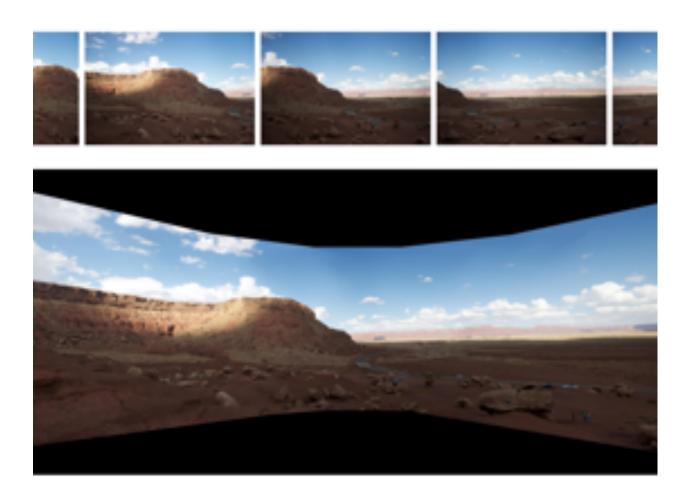
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CSE573: Project 2 - Image Stitching



The goal of this task was to experiment with image stitching methods. Given a set of photos, the program should be able to stitch them into a panoramic photo.

Algorithm for Image Stitching:

- Extract key points and descriptors from both the images.
- Find matches between the two sets of key points based on hamming distance of their descriptors.
- Using RANSAC algorithm eliminate all the outliers from the set of matching points and generate the best homography matrix.
- Finally warp one image to another using the tomography matrix obtained and stitch them.

Program Steps:

 The main function checks and adds all the image files from the specified source directory to the image queue.

```
def main():
    # Check if source directory is specified while running the script
    if len(sys.argv) < 2:
        raise ValueError("Insufficient number of arguments")

img_dir = sys.argv[1]
    image_queue = list()

# Only add images from source directory to the queue
    for image in os.listdir(os.path.abspath(img_dir)):
        if (image.lower().endswith('jpg') or image.lower().endswith('png') or image.lower().endswith('jpeg')) and image.lower()
        image_queue.append(os.path.join(os.path.abspath(img_dir), image))</pre>
```

- Two random images are picked up from the image queue and checked for a possible overlap (for images with overlap, minimum hamming distance between the descriptor bits should be less than 10).
- If they have, then use these images to proceed further else pick the next two.

```
# Select two images that can be stitched
for i in range(2):
    image1 = image_queue[i]
    image2 = image_queue[i+1]
    img1, gray1 = read_image(image1, img_dir)
    img2, gray2 = read_image(image2, img_dir)

    result_image, result_gray = stitcher(img1, img2, gray1, gray2)

if result_image is not None:
    image_queue.remove(image1)
    image_queue.remove(image2)
    break
```

Using ORB, key points and descriptors of images are extracted.

```
def get_keypoint_descriptors(img):
    ''' Uses ORB to extract keypoints and descriptors from an image '''
    return orb.detectAndCompute(img, None)
```

• Using find_matches function, we create a set of matching pairs from both the images using hamming distance calculated over bits of descriptors.

```
lef find_matches(keypoints1, descriptors1, keypoints2, descriptors2):
  results = list()
  descriptor_size = len(descriptors1[0])
  overall_min = sys.maxsize
  for i in range(len(descriptors1)):
      min_diff = sys.maxsize
       min_right_index = -1
       for j in range(len(descriptors2)):
           total_diff_bits = 0
           for k in range(descriptor_size):
              y = int( \{0:08b\} \cdot .format(descriptors2[j][k]))
              total_diff_bits += np.count_nonzero(x != y)
           if total_diff_bits < min_diff:
              min_diff = total_diff_bits
              min_right_index = j
               if min_diff < overall_min:
                  overall_min = min_diff
       results.append([list(keypoints1[i].pt), list(keypoints2[min_right_index].pt)])
   if overall_min > 10:
```

• Once we have the matches, ransac function is used to eliminate all the outliers from the matches set and compute the homography matrix.

```
ransac(source_points, destination_points, inlier_threshold):
"" Used to calculate the best homography matrix based on 4 randomnly chosen points and counting inliers ""
max_inliers = - sys.maxsize - 1
best_h = None
    random_pts = np.random.choice(range(0, len(source_points) - 1), 4, replace=False)
    for i in random_pts:
        src.append(source_points[i])
       dst.append(destination_points[i])
    h = find_homograph(src, dst)
    for index in range(len(source_points)):
        src_pt = np.append(source_points[index], 1)
       dest_pt = np.dot(h, src_pt.T)
       dest_pt = np.true_divide(dest_pt, dest_pt[2])[0: 2]
       if distance.euclidean(destination_points[index], dest_pt) <= inlier_threshold:
           count += 1
    if count > max_inliers:
       max_inliers = count
       best_h = h
return best_h, max_inliers
```

find_homograph and create_P_matrix are the helper functions to create the homography matrix.

```
fef create_P_matrix(source_points, destination_points):
    "" Helper matrix for creating homography matrix ""
   sub_matrices = []
   for i in range(len(source_points)):
       sub_matrix = np.zeros((2,9))
       sub_matrix[0][0] = -1 * source_points[i][0]
       sub_matrix[0][1] = -1 * source_points[i][1]
       sub_matrix[0][2] = -1
       sub_matrix[0][6] = source_points[i][0] * destination_points[i][0]
       sub_matrix[0][7] = source_points[i][1] * destination_points[i][0]
       sub_matrix[0][8] = destination_points[i][0]
       sub_matrix[1][3] = -1 * source_points[i][0]
       sub_matrix[1][4] = -1 * source_points[i][1]
       sub_matrix[1][5] = -1
       sub_matrix(1)(6) = source_points(i)(0) * destination_points(i)(1)
       sub\_matrix[1][7] = source\_points[i][1] + destination\_points[i][1]
       sub_matrix[1][8] = destination_points[i][1]
       sub_matrices.append(sub_matrix)
def stitcher(img1, img2, gray1, gray2):
   keypoints1, descriptors1 = get_keypoint_descriptors(gray1)
   keypoints2, descriptors2 = get_keypoint_descriptors(gray2)
   matches = find_matches(keypoints1, descriptors1, keypoints2, descriptors2)
   if matches is None:
   src = np.float32([match[0] for match in matches]).reshape(-1,2)
   dst = np.float32([match[1] for match in matches]).reshape(-1,2)
   homograph = None
   threshold = min(len(descriptors1), len(descriptors2))
       homograph, max_inliers = ransac(src, dst, i)
       if 0.1 * threshold <= max_inliers < 0.3 * threshold:
       elif max_inliers < 0.1 * threshold:
       else: i -= 1
   return warp_images(img2, img1, homograph), warp_images(gray2, gray1, homograph)
```

- The process of running the RANSAC algorithm is done until we get the inliers in the range of 10% to 30% the size of descriptors to avoid under fitting and over fitting of the images.
- This is done by increasing the inlier threshold distance by 1 starting from 1.

 Once we have the correct set of inliers and homography matrix, warp_images is called to warp the second image over the first one.

```
def warp_images(img1, img2, H):
    ''' Warps image 2 and stitches it to image 1 '''
    h1, w1 = img1.shape[ :2]
    h2, w2 = img2.shape[ :2]
    pts1 = np.float32([[0, 0], [0, h1], [w1, h1], [w1, 0]]).reshape(-1,1,2)
    pts2 = np.float32([[0, 0], [0, h2], [w2, h2], [w2, 0]]).reshape(-1,1,2)
    pts2_ = cv2.perspectiveTransform(pts2, H)
    pts = np.concatenate((pts1, pts2_), axis=0)
    [xmin, ymin] = np.int32(pts.min(axis=0).ravel() - 0.5)
    [xmax, ymax] = np.int32(pts.max(axis=0).ravel() + 0.5)
    t = [-xmin, -ymin]
    Ht = np.array([[1, 0, t[0]], [0, 1, t[1]], [0, 0, 1]])
    result = cv2.warpPerspective(img2, Ht.dot(H), (xmax-xmin, ymax-ymin))
    result[t[1]:h1+t[1],t[0]:w1+t[0]] = img1
    return result
```

To stitch a third image, we use the stitched two images and repeat the process with the third image.

```
# If there are more than two images then stitch this third image to the panorama obtained earlier
if len(image_queue) > 0:
    image = image_queue[-1]
    img, gray = read_image(image, img_dir)
    result_image, result_gray = stitcher(result_image, img, result_gray, gray)

# Finally write the panorama to disk
write_output_image(result_image, img_dir, 'panorama')
```

Results:

The following were the results of the stitcher script when ran on two sets of test images placed in "data" and "ubdata" directories.

• Nevada Images:









The above image is the result of stitching the three Nevada images into panorama style.

• UB Images:

The below image is the panorama stitching performed on images from the UB Capen library. This is the place I value the most at UB and spend most of my time learning and doing projects.







