Normalised DLT

```
Loading the image
```

```
kitchen1 = imread('kitchen2.jpg');
figure,imshow(kitchen1)
kitchen2 = imread('kitchen1.jpg');
figure,imshow(kitchen2)
```





Grayscaling and finding the points that match with each other from the two images.

```
graykitchen1 = rgb2gray(kitchen1);
graykitchen2 = rgb2gray(kitchen2);
points1 = detectSURFFeatures(graykitchen1)
points2 = detectSURFFeatures(graykitchen2)

[features1,valid_points1] = extractFeatures(graykitchen1,points1);
[features2,valid_points2] = extractFeatures(graykitchen2,points2);
indexPairs = matchFeatures(features1,features2,'MaxRatio',0.1);

%disp(indexPairs)
matchedPoints1 = valid_points1(indexPairs(:,1),:);
```

```
%disp(matchedPoints1)
matchedPoints2 = valid_points2(indexPairs(:,2),:);
%disp(matchedPoints2)
```

To get the minimum amount of points needed to do the homography (four points), I randomly selected 4 points from the first image, then I first transposed the points and normalized them with the algorithm given, obtaining normalisedFourPoints at the end.

```
fourPoints1 =
  3×4 single matrix
   1.0e+03 *
    2.4353 2.7005 2.8464
                                      2.3790
    2.9615 3.0575 2.8822
                                     2.8540
    0.0010 0.0010 0.0010 0.0010
normalisedFourPoints =
   -0.2711
              0.1928 0.4479 -0.3696
              0.2075 -0.0990 -0.1483
    0.0397
    1.0000
                        1.0000
                                   1.0000
              1.0000
      fourPoints1 = transpose(matchedPoints1([1 3 7 14],:).Location)
      fourPoints1(3,:) = 1;
      average = mean(fourPoints1,2)
      d1 = 0;
      for i = 1:size(fourPoints1,2)
        cal = sqrt((fourPoints1(1,i)-average(1,1)).^2+(fourPoints1(2,i)-average(2,1)).^2);
        d1 = d1 + cal;
      end
      T1 = [sqrt(2)/d1 \ 0 \ -sqrt(2)*average(1,1)/d1; 0 \ sqrt(2)/d1 \ -sqrt(2)*average(2,1)/d1; 0 \ 0 \ 1]
      normalisedFourPoints = zeros(size(fourPoints1))
      for i = 1:size(fourPoints1,2)
        temp = fourPoints1(:,i)
        temp = T1*temp
        normalisedFourPoints(:,i) = temp
      end
```

Doing the same thing to the **SAME** four points matched in the second image:

```
fourPoints2 = transpose(matchedPoints2([1 3 7 14],:).Location)
        fourPoints2(3,:) = 1;
        average2 = mean(fourPoints2,2)
        d2 = 0;
        for i = 1:size(fourPoints2,2)
          cal = sqrt((fourPoints2(1,i)-average2(1,1)).^2+(fourPoints1(2,i)-average2(2,1)).^2);
          d2 = d2 + cal:
        end
        T2 = [sqrt(2)/d2 \ 0 - sqrt(2)*average2(1,1)/d2; 0 \ sqrt(2)/d2 - sqrt(2)*average2(2,1)/d2; 0 \ 0 \ 1]
        normalisedFourPoints2 = zeros(size(fourPoints2))
        for i = 1:size(fourPoints2,2)
          temp = fourPoints2(:,i)
          temp = T2*temp
          normalisedFourPoints2(:,i) = temp
        end
After obtaining the normalisedPoints, I used the following function that I implemented called
calchomography to find the homography of the points.
        % Estimate the homography from a set of point correspondences.
        function H = calchomography(points1, points2)
          if size(points1, 2) < 4 | | size(points2, 2) < 4
            error('no 4 points passed');
          end
          A = [];
          for point = 1:size(points1, 2)
           x = points1(1, point);
           y = points1(2, point);
           u = points2(1, point);
```

```
v = points2(2, point);
          A = [A; ...
             -x -y -1 0 0 0 x * u y * u u; ...
             0 0 0 -x -y -1 x*v y*v v];
         end
         % Solve A * h = 0 for h.
         [U, S, V] = svd(A);
         h = V(:,end);
         % Reshape vectorised result back into matrix shape.
         H = reshape(h, 3, 3)';
         % Homogeneous normalisation.
         H = H . / H(3,3);
Calling the function above with H and denormalise it, then saving it as mymatrix.mat:
       H = calchomography(normalisedFourPoints,normalisedFourPoints2);
       H = inv(T2)*H*T1 %Denormalising
       disp(H)
       save ('mymatrix.mat', 'H');
```

To show that the homography we obtained indeed works, I have created a backwardWarping algorithm that allows me to warp the first image to the second image. I didn't include the codes in **week3matlab.m** but it is at **imageAlignment.m** since backward mapping isn't necessary, the result is below, notice the sinks in both images, they line up perfectly.

```
source = im2double(imread('kitchen2.jpg'));
source2 = im2double(imread('kitchen1.jpg'));
newLeft = zeros(size(source));
% The backward warping transformation (rotation + scale about an arbitrary point).
Matrix = load('mymatrix.mat');
M = Matrix.H;
for y = 1:size(newLeft, 1)
    for x = 1:size(newLeft, 2)
```

```
% Transform source pixel location (round to pixel grid).
    p = [x; y; 1];
    q = inv(M) * p;
    u = round(q(1) / q(3));
    v = round(q(2) / q(3));
   % Check if target pixel falls inside the image domain.
    if (u > 0 \&\& v > 0 \&\& u \le size(source, 2) \&\& v \le size(source, 1))
      % Sample the target pixel colour from the source pixel.
      newLeft(y, x, :) = source(v, u, :);
    end
  end
end
target = imfuse(source2,newLeft);
%overlay
subplot(141); imshow(source); title('Old Left');
subplot(142); imshow(newLeft); title('New Left');
subplot(143); imshow(source2); title('Right');
subplot(144); imshow(target); title('Stitch');
```

Results:

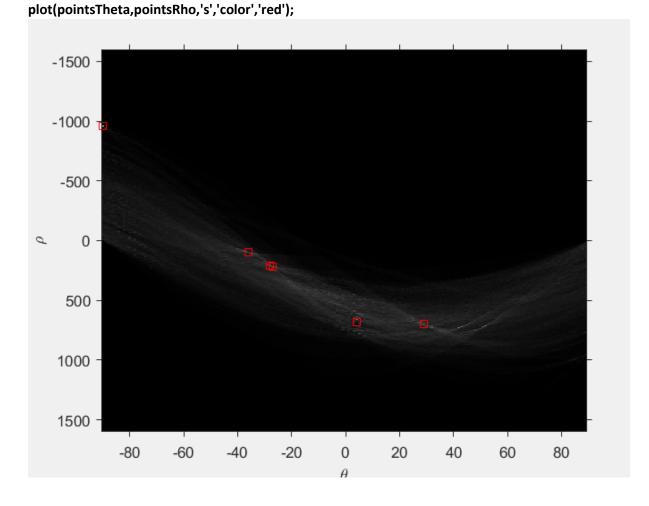


HoughLines detector

```
Reading and detecting the edge with matlab canny edge detector.
```

```
corridor = imread('Corridor1.jpg');
       grayCorridor = rgb2gray(corridor);
       Canny_img = edge(grayCorridor, 'Canny');
       [rows, cols] = size(Canny_img);
Initialising the hough space (Accumulator) to the size of theta_range and rho_range.
       theta_maximum = 90;
       rho_maximum = hypot(rows,cols) - 1;
       theta_range = -theta_maximum:theta_maximum-1;
       rho_range = -rho_maximum:rho_maximum;
       Hough = zeros(length(rho_range), length(theta_range));
Converting the edge found with r = x*cos(theta) + y*sin(theta).
       for j = 1:rows
         for i = 1:cols
           if Canny_img(j, i) ==1
             x = i - 1;
             y = j - 1;
             for T = theta_range
                R = round((x * cosd(T)) + (y * sind(T)));
                R_Index = R + rho_maximum + 1;
                T_Index = T + theta_maximum + 1;
                Hough(R_Index, T_Index) = Hough(R_Index, T_Index) + 1;
             end
           end
         end
       end
Hough peak with thresholding.
```

[X,Y] = find(Hough>0.45*max(Hough(:)))



Here is my attempt at dehoughing the peaks to image space. I first get the Rho and Theta pair into a list. I then created a set of binary matrices of the size imageRow x imageColumn x number of rhotheta pair. Then I substitute all the x and y pixel coordinates into the function xcos(theta) + ysin(theta) = rho, and if left - right == 0, I would mark the location of the binary matrix(x,y) as 1 and I run this for all the theta pairs against all the x and y value and store them separately in the set of binary matrices mentioned above.

In the second for loop, I just went through all of the set of binary matrices I had obtained and just aggregate them into one binary matrix.

```
for k = 1:length(pointsRho)
  for i = 1:1280
    for j = 1:960
      r = pointsRho(k);
      t = pointsTheta(k);
      threshold = abs(i*cosd(t)+j*sind(t)-r);
      if threshold < 0.5
         allLines(j,i,k) = 1;
      end
    end
  end
end
aggregatedImage = zeros(size(Canny_img));
for k = 1:length(pointsRho)
  for j = 1:1280
    for i = 1:960
      if allLines(i,j,k) ==1
         aggregatedImage(i,j) = 1;
      end
    end
  end
end
```

figure, imshow (imfuse (aggregated Image, corridor))

With houghpeaks = 0.45*max(Hough(:)) we get the following lines

