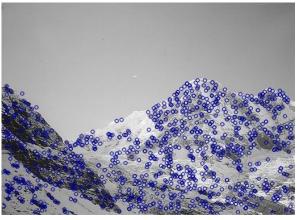
CVIP - Project 2

1 IMAGE FEATURES AND HOMOGRAPHY

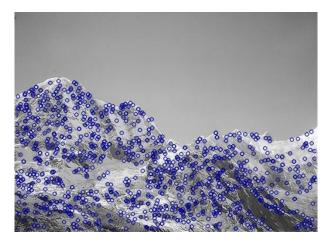
IN THIS TASK WE FIND THE IMAGE FEATURES AND TRY TO WARP THE TWO IMAGE WITH HOMOGRAPHY. ANY TWO IMAGES OF THE SAME PLANAR SURFACE IN SPACE ARE RELATED BY A HOMOGRAPHY

1.1

SIFT (Scale Invariant Feature Transform) is one of the widely used feature detector and discriptors. In this task we extract the SIFT features for the given two images. The SIFT function is called with the OpenCV libraries and source. After the detect and compute function call, the keypoints and descriptors are stored. Those points are drawn on the image. The output is shown below



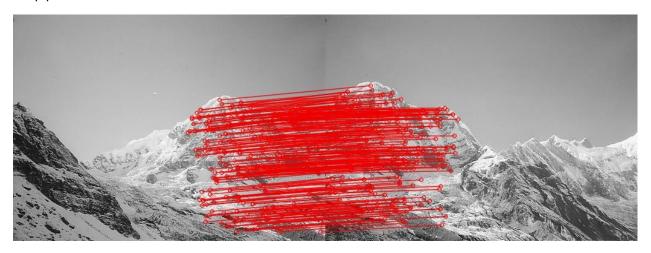
SIFT - Mountian1



SIFT - Mountain2

1.2 MATCH IMAGE

The keep points are matched with k means clustering of the both the images and the matching keep points are drawn.



Matching image with k-means

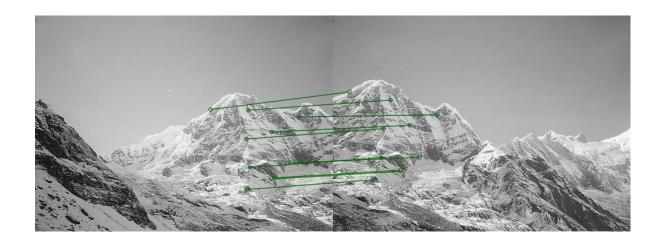
1.3 Homography Matrix

The homography matrix with first image to the second image.

```
[ 1.58930230e+00 -2.91559040e-01 -3.95969265e+02 ]
[ 4.49423930e-01 1.43110916e+00 -1.90613988e+02 ]
[ 1.21265043e-03 -6.28729364e-05 1.00000000e+00 ]
```

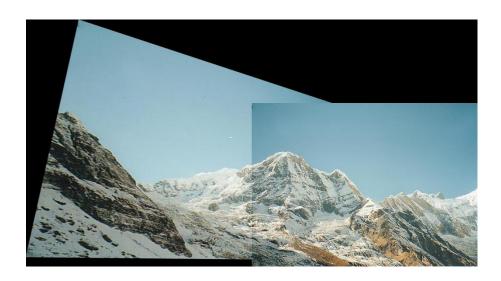
1.4 Draw keypoints for random 10 inliners

Random 10 numbers are selected in the range of 255. Only those which match with the random 10 numbers already selected are matched and selected.



1.5 WARP IMAGES

 $\hbox{`Warpperspective' function is used to warp the two images given.}\\$



```
import cv2
import numpy as np
from matplotlib import pyplot as plt
UBIT = 'jayanthk'
np.random.seed(sum([ord(c) for c in UBIT]))
import random
# Reading the images in grey scale
img1 = cv2.imread("D:\\sem1\\cvip\\proj2\\mountain1.jpg", 0)
img2 = cv2.imread("D:\\sem1\\cvip\\proj2\\mountain2.jpg", 0)
img1_color = cv2.imread("D:\\sem1\\cvip\\proj2\\mountain1.jpg", 1)
img2_color= cv2.imread("D:\\sem1\\cvip\\proj2\\mountain2.jpg", 1)
# calling the SIFT function
sift = cv2.xfeatures2d.SIFT create()
#computing the sift features for img1 and img2
(kp1, d1) = sift.detectAndCompute(img1, None)
(kp2, d2) = sift.detectAndCompute(img2, None)
#drawing keypoints
sift1 = cv2.drawKeypoints(img1, kp1, color=(150, 0, 0), outImage=np.array([]))
sift2 = cv2.drawKeypoints(img2, kp2, color=(150, 0, 0), outImage=np.array([]))
match c = 10
print("# keypoints: {}, descriptors: {}".format(len(kp1), d1.shape))
print("# keypoints: {}, descriptors: {}".format(len(kp2), d2.shape))
cv2.imwrite("task1 sift1.jpg", sift1)
cv2.imwrite("task1 sift2.jpg", sift2)
FLANN INDEX KDTREE = 0
#index parameters and search parameters with FLANN
indexp= dict(algorithm=FLANN INDEX KDTREE, trees=5)
searchp = dict(checks=50)
flann = cv2.FlannBasedMatcher(indexp, searchp)
matches = flann.knnMatch(d1, d2, k=2)
g = []
#knn matcher with the condition mentioned
for m, n in matches:
 if m.distance < 0.75 * n.distance:
    g.append(m)
# Fucntion to find inliners - random selection
if (len(g) > match_c):
 #source points
 src pt = np.float32([kp1[m.queryldx].pt for m in g]).reshape(-1, 1, 2)
 # destiantion points
 dst pt = np.float32([kp2[m.trainIdx].pt for m in g]).reshape(-1, 1, 2)
 H, mask = cv2.findHomography(src pt, dst pt, cv2.RANSAC, 5.0)
 matchesMask = mask.ravel().tolist()
 Mask1 = (mask.ravel()==1).tolist()
 #list to the random points
 list_t = []
```

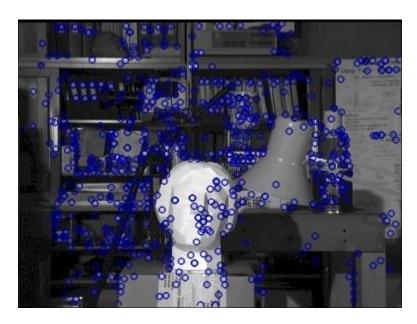
```
#randomly generated points for selecting the inliers
  for i in range(0, 10):
    x = random.randint(0, 244)
    list_t.append(x)
  for i in range(0,len(Mask1)):
    if (i in list t):
      y = 0
    else:
      Mask1[i]=False
  h, w = img1.shape
  pts = np.float32([[0, 0], [0, h - 1], [w - 1, h - 1], [w - 1, 0]]).reshape(-1, 1, 2)
  dst = cv2.perspectiveTransform(pts, H)
#function for wrapping the image - mostly made up of fucntion call
def warpImages(img1, img2, H):
  rows1, cols1 = img1.shape[:2]
  rows2, cols2 = img2.shape[:2]
  list_of_points_1 = np.float32([[0, 0], [0, rows1], [cols1, rows1], [cols1, 0]]).reshape(-1, 1, 2)
  temp_points = np.float32([[0, 0], [0, rows2], [cols2, rows2], [cols2, 0]]).reshape(-1, 1, 2)
  list_of_points_2 = cv2.perspectiveTransform(temp_points, H)
  list of points = np.concatenate((list of points 1, list of points 2), axis=0)
  [x_min, y_min] = np.int32(list_of_points.min(axis=0).ravel() - 0.5)
  [x_max, y_max] = np.int32(list_of_points.max(axis=0).ravel() + 0.5)
  translation_dist = [-x_min, -y_min]
  H translation = np.array([[1, 0, translation dist[0]], [0, 1, translation dist[1]], [0, 0, 1]])
  output_img = cv2.warpPerspective(img2, H_translation.dot(H), (x_max - x_min, y_max - y_min))
  output_img[translation_dist[1]:rows1 + translation_dist[1], translation_dist[0]:cols1 + translation_dist[0]] =
img1
  return output img
warp = warpImages(img2_color, img1_color, H)
draw params = dict(matchColor=(0, 0, 240),
          singlePointColor=None,
          matchesMask=matchesMask,
          flags=2)
params1 = dict(matchColor=(0, 100, 0),
          singlePointColor=None,
          matchesMask=Mask1,
img3 = cv2.drawMatches(img1, kp1, img2, kp2, g, None, **draw_params)
cv2.imwrite("warp.jpg", warp)
cv2.imwrite("task1_matches_knn.jpg", img3)
img4 = cv2.drawMatches(img1_color, kp1, img2_color, kp2, g, None, **params1)
cv2.imwrite("task1 matches.jpg", img4)
print("The homography matrix is")
print(H)
```

2 EPIPOLAR GEOMETRY

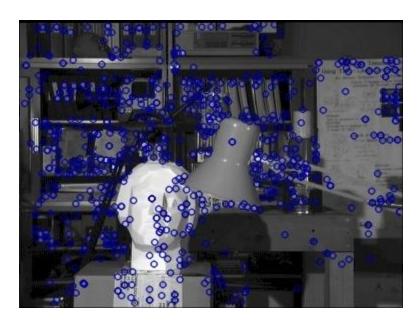
Epipolar geometry is the geometry of stereo vision. When two cameras view a 3D scene from two distinct positions, there are a number of geometric relations between the 3D points and their projections onto the 2D images that lead to constraints between the image points. These relations are derived based on the assumption that the cameras can be approximated by the pinhole camera model.

2.1 SIFT AND KNN FOR THE GIVEN IMAGE

The task 1.1 and 1.2 is repeated for both the images given, which gives us two SIFT images and one match image.



SIFT 1 Image



SIFT 2 Image



2.2 FUNDAMENTAL MATRIX

Fundamental Matrix with RANSAC.

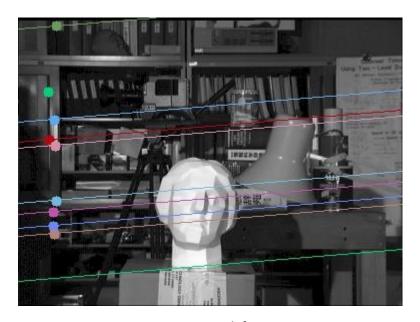
[2.58938824e-06 3.00178724e-05 2.05540959e-02]

[2.36594625e-06 1.58586353e-05 2.80750023e-01]

[-2.40895761e-02 -2.87750686e-01 1.00000000e+00]

2.3 RANDOM 10 INLINERS

Randomly 10 inliners have been selected and is drawn on the image and is shown in figure below. The color of both the lines are not same as the colors have been randomized.



Epi image left



Epi image right

2.4 DISPARITY IMAGE

Disparity refers to the distance between two corresponding points in the left and right image of a stereo pair. The Disparity map is computed with the two images using libraries and the output is shown as below. The image below which is written using imwrite is shown as pic1 and the image plotted with pyplot is pic2. The program also plots the disparity image



Pic1



Pic 2

```
import cv2
import numpy as np
from matplotlib import pyplot as plt
UBIT = 'jayanthk'
np.random.seed(sum([ord(c) for c in UBIT]))
import random
img1 = cv2.imread("D:\\sem1\\cvip\\proj2\\tsucuba left.png", 0)
img2 = cv2.imread("D:\\sem1\\cvip\\proj2\\tsucuba_right.png", 0)
sift = cv2.xfeatures2d.SIFT create()
(kp1, d1) = sift.detectAndCompute(img1, None)
(kp2, d2) = sift.detectAndCompute(img2, None)
sift img1 = cv2.drawKeypoints(img1, kp1, color=(150,0,0), outImage=np.array([]))
sift img2 = cv2.drawKeypoints(img2, kp2, color=(150,0,0), outImage=np.array([]))
print("# keypoints: {}, descriptors: {}".format(len(kp1), d1.shape))
print("# keypoints: {}, descriptors: {}".format(len(kp2), d2.shape))
cv2.imwrite("D:\\sem1\\cvip\\proj2\\task2_sift1.jpg", sift_img1)
cv2.imwrite("D:\\sem1\\cvip\\proj2\\task2_sift2.jpg", sift_img2)
FLANN INDEX KDTREE = 0
index_params = dict(algorithm = FLANN_INDEX KDTREE, trees = 5)
search params = dict(checks = 50)
flann = cv2.FlannBasedMatcher(index_params, search_params)
matches = flann.knnMatch(d1,d2,k=2)
g = []
for m,n in matches:
 if m.distance < 0.75*n.distance:
    g.append(m)
src_pt = np.float32([ kp1[m.queryldx].pt for m in g ])
dst pt = np.float32([kp2[m.trainIdx].pt for m in g])
F, mask = cv2.findFundamentalMat(src_pt, dst_pt, cv2.RANSAC,5.0)
matchesMask = mask.ravel().tolist()
draw_params = dict(matchColor = (150,0,0), # draw matches in green color
          singlePointColor = None,
          matchesMask = matchesMask,
          flags = 2
img3 = cv2.drawMatches(img1,kp1,img2,kp2,g,None,**draw params)
cv2.imwrite("D:\\sem1\\cvip\\proj2\\task2_matches_knn.jpg", img3)
FLANN INDEX KDTREE = 0
index_p = dict(algorithm = FLANN_INDEX_KDTREE, trees = 5)
search_p = dict(checks = 50)
flann = cv2.FlannBasedMatcher(index p, search p)
matches = flann.knnMatch(d1,d2,k=2)
print("Fundamental Matrix ")
print(F)
stereoMatcher = cv2.StereoBM_create()
stereoMatcher.setMinDisparity(16)
```

```
stereoMatcher.setNumDisparities(112)
stereoMatcher.setBlockSize(17)
stereoMatcher.setSpeckleRange(32)
stereoMatcher.setSpeckleWindowSize(120)
stereo = stereoMatcher.compute(img1, img2)
cv2.imwrite("D:\\sem1\\cvip\\proj2\\task2 disparity.jpg", stereo)
#showing plot of the disparity
plt.imshow(stereo)
src pt = np.int32(src pt)
dst_pt = np.int32(dst_pt)
pts1 = src_pt
pts2 = dst_pt
pts1 = pts1[mask.ravel()==1]
pts2 = pts2[mask.ravel()==1]
def drawlines(img1,img2,lines,pts1,pts2):
  r,c = img1.shape
  img1 = cv2.cvtColor(img1,cv2.COLOR_GRAY2BGR)
  img2 = cv2.cvtColor(img2,cv2.COLOR GRAY2BGR)
  for r,pt1,pt2 in zip(lines,pts1,pts2):
    color = tuple(np.random.randint(0,255,3).tolist())
    x0,y0 = map(int, [0, -r[2]/r[1]])
    x1,y1 = map(int, [c, -(r[2]+r[0]*c)/r[1]])
    img1 = cv2.circle(img1,tuple(pt1),5,color,-1)
    img2 = cv2.circle(img2,tuple(pt2),5,color,-1)
return img1,img2
list = []
for i in range(0,11):
  y = random.randint(0,271)
  list.append(y)
lines1 = cv2.computeCorrespondEpilines(pts2.reshape(-1,1,2), 2,F)
lines re = lines1.reshape(-1,3)
lines_co = np.copy(lines1)
count=0
for i in list:
  lines_co[count,:] = lines1[i,:]
line2 co = np.copy(lines11[0:10])
img5,img6 = drawlines(img1,img2,line2_co,pts1,pts2)
# epilines corresponding to points in left image (first image) and drawing its lines on right image
lines2 = cv2.computeCorrespondEpilines(pts1.reshape(-1,1,2), 1,F)
lines2 = lines2.reshape(-1,3)
lines2_co= np.copy(lines2)
count=0
for i in list:
  lines2 co[count,:] = lines2 co[i,:]
line2_rco= np.copy(lines2_co[0:10])
img3,img4 = drawlines(img2,img1,line2_rco,pts2,pts1)
```

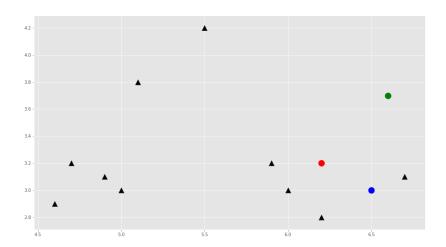
```
cv2.imwrite("D:\\sem1\\cvip\\proj2\\task2_epi1_left.jpg", img5)
cv2.imwrite("D:\\sem1\\cvip\\proj2\\task2_epi2_right.jpg", img3)
```

3 K-MEANS CLUSTERING

Implement K-Means clustering algorithm using only numpy with Euclidean distance.

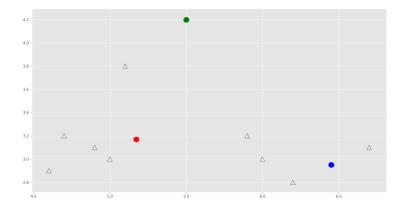
3.1 Classify N = 10 samples with the nearest meu points

The Classification vector and classification plot meu(i = 1; 2; 3).



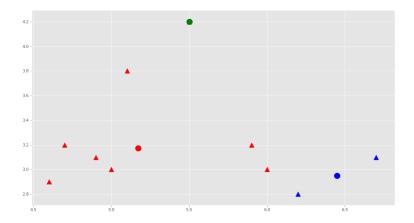
3.2 UPDATED MEU

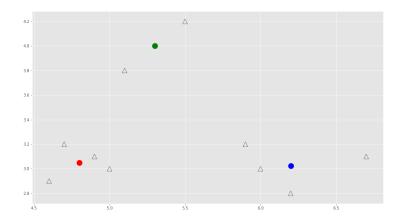
Plot with the updated MEU.



3.3 PLOTS FOR SECOND ITERATION

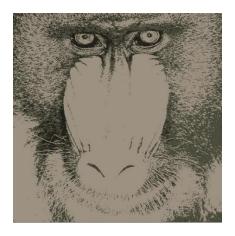
Plots after second iteration.





3.4 COLOR QUANTIZATION

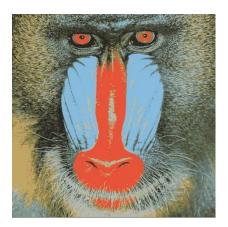
Color quantization or color image quantization is quantization applied to color spaces; it is a process that reduces the number of distinct colors used in an image, usually with the intention that the new image should be as visually similar as possible to the original image. We have applied k-means to for image quantization. The images are saved for $k = \{3,5,10,20\}$



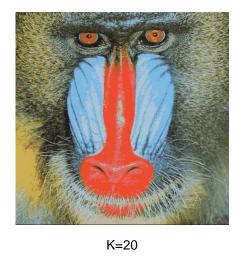
K= 3



K=5



K=10



```
import cv2
import numpy as np
from matplotlib import pyplot as plt
UBIT = 'jayanthk'
np.random.seed(sum([ord(c) for c in UBIT]))
import matplotlib.pyplot as plt
from matplotlib import style
import matplotlib
plt.rcParams['figure.figsize'] = (16, 9)
plt.style.use('ggplot')
colors = ['r', 'g', 'b', 'y', 'c', 'm']
MatchThreshold = 10
y1 = [5.9, 4.6, 6.2, 4.7, 5.5, 5.0, 4.9, 6.7, 5.1, 6.0]
y2 = [3.2, 2.9, 2.8, 3.2, 4.2, 3.0, 3.1, 3.1, 3.8, 3.0]
X = np.array(list(zip(y1, y2)))
plt.scatter(f1, f2, c='black', s=7)
# Euclidean Distance Caculator
def dist(a, b, ax=1):return np.linalg.norm(a - b, axis=ax)
# Number of clusters
k = 3
# X and Y coordinates of random centroids
C x = [6.2, 6.6, 6.5]
C_y = [3.2, 3.7, 3.0]
C = np.array(list(zip(C_x, C_y)), dtype=np.float32)
plt.scatter(f1, f2, c='#050505', marker='^', s=150)
plt.scatter(C[0, 0], C[0, 1], marker='o', s=200, c=colors[0])
plt.scatter(C[1, 0], C[1, 1], marker='o', s=200, c=colors[1])
plt.scatter(C[2, 0], C[2, 1], marker='o', s=200, c=colors[2])
plt.savefig('task3 iter1 a.jpg')
C old = np.zeros(C.shape)
clusters = np.zeros(len(X))
error = dist(C, C_old, None)
colors = ['r', 'g', 'b', 'y', 'c', 'm']
c = 0
while error != 0:
  c = c + 1
  for i in range(len(X)):
    distances = dist(X[i], C)
    cluster = np.argmin(distances)
    clusters[i] = cluster
  C_{old} = np.copy(C)
  for i in range(k):
  points = [X[j] for j in range(len(X)) if clusters[j] == i]
    C[i] = np.mean(points, axis=0)
  fig, b = plt.subplots()
  b.scatter(f1, f2, marker='^', s=150, edgecolor='black', facecolor='none')
  b.scatter(C[0, 0], C[0, 1], marker='o', s=200, c=colors[0])
  b.scatter(C[1, 0], C[1, 1], marker='o', s=200, c=colors[1])
```

```
b.scatter(C[1, 0], C[1, 1], marker='o', s=200, c=colors[1])
b.scatter(C[2, 0], C[2, 1], marker='o', s=200, c=colors[2])
plt.savefig('task3_iter{}_b.png'.format(c))
error = dist(C, C_old, None)
  fig, a = plt.subplots()
  if(c>1):
    break
  for i in range(3):
    points = np.array([X[j] for j in range(len(X)) if clusters[j] == i])
    a.scatter(points[:, 0], points[:, 1], marker='^', s=150, c=colors[i])
    a.scatter(C[0, 0], C[0, 1], marker='o', s=200, c=colors[1])
    a.scatter(C[1, 0], C[1, 1], marker='o', s=200, c=colors[2])
    a.scatter(C[2, 0], C[2, 1], marker='o', s=200, c=colors[3])
    plt.savefig('task3_iter{}_a.png'.format(c+1))
if (error == 0):
   print(C)
class KMeans custom():
       def _init_(self, num_clusters, tolerance=0.001, epoch=25, centroids={}):
    self.tolerance = tolerance
    self.epochs = epoch
    self.centroids = centroids
    self.num clusters = num clusters
  def predict(self, data):
    cl= list()
    for point in data:
       distances = [np.linalg.norm(point - self.centroids[centroid]) for centroid in self.centroids]
                                                                                                            С
cl.append(distances.index(min(distances)))
    return np.array(cl)
  def fit(self, data):
    if self.centroids == {}:
      for i in range(self.num clusters):
         self.centroids[i] = data[i]
    for i in range(self.epochs):
       self.classifications = {}
      for clu in range(self.num_clusters):
         self.classifications[clu] = []
      for f in data:
         distances = [np.linalg.norm(f - self.centroids[centroid]) for centroid in self.centroids]
         classification = distances.index(min(distances))
self.classifications[classification].append(features)
       prev centroid = dict(self.centroids)
       for classification in self.classifications:
         self.centroids[classification] = np.average(self.classifications[classification], axis=0)
       optimized = True
      for cent in self.centroids:
         original_centroid = prev_centroid[cent]
         current centroid = self.centroids.get(cent)
         if np.sum(((original_centroid - current_centroid) * 100) * original_centroid) > self.tolerance:
            opti = False
       if opti:
        break
```

```
def recreate(codebook, labels, w, h):
  img = np.zeros((w, h, 3))
  img = np.array(img, dtype=np.float64)
  label_idx = 0
  for i in range(w):
    for j in range(h):
      img[i][j] = codebook[labels[label_idx]]
      label idx += 1
  return img
def baboon(nc):
  imm = cv2.imread('D:\\sem1\\cvip\\proj2\\baboon.jpg')
  img matrix = cv2.cvtColor(imm, cv2.COLOR BGR2RGB)
  image = np.array(img_matrix, dtype=np.float64) / 255
  w, h, d = tuple(image.shape)
  image_array = np.reshape(image, (w * h, d))
  image_quantization = KMeans(num_clusters=nc, epoch=30)
  image_quantization.fit(image_array)
  lab = image_quantization.predict(image_array)
  print(image_quantization.centroids)
  imgn = recreate(image_quantization.centroids, lab, w, h)
matplotlib.image.imsave('task3_baboon_{}).png'.format(nc), imgn)
baboon(3)
```

4. REFERENCES

- 1) https://en.wikipedia.org/wiki/Color quantization
- 2) https://en.wikipedia.org/wiki/Epipolar geometry
- 3) https://opencv-python-tutroals.readthedocs.io/en/latest/py tutorials/py feature2d/py feature homography/py feature homography.html
- 4) https://mubaris.com/posts/kmeans-clustering/
- 5) https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4712040/