# CSE 252A Computer Vision I Fall 2021 - Assignment 2

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- Assignment Published On: Wed, Oct. 20, 2021.
- Due On: Wed, Nov. 3, 2021 11:59 PM (Pacific Time).

# **Instructions**

Please answer the questions below using Python in the attached Jupyter notebook and follow the guidelines below:

- This assignment must be completed individually. For more details, please follow the Academic Integrity Policy and Collaboration Policy on <u>Canvas (https://canvas.ucsd.edu/courses/29614)</u>.
- All the solutions must be written in this Jupyter notebook.
- After finishing the assignment in the notebook, please export the notebook as a PDF and submit both the notebook and the PDF (i.e. the . ipynb and the . pdf files) on Gradescope.
- You may use basic algebra packages (e.g. NumPy, SciPy, etc) but you are not allowed to use the packages that directly solve the problems. Feel free to ask the instructor and the teaching assistants if you are unsure about the packages to use.
- It is highly recommended that you begin working on this assignment early.

**Late Policy:** Assignments submitted late will receive a 15% grade reduction for each 12 hours late (i.e., 30% per day). Assignments will not be accepted 72 hours after the due date. If you require an extension (for personal reasons only) to a due date, you must request one as far in advance as possible. Extensions requested close to or after the due date will only be granted for clear emergencies or clearly unforeseeable circumstances.

# Problem 1 Image filtering [15 pts]

# **Problem 1.1 Implementing Convolution[5 pts]**

In this problem, you will implement the convolution filtering operation in NumPy.

As shown in the lecture, a convolution can be considered as a sliding window that computes a sum of the pixel values weighted by the flipped kernel. Your version will i) zero-pad an image, ii) flip the kernel horizontally and vertically, and iii) compute a weighted sum of the neighborhood at each pixel.

#### **Problem 1.1.1 [1 pts]**

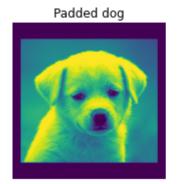
First you will want to implement the zero\_pad function.

```
In [1]: import numpy as np
    from time import time
    from skimage import io
    from heapq import *
    %matplotlib inline
    import matplotlib.pyplot as plt
```

```
def zero pad(image, pad top, pad down, pad left, pad right):
In \lceil 2 \rceil:
             """ Zero-pad an image.
             Ex: a 1x1 image [[1]] with pad top = 1, pad down = 1, pad left = 2, pad right = 2 become
                  [[0, 0, 0, 0, 0],
                   [0, 0, 1, 0, 0],
                                           of shape (3, 5)
                  [0, 0, 0, 0, 0]
             Args:
                  image: numpy array of shape (H, W)
                  pad left: width of the zero padding to the left of the first column
                 pad right: width of the zero padding to the right of the last column
                 pad top: height of the zero padding above the first row
                 pad down: height of the zero padding below the last row
             Returns:
                 out: numpy array of shape (H + pad top + pad down, W + pad left + pad right)
             YOUR CODE HERE
             _____ """
             height = np. size (image, 0)
             width = np. size (image, 1)
             out = np.zeros([height+pad top+pad down, width+pad left+pad right])
             out[pad top:pad top+height, pad left:pad left+width] = image
             return out
         # Open image as grayscale
         img = io.imread('dog.jpg', as gray=True)
         # Show image
         plt.imshow(img)
         plt.axis('off')
         plt. show()
         pad width = 20 # width of the padding on the left and right
         pad height = 40 # height of the padding on the top and bottom
         padded img = zero pad(img, pad height, pad height, pad width)
         # Plot your padded dog
         plt. subplot (1, 2, 1)
         plt.imshow(padded img)
         plt. title ('Padded dog')
         plt.axis('off')
         # Plot what you should get
         solution img = io.imread('padded dog.jpg', as gray=True)
         plt. subplot (1, 2, 2)
         plt.imshow(solution img)
         plt.title('What you should get')
         plt.axis('off')
```

plt.show()







Problem 1.1.2 [2 pts]

Now implement the function  $\operatorname{conv}$ , **using at most 2 loops**. This function should take an image f and a kernel h as inputs and output the convolved image (f\*h) that has the same shape as the input image (use zero padding to accomplish this). Depending on the computer, your implementation should take around a second or less to run.

```
def conv(image, kernel):
In [3]:
             """ An efficient implementation of a convolution filter.
             This function uses element-wise multiplication and np. sum()
             to efficiently compute a weighted sum of the neighborhood at each
             pixel.
             Hints:
                 - Use the zero pad function you implemented above
                 - You should need at most two nested for-loops
                 - You may find np.flip() and np.sum() useful
                 - You need to handle both odd and even kernel size
             Args:
                 image: numpy array of shape (Hi, Wi)
                 kernel: numpy array of shape (Hk, Wk)
             Returns:
                 out: numpy array of shape (Hi, Wi)
             Hi, Wi = image. shape
             Hk, Wk = kernel.shape
             out = np.zeros((Hi, Wi))
             """ ======
             YOUR CODE HERE
             _____ """
             kernel = np. flip(kernel, 0)
             kernel = np. flip(kernel, 1)
             pad top = 0
             pad down = 0
             pad left = 0
             pad right = 0
             if Hk \% 2 == 0:
                 pad top = Hk//2 - 1
                 pad down = Hk//2
             else:
                 pad_top = Hk//2
                 pad down = Hk//2
             if Wk \% 2 == 0:
                 pad left = Wk//2 - 1
                 pad right = Wk//2
             else:
                 pad left = Wk//2
                 pad right = Wk//2
             image pad = zero pad(image, pad top, pad down, pad left, pad right)
             for i in range(Hi):
                 for j in range(Wi):
                     out[i, j]=np. sum(image pad[i:i+Hk, j:j+Wk]*kernel)
```

```
return out
# Simple convolution kernel.
# Feel free to change the kernel and to see different outputs.
kernel = np. array(
    [1, 0, -1],
    [2, 0, -2],
    [1, 0, -1]
])
t1 = time()
out = conv(img, kernel)
t2 = time()
print("took %f seconds." % (t2 - t1))
# Plot original image
plt. subplot (2, 2, 1)
plt.imshow(img)
plt.title('Original')
plt.axis('off')
# Plot your convolved image
plt. subplot (2, 2, 3)
plt. imshow(out)
plt.title('Convolution')
plt.axis('off')
# Plot what you should get
solution_img = io.imread('convolved_dog.jpg', as_gray=True)
plt. subplot (2, 2, 4)
plt.imshow(solution img)
plt. title ('What you should get')
plt.axis('off')
plt.show()
```

took 0.887101 seconds.

#### Original



Convolution



What you should get



#### **Problem 1.1.3 [1 pt]**

Now let's filter some images! Here, you will apply the convolution function that you just implemented in order to bring about some interesting image effects. More specifically, we will use convolution to blur and sharpen our images.

First we will apply convolution for image blurring. To accomplish this, convolve the dog image with a 7x7 Gaussian filter for  $\sigma = 1.0$ . You can use the included function to obtain the Gaussian kernel.

```
In [4]:
         def gaussian2d(filter size=7, sig=1.0):
             Creates 2D Gaussian kernel with side length `filter_size` and a sigma of `sig`.
             ax = np. arange(-filter\_size // 2 + 1., filter\_size // 2 + 1.)
             xx, yy = np. meshgrid(ax, ax)
             kernel = np. exp(-0.5 * (np. square(xx) + np. square(yy)) / np. square(sig))
             return kernel / np. sum(kernel)
         def blur_image(img):
              """Blur the image by convolving with a Gaussian filter."""
             blurred img = np. zeros like(img)
             YOUR CODE HERE
              ______ """
             kernel = gaussian2d()
             blurred_img = conv(img, kernel)
             return blurred img
         # Plot original image
         plt. subplot (2, 2, 1)
         plt.imshow(img)
         plt.title('Original')
         plt.axis('off')
         # Plot blurred image
         plt. subplot (2, 2, 2)
         plt.imshow(blur image(img))
         plt. title('Blurred')
         plt.axis('off')
         plt.show()
```





**Problem 1.1.4 [1 pt]** 

Next, we will use convolution to sharpen the images. Convolve the image with the following filter to produce a sharpened result. For convenience, we have defined the filter for you:

```
In [6]:
         def sharpen image (img):
              """Sharpen the image by convolving with a sharpening filter."""
             sharpened img = np. zeros like(img)
             YOUR CODE HERE
              _____ """
             sharpened_img = conv(img, sharpening_kernel)
             return sharpened img
         # Plot original image
         plt. subplot (2, 2, 1)
         plt.imshow(img, vmin=0.0, vmax=1.0)
         plt.title('Original')
         plt.axis('off')
         # Plot sharpened image
         plt. subplot (2, 2, 2)
         plt.imshow(sharpen image(img), vmin=0.0, vmax=1.0)
         plt. title('Sharpened')
         plt.axis('off')
         plt.show()
```

# Original



Problem 1.2: Convolution Theory [5 pts]

### Problem 1.2.1 [2 pts]

Consider (1) smoothing an image with a 3x3 box filter and then computing the derivative in the x-direction. Also consider (2) computing the derivative first, then smoothing. What is a single convolution kernel that will simultaneously implement both (1) and (2)? Try to give a brief justification for how you arrived at the kernel.

Use the x-derivative filter

[-1/2, 0, 1/2]

for this problem.

Assume that the gradient convolution kernel is D, the smooth convolution kernel is K, and the image is  ${\rm I}$ 

so (1) can express as I\*(K\*D), (2) can express as (I\*D)\*K Convolution has commutative and associative laws

(I\*K)\*D=I\*(K\*D), (I\*D)\*K=I\*(D\*K), Due to commutative law K\*D=D\*K, so (I\*K)\*D=(I\*D)\*K, So we can define a new convolution kernel M,

M=K\*D=D\*K, so I\*M can process 1 and 2 operations at the same time

#### **Problem 1.2.2 [3 pts]**

Certain 2D filters can be expressed as a convolution of two 1D filters. Such filters are called separable filters. Give an example of a 3x3 separable filter and compare the number of arithmetic operations it takes to convolve an n x n image using that filter before and after separation. Count both, the number of multiplication and addition operations in each case.

Assume that the convolution of the image and filter is performed in "valid" mode, i.e., the image is not padded before convolution.

## Type *Markdown* and LaTeX: $\alpha^2$

For the picture of n\*n, the convolution of 3\*3 once requires 9 times of multiplication and 8 times of addition,

so a total of  $9 \text{n}^2$  times of multiplication is required and  $8 \text{n}^2$  times of addition is required

For the picture of n\*n, the convolution of 1\*3 once requires 3 times of multiplication and 2 times of addition,

so a total of 3n<sup>2</sup> times of multiplication is required and 2\*n<sup>2</sup> additions, performing a convolution of 3\*1 needs to do 3 times multiplications and 2 times additions, so a total of 3\*n<sup>2</sup> multiplications and

 $2*n^2$  Times addition. Therefore, a total of  $6*n^2$  times multiplications and  $4*n^2$  times additions are required.

It can be found that solving the convolution of into two one-dimensional convolutions can reduce the amount of calculation while reducing the amount of parameters

# Problem 1.3 Template Matching [5 pts]

Suppose that you are a clerk at a grocery store. One of your responsibilities is to check the shelves periodically and stock them up whenever there are sold-out items. You got tired of this laborious task and decided to build a computer vision system that keeps track of the items on the shelf.

Luckily, you have learned in CSE 252A (or are learning right now) that convolution can be used for template matching: a template g is multiplied with regions of a larger image f to measure how similar each region is to the template. Note that you will want to flip the filter before giving it to your

convolution function, so that it is overall not flipped when making comparisons. You will also want to subtract off the mean value of the image or template (whichever you choose, subtract the same value from both the image and template) so that our solution is not biased toward higher-intensity (white) regions.

The template of a product (template.jpg) and the image of the shelf (shelf.jpg) is provided. We will use convolution to find the product in the shelf.



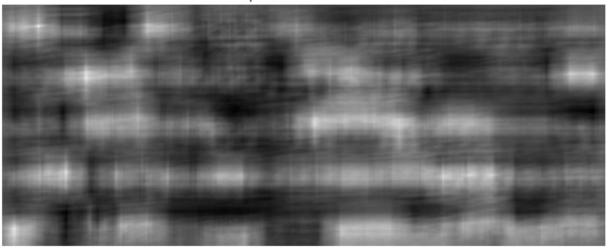


```
In [7]: # Load template and image in grayscale
         img = io.imread('shelf.jpg')
         img_gray = io.imread('shelf.jpg', as_gray=True)
         temp = io.imread('template.jpg')
         temp gray = io.imread('template.jpg', as gray=True)
         # Perform a convolution between the image and the template and store
         # the result in the out variable
         YOUR CODE HERE
         _____ """
         temp gray = np. flip(np. flip(temp gray, 0), 1)
         out = conv(img gray-np.mean(img gray), temp gray-np.mean(img gray))
         mx = np. max(out)
         h, w = out. shape
         X = []
         y = []
         for i in range(h):
             for j in range (w):
                 if out[i, j] == mx:
                      y. append(i)
                      x. append(j)
         # Display product template
         plt. figure (figsize= (20, 16))
         plt. subplot (3, 1, 1)
         plt.imshow(temp gray, cmap="gray")
         plt. title('Template')
         plt.axis('off')
         # Display convolution output
         plt. subplot (3, 1, 2)
         plt.imshow(out, cmap="gray")
         plt.title('Convolution output (white means more correlated)')
         plt.axis('off')
         # Display image
         plt. subplot (3, 1, 3)
         plt.imshow(img, cmap="gray")
         plt.title('Result (blue marker on the detected location)')
         plt.axis('off')
         # Draw marker at detected location
         plt.plot(x, y, 'bx', ms=40, mew=10)
         plt. show()
```

Template



Convolution output (white means more correlated)



Result (blue marker on the detected location)



# Problem 2: Edge detection [21 pts]

In this problem, you will write a function to perform Canny edge detection. The following steps need to be implemented.

# Problem 2.1 Smoothing [1 pt]

First, we need to smooth the images in order to prevent noise from being considered as edges. For this assignment, use a 9x9 Gaussian kernel filter with  $\sigma = 1.4$  to smooth the images.

```
In [8]: import numpy as np
    from skimage import io
    import matplotlib.pyplot as plt
    import matplotlib.cm as cm
    from scipy.signal import convolve
    %matplotlib inline

import matplotlib
matplotlib.rcParams['figure.figsize'] = [5, 5]
```

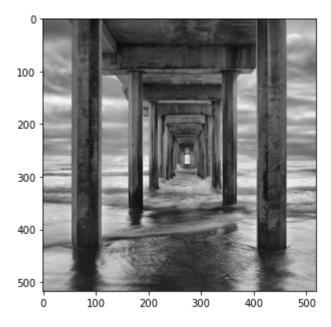
```
In [9]: def gaussian2d(filter_size=None, sig=None):
    """Creates a 2D Gaussian kernel with
    side length `filter_size` and a sigma of `sig`."""
    ax = np.arange(-filter_size // 2 + 1., filter_size // 2 + 1.)
    xx, yy = np.meshgrid(ax, ax)
    kernel = np.exp(-0.5 * (np.square(xx) + np.square(yy)) / np.square(sig))
    return kernel / np.sum(kernel)
```

```
In [10]: def smooth(image):
    """ ========
    YOUR CODE HERE
    ======== """
    kernel = gaussian2d(9, 1.4)
    out = conv(image, kernel)
    return out
```

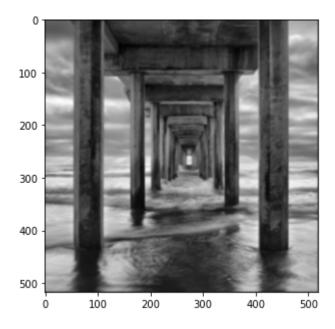
```
In [11]: # Load image in grayscale
    image = io.imread('sio_pier.jpg', as_gray=True)
    assert len(image.shape) == 2, 'image should be grayscale; check your Python/skimage versions smoothed = smooth(image)
    print('Original:')
    plt.imshow(image, cmap=cm.gray)
    plt.show()

print('Smoothed:')
    plt.imshow(smoothed, cmap=cm.gray)
    plt.show()
```

### Original:



#### Smoothed:

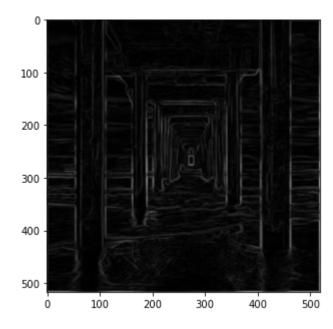


# **Problem 2.2 Gradient Computation [5 pts]**

After you have finished smoothing, find the image gradient in the horizontal and vertical directions. Compute the gradient magnitude image as  $|G| = \sqrt{G_x^2 + G_y^2}$ . The edge direction for each pixel is given by  $G_\theta = \tan^{-1}\left(\frac{G_y}{G_x}\right)$ .

```
In [13]: g_mag, g_theta = gradient(smoothed)
print('Gradient magnitude:')
plt.imshow(g_mag, cmap=cm.gray)
plt.show()
```

Gradient magnitude:



# Problem 2.3 Non-Maximum Suppression [7 pts]

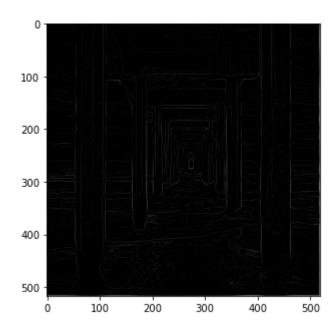
We would like our edges to be sharp, unlike the ones in the gradient image. Use non-maximum suppression to preserve all local maxima and discard the rest. You can use the following method to do so:

- For each pixel in the gradient magnitude image:
  - Round the gradient direction  $\theta$  to the nearest multiple of  $45^{\circ}$  (which we will refer to as ve).
  - Compare the edge strength at the current pixel to the pixels along the +ve and -ve gradient direction in the 8-connected neighborhood.
  - If the pixel does not have a larger value than both of its two neighbors in the +ve and -ve gradient directions, suppress the pixel's value (set it to 0). By following this process, we preserve the values of only those pixels which have maximum gradient magnitudes in the neighborhood along the +ve and -ve gradient directions.
- · Return the result as the NMS response.

```
In [14]:
         def nms(g_mag, g_theta):
              YOUR CODE HERE
               _____ """
              nms_response = np.zeros_like(g_mag)
              h, w = g \text{ theta. shape}
              for i in range(h):
                   for j in range(w):
                       cur theta = g theta[i, j]
                       cur theta \neq np. pi/2
                       cur theta = np. round(cur theta)
                       flag = True
                       if cur theta == 0.0:
                           if j+1 < w and g_mag[i, j] < g_mag[i, j+1]:
                               flag = False
                           if j-1 \ge 0 and g mag[i, j] < g mag[i, j-1]:
                               flag = False
                       elif cur theta == 1.0:
                           if i+1 \le h and j+1 \le w and g mag[i,j] \le g mag[i+1,j+1]:
                               flag = False
                           if i-1 \ge 0 and j-1 \ge 0 and g mag[i, j] < g mag[i-1, j-1]:
                               flag = False
                       elif cur theta == 2.0:
                           if i+1 < h and g_mag[i, j] < g_mag[i+1, j]:
                               flag = False
                           if i-1 \ge 0 and g mag[i, j] < g mag[i-1, j]:
                               flag = False
                       elif cur theta == -1.0:
                           if i-1 \ge 0 and j+1 \le w and g mag[i, j] \le g mag[i-1, j+1]:
                               flag = False
                           if i+1 < h and j-1 >= 0 and g mag[i, j] < g mag[i+1, j-1]:
                               flag = False
                       elif cur theta == -2.0:
                           if i-1 \ge 0 and g_mag[i,j] < g_mag[i-1,j]:
                               flag = False
                           if i+1 < h and g_mag[i,j] < g_mag[i+1,j]:
                               flag = False
                       if flag:
                           nms response[i, j] = g mag[i, j]
                       else:
                           nms_response[i, j] = 0
              return nms_response
```

```
In [15]: nms_image = nms(g_mag, g_theta)
    print('NMS:')
    plt.imshow(nms_image, cmap=cm.gray)
    plt.show()
```

NMS:



# Problem 2.4 Hysteresis Thresholding [8 pts]

Choose suitable values of thresholds and use the thresholding approach decribed in lecture 6. This will remove the edges caused by noise and color variations.

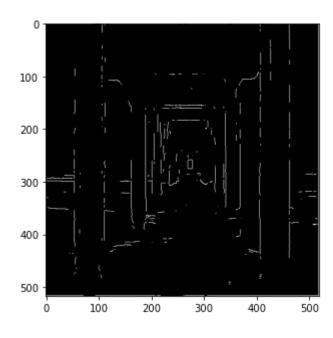
- Define two thresholds  $t_min$  and  $t_max$ .
- If the nms > t max, then we select that pixel as an edge.
- If nms < t min , we reject that pixel.
- If  $t_{min} < nms < t_{max}$ , we select the pixel only if there is a path from to another pixel with  $nms > t_{max}$ . (Hint: Think of all pixels with  $nms > t_{max}$  as starting points and run BFS/DFS from these starting points).
- The choice of value of low and high thresholds depends on the range of values in the gradient magnitude image. You can start by setting the high threshold to some percentage of the max value in the gradient magnitude image, e.g. thres\_high = 0.2 \* image.max(), and the low threshold to some percentage of the high threshold, e.g. thres\_low = 0.85 \* thres\_high. And then you can tune those values however you want.

```
In [16]:
           def hysteresis threshold(image, g theta, use g theta=False):
               YOUR CODE HERE
               ______ """
               h, w = image. shape
               result = np. zeros like(image)
               t max = np. max(image)*0.2
               t min = t max*0.85
               def valid coordinate(y, x):
                    if image[y, x] > t_min and <math>image[y, x] < t_max:
                        return True
                    else:
                        return False
               vis = np. zeros([h, w])
               for i in range(h):
                    for j in range(w):
                        if image[i, j] > t max and result[i, j] != 1:
                            result[i, j] = 1
                            que = [(i, j)]
                            while len(que)>0:
                                y, x = que. pop()
                                 theta = g_{theta}[y, x]
                                 theta = np. pi/2
                                 theta = np. round(theta)
                                 if theta == 0.0:
                                     if y+1 \le h and result[y+1, x]!=1 and vis[y+1, x]!=1:
                                         que. append ((y+1, x))
                                         vis[y+1, x]=1;
                                         if valid coordinate (y+1, x):
                                             result[y+1, x]=1;
                                     if y-1 \ge 0 and result[y-1, x]!=1 and vis[y-1, x]!=1:
                                         que. append ((y-1, x))
                                         vis[y-1, x]=1
                                         if valid coordinate (y-1, x):
                                             result[y-1, x]=1
                                elif theta == 1.0:
                                     if y+1 \le h and x-1 \ge 0 and result [y+1, x-1]!=1 and vis[y+1, x-1]!=1:
                                         que. append ((y+1, x-1))
                                         vis[y+1, x-1]=1
                                         if valid coordinate (y+1, x-1):
                                              result[y+1, x-1]=1;
                                     if y-1 \ge 0 and x+1 \le w and result [y-1, x+1] = 1 and vis[y-1, x+1] = 1:
                                         que. append ((y-1, x+1))
                                         vis[y-1, x+1]=1
                                         if valid coordinate (y-1, x+1):
                                             result[y-1, x+1]=1
                                elif theta == 2.0 or theta == -2.0:
                                     if x+1 \le w and result[y, x+1]!=1 and vis[y, x+1]!=1:
                                         que. append ((y, x+1))
                                         vis[y, x+1]=1
                                         if valid coordinate (y, x+1):
                                             result[y, x+1]=1;
                                     if x-1 \ge 0 and result[y, x-1]!=1 and vis[y, x-1]!=1:
                                         que. append ((y, x-1))
                                         vis[y, x-1]=1
                                         if valid coordinate (y, x-1):
```

```
result[y, x-1]=1
elif theta == -1.0:
    if y+1<h and x+1<w and result[y+1, x+1]!=1 and vis[y+1, x+1]!=1:
        que.append((y+1, x+1))
        vis[y+1, x+1]=1
        if valid_coordinate(y+1, x+1):
            result[y+1, x+1]=1;
        if y-1>=0 and x-1>=0 and result[y-1, x-1]!=1 and vis[y-1, x-1]!=1
        que.append((y-1, x-1))
        vis[y-1, x-1]=1
        if valid_coordinate(y-1, x-1):
            result[y-1, x-1]=1
elif image[i, j] < t_min:
        result[i, j]=0
return result</pre>
```

```
In [17]: thresholded = hysteresis_threshold(nms_image, g_theta)
    print('Thresholded:')
    plt.imshow(thresholded, cmap=cm.gray)
    plt.show()
```

#### Thresholded:



# Problem 3 Corner detection [13 pts]

# **Problem 3.1 [12 pts]**

In this problem, we are going to build a corner detector. This should be done according to the lecture slides. You should fill in the function <code>corner\_detect</code> below, which takes as input <code>image</code>, <code>nCorners</code>, <code>smoothSTD</code>, <code>windowSize</code> -- where <code>smoothSTD</code> is the standard deviation of the smoothing kernel and <code>windowSize</code> is the window size for corner detector and non-maximum suppression. In the lecture, the corner detector was implemented using a hard threshold. Do not do that; instead, return the <code>nCorners</code> strongest corners after non-maximum suppression. This way you can control exactly how

many corners are returned. Run your code on all four images (with nCorners = 20) and display outputs as shown below. You may find  $scipy.ndimage.filters.gaussian_filter helpful for smoothing.$ 

In this problem, try the following different standard deviation ( $\sigma$ ) parameters for the Gausian smoothing kernel: 0.5, 1, 2 and 4. For a particular  $\sigma$ , you should take the kernel size to be  $6 \times \sigma$  (add 1 if the kernel size is even). So for example if  $\sigma = 2$ , corner detection kernel size should be 13. This should be followed throughout all of the experiments in this assignment.

There will be a total of 16 images as outputs: 4 choices of smooth STD  $\times$  2 matrix images  $\times$  2 warrior images.



# inoCorner2

```
In [18]: import numpy as np
import matplotlib.pyplot as plt
from scipy.ndimage.filters import gaussian_filter
import imageio
from scipy.signal import convolve

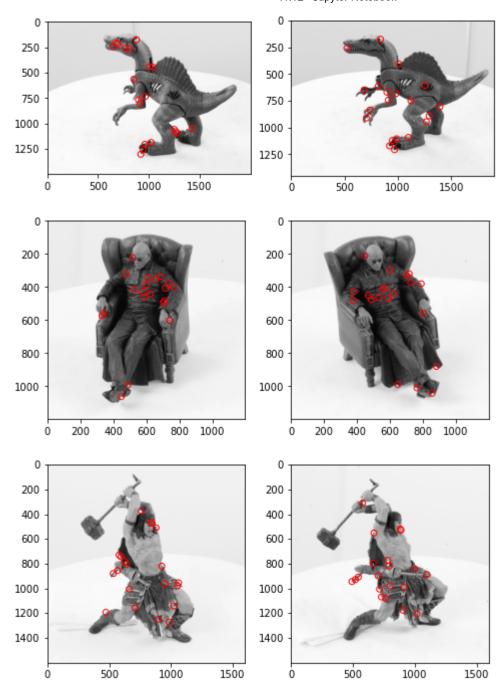
def rgb2gray(rgb):
    """ Convert rgb image to grayscale.
    """
    return np.dot(rgb[...,:3], [0.299, 0.587, 0.114])
```

```
def corner detect(image, nCorners, smoothSTD, windowSize):
In [19]:
               """Detect corners on a given image.
               Args:
                   image: Given a grayscale image on which to detect corners.
                   nCorners: Total number of corners to be extracted.
                   smoothSTD: Standard deviation of the Gaussian smoothing kernel.
                   windowSize: Window size for corner detector and non-maximum suppression.
               Returns:
                   Detected corners (in image coordinate) in a numpy array (n*2).
               n n n
               YOUR CODE HERE
               _____ """
               h, w = image. shape
               filter size = 6 * smoothSTD
               if filter size % 2 == 0:
                   filter size += 1
               kernel = gaussian2d(filter size, smoothSTD)
               image f = conv(image, kernel)
               corners = np. zeros ((nCorners, 2))
               ix = np. zeros((h, w))
               iy = np. zeros((h, w))
               for i in range(h):
                   for j in range(w):
                       if j > 0:
                            ix[i, j] = image[i, j] - image[i, j-1]
                       else:
                            ix[i, j] = image[i, j]
                       if i > 0:
                            iy[i, j] = image[i, j] - image[i-1, j]
                       else:
                           iy[i, j] = image[i, j]
               ixx = ix*ix
               ixy = ix*iy
               iyy = iy*iy
               def comput_i (mat, i, j):
                   if i > 0:
                       mat[i, j] += mat[i-1, j]
                   if i > 0 and j > 0:
                       mat[i, j] = mat[i-1, j-1]
                   if j > 0:
                       mat[i, j] += mat[i, j-1]
               for i in range(h):
                   for j in range(w):
```

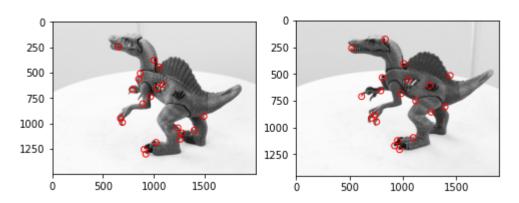
```
comput i(ixx, i, j)
        comput_i(iyy, i, j)
        comput_i(ixy, i, j)
def comput m(m, i, j, w):
    res = m[i+w-1, j+w-1]+0
    if i > 0:
        res = m[i-1, j+w-1]
    if j > 0:
        res = m[i+w-1, j-1]
    if i > 0 and j > 0:
        res += m[i-1, j-1]
    return res
score = np. zeros((h, w))
for i in range (h-windowSize):
    for j in range (w-windowSize):
        M = np. zeros((2, 2))
        M[0,0] = comput m(ixx, i, j, windowSize)
        M[0, 1]=comput m(ixy, i, j, windowSize)
        M[1,0]=comput m(ixy, i, j, windowSize)
        M[1, 1] = comput m(iyy, i, j, windowSize)
        1, v = np. linalg. eig(M)
        score[i, j] = 1[0]*1[1]-0.05*(1[0]+1[1])*(1[0]+1[1])
1st = []
for i in range (windowSize, h-2*windowSize):
    for j in range (windowSize, w-2*windowSize):
        flag = True
        for si in range (-1, 1):
             for sj in range (-1,1):
                 ci = si + i
                 cj = sj + j
                 if score[i, j] < score[ci, cj]:</pre>
                     flag = False
        if flag:
             1st. append((score[i, j], i, j))
1st. sort()
cnt = 0
while cnt < nCorners and len(1st) > 0:
    score, i, j = 1st.pop()
    flag = True
    for idx in range(cnt):
        if abs(j-corners[idx,0])<2*windowSize or abs(i-corners[idx,1])<2*windowSize:
            flag = False
            break
    if flag:
        corners[cnt, 0] = j
        corners[cnt, 1] = i
        cnt += 1
corners += windowSize / 2
return corners
```

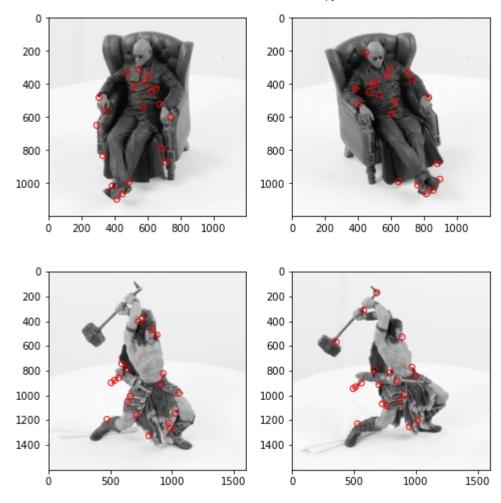
```
In [240]:
           def show corners result(imgs, corners):
               fig = plt. figure (figsize= (8, 8))
               ax1 = fig. add subplot (221)
               ax1. imshow(imgs[0], cmap='gray')
               ax1. scatter(corners[0][:, 0], corners[0][:, 1], s=35, edgecolors='r', facecolors='none
               ax2 = fig. add subplot (222)
               ax2. imshow(imgs[1], cmap='gray')
               ax2. scatter(corners[1][:, 0], corners[1][:, 1], s=35, edgecolors='r', facecolors='none
               plt.show()
           for smoothSTD in (0.5, 1, 2, 4):
               windowSize = int(smoothSTD * 6)
               if windowSize \% 2 == 0:
                   windowSize += 1
               print('smooth stdev: %r' % smoothSTD)
               print('window size: %r' % windowSize)
               nCorners = 20
               # read images and detect corners on images
               imgs din = []
               crns din = []
               imgs mat = []
               crns mat = []
               imgs war = []
               crns war = []
               for i in range (2):
                   img din = imageio.imread('dino/dino' + str(i) + '.png')
                   imgs din.append(rgb2gray(img din))
                   # downsize your image in case corner detect runs slow in test
                   # imgs din.append(rgb2gray(img din)[::2, ::2])
                   crns_din.append(corner_detect(imgs_din[i], nCorners, smoothSTD, windowSize))
                   img mat = imageio.imread('matrix/matrix' + str(i) + '.png')
                   imgs mat.append(rgb2gray(img mat))
                   # downsize your image in case corner detect runs slow in test
                   # imgs mat.append(rgb2gray(img mat)[::2, ::2])
                   crns mat.append(corner detect(imgs mat[i], nCorners, smoothSTD, windowSize))
                   img war = imageio.imread('warrior/warrior' + str(i) + '.png')
                   imgs war.append(rgb2gray(img war))
                   # downsize your image in case corner detect runs slow in test
                   # imgs war.append(rgb2gray(img war)[::2, ::2])
                   crns war.append(corner detect(imgs war[i], nCorners, smoothSTD, windowSize))
               show corners result (imgs din, crns din)
               show corners result (imgs mat, crns mat)
               show corners result (imgs war, crns war)
```

smooth stdev: 0.5
window size: 3

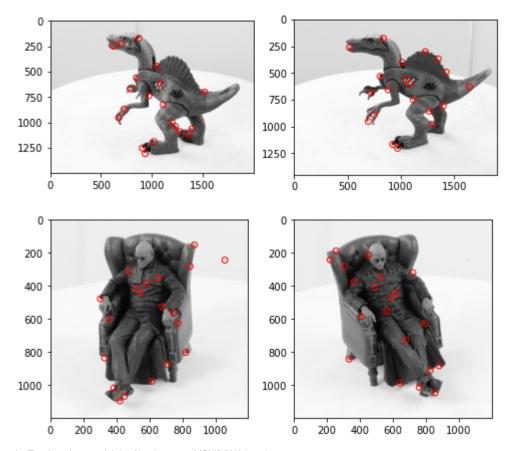


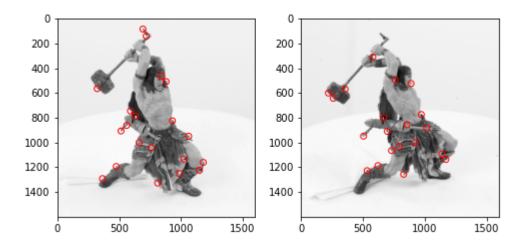
smooth stdev: 1
window size: 7



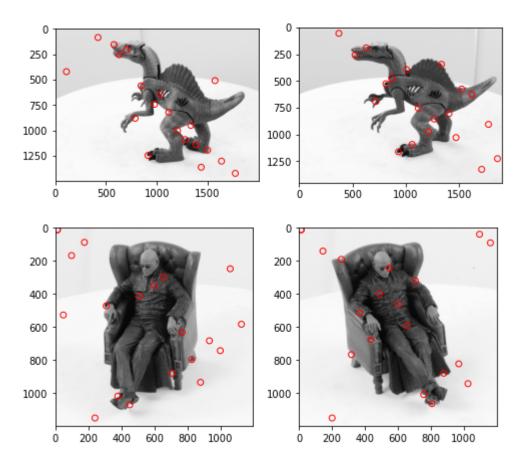


smooth stdev: 2
window size: 13





smooth stdev: 4
window size: 25





# Problem 3.2 [1 pts]

Comment on your results and observations. You don't need to comment per output; just discuss any trends you see for the detected corners as you change the windowSize and increase the smoothing w.r.t the two pairs of images (warrior and matrix). Also discuss whether you are able to find corresponding corners for the pairs of images.

Increase the window size, the detection range becomes larger, and the detection points are more scattered, but no small inflection points can be detected. When the variance reaches 4, only A

# Problem 4 Epipolar rectification and feature matching [43 pts]

# 4.1 Epipolar rectification [22 pts]

small number of inflection points were detected.

In this problem, we are going to perform epipolar rectification. Given calibrated stereo cameras (i.e., calibration matrices  $K_1$  and  $K_2$ , camera rotation matrices  $R_1$  and  $R_2$ , camera translation vectors  $t_1$  and  $t_2$ ), you are expected to determine the rotation matrix R and calibration matrix K of the virtual cameras. Your goal is to complete the function epipolarRecification, which determines the calibration matrix and rotation matrix of both cameras, the translation vector of each of the cameras, and matching planar transformations that epipolar rectify the two images acquired by the cameras. The destination virtual cameras have the same centers as the source real cameras.

#### 4.1.1 Camera translation matrices and Projective Transformation matrices [6 pts]

To calculate the camera translation from cameras with the same camera center, you will have to complete the cameraTranslation first. Another function you need to complete is calcProjectiveTransformation, which calculates the planar projective transformation from cameras with the same camera center. The camera calibration matrix (same for both cameras) will be calculated by calcDestinateK. This is provided for you. To get the rotation matrix R of the virtual camera, we usually interpolate halfway between the two 3D rotations embodied by  $R_1$  and  $R_2$ . For simplicity, this will be also given to you.

```
In [20]: from imageio import imread import matplotlib.pyplot as plt import numpy as np import math import pickle from math import floor, ceil
```

```
In [21]:
          def cameraTranslation(R1, t1, R2):
              Calculate the camera translation from cameras with the same camera center.
              Args:
              R1: The rotation matrix of the first camera.
              tl: The translation vector of the first camera.
              R2: The rotation matrix of the second camera.
              Returns:
              The translation vector of the second camera.
              """ -----
              YOUR CODE HERE
              _____ """
              sz = 1en(R1)
              I = np. eye(sz)
              M1 = I - R2
              M2 = np. linalg. inv (I-R1)
              return M1. dot(M2). dot(t1)
```

```
[22]:
       def calcProjectiveTransformation(K1, R1, K2, R2):
           Calculates the planar projective transformation from cameras with the same camera cent
           This function determines the planar projective transformation from the image of a 3D po
           Args:
           K1: The calibration matrix of the first camera.
           R1: The rotation matrix of the first camera.
           K2: The calibration matrix of the second camera.
           R2: The rotation matrix of the second camera.
           Returns:
           The transformation matrix.
           """ =======
           YOUR CODE HERE
           _____ """
           M1 = K1. dot(R1)
           M1 = np. linalg. inv (M1)
           M2 = K2. dot(R2)
           trans = M2. dot(M1)
           return trans
```

In

[23]:

def calcDestinateK(srcK1, srcK2):

Camera calibration matrix (same for both cameras)

```
alpha = (srcK1[0][0] + srcK2[0][0] + srcK1[1][1] + srcK2[1][1]) // 4
           x0 = (srcK1[0][2] + srcK2[0][2]) // 2
           y0 = (srcK1[1][2] + srcK2[1][2]) // 2
           dstK = np. zeros((3, 3))
           dstK[0][0] = alpha
           dstK[0][2] = x0
           dstK[1][1] = alpha
           dstK[1][2] = y0
           dstK[2][2] = 1
           return dstK
[24]: def epipolarRecification(srcK1, srcR1, src t1,
                                srcK2, srcR2, src t2, dstR):
           Given two calibrated cameras, this function determines the calibration matrix and rota
           Args:
           srcK1: The calibration matrix of the first source camera.
           srcR1: The rotation matrix of the first source camera.
           src tl: The translation vector of the first source camera.
           srcK2: The calibration matrix of the second source camera.
           srcR2: The rotation matrix of the second source camera.
           src t2: The translation vector of the second source camera.
           dstR: The rotation matrix of the destination cameras.
           Returns:
           dstK: The calibration matrix of the destination cameras.
           dst t1: The translation vector of the first destination camera.
           dst t2: The translation vector of the second destination camera.
           H1, H2: The image rectification transformation matrices.
           dst t1 = cameraTranslation(srcR1, src t1, dstR)
           dst t2 = cameraTranslation(srcR2, src t2, dstR)
           dstK = calcDestinateK(srcK1, srcK2)
           H1 = calcProjectiveTransformation(srcK1, srcR1, dstK, dstR)
           H2 = calcProjectiveTransformation(srcK2, srcR2, dstK, dstR)
           return dstK, dst t1, dst t2, H1, H2
```

Problem 4.1.2 Warp Image [10 pts]

After calling epipolarRectification, we can get the projective transformation matrices H1 and H2. Next, we will geometrically transform (i.e., 'warp') the image so that the epipolar lines are image rows. You have to complete warpImage using the backward method in Lecture 7. Note the destination images are required to be the same size as the source images.

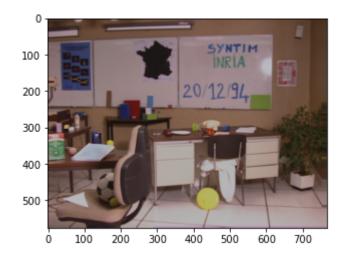
```
[25]:
       def warpImage(image, H, out height, out width):
           Performs the warp of the full image content.
           Calculates bounding box by piping four corners through the transformation.
           Args:
           image: Image to warp
           H: The image rectification transformation matrices.
           out_height, out_width: The shape of output image.
           Returns:
           Out: An inverse warp of the image, given a homography.
           min x, min y, max x, max y: The minimum/maxmum of warped image bound.
           """ =======
           YOUR CODE HERE
           _____ """
           out = np. zeros ((out height, out width, 3), np. int64)
           min x = out height
           min_y = out_width
           \max x = 0
           \max y = 0
           for i in range(out_height):
               for j in range (out width):
                   x = H. dot(np. array([[i], [j], [1]]))[:, 0]
                   \min x = \min(\min x, x[0])
                   \min y = \min(\min y, x[1])
                   \max x = \max(\max x, x[0])
                   \max_{y} = \max(\max_{y}, x[1])
                   cx = int(round(x[0]))
                   cy = int(round(x[1]))
                   if cy < 0:
                       cy = 0
                   elif cy >= out width:
                       cy = out width - 1
                   if cx < 0:
                       cx = 0
                   elif cx \ge out height:
                       cx = out height - 1
                   out[cx, cy, :] = image[i, j, :]
           return out, min_x, min_y, max_x, max_y
```

```
In [37]: file_param = open('param.pkl', 'rb')
    param = pickle.load(file_param)
    file_param.close()
    srcK1, srcR1, src_t1 = param['srcK1'], param['srcR1'], param['src_t1']
    srcK2, srcR2, src_t2 = param['srcK2'], param['srcR2'], param['src_t2']
    dstR = param['dstR']
```

```
In [38]: dstK, dst_t1, dst_t2, H1, H2 = epipolarRecification(srcK1, srcR1, src_t1, srcK2, srcR2, src_t2, dstR)
```

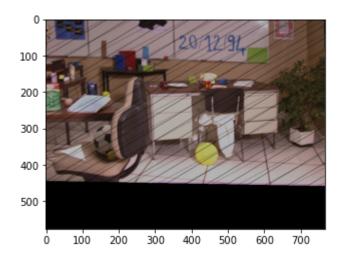
```
In [39]: src1 = imread('Sport0_0G0.bmp')
    plt. imshow(src1)
    print('Original image 1:')
```

#### Original image 1:



```
In [40]: height1, width1, _ = src1.shape
    rectified_im1_unbounded, min_x1, min_y1, max_x1, max_y1 = warpImage(src1, H1, height1, wid
    plt.imshow(rectified_im1_unbounded)
    print('Unbounded Rectified image 1:')
```

#### Unbounded Rectified image 1:



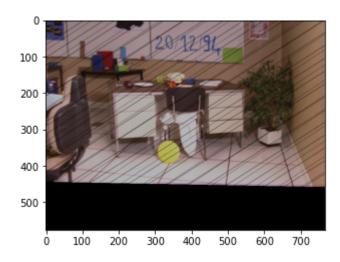
```
In [41]: src2 = imread('Sport1_0G0.bmp')
plt.imshow(src2)
print('Original image 2:')
```

#### Original image 2:



In [42]: height2, width2, \_ = src2.shape
 rectified\_im2\_unbounded, min\_x2, min\_y2, max\_x2, max\_y2 = warpImage(src2, H2, height2, wid
 plt.imshow(rectified\_im2\_unbounded)
 print('Unbounded Rectified image 2:')

Unbounded Rectified image 2:



#### 4.1.3 Partial bounded retification [3 pts]

In the resulting images, although they are epipolar rectified, you should observe portions of the source images being transformed "out of bounds" of the destination images. To fix this problem, we can introduced a 2D transformation containing a translation (i.e., T1 and T2).

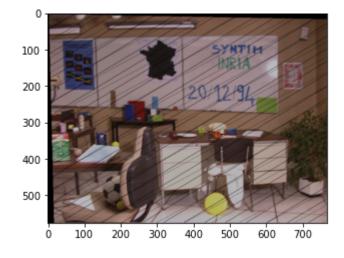
$$T1 = \begin{bmatrix} 1 & 0 & -min_{x}1 - 0.5 \\ 0 & 1 & -min(min_{y}1, min_{y}2) - 0.5 \\ 0 & 0 & 1 \\ \end{bmatrix}$$

$$T2 = \begin{bmatrix} 1 & 0 & -min_{x}2 - 0.5 \\ 0 & 1 & -min(min_{y}1, min_{y}2) - 0.5 \\ 0 & 0 & 1 \end{bmatrix}$$

H1, H2 can be updated by left multiplying T1, T2, respectivley. Again, geometrically tranform the images under the updated H1, H2. The destination image is required to be the same size as the source images. In the resulting images, although they are (still) epipolar rectified, you should observe the portions of the source images being transformed are no longer "out of bounds" on the top and left of the destination images.

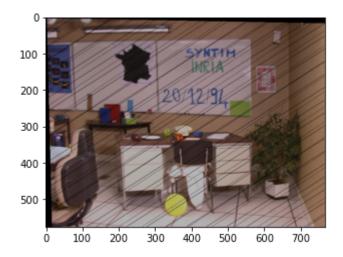
In [43]: H1\_bounded, H2\_bounded = partialboundedRetification(min\_x1, min\_y1, min\_x2, min\_y2, H1, H2 rectified\_im1\_bounded, min\_x1\_bounded, min\_y1\_bounded, max\_x1\_bounded, max\_y1\_bounded = waplt.imshow(rectified\_im1\_bounded) print('The partial bounded rectified image 1:')

The partial bounded rectified image 1:



```
In [44]: rectified_im2_bounded, min_x2_bounded, min_y2_bounded, max_x2_bounded, max_y2_bounded = warplt.imshow(rectified_im2_bounded) print('The partial bounded rectified image 2:')
```

The partial bounded rectified image 2:



## 4.1.4 Completely bounded rectification [3 pts]

Finally, determine the size of the destination images that completely bound the transformed images. Again geometrically transform the images under the updated 2D projective transformation matrices H1 and H2 (these are not updated a second time). You should complete the function completelyBoundedRectification. The destination images are required to be the size you just calculated. In the resulting images, you should observe the source images being transformed such that they are epipolar rectified and are completely bounded.

```
def completelyBoundedRectification(src1, src2, H1 bounded, H2 bounded,
                                     min x1 bounded, max x1 bounded,
                                     min y1 bounded, max y1 bounded):
    Determine the size of the destination images (same size for both) that completely bound
    """ =======
    YOUR CODE HERE
    _____ """
    min x1 bounded = int(min x1 bounded) - 1
    \max_{x_1} bounded = int(\max_{x_1} bounded) + 1
    min y1 bounded = int(min y1 bounded) - 1
    max_y1\_bounded = int(max_y1\_bounded) + 1
    h = max_x1\_bounded - min_x1\_bounded + 1
    w = max y1 bounded - min y1 bounded + 1
    rectified im1 final = np. zeros((h, w, 3), np. int64)
    rectified im2 final = np. zeros((h, w, 3), np. int64)
    h1, w1, _{-} = src1. shape
    for i in range(h1):
        for j in range(w1):
             x = H1 \text{ bounded. dot (np. array([[i], [j], [1]]))[:, 0]}
             cx = int(x[0]) - min x1 bounded
             cy = int(x[1]) - min y1 bounded
             rectified im1 final[cx, cy, :]=src1[i, j, :]
    h2, w2, \_ = src2.shape
    for i in range(h2):
        for j in range (w2):
             x = H2\_bounded. dot(np. array([[i], [j], [1]]))[:, 0]
             cx = int(x[0]) - min x1 bounded
             cy = int(x[1]) - min y1 bounded
             if cx \ge h:
                 cx = h-1
             if cy \ge w:
                 cy = w-1
             rectified im2 final[cx, cy, :]=src2[i, j, :]
    return rectified_im1_final, rectified_im2_final
```

```
In [65]: rectified_im1_final, rectified_im2_final = completelyBoundedRectification(src1, src2, H1_bounded, max_x1_bounded, min_y1_bounded)
```

# Problem 4.2 Feature matching [3 pts]

## 4.2.1 SSD (Sum Squared Distance) Matching [1 pts]

Complete the function ssdMatch : SSD =  $\sum_{x,y} |W_1(x,y) - W_2(x,y)|^2$ 

```
[28]:
          def ssdMatch(img1, img2, c1, c2, R):
In
               """Compute SSD given two windows.
              Args:
                   img1: Image 1.
                   img2: Image 2.
                  c1: Center (in image coordinate) of the window in image 1.
                   c2: Center (in image coordinate) of the window in image 2.
                  R: R is the radius of the patch, 2 * R + 1 is the window size
              Returns:
                  SSD matching score for two input windows.
               """ =======
              YOUR CODE HERE
               _____ """
              matching score = 0
              for i in range (-R, R+1):
                   for j in range (-R, R+1):
                       dist = img1[c1[1]+i, c1[0]+j]-img2[c2[1]+i, c2[0]+j]
                       matching score += dist * dist
              return matching score
```

```
In [19]: # Here is the code for you to test your implementation
    img1 = np.array([[1, 2, 3, 4], [4, 5, 6, 8], [7, 8, 9, 4]])
    img2 = np.array([[1, 2, 1, 3], [6, 5, 4, 4], [9, 8, 7, 3]])
    print(ssdMatch(img1, img2, np.array([1, 1]), np.array([1, 1]), 1))
    # should print 20
    print(ssdMatch(img1, img2, np.array([2, 1]), np.array([2, 1]), 1))
    # should print 30
    print(ssdMatch(img1, img2, np.array([1, 1]), np.array([2, 1]), 1))
# should print 46
```

#### Problem 4.2.2 NCC (Normalized Cross-Correlation) Matching [2 pts]

Write a function <code>ncc match</code> that implements the NCC matching algorithm for two input windows.

NCC = 
$$\sum_{i,j} \tilde{W_1}(i,j) \cdot \tilde{W_2}(i,j)$$

30 46 where  $ilde{W}=rac{W-\overline{W}}{\sqrt{\sum_{k,l}(W(k,l)-\overline{W})^2}}$  is a mean-shifted and normalized version of the window and  $\overline{W}$  is

the mean pixel value in the window W.

```
In [85]:
          def normalize window(window):
              mean = np. mean (window)
              stdev = np. sqrt(np. sum((window - mean) ** 2))
              return (window - _mean) / (_stdev + 1e-6)
          def ncc match(img1, img2, c1, c2, R):
              """Compute NCC given two windows.
              Args:
                  img1: Image 1.
                   img2: Image 2.
                  c1: Center (in image coordinate) of the window in image 1.
                  c2: Center (in image coordinate) of the window in image 2.
                  R: R is the radius of the patch, 2 * R + 1 is the window size
              Returns:
                  NCC matching score for two input windows.
              n n n
              """ =======
              YOUR CODE HERE
              ______ """
              mean1 = 0
              mean2 = 0
              sz = 2*R+1
              for i in range (-R, R+1):
                  for j in range (-R, R+1):
                      mean1 += img1[c1[1]+i, c1[0]+j]/(sz*sz)
                      mean2 += img2[c2[1]+i, c2[0]+j]/(sz*sz)
              d 1 = 0
              d 2 = 0
              for i in range (-R, R+1):
                   for j in range (-R, R+1):
                      t 1 = img1[c1[1]+i, c1[0]+j]-mean1
                      d 1 += t 1 * t 1
                      t 2 = img2[c2[1]+i, c2[0]+j]-mean2
                      d 2 += t 2 * t 2
              d 1 = np. sqrt(d 1)
              d = np. sqrt(d = 2)
              w 1 = (img1[-R+c1[1]:R+c1[1]+1,-R+c1[0]:R+c1[0]+1]-mean1)/d 1
              w 2 = (img2[-R+c2[1]:R+c2[1]+1,-R+c2[0]:R+c2[0]+1]-mean2)/d 2
              ans = 0
              for i in range(sz):
                   for j in range(sz):
                      ans += w_1[i, j] * w_2[i, j]
              return ans
```

```
In [21]: # test NCC match
    img1 = np.array([[1, 2, 3, 4], [4, 5, 6, 8], [7, 8, 9, 4]])
    img2 = np.array([[1, 2, 1, 3], [6, 5, 4, 4], [9, 8, 7, 3]])

print (ncc_match(img1, img2, np.array([1, 1]), np.array([1, 1]), 1))
# should print 0.8546

print (ncc_match(img1, img2, np.array([2, 1]), np.array([2, 1]), 1))
# should print 0.8457

print (ncc_match(img1, img2, np.array([1, 1]), np.array([2, 1]), 1))
# should print 0.6258
```

- 0.8546547739343037
- 0.845761528217442
- 0.6258689611426175

## Problem 4.3 Naive Matching [8 pts]

Equipped with the corner detector and the NCC matching function, we are ready to start finding correspondences. One naive strategy is to try and find the best match between the two sets of corner points. Write a script that does this, namely, for each corner in image1, find the best match from the detected corners in image2 (or, if the NCC match score is too low, then return no match for that point). You will have to figure out a good threshold (NCCth) value by experimentation.

Write a function <code>naive\_matching</code> and call it as below. Examine your results for 10, 20, and 30 detected corners in each image. Choose the number of detected corners to maximize the number of correct matching pairs. <code>naive\_matching</code> will call your NCC matching code.

Properly label or mention which output corresponds to which choice of number of corners.

The total number of outputs is 6 images: (3 choices of number of corners for each of matrix and warrior), where each figure might look like the following:

Number of corners: 10

dino match

```
def naive matching(img1, img2, corners1, corners2, R, NCCth):
In [30]:
               """Compute NCC given two windows.
              Args:
                   img1: Image 1.
                   img2: Image 2.
                   corners1: Corners in image 1 (nx2)
                   corners2: Corners in image 2 (nx2)
                   R: NCC matching radius
                   NCCth: NCC matching score threshold
              Returns:
                   NCC matching result a list of tuple (c1, c2),
                   c1 is the 1x2 corner location in image 1,
                   c2 is the 1x2 corner location in image 2.
               n n n
               """ =======
              YOUR CODE HERE
               _____ """
              matching = []
              h1, w1 = img1. shape
              h2, w2 = img2. shape
              def is valid(h, w, x, y):
                   if x-R<0 or x+R+1>w:
                       return False
                   if y-R<0 or y+R+1>h:
                       return False
                   return True
              for x1, y1 in corners1:
                   if not is valid(h1, w1, x1, y1):
                       continue
                   bx = 0
                   by = 0
                   bscore = 0
                   for x2, y2 in corners2:
                       if not is valid(h2, w2, x2, y2):
                           continue
                       score = ncc match (img1, img2, np. array([x1, y1], np. int64), np. array([x2, y2], np. int
                       if bscore < score:
                           bscore = score
                           bx = x2
                           by = y2
                   if bscore > NCCth:
                       matching. append (([x1, y1], [bx, by]))
              return matching
```

```
In [123]: # detect corners on warrior and matrix sets
# you are free to modify code here, create your helper functions, etc.

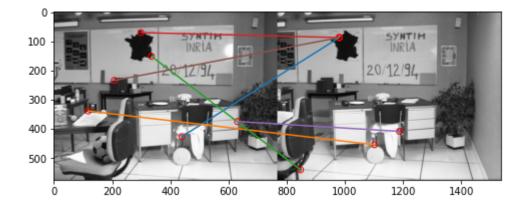
nCorners = 50  # do this for 10, 20 and 30 corners
smoothSTD = 1
windowSize = 17

# read images and detect corners on images

imgs_sport = []
crns_sport = []
for i in range(2):
    img_sport = imageio.imread('Sport' + str(i) + '_0GO.bmp')
    imgs_sport.append(rgb2gray(img_sport))
    # downsize your image in case corner_detect runs slow in test
# imgs_mat.append(rgb2gray(img_mat)[::2, ::2])
    crns_sport.append(corner_detect(imgs_sport[i], nCorners, smoothSTD, windowSize))
```

```
[126]:
         # plot matching result
         def show matching result (img1, img2, matching):
             YOUR CODE HERE
              _____ """
             img = np. hstack([img1, img2])
             h1, w1 = img1. shape
             h2, w2 = img2. shape
             print (img. shape)
             fig = plt.figure(figsize=(8, 8))
             ax = fig. add subplot()
             X = \begin{bmatrix} 1 \end{bmatrix}
             y = []
             for node1, node2 in matching:
                  x. append (node1[0])
                  y. append (node1[1])
                  x. append (node2\lceil 0 \rceil + w1)
                  y. append (node2[1])
                  ax. plot ([node2[0]+w1, node1[0]], [node2[1], node1[1]])
             ax. imshow(img, cmap='gray')
             ax.scatter(x, y, s=35, edgecolors='r', facecolors='none')
         print("Number of Corners:", nCorners)
         show matching result(imgs sport[0], imgs sport[1], matching sport)
```

Number of Corners: 50 (576, 1536)



# Problem 4.4 Matching using epipolar geometry [10 pts]

Next, we will use the epipolar geometry constraint on the rectified images and updated corner points to build a better matching algorithm. First, detect 10 corners in image1. Then, for each corner, do a line search along the corresponding parallel epipolar line in image2.

Evaluate the NCC score for each point along this line and return the best match (or no match if all scores are below the NCCth). R is the radius (size) of the NCC patch in the code below.

You do not have to run this in both directions. Show your result as in the naive matching part.

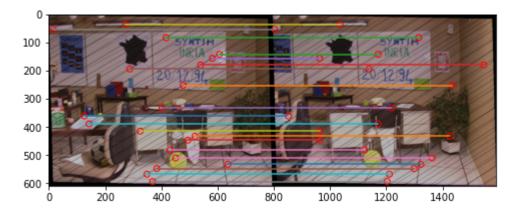
```
In [125]:
           def display correspondence (img1, img2, corrs):
               """Plot matching result on image pair given images and correspondences
               Args:
                   img1: Image 1.
                   img2: Image 2.
                   corrs: Corner correspondence
               """ ======
               YOUR CODE HERE
               You may want to refer to the `show_matching_result` function.
               img = np.hstack([img1, img2])
               h1, w1, = img1. shape
               h2, w2, = img2. shape
               fig = plt.figure(figsize=(8, 8))
               ax = fig. add subplot()
               X = []
               y = []
               for node1, node2 in corrs:
                   x. append (node1[0])
                   y. append (node1[1])
                   x. append (node2[0]+w1)
                   y. append (node2[1])
                   ax. plot([node2[0]+w1, node1[0]], [node2[1], node1[1]])
               ax.imshow(img, cmap='gray')
               ax.scatter(x, y, s=35, edgecolors='r', facecolors='none')
           def correspondence matching epipole(img1, img2, corners1, R, NCCth):
               """Find corner correspondence along epipolar line.
               Args:
                   img1: Image 1.
                   img2: Image 2.
                   corners1: Detected corners in image 1.
                   F: Fundamental matrix calculated using given ground truth corner correspondences.
                   R: NCC matching window radius.
                   NCCth: NCC matching threshold.
                   Matching result to be used in display_correspondence function
               """ _____
               YOUR CODE HERE
               _____ """
               h2, w2, = img2. shape
               matching = []
               for x, y in corners1:
                   if x < R or x >= w2 + R - 1 or y < R or y >= h2 + R - 1:
                       continue
                   bscore = 0
```

```
bx = 0
by = 0
for i in range(R, w2-R-1,R):
    score = ncc_match(rgb2gray(img1), rgb2gray(img2), np.array([x,y],np.int64), np
    if score > bscore:
        bscore = score
        bx = i
        by = y
if bscore < NCCth:
    matching.append(([x,y],[bx,by]))</pre>
return matching
```

```
In [45]: rectified_im1_final, rectified_im2_final = completelyBoundedRectification(src1, src2, H1_bounded, H2_bounded, min_x1_bounded, max_x1_bounded, min_y1_bounded, max_y1_bounded, max_y1_bounded)
```

```
In [135]: # replace black pixels with white pixels
    _black_idxs = (rectified_iml_final[:, :, 0] == 0) & (rectified_iml_final[:, :, 1] == 0) &
    rectified_iml_final[:, :][_black_idxs] = [1.0, 1.0, 1.0]
    _black_idxs = (rectified_im2_final[:, :, 0] == 0) & (rectified_im2_final[:, :, 1] == 0) &
    rectified_im2_final[:, :][_black_idxs] = [1.0, 1.0, 1.0]

nCorners = 20
# Choose your threshold and NCC matching window radius
smoothSTD = 1
windowSize = 7
NCCth = 0.9
R = 8
# detect corners using corner detector here, store in corners1
corners1 = corner_detect(rgb2gray(rectified_im1_final), nCorners, smoothSTD, windowSize)
corrs = correspondence_matching_epipole(rectified_im1_final, rectified_im2_final, corners1
display_correspondence(rectified_im1_final, rectified_im2_final, corrs)
```



In [ ]: