50.039 Theory and Practice of Deep Learning

W4-S1 Introduction to Computer Vision and Convolutional Neural Networks

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- 1. What are **images** and how is this datatype represented?
- 2. What is a pixel and how can its information be interpreted?
- 3. What is the **spatial dependence property** of pixels?
- 4. What is the **homophily property** of pixels?
- 5. Why is the linear processing operation failing on images?
- 6. What is the **convolution** operation?
- 7. How can we perform **image processing using convolutions**?
- 8. What is **padding** in convolutions?

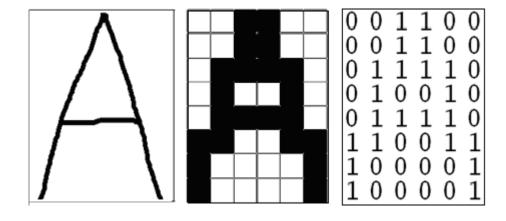
- 9. What is **stride** in convolution?
- 10. What is **dilation** in convolution?
- 11. How does convolution apply to higher dimensional images?
- 12. How to write our own convolutional processing layer in PyTorch?
- 13. What is Conv2d in PyTorch?
- 14. What is a **Convolutional Neural Network** (CNN)?
- 15. What is the **intuition** behind using Convolutional layers in a Convolutional Neural Network?

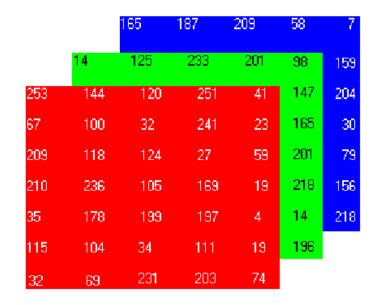
- 16. How to train our first Convolutional Neural Network on MNIST?
- 17. What are **other typical image processing layers** in Computer Vision and how are they implemented?
- 18. What is the **pooling layer**?
- 19. What is the **batchnorm layer**?
- 20. What is the **dropout layer**?
- 21. What is data augmentation in Computer Vision?
- 22. What are some **milestone Computer Vision models** and their **contributions** to the field of Deep Learning?

- 23. What are the **AlexNet** and **VGG models**?
- 24. What is a **skip connection/residual**? What is its effect on a Neural Network and **how does it help with vanishing gradient problems**?
- 25. What are **ResNet** and **DenseNet models**?
- 26. What is the **Inception model**?
- 27. What is the **EfficientNet model**?
- 28. What is **transfer learning** and its uses?
- 29. How to **freeze** and **fine-tune layers** in a Neural Network?

 Both grayscale and RGB (Red Green Blue) images can be represented as matrices or tensors.

• Typically, our sample images in the MNIST dataset were 2D matrices of shape 28×28 .

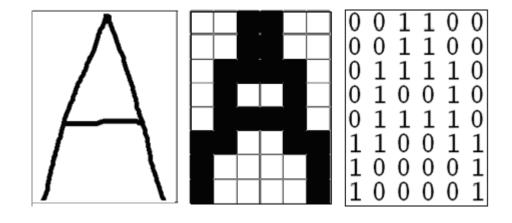




Definition (encoding a grayscale image):

Grayscale images can be represented as a **2D matrix** of shape $height \times width$, where each element is an 8-bit integer.

- Each **element** of the matrix represents the **intensity of light** at that pixel.
- The intensity of light can be represented as a single number between 0 and 255, where **0** represents no light (**black**) and **255** represents maximum light intensity (**white**). Sometimes normalized to [0, 1].

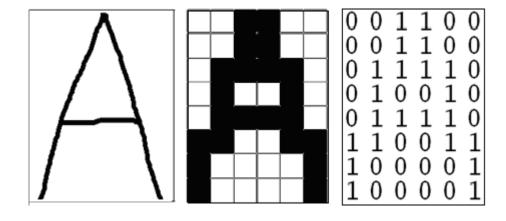


		165	187	209	58	7
	14	1 25	233	201	98	159
253	144	120	251	41	147	204
67	100	32	241	23	165	30
209	118	124	27	59	201	79
210	236	105	169	19	21 8	156
35	178	199	197	4	14	218
115	104	34	111	19	196	
32	69	231	203	74		

Definition (encoding a RGB image):

RGB images are more complex because each pixel has three values representing the intensity of light for each color channel (red, green, and blue).

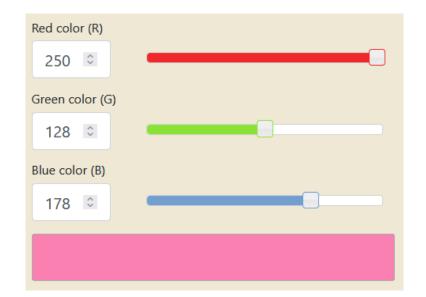
- An RGB image is represented as a 3D tensor of shape height × width × 3.
- Each element in the tensor would be an integer representing the intensity of the corresponding color channel.

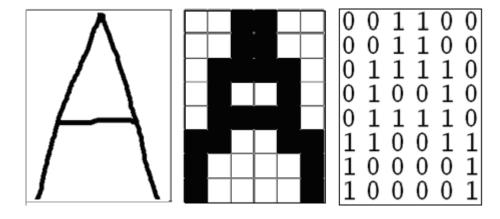


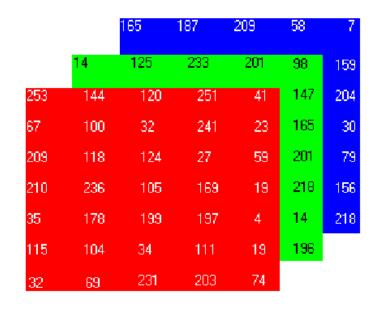
		165	187	209	58	7
	14	125	233	201	98	159
253	144	120	251	41	147	204
67	100	32	241	23	165	30
209	118	124	27	59	201	79
210	236	105	169	19	218	156
35	178	199	197	4	14	218
115	104	34	111	19	196	
32	69	231	203	74		

 Combining the red, green and blue values allows to generate any color. Try it!

https://www.rapidtables.com/convert/color/rgb-to-hex.html







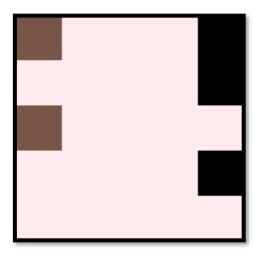
- Each element in a Greyscale matrix or RGB matrix corresponds to a pixel, representing something in the image.
- The problem, however, is that it is often impossible to guess the meaning of a given pixel, by looking at this pixel alone.

→ Can you guess what this pixel represents and what object it belongs to?

- 1. Flower petal
- 2. Piece of clothing
- 3. Skin of a human person
 - 4. Feather of a bird

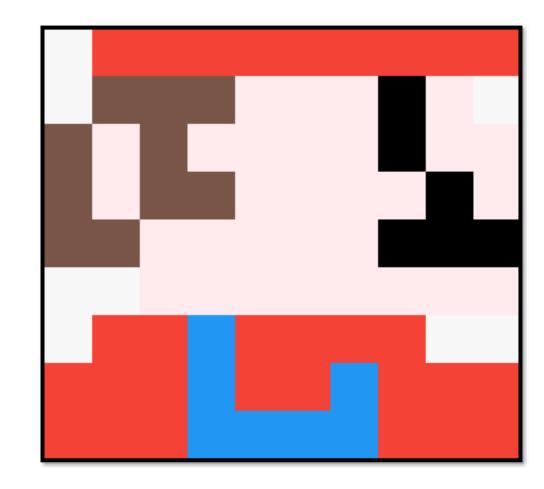
→ What if I show you <u>some</u> neighbouring pixels now?

- 1. Flower petal
- 2. Piece of clothing
- 3. Skin of a human person
 - 4. Feather of a bird



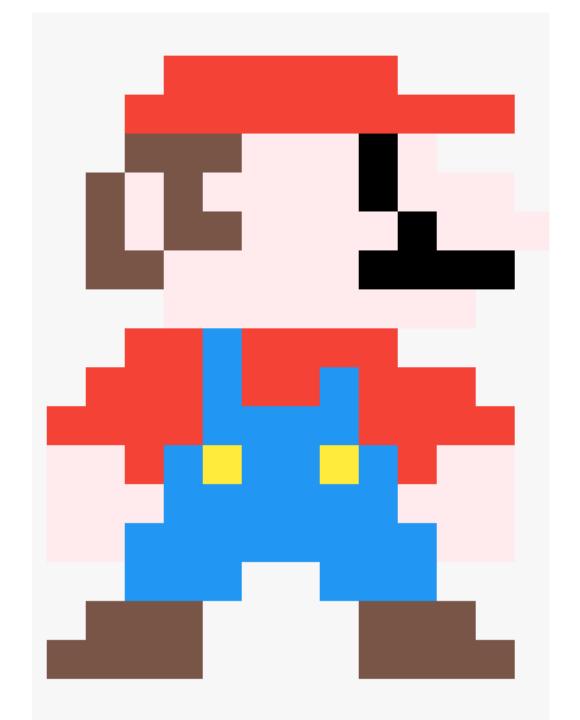
→ What if I show you MORE neighbouring pixels now?

- 1. Flower petal
- 2. Piece of clothing
- 3. Skin of a human person
 - 4. Feather of a bird

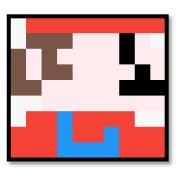


→ What if I show you MORE neighbouring pixels now?

- 1. Flower petal
- 2. Piece of clothing
- 3. Skin of a human person
 - 4. Feather of a bird



The spatial dependence property



Definition (the spatial dependence property of pixels):

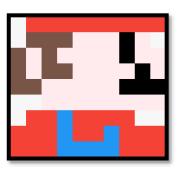
The spatial dependence property of pixels states that the meaning or interpretation of a single pixel in an image can often only be understood in the context of the pixels around it.

This is because the information in a single pixel is often not enough to describe the complete picture, or even just the pixel itself.

The relationship between the given pixel and its neighbouring pixels is necessary to provide a full understanding of this particular pixel, or even the image as a whole.

Images have "structure", which is something we observed in the Mario pixel art example.

The homophily property



Definition (the homophily property of pixels):

Homophily in images refers to the phenomenon where pixels that are close to each other in space tend to have similar properties, such as color, texture, or intensity. In other words, is the tendency for similar pixels to be grouped together.

For example, in an image of a sky, pixels that are close to each other and have a similar blue color can often be grouped together and be considered part of the sky region.

This property is a **key factor in image processing** and **computer vision** algorithms, as it provides a way to segment images into meaningful regions and perform tasks like object recognition and image classification.

The problem with Linear processing

So far we have only used Linear operations in our neural networks, even when playing with images, as in the MNIST dataset.

- We would flatten the image, assembling all pixels in a 1D vector, essentially losing the "structure" or spatial information of the image.
- Then, Linear processing layers would simply process pixels independently, applying individual coefficients to each pixel separately.

$$Z = \sum_{i} \sum_{j} w_{i,j} x_{i,j} + b$$

With $x_{i,j}$ the pixel value at location (i,j), $w_{i,j}$ the weight coefficient for the Linear layer assigned to this pixel, and b the bias.

The problem with Linear processing

Linear layers are <u>not great at processing images</u> because they do not take into account the <u>spatial dependence</u> of pixels.

- Pixels in images are typically related to each other in a spatial manner, i.e. the meaning of a pixel depends on the neighbour pixels.
- Unfortunately, linear models would assume each pixel is independent of the others and process them independently, which is not correct.
- As a result, linear layers can miss important spatial relationships between pixels. This can, in turn, result in poor image representation and interpretation.
- → Need for <u>a more specific processing operation for images</u>, to use in place of the Linear one!

Definition (convolution):

Convolution is a **mathematical operation** that is widely used in image processing, computer vision, and deep learning.

It involves applying a small matrix called a **convolution kernel** or **filter** K, over a given image X, element-wise multiplying each overlapping set of values in the image with the kernel, and then summing the results.

The result of this operation is a new image Y, where each output pixel represents the sum of the product of the corresponding input pixel and the filter.

$$Y = f(X, K)$$

Let us consider

- A grayscale image, defined as a 2D matrix X of size $h \times w$, with pixel values $(X_{i,j}) \in [0, 255]$.
- A convolution kernel K of size $k \times k$, with values $(K_{i,i}) \in \mathbb{R}$.

First, let us discuss sizes...

The convolution operation defined as Y = f(X, K), produces an image Y of size $h' \times w'$,

- With h' = h k + 1,
- And w' = w k + 1.

Note: we often use a kernel defined as a square matrix, with odd sizes (i.e. 3×3 , 5×5 , etc.). The operation, however, remains valid for other kernel sizes (but some notations showed in next slide would have to be adjusted).

Second, the pixel values $y_{i,i}$ for image Y are calculated using the convolution operation, defining $Y_{i,i}$, $\forall i \in [1, h'], j \in [1, w']$ as:

$$Y_{i,j} = \frac{1}{k^2} \sum_{m=1}^{k} \sum_{n=1}^{k} X_{i+m-1, j+n-1} K_{m,n}$$

This operation might seem complicated, but is easily visualised!

1	2	4	2	1
4	7	3	2	1
4	5	2	3	1
2	1	7	8	4
3	2	4	7	8

X is 5 by 5

K is 3 by 3

(5-3+1=3)

1	0	
0	2	
1	0	

 :	:

Second, the pixel values $y_{i,j}$ for image Y are calculated using the convolution operation, defining $Y_{i,j}$, $\forall i \in [1,h'], j \in [1,w']$ as:

$$Y_{i,j} = \frac{1}{k^2} \sum_{m=1}^{k} \sum_{n=1}^{k} X_{i+m-1, j+n-1} K_{m,n}$$

This operation is a weighted average of a pixel and its neighbouring pixels!

1	2	4	2	1
4	7	3	2	1
4	5	2	3	1
2	1	7	8	4
3	2	4	7	8

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+4	×	1
+4	×	0
+7	×	2
+3	×	1
+4	×	1
+5	×	0
+2	×	1
=	28	3

1	0	1
0	2	1
1	0	1

V

28 9		
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Second, the pixel values $y_{i,j}$ for image Y are calculated using the convolution operation, defining $Y_{i,j}$, $\forall i \in [1,h'], j \in [1,w']$ as:

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1	2	4	2	1
4	7	3	2	1
4	5	2	3	1
2	1	7	8	4
3	2	4	7	8

+2	X	U
+3	×	1
+1	×	0
+7	×	2
+8	×	1
+2	×	1
+4	×	0
+7	×	1
=	39)

K

1	0	1
0	2	1
1	0	1

Y

$\frac{28}{9}$	•••	•••
		:
	39 9	

Second, the pixel values $y_{i,j}$ for image Y are calculated using the convolution operation, defining $Y_{i,j}$, $\forall i \in [1, h'], j \in [1, w']$ as:

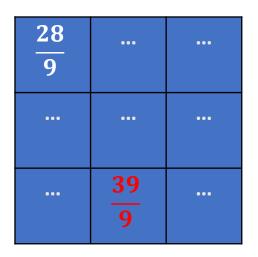
$$Y_{i,j} = \sum_{m=1}^{k} \sum_{n=1}^{k} X_{i+m-1, j+n-1} K_{m,n}$$

Note: Sometimes, the **normalization term** is removed and we may instead **normalize the kernel** (ensuring elements in kernel matrix sum up to 1), or not.

1	2	4	2	1
4	7	3	2	1
4	5	2	3	1
2	1	7	8	4
3	2	4	7	8

1	0	1
0	2	1
1	0	1

5 :	× 1
+2	× (
+3	X
+1	\times (
+7	X
+8	X
+2	X
	× (
+7	X
=	39



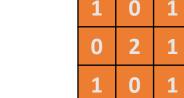
Second, the pixel values $y_{i,j}$ for image Y are calculated using the convolution operation, defining $Y_{i,j}$, $\forall i \in [1, h'], j \in [1, w']$ as:

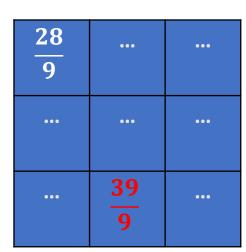
$$Y_{i,j} = \sum_{m=1}^{1} \sum_{m=1}^{k} X_{i+m-1, j+n-1} K_{m,n}$$

→ At the end of the day, convolution is just another type of matrix multiplication!

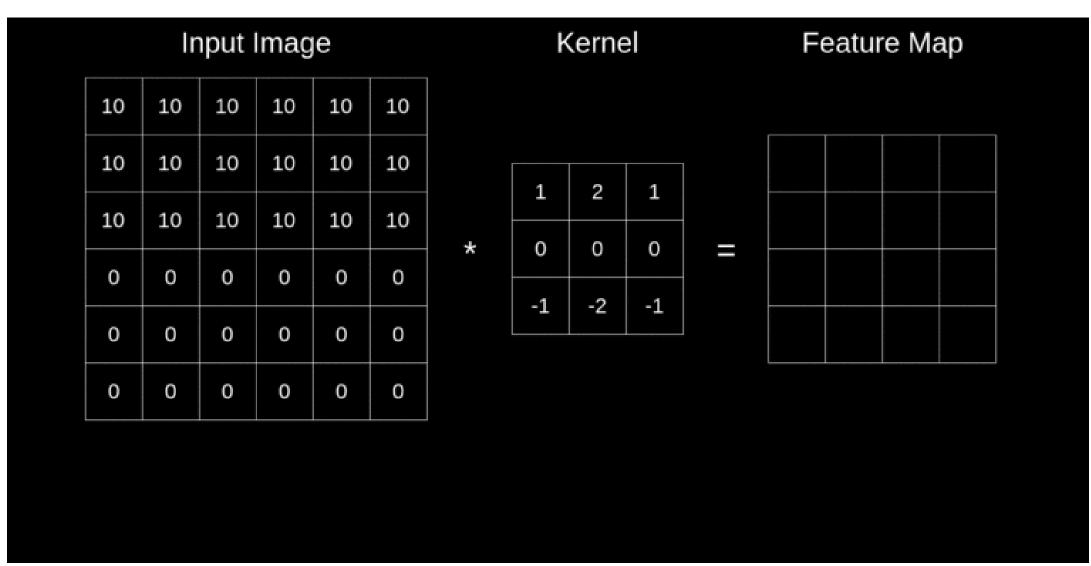
1	2	4	2	1
4	7	3	2	1
4	5	2	3	1
2	1	7	8	4
3	2	4	7	8

5 >	X [L
+2	×	0
+3	×	1
+1	×	0
+7	×	2
+8	×	1
+2	×	1
+4	×	0
+7	×	1
=	39)





An animated visualisation of Convolution



Convolution and Image Processing courses

Convolution is a **key mathematical operation** that is widely used in computer vision and image processing to apply a wide range of **transformations on an image** (enhancing contrast, blurring, edge detection, Instagram filters, etc).

- If **Computer Vision** is a direction that interests you, I would strongly advise to take a course on Image Processing (with implementations in OpenCV, Photoshop, etc.) during your continued learning.
- For instance: https://www.udemy.com/course/digital-image-processing-from-ground-up-in-python/

Let us try the convolution operation on images to see their effect.

• First, we need a **greyscale image**.

Restricted

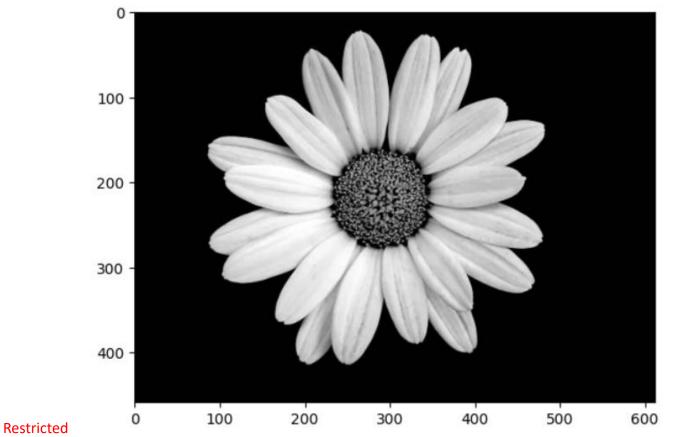
```
# Open the image and convert it to grayscale
im = Image.open('flower.jpg').convert('L')

# Convert the image to a NumPy array
im_array = np.array(im)

# Print the shape of the array
print(im_array.shape)
```

(459, 612)

```
# Display image in matplotlib
plt.imshow(im_array, cmap = 'gray')
plt.show()
```



Restricted

Testing convolution on images

Let us try the convolution operation on images to see their effect.

- First, we need a greyscale image.
- Second, we need a convolution function.

```
# Our convolution function
def convolution(image, kernel):
   # Flip the kernel (optional)
   kernel = np.flipud(np.fliplr(kernel))
   # Get the dimensions of the image and kernel
   image rows, image cols = image.shape
   kernel rows, kernel cols = kernel.shape
   # Convolve using Numpy
   output = correlate(image, kernel, mode = 'valid')
   # Note that this is equivalent to this
   # Loop through the image, applying the convolution
   output = np.zeros like(image)
   for x in range(image rows - kernel rows + 1):
       for y in range(image cols - kernel cols + 1):
            output[x, y] = (kernel * image[x:x+kernel_rows, y:y+kernel_cols]).sum()
   return output
```

What is the purpose of flipping the kernel?

Optional, but in short, flipping the kernel has mathematical and practical advantages that make it a common practice in deep learning.

(It has to do with signal processing, Fast Fourier Transform and the convolution theorem, out-of-scope).

```
# Our convolution function
def convolution(image, kernel):
  # Flip the kernel (optional)
   kernel = np.flipud(np.fliplr(kernel))
    # Get the dimensions of the image and kernel
   image rows, image cols = image.shape
   kernel rows, kernel cols = kernel.shape
   # Convolve using Numpy
   output = correlate(image, kernel, mode = 'valid')
   # Note that this is equivalent to this
   # Loop through the image, applying the convolution
   output = np.zeros like(image)
   for x in range(image rows - kernel rows + 1):
       for y in range(image cols - kernel cols + 1):
            output[x, y] = (kernel * image[x:x+kernel_rows, y:y+kernel_cols]).sum()
   return output
```

https://stackoverflow.com/questions/ /45152473/why-is-theconvolutional-filter-flipped-inconvolutional-neural-networks

Let us try the convolution operation on images to see their effect.

- First, we need a greyscale image.
- Second, we need a convolution function.
- The magic for the transformation then happens when deciding which kernel to use.

```
# Our convolution function
def convolution(image, kernel):
   # Flip the kernel (optional)
   kernel = np.flipud(np.fliplr(kernel))
   # Get the dimensions of the image and kernel
   image rows, image cols = image.shape
   kernel rows, kernel cols = kernel.shape
   # Convolve using Numpy
   output = correlate(image, kernel, mode = 'valid')
   # Note that this is equivalent to this
   # Loop through the image, applying the convolution
   output = np.zeros like(image)
   for x in range(image rows - kernel rows + 1):
       for y in range(image cols - kernel cols + 1):
            output[x, y] = (kernel * image[x:x+kernel_rows, y:y+kernel_cols]).sum()
   return output
```

Definition (the blur kernel):

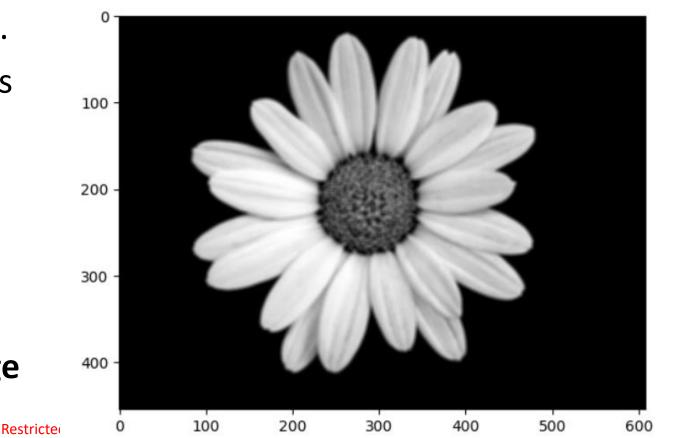
The **blur kernel** is a matrix with constant values, summing up to 1.

For instance, a 5×5 blur kernel is defined as:

Its effect is to transform the image into a more blurred version of it.

(455, 608)

```
# Display image in matplotlib
plt.imshow(image_conv, cmap = 'gray')
plt.show()
```



```
Restricted
 1 # Open the image and convert it to grayscale
                                                                                     # Blur kernel
 2 im = Image.open('flower.jpg').convert('L')
                                                                                      kernel = np.array([[1, 1, 1, 1, 1],
                                                                                                          [1, 1, 1, 1, 1],
 4 # Convert the image to a NumPy array
                                                                                                          [1, 1, 1, 1, 1],
 5 im array = np.array(im)
                                                                                                          [1, 1, 1, 1, 1],
                                                                                                          [1, 1, 1, 1, 1]])/25
   # Print the shape of the array
                                                                                     image_conv = convolution(im_array, kernel)
 8 print(im_array.shape)
                                                                                     print(image conv.shape)
(459, 612)
                                                                                 (455, 608)
 1 # Display image in matplotlib
                                                                                   1 # Display image in matplotlib
 2 plt.imshow(im_array, cmap = 'gray')
                                                                                     plt.imshow(image_conv, cmap = 'gray')
 3 plt.show()
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```

```
Restricted
 1 # Open the image and convert it to grayscale
                                                                                   # Blur kernel
 2 im = Image.open('flower.jpg').convert('L')
                                                                                    kernel = np.array([[1, 1, 1, 1, 1],
                                                                                                        [1, 1, 1, 1, 1],
 4 # Convert the image to a NumPy array
                                                                                                        [1, 1, 1, 1, 1],
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                                                                                                        [1, 1, 1, 1, 1],
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(459, 612)
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 2 plt.imshow(im_array, cmap = 'gray')
                                                                                    plt.imshow(image_conv, cmap = 'gray')
 3 plt.show()
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```

Definition (the edge detection kernel):

We can also perform **edge detection**, using, for instance the 3×3 **Prewitt horizontal kernel**, defined as:

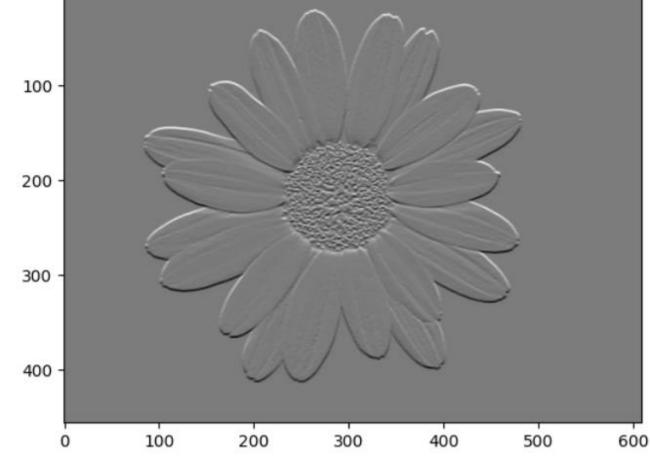
$$K = \begin{pmatrix} 1 & 1 & 1 \\ 0 & 0 & 0 \\ -1 & -1 & -1 \end{pmatrix}$$

This kernel highlights the horizontal edges in the image.

```
# Trying one kernel (Prewitt kernel for horizontal edges)
kernel1 = np.array([[1, 1, 1],[0, 0, 0],[-1, -1, -1]])
image_conv1 = convolution(im_array, kernel1)
print(image_conv1.shape)

(457, 610)

# Display image in matplotlib
plt.imshow(image_conv1, cmap = 'gray')
plt.show()
```



Definition (the edge detection kernel):

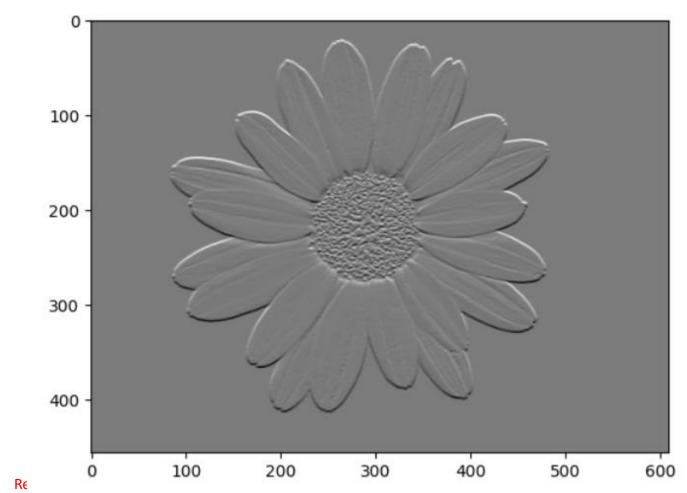
This kernel produces

- a value close to zero when all pixels are similar,
- and a non-zero value if there is a drastic change of values in the pixel and its neighbouring pixels.

Typically, it detects scenarios where the homophily property does not hold (i.e. an edge!).

```
# Trying one kernel (Prewitt kernel for horizontal edges)
   kernel1 = np.array([[1, 1, 1],[0, 0, 0],[-1, -1, -1]])
    image conv1 = convolution(im array, kernel1)
   print(image conv1.shape)
(457, 610)
    # Display image in matplotlib
    plt.imshow(image conv1, cmap = 'gray')
```

```
plt.show()
```



```
R€
 1 # Open the image and convert it to grayscale
                                                                         1 # Trying one kernel (Prewitt kernel for horizontal edges)
 2 im = Image.open('flower.jpg').convert('L')
                                                                         2 kernel1 = np.array([[1, 1, 1],[0, 0, 0],[-1, -1, -1]])
                                                                            image_conv1 = convolution(im_array, kernel1)
 4 # Convert the image to a NumPy array
                                                                         4 print(image_conv1.shape)
 5 im_array = np.array(im)
                                                                       (457, 610)
   # Print the shape of the array
 8 print(im_array.shape)
                                                                         1 # Display image in matplotlib
(459, 612)
                                                                            plt.imshow(image_conv1, cmap = 'gray')
                                                                           plt.show()
 1 # Display image in matplotlib
 2 plt.imshow(im_array, cmap = 'gray')
 3 plt.show()
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              100
                       200
                                  300
                                            400
                                                      500
                                                                600
```

Testing convolution on images

Definition (the sharpen, or contrast enhancer kernel):

We can also perform **contrast enhancement** (or **sharpening**), using, the following 3×3 **sharpen kernel**, defined as:

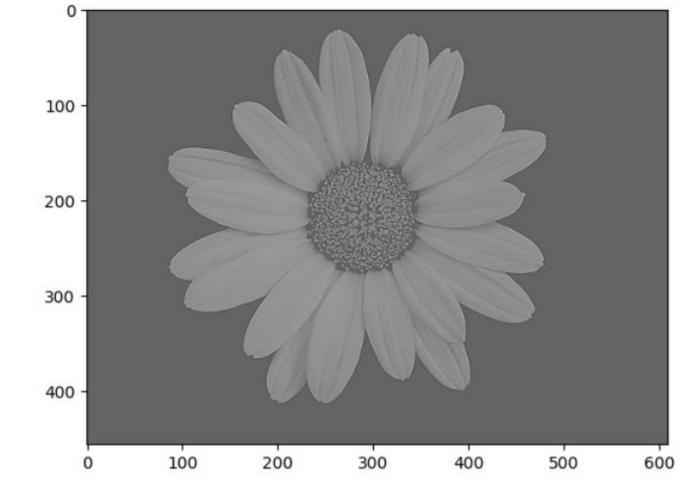
$$K = \begin{pmatrix} 0 & -1 & 0 \\ -1 & 5 & -1 \\ 0 & -1 & 0 \end{pmatrix}$$

This kernel typically enhances the edges and high-frequency details

```
# Sharpen (= improve contrast)
kernel5 = np.array([[0, -1, 0], [-1, 5, -1], [0, -1, 0]])
image_conv5 = convolution(im_array, kernel5)
print(image_conv5.shape)
```

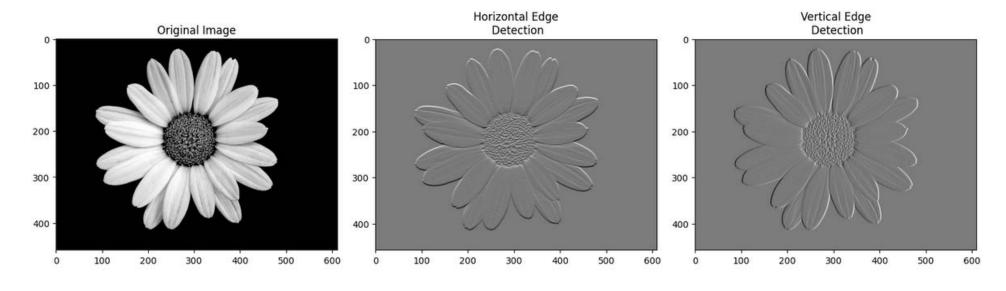
```
(457, 610)
```

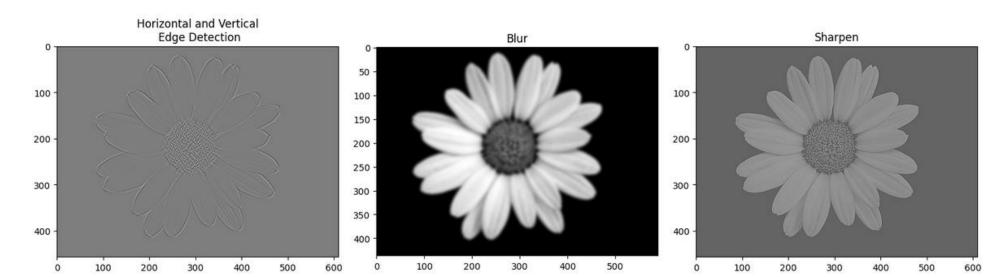
```
# Display image in matplotlib
plt.imshow(image_conv5, cmap = 'gray')
plt.show()
```



```
1 # Open the image and convert it to grayscale
                                                                         # Sharpen (= improve contrast)
 2 im = Image.open('flower.jpg').convert('L')
                                                                         kernel5 = np.array([[0, -1, 0], [-1, 5, -1], [0, -1, 0]])
                                                                          image_conv5 = convolution(im_array, kernel5)
 4 # Convert the image to a NumPy array
                                                                       4 print(image conv5.shape)
 5 im_array = np.array(im)
                                                                     (457, 610)
   # Print the shape of the array
 8 print(im_array.shape)
                                                                         # Display image in matplotlib
(459, 612)
                                                                          plt.imshow(image_conv5, cmap = 'gray')
                                                                          plt.show()
 1 # Display image in matplotlib
 2 plt.imshow(im_array, cmap = 'gray')
                                                                         0
 3 plt.show()
                                                                       100 -
 100 -
                                                                       200 -
 200 -
                                                                       300
 300 -
                                                                       400
 400 -
              100
                       200
                                  300
                                            400
                                                      500
                                                                                     100
                                                                                                200
                                                                                                            300
                                                                                                                       400
                                                                                                                                  500
                                                                                                                                             600
                                                                600
```

Many transformation kernels exist





More on convolution kernels and their effects



Downsizing

Remember: The convolution operation defined as Y = f(X, K), produces an image Y of size $h' \times w'$,

- With h' = h k + 1,
- And w' = w k + 1.

A typical problem with the convolution operation we defined earlier is that it produces a new image Y whose size/resolution has been reduced.

This has to do with the kernel being used, which sums several pixels together, but only produces one pixel as a result.

This effect is even increased with the size of the kernel being used.

Definition (padding in convolution):

A typical way to address this issue consists of **padding** the image.

Padding consists of adding some extra pixels on the outer part of the original image *X*.

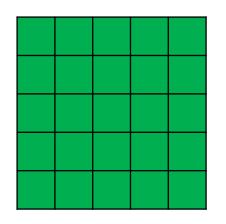
Padding artificially increases the size of the original image X, so that the convolution produces an image Y matching the size of X.

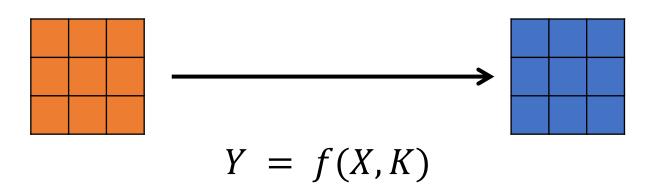
Three possible patterns for padding exist:

- Valid/Same padding,
- Zero padding,
- Other types of padding...

This is typically be used to maintain the spatial dimensions of the input image and prevent the spatial dimensions of the output from getting too small.

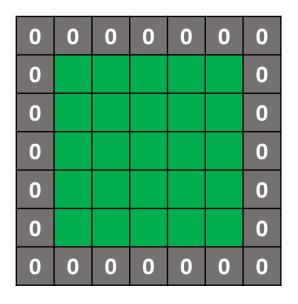
Valid padding: default configuration, no padding is applied to the input data. The convolution operation is only performed only on the valid parts of the input and the output size is then smaller than the input size.

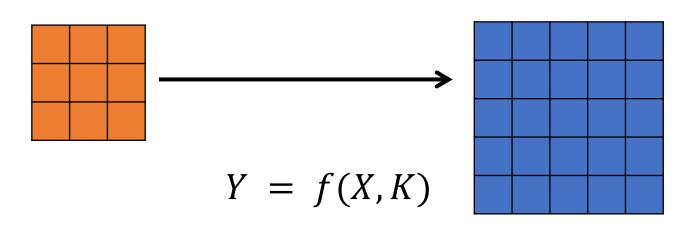




Same padding: The number of elements added to the image (i.e. padding size p) is defined so that image Y has the same size as X.

Zero padding: added pixels can simply be set to have the value 0.









When using a padding with size p, the convolution operation defined as Y = f(X, K), produces an image Y of size $h' \times w'$,

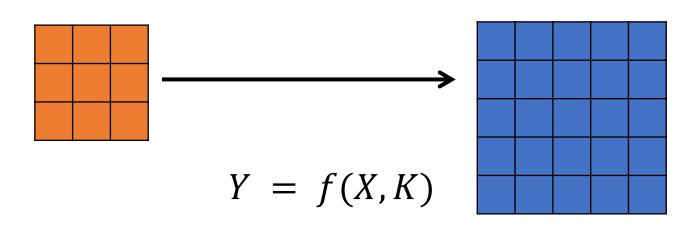
- With h' = h + 2p k + 1,
- And w' = w + 2p k + 1.

Same padding: to ensure the image Y has the same dimension as X, we need to use a padding size $p = \frac{k-1}{2}$. It is therefore adjusted to the kernel size k.

If we are not interested in Y having the same size as X, we are free to use any padding size.

Interpolation padding: instead of zeros, added pixels have the value of the closest pixel. Can also "zoom in" the original image and resize it to the expected number of pixels we want for image X after padding.

1	1	5	5	2	4	4
1	1	5	5	2	4	4
2	2	4	5	2	2	2
3	3	1	1	1	1	1
4	4	4	7	2	3	3
5	5	0	0	3	1	1
5	5	0	0	0	1	1





Padding is simply implemented using the **pad()** function.

It adds a layer of zeroes to the matrix image, of size *padding* on top, bottom, right and left side of the image matrix.

```
def convolution with padding(image, kernel, padding = 0):
   # Flip the kernel (optional)
    kernel = np.flipud(np.fliplr(kernel))
   # Get the dimensions of the image and kernel
    image rows, image cols = image.shape
    kernel rows, kernel cols = kernel.shape
   # Add padding to the image
    image = np.pad(image, ((padding, padding), (padding, padding)), 'constant')
    # Set the output image to the correct size
    output rows = image rows - kernel rows + 1
    output cols = image cols - kernel cols + 1
    output = np.zeros((output rows, output cols))
   # Convolve using Numpy
   output = correlate(image, kernel, mode = 'valid')
    return output
```

- When using no padding, the resulting image is downsized.
- When using same padding, i.e.

$$p = \frac{k-1}{2} = \frac{5-1}{2} = 2$$

the resulting image has the same size as the original one.

```
# Blur (no padding)
    kernel = np.array([[1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1])/25
    image conv = convolution basic(im array, kernel)
    # Print the shape of the array
    print(im array.shape)
    print(image conv.shape)
(459, 612)
(455, 608)
    # Blur (same padding)
    kernel = np.array([[1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1],
                        [1, 1, 1, 1, 1])/25
    image_conv_pad = convolution_with_padding(im_array, kernel, padding = 2)
    # Print the shape of the array
    print(im array.shape)
    print(image conv pad.shape)
(459, 612)
(459, 612)
```

Definition (stride in convolution):

The purpose of **stride** in convolution is to **control the movement or step size of the convolution filter as it slides over the input image**.

A larger stride results in a smaller output feature map, while a smaller stride results in a larger output feature map.

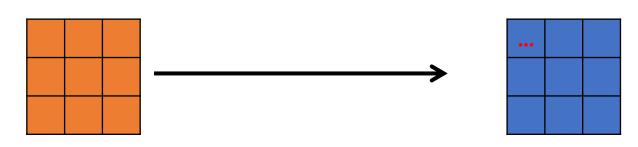
Stride can also be used to reduce the spatial dimensions of the feature map, which can help to reduce the number of parameters and computation in the network.

Note: Using a stride with a size *s* that is not 1, will drastically reduce the size of the output image, so beware!

Below, we demonstrate what a stride of size s = 2 does (no padding).

First, use the convolution as before, next elements to be used are "two steps" away from the first position, and so on.

1	1	5	5	2	4	4
1	1	5	5	2	4	4
2	2	4	5	2	2	2
3	3	1	1	1	1	1
4	4	4	7	2	3	3
5	5	0	0	3	1	1
5	5	0	0	0	1	1

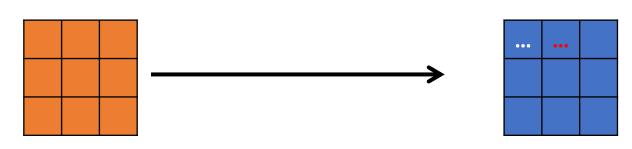


$$Y = f(X, K)$$

Below, we demonstrate what a stride of size s = 2 does (no padding).

First, use the convolution as before, next elements to be used are "two steps" away from the first position, and so on.

1	1	5	5	2	4	4
1	1	5	5	2	4	4
2	2	4	5	2	2	2
3	3	1	1	1	1	1
4	4	4	7	2	3	3
5	5	0	0	3	1	1
5	5	0	0	0	1	1



$$Y = f(X, K)$$

- Stride is easily implemented by using a slicing in the result of the correlation.
- This is not the most optimized way to do it, however!
- (We do not care so much, as we will use the PyTorch version of convolution after this!)

```
def convolution with stride and padding(image, kernel, stride = 1, padding = 0):
    # Flip the kernel (optional)
    kernel = np.flipud(np.fliplr(kernel))
    # Get the dimensions of the image and kernel
    image rows, image cols = image.shape
    kernel rows, kernel cols = kernel.shape
    # Add padding to the image
    image = np.pad(image, ((padding, padding), (padding, padding)), 'constant')
   # Set the output image to the correct size
    output rows = (image rows - kernel rows) // stride + 1
    output cols = (image cols - kernel cols) // stride + 1
    output = np.zeros((output rows, output cols))
    # Convolve using Numpy
   # (not the most optimal way to implement it but good enough!)
    output = correlate(image, kernel, mode = 'valid')[::stride, ::stride]
    return output
```

The (final) magic formula for convolution

Magic formula for convolution: let us consider

- An input image, defined as a 3D tensor X of size h × w × c, with h the height, w the weight and c the number of channels.
- A convolution kernel K of size $k \times k$.
- A padding of size p, a stride of size s.

The resulting image Y will have a size $h' \times w' \times c$, with:

$$h' = \left| \frac{h + 2p - k}{s} + 1 \right|$$

And

$$w' = \left| \frac{w + 2p - k}{s} + 1 \right|$$

Note: the floor function [...] is used in case the division result is an integer.

```
# Blur (with stride and padding)
    kernel = np.array([[1, 1, 1, 1, 1],
                           [1, 1, 1, 1, 1],
                           [1, 1, 1, 1, 1],
                           [1, 1, 1, 1, 1],
 6
                           [1, 1, 1, 1, 1]])/25
    image_conv_pad_stride = convolution_with_stride_and_padding(im_array, \
                                                                        kernel, \
 8
                                                                        stride = 2, \
 9
10
                                                                        padding = 2)
    # Print the shape of the array
12
    print(im_array.shape)
                                                             h' = \left| \frac{h + 2p - k}{s} + 1 \right|
    print(image conv pad stride.shape)
(459, 612)
                                                     h' = \left| \frac{459 + 2 \times 2 - 5}{2} + 1 \right| = 230
(230, 306)
```

Adding a dilation

Definition (dilation in convolution):

Dilation in image convolution determines the spacing between the values of the original image multiplying the kernel.

By default, the value for dilation is d = 1.

With a dilation greater than 1, the pixel values of the original image multiplying the kernel are **spread apart by a specified number of pixels**.

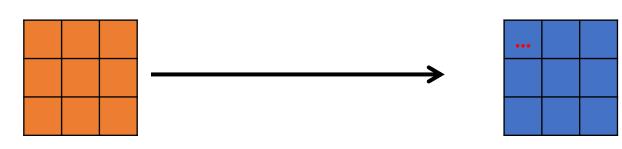
This is a very niche feature of the convolution operation. It is easier to visualise it than explain it.

(We leave its implementation as practice to the reader.)

Adding a dilation

Below, we demonstrate what a dilation of size d=2 does. Our convolution uses a stride with default value set to 1, padding is also set to its default value 0. Elements being computed together at then "2 steps away" from each other.

1	1	5	5	2	4	4
1	1	5	5	2	4	4
2	2	4	5	2	2	2
3	3	1	1	1	1	1
4	4	4	7	2	3	3
5	5	0	0	3	1	1
5	5	0	0	0	1	1



$$Y = f(X, K)$$



The (final) magic formula for convolution

Magic formula for convolution version 2.0: let us consider

- An input image, defined as a 3D tensor X of size h × w × c, with h the height, w the weight and c the number of channels.
- A convolution kernel K of size $k \times k$.
- A padding of size p, a stride of size s, and a dilation of size d.

The resulting image Y will have a size $h' \times w' \times c$, with:

$$h' = \left[\frac{h + 2p - d(k - 1) - 1}{s} + 1 \right]$$

And

$$w' = \left| \frac{w + 2p - d(k-1) - 1}{s} + 1 \right|$$

Note: the floor function [...] is used in case the division result is an integer.

Implementing a custom Conv in PyTorch (with stride and padding, no dilation)

```
def convolution_with_stride_and_padding_torch(image, kernel, stride = 1, padding = 0):
    # Convert image and kernel to PyTorch tensors
    image = torch.from numpy(image)
    kernel = torch.from numpy(kernel)
   # Flip the kernel (optional)
    kernel = torch.flip(torch.flip(kernel, [0]), [1])
    # Add padding to the image
    image = torch.nn.functional.pad(image, (padding, padding, padding, padding))
   # Set the output image to the correct size
    output rows = (image.shape[0] - kernel.shape[0]) // stride + 1
    output_cols = (image.shape[1] - kernel.shape[1]) // stride + 1
    output = torch.zeros((output rows, output cols))
   # Convolve using PyTorch
    for i in range(0, output_rows, stride):
       for j in range(0, output_cols, stride):
            output[i, j] = (kernel * image[i:i + kernel.shape[0], j:j + kernel.shape[1]]).sum()
    return output
```

Implementing a custom Conv in PyTorch (with stride and padding, no dilation)

The PyTorch and the Numpy implementations are indeed equivalent.

torch.Size([230, 306])

The Conv2d in PyTorch

PyTorch has a Conv2d function implementing said convolution.

We will rely on it from now on.

```
def convolution batch torch conv2d(images, kernel, stride = 1, padding = 0):
       # Convert kernel to PyTorch tensor, if needed
       kernel = torch.from numpy(kernel)
       kernel = kernel.view(1, 1, kernel.shape[0], kernel.shape[1])
       kernel = kernel.float()
       # Flip the kernel (optional)
       kernel = torch.flip(torch.flip(kernel, [2]), [3])
 9
10
       # Create a convolutional layer
11
       conv = torch.nn.Conv2d(in channels = images.shape[1], \
12
13
                               out_channels = 1, \
                               kernel size = kernel.shape[2:], \
14
                               stride = stride, \
15
                               padding = padding)
16
17
       # Assign the kernel to the layer
18
19
       conv.weight = torch.nn.Parameter(kernel)
       conv.bias = torch.nn.Parameter(torch.tensor([0.0]))
20
22
       # Perform convolution
       output = conv(images)
23
24
25
       return output
```

The Conv2d in PyTorch

Note, however that this layer has **two trainable parameters**:

- Weight (which is our kernel, whose values will be decided later during training).
- Bias (which was not there before?)

```
def convolution batch torch conv2d(images, kernel, stride = 1, padding = 0):
       # Convert kernel to PyTorch tensor, if needed
       kernel = torch.from numpy(kernel)
       kernel = kernel.view(1, 1, kernel.shape[0], kernel.shape[1])
       kernel = kernel.float()
       # Flip the kernel (optional)
       kernel = torch.flip(torch.flip(kernel, [2]), [3])
 9
10
11
       # Create a convolutional layer
12
       conv = torch.nn.Conv2d(in channels = images.shape[1], \
13
                               out channels = 1, \
                               kernel_size = kernel.shape[2:], \
14
                               stride = stride, \
15
16
                               padding = padding)
17
       # Assign the kernel to the layer
18
19
       conv.weight = torch.nn.Parameter(kernel)
        conv.bias = torch.nn.Parameter(torch.tensor([0.0]))
20
21
22
       # Perform convolution
       output = conv(images)
23
24
25
       return output
```

The Conv2d in PyTorch

Note, however that this layer has two trainable parameters:

- Weight (which is our kernel, whose values will be decided later during training).
- Bias (which was not there before?)

Bias is simply added to our convolution operation, following the intuition of the WX + b operation from earlier.

$$Y_{i,j} = \sum_{m=1}^{k} \sum_{m=1}^{k} X_{i+m-1, j+n-1} K_{m,n} + b$$