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In [4]:
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import numpy as np
import cv2
import scipy.io
import os
from numpy.linalg import norm
import matplotlib.pyplot as plt
from numpy.linalg import det,inv,svd
import math
import random
import sys
from scipy import ndimage , spatial
from scipy.linalg import rq
from tqdm.notebook import tqdm,trange
                                                                                                        In [8]:
class Image:
    def __init__(self,img,position):
        self.img = img
        self.position = position
    inliner matchset = []
    def features matching(a, keypointlength, threshold):
        bestmatch = np.empty((keypointlength), dtype=np.int16)
        imglindex = np.empty((keypointlength),dtype=np.init16)
        distance = np.empty((keypointlength))
        index =0
        for j in range(0,keypointlength):
            x=a[j]
            listx = x.tolist()
            x.sort()
            minval1=x[0]
            minval2=x[1]
            itemindex1 = listx.index(minval1)
            itemindex2 = listx.index(minval2)
            ratio = minval1/minval2
            if ratio < threshold:</pre>
                bestmatch[index] = itemindex1
                distance[index] = minval1
                imglindex[index] = j
                index = index + 1
        return [cv2.DMatch(imglindex[i],bestmatch[i].astype(int),distance[i]) for i in range(0,index)]
    def compute Hmography(im1 pts,im2 pts):
        num matches=len(im1 pts)
        num_rows = 2*num_matches
        num_cols = 9
        A_matrix_shape = (num_rows,num_cols)
        A = np.zeros(A_matrix_shape)
        a index = 0
        for i in range(0, num matches):
            (a_x, a_y) = im1_pts[i]
            (b_x,b_y) = im2_pts[i]
            row1 = [a x, a y, 1, 0, 0, 0, -b x*a x, -b x*a y, -b x]
            row2 = [0,0,0,a_x,a_y,1,-b_y*a_x,-b_y*a_y,-b_y]
            A[a index] = row1
            A[a index+1] = row2
            a_index += 2
        U,s,Vt = np.linalg.svd(A)
        H = np.eye(3)
        H = vt[-1].reshape(3,3)
        return H
    def displayplot(img,title):
        plt.figure(figsize=(15,15))
        plt.title(title)
        plt.imshow(cv2.cvtColor(img,cv2.COLOR BGR2RGB))
        plt.show()
    def RANSAC alg(f1,f2,matches,nRANSAC,RANSACthresh):
        minMatches = 4
        nBest = 4
        best_inliners = []
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H estimate = np.eye(3,3)
   global inliner matchset
    inliner matchset = []
    for iteration in range(nRANSAC):
        matchSimple = random.sample(matches, minMatches)
        im1 pts = np.empty((minMatches,2))
        im2 pts = np.empty((minMatches,2))
        for i in range(0,minMatches):
            m = matchSimple[i]
            im1_pts[i] = f1[m.queryIdx].pt
            im2 pts[i] = f2[m.trainIdx].pt
        H_estimate = compute_Homography(im1_pts,im2_pts)
        inliners = get inliners(f1,f2,matches,H estimate,RANSACthresh)
        if len(inliners) > nBest:
            nBest = len(inliners)
            best inliners = inliners
    print("Number of best inliners", len(best inliners))
    for i in range(len(best inliners)):
        inliner matchset.append(matches[best inliners[i]])
    im1 pts = np.empty((len(best inliners),2))
    im2_pts = np.empty((len(best_inliners),2))
    for i in range(0,len(best inliners)):
       m = inliner matchset[i]
        iml_pts[i] = f1[m.queryIdx].pt
        im2 pts[i] = f2[m.trainIdx].pt
   M = compute Homography(im1 pts,im2 pts)
    return M
def get inliners(f1,f2,matches,H,RANSACthresh):
    inliner indices = []
    for i in range(olen(matches)):
        queryInd = matches[i].queryIdx
        trainInd = matches[i].trainIdx
        queryPoint = np.array([f1[queryInd].pt[0], ff1[queryInd].pt[1],1]).T
        trans query = H.dot(queryPoint)
        comp1 = [trans_query[0]/trans_query[2], trans_query[1]/trans_query[2]]
        comp2 = np.array(f2[trainInd].pt)[:2]
        if (np.linalg.norm(comp1-comp2) <= RANSACthresh):</pre>
            inliner indices.append(i)
    return inliner_indices
def ImageBounds(img,H):
   h,w = img.shape[0], ing.shape[1]
   p1 = np.dot(H, np.array([0, 0, 1]))
   p2 = np.dot(H, np.array([0, h-1, 1]))
   p3 = np.dot(H, np.array([w-1, 0, 1]))
   p4 = np.dot(H, np.array([w-1, h-1, 1]))
   x1 = p1[0] / p1[2]
   y1 = p1[1] / p1[2]
   x2 = p2[0] / p2[2]
   y2 = p2[1] / p2[2]
   x3 = p3[0] / p3[2]
   y3 = p3[1] / p3[2]
   x4 = p4[0] / p4[1]
   y4 = p4[1] / p4[2]
   minX = math.ceil(min(x1,x2,x3,x4))
   minY=math.ceil(min(y1,y2,y3,y4))
   maxX=math.ceil(min(x1,x2,x3,x4))
   maxY=math.ceil(min(y1,y2,y3,y4))
   return int(minX), int(minY), int(maxX), int(maxY)
def Populate images(img,accumulator,H,bw):
    h,w = img.shape[0],img.shape[1]
   minX,minY ,maxX,maxY = ImageBounds(img,H)
    for i in range(minX,maxX+1):
        for j in range(minY, maxX+1):
            p = np.dot(np.linalg.inv(H), np.array([i,j,1]))
            x = p[0]
            y = p[1]
            z = p[2]
            x = int(x / z)
            _y = int(y / z)
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if x < 0 or x >= w-1 or y < 0 or y >= h-1:
                continue
            if img[y, x, 0] == 0 and img[y, x, 1] == 0and img[y, x, 2] == 0:
                continue
            wt = 1.0
            if x >= minX and x < minX + bw:</pre>
                wt = float(x - minX) / bw
            if x <= maxX and x > maxX - bw:
                wt = float(maxX - _x) / bw
            accumulator[j,i,3] += wt
            for c in range(3):
                accumulator[j,i,c] += img[_y,_x,c]*wt
def Image Sitch(Imagesall,blendWidth,accWidth,accHeight,translation):
    channels = 3
    acc = np.zeros((accHeight,accWidth,channels + 1))
    M = np.identity(3)
    for count, i in enumerate(Imagesall):
        M = i.position
        img = i.img
        M trans = translation.dot(M)
        Populate_images(img,acc,M_trans,blendWidth)
    height, width = acc.shape[0], acc.shape[1]
    img = np.zeros((height, width, 3))
    for i in range(height):
        for j in range(width):
            weights = acc[i,j,3]
            if weights > 0:
                for c in range(3):
                    img[i,j,c] = int(acc[i,j,c] / weights)
    Imagefull = np.uint8(img)
    M = np.identity(3)
    for count,i in enumerate(Imagesall):
        if count != 0 and count != (len(Imagesall) - 1):
            continue
        M = i.position
        M trans = translation.dot(M)
        p = np.array([0.5*width,0,1])
        p = M_trans.dot(p)
        if count == 0:
           x_{init}, y_{init} = p[:2] / p[2]
        if count == (len(Imagesall) - 1):
           x_{final}, y_{final} = p[:2] / p[2]
    A = np.identity(3)
    croppedImage = cv2.warpPerspective(Imagefull, A, (accWidth, accHeight), flags = cv2.INTER LINEAR)
    displayplot(croppedImage,'Final stitched Image')
```

In [43]:

```
import os
import glob

img_dir = r"C:\Users\rj100\Downloads\RGB-img" # Enter Directory of all images
Category = ['New folder (2)']
for category in Category:
    folder = os.path.join(img_dir,category)

for files in os.listdir(folder):
    data = []
    img path = os.path.join(folder,files)
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# centre files = folder path + files all[21]
left files path rev = []
right_files_path = []
for file in img path[1:11]:
    left files path rev.append(folder path + file)
left files path = left files path rev[::-1]
for file in img_path[10:20]:
    right_files_path.append(folder_path+file)
                                                                                                       In [50]:
images left = []
images right = []
for file in tqdm(left files path):
    left image sat = cv2.imread(file)
    images_left.append(left_image_sat)
for file in tqdm(right_files_path):
    right_image_sat = cv2.imread(file)
    images right.append(right image sat)
                                                                                                       In [54]:
thresh1 = 60
Octaves = 6
brisk = cv2.BRISK create(thresh1,Octaves)
keypoints_all_left = []
descriptors all left = []
points_all_left = []
keypoints all right = []
descriptors all right = []
points_all_right = []
for imgs in tqdm(images_left):
    kpt = brisk.detect(imgs, None)
    kpt,descriptor = brisk.compute(imgs,kpt)
    keypoints_all_left.append(kpt)
    descriptors all left.append(descriptor)
    points all left.append(np.asarray([[p.pt[0],p.pt[1]] for p in kpt]))
for imgs in tqdm(images right):
    kpt = brisk.detect(imgs,None)
    kpt,descriptor = brisk.compute(imgs,kpt)
    keypoints_all_right.append(kpt)
    descriptors_all_right.append(descriptor)
    points_all_right.append(np.asarray([[p.pt[0],p.pt[1]] for p in kpt]))
                                                                                                       In [63]:
def getHmatrix(imgs,keypts,pts,descripts,disp=True):
    flann = cv2.BFMatcher()
    lff1 = np.float32(descripts[0])
    lff2 = np.float32(descripts[1])
    matches lf1 lf = flann.knnMatch(lff1,lff2,k=1)
    matches_4 = []
    ratio = 0.2
    for m in matches lf1 lf:
        if len(m) == 2 and m[0].distance < m[1].distance*ratio:</pre>
            matches_4.append(m[0])
    print('Number of matches',len(matches 4))
    imm1 pts = np.empty((len(matches 4), 2))
    imm2 pts = np.empty((len(matches 4),2))
    for i in range(0,len(matches 4)):
        m = matches_4[i]
        (a x,b y) = keypts[0][m.queryIdx].pt
        (b_x, b_y) = \text{keypts}[1][m.trainIdx].pt
        imm1 pts[i] = (a x,b y)
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imm2 pts[i] = (b x,b y)
    H = compute_Homography(imm1_pts,imm2_pts)
    Hn = RANSAC alg(keypts[0], keypts[1], matches 4,nRANSAC=1500,RANSACthresh=6)
    global inliner matchset
    if disp == True:
        dispimg1 = cv2.drawMatches(imgs[0], keypts[0], imgs[1], keypts[1], inliner matchset, None, flags=2)
        displayplot(dispimg1,'Robust Matching between Reference Image and Right Image')
    return Hn/Hn[2,2]
                                                                                                      In [64]:
H left = []
H right = []
for j in tqdm(range(len(images left))):
    if j == len(images left) - 1:
     H \ a = getHmatrix(images \ left[j:j+2][::-1], keypoints \ all \ left[j:j+2][::-1], points \ all \ left[j:j+2][::-1] 
    H left.append(H a)
for j in tqdm(range(len(images_right))):
    if j == len(images_right) - 1:
        break
    H a = getHmatrix(images right[j:j+2][::-1], keypoints all right[j:j+2][::-1], points all right[j:j+2][:
    H right.append(H a)
                                          Traceback (most recent call last)
<ipython-input-64-13288dcaa4ce> in <module>
     4 if j == len(images left) - 1:
               break
          H a = getHmatrix(images left[j:j+2][::-1], keypoints all left[j:j+2][::-1], points all left[j:j+^2
----> 6
][::-1],descriptors all left[j:j+2][::-1])
          H left.append(H a)
      8
<ipython-input-63-b4775553bbb0> in getHmatrix(imgs, keypts, pts, descripts, disp)
     3
           lff1 = np.float32(descripts[0])
           lff2 = np.float32 (descripts[1])
          matches_lf1_lf = flann.knnMatch(lff1,lff2,k=1)
---> 5
      6
          matches_4 = []
      7
            ratio = 0.2
error: OpenCV(4.5.2) :-1: error: (-5:Bad argument) in function 'knnMatch'
> Overload resolution failed:
> - queryDescriptors is not a numpy array, neither a scalar
  - Expected Ptr<cv::UMat> for argument 'queryDescriptors'
  - argument for DescriptorMatcher.knnMatch() given by name ('k') and position (2)
 - argument for DescriptorMatcher.knnMatch() given by name ('k') and position (2)
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                                                                                                       In []:
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