

## Creating Image Mosaics using Homographies (Due Monday, 2/4/2019)

In this assignment, you will extend the field of view of a camera by forming a panorama mosaic from multiple photographs.

When images are taken with same optical center, they can be aligned with a homography transformation. To estimate the correct warping, you'll need at least 4 pairs of corresponding points in the overlap region. You'll mark these points manually (later in the quarter we will discuss some methods for finding such points automatically without user input).

For the assignment you will submit two files, (1) a completed version of this python notebook "2\_panorama.ipynb" which contains the python script that calls your functions and displays various inputs and outputs, and (2) a pdf printout of your python notebook. As in the previous assignment, you can generate the pdf by executing your whole notebook and then printing to a pdf from your web browser.

The source images for alignment are available via Canvas in the assignment directory. In addition I have provided a module **selectpoints.py** which contains some code for capturing user clicks. You should download all three files and unzip them into the same directory as this notebook.

**To start with, please enter your name and student ID # in the cell below**

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In [1]:

```
#tell ipython that we are running in a notebook.
#this is necessary for the interactive user interface in part 1 below
%matplotlib notebook

#import a few key modules, these should be sufficient to complete the assignment
import numpy as np
import matplotlib.pyplot as plt
from scipy.interpolate import griddata
from scipy.ndimage import gaussian_filter
import pickle

#import the provided functions to support point selection in the notebook
import selectpoints
```

## 1. Point Correspondence [5 pts]

The first image in each example provided is the central image. It's simplest to construct a mosaic from a central image and a set of peripheral images, since we then need to find just one homography for each peripheral image. You should flesh out the code below to allow the user to manually select at least 4 pairs of corresponding points between each central and peripheral image. These points should be located on distinctive locations that you can easily identify between the images such as high contrast corners.

Write code below which loads in the central image and then loops over the remaining images and for each image allows you to select four or more points. Save out the resulting points to a pckl file which you will include as part of your upload. You should encapsulate the basic procedure of getting the point correspondences in a function called **get\_correspondences** to make it easier to run on different sets of images to produce your final results

I have provided some code which takes care of getting user clicks in the file **selectpoints.py**. It takes as input the plot axis and the number of points you want from the user. As you click, the points are numbered so that you can make sure points correspond in the two images you are trying to align.

```
sp = select_k_points(ax,npaints)
```

Once you have finished marking **npaints** locations in the plot, you can access their coordinates via the fields **sp.xs** and **sp.ys**

NOTES:

1. Make sure you read through the code in **selectpoints.py** so you understand how it works before proceeding.
2. In an interactive application you would need to write code to wait until the user finished clicking before accessing **sp.xs**, **sp.ys** but here in the notebook environment we will just split the code into two different cells and only execute the second one after clicking on all the points
3. If you are running into memory limits, you should downsample the images when you load them in (e.g. using use **scipy.ndimage.zoom**) to make things more manageable and run faster.

In [2]:

```

from scipy.ndimage.interpolation import zoom
def get_correspondences(imnames,baseim,npoints=4):

"""
The function loads in a set of images which are going to be assembled
in to a mosaic. It displays the central (base) image along with each
peripheral image and allows the user to click to select corresponding
points between the base image and the peripheral image. It returns the
images along with the user input. If the original image files were in
color the function converts them to

Parameters
-----
imnames : list of str
    Filenames of image files that are going in to the mosaic

baseim : int
    An index which specifies which of the image files is the base image

npoints : int
    How many points are required from the user. Defaults to 4

Returns
-----
imgs : list of 2D float arrays
    The arrays for the corresponding images given in imnames. These
    are gray scale images represented as floats.

pointmatches : list of SelectPoints objects
    Returns an object whose fields xs and ys contain the point
    coordinates once the user has clicked (see selectpoints.py)

"""

nimages = len(imnames)

#loop over images and load in each one and convert to grayscale
imgs = list()
for fname in imnames:
    print('loading...',fname)
    I = plt.imread(fname)
    #
    #convert to float data type if necessary
    if (I.dtype == np.uint8):
        I = I.astype(float) / 256
    #convert from color to grayscale if necessary
    if len(I.shape) != 2:
        I = np.dot(I[...,:3], [0.299, 0.587, 0.114])
    #finally, store the array in our list of images
    I = zoom(I, 0.15)
    imgs.append(I)

#loop over each pair of overlapping images and have the user
#click to specify corresponding points
pointmatches = list()
for i in range(nimages):
    if (i==baseim):
        continue

```

```
fig = plt.figure()

#select points in base image
ax1 = fig.add_subplot(2,1,1)
ax1.imshow(imgs[baseim],cmap=plt.cm.gray)
sp1 = selectpoints.select_k_points(ax1,npoints)

#corresponding points in overlapping image
ax2 = fig.add_subplot(2,1,2)
ax2.imshow(imgs[i],cmap=plt.cm.gray)
sp2 = selectpoints.select_k_points(ax2,npoints)

pointmatches.append((sp1,sp2))

return imgs,pointmatches
```

The following three cells demonstrate the functionality of your **get\_correspondences** function. As noted above, they should be executed one at a time. First execute the first cell which should display two pairs of images and get the user clicks. Once you have finished clicking, execute the second cell which grabs the point coordinates where the user clicked and saves the results out to disk. The third cell demonstrates loading the data back in from disk and visualizing the points again. Once you have finished implementing **get\_correspondences** you should execute these cells including the final visualization step so that the results are visible in your saved notebook pdf.

In [5]:

```
imnames = ('atrium/IMG_1347.JPG','atrium/IMG_1348.JPG','atrium/IMG_1349.JPG')
baseim = 0    #index of the central base image that we will align everything to

#make the default figure size larger to make clicking points easier
#feel free to adjust this to suit your monitor / display size
plt.rcParams['figure.figsize'] = [9, 9]

#call your function to get user clicks
imgs,pointmatches = get_correspondences(imnames,baseim)

#reduce the figure size a bit for visualizations later in the notebook
plt.rcParams['figure.figsize'] = [9, 9]

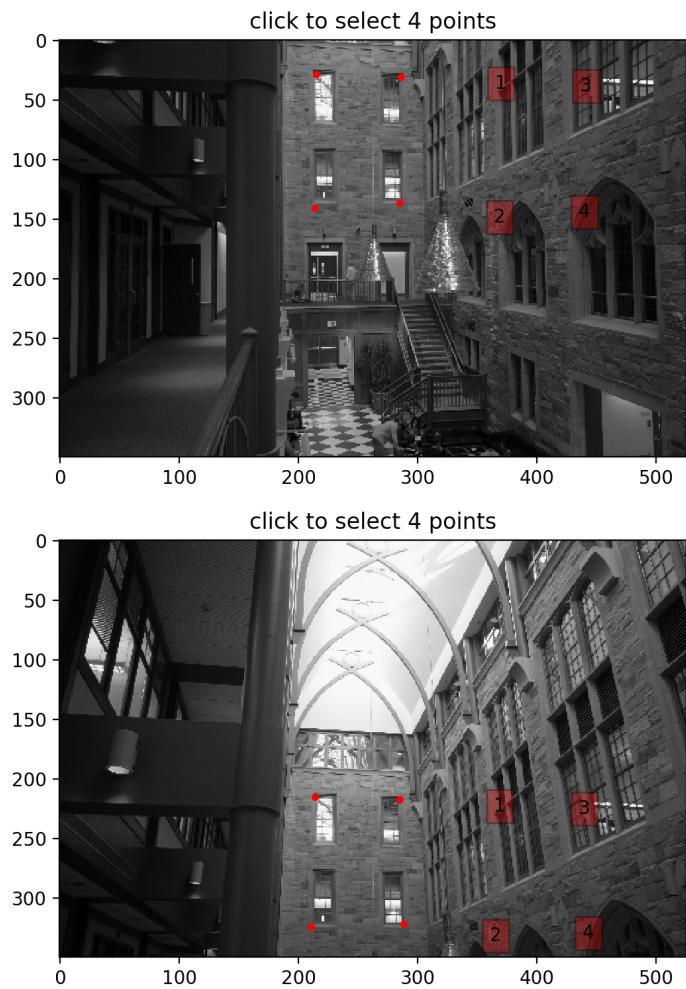
# after executing this cell, we need to stop and click before running to the next cell
```

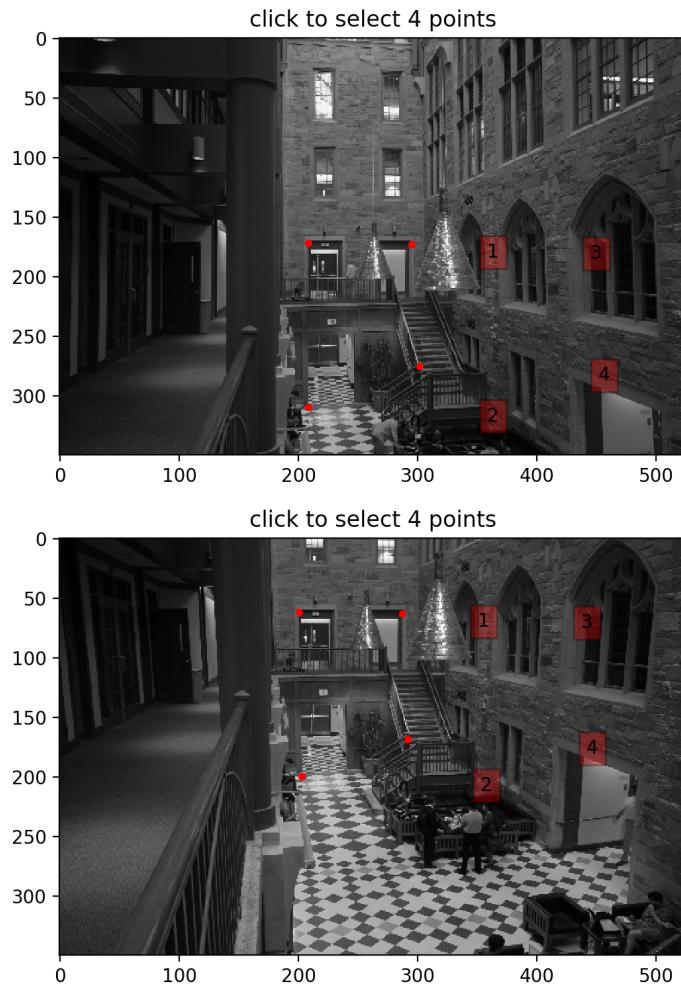
loading... atrium/IMG\_1347.JPG

/Users/alexhu/anaconda/lib/python3.6/site-packages/scipy/ndimage/interpolation.py:605: UserWarning: From scipy 0.13.0, the output shape of zoom() is calculated with round() instead of int() - for these inputs the size of the returned array has changed.  
"the returned array has changed.", UserWarning)

loading... atrium/IMG\_1348.JPG

loading... atrium/IMG\_1349.JPG





In [6]:

```
# collect together all the point coordinates into numpy arrays
nimages = len(imnames)
srcpts = list()
basepts = list()
for i in range(nimages-1):
    basepts.append(np.array((pointmatches[i][0].xs,pointmatches[i][0].ys)))
    srcpts.append(np.array((pointmatches[i][1].xs,pointmatches[i][1].ys)))

# save everything out to a file so we don't have to click again.
f = open('atrium_correspondeces.pkl','wb')
pickle.dump((baseim,imgs,srcpts,basepts),f)
f.close()
```

In [7]:

```
# later on we can quickly read back in the variables and visualize them
f = open('atrium_correspondeces.pkl','rb')
baseim,imgs,srcpts,basepts = pickle.load(f)
f.close()
fig = plt.figure()

#plot all the base points on the base image
ax1 = fig.add_subplot(2,2,1)
ax1.imshow(imgs[0],cmap=plt.cm.gray)
for i in range(2):
    x_l = basepts[i][0]
    y_l = basepts[i][1]
    for p in range(len(x_l)):
        ax1.scatter([x_l[p]],[y_l[p]],color = 'red',alpha = 0.5)

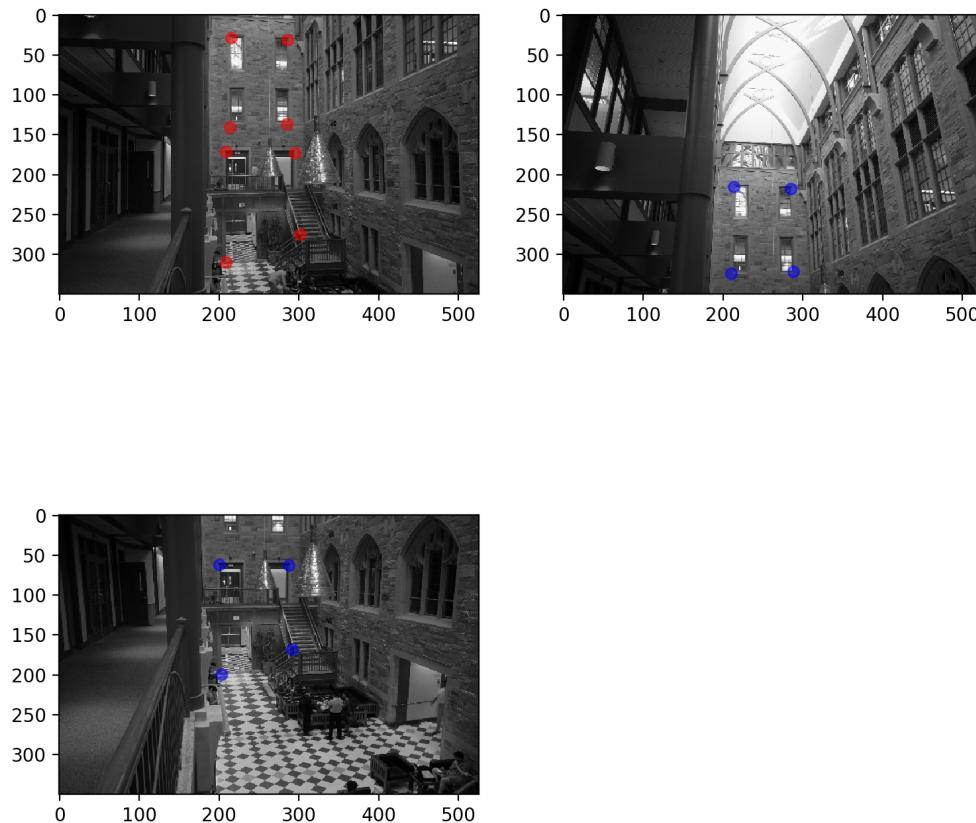
#plot the src points on the peripheral images
ax2 = fig.add_subplot(2,2,2)
ax2.imshow(imgs[1],cmap = plt.cm.gray)

ax3 = fig.add_subplot(2,2,3)
ax3.imshow(imgs[2],cmap = plt.cm.gray)

for i in range(2):
    x_l = srcpts[i][0]
    y_l = srcpts[i][1]
    if i == 0:
        for p in range(len(x_l)):
            ax2.scatter(x_l[p],y_l[p],color = 'blue', alpha = 0.5)
    elif i == 1:
        for p in range(len(x_l)):
            ax3.scatter(x_l[p],y_l[p],color = 'blue', alpha = 0.5)

## your code here

plt.show()
```



## 2. Homography Transformations [20 pts]

For each image, we will need to compute the homography (3x3 transformation matrix) using linear least squares. This transformation should map the points in the peripheral image that you clicked to their corresponding points in the base "central" image. For the central image itself, this transformation would just be the identity matrix.

For this part you should write two functions. The first function, **compute\_homography** should estimate a transformation matrix  $H$  given the pairs of points. The second function, **apply\_homography** should take as input an array of point coordinates and a 3x3 matrix and return the transformed coordinates.

Note that if the matrix  $H$  maps  $(x_1, y_1)$  to  $(x_2, y_2)$ , then the inverse mapping is given simply by inverting the matrix. So applying the homography

`inv(H)` will map  $(x_2, y_2)$  back to  $(x_1, y_1)$ .

You will want to use `np.linalg.lstsq` to solve for the coefficients of  $H$ .

In [8]:

```

def compute_homography(pts1,pts2):
    """
    Computes the 3x3 transformation matrix (homography) that when applied to pts1 yields pts2.

    Parameters
    -----
    pts1 : 2xN array
        2D coordinates of points (to warp)

    pts2 : 2xN array
        target 2D coordinates where points should end up

    Returns
    -----
    numpy.array (dtype=float)
        3x3 array containing the matrix H
    """

#expected dimensions of input
assert(pts1.shape[0]==2)
assert(pts2.shape[0]==2)
assert(pts1.shape[1]>=4)
assert(pts2.shape[1]>=4)

x = pts1[0]
y = pts1[1]
X = pts2[0]
Y = pts2[1]
A = np.zeros(shape=(2*len(x),8))
L = list()
for i in range(len(x)):
    a = np.array([x[i],y[i],1])
    b = np.array([0,0,0])
    c = np.array([-x[i]*X[i],-y[i]*X[i]])
    d = np.array([-x[i]*Y[i],-y[i]*Y[i]])
    A[i*2:i*2+1, :3] = a
    A[i*2:i*2+1, 3:6] = b
    A[i*2:i*2+1, 6:] = c
    A[i*2+1:i*2+2, :3] = b
    A[i*2+1:i*2+2, 3:6] = a
    A[i*2+1:i*2+2, 6:] = d
    L.append(X[i])
    L.append(Y[i])

pts2_array = np.array(L)
H = np.linalg.solve(A,pts2_array)
H = np.insert(H,len(H),1)

H = H.reshape(3,3)
#
#your code goes here
#
#expected dimension of output
assert (H.shape==(3,3))

```

```
return H
```

In [9]:

```
def apply_homography(H,pts):
    """
    Apply a specified homography H to a set of 2D point coordinates

    Parameters
    -----
    H : 3x3 array
        matrix describing the transformation

    pts : 2xN array
        2D coordinates of points to transform

    Returns
    -----
    numpy.array (dtype=float)
        2xN array containing the transformed points
    """

#assert expected dimensions of input
assert(H.shape==(3,3))
assert(pts.shape[0]==2)
assert(pts.shape[1]>=1)
x_l = list()
y_l = list()
for i in range(len(pts[0])):
    x = pts[0][i]
    y = pts[1][i]
    X = ((H[0][0]*x)+(H[0][1]*y) + H[0][2])/((H[2][0]*x)+(H[2][1]*y+1))
    Y = ((H[1][0]*x)+(H[1][1]*y) + H[1][2])/((H[2][0]*x)+(H[2][1]*y+1))
    x_l.append(X)
    y_l.append(Y)

x_a = np.array(x_l)
y_a = np.array(y_l)
tpts = np.vstack((x_a,y_a))
#your code goes here

#make sure transformed pts are correct dimension
assert(tpts.shape[0]==2)
assert(tpts.shape[1]==pts.shape[1])

return tpts
```

## Testing

You should enter some test cases in the cell below to verify that your code is working as expected. I've given a couple examples but you should fill out the others and add your own.

In [10]:

```
# since there may be some numerical imprecision, specify a small error tolerance
tol = 1e-07

### testing apply homography

#
# If you apply the identity transformation you should get back the same points
#
pts1 = np.random.random((2,4))
H = np.array([[1.,0,0],[0,1,0],[0,0,1]])
pts2 = apply_homography(H,pts1)
err = np.mean(np.abs(pts1-pts2))
assert(err<tol)

#
# If you apply H to some points and then inv(H) to the results, you should get back
#

pts3 = np.random.random((2,4))#source points(points be applied H)
pts4 = np.random.random((2,4))#destinata point
H = compute_homography(pts3,pts4)
pts5 = apply_homography(H,pts3)#after apply H
pts6 = apply_homography(np.linalg.inv(H),pts5)
err1 = np.mean(np.abs(pts6-pts5))
assert(err1-tol)

### testing computeHomography

#
# If you compute the homography between a set of points and itself, you should get I
#

pts7 = np.random.random((2,4))
H = np.array([[1.,0,0],[0,1,0],[0,0,1]])
IDM = compute_homography(pts7,pts7)
err2 = np.mean(np.abs(IDM-H))
assert(err2-tol)

#
# If you compute the homography between a set of points and those points multiplied I
# back a matrix which looks like a simple scaling (i.e. 2 2 1 on the diagonal)
#
pts1 = np.array([[0.,0,1,1],[0.,1,1,0]])
pts2 = 2*pts1
H = compute_homography(pts1,pts2)
Htrue = np.array([[2.,0.,0.],[0.,2.,0],[0.,0.,1]])
err = np.mean(np.abs(H-Htrue))
assert(err<tol)

#
# If you generate two random sets of points and compute the homography between them,
# estimated homography to the first set of points, you should get back the second se
#

pts9 = np.random.random((2,4))
pts10 = np.random.random((2,4))

H = compute_homography(pts9,pts10)
```

```
pts11 = apply_homography(H,pts9)
err = np.mean(np.abs(pts11-pts10))
assert(err<tol)
```

## 3. Warping [20pts]

Implement a function called **warp\_images** which takes the collection of correspondences and generates warped versions of all the input images to align them with the final mosaic.

1. We will use the central image's coordinate system for the final mosaic. You first need to figure out how big the final mosaic will be. You can accomplish this by determining where the corners of each source image will be mapped to in the final mosaic (using your **apply\_homography** function) and then use min/max to determine the left-most, right-most, top-most and bottom-most points across all of the warped images. After this step you will have determined that all the warped image pixels from all the images will fall inside some rectangular region  $(xmin,ymin)-(xmax,ymax)$ . Note that these coordinates will be expressed with respect to your central image. For example,  $xmin$  will be a negative value if some of the images in your mosaic are mapped to the left of your central image.
2. Generate the coordinates of all the pixels for your final mosaic as well as the coordinates of pixels in each source image (I suggest using **np.mgrid** for this). To get the warped image coordinates, apply the estimated homography to the source image pixel coordinates to determine where they will fall in the output mosaic.
3. To produce the warped image, we will use **scipy.interpolate.griddata** to perform interpolation gray values onto a regular grid. We need to provide **griddata** with three pieces of information: the coordinates of the pixels after we have warped them with the appropriate homography, the gray value for each of those pixels, and the grid of pixel coordinates for our final mosaic. We will ultimately call this warping function for each source image, resulting in a new warped image the size of the final mosaic containing. By default, **griddata** will set the value of any pixels that are outside the source image to NaN. The figure below shows examples of warped images where the white pixels correspond to regions outside the source image (i.e. filled with NaNs).

In [11]:

```
def warp_images(imgs,baseim,srcpts,basepts):
    """
    Warp a collection of images to align them with a base image using
    provided point correspondences to estimate the warping.

    Parameters
    -----
    imgs : list of 2D numpy.array (dtype=float)
        The arrays containing the input grayscale images

    baseim : int
        The index of the image which is the base image

    srcpts: list of 2D numpy.array, each of size 2xNclicks
        x,y coordinates of points clicked by the user for each peripheral image

    basepts : list of 2D numpy.array, each of size 2xNclicks
        x,y coordinates of corresponding locations in the base image

    Returns
    -----
    imgs_warped: list of 2D numpy.array (dtype=float)
        list of the resulting warped images, all of which will have the same shape
    """

# first estimate all the homographies
homs = list()
base_check = False
for i in range(len(imgs)):
    if (i == baseim):
        H = np.array([[1.,0,0],[0,1,0],[0,0,1]])#identity matrix
        base_check = True
    else:
        if base_check == True:
            H = compute_homography(srcpts[i-1],basepts[i-1]) #compute homography
        else:
            H = compute_homography(srcpts[i],basepts[i])
    homs.append(H)

# compute where the corners of each warped image end up
# and the upper left and lower-right corners of a rectangle
# that will enclose all of them

xul_min = 99999999
yul_min = 99999999
xlr_max = -99999999
ylr_max = -99999999
for i in range(len(imgs)):
    #original corner coordinates
    h,w = imgs[i].shape
    cx = np.array([0,w,0,w])
    cy = np.array([0,0,h,h])
    cc = np.vstack((cx,cy))
    #warped coordinates
    cx_warped = apply_homography(homs[i],cc)
    xul = np.min(cx_warped[0])
    yul = np.min(cx_warped[1])
```

```

xlr = np.max(cx_warped[0])
ylr = np.max(cx_warped[1])
if xul < xul_min:
    xul_min = xul
if yul < yul_min:
    yul_min = yul
if xlr > xlr_max:
    xlr_max = xlr
if ylr > ylr_max:
    ylr_max = ylr
#round these to integer valued coordinates
xul_min = np.floor(xul_min)
yul_min = np.floor(yul_min)
xlr_max = np.ceil(xlr_max)
ylr_max = np.ceil(ylr_max)
ul = np.array([xul_min,yul_min])
lr = np.array([xlr_max,ylr_max])

# for debugging purposes, you may find it very useful at
# this point to use plt.plot to plot out the warped
# corner locations and the bounding rectangle in the
# code above.

# generate the coordinates of pixels in the output image that
# cover the bounding rectangle determined above
ogrid_y,ogrid_x = np.mgrid[ul[1]:lr[1],ul[0]:lr[0]]

# loop over each input and warp it
imgs_warped = list()
for i in range(len(imgs)):
    #input pixel grid
    igrid_y,igrid_x = np.mgrid[...]
    h,w = imgs[i].shape
    igrid_y,igrid_x = np.mgrid[0:h,0:w]
    # reshape igrid_y and igrid_x into a 2xN array so we can use
    # our applyHomography function on them. you can do
    # this with some combination of flatten() and stack()
    gy = igrid_y.flatten()
    gx = igrid_x.flatten()

    #incoords = ....
    incoords = np.vstack([gx,gy])

    # transform pixel coordinates
    warpedcoords = apply_homography(homs[i],incoords)

    # note: griddata expects coords to be Nx2 and values to be Nx1 so
    # make sure you transpose / flatten as necessary

    # brightness values reshaped as a Nx1 array
    ##bvalues =
    ##wcoords =
    wcoords = np.transpose(warpedcoords)
    bvalues = imgs[i].flatten()
    # use griddata to warp
    #img_warped = griddata(....)
    img_warped = griddata(wcoords,bvalues,(ogrid_x,ogrid_y))
    imgs_warped.append(img_warped)

return imgs_warped

```

## 4. Blending [35 pts]

Now that you have generated the individual warped images, we need to blend them together into the final mosaic image. The simplest approach is to paste down the pixels from each warped image in some order into the output image. However, as we discussed in class this can lead to bad artifacts. Instead you should create a smooth blend between the images in the regions where they overlap. To receive full credit, you must implement some type of weighted blending.

To create a blend, first compute an alpha mask for each image which is the same size as the target output with 1s where you have values from that image. You can start by using `np.isnan` to get a boolean mask for each warped image that tells you which pixels are valid and which are invalid. In order to turn the binary mask into an alpha mask, you can feather the edges by blurring them with a Gaussian filter. I suggest using `scipy.ndimage.gaussian_filter`. Since more than one image can overlap at a given location, the final step you will need to carry out is to normalize these alpha maps by the sum of the alphas across all images at that location.

To create the Gaussian filter, `gaussian_filter` takes an argument sigma which specifies the width of the Gaussian. You will need to experiment with the parameter sigma in order to get good feathering of the edges. I recommend visualizing the alpha masks as images so you can see what they look like.

NOTE: If you simply blur the binary alpha mask for an image, you will end up with non-zero values outside of the support of a warped image which will cause problems when you composite them (e.g., black bands around the edge of the image). To get a good result, you will need to find a way to fix this so that the alpha is non-zero only where you actually have color values for a given warped image.

In [12]:

```

import copy
def blend_images(imgs_warped):
    """
    Blend together a collection of warped images into a mosaic
    using alpha masks to smooth transitions between individual photos

Parameters
-----


```

```
#You will need to modify this code to deal with the fact that imgs_warped
#contains NaNs. Your output mosaic should not contain any NaNs, just 0s
#in the places which are not covered by any input image
for i in range(len(imgs_warped)):
    img_w = copy.deepcopy(imgs_warped[i])
    is_nan = np.isnan(img_w)
    img_w[is_nan] = 0
    mosaic = mosaic + (alphas[i]*img_w)

return mosaic, alphas
```

## 5. Results [20pts]

Now you need to put together all the pieces. In the space below, write code which assembles mosaics and visualizes the results in the notebook using your functions.

For the two provided example image sets, visualize (1) the individual source images after they have been warped but before they are blended together into the final mosaic, (2) the computed alpha maps, and (3) the final mosaic.

In addition, show one mosaics of your own creation, each with at least 3 source images. You can use your own camera to take images. Remember that you want to keep the camera center in the same location and simply rotate the camera. For your writeup below, show also the warped source images before they are blended into each final mosaic.

NOTES:

1. If you are running into memory limitations, you should downsample the images when you first load them in (but before you start clicking correspondences).
2. For the two provided image sets, you do not need to upload them. Just write your code so that it assumes that you have unzipped them as folders in the subdirectory containing your notebook.
3. You do not need to include the images you used to make your mosaic, you just need to make sure that they appear visible in the pdf version of your notebook.

### 5.1 Atrium Result

In [13]:

```
imnames = ('atrium/IMG_1347.JPG','atrium/IMG_1348.JPG','atrium/IMG_1349.JPG')
baseim = 0    #index of the central base image that we will align everything to

#make the default figure size larger to make clicking points easier
#feel free to adjust this to suit your monitor / display size
plt.rcParams['figure.figsize'] = [9, 9]

#call your function to get user clicks
imgs,pointmatches = get_correspondences(imnames,baseim)

#reduce the figure size a bit for visualizations later in the notebook
plt.rcParams['figure.figsize'] = [9, 9]

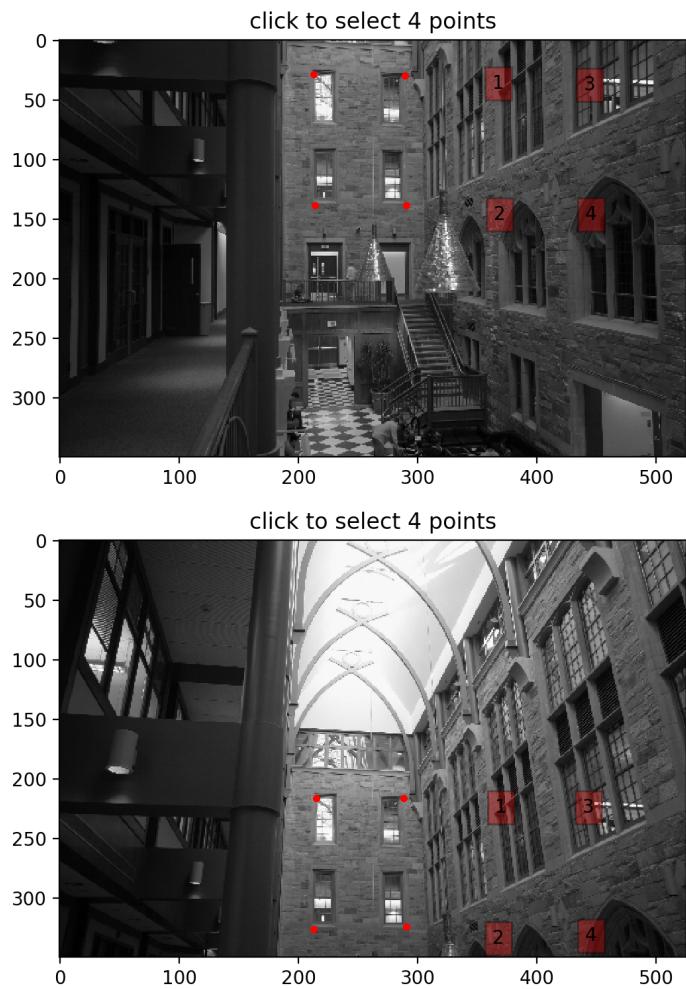
# after executing this cell, we need to stop and click before running to the next cell
```

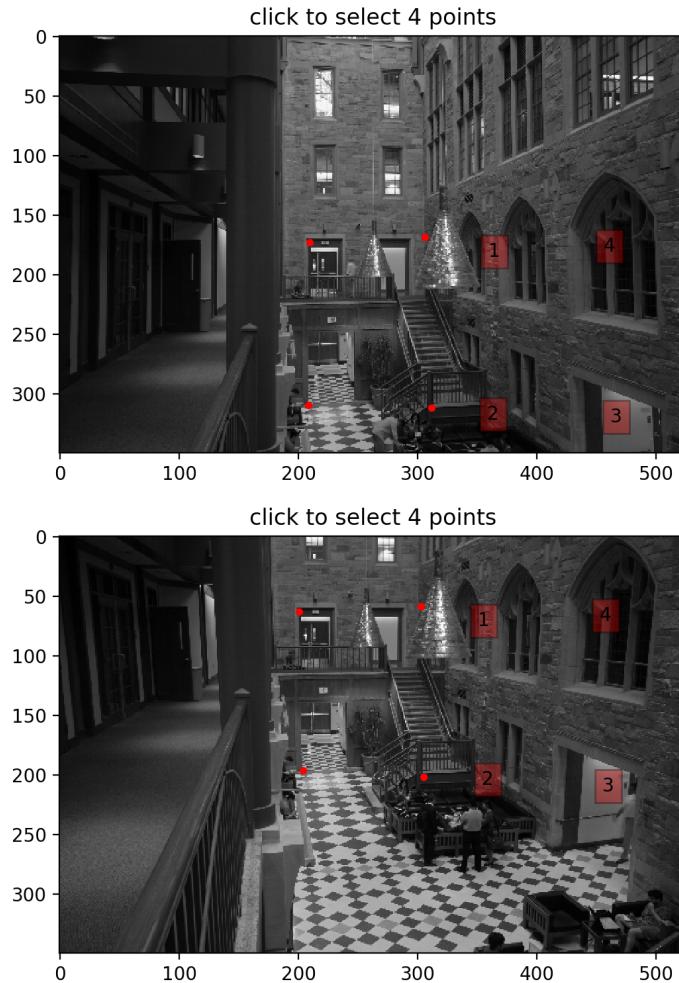
loading... atrium/IMG\_1347.JPG

/Users/alexhu/anaconda/lib/python3.6/site-packages/scipy/ndimage/interpolation.py:605: UserWarning: From scipy 0.13.0, the output shape of zoom() is calculated with round() instead of int() - for these inputs the size of the returned array has changed.  
"the returned array has changed.", UserWarning)

loading... atrium/IMG\_1348.JPG

loading... atrium/IMG\_1349.JPG





In [14]:

```
# collect together all the point coordinates into numpy arrays
nimages = len(imnames)
srcpts = list()
basepts = list()
for i in range(nimages-1):
    basepts.append(np.array((pointmatches[i][0].xs,pointmatches[i][0].ys)))
    srcpts.append(np.array((pointmatches[i][1].xs,pointmatches[i][1].ys)))
imgs_warped = warp_images(imgs,baseim,srcpts,basepts)

mosaic,alphas = blend_images(imgs_warped)

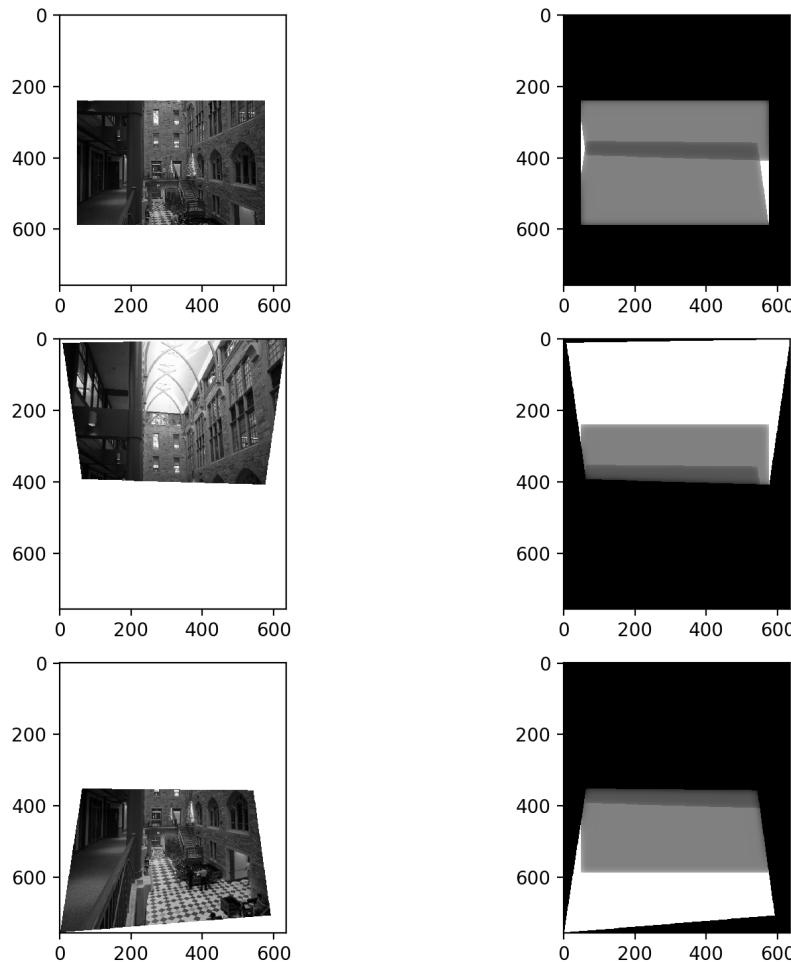
# add code here to display imgs_warped, alphas and the final mosaic in the notebook
```

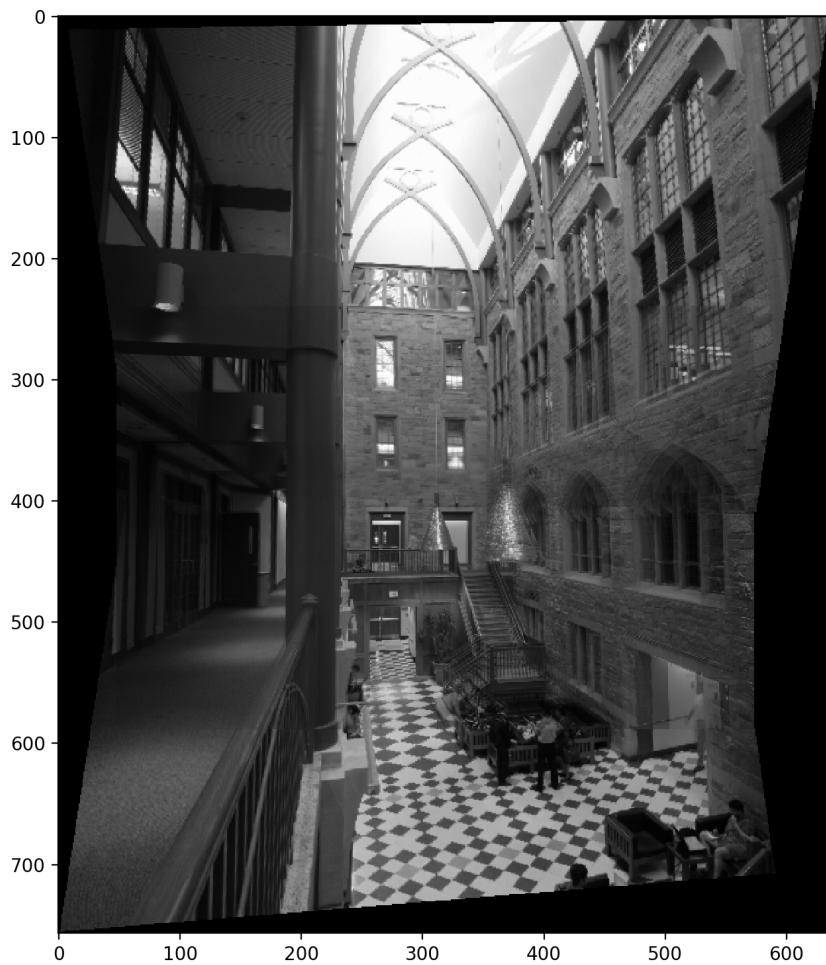
```
/Users/alexhu/anaconda/lib/python3.6/site-packages/ipykernel_launcher.py:52: RuntimeWarning: invalid value encountered in true_divide
```

In [15]:

```
###imgs_warped and alphas
fig = plt.figure()
for i in range(nimages):
    fig.add_subplot(nimages,2,2*i+1).imshow(imgs_warped[i],cmap=plt.cm.gray)
    fig.add_subplot(nimages,2,2*i+2).imshow(alphas[i],cmap=plt.cm.gray)
###final mosaic
fig,ax = plt.subplots()
ax.imshow(mosaic,cmap=plt.cm.gray)

plt.show()
```





## 5.2 Window Result

In [16]:

```
imnames = ('window/IMG_1341.JPG','window/IMG_1342.JPG','window/IMG_1343.JPG','window/IMG_1344.JPG','window/IMG_1345.JPG')
baseim = 0 #index of the central base image that we will align everything to

#make the default figure size larger to make clicking points easier
#feel free to adjust this to suit your monitor / display size

plt.rcParams['figure.figsize'] = [9, 9]

#call your function to get user clicks
imgs,pointmatches = get_correspondences(imnames,baseim)

#reduce the figure size a bit for visualizations later in the notebook
plt.rcParams['figure.figsize'] = [9, 9]
# after executing this cell, we need to stop and click before running to the next cell
```

loading... window/IMG\_1341.JPG

```
/Users/alexhu/anaconda/lib/python3.6/site-packages/scipy/ndimage/interpolation.py:605: UserWarning: From scipy 0.13.0, the output shape of zoom() is calculated with round() instead of int() - for these inputs the size of the returned array has changed.
```

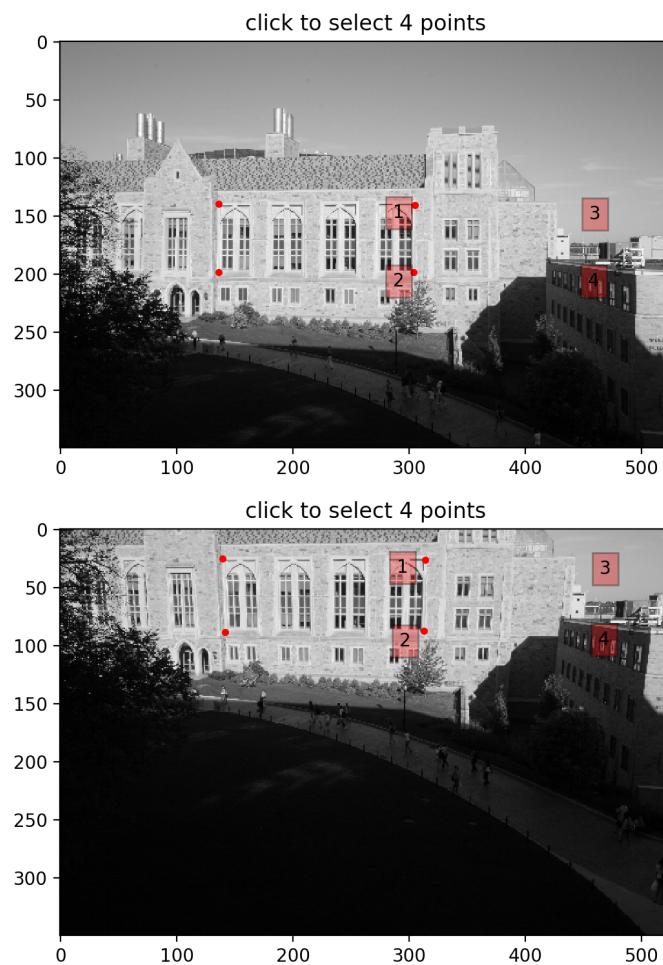
"the returned array has changed.", UserWarning)

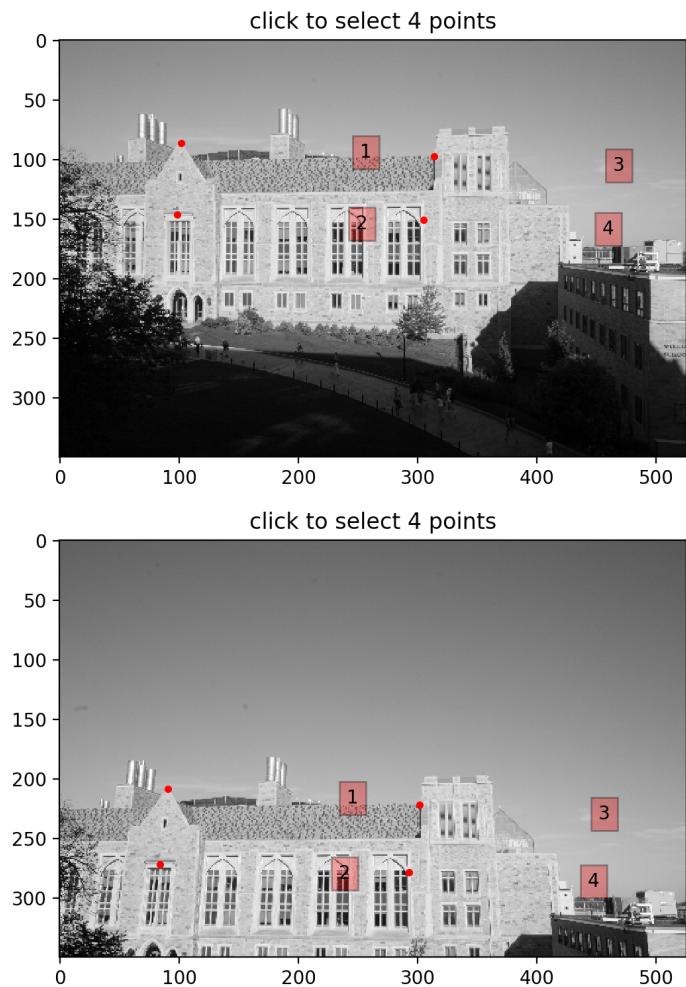
loading... window/IMG\_1342.JPG

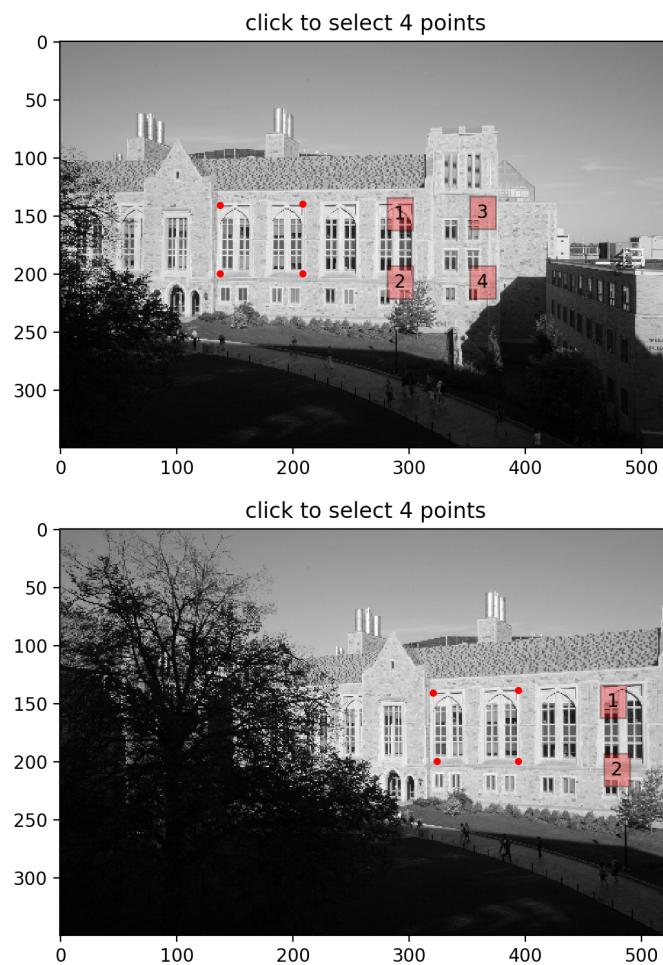
loading... window/IMG\_1343.JPG

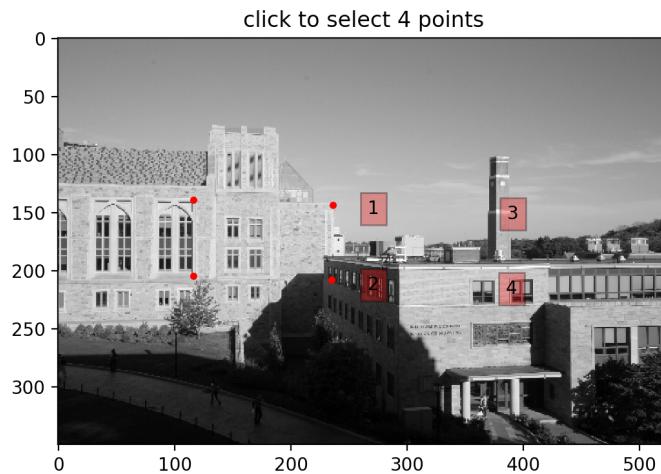
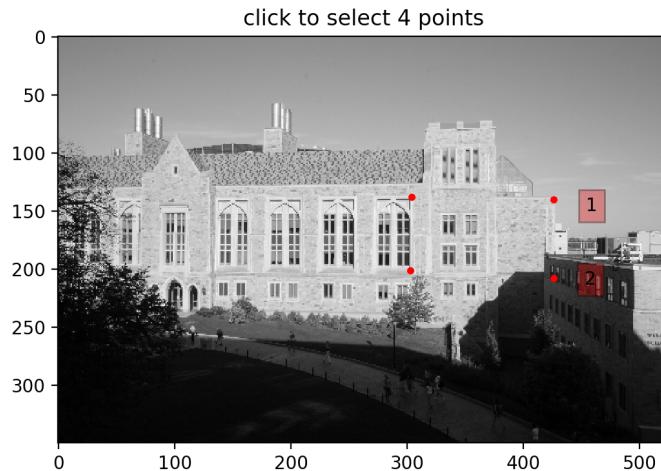
loading... window/IMG\_1344.JPG

loading... window/IMG\_1345.JPG









In [17]:

```
# collect together all the point coordinates into numpy arrays
nimages = len(imnames)
srcpts = list()
basepts = list()
for i in range(nimages-1):
    basepts.append(np.array((pointmatches[i][0].xs, pointmatches[i][0].ys)))
    srcpts.append(np.array((pointmatches[i][1].xs, pointmatches[i][1].ys)))
imgs_warped = warp_images(imgs, baseim, srcpts, basepts)

mosaic, alphas = blend_images(imgs_warped)

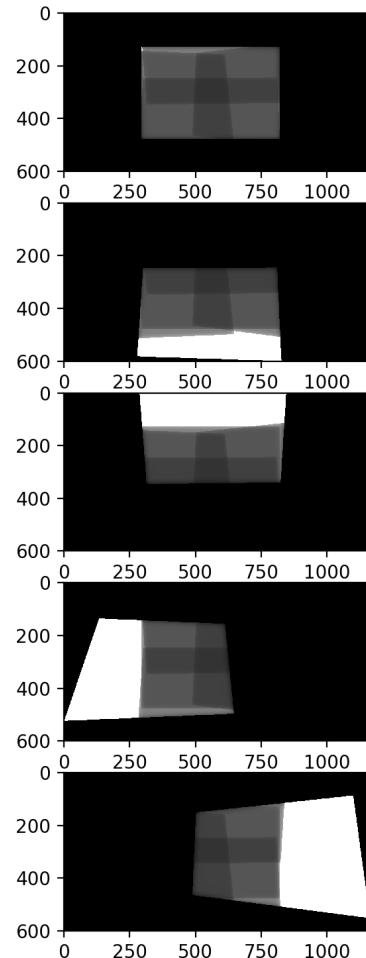
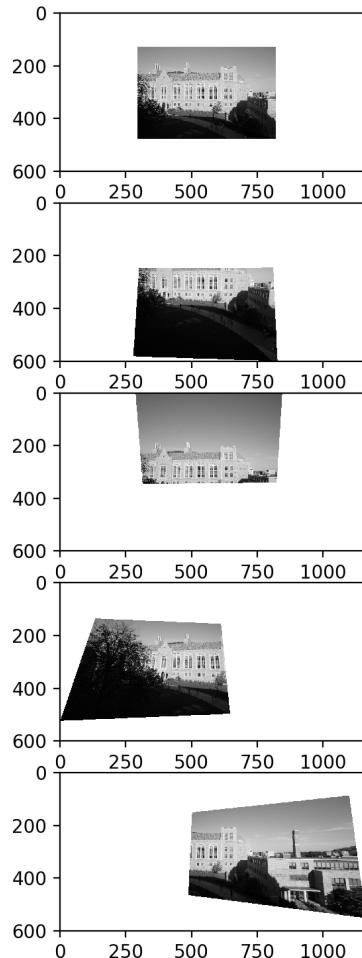
# add code here to display imgs_warped, alphas and the final mosaic in the notebook
```

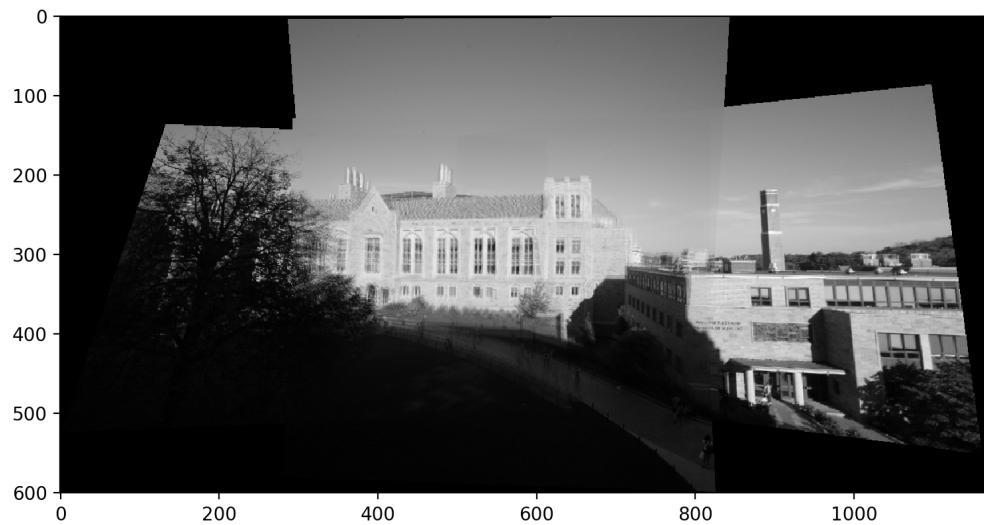
/Users/alexhu/anaconda/lib/python3.6/site-packages/ipykernel\_launcher.py:52: RuntimeWarning: invalid value encountered in true\_divide

In [18]:

```
###imgs_warped and alphas
fig = plt.figure()
for i in range(nimages):
    fig.add_subplot(nimages,2,2*i+1).imshow(imgs_warped[i],cmap=plt.cm.gray)
    fig.add_subplot(nimages,2,2*i+2).imshow(alphas[i],cmap=plt.cm.gray)
###final mosaic
fig,ax = plt.subplots()
ax.imshow(mosaic,cmap=plt.cm.gray)

plt.show()
```





## 5.3 Result on your own photos

In [19]:

```
imnames = ('1.JPG','2.JPG','3.JPG')
baseim = 0 #index of the central base image that we will align everything to

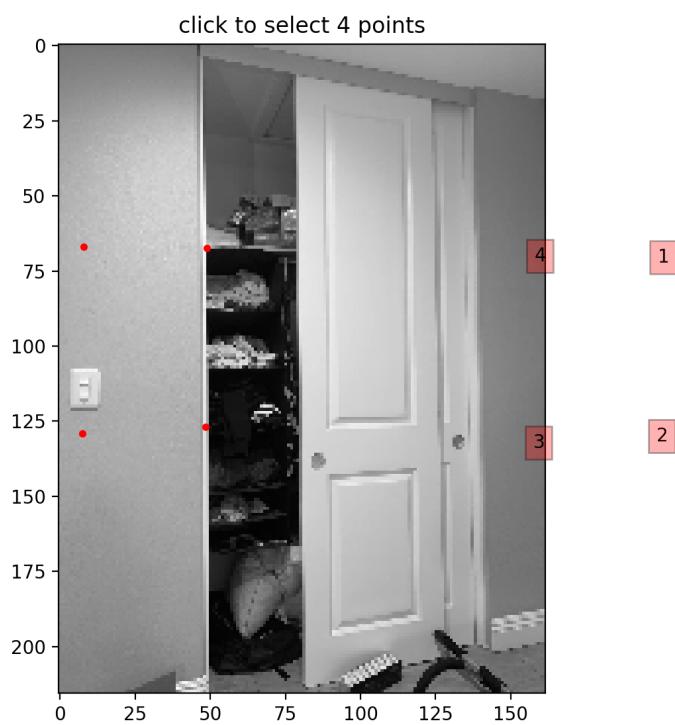
#make the default figure size larger to make clicking points easier
#feel free to adjust this to suit your monitor / display size

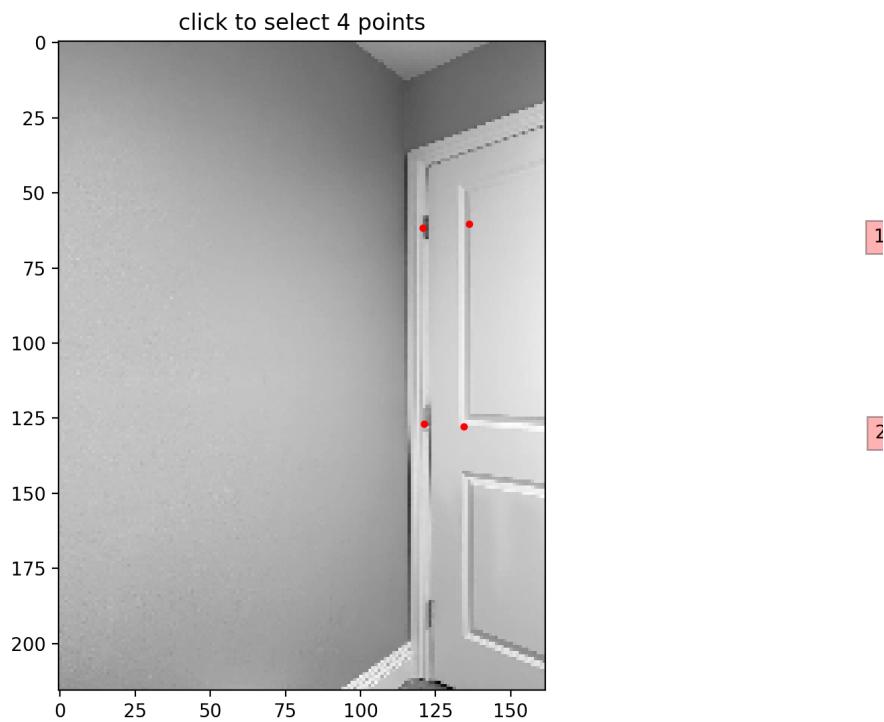
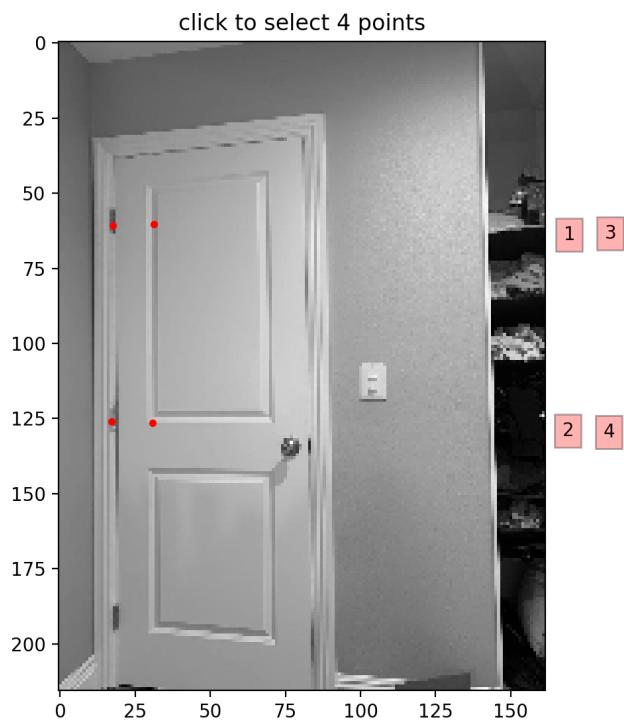
plt.rcParams['figure.figsize'] = [9, 14]

#call your function to get user clicks
imgs,pointmatches = get_correspondences(imnames,baseim)

#reduce the figure size a bit for visualizations later in the notebook
plt.rcParams['figure.figsize'] = [9, 9]
# after executing this cell, we need to stop and click before running to the next cell
```

loading... 1.JPG  
loading... 2.JPG  
loading... 3.JPG





In [20]:

```
# collect together all the point coordinates into numpy arrays
nimages = len(imnames)
srcpts = list()
basepts = list()
for i in range(nimages-1):
    basepts.append(np.array((pointmatches[i][0].xs,pointmatches[i][0].ys)))
    srcpts.append(np.array((pointmatches[i][1].xs,pointmatches[i][1].ys)))
imgs_warped = warp_images(imgs,baseim,srcpts,basepts)

mosaic,alphas = blend_images(imgs_warped)

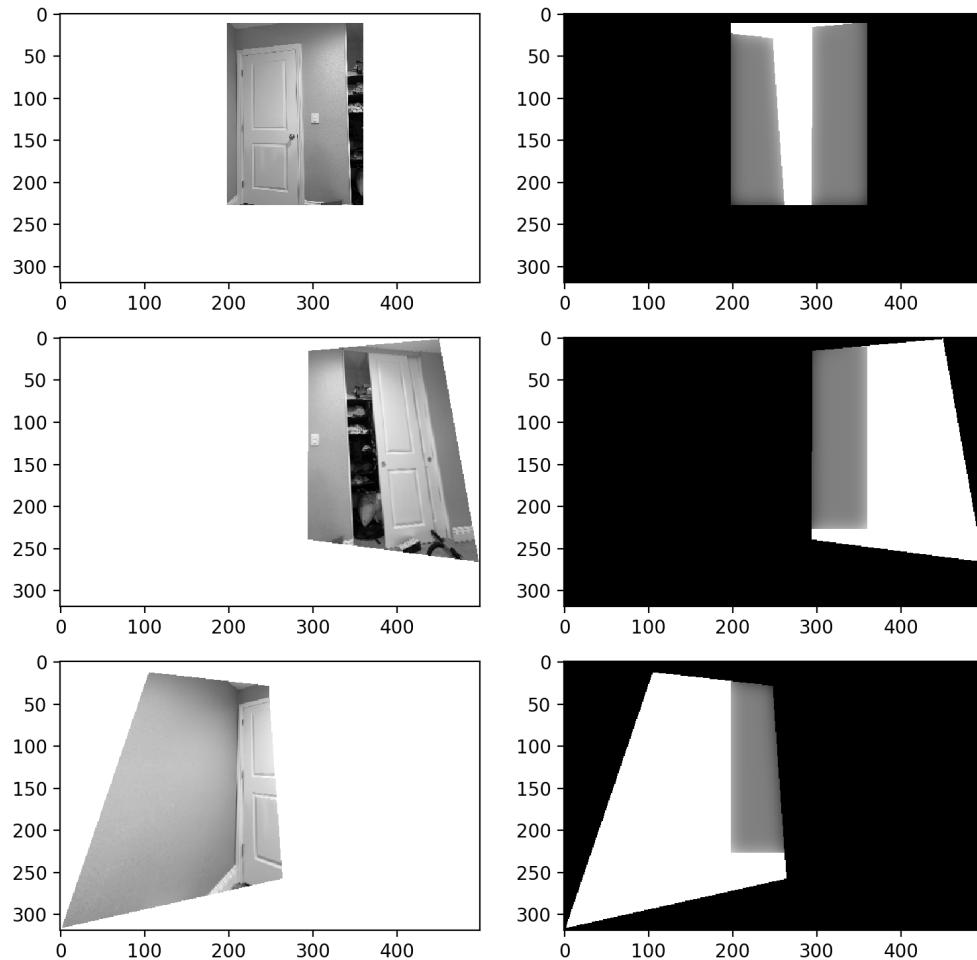
# add code here to display imgs_warped, alphas and the final mosaic in the notebook
```

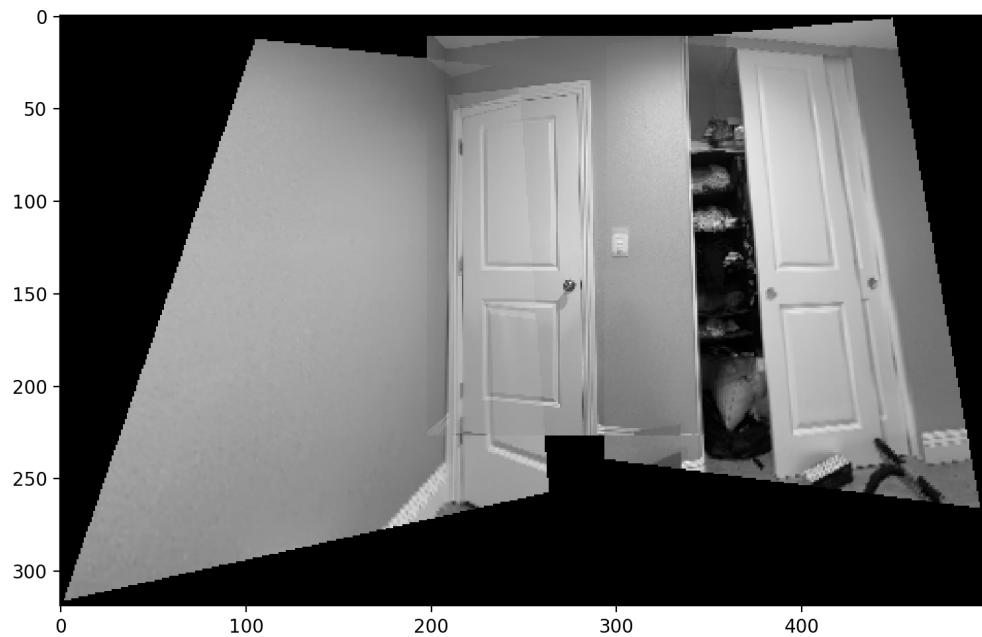
/Users/alexhu/anaconda/lib/python3.6/site-packages/ipykernel\_launcher.py:52: RuntimeWarning: invalid value encountered in true\_divide

In [21]:

```
###imgs_warped and alphas
fig = plt.figure()
for i in range(nimages):
    fig.add_subplot(nimages,2,2*i+1).imshow(imgs_warped[i],cmap=plt.cm.gray)
    fig.add_subplot(nimages,2,2*i+2).imshow(alphas[i],cmap=plt.cm.gray)
###final mosaic
fig,ax = plt.subplots()
ax.imshow(mosaic,cmap=plt.cm.gray)

plt.show()
```





In [ ]: