CZ4003 COMPUTER VISION - 2020

Lab 2: Edge Detection + Hough Transform + 3D Stereo + Spatial Pyramid Matching

by BL U-H

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1 Edge Detection (Part 3.1a,b,c,d,e)

1.1 Edge Detection: Sobel (Part 3.1a,b,c,d)

Sobel filter is an edge detector that works by determining the gradient ("the jump") of image intensity. Sobel filter edge detector works because edges typically has the largest increase from light to dark.

Sobel filter works by using a vertical and horizontal edge detector:

Vertical Gradient Filter (detect Horizontal edge)	Horizontal Gradient Filter (detect Verticle edge)			
$\begin{bmatrix} -1 & -2 & -1 \\ 0 & 0 & 0 \\ 1 & 2 & 1 \end{bmatrix}$	$\begin{bmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{bmatrix}$			
Results: G_y =Magnitude of Verticle Gradient	Results: G_x =Magnitude of Horizontal Gradient			
Resultant Gradient Magnitude				
$G = \sqrt{ G_y ^2 + G_x ^2}$				
Resultant Gradient Direction				
$\theta = \tan^{-1}(\frac{G_y}{G_x})$				

After convolution with the filter separately, we will have a map of the magnitudes of verticle gradient (from Vertical Edge Filter) and horizontal gradient (from Horizontal Edge Filter). The resultant gradient magnitude and orientation of the gradient can be determined as shown in the formula above. [1]

1.1.1 Code (Part 3.1b,c)

1.1.2 Results (Part 3.1b, Part 3.1c)

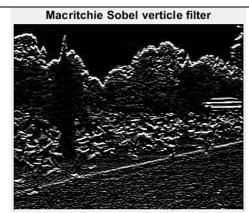


Figure 1: Convolution with Vertical Filer (Part 3.1b)

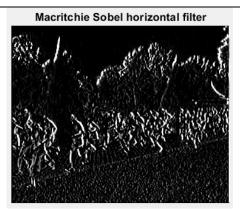


Figure 2:Convolution with Horizontal Filer (Part 3.1b)

Macritchie Sobel edge detector

Figure 3: Map of the Magnitude of the Resultant Gradient

1.1.2.1 (Part 3.1b) What happens to edges which are not strictly vertical nor horizontal, i.e. diagonal?

After convolution with the **verticle gradient filter**, the image is a map of the magnitudes of verticle gradient (Figure 1).

Horizontal lines appear brighter while perfectly vertical lines are dark. The intensity of the **resultant verticle gradient map** changes from brightest when the line is horizontal to darkest when the line is verticle or if there is no change intensity of the original map. Edges that are diagonal appear less bright than a horizontal line.

After convolution with the **horizontal gradient filter**, the image is a map of the magnitudes of horizontal gradient (Figure 2).

Vertical lines appear brighter while perfectly horizontal lines are dark. The intensity of the **resultant verticle gradient map** changes from brightest when the line is Vertical to darkest when the line is horizontal or if there is no change intensity of the original map. Edges that are diagonal appear less bright than a Verticle line.

In the final resultant gradient magnitude map (Figure 3), all edges are detected both vertical and horizontal. Edges that are diagonal are also detected.

1.1.2.2 (Part 3.1c) Suggest a reason why a squaring operation is carried out.

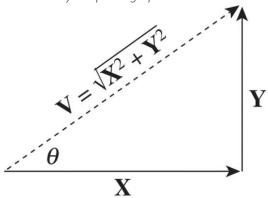


Figure 4: Resultant Vectors derived from horizontal and vertical components

The Squaring operation is carried and summed to obtain the resultant magnitude of the gradient. Because convolution with the **horizontal gradient filter** give the horizontal gradient vector. Convolution with the **verticle gradient filter** gives the verticle gradient vector. Therefore, we can use Pythagorean theorem to determine the resultant gradient (Figure 4).

1.1.2.3 (Part 3.1d) Vary Threshold: Try different threshold values and display the binary edge images.

1.1.2.3.1 Code

1.1.2.3.1 Code	
<pre>% 3.1 d) Thresholding min_P2=double(min(P2(:)))%Min intensity=13 max_P2=double(max(P2(:)))%Max intensity=204 %contrast stretching P2C = (double(P2(:,:))-min_P2).*(255/(max_P2-min_P2)); %checking min max of P2 min(P2C(:)), max(P2C(:)) %0, 255</pre>	Contrast stretch to make sure resultant gradient magnitude map is between 0 to 255
t=OTSU_B(P2C,true); P2t=P2C>t; [count,bins]=imhist(uint8(P2C),256);	Use OTSU to determine optimal threshold with minimal intra class variance/ maximum interclass variance Refer to Appendix OTSU_B.m.
<pre>figure('Position', [10 10 900 600]); colormap('gray');imshow(P2t);title("Macritchie Sobel edge detector") hold off; t_varry=linspace(10,220,8) t_varry=[t_varry(1:2),t,t_varry(3:7)] figure('Position', [10 10 1440 960]); for i = 1 : size(t_varry,2)</pre>	Plot out resulting sober filter after using different level of thresholding for binarization
<pre>subplot(2, size(sigma_varry,2)/2,i);colormap('gray');imshow(P2 t); if i==3</pre>	

1.1.2.3.2 Results

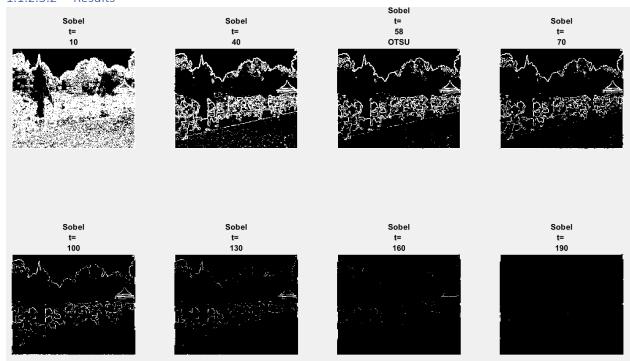


Figure 5: Sobel with different threshold

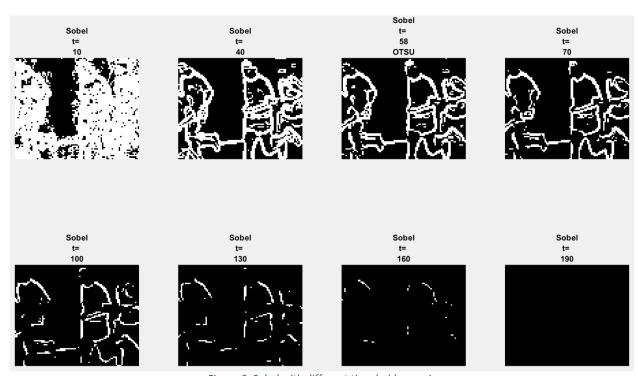


Figure 6: Sobel with different threshold zoom in

1.1.2.4 (Part 3.1 d)What are the advantages and disadvantages of using different thresholds?

Thresholding will help us to binarize the gradient map and help us identify likely edges. At lower threshold we obtain thicker lines (Figure 5, t=10) (Figure 6, t=10) while at higher thresholds we get thinner lines, and more lines is not detected (Figure 5, t=190) (Figure 6, t=190). To get the 'optimal' thresholding OTSU thresholding (Figure 5, t=58) (Figure 6, t=58). is implemented as a function

1.2 Edge Detection: Canny edge detector (Part 3.1 e i ii)

1.2.1 Code (Part 3.1e)

```
% 3.1 e)
tl=0.04, th=0.1, sigma=1.0;

E = edge(P,'canny',[tl th],sigma);
figure;colormap('gray');imshow(E);title("Macritchie Sobel edge detector")
```

1.2.2 Results (Part 3.1e)

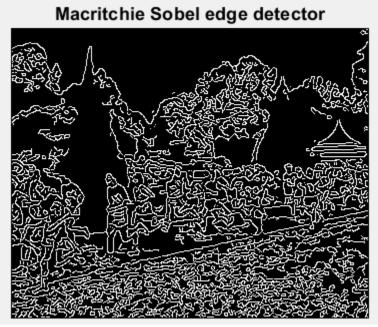


Figure 7: Canny edge Detector tl = 0.04, th = 0.1, $\sigma = 1.0$;

1.2.3 Vary σ : Code (Part 3.1ei)

```
% 3.1 e)i) Varry Sigma
sigma_varry=linspace(1,5,5);
figure( 'Position', [10 10 1440 960]);
for i = 1 : size(sigma_varry,2)
    E = edge(P,'canny',[t1 th],sigma_varry(i));
    hold on
    subplot(2,3,i);colormap('gray');imshow(E);title(["Canny" "\sigma= " sigma_varry(i)])
end
subplot(2,3,6);colormap('gray');imshow(P);title(["Original"])
hold off
```

1.2.4 Vary σ : Results (Part 3.1ei)

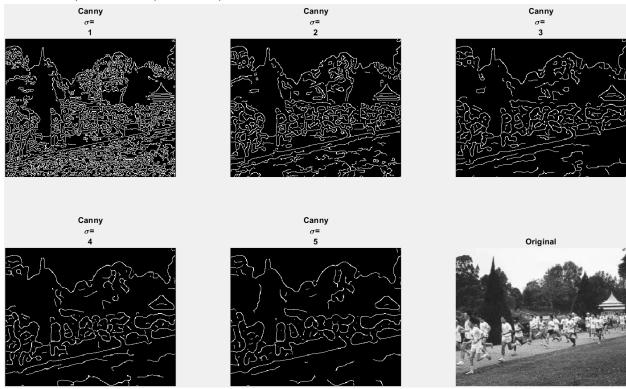


Figure 8: Canny Edge Detector varying σ

1.2.4.1 (Part 3.1 e i) What do you see and can you give an explanation for why this occurs? Discuss how different sigma are suitable for (a) noisy edgel removal, and (b) location accuracy of edgels.

As σ increases from 1 to 5 the sensitivity towards the edgel decreases. In other words, less edgel is detected. In terms of noise, less noisy edgel is detected. The location accuracy of the edgel also decreases. The reason for this is because gaussian edge filtering is able to remove noise. As, σ increases more noise is removed(less noisy edgel is detected) but it also means more details such as edges is lost (location accuracy of the edgel also decreases).

If the original image is noisy and a lot of noisy edgel is detected, a larger sigma should be used to remove noisy edgel.

If the original image is not noisy and we want to detect more edges with higher location accuracy, a σ of 1 should be sufficient.

1.2.5 Vary *tl*: Code (Part 3.1eii)

```
% 3.1 e)i)Varry tl
tl_varry=linspace(0.01,0.09,8);
figure( 'Position', [10 10 1440 960]);
for i = 1 : size(tl_varry,2)
    E = edge(P,'canny',[tl_varry(i) th],sigma);
    hold on
    subplot(2,size(tl_varry,2)/2,i);colormap('gray');imshow(E);title(["Canny" "th= "tl_varry(i)])
end
hold off
```

1.2.6 Vary *tl*: Results (Part 3.1eii)

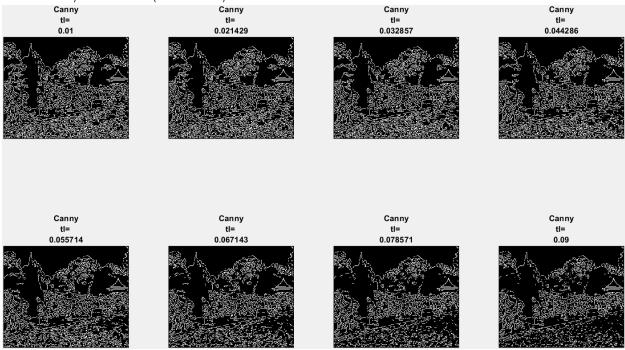


Figure 9: Canny Edge Detector varying tl

1.2.6.1 (Part 3.1 e ii) Try raising and lowering the value of tl. What does this do? How does this relate to your knowledge of the Canny algorithm?

Raising the tl decreases the number of edges detected (Figure 9)

In Canny's edge detector, hysteresis thresholding is employed.

$$f_t = \begin{cases} 1, & f > t_H \\ 1, & t_L < f < t_H \text{ and Condition} \\ 0, & f < t_L \end{cases}$$

- If a pixel is bellow tl, the pixel is set to 0.
- If a pixel is above tl and bellow th, the pixel is set to 1 if (Condition) its neighboring pixel perpendicular to the edge gradient have already been set to 1.
- If a pixel is above th, the pixel is set to 0.

Therefore, increasing tl increases the threshold for edge detection and thus lesser edge is detected and lesser noisy edge is detected as well.

2 Line Finding using Hough Transform (Part 3.2)

2.1 (Part 3.2 b) Read the help manual on Radon transform and explain why the transforms are equivalent in this case. When are they different? [2, 3, 4]

Radon Transform and Hough transform are mathematically similar

Radon Transform

The Radon Projection is computed for a set of θ angle. "In general, the Radon transform of f(x,y) is the line integral of f parallel to the y'-axis." [2]

$$R\theta(x') = \int_{\infty}^{\infty} f(x'\cos\theta - y'\sin\theta, x'\sin\theta + y'\cos\theta)dy'$$

$$where, \begin{bmatrix} x' \\ y' \end{bmatrix} = \begin{bmatrix} \cos\theta & \sin\theta \\ -\sin\theta & \cos\theta \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix}$$

Geometry of the Radon Transform

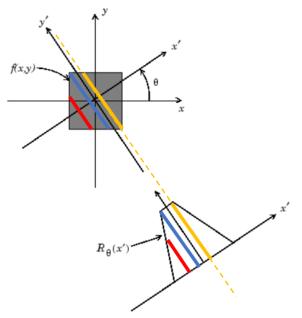


Figure 10: Geometry for Radon Transform

In other words, Radon Projection is like the **number of votes for a specific line represented in the parameter space in Hough Transform**.

"The Radon transform of an image is the sum of the Radon transforms of each individual pixel." [2] The magnitude of radon projection $R\theta(x')$ tell us the number of points that lie on the line perpendicular to x'

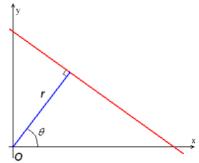
For instance, from Figure 10, "yellow amount" of points lie on the yellow dotted line which is a line perpendicular to x'.

It is important to note that

Hough transform

 $\rho = x\cos\theta + y\sin\theta$

"where ρ is the distance from the origin to the closest point on the straight line, and θ (theta) is the angle between the x axis and the line connecting the origin with that closest point." [5]



The fundamental idea is that any point (x, y) can be described using a set of (ρ, θ) that describes a set of lines that intersects the point (x, y).

"Given a single point in the plane, then the set of all straight lines going through that point corresponds to a sinusoidal curve in the (ρ, θ) plane, which is unique to that point. A set of two or more points that form a straight line will produce sinusoids which cross at that specific (ρ, θ) for that line. Thus, the problem of detecting collinear points can be converted to the problem of finding concurrent curves." [5]

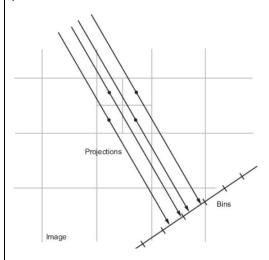
 θ correspond to θ in Hough transform x' correspond to ρ in Hough transform $R\theta(x')$ correspond to

number of votes for a specific line represented in the parameter space in Hough transform

Algorithm

Step 1: The Radon transform is computed for a set of θ angle.

For each θ angle, the Radon transform of an image is the sum of the Radon transforms of each individual pixel.



The algorithm first divides pixels in the image into four subpixels and projects each subpixel separately, as shown in the following figure.

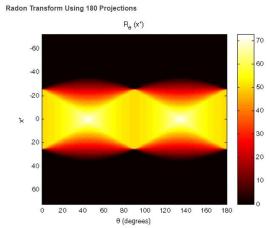


Figure 11: Radon projection magnitude map in a theta and x' array

Algorithm

The linear Hough transform algorithm uses a two-dimensional array, called an accumulator, to detect the existence of a line described by $\rho = x\cos\theta + y\sin\theta$

The dimension of the accumulator equals the two quantized values of ρ and θ in the pair (ρ, θ) .

Step 1: For each pixel at (x,y) the Hough Algorithm determines if there is enough evidence of a straight line at that pixel.

Step 2: If there is a line, it will calculate the parameters (ρ, θ) of lines that pass through that pixel and then increment the value of that corresponding bin in the accumulator

Step 3: Highest Values in the accumulator bin likely indicates a line can be extracted. The (approximate) geometric definition of that line in the x,y space can be derived from (ρ, θ) and $\rho = x cos \theta + y sin \theta$

Due to **imperfection errors** in the edge detection step **(Step 2)**, there will usually be errors in the accumulator space, which may make it not easy to find the appropriate peaks, and thus the appropriate lines.

"The final result of the linear Hough transform is a two-dimensional array (matrix) similar to the accumulator—one dimension of this matrix is the quantized angle θ and the other dimension is the quantized distance ρ .

Each element of the matrix has a value equal to the sum of the points or pixels that

Step 2: Radon projection magnitude can be map in a theta and x' array [2]

are positioned on the line represented by quantized parameters (ρ, θ) . So the element with the highest value indicates the straight line that is most represented in the input image." [5]

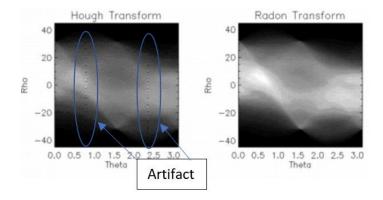
From this we see that magnitude in the parameter space is derived continuously (implemented discretely) using integration for Radon while, magnitude in the parameter space is derived discretely when using Hough Transform.

The difference in the Algorithm approach of mapping from the image space to a parameter space of ρ and θ is in their point of view.

- While the Radon transform derives the magnitude of a point in parameter space from image space (the *reading paradigm*) by using the radon projection transform on the image space,
- the Hough transform explicitly maps data points from image space to parameter space (the *writing paradigm*) when a line is detected. Magnitude of a point in a parameter space is derived by the accumulation of mapped points.

"Because of this, it is clear by comparing their (discrete) algorithms. For each θ in parameter space

- Radon projects the image points on a line described by (its angle) θ using buckets of size Δρ.
- Hough on the other hand takes each image point (x,y) and adds the appropriate intensity to all corresponding parameter space bins.



As a result Hough suffers artifacts whereas Radon allows for high resolution in parameter space (by adjusting $\Delta\theta$ and $\Delta\rho$ and dividing pixels into subpixels)." [4]

Therefore, in this case the transform is essentially equivilant because X_p is discretized by a unit of 1. They will differ slightly if X_p is discretized to smaller units.

(Note at the background, X_p is discretized to smaller units but the radon Transform is presented with X_p discretized to a unit of 1).

2.2 Radon transform Code (Part 3.2 b and c)

```
%3.2 b
theta = 0:180;
[H, xp] = radon(E, theta);
figure( 'Position', [10 10 900 350]);
subplot(1,2,1);imagesc(theta,xp,H),colormap(gca), colorbar;
title('R_{\theta} (X\prime)');
xlabel('\theta (degrees)')
ylabel('x''')
hold on;
%3.2 c
H_max = max(H,[],'all')
[y,x] = find(H == H max)
hold off;
zoom=3
subplot(1,2,2);imagesc(theta(x-zoom:x+zoom),xp(y-zoom:y+zoom),H(y-zoom:y+zoom,x-zoom:x+zoom))
colormap(gca), colorbar
title({'Max R {\theta} (X\prime) = num2str(H max),strcat("@ \theta= ", num2str(theta(x)), "
and xp = ", num2str(xp(y)));
xlabel('\theta (degrees)')
ylabel('x''')
hold on;
plot(theta(x), xp(y), 'o',...
             'MarkerEdgeColor','red',...
             'MarkerFaceColor',[1 .6 .6])
```

2.3 Radon transform Results (Part 3.2 b and c)

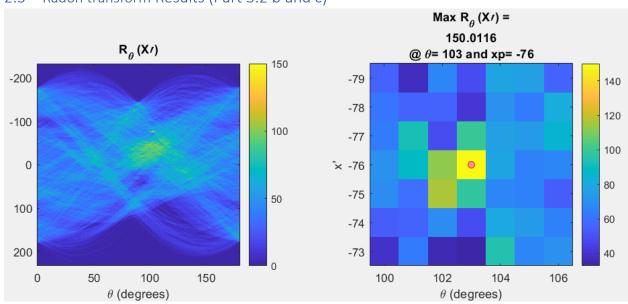


Figure 12: Results from Radon Transform (left) max intensity (right)

2.3.1 (Part 3.2 c) Find the location of the maximum pixel intensity in the Hough image in the form of [theta, radius].

From the graph Figure 12, the maximum pixel intensity is 150.0116 at $\theta = 103$ and xp = -76.

2.4 (Part 3.2 d) Derive the equations to convert the [theta, radius] line representation to the normal line equation form Ax + By = C in image coordinates.

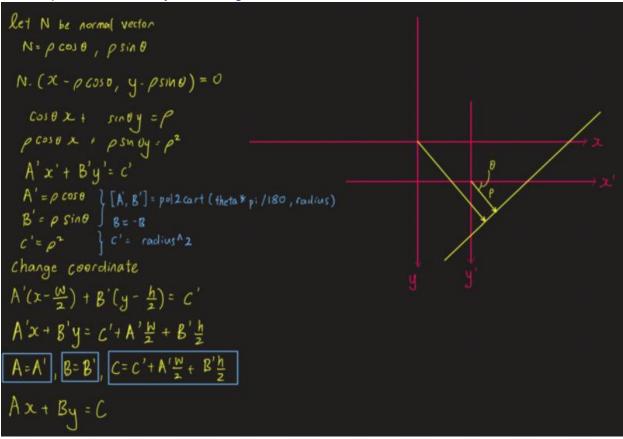
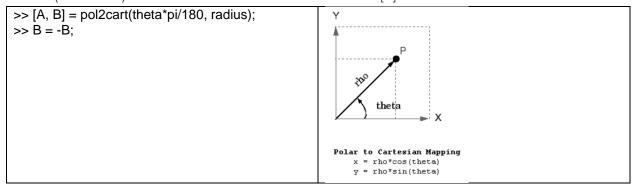


Figure 13: Derive the equations to convert the [theta, radius] line representation to the normal line equation form Ax + By = C in image coordinates.

2.4.1 (Part 3.2 d) Show that A and B can be obtained via [6]



Since pol2cart(theta, radius) converts polar coordinates (ρ, θ) to cartesian coordinate (x, y) and it is shown in Figure 13 that A and B is the vector coordinates of P, therefore A and B can be obtained via the code above.

2.4.2 Code: Line in Image plane (Part 3.2 d, e, f)

```
[A, B] = pol2cart(theta(x)*pi/180, xp(y));
B=-B;
C=xp(y)^2+A*size(P,2)/2+B*size(P,1)/2;
y_line=@(x) -A/B.*x+C/B;
xl = 0,xr = size(P,2) - 1;
yl=y_line(xl)
yr=y_line(xr)

figure;colormap('gray');imshow(uint8(P));title("Macritchie (Finding Line)")
line([xl xr], [yl yr],'LineWidth',2);
```

2.4.3 Results: Line in Image plane (Part 3.2 d, e, f)





Figure 14: Line in Image Plane

2.4.3.1 (Part 3.2 f) Results: Does the line match up with the edge of the running path? What are, if any, sources of errors? Can you suggest ways of improving the estimation?

Yes. it matches up almost perfectly with the edge of the running path. But on closer inspection there is some deviation from the line on the right-hand side of the Image. This is likely because the running path is not a perfect straight line and therefore does will not match up with the perfect line.

The possible sources of error and suggested improvements are

- The use of Radon to identify straight line introduced error in the detection of the straight line and this can be avoided by using a Generalized Hough Transform that can be used to detect curves
- 2. Another source of error may come from the discretization of the X_p . This can be improved by using Radon transform with Xp discretized to smaller units rather than Hough transform.

3 3D Stereo (Part 3.2)

3.1 Disparity Map Function (Part 3.2a)

$$S(x,y) = \sum_{j=0}^{M} \sum_{k=0}^{N} I^{2}(x+j,y+k) \cdot 1 + \sum_{j=0}^{M} \sum_{k=0}^{N} T^{2}(j,k) - 2\sum_{j=0}^{M} \sum_{k=0}^{N} I(x+j,y+k) T(j,k)$$

```
function map = disparity map Barn(Pl, Pr, tempx,tempy)
[h, w] = size(Pl);
% Empty Map
map = ones(h, w);
%Half the size of template
half tx=floor(tempx/2);
half_ty=floor(tempy/2);
max disparity=20;
for row = half tx+1:h-half tx
    for xl = half ty+1:w-half ty
        T = Pl(row-half_ty:row+half_ty,xl-half_tx:xl+half_tx);
        %Since right point image will be more left than left point image
        left = xl-max disparity;
        right = xl;
        if left<half tx+1</pre>
            left = half tx+1;
        end
        ssd min = Inf;
        xr \overline{min} = left;
        for xr = left:right
            I = Pr(row-half ty:row+half ty,xr-half tx:xr+half tx);
            ssd=sum(double(I).*double(I),"All")-2*sum(double(I).*double(T),"All");
            if ssd<ssd min
                ssd min=ssd;
                xr_min = xr;
            end
        end
        d = xl - xr min;
        map(row, xl) = -d;
map=map(half tx+1:h-half tx, half ty+1:w-half ty);
```

Function parameter

- PI: gray scale left stereo image
- Pr: gray scale right stereo image
- tempx: width of template (left image)
- tempy: height of template (right image)

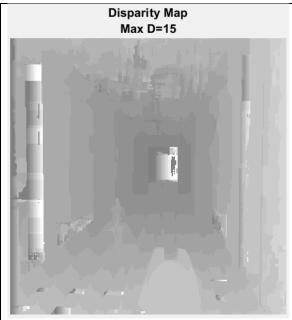
Avoided additional computation from fft and conv2 by using sum(, "All") and element wise multiplication. Didn't compute middle term because the middle term is a constant.

3.2 Disparity Map Function on 'corridorl.jpg' and 'corridorr.jpg' (Part 3.2b, c, d)

3.2.1 Code

```
%3.3 3D stereo
clear all;
%3.3 b
Pl = rgb2gray(imread('resource\corridorl.jpg'));
Pr = rgb2gray(imread('resource\corridorr.jpg'));
temp_size=11;
half=floor(temp_size/2);
whos Pl
whos Pr
figure;
subplot(1,2,1);colormap('gray');imshow(uint8(Pl));title("corridor left")
subplot(1,2,2);colormap('gray');imshow(uint8(Pr));title("corridor right")
%3.3 c
map=disparity_map_Barn(Pl,Pr,11,11);
figure;imshow(-map,[-15 15]);title({"Disparity Map","Max D=15"})
```

3.2.2 Results (Part 3.2 c)





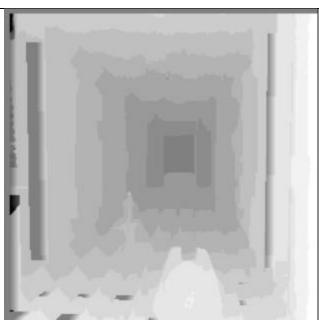


Figure 16: corridor_disp.jpg



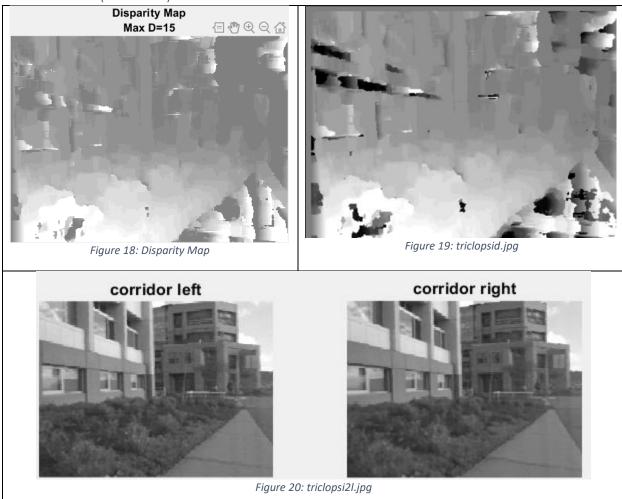
Figure 17: corridorl.jpg

3.2.3 (Part 3.2 c) Comment on how the quality of the disparities computed varies with the corresponding local image structure.

Generally, as we move further away from the corridor, the disparity map becomes darker. This corresponds to reality because the distance is further away from the lens and thus disparity is smaller. The cone and sphere can be identified with the sphere being lighter and having a larger disparity than the cone. For the furthest most wall, because it seems the most homogeneous with even lighting, there seems to be some error in the where there is large disparity detected. The disparity map derived from SSD works best when there are more details or texture on the object.

3.3 Rerun your algorithm on the real images of 'triclops-i2l.jpg' and triclops-i2r.jpg'.

3.3.1 Results (Part 3.2 d)



3.3.2 How does the image structure of the stereo images affect the accuracy of the estimated disparities?

Generally, the results show that objects closer to the lens have larger disparity and thus appear brighter while objects further away from the lens have smaller disparity and thus appear darker. There pavement does not seem to be properly mapped. One possible reason for this because we constrain the algorithm to only horizontal search. As you can see, there is **some rotation** between the left and the right image. Another possible reason for the poor disparity map of the pavement is due to the lack of very distinguishable features or edges that can help improve the robustness of the SSD this is in contranst to the vegetation on the left that seems to have a proper disparity map.

Also due to minimum disparity limit due to pixel size, the any object that is very far away will not be differentiated on the disparity map.

3.4 Optional Bag of Words and Spatial Pyramid Matching

The section will reference the original article on SPM [7] and reference code from [8] to implement BOW and SPM.

3.4.1 Summary

Algorithm	BOW+ KNN	BOW+SVM	SPM (L=2) +SVM
Accuracy	60%	67%	67-68%

Refer to the sections bellow for proof of results

- Results Bag of SIFT Representation + (KNN) Nearest Neighbor Classifier
- Results Bag of SIFT Representation + one-vs-all SVMs
- Results for Spatial Pyramid Matching + one-vs-all SVMs

3.4.2 Algorithm

3.4.2.1 Bag of words

"Bag of words models are a popular technique for image classification inspired by models used in natural language processing. The model ignores or downplays word arrangement (spatial information in the image) and classifies based on a histogram of the frequency of visual words. The visual word "vocabulary" is established by clustering a large corpus of local features. See Szeliski chapter 14.4.1 for more details on category recognition with quantized features. In addition, 14.3.2 discusses vocabulary creation and 14.1 covers classification techniques." [8]

Commonly in bag of words, huge data images is first vectorized using **SIFT. The vocabulary (features)** is extracted using K means clustering of the SIFT. Each K means represents a feature. Note the number of K means is selected by the programmer. In the code bellow we study how K means affect the accuracy of prediction. (Typically larger K, better accuracy because there is more features for prediction)

Using the set of K means/ features, feature histogram is formed for each image.

Using label training data set feature histogram and their corresponding label, a model can be trained either using KNN(K nearest neighbour) or SVM(Support vector Machine). Using the Trained Model, a prediction can be made on any input image. Note Typically (SVM models perform better than KNN)

3.4.2.2 Spatial Pyramid Matching

SPM builds on the same concepts of back of words by considering location of features. This is different from Bag of Words that sole depend on the histogram of features. In SPM, the same image is further sub divided into smaller images (Level 0=1 image, Level 1=4 image, Level 2=16 image, total 21 images). Each image is vectorized using SIFT and using K mean clustering, histogram features is formed for them. The histogram of features for the different level is then appended in a specific order.

3.4.3 CODE

Note

- Code was fully extracted from [8] with minor edits to reproduce accuracy of SPM for different levels L=0, L=1, L=2.
- Data was taken from [9] (source Caltect 101 [10])
- Explanations of flow of code for both BOW and SP and evaluations were added by me.

In the following code

- BOW was implemented with KNN(K nearest neighbour) and SVM(Support vector Machine)
 - o (Not related of KNN) Varying number of K-means clustering/ feature selected
 - Varying C in SVM to find optimal parameter for SVM
- Spatial Pyramid Matching was implemented with SVM(Support vector Machine)

Source	: [8]	-	itions of flow of code for both BOW and evaluations by me
1. 2. 3. 4. 5. 6. 7. 8. 9. 10.	<pre># import packages here import cv2 import numpy as np import matplotlib.pyplot as plt import glob #from scipy.misc import imresize # resize images import copy print('OpenCv Version:',cv2version) class_names = [name[11:] for name in glob.glob('data/train/*')] class_names = dict(zip(range(0,len(class_names)), class_names)) print (class_names)</pre>		,
1.	***************************************	This se	ction load data
2. 3. 4. 5.	<pre>####################################</pre>		train data, train label test data, test label
7. 8. 9.	<pre>if num_per_class > 0: img_path_class = img_path_class[:num_per_class] labels.extend([id]*len(img_path_class)) for filename in img_path_class:</pre>		

```
15. train_data, train_label = load_dataset('data/train/', 100)
    train_num = len(train_label)
17.
    print (train num)
    # load testing dataset
19. test_data, test_label = load_dataset('data/test/', 100)
20. test_num = len(test_label)
    ###############
    from sklearn.neighbors import KNeighborsClassifier
2.
3.
                                                              1.
    # train model
4.
5. def trainKNN(data, labels, k):
        neigh = KNeighborsClassifier(n_neighbors=k, p=2)
6.
7.
        neigh.fit(data, labels)
        return neigh
8.
9.
    10.
    #Bag of SIFT Representation + Nearest Neighbor Classifer
11.
12.
    from sklearn.cluster import KMeans
13. from sklearn import preprocessing
14.
15. # compute dense SIFT
16.
    def computeSIFT(data):
17.
     x = []
18.
        for i in range(0, len(data)):
           sift = cv2.xfeatures2d.SIFT_create()
19.
            img = data[i]
21.
            step size = 15
            kp = [cv2.KeyPoint(x, y, step_size) for x in range(0,
     img.shape[0], step_size) for y in range(0, img.shape[1], step_
23.
            dense_feat = sift.compute(img, kp)
            x.append(dense_feat[1])
25.
26.
        return x
27.
28. # extract dense sift features from training images
29. x_train = computeSIFT(train_data)
30. x_test = computeSIFT(test_data)
31.
32. all_train_desc = []
33. for i in range(len(x_train)):
34.
        for j in range(x_train[i].shape[0]):
     all_train_desc.append(x_train[i][j,:])
35.
36.
37. all_train_desc = np.array(all_train_desc)
##############
39. # build BoW presentation from SIFT of training images
40. def clusterFeatures(all train desc. k):
41.
        kmeans = KMeans(n_clusters=k, random_state=0).fit(all_trai
    n desc)
       return kmeans
```

data.append(cv2.imread(filename, 0))

12.

13.

return data, labels

load training dataset

3.4.3.1 Bag of SIFT Representation + Nearest Neighbor Classifier

- Vectorise Image by converting image to dense SIFT
 - computeSIFT(data)
 - return SHIFT data x
- build BoW presentation from SIFT of training images
 - clusterFeatures(all_train_desc, k)
 - return k means
 - K means clustering to for k number of means (k features) from SIFT
- form training set histograms for each training image using BoW representation
 - def formTrainingSetHistogram(x_train, kmeans, k):
 - return histogram of features
 - using SIFT predict closest k mean and fill up feature histogram
- build histograms for test set and predict using KNN
 - def predictKMeans(kmeans, scaler, x_test, train_hist, train_label, k):
 - return test set prediction
 - Use KNN(K nearest neighbours) different from K mean clustering.
 - k-NN is a supervised algorithm used for classification
 - For each evaluation matrix, Decide upon the value of k. Here k refers to the number of closest neighbors we will consider while doing the majority voting of target labels.
 - Run k-NN a few times, changing k and checking the evaluation measure.
 - In each iteration, k neighbors vote, majority vote wins and becomes the ultimate prediction
 - Optimize k by picking the one with the best evaluation measure.
 - Once you've chosen k, use the same training set and now create a new test set with the people's ages and incomes that you have no labels for, and want to predict.
 - Train KNN using train_hist and train_label
 - Use trained KNN to predict test_hist's label
- . evaluate accuracy of predict test_hist's label

```
44. # form training set histograms for each training image using B
45. def formTrainingSetHistogram(x_train, kmeans, k):
        train_hist = []
     for i in range(len(x_train)):
47.
48.
            data = copy.deepcopy(x_train[i])
         predict = kmeans.predict(data)
49.
50.
           train_hist.append(np.bincount(predict, minlength=k).re
     shape(1,-1).ravel())
51.
52.
        return np.array(train_hist)
53.
54. # build histograms for test set and predict
55. def predictKMeans(kmeans, scaler, x_test, train_hist, train_la
       # form histograms for test set as test data
    test_hist = formTrainingSetHistogram(x_test, kmeans, k)
57.
    # make testing histograms zero mean and unit variance
59.
    test_hist = scaler.transform(test_hist)
60.
61.
62.
       # Train model using KNN
63. knn = trainKNN(train_hist, train_label, k)
64.
        predict = knn.predict(test_hist)
    return np.array([predict], dtype=np.array([test_label]).dt
65.
 ype)
66.
67.
68. def accuracy(predict_label, test_label):
69.
       return np.mean(np.array(predict_label.tolist()[0]) == np.a
 rray(test_label))
#############
71.
72. k = [10, 15, 20, 25, 30, 35, 40]
73. for i in range(len(k)):
74.
        kmeans = clusterFeatures(all_train_desc, k[i])
75.
        train_hist = formTrainingSetHistogram(x_train, kmeans, k[i
 ])
76.
77. # preprocess training histograms
78.
        scaler = preprocessing.StandardScaler().fit(train hist)
79. train_hist = scaler.transform(train_hist)
80.
81. predict = predictKMeans(kmeans, scaler, x_test, train_hist
 , train_label, k[i])
        res = accuracy(predict, test_label)
82.
83. print("k =", k[i], ", Accuracy:", res*100, "%")
```

- def accuracy(predict_label, test_label):
 - return correct label/ total test label

```
    #Bag of SIFT Representation + one-vs-all SVMs
    from sklearn.svm import LinearSVC
    k = 60
    kmeans = clusterFeatures(all_train_desc, k)
    # form training and testing histograms
```

3.4.3.2 Bag of SIFT Representation + one-vsall SVMs

- (Same) Vectorise Image by converting image to dense SIFT
 - computeSIFT(data)
 - return SHIFT data x

```
train_hist = formTrainingSetHistogram(x_train, kmeans, k)
    test_hist = formTrainingSetHistogram(x_test, kmeans, k)
11.
    # normalize histograms
14. scaler = preprocessing.StandardScaler().fit(train_hist)
    train_hist = scaler.transform(train_hist)
    test_hist = scaler.transform(test_hist)
18. #Train one-vs-all SVMs using sklearn
19.
20. for c in np.arange(0.0001, 0.1, 0.00198):
21.
       clf = LinearSVC(random_state=0, C=c)
    clf.fit(train_hist, train_label)
22.
23.
        predict = clf.predict(test_hist)
        print ("C =", c, ",\t Accuracy:", np.mean(predict == test_
24.
    label)*100, "%")
##############
26. #We can train 15 SVM classifiers manually and get same result
27.
28. y_train_global = np.zeros((len(train_label), 1))
29. y = copy.deepcopy(y_train_global)
30.
31. y_predict = np.zeros((len(test_label), 1))
32. for i in range(len(test label)):
33.
       index = 0
    test = np.array([test_hist[i,:]]).T
34.
35.
        for j in range(len(class_names)):
       y = copy.deepcopy(y_train_global)
36.
           y[index:index+100, 0:1] = np.ones((100,1))
37.
           clf = LinearSVC(random_state=0, C=0.06148)
38.
           clf.fit(train_hist, y.ravel())
39.
           if j == 0:
40.
41.
              maxScore = np.dot(clf.coef_, test) + clf.intercept
42.
              y_predict[i, 0:1] = j
43.
           elif np.dot(clf.coef_, test) + clf.intercept_ > maxSco
44.
              maxScore = np.dot(clf.coef_, test) + clf.intercept
45.
               y_predict[i, 0:1] = j
46.
           index = index + 100
47. print ("Accuracy:", np.mean(y_predict.ravel() == test_label)*1
```

- (Same) build BoW presentation from SIFT of training images
 - clusterFeatures(all_train_desc, k)
 - return k means
 - K means clustering to for k number of means (k features) from SIFT
- (Same) form training and test set histograms for each training image using BoW representation
 - def formTrainingSetHistogram(x_train, kmeans, k):
 - return histogram of features
 - using SIFT predict closest k mean and fill up feature histogram
- 4. Use 1 v All SVM (Support Vector Machine) to train and predict
 - clf = LinearSVC(random_state=0, C=c)
 - clf.fit(train_hist, train_label)
 - predict = clf.predict(test_hist)
 - Large Value of parameter C => small margin
 - Small Value of parameter C => Large margin
 - offind optimal smallest c (largest margin) value for training data set so that misclassification of test data set is lower. In order words, it gerneralised the SVM model for other data sets other than training data set source: Medium
- (Same)evaluate accuracy of predict test_hist's label
 - def accuracy(predict_label, test_label):
 - return correct label/ total test label

rix'

3.4.3.3 Plot Confusion Matrix for Bag of SIFT Representation + one-vs-all SVMs

REFER TO RESULTS: "Results Bag of SIFT Representation + one-vs-all SVMs"

```
8.
                               cmap=plt.cm.Blues):
        ....
9.
        This function prints and plots the confusi
10.
    on matrix.
        Normalization can be applied by setting `n
11.
    ormalize=True`.
12.
        if normalize:
13.
            cm = cm.astype('float') / cm.sum(axis=
14.
    1)[:, np.newaxis]
15.
            #print("Normalized confusion matrix")
16.
17.
        #print(cm)
18.
19.
        plt.imshow(cm, interpolation='nearest', cm
    ap=cmap)
20.
        plt.title(title)
21.
        plt.colorbar()
22.
        tick_marks = np.arange(len(classes))
23.
        plt.xticks(tick marks, classes, rotation=4
    5)
24. plt.yticks(tick_marks, classes)
25.
26.
        fmt = '.2f' if normalize else 'd'
        thresh = cm.max() / 2.
27.
28.
        for i, j in itertools.product(range(cm.sha
    pe[0]), range(cm.shape[1])):
            plt.text(j, i, format(cm[i, j], fmt),
29.
30.
                     horizontalalignment="center",
                     color="white" if cm[i, j] > t
31.
    hresh else "black")
32.
33.
        plt.tight_layout()
        plt.ylabel('True label')
35.
        plt.xlabel('Predicted label')
36.
37. # Compute confusion matrix
38. cnf_matrix = confusion_matrix(np.array([test_l
    abel]).T, y_predict)
39. np.set_printoptions(precision=2)
41. # Plot non-normalized confusion matrix
42. plt.figure(figsize=(18, 6))
43. plot_confusion_matrix(cnf_matrix, classes=clas
44.
                          title='Confusion matrix,
     without normalization')
45.
46. # Plot normalized confusion matrix
47. plt.figure(figsize=(18, 6))
48. plot_confusion_matrix(cnf_matrix, classes=clas
    s_names, normalize=True,
49.
                          title='Normalized confus
    ion matrix')
50.
51. plt.show()
```

```
#Improve performance with Spatial Pyramid Matching
3.
4.
     def extract_denseSIFT(img):
5.
         DSIFT_STEP_SIZE = 2
6.
7.
         sift = cv2.xfeatures2d.SIFT_create()
8.
         disft_step_size = DSIFT_STEP_SIZE
         keypoints = [cv2.KeyPoint(x, y, disft_step_size)
10.
                 for y in range(0, img.shape[0], disft_step_size)
11.
                     for x in range(0, img.shape[1], disft_step_siz
     e)]
12.
13.
         descriptors = sift.compute(img, keypoints)[1]
14.
         #keypoints, descriptors = sift.detectAndCompute(gray, None
16.
         return descriptors
17.
18.
     # form histogram with Spatial Pyramid Matching upto level L wi
     th codebook kmeans and k codewords
     def getImageFeaturesSPM(L, img, kmeans, k):
21.
     W = img.shape[1]
22.
         H = img.shape[0]
     h = []
23.
24.
         for l in range(L+1):
25.
             w_step = math.floor(W/(2**1))
26.
             h_{step} = math.floor(H/(2**1))
27.
            x, y = 0, 0
28.
             for i in range(1,2**l + 1):
29.
               x = 0
30.
                 for j in range(1, 2**l + 1):
31.
                     desc = extract_denseSIFT(img[y:y+h_step, x:x+w
     _step])
32.
                     #print("type:",desc is None, "x:",x,"y:",y, "d
     esc_size:",desc is None)
33.
                     predict = kmeans.predict(desc)
                     histo = np.bincount(predict, minlength=k).resh
34.
     ape(1,-1).ravel()
35.
                     weight = 2**(1-L)
36.
                     h.append(weight*histo)
37.
                     x = x + w step
38.
                 y = y + h_step
39.
40.
         hist = np.array(h).ravel()
41.
      # normalize hist
42.
         dev = np.std(hist)
       hist -= np.mean(hist)
43.
44.
         hist /= dev
45.
       return hist
46.
47.
48. # get histogram representation for training/testing data
49. def getHistogramSPM(L, data, kmeans, k):
50.
         x = []
51.
        for i in range(len(data)):
             hist = getImageFeaturesSPM(L, data[i], kmeans, k)
             x.append(hist)
```

3.4.3.4 Improve performance with Spatial Pvramid Matchina

- (Same done) Vectorise Image by converting image to dense SIFT
 - computeSIFT(data)
 - return SHIFT data x
- (Same 200 features SHIFT data) build BoW presentation from SIFT of training images
 - clusterFeatures(all_train_desc, k)
 - return k means
 - K means clustering to for k number of means (k features) from SIFT
- (different) form training and test set feature histograms for each training image using BoW representation
 - getHistogramSPM(L, data, kmeans, k)
 - return SPM Histogram
 - return combined SPM Histogram
 † getImageFeaturesSPM(L, img,
 - kmeans, k)
 - return SPM Histogram for each all level of spaces in the spacial pyramid.
 - For L=0: 1 image, L=1: 4image,L=2: 16image
 - † extract_denseSIFT(img)
- 4. (same) **Use 1 v All SVM (Support Vector** *Machine) to train and predict*
 - clf = LinearSVC(random_state=0, C=c)
 - clf.fit(train_hist, train_label)
 - predict = clf.predict(test_hist)
 - Large Value of parameter C => small margin
 - Small Value of parameter C => Large margin
 - o find optimal smallest c (largest margin) value for training data set so that misclassification of test data set is lower. In order words, it gerneralised the SVM model for other data sets other than training data set source: Medium
- (Same) evaluate accuracy of predict test_hist's label
 - def accuracy(predict_label, test_label):
 - return correct label/ total test label

```
return np.array(x)
55.
58. kmeans = clusterFeatures(all_train_desc, k)
60. train_histo = getHistogramSPM(2, train_data, kmeans, k)
61. test_histo = getHistogramSPM(2, test_data, kmeans, k)
#############
63. # train SVM
64. for c in np.arange(0.000307, 0.001, 0.0000462):
65. clf = LinearSVC(random_state=0, C=c)
     clf.fit(train_histo, train_label)
67. predict = clf.predict(test_histo)
68. print ("C =", c, ",\t\t Accuracy:", np.mean(predict == tes
   t_label)*100, "%")
```

3.4.4 Results Bag of SIFT Representation + (KNN) Nearest Neighbor Classifier

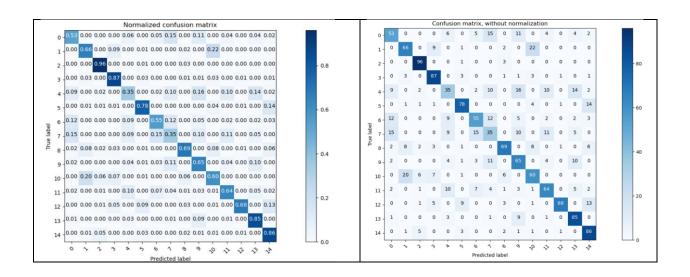
```
k = 10 , Accuracy: 52.8000000000000004 %
k = 15 , Accuracy: 51.933333333333333 %
k = 20 , Accuracy: 55.266666666666666 %
k = 25 , Accuracy: 55.600000000000000 %
k = 30 , Accuracy: 58.4 %
k = 35 , Accuracy: 58.86666666666667 %
k = 40 , Accuracy: 60.0 %
```

3.4.5 Results Bag of SIFT Representation + one-vs-all SVMs C = 0.00208, C = 0.00406, Accuracy: 63.0 % C = 0.00604, Accuracy: 63.4666666666666 % C = 0.00802, Accuracy: 63.933333333333333 % Accuracy: 64.4 % C = 0.01396 , Accuracy: 65.0666666666666 % C = 0.01594 , Accuracy: 65.4666666666666 % Accuracy: 65.8 % C = 0.01792, Accuracy: 65.933333333333334 % C = 0.02188 , Accuracy: 65.866666666666 % C = 0.02386 , Accuracy: 65.8666666666666 % Accuracy: 65.93333333333334 % C = 0.02584 , C = 0.02782, C = 0.0298 , C = 0.03178, Accuracy: 66.26666666666667 % Accuracy: 66.2 % C = 0.03574 , Accuracy: 66.4 % C = 0.0377200000000000004, Accuracy: 66.4 % C = 0.0397, C = 0.04168, Accuracy: 66.8 % C = 0.04564 , Accuracy: 66.866666666666 % C = 0.04762Accuracy: 66.933333333333334 % Accuracy: 66.93333333333334 % C = 0.05158 , Accuracy: 66.933333333333334 % C = 0.05356, Accuracy: 66.93333333333334 % Accuracy: 67.0 C = 0.05752 , Accuracy: 67.066666666666 % C = 0.0595000000000000000004, Accuracy: 66.8666666666666 C = 0.06148 , Accuracy: 66.8 % Accuracy: 66.8666666666666 Accuracy: 66.933333333333334 % C = 0.06544, C = 0.0674200000000001, Accuracy: 66.9333333333333 % C = 0.0694 , Accuracy: 66.93333333333333 % C = 0.07138, Accuracy: 67.0 % C = 0.07336000000000001, C = 0.07534, Accuracy: 67.0 % C = 0.07732, C = 0.0793 , C = 0.08128 , C = 0.08326 , Accuracy: 67.2 % C = 0.08524, Accuracy: 67.133333333333334 % C = 0.08722, Accuracy: 67.2 % C = 0.0892 , Accuracy: 67.2 % C = 0.09118, Accuracy: 67.2 % C = 0.09316 , Accuracy: 67.2 % C = 0.09514, C = 0.09712, C = 0.099100000000000000001,

```
{0: 'Bedroom', 1: 'Coast', 2: 'Forest', 3: 'Highway', 4: 'Indus trial', 5: 'InsideCity', 6: 'Kitchen', 7: 'LivingRoom', 8: 'Mou ntain', 9: 'Office', 10: 'OpenCountry', 11: 'Store', 12: 'Stree t', 13: 'Suburb', 14: 'TallBuilding'}
```

Accuracy: 66.8 %

0	Bedroom	8	Mountain
1	Coast	9	Office
2	Forest	10	OpenCountry
3	Highway	11	Store
4	Industrial	12	Street
5	InsideCity	13	Suburb
6	Kitchen	14	TallBuilding
7	LivingRoom		



3.4.5.1 Results for Spatial Pyramid Matching + one-vs-all SVMs

```
Accuracy: 67.933333333
Accuracy: 68.0 %
C = 0.000445599999999999999999
                         Accuracy: 67.933333333
C = 0.00049179999999999999999
                         Accuracy: 67.733333333
Accuracy: 67.666666666
C = 0.00058419999999999999999
                         Accuracy: 67.733333333
C = 0.00063039999999999999999
                         Accuracy: 67.800000000
C = 0.00067659999999999999999
                         Accuracy: 67.733333333
Accuracy: 67.466666666
Accuracy: 67.4 %
C = 0.00081519999999999999999
                          Accuracy: 67.26666666
66667 %
Accuracy: 67.4 %
Accuracy: 67.26666666
66667 %
Accuracy: 67.2 %
Accuracy: 67.333333333
33333 %
```

4 Appendix

4.1 OTSU B.m

```
function threshold =OTSU_B(gray, Analysis)
                                                               Maximize intraclass variance
%OTSU Thresholding
                                                                          \sigma^2 - \sigma_w^2 = W_L W_H (\mu_L + \mu_H)^2
%Maximise interclass variance
size(count)
inter class var=zeros(size(bins,1),1);
for n=1:size(bins,1)
    %Mean of L(backgnd) and H(foregnd)
mean L=dot(count(1:n),bins(1:n))/sum(count(1:n));
mean H=dot(count(n+1:256), bins(n+1:256))/sum(count(n))
+1:256));
    weight L=sum(count(1:n))/sum(count);
    weight H=sum(count(n+1:256))/sum(count);
    inter class var(n)=weight L*weight H*(mean L-
mean H)^2;
[M,I] = max(inter_class_var)
%Analysis Report
if Analysis ==true
                                                               Analysis produce
         %Plot Interclass variance
                                                                        interclass variance vs threshold
         figure( 'Position', [10 10 900 600]);
                                                                        plot and
         subplot(1,2,1);plot(bins,inter class var);
                                                                        image histogram with threshold
         plot(bins(I),inter_class_var(I),'o',...
                                                                        Inter class variance \sigma^2 \text{-} \sigma_{\text{ w}}^{2} \text{=W}_{\text{L}} \text{W}_{\text{H}} (\mu_{\text{L}} \text{+} \mu_{\text{H}})^2
              'MarkerEdgeColor', 'red',...
              'MarkerFaceColor',[1 .6 .6])
        title({'Inter class variance','{\sigma}^2-
{\sigma w}^2={W} L{W}_H({\mu _L}+{\mu _H})^2'}
        xlabel('Threshold, t');
        ylabel('Inter class variance, {\sigma}^2-
{\langle w \rangle^2 };
         %Plot histogram with OTSU threshold
subplot(1,2,2);imhist(uint8(gray),256);title("Histog
ram with OTSU Threshold");
        hold on;
        xline(bins(I),'-r',{'OTSU
Threshold',strcat('t= ',num2str(bins(I)))})
    end
threshold=bins(I)
```

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