Image Processing HW\_5

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### Problem 1:

### Section a:

Implementing scale\_down function by applying our knowledge on Fourier, we know that a scaled down image by ratio is the image smaller in that ratio and has and without the high frequencies in the same ratio, so we take the crop the Fourier transform by the resize\_ratio starting from the lower frequencies, and then we inverse the cropped window to get the sacled\_down image.

```
# Section a
def scale_down(image, resize_ratio):
    fourier_spectrum = fftshift(fft2(image))
    h, w = image.shape
    new_h, new_w = int(h / resize_ratio), int(w / resize_ratio)

start_row = (h - new_h) // 2
    start_col = (w - new_w) // 2
    end_row = start_row + new_h
    end_col = start_col + new_w

cropped_spectrum = fourier_spectrum[start_row:end_row, start_col:end_col]

return np.abs(ifft2(ifftshift(cropped_spectrum)))
```

## Section b:

The same thing we did in scale\_down, butt this time we need to strentgth the freuquencies because now we have a bigger image, and we do that in the last line

```
# Section b

def scale_up(image, resize_ratio):
    fourier_spectrum = fftshift(fft2(image))
    h, w = image.shape
    new_h, new_w = int(h * resize_ratio), int(w * resize_ratio)
    fourier_spectrum_zero_padding = np.zeros((new_h, new_w), dtype=complex)

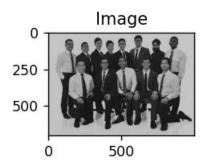
start_row = (new_h - h) // 2
    start_col = (new_w - w) // 2
    end_row = start_row + h
    end_col = start_col + w

fourier_spectrum_zero_padding[start_row:end_row, start_col:end_col] = fourier_spectrum

return np.abs(ifft2(ifftshift(fourier_spectrum_zero_padding))) * (resize_ratio ** 2)
```

## Example:

The crew image scaled up with resize\_ratio = 4:



We can see now the image contains more pixels than the original and in the same quality

### Section c:

This function calculates the ncc (Nomalized Cross Correlation) between an image and a pattern, using sliding windows, mean and variance. This fuction used in "find the pattern in the image problem".

The following implementation is based on this formula:

$$\sum_{x,y\in N} \left[\!\!\left[\!\!\left(\!\!\left(u+x,v+y\right)\!\!-\!\bar{l}_{\!\scriptscriptstyle uv}\right)\!\!\right]\!\!\left[\!\!\left[\!\!P\!\!\left(x,y\right)\!\!-\!\overline{P}\right]\!\!\right]$$

$$\left[\sum_{x,y\in N}\left[I\!\!\left(u+x,v+y\right)\!-\bar{I}_{uv}\right]^{\!2}\!\sum_{x,y\in N}\left[P\!\left(x,y\right)\!-\overline{P}\right]^{\!2}\right]^{\!1/2}$$

Where:

P – pattern

I – Image

 ${\rm I}^-$  -image mean

P<sup>-</sup>-pattern mean

```
# Section c
def ncc_2d(im, patt):
    windows = np.lib.stride_tricks.sliding_window_view(im, patt.shape)
    ncc_im = np.zeros(windows.shape[:2])
    patt_mean = np.mean(patt)
    patt_var = np.sum((pattern - patt_mean) ** 2)

# implementing the formula for calculating ncc
for row in range(len(windows)):
    for col in range(len(windows[0])):
        window_mean = np.mean(windows[row, col])
        window_var = np.sum((windows[row, col] - window_mean) ** 2)
        means_sum = np.sum((windows[row, col] - window_mean) * (patt - patt_mean))
        denominator = np.sqrt(window_var * patt_var)
        if denominator == 0:
            continue
        ncc_im[row, col] = means_sum / denominator

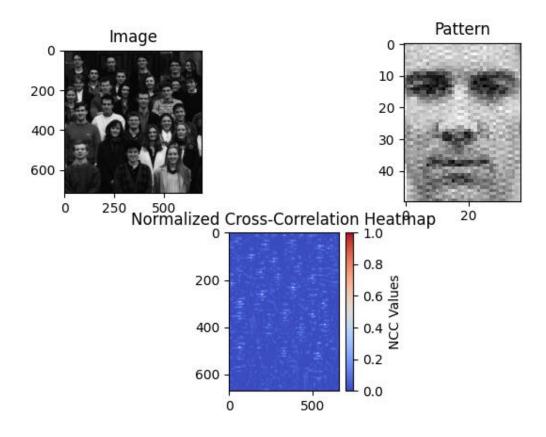
return ncc_im
```

## Section d:

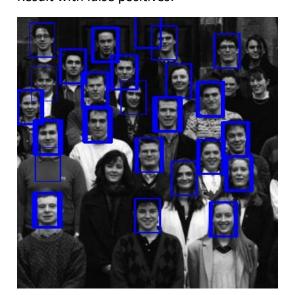
Students image:

Display:

image\_scale\_ratio = 1.8
pattern\_scale\_ratio = 1
threshold = 0.46



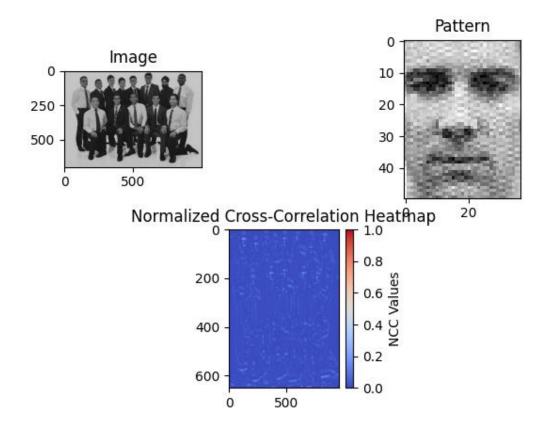
# Result with false positives:



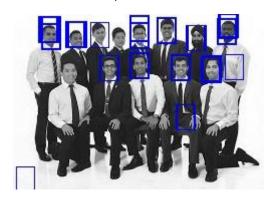
# Thecrew image:

# Display:

image\_scale\_ratio = 4
pattern\_scale\_ratio = 1
threshold = 0.37



# Result with false positives:



# Section e:

## **Our best Results**

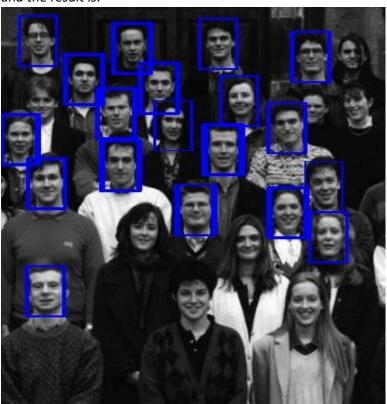
After search and trial and error, here are our best results

Students:

we chose these params:

image\_scale\_ratio = 2
pattern\_scale\_ratio = 1
threshold = 0.55

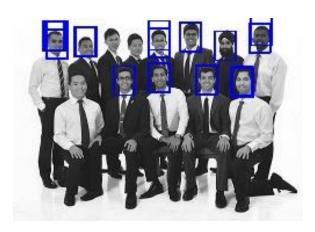
# and the result is:



# thecrew:

we chose these params:

image\_scale\_ratio = 4
pattern\_scale\_ratio = 1
threshold = 0.4



### Problem 2:

First I want to mention that I implemented these aid functions in addition to the functions required in the question:

```
def get_gaussian_pyramid(image, levels):
    """
    this function takes an image and returns the gaussain pyramid of the image
        :param image: grey-scale image
        :param levels: the levels of the pyramid
        :return: list contains the layers of the gaussian pyramid of the image
    """
        pyramid = [image]
        current_layer = image
    for _ in range(levels - 1):
            current_layer = cv2.GaussianBlur(current_layer, (7, 7), 0)
            current_layer = current_layer[::2, ::2]
            pyramid.append(current_layer)
```

And i used scale\_down() and scale\_up() from Question 1

### Section a:

```
def get_laplacian_pyramid(image, levels):

"""

this function takes an image and levels, and it returns the laplacian pyramid of the image as a list,
it does that by applying what we learned in the class, and by using the gaussian pyramid, and a simple formula
we learned in class
:param image: the image we want to create the laplacian pyramid for
:param levels: levels of the pyramid
:return: the laplacian pyramid of the given image

"""

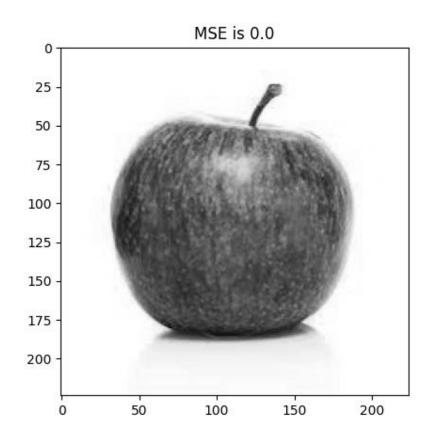
gaussian_pyramid = get_gaussian_pyramid(image, levels)
laplacian_pyramid = []
# Running on the levels and doing the needed
for i in range(levels - 1):
    expanded = scale_up(gaussian_pyramid[i] - expanded)
laplacian_pyramid.append(gaussian_pyramid[i]) # Append the smallest level

return laplacian_pyramid
```

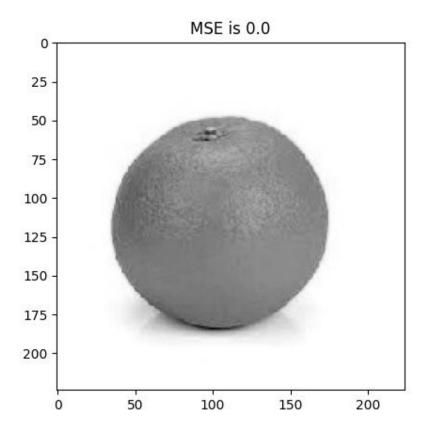
### Section b:

By using the given function validate\_operation, we created Laplacian pyramids for both orange and apple and we restored it and we got this results:

Apple:



# Orange:



Less than 1, so NICE!

### Section c:

```
def blend_pyramids(levels):
   blend_pyr = []
    for curr_level in range(levels):
       mask = np.zeros(pyr_apple[curr_level].shape)
       width = mask.shape[1]
       mask[:, :int((0.5 * width) - (curr_level + 1))] = 1.0
       for i in range(2 * (curr_level + 1)):
            mask[:, (width // 2) - (curr_level + 1) + i] = 0.9 - 0.9 * i / (2 * (curr_level + 1))
       blend_pyr.append((pyr_orange[curr_level] * mask) + (pyr_apple[curr_level] * (1 - mask)))
   return blend_pyr
```

### Section d:

We created Laplacian pyramids for both images Apple and Orange, and then we blended the two images using our blend function:

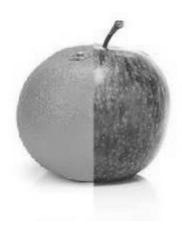
```
# Creating laplacian pyramids for both images Orange and Apple
pyr_apple = get_laplacian_pyramid(apple, levels)
pyr_orange = get_laplacian_pyramid(orange, levels)

# Blending Pyramids
pyr_result = blend_pyramids(levels)

# Getting and plotting the blended image
final = restore_from_pyramid(pyr_result)
plt.imshow(final, cmap='gray')
plt.show()
```

But before that we needed to choose the best level value that gives the best image we can get from blending the two images, after trying all the possible levels, we found out that level=3 is the perfect match in the trade-off between the smoothens in the cross-dissolve and between the "showing" that this is blended image (there is dominant part in each side has its own colors)

Level = 1:



Level = 2:



Level = 3: our best result



Level = 4:



Level = 5



in levels = 1\ levels = 2, we clearly can see the line in the middle that separates the two images, the transition between the images is not smooth and it is not good.

While in levels = 4\ levels = 5, we can see also the fainting in the colors of each side, which is also not good.

And finally in levels=3 we find the best results, we have a good blending and also we have great colors for each side.

## **Bonus Question:**

### Section a:

I completed this section in the code file in many different places

### Section b:

I used the following canny implementation to find edges in the image:

### Section c:

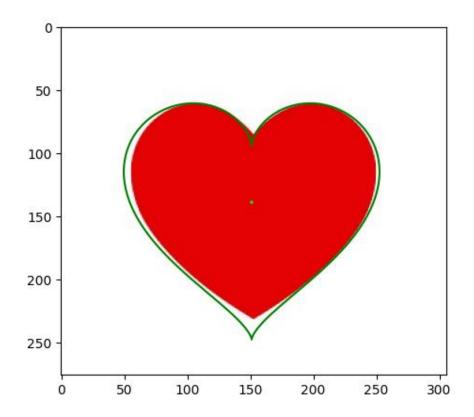
After many trail and error I found these parameters that led me to the best result I had:

## simple:

Parmeters:

```
r_min = 6.5
r_max = 7.5
bin threshold = 0.18
```

## Result

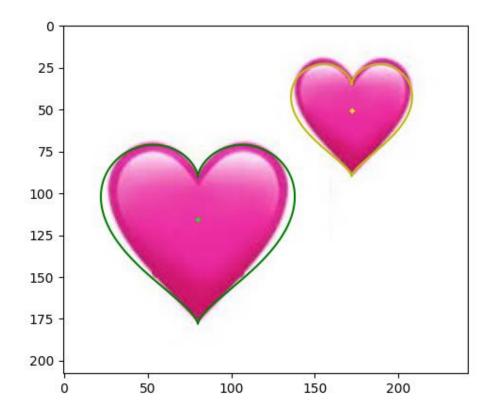


# med:

# Parameters:

```
r_min = 2
r_max = 5
bin_threshold = 0.3
```

# Result:

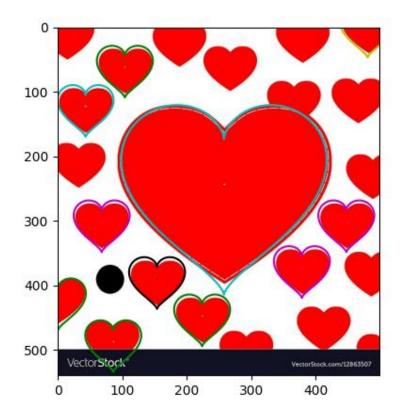


### hard:

### Parameters:

```
r_min = 3
r_max = 12
bin threshold = 0.26
```

## Result:



We can see that we succeeded to find all the hearts in simple and med image, but we have a problem to do that in the hard image, and that is because we can see that in simple and med image we only have hearts and few of them (simple contains 1 and med contains 2) while in the hard image we not only have a much more hearts but also we have also different shapes in the image itself, like the black circle and the black rectangle below, these factors makes it hard to detect all the hearts in the image.