## CSC420 Assignment 3

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## 1

From wikipedia: Size of the 5 dollar canadian bill is height:  $69.85~\mathrm{mm}$  and width:  $152.4~\mathrm{mm}$ 

We round to  $700 \times 1524$  pixels, 1 per 1/10 of a mm.

From the output of q1, we can see that the bill is roughly 45% of the shoe in length. We can also so that it is roughly 80% of the shoe in width. Measuring the pixels, we can see that the estimated shoe size is roughly 26cm in length 10.5cm in width

Figure 1: Projection matlab code for q1



Figure 2: Output

2

(a + b)

```
affineX.m × RANSAC.m × +
            function [ransac_pts, ransac_affine] = affineX(i1, i2, ~)
4
5
6
7
8 -
9
                \label{eq:simgl} $$ \lim_{n\to\infty} 1 = \inf_{n\to\infty} (-h/u9/g6/00/changkao/csc420/assignments/Assignment2/book.jpg'); $$ \lim_{n\to\infty} 1 = \inf_{n\to\infty} (-h/u9/g6/00/changkao/csc420/assignments/Assignment2/findBook.JPG'); $$$ img2 = imread(-h/u9/g6/00/changkao/csc420/assignments/Assignment2/findBook.JPG'); $$$$ img2 = imread(-h/u9/g6/00/changkao/csc420/assignments/Assignment2/findBook.JPG'); $$$$$$$$
                threshold = 0.5;
                imgl = single(rgb2gray(i1));
11 -
                img2 = single(rgb2gray(i2));
12
13
14 -
15
               [f,d] = vl_sift(imgl) ;
15
16 -
17
18
               [f2, d2] = vl_sift(img2) ;
               e_dist= pdist2(double(d)', double(d2)', 'euclidean');
ascending_l = sort(e_dist, 2);
r=ascending_l(:,1)./ascending_l(:,2);
19 -
20 -
21 -
22
23 -
                t_index = find(threshold>r);
24 -
25
                num_matches = zeros(size(t_index,1), 3);
26
27 -
            for i = 1:size(e_dist, 1)
29 -
                        if threshold > r(i)
30
31 -
32 -
33 -
                               num_matches(i,1) = r(i);
num_matches(i,2) = i;
t_matches = (e_dist(i,:)==ascending_l(i,1));
num_matches(i,3)=find(t_matches);
34 -
35 -
36 -
37
               - end
38 -
39 -
40 -
41 -
42 -
               tesmp = any(num_matches, 2);
num_matches( ~tesmp, : ) = [];
mscores = zeros(size(num_matches,1), 1);
               x_coordinates = zeros(size(num_matches,1),2);
y_coordinates = zeros(size(num_matches,1),2);
44 -
45
46 -
47 -
48 -
            for i = 1:size(num_matches,1)
                        \label{eq:mscores} \begin{split} & \mathsf{mscores}(i) = \mathsf{num\_matches}(i,1); \\ & \mathsf{x\_coordinates}(i,1:2) = [\mathsf{f}(1,\mathsf{num\_matches}(i,2)) \ \mathsf{f}2(1,\mathsf{num\_matches}(i,3))]; \\ & \mathsf{y\_coordinates}(i,1:2) = [\mathsf{f}(2,\mathsf{num\_matches}(i,2)) \ \mathsf{f}2(2,\mathsf{num\_matches}(i,3))]; \end{split}
```

Figure 3: main function - RANSAC code matlab part 1

```
43
     for i = 1:size(num_matches,1)
44 -
45
           mscores(i) = num_matches(i,1);
46 -
           x_{coordinates(i,1:2)} = [f(1,num_matches(i,2)) f2(1,num_matches(i,3))];
47 -
            y_{coordinates(i,1:2)} = [f(2,num_matches(i,2)) f2(2,num_matches(i,3))];
48 -
49
50 -
       end
51
           k = size(mscores,1)
52 -
53
           %helper to get k top matches and their coordinates
54
            top = topmatches(mscores, x_coordinates, y_coordinates, k);
55 -
56 -
            trials = 100;
57 -
           num perm = 3;
58 -
            threshold2 = 2.21;
59
           %RANSAC ( matches, num of trials, num or pairs )
60
61 -
            [ransac_index, ransac_affine] = RANSAC(top, trials, num_perm, threshold2);
62 -
            ransac_pts = top(ransac_index,:);
63
64 -
       end
```

Figure 4: main function - RANSAC code matlab part 2

```
Q2bShredded.m
                     affine.m × RANSAC.m* × +
      □ function [indexs, max_affine] = RANSAC(top_pts, trials, num_perm, threshold)
 1
 2
 3 -
            indexs = [];
 4 -
            max_matches = 0;
 5
 6
            P = [];
 7 -
 8 -
            Pa = [];
            for i = 1:size(top_pts, 1);
 9 -
10
11 -
                x1 = top pts(i,2);
12 -
                yl = top_pts(i,4);
13
14
15 -
                x2 = top_pts(i,3);
16 -
                y2 = top_pts(i,5);
17
18 -
                P(size(P,1)+1,:) = [x1 y1 0 0 1 0];
19 -
                P(size(P,1)+1,:) = [0 0 x1 y1 0 1];
20
21 -
                Pa(size(Pa,1)+1,:) = x2;
22 -
                Pa(size(Pa,1)+1,:) = y2;
23 -
24
27 -
           while trials > 0
28
29 -
                random choices = randperm(size(top pts,1), num perm);
30 -
                random_pts = top_pts(random_choices,:);
31
                trials = trials-1;
32 -
33 -
                Rp = [];
34 -
                Rpa = [];
35
                for i = 1:3
36 -
37
                    % random points from img 1
38
39 -
                    x1 = random_pts(i,2);
40 -
                    y1 = random_pts(i,4);
41
42
                    % random points from img 2
43 -
                    x2 = random_pts(i,3);
44 -
                    y2 = random_pts(i,5);
45
                    Rp(size(Rp,1)+1,:) = [x1 y1 0 0 1 0];
46 -
47 -
                    Rp(size(Rp,1)+1,:) = [0 0 x1 y1 0 1];
48
                    Bpa(size(Rpa,1)+1,:) = x2;
49 -
50 -
                    Rpa(size(Rpa,1)+1,:) = y2;
```

Figure 5: RANSAC code matlab part 1

```
2T
                 end
52 -
53
54
                 %affine = P'*inv(P*P')*Pa;
                 random_affine = inv(Rp'*Rp)*Rp'*Rpa;
56 -
57
                 transform = P*random_affine;
58 -
59
                 T_x = transform(1:2:length(transform));
60 -
61 -
                 T_y = transform(2:2:length(transform));
62 -
                 Pa_x = Pa(1:2:length(Pa));
63 -
                 Pa_y = Pa(2:2:length(Pa));
64
                 diff_x = abs(Pa_x - T_x);
65 -
                 diff_y = abs(Pa_y - T_y);
diff_total = abs(diff_x+diff_y);
66 -
67 -
                 closest = find(diff_total < threshold);</pre>
68 -
69
70 -
                 if max_matches < size(closest, 1);</pre>
                     max_matches = size(closest,1);
71 -
72 -
                     max_affine = random_affine;
73 -
                     indexs = closest;
74 -
                 end
            end
75 -
76 -
      end
```

Figure 6: RANSAC code matlab part 2

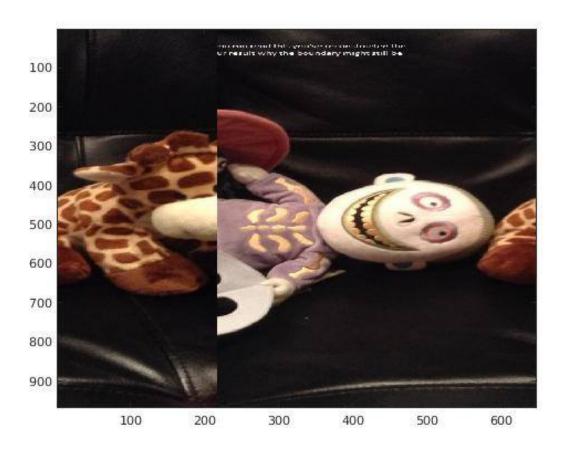


Figure 7: Output - 100 trials, 0.68 threshold