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Plotting NumPy series

As an example, let us import a time series data. This represent human electroencephalogram (EEG) as recorded during normal background activity.

```
1 from pandas import read_csv
2
3 from matplotlib.pyplot import subplots, show
4
5 from numpy import arange, linspace, zeros
```

```
1 df = read_csv("data/EEG_background.txt", delim_whitespace=True)
2
3 df.head()
```

1	FP1	FP2	F3	F4		E02	EM1	EM2
		PHO						
2	0 -7.4546	22.8428	6.28159	15.6212		13.7021	12.9109	13.7034
	9.37573	3						
3	1 -11.1060	21.4828	6.89088	15.0562		13.7942	13.0194	13.7628
	9.44731	L						
4	2 -14.4000	20.0907	7.94856	14.1624		13.8982	13.1116	13.8239
	9.51796	5						
5	3 -17.2380	18.7206	9.36857	13.0093		14.0155	13.1927	13.8914
	9.58770)						
6	4 -19.5540	17.4084	11.06040	11.6674		14.1399	13.2692	13.9652
	9.65654	1						
7								
8 [5 rows x 28 columns]								

To see the names of the channels (or recording sensors) we can use head function.

```
1 df.shape
```

```
1 (2373, 28)
```

Numpy Plot

The data in the above dataframe df is converted to Numpy arrays, here called df_np.

Time in the rows, sensors in the columns

```
1 sr = 256  # Sampling rate: 1 / seconds
2
3 duration = 5  # seconds
4
5 df_np = df.to_numpy()
6
7 data = df_np[:duration*sr, :19] ## SF Comment, needs to explain this above
8
9 data.shape
```

```
1 (1280, 19)
```

Python Function

Please execute the following function definition before proceeding. The function code takes data and creates a plot of all columns as time series, one above the other. When you execute the function code nothing happens. Similar to the import, running a function code will only activate it and make it available for later use.

```
1 def plot_series(data, sr):
2
3
       Time series plot of multiple time series
4
       Data are normalised to mean=0 and var=1
5
       data: nxm numpy array. Rows are time points, columns are channels
6
7
       sr: sampling rate, same time units as period
8
9
       from numpy import flip
10
11
       samples = data.shape[0]
       sensors = data.shape[1]
12
13
14
       period = samples // sr
15
       time = linspace(0, period, period*sr)
16
17
       offset = 5 # for mean=0 and var=1 normalised data
18
19
       # Calculate means and standard deviations of all columns
21
       means = data.mean(axis=0)
22
       stds = data.std(axis=0)
```

```
23
24
       # Plot each series with an offset of 2 times the standard
           deviations
25
       fig, ax = subplots(figsize=(7, 8))
26
       ax.plot(time, (data - means)/stds + offset*arange(sensors-1,-1,-1))
27
           ;
28
29
       ax.plot(time, zeros((samples, sensors)) + offset*arange(sensors
           -1,-1,-1),'--',color='gray');
30
31
       ax.set(xlabel='Time')
       ax.set_yticks(offset*arange(sensors))
32
       ax.set_yticklabels(flip(arange(sensors)+1))
```

```
plot_series(data, sr);
show()
```

How to create a function

```
1 def my_plot1(data):
2
3    fig, ax = subplots()
4
5    ax.plot(data)
```

```
1 my_plot1(data)
2 show()
```

```
def my_plot2(data, factor):
3
       this is just a test
4
5
       columns = data.shape[1]
6
7
       offset = arange(columns)
8
9
       fig, ax = subplots()
10
11
       ax.plot(data + offset*factor)
12
```

```
1 my_plot2(data, 100)
2 show()
```

FourierSpectrum

The Fourier spectrum decomposes the time series into a sum of sine waves. The spectrum gives the coefficients of each of the sine wave components. The coefficients are directly related to the amplitudes needed to optimally fit the sum of all sine waves to recreate the original data.

However, the assumption behind the Fourier transform is that the data are provided as in infinitely long stationary time series. These assumptions are not fulfilled as the data are finite and stationarity of a biological system can typically not be guaranteed. Thus, interpretation needs to be cautious.

We import the Fourier transform function fft from scipy. fftpack and can use it to transform all columns at the same time.

```
from pandas import read_csv
from matplotlib.pyplot import subplots, yticks, legend, rcParams, show
from numpy import arange, linspace, zeros
from scipy.fftpack import fft
```

```
1  df = read_csv("data/EEG_absence.txt", delim_whitespace=True)
2  3  sr = 256
4  duration = 5
5  df_np = df.to_numpy()
7  8  data = df_np[:duration*sr, :2] ## SF, needs explanation
9  df.head()
```

```
FP1
                   FP2
1
                             F3
                                     F4
                                                  E02
                                                           EM1
                                                                    EM<sub>2</sub>
                 PHO
2 0 -6.9732 30.00060 60.9815 -23.047
                                              20.8242
                                                      20.3583 21.1760
                                         . . .
     14.5002
3 1 -15.1590 22.85930 62.2845 -24.359
                                             20.8289
                                                      20.3292 21.1118
                                         . . .
     14.5056
4 2 -23.3680 15.85860 63.2742 -25.353
                                             20.8337 20.3120 21.0367
                                         . . .
     14.5109
5 3 -31.5560
               9.05790 63.9646 -26.034
                                             20.8327 20.3002 20.9580
     14.5161
               2.45328 64.4026 -26.451 ... 20.8248 20.2862 20.8843
  4 -39.6840
6
     14.5212
7
8 [5 rows x 28 columns]
```

```
1 data.shape
```

```
1 (1280, 2)
```

```
1
   def plot_series(data, sr):
2
       Time series plot of multiple time series
3
       Data are normalised to mean=0 and var=1
4
5
6
       data: nxm numpy array. Rows are time points, columns are channels
       sr: sampling rate, same time units as period
8
9
       leg: Legend of figure, uses column index
10
11
12
       samples = data.shape[0]
13
       sensors = data.shape[1]
14
15
       period = samples // sr
16
       time = linspace(0, period, period*sr)
17
18
       offset = 5 # for mean=0 and var=1 normalised data
19
20
21
       # Calculate means and standard deviations of all columns
22
       means = data.mean(axis=0)
23
       stds = data.std(axis=0)
24
       # Plot each series with an offset of 2 times the standard
25
           deviations
26
       fig, ax = subplots(figsize=(7, 5))
27
       ax.plot(time, (data - means)/stds + offset*arange(sensors-1,-1,-1))
28
           ;
29
       ax.plot(time, zeros((samples, sensors)) + offset*arange(sensors
           -1,-1,-1),'--',color='gray');
31
       yticks([]);
32
34
       ax.set(xlabel='Time')
```

```
plot_series(data[:, :2], sr)
show()
```

```
1 data_fft = fft(data, axis=0)
2
3 data_fft.shape
```

```
1 (1280, 2)
```

```
1 rows = data.shape[0]
2 freqs = (sr/2)*linspace(0, 1, rows//2)
```

Time Series (EEG)

```
amplitude = (2.0 / rows) * abs(data_fft[:rows//2, :])

fig, ax = subplots()

ax.plot(freqs, amplitude);

show()
```

```
fig, ax = subplots()
ax.plot(freqs, amplitude);
ax.set_xlim(0, 10);
ax.set_xlabel('Frequency (Hz)', fontsize=20)
ax.set_ylabel('Amplitude (abs)', fontsize=20);
show()
```

Keypoints

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