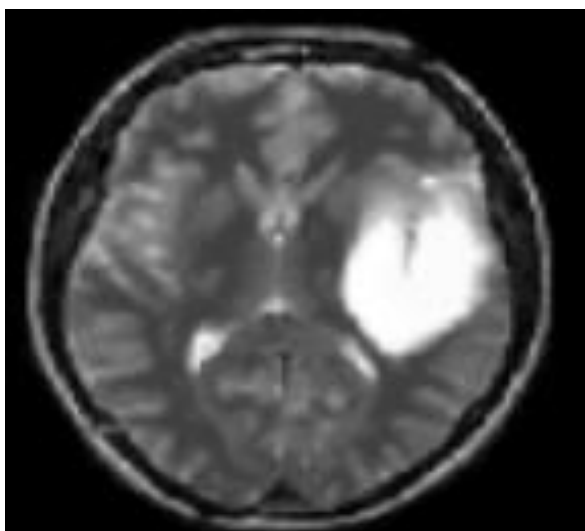
subplot(2,3,2);title('Filtered image','FontSize',20);
subplot(2,3,3);title('Bounding Box','FontSize',20);
subplot(2,3,4);title('Tumor alone','FontSize',20);
subplot(2,3,5);title('Tumor Outline','FontSize',20);
subplot(2,3,6);title('Detected Tumor','FontSize',20);
subplot(2,3,2);imshow(inp);title('Filtered image','FontSize',20);
subplot(2,3,3);imshow(inp);title('Bounding Box','FontSize',20);
hold on;rectangle('Position',wantedBox,'EdgeColor','y');hold off;
subplot(2,3,4);imshow(tumor);title('Tumor alone','FontSize',20);
subplot(2,3,5);imshow(tumorOutline);title('Tumor
Outline','FontSize',20);
subplot(2,3,7);imshow(img);title('100 iterations','FontSize',20);
subplot(2,3,8);imshow(img);title('200 iterations','FontSize',20);
subplot(2,3,9);imshow(img);title('300 iterations','FontSize',20);
subplot(2,3,10);imshow(img);title('400 iterations','FontSize',20);
subplot(2,3,11);imshow(img);title('500 iterations','FontSize',20);
subplot(2,3,12);imshow(img);title('Final contour, 617
iterations','FontSize',20);
subplot(2,3,13);imshow(tumorOutlineInserted);title('Detected
Tumor','FontSize',20);

```

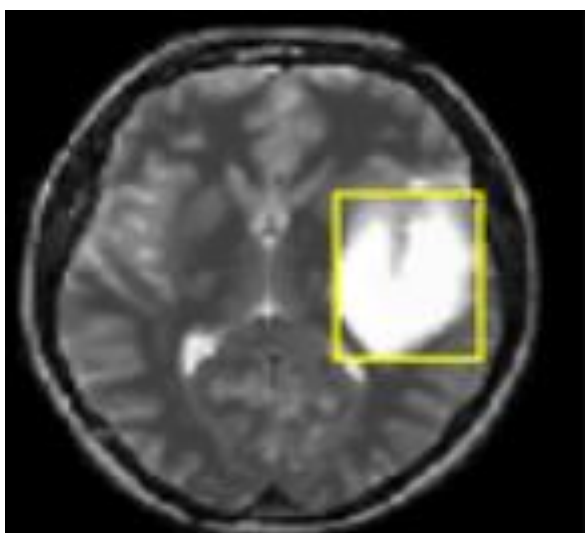
INPUT IMAGE:



Filtered Image:



Bounding Box:



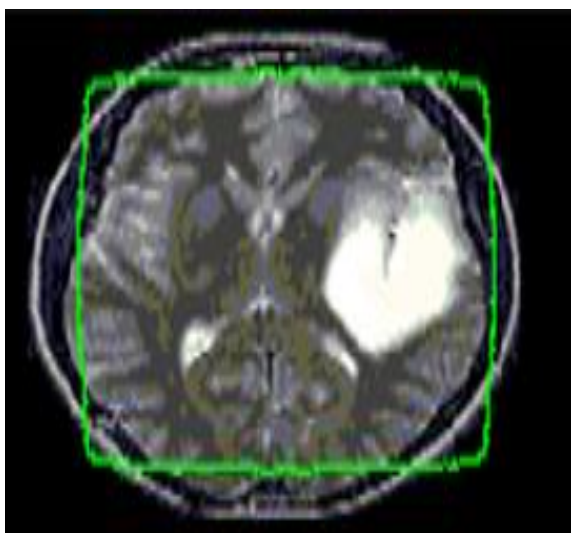
Tumor Alone:



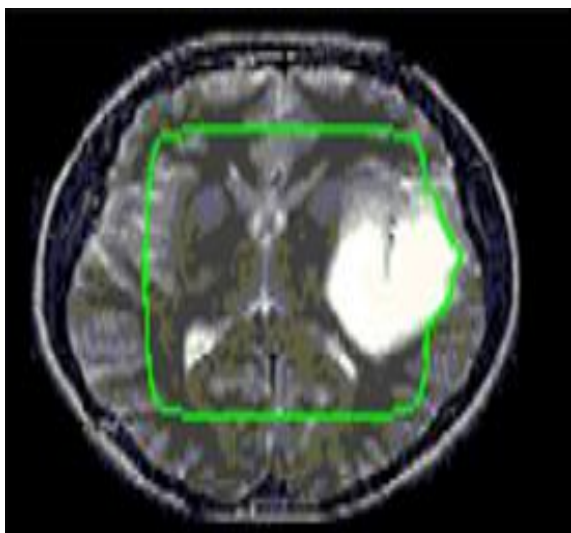
Tumor Outline:



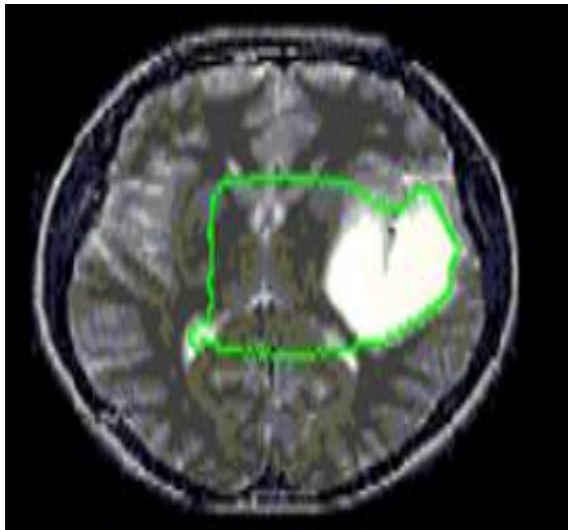
100 iterations:



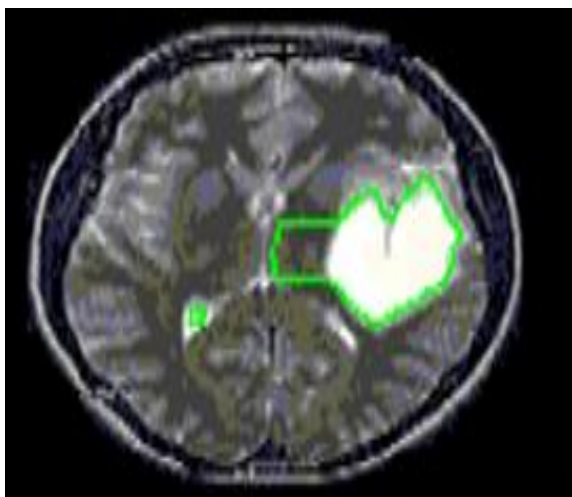
200 iterations:



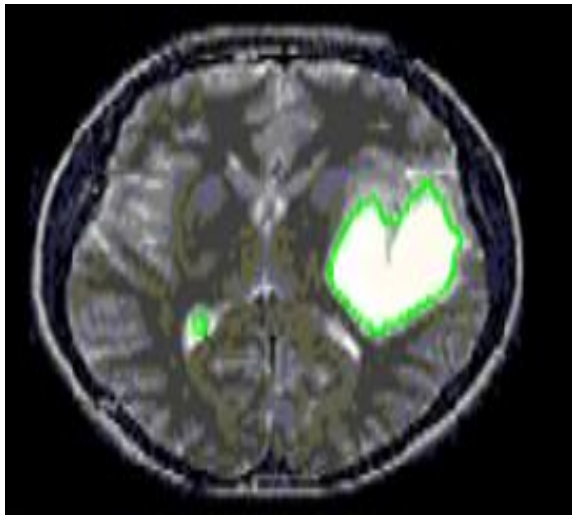
300 iterations:



400 iterations:



500 iterations:



617 iterations:

Tumor Detection

