

Occlusion Detection

Importing Libraries

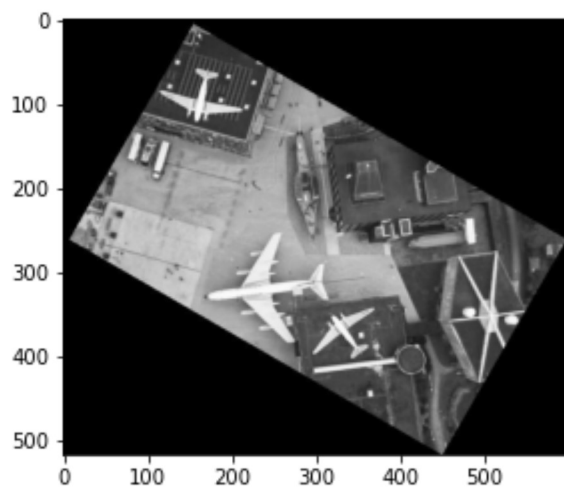
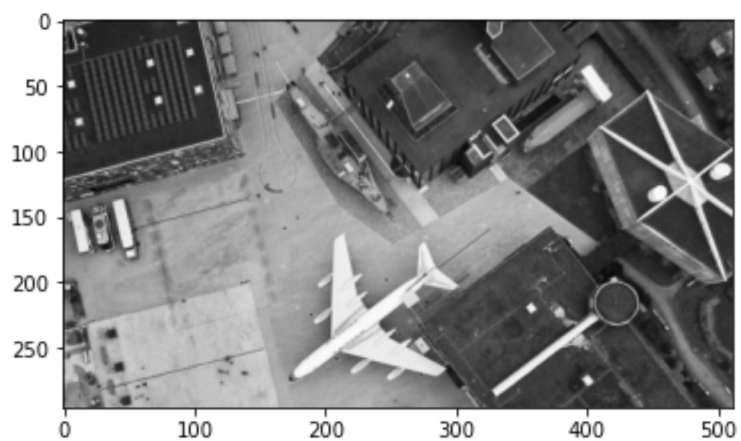
```
In [1]: #%matplotlib inline
import numpy as np
from matplotlib import pyplot as plt
import cv2
import imageio
import SimpleITK
import sys
from pylab import *
```

Reading and Visualising the Images

```
In [2]: img1=cv2.imread('IMG1.png',0)
img2=cv2.imread('IMG2.png',0)

plt.imshow(img1,cmap='gray')
plt.show()

plt.imshow(img2,cmap='gray')
plt.show()
```



Corresponding Points

```
In [3]: points = np.array([[29, 124], [157, 372]])
        corresponding_points = np.array([[93, 248], [328, 399]])
```

Bilinear Interpolation

```
In [4]: def bilinear_interpolation(src, si, sj):
        #si , sj = src_pt
        #si= ti-ty
        #sj= tj-tx
        si,sj=si+1,sj+1
        i=int(np.floor(si))
        j=int(np.floor(sj))      ## Here i,j are the co-ordinate points of the top left
        t_l = i , j

        ##Now the remaining three co-ordinates with respect to i,j will be
        t_r = i , j+1      # Top right
        b_l = i+1 , j      # Bottom Left
        b_r = i+1 , j+1    # Bottom Right

        ## distance of source point from the top left corner would be
        di = si - i
        dj = sj - j

        """
        ## Now calculating the pixel value at the source point by using bilinear interpolation
        ## Create a variable pxl_val and assign the pixel value obtained by bilinear interpolation
        ## b_l,b_r
        ## di,dj that we got is used to obtain the weights for interpolation
        ## We ignore all the target points whose source points lies outside the source image
        ## these pixel values as 0.
        """

        if t_l[0] >= np.shape(src)[0]-1 or t_l[1] >= np.shape(src)[1]-1 or t_l[0] < 0 or t_l[1] < 0:
            pxl_val = 0
        else :
            pxl_val = (1-di)*(1-dj)*src[t_l] + (1-di)*(dj)*src[t_r] + (di)*(1-dj)*src[b_l] + (di)*(dj)*src[b_r]

        return pxl_val
```

Calculating A matrix and Computing the H matrix

We know from our discussion in class that the solution for h vector is the vector corresponding to the smallest singular value of A . From the decomposition mentioned above, this is the last row of V Transpose when the singular values are in decreasing order. We then arrange the values appropriately to get the H matrix.

```
In [5]: n = len(points)  ## number of correspondences
def homography(corresponding_points, points):
    # from each point correspondance we get two values
    A = np.zeros((2*n, 5))  ## n=len(points)=2 ## A will be 4 X 5 matrix
    for r in range(n):
        i, j = points[r]
        si, sj = corresponding_points[r]
        # As defined above
        A[2*r] = [si, sj, 1, 0, -i]
        A[2*r+1] = [sj, -si, 0, 1, -j]
    # NumPy SVD gives singular values in decreasing order
    u, s, v_transpose = np.linalg.svd(A)
    # take the last row of v_transpose
    a, b, c, d, h = v_transpose[-1]
    # construct the appropriate 3x3 matrix
    H = np.array([[a, b, c],
                  [-b, a, d],
                  [0, 0, h]])

    return H
```

Transforming the image

Homogenous coordinates of the form $[a, b, c]$ can be converted to non homogenous only if $c \neq 0$. In case that condition is violated, we set the non homogenous coordinates to $(0,0)$ even though in reality such a point does not have a finite representation in non homogenous coordinates

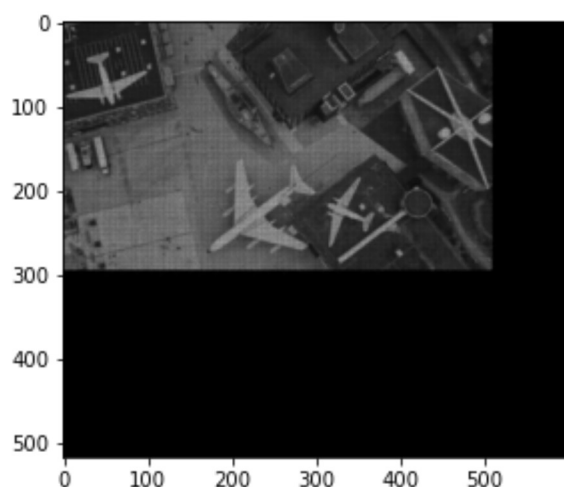
```
In [6]: def transform(src, H):
    r, c = np.shape(src)
    trg = np.zeros((r, c))
    H_inv = np.linalg.inv(H)

    ## iterating over the target image and assign all the pixel values to them
    for ti in range(r):
        for tj in range(c):
            # convert to homogenous coordinates
            t = np.array([ti, tj, 1])
            s = H_inv @ t
            if s[2] != 0:
                si, sj = s[0]/s[2], s[1]/s[2]
            else:
                si, sj = 0, 0
            # calculate the corresponding points for (ti, tj) in the source image
            # si, sj are the corresponding points for (ti, tj)
            # Assign the value using bilinear interpolation
            # Assigning the intensity values of the target image at (ti, tj) using
            trg[ti][tj] = bilinear_interpolation(src, si, sj)

    return trg
```

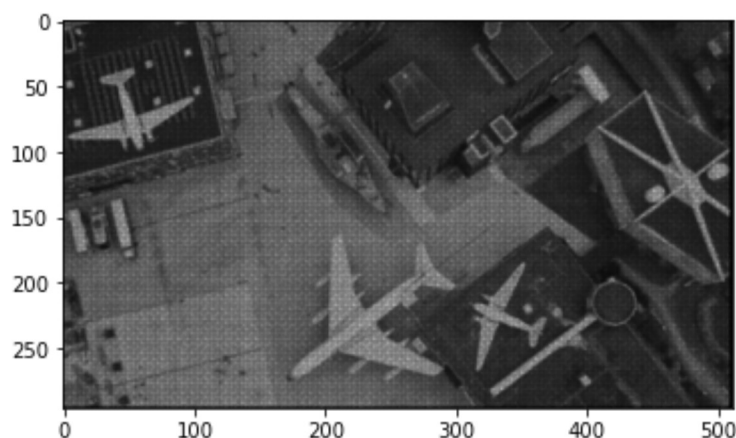
Transformed IMG2

```
In [7]: H= homography(corresponding_points,points)
#H = homography(A)
img2_new = transform(img2, H)
plt.imshow(img2_new,cmap='gray')
plt.show()
```



Cropping the transformed image to the shape of img1

```
In [8]: x,y=img1.shape
img2_new_cropped = img2_new[:x, :y]
plt.imshow(img2_new_cropped,cmap='gray')
plt.show()
```



Calculating the difference

Subtract the aligned transformed img2 and img1 to notice any changes.

```
In [9]: diff=img1-img2_new_cropped
plt.imshow(diff,cmap='gray')
plt.show()
```

