



# **ARTIFICIAL AND COMPUTATIONAL INTELLIGENCE AND KNOWLEDGE ENGINEERING**

## **Cutting-Edge Python Computational Tools for AI, CI, KE and Data Mining**



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# Scope



- ✓ **Python** – a modern language often used for AI, CI, KE, and DM computing
- ✓ **Jupyter Notebook** – a modern and intuitive programming environment with linked libraries (like Tensorflow, Keras) that allow to effectively process deep learning algorithms and show results quickly.
- ✓ **Tensorflow and Keras** libraries produced by leading IT companies, like Google.

## The Jupyter Notebook:

- is an open-source web application that allows you to create and share documents that contain live code, equations, visualizations and narrative text;
- includes data cleaning and transformation, numerical simulation, statistical modeling, data visualization, machine learning, and much more.



**We will use it to demonstrate various algorithms, so you are asked to install it.**

Jupyter in your browser

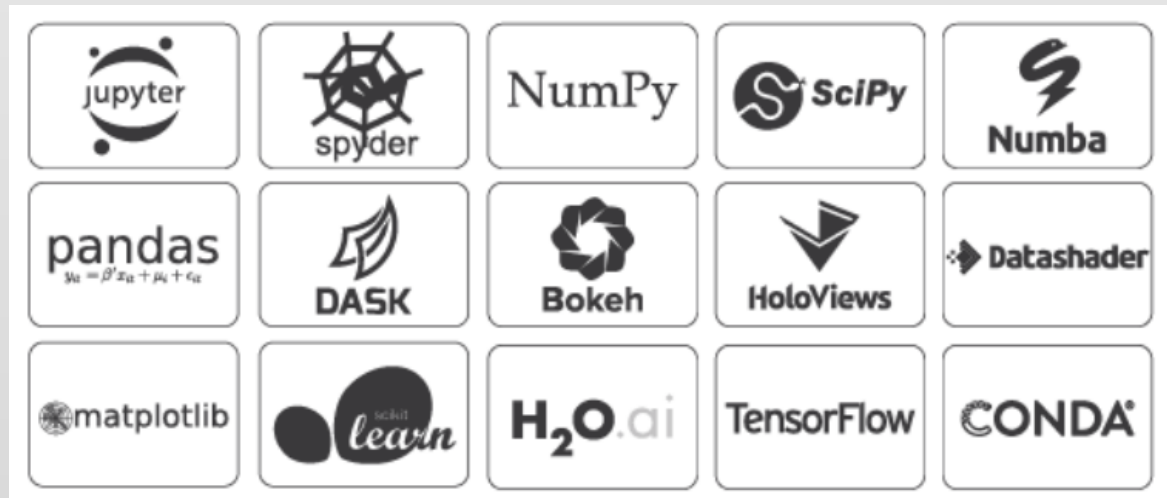
Install a Jupyter Notebook

## Install Jupyter using [Anaconda](#) with built in Python 3.7+

- It includes many other commonly used packages for scientific computing, data science, machine learning ,and computational intelligence libraries.
- It manages libraries, dependencies, and environments with Conda.
- It allows developing and training various machine learning and deep learning models with scikit-learn, TensorFlow, and Theano etc.
- It supplies us with data analysis including scalability and performance with Dask, NumPy, pandas, and Numba.
- It quickly visualizes results with Matplotlib, Bokeh, Datashader, and Holoviews.

And [run it](#) at the Terminal (Mac/Linux) or Command Prompt (Windows):

```
jupyter notebook
```





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# Jupyter Notebook Dashboard



## Running a Jupyter Notebook in your browser:

- When the Jupyter Notebook opens in your browser, you will see the Notebook Dashboard, which will show you a list of the notebooks, files, and subdirectories in the directory where the notebook server was started by the command line „jupyter notebook”.
- Most of the time, you will wish to start a notebook server in the highest level directory containing notebooks. Often this will be your home directory.

localhost:8888/tree

jupyter

Quit Logout

Files Running Clusters

Select items to perform actions on them. Upload New ↻

<input type="checkbox"/> 0	Name ↓	Last Modified	File size
<input type="checkbox"/>	3D Objects	4 miesiące temu	
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<input type="checkbox"/>	Contacts	4 miesiące temu	
<input type="checkbox"/>	Desktop	miesiąc temu	
<input type="checkbox"/>	Documents	4 miesiące temu	
<input type="checkbox"/>	Downloads	10 minut temu	
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<input type="checkbox"/>	Favorites	4 miesiące temu	
<input type="checkbox"/>	Links	4 miesiące temu	
<input type="checkbox"/>	miniconda3	12 minut temu	
<input type="checkbox"/>	Music	3 miesiące temu	
<input type="checkbox"/>	OneDrive	16 dni temu	
<input type="checkbox"/>	OpenVPN	2 lata temu	
<input type="checkbox"/>	Pictures	2 miesiące temu	
<input type="checkbox"/>	PycharmProjects	10 minut temu	
<input type="checkbox"/>	Saved Games	4 miesiące temu	
<input type="checkbox"/>	Searches	4 miesiące temu	
<input type="checkbox"/>	source	9 miesięcy temu	
<input type="checkbox"/>	Tracing	rok temu	
<input type="checkbox"/>	Videos	2 miesiące temu	

It is recommended to install [PyCharm](#) for Anaconda:



ANACONDA

Anaconda3 2019.03 (64-bit)

Anaconda + JetBrains

Anaconda and JetBrains are working together to bring you Anaconda-powered environments tightly integrated in the PyCharm IDE.

PyCharm for Anaconda is available at:

<https://www.anaconda.com/pycharm>



ANACONDA®





# Jupyter Notebook



## PyCharm is a python IDE for Professional Developers

- It includes scientific mode to interactively analyze your data.

The screenshot displays the PyCharm IDE interface with a Jupyter Notebook open. The notebook file is named 'sine.ipynb'. The code in the notebook is as follows:

```
#%%  
import numpy as np  
import matplotlib.pyplot as plt  
  
x = np.linspace(0, 2*np.pi)  
  
#%%  
plt.plot(x, np.sin(x))  
plt.plot(x, np.sin(x-1/4*np.pi))
```

The output of the notebook shows the execution of the code, resulting in a plot of two sine waves. The first wave is blue and the second is orange. The x-axis ranges from 0 to 6, and the y-axis ranges from -1.00 to 1.00. The plot is displayed in the 'SciView' tab on the right side of the IDE.

The IDE interface includes a top toolbar with various icons for file operations, a left sidebar with '1: Project', '2: Structure', and '3: Favorites' views, and a bottom status bar with tabs for '4: Run', '5: Debug', '6: TODO', '7: Docker', '8: Version Control', '9: Jupyter', 'Terminal', 'Python Console', and 'Event Log'.

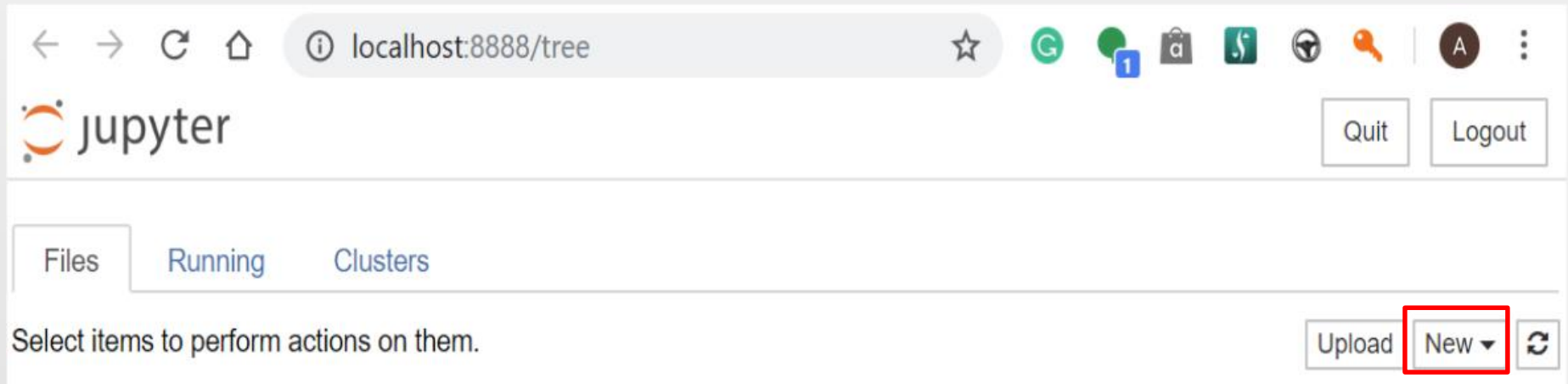


# Starting a new Python notebook

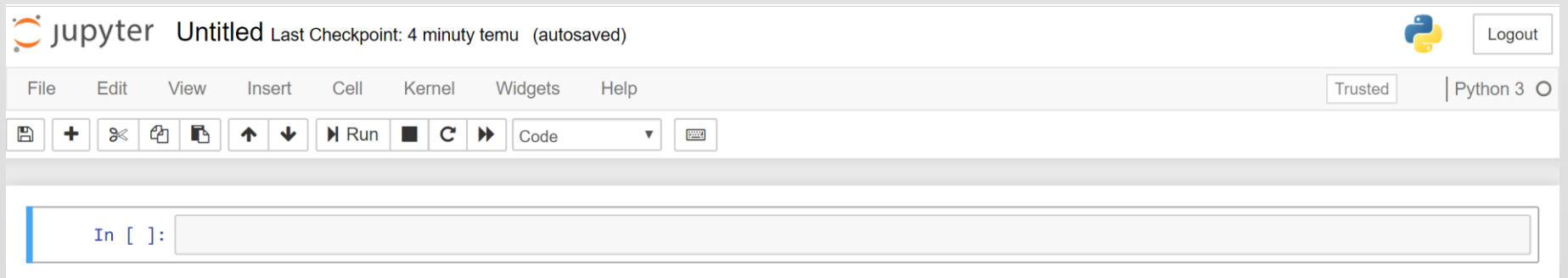


## Start a new Python notebook:


- Clicking New → Python 3



- And a new Python project in the Jupyter Notebook will be started:



**When dealing with big data collections and big data vectors, we should use vectorization to proceed computations faster:**

Jupyter Comparison of for-looped and vectorized efficiency of computations Last Checkpoint: 3 godziny temu (unsaved changes)  Logout

File Edit View Insert Cell Kernel Widgets Help Trusted Python 3

Run Code

In [4]:

```
import numpy as np

a = np.array([1,2,3,4])
print (a)

import time

a = np.random.rand(1000000)
b = np.random.rand(1000000)

tic = time.time()
c = np.dot(a,b)
toc = time.time()

print ("Vectorized dot product computation time:" + str(1000 * (toc-tic)) + "ms")

tic = time.time()
for i in range(1000000):
    d += a[i]*b[i]
toc = time.time()

print ("For-Looped dot product computation time:" + str(1000 * (toc-tic)) + "ms")
```

## Conclusion:

Whenever possible, avoid explicit for-loops and use vectorization:  
`np.dot(A,v)`, `np.dot(x,w)`, `np.log(v)`, `np.abs(v)`,  
`np.zeros(v)`, `np.sum(v)` etc.

```
[1 2 3 4]
Vectorized dot product computation time:0.9996891021728516ms
For-Looped dot product computation time:420.8984375ms
```

In [ ]:

We use numpy vectorization to compute `sigmoid(x)` and `sigmoid_derivative` for any input vector `x`:

$$\text{For } x \in \mathbb{R}^n, \text{sigmoid}(x) = \text{sigmoid} \begin{pmatrix} x_1 \\ x_2 \\ \vdots \\ x_n \end{pmatrix} = \begin{pmatrix} \frac{1}{1+e^{-x_1}} \\ \frac{1}{1+e^{-x_2}} \\ \vdots \\ \frac{1}{1+e^{-x_n}} \end{pmatrix} \quad (1)$$

$$\text{sigmoid\_derivative}(x) = \sigma'(x) = \sigma(x)(1 - \sigma(x)) \quad (2)$$

```
In [14]: import numpy as np # You can access numpy functions by writing np.function() instead of numpy.function()

def sigmoid(x):
    s = 1 / (1 + np.exp(-x)) # It computes the sigmoid of x, where x can be a scalar or numpy array of any size
    return s

def sigmoid_derivative(x):
    s = sigmoid(x)
    ds = s * (1 - s) # It computes the gradient (slope or derivative) of the sigmoid function with respect to its input x.
    return ds
```

```
In [15]: x = np.array([-2,-1,0,1, 2])
print ("sigmoid(x) = " + str(sigmoid(x)))
print ("sigmoid_derivative(x) = " + str(sigmoid_derivative(x)))

sigmoid(x) = [0.11920292 0.26894142 0.5          0.73105858 0.88079708]
sigmoid_derivative(x) = [0.10499359 0.19661193 0.25          0.19661193 0.10499359]
```

**We use normalization to achieve a better performance because gradient descent converges faster after normalization:**

Normalization is changing  $x$  to  $\frac{x}{\|x\|}$  (dividing each row vector of  $x$  by its norm), e.g.

If

$$x = \begin{bmatrix} 3 & 2 & 4 \\ 1 & 8 & 2 \end{bmatrix} \quad (3)$$

then

$$\|x\| = \text{np.linalg.norm}(x, \text{axis} = 1, \text{keepdims} = \text{True}) = \begin{bmatrix} \sqrt{29} \\ \sqrt{69} \end{bmatrix} \quad (4)$$

and

$$x_{\text{normalized}} = \frac{x}{\|x\|} = \begin{bmatrix} \frac{3}{\sqrt{29}} & \frac{2}{\sqrt{29}} & \frac{4}{\sqrt{29}} \\ \frac{1}{\sqrt{69}} & \frac{8}{\sqrt{69}} & \frac{2}{\sqrt{69}} \end{bmatrix} \quad (5)$$

```
In [25]: def normalizeRows(x):
# This function normalizes each row of the matrix x, where x is a numpy matrix of shape (n, m)
x_norm = np.linalg.norm(x,ord=2,axis=1,keepdims=True)
print("x_norm = " + str(x_norm))
x = x/x_norm
return x
```

```
In [26]: x = np.array([
[3, 2, 4],
[1, 8, 2]])
print("normalizeRows(x) = " + str(normalizeRows(x)))

x_norm = [[5.38516481]
[8.30662386]]
normalizeRows(x) = [[0.55708601 0.37139068 0.74278135]
[0.12038585 0.96308682 0.24077171]]
```

**Broadcasting** is very useful for performing mathematical operations between arrays of different shapes.

A softmax function is a normalizing function often used in the output layers of neural networks when you need to classify two or more classes:

- for  $x \in \mathbb{R}^{1 \times n}$ ,  $\text{softmax}(x) = \text{softmax}([x_1 \quad x_2 \quad \dots \quad x_n]) = \left[ \frac{e^{x_1}}{\sum_j e^{x_j}} \quad \frac{e^{x_2}}{\sum_j e^{x_j}} \quad \dots \quad \frac{e^{x_n}}{\sum_j e^{x_j}} \right]$
- for a matrix  $x \in \mathbb{R}^{m \times n}$ ,  $x_{ij}$  maps to the element in the  $i^{\text{th}}$  row and  $j^{\text{th}}$  column of  $x$ , thus we have:

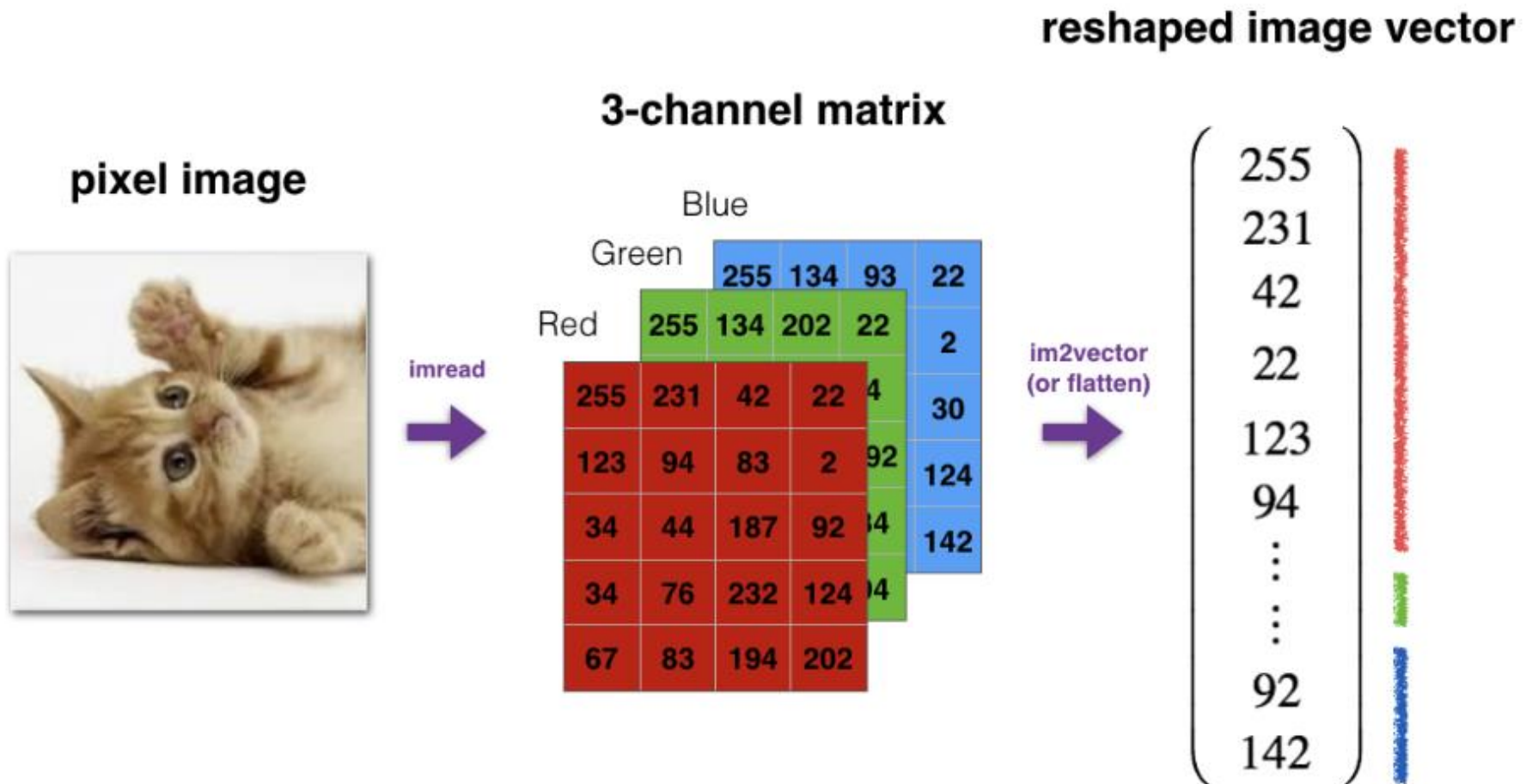
$$\text{softmax}(x) = \text{softmax} \begin{bmatrix} x_{11} & x_{12} & x_{13} & \dots & x_{1n} \\ x_{21} & x_{22} & x_{23} & \dots & x_{2n} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ x_{m1} & x_{m2} & x_{m3} & \dots & x_{mn} \end{bmatrix} = \begin{bmatrix} \frac{e^{x_{11}}}{\sum_j e^{x_{1j}}} & \frac{e^{x_{12}}}{\sum_j e^{x_{1j}}} & \frac{e^{x_{13}}}{\sum_j e^{x_{1j}}} & \dots & \frac{e^{x_{1n}}}{\sum_j e^{x_{1j}}} \\ \frac{e^{x_{21}}}{\sum_j e^{x_{2j}}} & \frac{e^{x_{22}}}{\sum_j e^{x_{2j}}} & \frac{e^{x_{23}}}{\sum_j e^{x_{2j}}} & \dots & \frac{e^{x_{2n}}}{\sum_j e^{x_{2j}}} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ \frac{e^{x_{m1}}}{\sum_j e^{x_{mj}}} & \frac{e^{x_{m2}}}{\sum_j e^{x_{mj}}} & \frac{e^{x_{m3}}}{\sum_j e^{x_{mj}}} & \dots & \frac{e^{x_{mn}}}{\sum_j e^{x_{mj}}} \end{bmatrix} = \begin{pmatrix} \text{softmax}(\text{first row of } x) \\ \text{softmax}(\text{second row of } x) \\ \dots \\ \text{softmax}(\text{last row of } x) \end{pmatrix}$$

```
In [27]: def softmax(x):
# This function calculates the softmax for each row of the input x, where x is a row vector or a matrix of shape (n, m).
x_exp = np.exp(x)
x_sum = np.sum(x_exp,axis=1,keepdims=True)
s = x_exp/x_sum # It automatically uses numpy broadcasting.
return s
```

```
In [29]: x = np.array([
[0, 9, 3, 0],
[3, 0, 8, 1]])
print("softmax(x) = " + str(softmax(x)))

softmax(x) = [[1.23074356e-04 9.97281837e-01 2.47201452e-03 1.23074356e-04]
[6.68456877e-03 3.32805082e-04 9.92077968e-01 9.04658008e-04]]
```

When working with images in deep learning, we typically reshape them into vector representation using [`np.reshape\(\)`](#) :



We commonly use the numpy functions [`np.shape\(\)`](#) and [`np.reshape\(\)`](#) in deep learning:

- `X.shape` is used to get the shape (dimension) of a vector or a matrix `X`.
- `X.reshape(...)` is used to reshape a vector or a matrix `X` into some other dimension(s).

Images are usually represented by 3D arrays of shape (*length, height, depth* = 3). Nevertheless, when you read an image as the input of an algorithm you typically convert it to a vector of shape (*length \* height \* 3, 1*), so you "unroll" (reshape) the 3D arrays into 1D vectors for further processing:

**Example 1:** If you would like to reshape an array `v` of shape (`a, b, c`) into a vector of shape (`a*b,c`) you would do:

```
v = v.reshape((v.shape[0] * v.shape[1], v.shape[2])) # where v.shape[0] = a ; v.shape[1] = b ; v.shape[2] = c
```

**Example 2:** If you would like to reshape an array `v` of shape (`a, b, c`) into a vector of shape (`abc`) you would do:

```
v = v.reshape((v.shape[0] * v.shape[1] * v.shape[2], 1)) # where v.shape[0] = a ; v.shape[1] = b ; v.shape[2] = c
```

- Never hard-code the dimensions of the image as a constant but use the quantities you need with `image.shape[0]`, etc.

```
In [30]: def image2vector(image):
# This function reshapes a numpy array of shape (length, height, depth) to a vector of shape (length*height*depth, 1)
v = image.reshape((image.shape[0]*image.shape[1]*image.shape[2]),1)
return v
```

```
In [33]: # Images usually are (num_px_x, num_px_y, 3) where 3 represents the RGB values: red, green, and blue
# This is an exemplary 3 by 3 by 3 array:
image = np.array([[[ 0.139,  0.381],
[ 0.982,  0.647],
[ 0.251,  0.551]],
[[ 0.219,  0.647],
[ 0.703,  0.845],
[ 0.397,  0.313]],
[[ 0.855,  0.165],
[ 0.313,  0.937],
[ 0.279,  0.077]]])

image2vector(image) = [[0.139]
[0.381]
[0.982]
[0.647]
[0.251]
[0.551]
[0.219]
[0.647]
[0.703]
[0.845]
[0.397]
[0.313]
[0.855]
[0.165]
[0.313]
[0.937]
[0.279]
[0.077]]

print ("image = " + str(image))
print ("image2vector(image) = " + str(image2vector(image)))
```

In deep learning, you deal with very large datasets. Non-computationally-optimal functions become a huge bottleneck in your algorithms and can result in models that take ages to run. To make sure that your code is computationally efficient, you should use vectorization. Compare the following codes:

```
In [4]: import time
import numpy as np

x1 = [2, 6, 3, 9, 1, 0, 2, 0, 6, 1, 9, 3, 5, 2, 1]
x2 = [6, 3, 1, 1, 7, 8, 2, 5, 2, 4, 0, 2, 3, 0, 9]

### CLASSIC DOT PRODUCT OF VECTORS IMPLEMENTATION ###
tic = time.process_time()
dot = 0
for i in range(len(x1)):
    dot += x1[i]*x2[i]
toc = time.process_time()
print ("dot = " + str(dot))
print ("Computation time of the classic dot product of vector implementation = " + str(1000 * (toc - tic)) + "ms")

### VECTORIZED DOT PRODUCT OF VECTORS IMPLEMENTATION ###
tic = time.process_time()
dot = np.dot(x1,x2)
toc = time.process_time()
print ("dot = " + str(dot))
print ("Computation time of the vectorized dot product of vectors implementation = " + str(1000 * (toc - tic)) + "ms")

dot = 99
Computation time of the classic dot product of vector implementation = 0.0ms
dot = 99
Computation time of the vectorized dot product of vectors implementation = 0.0ms
```

In deep learning, you deal with very large datasets. Non-computationally-optimal functions become a huge bottleneck in your algorithms and can result in models that take ages to run. To make sure that your code is computationally efficient, you should use vectorization. Compare the following codes:

```
In [5]: import time
import numpy as np

x1 = [2, 6, 3, 9, 1, 0, 2, 0, 6, 1, 9, 3, 5, 2, 1]
x2 = [6, 3, 1, 1, 7, 8, 2, 5, 2, 4, 0, 2, 3, 0, 9]

### CLASSIC OUTER PRODUCT IMPLEMENTATION ###
tic = time.process_time()
outer = np.zeros((len(x1),len(x2))) # we create a len(x1)*len(x2) matrix with only zeros
for i in range(len(x1)):
    for j in range(len(x2)):
        outer[i,j] = x1[i]*x2[j]
toc = time.process_time()
print ("outer = " + str(outer))
print ("Computation time of the classic outer product of vector implementation = " + str(1000 * (toc - tic)) + "ms")

### VECTORIZED OUTER PRODUCT IMPLEMENTATION ###
tic = time.process_time()
outer = np.outer(x1,x2)
toc = time.process_time()
print ("outer = " + str(outer))
print ("Computation time of the vectorized outer product of vectors implementation = " + str(1000 * (toc - tic)) + "ms")
```

```
outer = [[12.  6.  2.  2. 14. 16.  4. 10.  4.  8.  0.  4.  6.  0. 18.]
 [36. 18.  6.  6. 42. 48. 12. 30. 12. 24.  0. 12. 18.  0. 54.]
 [18.  9.  3.  3. 21. 24.  6. 15.  6. 12.  0.  6.  9.  0. 27.]
 [54. 27.  9.  9. 63. 72. 18. 45. 18. 36.  0. 18. 27.  0. 81.]
 [ 6.  3.  1.  1.  7.  8.  2.  5.  2.  4.  0.  2.  3.  0.  9.]
 [ 0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.]
 [12.  6.  2.  2. 14. 16.  4. 10.  4.  8.  0.  4.  6.  0. 18.]
 [ 0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.  0.]
 [36. 18.  6.  6. 42. 48. 12. 30. 12. 24.  0. 12. 18.  0. 54.]
 [ 6.  3.  1.  1.  7.  8.  2.  5.  2.  4.  0.  2.  3.  0.  9.]
 [54. 27.  9.  9. 63. 72. 18. 45. 18. 36.  0. 18. 27.  0. 81.]
 [18.  9.  3.  3. 21. 24.  6. 15.  6. 12.  0.  6.  9.  0. 27.]
 [30. 15.  5.  5. 35. 40. 10. 25. 10. 20.  0. 10. 15.  0. 45.]
 [12.  6.  2.  2. 14. 16.  4. 10.  4.  8.  0.  4.  6.  0. 18.]
 [ 6.  3.  1.  1.  7.  8.  2.  5.  2.  4.  0.  2.  3.  0.  9.]]
Computation time of the classic outer product of vector implementation = 0.0ms
```

```
outer = [[12  6  2  2 14 16  4 10  4  8  0  4  6  0 18]
 [36 18  6  6 42 48 12 30 12 24  0 12 18  0 54]
 [18  9  3  3 21 24  6 15  6 12  0  6  9  0 27]
 [54 27  9  9 63 72 18 45 18 36  0 18 27  0 81]
 [ 6  3  1  1  7  8  2  5  2  4  0  2  3  0  9]
 [ 0  0  0  0  0  0  0  0  0  0  0  0  0  0  0]
 [12  6  2  2 14 16  4 10  4  8  0  4  6  0 18]
 [ 0  0  0  0  0  0  0  0  0  0  0  0  0  0  0]
 [36 18  6  6 42 48 12 30 12 24  0 12 18  0 54]
 [ 6  3  1  1  7  8  2  5  2  4  0  2  3  0  9]
 [54 27  9  9 63 72 18 45 18 36  0 18 27  0 81]
 [18  9  3  3 21 24  6 15  6 12  0  6  9  0 27]
 [30 15  5  5 35 40 10 25 10 20  0 10 15  0 45]
 [12  6  2  2 14 16  4 10  4  8  0  4  6  0 18]
 [ 6  3  1  1  7  8  2  5  2  4  0  2  3  0  9]]
Computation time of the vectorized outer product of vectors implementation = 0.0ms
```

In deep learning, you deal with very large datasets. Non-computationally-optimal functions become a huge bottleneck in your algorithms and can result in models that take ages to run. To make sure that your code is computationally efficient, you should use vectorization. Compare the following codes:

```
In [8]: import time
import numpy as np

x1 = [2, 6, 3, 9, 1, 0, 2, 0, 6, 1, 9, 3, 5, 2, 1]
x2 = [6, 3, 1, 1, 7, 8, 2, 5, 2, 4, 0, 2, 3, 0, 9]

### CLASSIC ELEMENTWISE MULTIPLICATION IMPLEMENTATION ###
tic = time.process_time()
mul = np.zeros(len(x1))
for i in range(len(x1)):
    mul[i] = x1[i]*x2[i]
toc = time.process_time()
print ("elementwise multiplication = " + str(mul))
print ("Computation time of the classic element-wise implementation = " + str(1000 * (toc - tic)) + "ms")

### VECTORIZED ELEMENTWISE MULTIPLICATION IMPLEMENTATION ###
tic = time.process_time()
mul = np.multiply(x1,x2)
toc = time.process_time()
print ("elementwise multiplication = " + str(mul))
print ("Computation time of the vectorized element-wise implementation = " + str(1000 * (toc - tic)) + "ms")

elementwise multiplication = [12. 18.  3.  9.  7.  0.  4.  0. 12.  4.  0.  6. 15.  0.  9.]
Computation time of the classic element-wise implementation = 0.0ms
elementwise multiplication = [12 18  3  9  7  0  4  0 12  4  0  6 15  0  9]
Computation time of the vectorized element-wise implementation = 0.0ms
```

In deep learning, you deal with very large datasets. Non-computationally-optimal functions become a huge bottleneck in your algorithms and can result in models that take ages to run. To make sure that your code is computationally efficient, you should use vectorization. Compare the following codes:

```
In [7]: import time
import numpy as np

x1 = [2, 6, 3, 9, 1, 0, 2, 0, 6, 1, 9, 3, 5, 2, 1]
x2 = [6, 3, 1, 1, 7, 8, 2, 5, 2, 4, 0, 2, 3, 0, 9]

### CLASSIC GENERAL DOT PRODUCT IMPLEMENTATION ###
W = np.random.rand(3,len(x1)) # Random 3*len(x1) numpy array
tic = time.process_time()
gdot = np.zeros(W.shape[0])
for i in range(W.shape[0]):
    for j in range(len(x1)):
        gdot[i] += W[i,j]*x1[j]
toc = time.process_time()
print ("gdot = " + str(gdot));
print ("Computation time of the classic general dot product of vector implementation = " + str(1000 * (toc - tic)) + "ms")

### VECTORIZED GENERAL DOT PRODUCT IMPLEMENTATION ###
tic = time.process_time()
dot = np.dot(W,x1)
toc = time.process_time()
print ("gdot = " + str(dot))
print ("Computation time of the vectorized general dot product of vectors implementation = " + str(1000 * (toc - tic)) + "ms")

gdot = [29.7697031  21.47839099 29.65064192]
Computation time of the classic general dot product of vector implementation = 0.0ms
gdot = [29.7697031  21.47839099 29.65064192]
Computation time of the vectorized general dot product of vectors implementation = 0.0ms
```

The loss functions are used to evaluate the performance of your models. The bigger your loss is, the more different your predictions ( $\hat{y}$ ) are from the true values ( $y$ ). In deep learning, we use optimization algorithms like Gradient Descent to train models and minimize the cost.

L1 loss function using absolute distance of  $y$  and  $\hat{y}$  is defined as:

$$L_1(\hat{y}, y) = \sum_{i=0}^m |y^{(i)} - \hat{y}^{(i)}| \quad (6)$$

L2 loss function using squared distance of  $y$  and  $\hat{y}$  is defined as:

$$L_2(\hat{y}, y) = \sum_{i=0}^m (y^{(i)} - \hat{y}^{(i)})^2 \quad (7)$$

```
In [40]: def L1(yhat, y):
# This function returns the the L1 loss function where
# yhat - vector of size m (predicted labels)
# y - vector of size m (true labels)
loss = np.sum(np.abs(y-yhat))
return loss

def L2(yhat, y):
# This function returns the the L2 loss function where
# yhat - vector of size m (predicted labels)
# y - vector of size m (true labels)
loss = np.sum(np.dot(y-yhat,y-yhat))
return loss
```

```
In [41]: yhat = np.array([.34, 0.98, 0.21, .15, .12])
y = np.array([0, 1, 0, 0, 0])
print("L1 = " + str(L1(yhat,y)))
print("L2 = " + str(L2(yhat,y)))
```

```
L1 = 0.8400000000000001
L2 = 0.19700000000000004
```



# Let's start with powerful computations!



- ✓ Questions?
- ✓ Remarks?
- ✓ Suggestions?
- ✓ Wishes?



# Bibliography and Literature

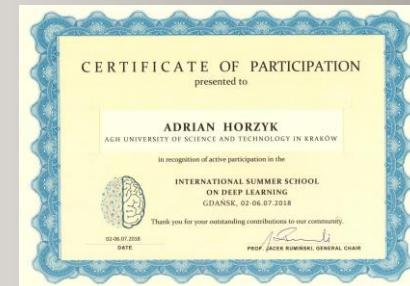
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