# HW1-9822762040

November 25, 2023

# 1 Image Processing

### 1.1 Libraries

First of all, we include all libraries needed for this exercise.

```
[3]: import numpy as np
import cv2 as cv
import math
import pywt
from matplotlib import pyplot as plt

Lena_img_path = "Lena.bmp"
```

/usr/lib/python3/dist-packages/scipy/\_\_init\_\_.py:146: UserWarning: A NumPy version >=1.17.3 and <1.25.0 is required for this version of SciPy (detected version 1.26.2

warnings.warn(f"A NumPy version >={np\_minversion} and <{np\_maxversion}"
/home/majiddrn/.local/lib/python3.10/site-</pre>

packages/matplotlib/projections/\_\_init\_\_.py:63: UserWarning: Unable to import Axes3D. This may be due to multiple versions of Matplotlib being installed (e.g. as a system package and as a pip package). As a result, the 3D projection is not available.

warnings.warn("Unable to import Axes3D. This may be due to multiple versions of "

## 1.2 Exercise 1

The method discussed in the exercise, LoG, is mainly used as an approach which removes the effect of noise before laplacian application. By reversing the process, the result will be different. But to elaborate, We apply these two different approaches in several different ways. **All approaches are by (5, 5) filter sizes** 

First, we apply these two different approaches on grayscale image of Lena. We measure SSIM, PSNR and subtraction of two approaches.

```
[4]: lena_gray = cv.imread(Lena_img_path, cv.IMREAD_GRAYSCALE)
```

Well, computation is over. Let's plot the result images.

```
[5]: fig = plt.figure(figsize=(10, 7))
    fig.add_subplot(1, 3, 1)
    plt.imshow(lena_gray, cmap='gray')
    plt.axis("off")
    plt.title("The Lena itself")
    fig.add_subplot(1, 3, 2)
    plt.imshow(log_lena_gray, cmap='gray')
    plt.axis('off')
    plt.title('LoG approach')
    fig.add_subplot(1, 3, 3)
    plt.imshow(gol_lena_gray, cmap='gray')
    plt.axis('off')
    plt.axis('off')
    plt.axis('off')
    plt.title('First laplacian and then gaussian')
```

[5]: Text(0.5, 1.0, 'First laplacian and then gaussian')

The Lena itself



LoG approach



First laplacian and then gaussian



Great, we saw some differences between two approaches in two stated different implementations. But to make sure about it, let's measure PSNR, SSIM and difference of LoG and GoL.

The implementation of PSNR in code and mathematics is as follow.

$$PSNR = 10 \cdot \log_{10} \left( \frac{MAX^2}{MSE} \right)$$

Which MSE is mean square error.

$$SSIM(x,y) = \frac{(2\mu_x \mu_y + C_1)(2\sigma_{xy} + C_2)(\sigma_x^2 + \sigma_y^2 + C_3)}{(\mu_x^2 + \mu_y^2 + C_1)(\sigma_x^2 + \sigma_y^2 + C_2)(\sigma_x \sigma_y + C_3)}$$

Code implementation credit: CV Notes

```
[6]: def calculate_psnr(img1, img2):
    # img1 and img2 have range [0, 255]
    img1 = img1.astype(np.float64)
    img2 = img2.astype(np.float64)
    mse = np.mean((img1 - img2)**2)
    if mse == 0:
        return float('inf')
    return 20 * math.log10(255.0 / math.sqrt(mse))
```

```
[7]: def ssim(img1, img2):
        C1 = (0.01 * 255)**2
        C2 = (0.03 * 255)**2

        img1 = img1.astype(np.float64)
        img2 = img2.astype(np.float64)
        kernel = cv.getGaussianKernel(11, 1.5)
        window = np.outer(kernel, kernel.transpose())

        mu1 = cv.filter2D(img1, -1, window)[5:-5, 5:-5] # valid
        mu2 = cv.filter2D(img2, -1, window)[5:-5, 5:-5]
        mu1_sq = mu1**2
```

```
mu2_sq = mu2**2
    mu1_mu2 = mu1 * mu2
    sigma1_sq = cv.filter2D(img1**2, -1, window)[5:-5, 5:-5] - mu1_sq
    sigma2_sq = cv.filter2D(img2**2, -1, window)[5:-5, 5:-5] - mu2_sq
    sigma12 = cv.filter2D(img1 * img2, -1, window)[5:-5, 5:-5] - mu1_mu2
    ssim_map = ((2 * mu1_mu2 + C1) * (2 * sigma12 + C2)) / ((mu1_sq + mu2_sq + L))
 →C1) *
                                                             (sigma1_sq +
 ⇒sigma2_sq + C2))
    return ssim_map.mean()
def calculate_ssim(img1, img2):
    '''calculate SSIM
    the same outputs as MATLAB's
    img1, img2: [0, 255]
    111
    if not img1.shape == img2.shape:
        raise ValueError('Input images must have the same dimensions.')
    if img1.ndim == 2:
        return ssim(img1, img2)
    elif img1.ndim == 3:
        if img1.shape[2] == 3:
            ssims = []
            for i in range(3):
                ssims.append(ssim(img1, img2))
            return np.array(ssims).mean()
        elif img1.shape[2] == 1:
            return ssim(np.squeeze(img1), np.squeeze(img2))
    else:
        raise ValueError('Wrong input image dimensions.')
```

Well, let's calculate the PSNR and SSIM factors. First to compare the LoG approach with the Lena image and then for GoL approach and Lena image itself comparsion.

For Lena and its LoG approach comparsion: PSNR:5.814888863609525 and SSIM:-0.00022580376884029653

\_\_\_\_

For Lena and its GoL approach comparsion: PSNR:5.833012211438813 and SSIM:0.0008247144778262229

While there is not much difference in PSNR factor, we can see a noteable difference in their SSIM factor which obviously shows us the difference between these two approaches.

Let's simply compute the subtraction of two approach results.

```
[9]: log_gol_lena_diff = log_lena_gray - gol_lena_gray

plt.imshow(log_gol_lena_diff, cmap='gray')
plt.axis('off')
```

[9]: (-0.5, 511.5, 511.5, -0.5)



The above difference, simply shows the result of LoG and GoL(Gaussian of Laplacian) are not the same.

Anyway, let's examine the computation on color image of Lena as well.

```
[10]: lena_color = cv.imread(Lena_img_path)

# LoG approach for colored Lena

lena_color_gaussian = cv.GaussianBlur(lena_color, (5, 5), 0)
log_lena_color = cv.Laplacian(lena_color_gaussian, cv.CV_64F, laplacian_kernel)

# We have implemented the kernel above

# GoL approach for colored Lena

lena_color_laplacian = cv.Laplacian(lena_color, cv.CV_64F, laplacian_kernel)
gol_lena_color = cv.GaussianBlur(lena_color_laplacian, (5, 5), 0)
```

Computation is over now, let's plot the results.

```
[11]: fig = plt.figure(figsize=(10, 7))
      fig.add_subplot(1, 4, 1)
      lena_color_rgb = cv.cvtColor(lena_color, cv.COLOR_BGR2RGB) # To have the_
       ⇔lena_color in RGB space
      plt.imshow(lena_color_rgb)
      plt.axis("off")
      plt.title("The Lena itself")
      fig.add_subplot(1, 4, 2)
      plt.imshow(log_lena_color)
      plt.axis('off')
      plt.title('LoG approach')
      fig.add_subplot(1, 4, 3)
      plt.imshow(gol_lena_color)
      plt.axis('off')
      plt.title('GoL approach')
      fig.add_subplot(1, 4, 4)
      plt.imshow(log_lena_color - gol_lena_color)
      plt.axis('off')
      plt.title('difference')
```

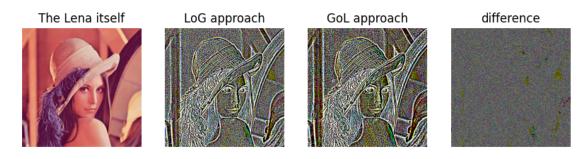
Clipping input data to the valid range for imshow with RGB data ([0..1] for floats or [0..255] for integers).

Clipping input data to the valid range for imshow with RGB data ([0..1] for floats or [0..255] for integers).

Clipping input data to the valid range for imshow with RGB data ([0..1] for

floats or [0..255] for integers).

[11]: Text(0.5, 1.0, 'difference')



Well, we saw the two approaches difference in color images as well.

### 1.3 Exercise 2

Clearly, applying the first derivation on the image gives us the edges. Horizontal derivation gives the vertical edges and vertical derivation gives us the horizontal edges. As stated, I myself will choose the first derivation vertically when I need the horizontal edges and I will use the first derivation horizontally when I need the vertical edges.

```
[12]: vertical_derivation_result = cv.imread('horizontal.png')
horizontal_derivation_result = cv.imread('vertical.png')

fig = plt.figure(figsize=(10, 7))

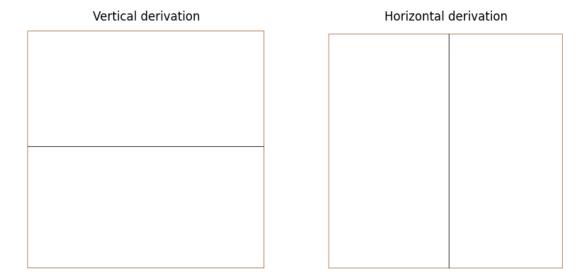
fig.add_subplot(1, 2, 1)

plt.imshow(vertical_derivation_result, cmap='gray')
plt.axis("off")
plt.title("Vertical derivation")

fig.add_subplot(1, 2, 2)

plt.imshow(horizontal_derivation_result, cmap='gray')
plt.axis("off")
plt.title("Horizontal derivation")
```

[12]: Text(0.5, 1.0, 'Horizontal derivation')



To elaborate, let's examine in code.

First of all we should implement the kernel needed for first derivative application and a function for convolution.

```
[13]: def convolution(img, kernel):
    width, height = img.shape
    width_k, height_k = kernel.shape

    output_height = height - height_k + 1
    output_width = width - width_k + 1

    output = np.zeros((output_height, output_width))

for i in range(output_height):
    for j in range(output_width):
        summation = np.sum(img[i:i + height_k, j:j + width_k] * kernel)
        output[i, j] = summation

return output
```

Well, let's apply the horizontal derivation and vertical derivation on the image.

Now, let's simply plot the results.

```
fig = plt.figure(figsize=(10, 7))
fig.add_subplot(1, 3, 1)

plt.imshow(square_gray, cmap='gray')
plt.axis("off")
plt.title("Image itself")

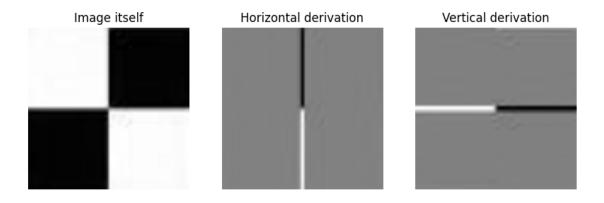
fig.add_subplot(1, 3, 2)

plt.imshow(first_derivative_horizontal_output, cmap='gray')
plt.axis("off")
plt.title("Horizontal derivation")

fig.add_subplot(1, 3, 3)

plt.imshow(first_derivative_vertical_output, cmap='gray')
plt.axis("off")
plt.title("Vertical derivation")
```

# [15]: Text(0.5, 1.0, 'Vertical derivation')



Well, We saw the results are as expected.

## 1.4 Exercise 3

First of all we should compute the frequency of each gray scale in the image. After that, the probablity and PDF of each. Finally, by multiplying each PDF by 255(which is L-1), we can have new intensities.

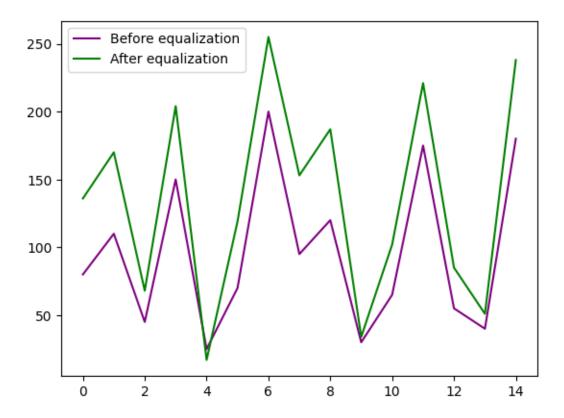
				$PDF \times 255 = $
Gray Level	Frequency	Probability	PDF	new gray level
25	1	1/15	1 15	17
30	1	$\frac{1}{15}$	$ \begin{array}{c} 1\\15\\2\\15\\3\\15\\4\\15\\15\\6\\15\\7\\15\\8\\15\\9\\15\\10\\15\\15\\15\\15\\15\\15\\15\\15\\15\\15\\15\\15\\15\\$	34
40	1	$\frac{1}{15}$	$\frac{\frac{3}{3}}{15}$	51
45	1	$\frac{\widetilde{1}}{15}$	$\frac{\frac{3}{4}}{15}$	68
55	1	$\frac{\widetilde{1}}{15}$	$\frac{5}{15}$	85
65	1	$\frac{\overline{1}}{15}$	$\frac{6}{15}$	102
70	1	$\frac{1}{15}$	$\frac{7}{15}$	119
80	1	$\frac{\overline{1}}{15}$	$\frac{8}{15}$	136
95	1	$\frac{\overline{1}}{15}$	$\frac{9}{15}$	153
110	1	$\frac{1}{15}$	$\frac{\overline{10}}{15}$	170
120	1	$\frac{1}{15}$	$\frac{11}{15}$	187
150	1	$\frac{1}{15}$	$\frac{12}{15}$	204
175	1	$\frac{1}{15}$	$\frac{13}{15}$	221
180	1	$\frac{1}{15}$	$\frac{14}{15}$	238
200	1	$\begin{array}{c} \frac{1}{15} \\ \frac{1}$	$\frac{15}{15} = 1$	255

The image before and after equalization is as follow.

```
[16]: image_before_equal = np.array([80, 110, 45, 150, 25, 70, 200, 95, 120, 30, 65, u 4175, 55, 40, 180])
image_after_equal = np.array([136, 170, 68, 204, 17, 119, 255, 153, 187, 34, u 4102, 221, 85, 51, 238])

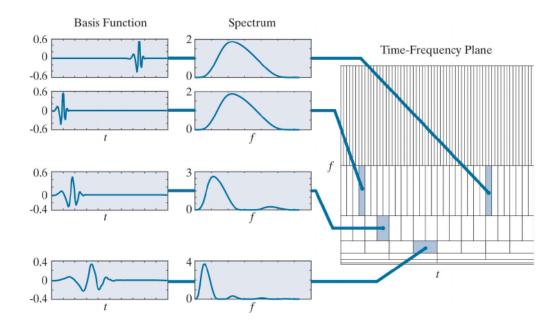
plt.plot(image_before_equal, color='purple', label='Before equalization')
plt.plot(image_after_equal, color='green', label='After equalization')
plt.legend(loc='upper left')
```

[16]: <matplotlib.legend.Legend at 0x7f19e1d3ba00>



## 1.5 Exercise 4

In wavelet transform, we try to have both frequency and spatial information of the image with the least cost. As a result, there would be a kernel with a variable size, moving across the signal (here, the signal is our image). The kernel computes the frequency of each area it consists in every iteration. One important factor in the iteration is that the size of wavelet window is dependent on the resolution and frequency of that area. If the area has high frequencies, the window size will be wider. Otherwise, if the area has lower frequencies, the spatial information to be remained, window size will be narrower. Well, It was simply the decomposition phase.



After decomposition, we have a set of wavelet coefficients. Quantization is the process of mapping these coefficients to a smaller set of values, which can be represented with fewer bits.

### 1.6 Exercise 5

To downsample the image, we simply drop rows/columns decussately. To acheive this goal, we implement a function for the rest.

```
[17]: def downsample2(img):
    rows_keep = img.shape[0] // 2
    cols_keep = img.shape[1] // 2

    downsampled_img = img[:rows_keep * 2, :cols_keep * 2]
    downsampled_img = downsampled_img.reshape((rows_keep, 2, cols_keep, 2))
    downsampled_img = downsampled_img.mean(axis=(1, 3))

    return downsampled_img
```

Now, let's impelement a function for upsampling the image with the replication or bilinear interpolation method.

```
[18]: def upsample2(img, method):
    upsampled_img = np.zeros((img.shape[0] * 2, img.shape[1] * 2), dtype=img.
    dtype)
    img_ = img.astype(np.float64)

if method == 'replication':
    for i in range(img_.shape[0]):
```

```
for j in range(img_.shape[1]):
              upsampled_img[i * 2, j * 2] = img_[i, j]
              upsampled_img[i * 2 + 1, j * 2] = img_[i, j]
              upsampled_img[i * 2, j * 2 + 1] = img_[i, j]
              upsampled_img[i * 2 + 1, j * 2 + 1] = img_[i, j]
  elif method == 'bilinear':
      for i in range(img_.shape[0]):
          for j in range(img_.shape[1]):
              upsampled_img[i * 2, j * 2] = img_[i, j]
              # Handle the last column and last row separately
              if j < img_.shape[1] - 1:</pre>
                  upsampled_img[i * 2, j * 2 + 1] = (img_[i, j] + img_[i, j +_{\sqcup}
→1]) / 2
              else:
                  upsampled_img[i * 2, j * 2 + 1] = img_[i, j]
              if i < img_.shape[0] - 1:</pre>
                  ⇒j]) / 2
              else:
                  upsampled_img[i * 2 + 1, j * 2] = img_[i, j]
              if i < img_.shape[0] - 1 and j < img_.shape[1] - 1:
                  upsampled_img[i * 2 + 1, j * 2 + 1] = (img_[i, j] + img_[i_\square
\rightarrow + 1, j] + img_[i, j + 1] + img_[i + 1, j + 1]) / 4
              elif i < img .shape[0] - 1:</pre>
                  upsampled_img[i * 2 + 1, j * 2 + 1] = (img_[i, j] + img_[i_\square
\hookrightarrow+ 1, j]) / 2
              elif j < img_.shape[1] - 1:</pre>
                  \rightarrowj + 1]) / 2
              else:
                  upsampled_img[i * 2 + 1, j * 2 + 1] = img_[i, j]
  return upsampled_img
```

Well, let's pass the Lena image to the functions above.

Now, let's plot the results.

```
fig = plt.figure(figsize=(10, 7))
fig.add_subplot(1, 3, 1)

plt.imshow(lena_gray, cmap='gray')
plt.axis("off")
plt.title("The Lena itself")

fig.add_subplot(1, 3, 2)

plt.imshow(lena_upsampled2_replication, cmap='gray')
plt.axis('off')
plt.title('Replication method')

fig.add_subplot(1, 3, 3)

plt.imshow(lena_upsampled2_bilinear, cmap='gray')
plt.axis('off')
plt.title('Bilinear interpolation method')
```

[20]: Text(0.5, 1.0, 'Bilinear interpolation method')







Let's simply compute the difference of two method results.

```
[21]: plt.imshow(lena_upsampled2_replication - lena_upsampled2_bilinear, cmap='gray')
   plt.axis("off")

[21]: (-0.5, 1023.5, 1023.5, -0.5)
```



The difference, obviously shows that the two methods have noteable differences.

## 1.7 Exercise 6

In this exercise, we simply use the previous convolution function implemented for the spatial domain. And for the fourier computations, we simply implement the following function to calculate the fourier transform of any filter.

```
[22]: def fourier__magnitude(img_in, shape=(0, 0)):
    if (shape == (0, 0)):
        fdft = np.fft.fft2(img_in)
    else:
        fdft = np.fft.fft2(img_in, shape)
    fdftshift = np.fft.fftshift(fdft)
    magnitude_spectrum_fdft = np.log(np.abs(fdftshift))
    return (fdftshift, magnitude_spectrum_fdft)

def fourier__pad_kernel(kernel, new_size):
    fdftshift__kernel, = fourier__magnitude(kernel, new_size)
    return fdftshift__kernel
```

Now, let's implement a function to compute the application of a the filters on the image in frequency domain.

```
[23]: def apply_filter_freq(img, kernel):
    img__, = fourier__magnitude(img)

    result_freq = img__ * kernel

    return np.uint8(np.abs(np.fft.ifft2(np.fft.ifftshift(result_freq))))
```

Now, with the help of the functions we implemented before, let's apply the filters on the image.

```
[24]: # We simply implement the kernels we need
      kernel_A = np.array([[1/9, 1/9, 1/9],
                            [1/9, 1/9, 1/9],
                            [1/9, 1/9, 1/9]])
      kernel_B = np.array([[-1, -1, -1],
                            [-1, 8, -1],
                            [-1, -1, -1]
      kernel_C = np.array([[0, -1, 0],
                            [-1, 5, -1],
                            [0, -1, 0]
      # Now, let's apply the kernel A on the Lena image with two spatial and fourier
       \hookrightarrowapproaches
      kernel_A_fourier = fourier__pad_kernel(kernel_A, lena_gray.shape)
      lena_A_spatial_builtin = cv.filter2D(lena_gray, -1, kernel_A)
      lena__A_fourier = apply_filter_freq(lena_gray, kernel_A_fourier)
      # Now, let's apply the kernel B on the Lena image with two spatial and fourier
       \hookrightarrowapproaches
      kernel_B_fourier = fourier__pad_kernel(kernel_B, lena_gray.shape)
      lena_B_spatial_builtin = cv.filter2D(lena_gray, -1, kernel_B)
      lena_B_fourier = apply_filter_freq(lena_gray, kernel_B_fourier)
      # Now, let's apply the kernel C on the Lena image with two spatial and fourier
      \rightarrowapproaches
      kernel_C_fourier = fourier__pad_kernel(kernel_C, lena_gray.shape)
      lena__C_spatial_builtin = cv.filter2D(lena_gray, -1, kernel_C)
```

```
lena__C_fourier = apply_filter_freq(lena_gray, kernel_C_fourier)
```

 $\label{lem:log-continuous} $$ / tmp/ipykernel_4642/1749126714.py:7: RuntimeWarning: divide by zero encountered in log$ 

```
magnitude_spectrum_fdft = np.log(np.abs(fdftshift))
```

Now, let's plot the result of applying kernel A on the image in both spatial and frequency domain.

```
[25]: fig = plt.figure(figsize=(10, 7))
    fig.add_subplot(1, 3, 1)

plt.imshow(lena_gray, cmap='gray')
plt.axis("off")
plt.title("The Lena itself")

fig.add_subplot(1, 3, 2)

plt.imshow(lena__A_spatial_builtin, cmap='gray')
plt.axis('off')
plt.title('Box filter builtin conv')

fig.add_subplot(1, 3, 3)

plt.imshow(lena__A_fourier, cmap='gray')
plt.axis('off')
plt.title('Box filter freq')
```

[25]: Text(0.5, 1.0, 'Box filter freq')







Now, it's the kernel B turn.

```
[26]: fig = plt.figure(figsize=(10, 7))
```

```
fig.add_subplot(1, 3, 1)

plt.imshow(lena_gray, cmap='gray')
plt.axis("off")
plt.title("The Lena itself")

fig.add_subplot(1, 3, 2)

plt.imshow(lena__B_spatial_builtin, cmap='gray')
plt.axis('off')
plt.title('Laplacian builtin conv')

fig.add_subplot(1, 3, 3)

plt.imshow(lena__B_fourier, cmap='gray')
plt.axis('off')
plt.title('Laplacian freq')
```

# [26]: Text(0.5, 1.0, 'Laplacian freq')







And finally, it's the kernel C's turn which is a simple sharpenning kernel.

```
[27]: fig = plt.figure(figsize=(10, 7))
    fig.add_subplot(1, 3, 1)

plt.imshow(lena_gray, cmap='gray')
    plt.axis("off")
    plt.title("The Lena itself")

fig.add_subplot(1, 3, 2)

plt.imshow(lena__C_spatial_builtin, cmap='gray')
    plt.axis('off')
```

```
plt.title('Sharpenning builtin conv')

fig.add_subplot(1, 3, 3)

plt.imshow(lena__C_fourier, cmap='gray')
plt.axis('off')
plt.title('Sharpenning freq')
```

[27]: Text(0.5, 1.0, 'Sharpenning freq')

The Lena itself







#### 1.8 Exercise 7

We have the box filter as follow.

$$K = \begin{pmatrix} \frac{1}{4} & \frac{1}{4} \\ \frac{1}{4} & \frac{1}{4} \end{pmatrix}$$

Let's make the pyramid as wanted.

Computation is over, let's plot the pyramid.

```
[29]: fig, axes = plt.subplots(nrows=3, ncols=2, figsize=(8, 8))

approximation_pyramid = [approx__L1, approx__L2, approx__L3]
prediction_residual_pyramid = [prediction__L1, prediction__L2, prediction__L3]

for level in range(3):
    axes[level, 0].imshow(approximation_pyramid[level], cmap='gray')
    axes[level, 0].set_title(f'Level {level+1} Approximation')

axes[level, 1].imshow(prediction_residual_pyramid[level], cmap='gray')
    axes[level, 1].set_title(f'Level {level+1} Prediction Residual')

for ax in axes.flat:
    ax.axis('off')

plt.tight_layout()
plt.show()
```

Level 1 Approximation



Level 2 Approximation



Level 3 Approximation



Level 1 Prediction Residual



Level 2 Prediction Residual



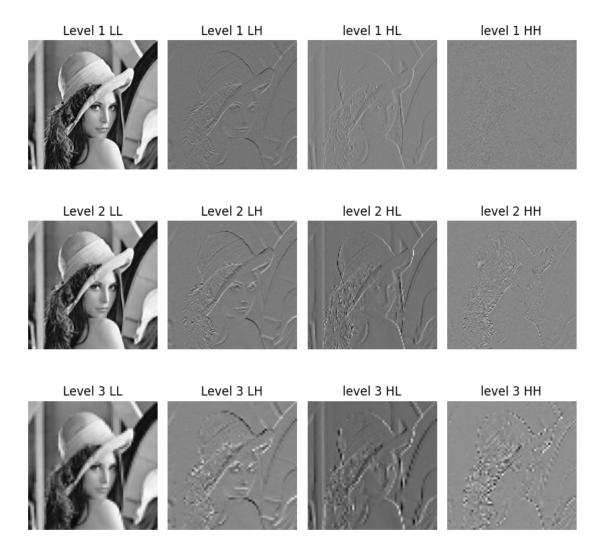
Level 3 Prediction Residual



## 1.9 Exercise 8

To acheive this goal, we simply install the library pywt. After the installation, usage is so easy. We store every level as follow.

```
[48]: fig, axes = plt.subplots(nrows=3, ncols=4, figsize=(8, 8))
      img__LL = lena_gray
      for level in range(3):
          coeffs = pywt.dwt2(img__LL, 'haar')
          cA, (cH, cV, cD) = coeffs
          axes[level, 0].imshow(cA, cmap='gray')
          axes[level, 0].set_title(f'Level {level+1} LL')
          axes[level, 1].imshow(cH, cmap='gray')
          axes[level, 1].set_title(f'Level {level+1} LH')
          axes[level, 2].imshow(cV, cmap='gray')
          axes[level, 2].set_title(f'level {level+1} HL')
          axes[level, 3].imshow(cD, cmap='gray')
          axes[level, 3].set_title(f'level {level+1} HH')
          img_{LL} = cA
      for ax in axes.flat:
          ax.axis('off')
      plt.tight_layout()
      plt.show()
```



Well, let's discuss on the differences.

- The wavelet transform, gives more infromation which contains vertical, horizontal and diagonal edges. While, the pyramid in the 7th question, doesn't give us horizontal or vertical edges seperately.
- The Haar wavelet transform provides a multi-resolution representation of the image, prepares both low-frequency and high-frequency details at different scales. This is useful for some image processing tasks, such as compression, denoising and feature extraction.
- Clearly, the Haar wavelet transform offers a more sophisticated analysis of the image content compared to a simple 2x2 filter.

## 1.10 Exercise 9

After some little changes in the upsample2(img, method) function implemented in the exercise 5, I added two other interpolation techniques, including bilinear and nearest neighbor interpolation.

There has been added another argument named scale\_factor which upsamples the image in any scale needed.

```
[88]: def upsample(img, method, scale_factor):
          upsampled_img = np.zeros((int(img.shape[0] * scale_factor), int(img.
       ⇒shape[1] * scale_factor)), dtype=img.dtype)
          img_ = img.astype(np.float64)
          for i in range(img_.shape[0]):
              for j in range(img_.shape[1]):
                  for p in range(int(scale_factor)):
                      for q in range(int(scale_factor)):
                          if method == 'replication':
                              upsampled_img[i * int(scale_factor) + p, j *__
       sint(scale_factor) + q] = img_[i, j]
                          elif method == 'bilinear':
                              upsampled_img[i * int(scale_factor) + p, j *__
       int(scale_factor) + q] = bilinear_interpolation(img_, i, j, p /__
       ⇔scale_factor, q / scale_factor)
                          elif method == 'nearest':
                              upsampled_img[i * int(scale_factor) + p, j *__
       →int(scale_factor) + q] = nearest_neighbor_interpolation(img_, i + p / _ _
       ⇔scale_factor, j + q / scale_factor)
          return upsampled_img
      def bilinear_interpolation(img, i, j, p, q):
          i_floor, j_floor = int(np.floor(i)), int(np.floor(j))
          i_ceil, j_ceil = min(i_floor + 1, img.shape[0] - 1), min(j_floor + 1, img.
       ⇒shape[1] - 1)
          top_left = img[i_floor, j_floor]
          top_right = img[i_floor, j_ceil]
          bottom_left = img[i_ceil, j_floor]
          bottom_right = img[i_ceil, j_ceil]
          interpolated_value = (1 - p) * (1 - q) * top_left + (1 - p) * q * top_right_u
       \rightarrow+ p * (1 - q) * bottom_left + p * q * bottom_right
          return interpolated_value
      def nearest_neighbor_interpolation(img, i, j):
          i round, j round = int(round(i)), int(round(j))
          i_round = max(0, min(img.shape[0] - 1, i_round))
          j_round = max(0, min(img.shape[1] - 1, j_round))
          return img[i_round, j_round]
```

Implementation is over. Let's first downscale the image and then upscale the image by factor 5. First by bilinear interpolation and then by nearest neighbor interpolation.

```
[99]: lena_up5_bilinear = upsample(lena_gray, method='replication', scale_factor=5) lena_up5_nearest = upsample(lena_gray, method='nearest', scale_factor=5)
```

Application is over. Let's plot the result.

```
[100]: fig = plt.figure(figsize=(10, 7))

fig.add_subplot(1, 2, 1)

plt.imshow(lena_up5_bilinear, cmap='gray')
plt.axis("off")
plt.title("Upscaled 3 times- bilinear interpolation")

fig.add_subplot(1, 2, 2)

plt.imshow(lena_up5_nearest, cmap='gray')
plt.axis('off')
plt.title('Upscaled 3 times- nearest neighbor interpolation')
```

[100]: Text(0.5, 1.0, 'Upscaled 3 times- nearest neighbor interpolation')

Upscaled 3 times- bilinear interpolation



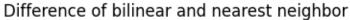
Upscaled 3 times- nearest neighbor interpolation



Well, to compare the results, let's subtract one from the other.

```
[101]: plt.imshow(lena_up5_bilinear - lena_up5_nearest, cmap='gray')
plt.axis("off")
plt.title("Difference of bilinear and nearest neighbor")
```

[101]: Text(0.5, 1.0, 'Difference of bilinear and nearest neighbor')





Well, the difference is clear, let's examine again by builtin OpenCV functions.

Let's plot

```
[106]: fig = plt.figure(figsize=(10, 7))

fig.add_subplot(1, 2, 1)

plt.imshow(lena_up5_nearest_builtin, cmap='gray')
plt.axis("off")
plt.title("-builtin function-bilinear interpolation")

fig.add_subplot(1, 2, 2)

plt.imshow(lena_up5_bilinear_builtin, cmap='gray')
```

```
plt.axis('off')
plt.title('-builtin function- nearest neighbor interpolation')
```

[106]: Text(0.5, 1.0, '-builtin function- nearest neighbor interpolation')

-builtin function- bilinear interpolation



-builtin function- nearest neighbor interpolation

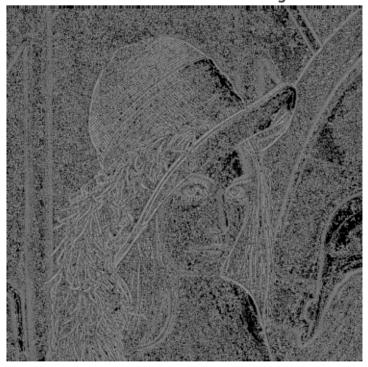


And now, let's subtract one from the other.

```
[107]: plt.imshow(lena_up5_bilinear_builtin - lena_up5_nearest_builtin,u cmap='gray')
plt.axis("off")
plt.title("Difference of bilinear and nearest neighbor -builtin-")
```

[107]: Text(0.5, 1.0, 'Difference of bilinear and nearest neighbor -builtin-')

Difference of bilinear and nearest neighbor -builtin-



Generally, if speed is most important, nearest neighbor interpolation can be a good choice. However, if image quality is paramount, bilinear interpolation will generally yield better results.