

Lecture 3: An Introduction to Scientific Computing with Python

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von KARMAN INSTITUTE
FOR FLUID DYNAMICS

Contents

- Numpy
- Matplotlib
- Scipy

Exercises

- Read and plot data
- Random numbers and histograms

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Numpy: the matrix manipulator

- Provides high-performance vector, matrix and higher-dimensional data structures for Python
- Implemented in C and Fortran for vectorized calculations (formulated with vectors and matrices) to have good performances
- Numpy is imported using:

In [1]:

```
import numpy as np # by convention alias np
```

Check here the list of almost all useful functions:

<https://numpy.org/devdocs/user/numpy-for-matlab-users.html>

Multiple ways of creating numpy arrays

- From lists:

In [2]:

```
m = np.array([[2,7,9], [4,3,3],[8,1,0]]); print(m)  
type(m)
```

```
[[2 7 9]  
 [4 3 3]  
 [8 1 0]]
```

Out[2]:

```
numpy.ndarray
```


Multiple ways of creating numpy arrays

- From lists:

In [2]:

```
m = np.array([[2,7,9], [4,3,3],[8,1,0]]); print(m)  
type(m)
```

```
[[2 7 9]  
 [4 3 3]  
 [8 1 0]]
```

Out[2]:

```
numpy.ndarray
```

- Using functions:

In [3]:

```
print(np.zeros((3, 2)))  
print(np.ones((2, 3)))  
print(np.linspace(0.0, 10.0, 3))  
print(np.logspace(1.0, 3.0, 3))
```

```
[[0.  0.]  
 [0.  0.]  
 [0.  0.]]  
[[1.  1.  1.]  
 [1.  1.  1.]]
```

```
[ 0.  5. 10.]  
[ 10. 100. 1000.]
```

In [4]:

```
print(np.random.rand(2, 4))
```

```
[[0.66654839 0.15653709 0.89851319 0.31345401]  
 [0.09526442 0.74339405 0.56887685 0.64758033]]
```


Matrix properties

In [5]:

```
print(m)
```

```
[[2 7 9]
 [4 3 3]
 [8 1 0]]
```

In [6]:

```
print(m.shape) # shape of matrix
print(np.shape(m))
print(m.size) # number of elements
print(np.size(m))
```

```
(3, 3)
(3, 3)
9
9
```

Indexing and slicing arrays as for lists

In [7]:

```
print(m)
```

```
[[2 7 9]
 [4 3 3]
 [8 1 0]]
```

In [8]:

```
print(m[0, 0])
```

```
2
```

In [9]:

```
print(m[:, -1]) # get the last column
```

```
[9 3 0]
```

In [10]:

```
print(m[0:2, 1]) # get two first elements of first column
```

```
[7 3]
```

In [11]:

```
print(m[[0, 2], 0]) # from column 0, get elements 0 and 2
```

```
[2 8]
```

Linear algebra

- Element-wise operations:

In [12]:

```
m = np.array([[0, 1, 2], [3, 4, 5]]) # our initial matrix  
m*2
```

Out[12]:

```
array([[ 0,  2,  4],  
       [ 6,  8, 10]])
```

In [13]:

```
m+2
```

Out[13]:

```
array([[2, 3, 4],  
       [5, 6, 7]])
```

In [14]:

```
m**2
```

Out[14]:

```
array([[ 0,  1,  4],  
       [ 9, 16, 25]])
```


Linear algebra

- Element-wise operations:

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```
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m*2
```

Out[12]:

```
array([[ 0,  2,  4],  
       [ 6,  8, 10]])
```

In [13]:

```
m+2
```

Out[13]:

```
array([[2, 3, 4],  
       [5, 6, 7]])
```

In [14]:

```
m**2
```

Out[14]:

```
array([[ 0,  1,  4],  
       [ 9, 16, 25]])
```

- Matrix algebra:

In [15]:

```
v = (np.array([1, 2, 3]))
print(m)
print(m.shape)
print(v)
print(v.shape)
```

```
[[0 1 2]
 [3 4 5]]
(2, 3)
[1 2 3]
(3,)
```

In [16]:

```
print(m.dot(v)) # matrix multiplication
print(np.dot(m, v))
print(m @ v)
```

```
[ 8 26]
[ 8 26]
[ 8 26]
```

In [17]:

```
print(m.transpose()) # transpose
print(m.T) # transpose, shorter
print(m.flatten()) # convert to 1D vector
```

```
[[0 3]
 [1 4]
 [2 5]]
```



```
[[0 3]
 [1 4]
 [2 5]]
[0 1 2 3 4 5]
```

In [18]:

```
# reshape
m_reshape = m.reshape(6,1)
print(m_reshape.shape)
```

```
(6, 1)
```

About dimensions in numpy...

This is used sometimes in sklearn

In [19]:

```
print(m_reshape.shape)  
print(v.shape)
```

```
(6, 1)  
(3, )
```

In [20]:

```
v = np.expand_dims(v, axis=1)  
print(v.shape)
```

```
(3, 1)
```

Numpy does not change the original matrix
inplace

Assign to a new variable if needed

Numpy does not change the original matrix inplace

Assign to a new variable if needed

In [21]:

```
print(m)
print(m.T)
print('---Square matrix did not change---')
print(m)
```

```
[[0 1 2]
 [3 4 5]]
[[0 3]
 [1 4]
 [2 5]]
---Square matrix did not change---
[[0 1 2]
 [3 4 5]]
```

Numpy matrix algebra is faster than loops!

To check this, let's calculate the modulus of a vector in three ways:

1. Loops
2. Matrix multiplication
3. Built-in function

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To check this, let's calculate the modulus of a vector in three ways:

1. Loops
2. Matrix multiplication
3. Built-in function

In [23]:

```
import time # to compute the time needed
# Vector of n points between [-10, 10]
n = 10000000
v = np.random.normal(loc=0, scale=20, size=n); # use of ; avoids print of output in jupyter
```

With loops: $|\mathbf{v}| = \sqrt{\sum v_i^2}$

In [24]:

```
start = time.time() # get start time

sum_vectors_squared = 0
for vi in v:
    sum_vectors_squared += vi**2
module = np.sqrt(sum_vectors_squared)

end = time.time() # get end time

print("Module is:", module)
print("Execution time :", end-start, " s")
```

```
Module is: 63217.60994740732
Execution time : 5.882690906524658 s
```

With matrices: $|\mathbf{v}| = \sqrt{v^T \cdot v}$

In [25]:

```
start = time.time()

product = v.transpose().dot(v)
module = np.sqrt(product)

end = time.time()

print("Module is:", module)
print("Execution time :", end-start, " s")
```

```
Module is: 63217.60994740611
Execution time : 0.0074841976165771484 s
```


With built-in function norm: $|\mathbf{v}| =$
`np.linalg.norm(v)`

In [26]:

```
start = time.time()
module = np.linalg.norm(v)
end = time.time()

print("Module is:", module)
print("Execution time :", end-start, " s")
```

```
Module is: 63217.60994740611
Execution time : 0.0038220882415771484 s
```

With built-in function norm: $|\mathbf{v}| =$
`np.linalg.norm(v)`

In [26]:

```
start = time.time()
module = np.linalg.norm(v)
end = time.time()

print("Module is:", module)
print("Execution time : ", end-start, " s")
```

```
Module is: 63217.60994740611
Execution time : 0.0038220882415771484 s
```

Conclusion: ALWAYS search for built-in functions

They are faster, cleaner and easy to use

You will save a lot of time

Extra trick: A better way to time your functions...use a decorator!

In [27]:

```
from time import time

def timer(func):
    # This function shows the execution time of
    # the function object passed
    def wrapper(*args, **kwargs):
        t1 = time()
        result = func(*args, **kwargs)
        t2 = time()
        print(f'Function {func.__name__!r} executed in {(t2-t1):.4f}s')
        return result
    return wrapper
```

In [28]:

```
import numpy

@timer
def compute_module(v):
    module = np.linalg.norm(v)
    return module

v = [1, 3, 4, 5, 6]
compute_module(v)

v = [1, 3, 4, 5, 6]*2000
compute_module(v)
```

```
Function 'compute_module' executed in 0.0001s
Function 'compute_module' executed in 0.0029s
```

Out[28]:

```
417.1330722922842
```


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Matplotlib: publication quality plots

- Provides capabilities for plotting similar to Matlab
- Interactive and animated figures
- Possibility for LaTeX rendering for improved quality
- Most other plotting libraries rely on Matplotlib or use a similar syntax

Matplotlib: publication quality plots

- Provides capabilities for plotting similar to Matlab
- Interactive and animated figures
- Possibility for LaTeX rendering for improved quality
- Most other plotting libraries rely on Matplotlib or use a similar syntax

The pyplot sub-module is the most useful for us

In [29]:

```
# # for interactive plotting in jupyter  
# %matplotlib widget  
# for static plotting in jupyter  
%matplotlib inline  
  
import matplotlib.pyplot as plt # import as an alias for easier typing.
```

Let's make a simple parabola plot

Let's make a simple parabola plot

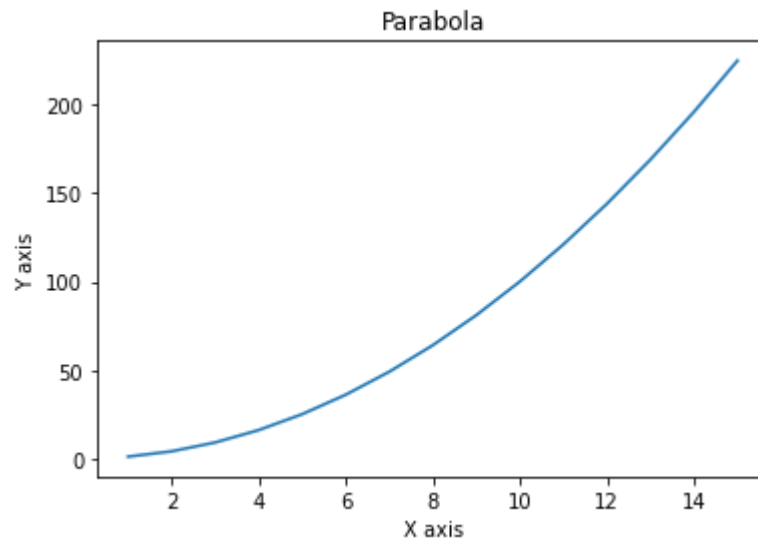
In [30]:

```
x = [1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15]
y = [1, 4, 9, 16, 25, 36, 49, 64, 81, 100, 121, 144, 169, 196, 225]

fig, ax = plt.subplots()
ax.set_xlabel('X axis')
ax.set_ylabel('Y axis')
ax.set_title('Parabola')
ax.plot(x, y)
```

Out[30]:

```
[<matplotlib.lines.Line2D at 0x7f8fd4a27a00>]
```



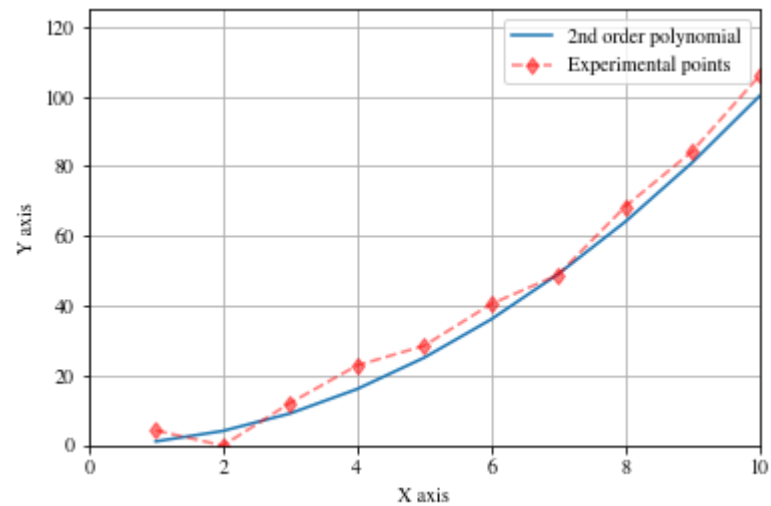
Customize your plots

In [31]:

```
# Use LaTeX fonts:
plt.style.use({'font.family': 'STIXGeneral',
              'font.serif': 'Computer Modern',
              'font.sans-serif': 'Computer Modern Sans serif',})

y_noisy = np.random.normal(y, 5) # create some noisy data to simulate the experimental points

# create a new plot
fig2, ax2 = plt.subplots()
ax2.set_xlabel('X axis')
ax2.set_ylabel('Y axis')
ax2.plot(x, y, label='2nd order polynomial')
ax2.plot(x, y_noisy, 'rd--', label='Experimental points', alpha=0.5)
ax2.grid()
ax2.set_xlim([0, 10])
ax2.set_ylim([0, 125])
ax2.legend()
fig2.savefig('figure.png')
```



Creating an animation

Generating the parabola step by step

In [32]:

```
import os

temp_folder = 'gif_temp'
if not os.path.exists(temp_folder):
    os.mkdir(temp_folder)
```

In [33]:

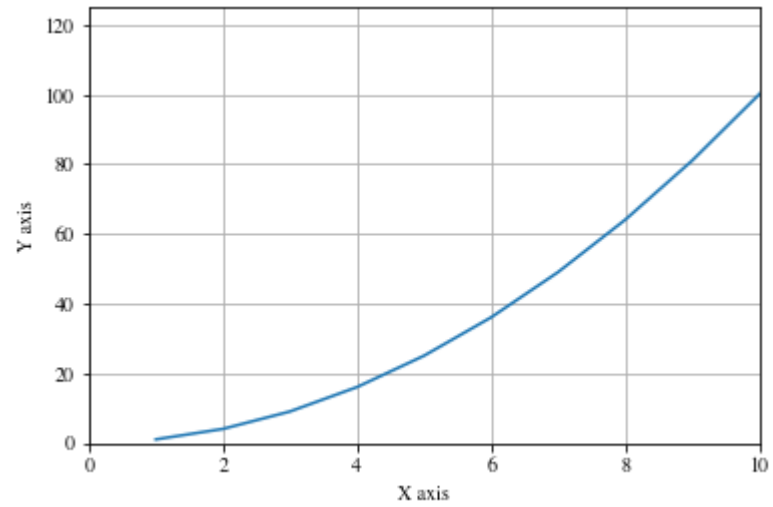
```
figAnim, axanim = plt.subplots()

# set plot properties that do not change.
axanim.set_xlabel('X axis')
axanim.set_ylabel('Y axis')
axanim.set_xlim([0, 10])
axanim.set_ylim([0, 125])
axanim.plot(x,y, label='2nd order polynomial')
axanim.grid()

for i,_ in enumerate(x):
    current_points = axanim.plot(x[:i],y_noisy[:i],'rd--', label='Experimental points', alpha=0.5)
    figAnim.savefig(f'{temp_folder}/Im_{str(i).zfill(2)}.png') # the use of zfill allows to recover them ordered
    print(f'Image {i} saved')
    # remove the current points for a clean plot
    points = current_points.pop(0)
    points.remove()
```

```
Image 0 saved
Image 1 saved
Image 2 saved
Image 3 saved
Image 4 saved
Image 5 saved
Image 6 saved
Image 7 saved
```

Image 8 saved
Image 9 saved
Image 10 saved
Image 11 saved
Image 12 saved
Image 13 saved
Image 14 saved



Building the GIF

In [34]:

```
import imageio # This is used to read the dumped images and create the animation
import shutil # package to manage files

GIF_name = 'animated_parabola.gif'
cleanup = True # delete the temporary folder

images = []
for filename in os.listdir(temp_folder):
    images.append(imageio.imread(f'{temp_folder}/{filename}'))

imageio.mimsave(GIF_name, images, duration=0.2)
```

In [35]:

```
if cleanup:
    shutil.rmtree(temp_folder)
```

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Scipy

- Typical function and constants (`scipy.pi`, `scipy.sin`, etc)
- Integrators (ODE45 --> `scipy.integrate.RK45`, `scipy.integrate.odeint`)
- Curve fitting (`scipy.optimize.curve_fit`)
- Interpolation (`scipy.interpolate`)
- Statistics (`scipy.stats`)

Example: Solving an ODE with Scipy

$$y'(x) = y \cdot \ln(y)/x$$

$$y(2) = e$$

In [37]:

```
from scipy.integrate import odeint
import numpy as np

X0 = [2.0, np.e] # initial value

t = np.linspace(2, 10, 10) # solution points
# Function to integrate
df = lambda y, X: y*np.log(y)/X
# Scipy has a very powerful ODE integrator: - BUILD-IN FUNCTION
y_P = odeint(df, np.e, t)
```


Example: Solving an ODE with Scipy

$$y'(x) = y \cdot \ln(y)/x$$

$$y(2) = e$$

In [37]:

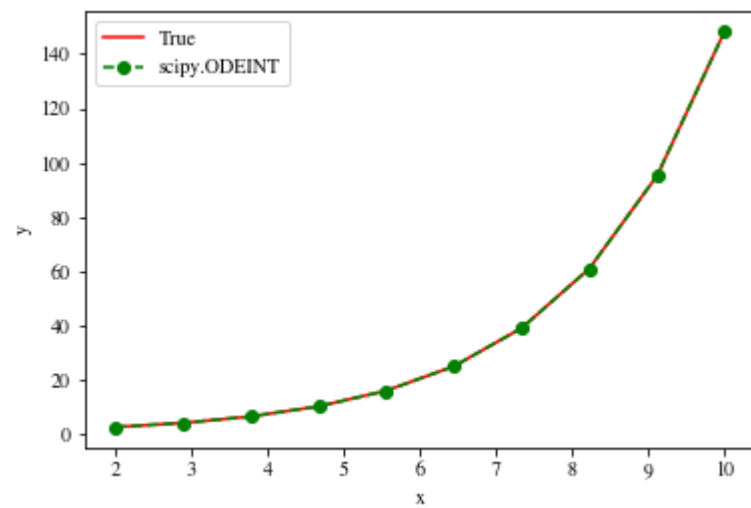
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# Function to integrate
df = lambda y, X: y*np.log(y)/X
# Scipy has a very powerful ODE integrator: - BUILD-IN FUNCTION
y_P = odeint(df, np.e, t)
```

In [38]:

```
import matplotlib.pyplot as plt
plt.figure
plt.plot(t, np.e**(t/2), color='r', label='True')
plt.plot(t, y_P, color='g', linestyle='--', label='scipy.ODEINT', marker='o' )
plt.legend()
plt.xlabel('x')
plt.ylabel('y')
plt.show()
```



Read data from a file

In [36]:

```
import numpy as np

file_to_read = 'alumina_data_raw.txt'
data = np.genfromtxt(file_to_read, skip_header=1, delimiter=';')
```

Exercise: plot data

- From the previous dataset read column 0 (temperature) and column 3 (mass)
- Add appropriate labels

Solution

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Solution

In [39]:

```
import matplotlib.pyplot as plt

temperature = data[:,0]
mass = data[:,3]

fig, ax = plt.subplots()

ax.plot(temperature, mass)
ax.set_xlabel('Temperature, °C')
ax.set_ylabel('Mass, %')
```

Out[39]:

```
Text(0, 0.5, 'Mass, %')
```

Exercise: gaussian histogram (part 1)

- Generate random data normally distributed
- Plot it in a histogram
- Include the theoretical normal distribution
- Add a legend
- Save the figure

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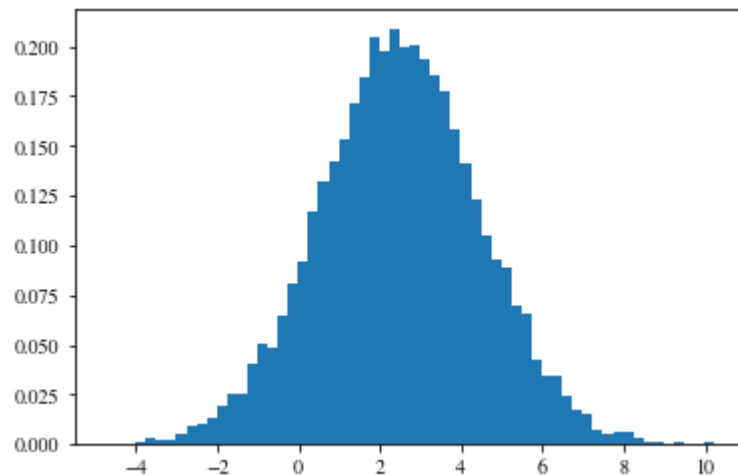
In [40]:

```
# %matplotlib widget
import numpy as np
import matplotlib.pyplot as plt

mu = 2.5
sigma = 2

random_data = np.random.normal(mu, sigma, 10000);

fig_hist, ax_hist = plt.subplots()
ax_hist.hist(random_data, density=True, bins=60, label='Data');
```



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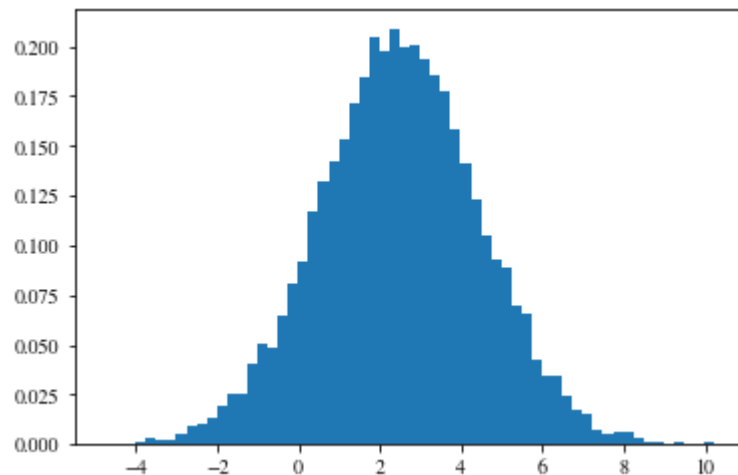
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```



In [41]:

```
xmin = mu - 3*sigma
xmax = mu + 3*sigma

x = np.linspace(xmin, xmax, num=300)
y = 1/(sigma * (2*np.pi)**0.5)*np.e**(-0.5*((x-mu)/sigma)**2)
ax_hist.plot(x, y, label='Normal distr.')
```

Out[41]:

```
[<matplotlib.lines.Line2D at 0x7f8fd321b3d0>]
```

Exercise: gaussian histogram (part 1)

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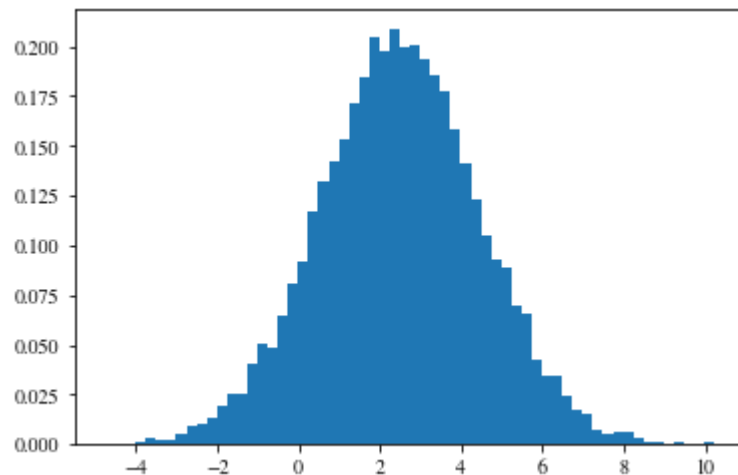
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ax_hist.plot(x, y, label='Normal distr.')
```

Out[41]:

```
[<matplotlib.lines.Line2D at 0x7f8fd321b3d0>]
```

In [42]:

```
ax_hist.legend()
fig_hist.savefig('Normal distribution.png')
```


Exercise: gaussian histogram (part 2)

- Generate random data normally distributed with different number of sample points
- Compare the mean and the std of your dataset to the provided values
- Plot it in a histogram
- Plot the normal distribution with your parameters
- Include the theoretical normal distribution
- Plot the convergence curves for mean and std
- Add a legend(s)

Exercise: gaussian histogram (part 2)

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- Add a legend(s)

In [43]:

```
# %matplotlib widget

import numpy as np
import matplotlib.pyplot as plt
import os

def compute_normal(x, mu, std):
    return 1/(sigma * (2*np.pi)**0.5)*np.e**(-0.5*((x-mu)/sigma)**2)

# Set temp folder to save figures
temp_folder = 'temp_histogram'
if not os.path.exists(temp_folder):
    os.mkdir(temp_folder)

## theoretical part
mu = 4.5
sigma = 7

xmin = mu - 3*sigma
xmax = mu + 3*sigma

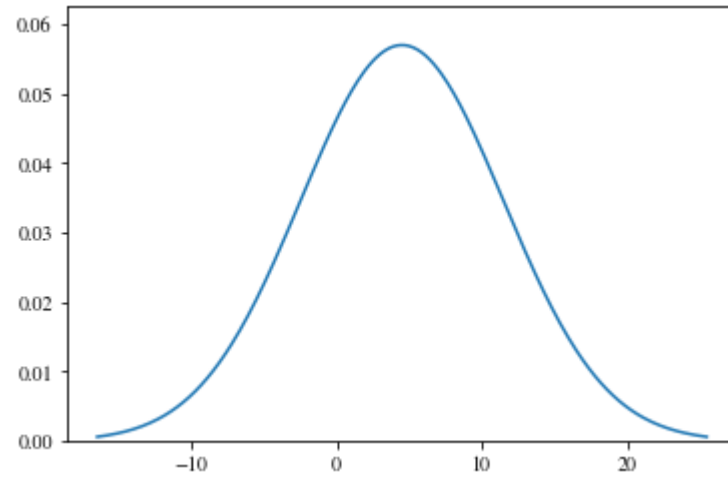
x = np.linspace(xmin, xmax, num=300)
y_theoretical = compute_normal(x, mu, sigma)

fig_hist, ax_hist = plt.subplots()
ax_hist.plot(x, y_theoretical, label='Normal distr. Theoretical', color='C0')

# set the limits
ax_hist.set_ylim(0, max(y_theoretical)*1.1)
```

Out[43]:

(0.0, 0.06268777430836729)



In [44]:

```
##

N_samples = np.geomspace(5, 20000, 100, dtype=int)

means = []
stds = []
for n in N_samples:
    random_data = np.random.normal(mu, sigma, n)

    mean = np.mean(random_data)
    std = np.std(random_data)

    means.append(mean)
    stds.append(std)

count, bins, bars = ax_hist.hist(random_data, color='C1', density=True, bins=60, label='Data', alpha=0.7)

y_computed = compute_normal(x, mean, std)
current_curve = ax_hist.plot(x, y_computed, color='C2', label='Normal distr. Computed')

# put the title
ax_hist.set_title(f'Samples: {n}, mean={mean:.2f}, std={std:.2f}')
ax_hist.legend(loc='upper left')

# save the figure
fig_hist.savefig(f'{temp_folder}/Im_{str(n).zfill(2)}.png')

# remove the current points for a clean plot
t = [b.remove() for b in bars]

line = current_curve.pop(0)
line.remove()
```

Exercise: gaussian histogram (part 2)

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- Compare the mean and the std of your dataset to the provided values
- Plot it in a histogram
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In [43]:

```
# %matplotlib widget

import numpy as np
import matplotlib.pyplot as plt
import os

def compute_normal(x, mu, std):
    return 1/(sigma * (2*np.pi)**0.5)*np.e**(-0.5*((x-mu)/sigma)**2)

# Set temp folder to save figures
temp_folder = 'temp_histogram'
if not os.path.exists(temp_folder):
    os.mkdir(temp_folder)

## theoretical part
mu = 4.5
sigma = 7

xmin = mu - 3*sigma
xmax = mu + 3*sigma

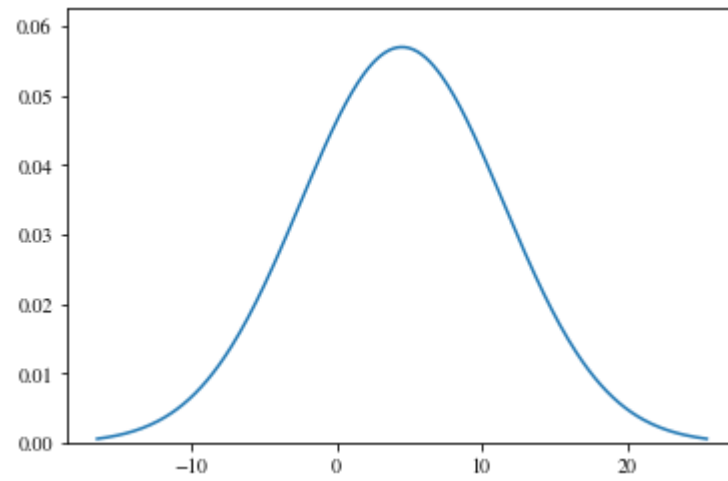
x = np.linspace(xmin, xmax, num=300)
y_theoretical = compute_normal(x, mu, sigma)

fig_hist, ax_hist = plt.subplots()
ax_hist.plot(x, y_theoretical, label='Normal distr. Theoretical', color='C0')

# set the limits
ax_hist.set_ylim(0,max(y_theoretical)*1.1)
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Out[43]:

(0.0, 0.06268777430836729)



In [44]:

```
##

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means = []
stds = []
for n in N_samples:
    random_data = np.random.normal(mu, sigma, n)

    mean = np.mean(random_data)
    std = np.std(random_data)

    means.append(mean)
    stds.append(std)

count, bins, bars = ax_hist.hist(random_data, color='C1', density=True, bins=60, label='Data', alpha=0.7)

y_computed = compute_normal(x, mean, std)
current_curve = ax_hist.plot(x, y_computed, color='C2', label='Normal distr. Computed')

# put the title
ax_hist.set_title(f'Samples: {n}, mean={mean:.2f}, std={std:.2f}')
ax_hist.legend(loc='upper left')

# save the figure
fig_hist.savefig(f'{temp_folder}/Im_{str(n).zfill(2)}.png')

# remove the current points for a clean plot
t = [b.remove() for b in bars]

line = current_curve.pop(0)
line.remove()
```

In [45]:

```
# We can animate the previous plot

import imageio # This is used to read the dumped images and create the animation
import shutil # package to manage files

GIF_name = 'histogram.gif'
```

Convergence of mean and std

In [46]:

```
fig, ax = plt.subplots()

ax.plot(N_samples, means, label='mean')
ax.plot(N_samples, stds, label='std')

ax.set_xlabel('Num. Samples')

ax.legend()
ax.set_xscale('log')

fig.savefig('convergence.pdf')
```

