Labb 3

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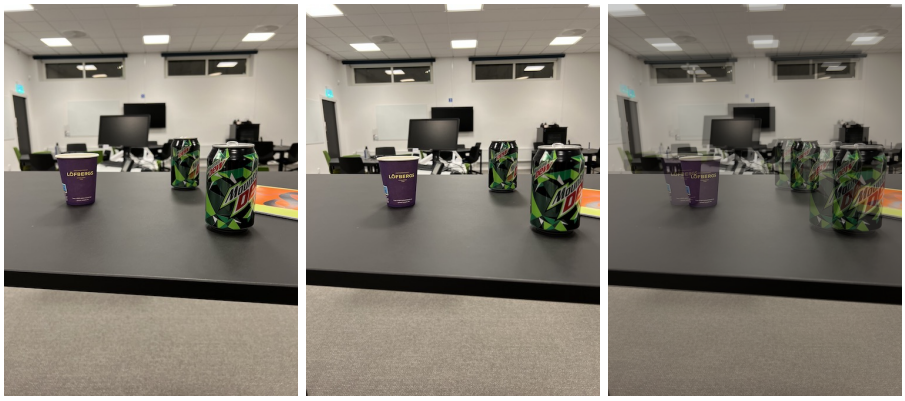
1 Week 5

1.1 Assignment 5

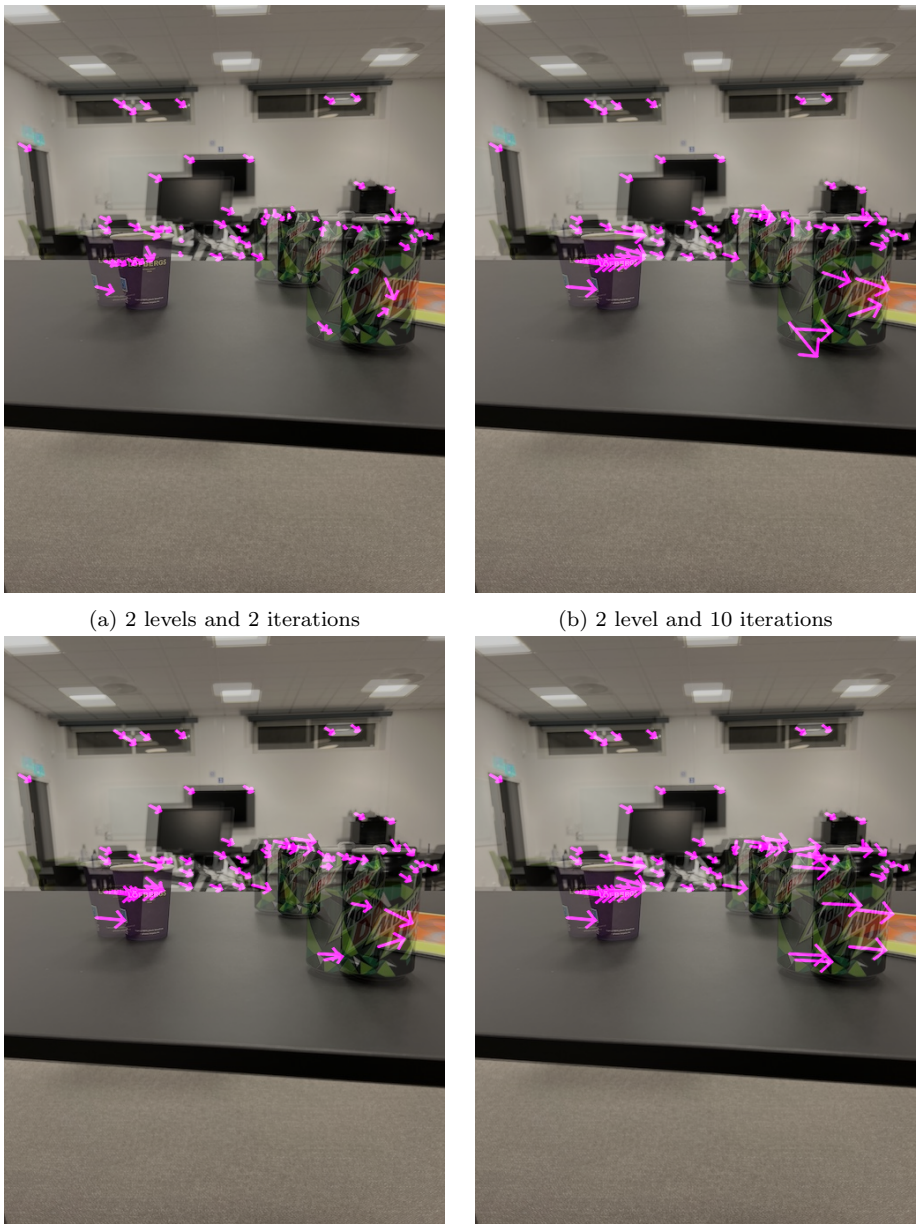
1.1.1 Optical Flow

I used OpenCV's implementation for Optical Flow. This was originally used for videos but was adopted to be used with images instead. Coarse-to-fine algorithm improves our result on larger deltas by creating different layers of the image in different sizes. The smaller sizes helps the algorithm find features points. The itterations helps the algorithm to improve our results by correcting errors. In figure 1 we can see the input images and in figure 2 we can see the different results. We can see that the arrows correct them selves with more iterations and the points getting more accurate with more layers. The Python code can be found in appendix A



(a) First image (b) Second image (c) Images overlayed

Figure 1: The two input images for Optical Flow and the the images overlayed



(a) 2 levels and 2 iterations

(b) 2 level and 10 iterations

(c) 3 layers and 2 iterations

(d) 3 layers and 10 iterations

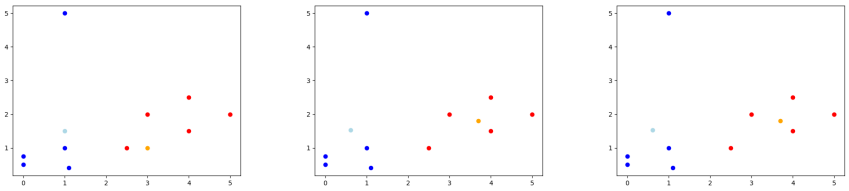
Figure 2: Resulting vectors from Lucas-Kanade method with different amount of pyramids and iterations

1.1.2 K-means

The K-means function can be found in equation 1. This algorithm assumes that all the datasets belong to a category μ.

$$\min_{\mu,y} \sum_{i} \|x_i - \mu_{y_i}\|^2 \tag{1}$$

In figure 3 we can see the different iterations. In figure 3a is the original data categorized. Then first iteration in figure 3b and then when we compare that to 3c we see that nothing changes and thus our final categorized are decided. These images were produced with a python script that can be found in appendix B.



(a) 0 Iterations (b) 1 Iteration (c) 2 Iteration

Figure 3: Iterations of K-means

1.1.3 Segmentation

I chose to implement K-means for color segmentation. My code can be found in appendix C. Input image can be found in figure 4 and output can be found in figure 5. The only parameters I can change is the max number of iterations and the number of clusters. The implementation I created stops the process if 70% of the clusters have moved less than 1 unit since the last iteration. So the number of max iterations just gives the algorithm more time to reach this criteria. The number of clusters will define how many colors the resulting image will have.

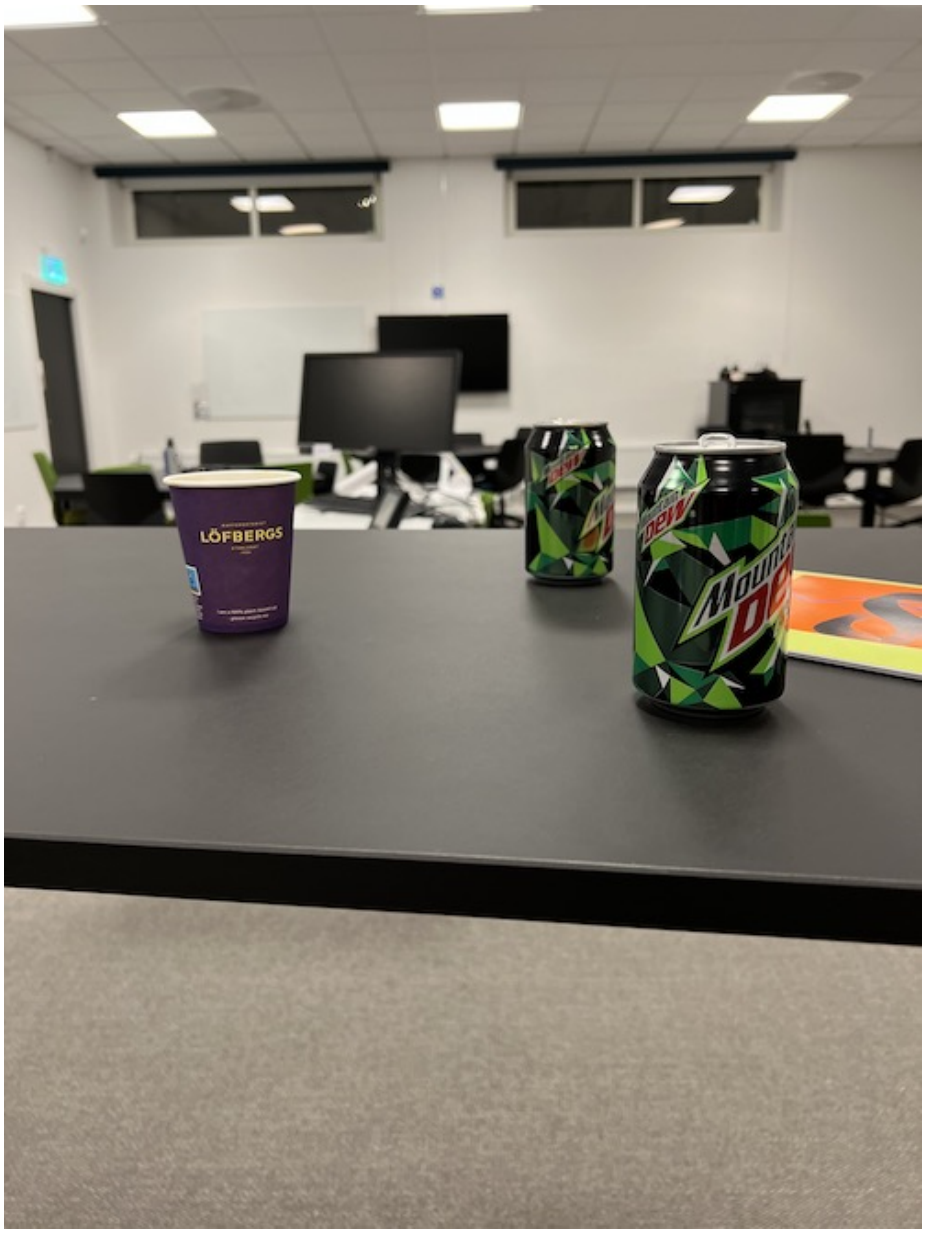


Figure 4: Input for K-means segmentation

Figure 5: Output from segmentation with 12 clusters and 27 iterations

2 Week 6

2.1 Assignment 6

2.1.1 Theory

The library I choose for Structure from Motion (SFM) was OpenMVG [1]. The library is written in C++ but is run through Python scripts. The Script takes input arguments for both a directory where the input images are located and an output directory. The output consist of a lot of different data. Most importantly it provides a number of .ply files that contain the resulting point cloud. The OpenMVG consists of several submodules for example we have a submodule for running SIFT. To run SFM I used the provided python script, and it goes through the following steps

1. Intrinsic analysis
2. Feature detection
3. Feature Matching
4. Filter Matches
5. Global Reconstruction
6. Coloring

The first step goes through the metadata of the images extracting the camera sensor that took the image and compares that to a pre-defined database. From this it goes through all the images and extracts data and compiles the necessary data to a JSON file to be used by the rest of the operations.

```
print ("1. Intrinsics analysis")
pIntrisics = subprocess.Popen( [os.path.join(OPENMVG_SFM_BIN, "
    openMVG_main_SfMInit_ImageListing"),  "-i", input_dir, "-o",
    matches_dir, "-d", camera_file_params] )
pIntrisics.wait()
```

Feature detection uses SIFT to extract features from the images and saves the extracted features to files corresponding to the images.

```
print ("2. Compute features")
pFeatures = subprocess.Popen( [os.path.join(OPENMVG_SFM_BIN, "
    openMVG_main_ComputeFeatures"),  "-i", matches_dir+"/sfm_data.
    json", "-o", matches_dir, "-m", "SIFT"] )
pFeatures.wait()
```

Next we have feature matching and pair generation. The pair generation generates all the possible pairs of images and puts it in a set. The set is then used to pair images and find matches. The matcher file contains multiple matching algorithm, but it prefers to use Cascade Hashing Matcher.

```
print ("3. Compute matching pairs")
```

```
pPairs = subprocess.Popen( [os.path.join(OPENMVG_SFM_BIN, "
    openMVG_main_PairGenerator"), "-i", matches_dir+"/sfm_data.json
    ", "-o" , matches_dir + "/pairs.bin" ] )
pPairs.wait()

print ("4. Compute matches")
pMatches = subprocess.Popen( [os.path.join(OPENMVG_SFM_BIN, "
    openMVG_main_ComputeMatches"),  "-i", matches_dir+"/sfm_data.
    json", "-p", matches_dir+ "/pairs.bin", "-o", matches_dir + "/
    matches.putative.bin" ] )
pMatches.wait()
```

Next it performes geometric filtering on the matches to filter out bad matches.

```
print ("5. Filter matches" )
pFiltering = subprocess.Popen( [os.path.join(OPENMVG_SFM_BIN, "
    openMVG_main_GeometricFilter"), "-i", matches_dir+"/sfm_data.
    json", "-m", matches_dir+"/matches.putative.bin" , "-g" , "f" ,
     "-o" , matches_dir+"/matches.f.bin" ] )
pFiltering.wait()
```

The last two steps comes down to the actual structure from motion and then coloring the 3D points. The method used here is sequential method. In their library I can see there are three methods for 3D reconstruction. Sequential, Sequential v2 and Global. They provide python script for both sequential and global reconstruction, so that is what I have tried. But I will only look into the sequential part due to the superior result it provided. The codebase for this project is massive, so I will only go through the processing steps.

1. Make sure we have a correct initial pair.
2. Create an essential matrix from these pairs.
3. Loop through possible images for reconstruction.
   (a) Add possible images to reconstruction.
   (b) Do bundle adjustment until all points are within a certain precision.
   (c) Remove unstable poses and observations.
4. Ensure there are no more outliers.
5. Done

```
print ("6. Do Sequential/Incremental reconstruction")
pRecons = subprocess.Popen( [os.path.join(OPENMVG_SFM_BIN, "
    openMVG_main_SfM"), "--sfm_engine", "INCREMENTAL", "--
    input_file", matches_dir+"/sfm_data.json", "--match_dir",
    matches_dir, "--output_dir", reconstruction_dir] )
pRecons.wait()

print ("7. Colorize Structure")
pRecons = subprocess.Popen( [os.path.join(OPENMVG_SFM_BIN, "
    openMVG_main_ComputeSfM_DataColor"),  "-i", reconstruction_dir+
    "/sfm_data.bin", "-o", os.path.join(reconstruction_dir,"
    colorized.ply")] )
```

```
pRecons.wait()
```

2.1.2 Implementation

I used openMVG [1] and ran both Sequential and Global reconstruction. Due to some errors whit camera calibration when using my own images I used a dataset provided by them with images of Échillais church [2]. Figure 6 shows an example of images in the dataset.

Figure 6: Example images from the Échillais church dataset[2]

The first reconstruction I did was through the sequential pipeline. The resulting

data points are shown in Figure 7.

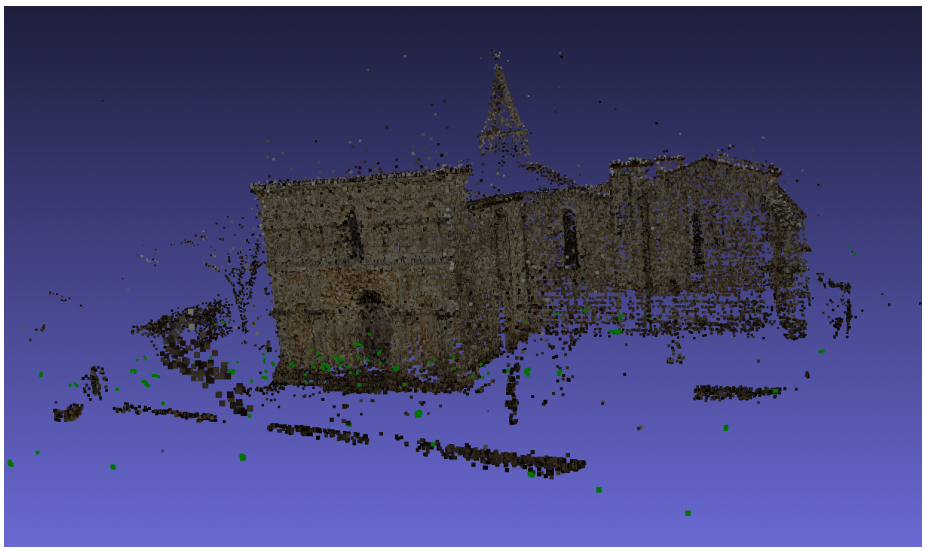


Figure 7: Resulting data points from 3D reconstruction using the sequential SFM pipeline

To compare the different pipelines I also ran the global reconstruction pipeline and the resulting data points ca be found in figure 8.

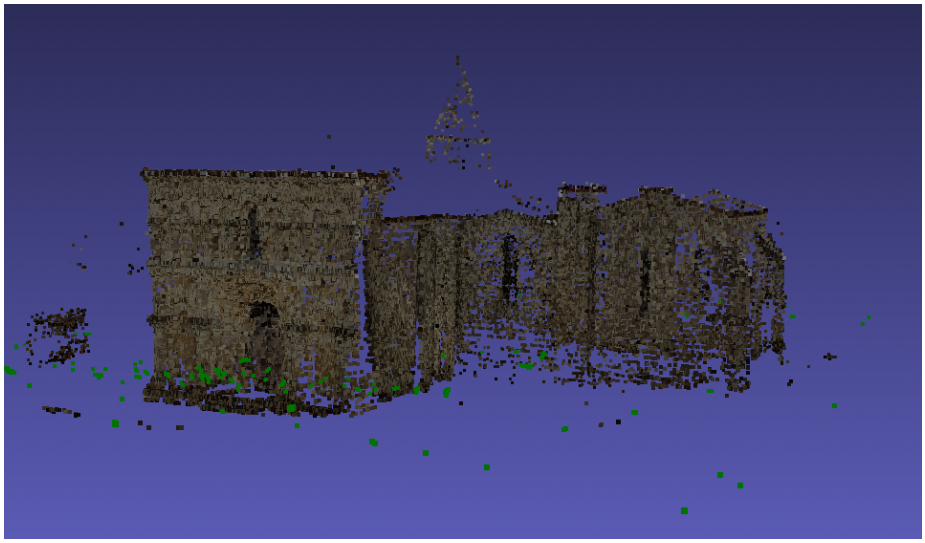


Figure 8: Resulting data points from 3D reconstruction using the sequential SFM pipeline

The global pipeline manages to produce is close to the sequential, but it is not as detailed as the sequential pipeline. The easiest way to see this is to look at the tower of the church as it is more detailed on the sequential pipeline.

3 Week 7

3.1 Assignment 7

3.1.1 Triangle

The triangle I choose to do had the following points:

1. (3,2)
2. (2,5)
3. (7,3)

I then chose the inside point (3,2) and the outside point (2,1). I created a python script to calculate my points. The script can be found in appendix D. And the plotted result can be found in figure 9.

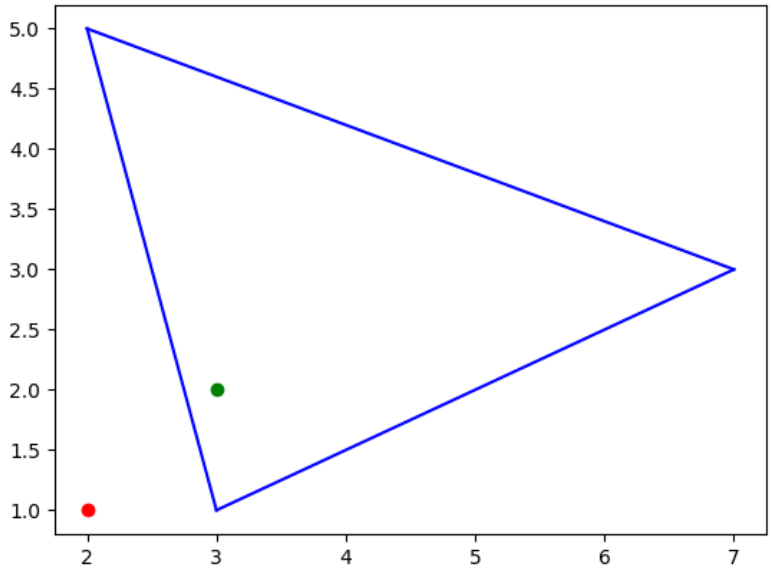


Figure 9: Resulting data points from 3D reconstruction using the sequential SFM pipeline

3.1.2 Phong

I used python to plot the model. For the constant values I just picket random numbers. The resulting plot can be seen in figure 10 and the code can be found in appendix E

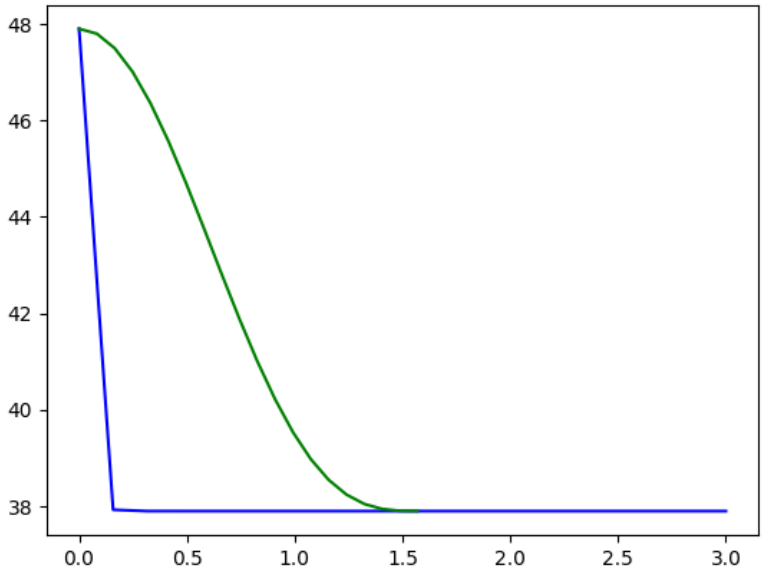


Figure 10: Plot of Phong lighting, blue is plot based on α and green is plot based on φ

References

[1] Pierre Moulon, Pascal Monasse, Romuald Perrot, et al. "OpenMVG: Open multiple view geometry". In: *International Workshop on Reproducible Research in Pattern Recognition*. Springer. 2016, pp. 60–74.

[2] Romuald Perrot. *Reconstruction Dataset*. 2017. URL: `https://github.com/rperrot/ReconstructionDataSet/tree/master/EchillaisChurch` (visited on 12/18/2021).

Appendix A Python code for optical flow

```
import numpy as np
import cv2
import os

frame1 = cv2.imread("im1.jpg")
frame2 = cv2.imread("im2.jpg")

frame1g = cv2.cvtColor(frame1, cv2.COLOR_BGR2GRAY)
frame2g = cv2.cvtColor(frame2, cv2.COLOR_BGR2GRAY)

combined = cv2.addWeighted(frame1, 0.3, frame2, 0.5, 0)
cv2.imwrite("pyramid/combined.png", combined)

parameters = dict(maxCorners=100, qualityLevel=0.3, minDistance=7,
    blockSize=7)

p0 = cv2.goodFeaturesToTrack(frame1g, mask=None, **parameters)

maxLevels = [2,3]
maxIterations = [2, 10]

for level in maxLevels:
    print("On Max Level: ", level)
    for iter in maxIterations:
        OFparams = dict(
            winSize=(15, 15),
            maxLevel=level,
            criteria=(cv2.TERM_CRITERIA_EPS |
                      cv2.TERM_CRITERIA_COUNT, iter, 0.03),
        )
        p1, st, err = cv2.calcOpticalFlowPyrLK(
            frame1g, frame2g, p0, None, **OFparams)
        good_new = p1[st == 1]
        good_prev = p0[st == 1]

        arrows = np.zeros_like(combined)

        for i, (n, p) in enumerate(zip(good_new, good_prev)):
            nx, ny = n.ravel()
            px, py = p.ravel()

            nx = int(nx)
            ny = int(ny)
            px = int(px)
            py = int(py)

            arrows = cv2.arrowedLine(arrows, (px, py), (nx, ny), [
                                255, 0, 255], 2, cv2.LINE_AA,
                                    tipLength=0.4)

            output = cv2.add(combined, arrows)
            if(not os.path.exists("pyramid/"+str(level))):
                os.mkdir("pyramid/"+str(level))
            outName = "pyramid/" + str(level)+"/" + \
                str(level) + "levels-" + str(iter)+"iterations.png"
            cv2.imwrite(outName, output)
```

Appendix B K-means

```
from typing import Tuple
from typing import List
import matplotlib.pyplot as plt
from numpy import sqrt
import numpy as np

def centerToLists(center1, center2):
    return ([center1[0], center2[0]], [center1[1], center2[0]])


def drawPlot(center1, center2, gc1, gc2):
    plt.scatter([i[0] for i in gc1], [i[1] for i in gc1], c='blue')
    plt.scatter([i[0] for i in gc2], [i[1] for i in gc2], c='red')
    plt.scatter(center1[0], center1[1],c='lightblue')
    plt.scatter(center2[0], center2[1],c='orange')

def calcDistance(p1, p2):
    return sqrt((p1[0] -p2[0])**2 + (p1[1] - p2[1])**2)

def categorizePoints(points, center1, center2) -> Tuple[List, List
    ]:
    gc1 = []
    gc2 = []
    for point in points:
        d1 = calcDistance(point, center1)
        d2 = calcDistance(point, center2)
        if d1 < d2:
            gc1.append(point)
        else:
            gc2.append(point)
    return (gc1, gc2)

def calculateNewCenter(gc1: List, gc2: List) -> Tuple[Tuple, Tuple
    ]:
    g1x = np.asarray([i[0] for i in gc1]).sum()/len(gc1)
    g1y = np.asarray([i[1] for i in gc1]).sum()/len(gc1)

    g2x = np.asarray([i[0] for i in gc2]).sum()/len(gc2)
    g2y = np.asarray([i[1] for i in gc2]).sum()/len(gc2)

    return ((g1x, g1y), (g2x, g2y))



points = [
    (0, 0.5),
    (0, 0.75),
    (1, 1),
    (1.1, 0.4),
    (1, 5, 0.75),
    (2.5, 1),
    (3, 2),
    (4, 1.5),
    (4, 2.5),
    (5, 2)
]
```

```
55
56 center1 = (1,1.5)
57 center2 = (3,1)
58
59 (gc1, gc2) = categorizePoints(points, center1, center2)
60 drawPlot(center1, center2, gc1, gc2)
61 plt.show()
62 (center1, center2) = calculateNewCenter(gc1, gc2)
63 (gc1, gc2) = categorizePoints(points, center1, center2)
64 drawPlot(center1, center2, gc1, gc2)
65 plt.show()
66 (center1, center2) = calculateNewCenter(gc1, gc2)
67 (gc1, gc2) = categorizePoints(points, center1, center2)
68 drawPlot(center1, center2, gc1, gc2)
69 plt.show()
```

Appendix C Color segmentation

```
1  import numpy as np
2  import cv2
3  from typing import List, Tuple
4  import collections
5  import random
6  import math
7  from PIL import Image
8
9  class Point:
10     position = (0,0)
11     color = [0,0,0]
12     category = [0,0,0]
13
14
15 def convImForShow(im):
16     return cv2.cvtColor(im, cv2.COLOR_RGB2BGR)
17
18 def calcDistance(p1, p2) -> float:
19     x = np.square(p1[0] - p2[0])
20     y = np.square(p1[1] - p2[1])
21     z = np.square(p1[2] - p2[2])
22     sum = x+y+z
23     return math.sqrt(sum)
24
25 def categorizePoints(points: List[Point], centers: List) -> Tuple[
       List[List], List[Point]]:
26     reee = set()
27     ret = [[] for i in range(len(centers))]
28     dist = np.zeros(len(centers))
29     l = []
30     for p in points:
31         for i, c in enumerate(centers):
32             dist[i] = calcDistance(p.color, c)
33         index = np.argmin(dist)
34         reee.add(index)
35         ret[index].append(p.color)
36         p.category = centers[index]
37         l.append(p)
```

```
38      return (ret, l)
39
40  def calcCenter(category: List) -> List:
41      x = np.asarray([i[0] for i in category]).sum()/len(category)
42      y = np.asarray([i[1] for i in category]).sum()/len(category)
43      z = np.asarray([i[2] for i in category]).sum()/len(category)
44      return [x,y,z]
45
46
47  def calcNewCenters(categories: List[List]) -> List[List]:
48      newCenters = [[] for i in range(len(categories))]
49      for i, cat in enumerate(categories):
50          newCenters[i] = calcCenter(cat)
51      return newCenters
52
53  def colorPoints(points: List[List], categories: List[List], centers
        : List[List]) -> List[List]:
54      compare = lambda x, y: collections.Counter(x) == collections.
            Counter(y)
55      for i, cat in enumerate(categories):
56          for p in cat:
57              indexes = [i for i,x in enumerate(points) if compare(x,
                    p)]
58              for j in indexes:
59                  points[j] = centers[i]
60      return points
61
62  def imageToPoints(img) -> List[Point]:
63      ret = []
64      for y in range(len(img)):
65          for x in range(len(img[0])):
66              temp = Point()
67              temp.position = (x, y)
68              temp.color = img[y][x]
69              ret.append(temp)
70      return ret
71
72  def pointsToImage(points: List[Point], shape):
73      p = [x.category for x in points]
74      return np.asarray(p, np.uint8).reshape(shape)
75
76  def compareCenters(oldCenter, newCenter):
77      underLimit = 0
78      for i, _ in enumerate(oldCenter):
79          distance = calcDistance(oldCenter[i], newCenter[i])
80          if distance < 1:
81              underLimit += 1
82
83      if underLimit > (len(oldCenter)*0.7):
84          return True
85      return False
86
87
88  im = Image.open("im1.jpg")
89  im = np.asarray(im)
90  numberOfClusters = 4
91
```

```
numberOfIterations = 50

shape = im.shape
points = imageToPoints(im)
randPoints = random.sample(points, numberOfClusters)
centers = [p.color for p in randPoints]

cat, points = categorizePoints(points, centers)

iters = 0
for i in range(numberOfIterations):
    cat, points = categorizePoints(points, centers)
    newCenters = calcNewCenters(cat)
    iters = i
    if compareCenters(centers, newCenters):
        break
    centers = newCenters

print("Did ", iters, " iterations")

temp = pointsToImage(points, shape)
temp = Image.fromarray(temp)
temp.show("Iter 0")
temp.save("segmented.png")
```

Appendix D Triangle code

```
import matplotlib.pyplot as plt

def subtract(p0, p1):
    return (p0[0] - p1[0], p0[1], p1[1])

def dotProduct(p0, p1):
    return (p0[0] * p1[0]) + (p0[1] * p1[1])

def isInside(triangle, point):
    p0, p1, p2 = triangle[0], triangle[1], triangle[2]
    dxdy = subtract(p1, p0)
    normal = (dxdy[1], -dxdy[0])
    line1_check = dotProduct(subtract(point, p0), normal) > 0

    dxdy = subtract(p2,p1)
    normal = (dxdy[1], -dxdy[0])
    line2_check = dotProduct(subtract(point, p1), normal) <0

    dxdy = subtract(p0,p2)
    normal = (dxdy[1], -dxdy[0])
    line3_check = dotProduct(subtract(point, p2), normal) > 0

    return line1_check and line2_check and line3_check

triangle = [
    (3, 1),
    (2, 5),
    (7, 3),
]
```

```
30
31 inside_point = (3,2)
32 outside_point = (2,1)
33
34 points = [inside_point, outside_point]
35
36 for point in points:
37     color = 'go' if isInside(triangle, point) else 'ro'
38     plt.plot([point[0]], [point[1]], color)
39
40 plt.plot([triangle[0][0], triangle[1][0]], [triangle[0][1],
       triangle[1][1]], 'b')
41 plt.plot([triangle[1][0], triangle[2][0]], [triangle[1][1],
       triangle[2][1]], 'b')
42 plt.plot([triangle[0][0], triangle[2][0]], [triangle[0][1],
       triangle[2][1]], 'b')
43 plt.show()
```

Appendix E Phong plot

```
1  import numpy as np
2  import matplotlib.pyplot as plt
3
4
5  def phong(kA, IA, kD, ID, kS, phi, alpha, IL):
6      return (kA*IA) + (kD+ID) + (kS * (np.cos(phi)**alpha)*IL)
7
8
9  IA = 6.4
10 ID = 3.9
11 IL = 2.5
12
13 kA = 5.
14 kD = 2.
15 kS = 4.
16
17 alpha = np.linspace(0.0, 3.0, num=20)
18 phi = np.linspace(0.0, np.pi/2, num=20)
19
20 ya = [phong(kA, IA, kD, ID, kS, np.pi/2, x, IL) for x in alpha]
21 yp = [phong(kA, IA, kD, ID, kS, x, 3, IL) for x in phi]
22
23 plt.plot(alpha, ya, 'b')
24 plt.plot(phi, yp, 'g')
25 plt.show()
```