

Analyzing the time-frequency content of EEG data

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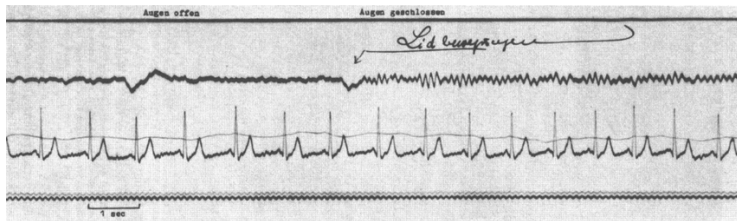
25/05/2022

Today

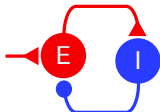
- ① Why study neural oscillations?
 - Evoked vs. induced activity
- ② Frequency analysis
 - Sine waves, Fourier transform
- ③ Time-frequency analysis
 - Morlet wavelets
- ④ Example workflow
 - MNE-Python style, hu-neuro-pipeline style

Why study neural oscillations?

Empirical data



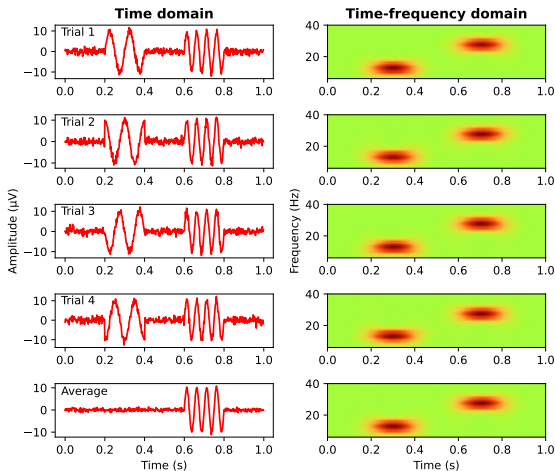
Algorithm



Computation

- Working memory
- Language
- Consciousness

Evoked vs. induced activity

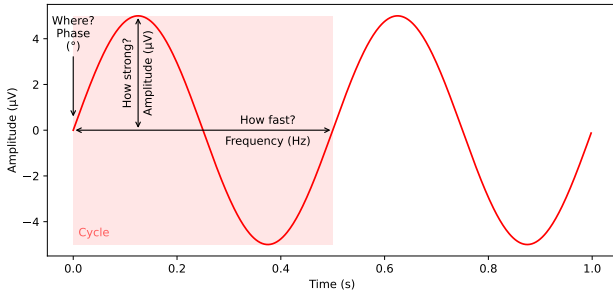


Frequency analysis

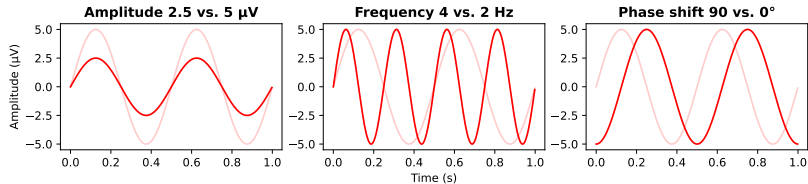
Goal: Examine which frequencies (oscillations) contribute to a stretch of continuous EEG

Approach: Decompose the continuous EEG into a set of sine waves → Fourier transform

Sine waves



Sine waves



Fourier transform



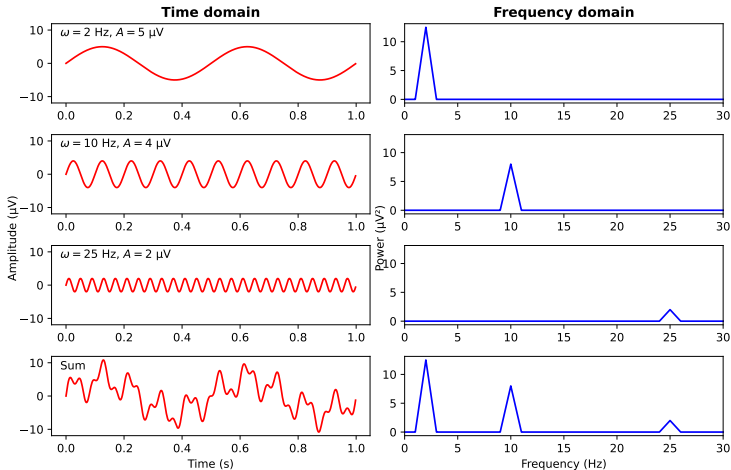
Joseph Fourier (1768–1830):

Any signal can be expressed as sum of weighted sine waves, each with its own frequency, amplitude, and phase

$$f(t) = A_0 + A_1 \cos(\omega t + \varphi_1) + A_2 \cos(\omega t + \varphi_2) + \dots + A_N \cos(\omega t + \varphi_N)$$

$$= \sum_{n=0}^N A_n \cos(\omega t + \varphi_n) \quad \text{where } A = \text{amplitude, } \omega = \text{frequency, } \varphi = \text{phase}$$

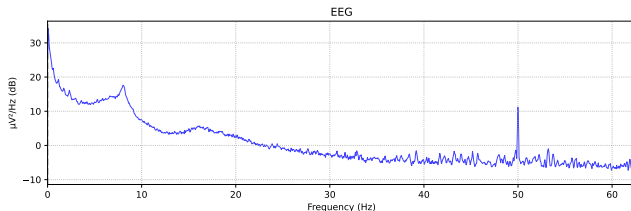
Fourier transform



Frequency analysis in MNE

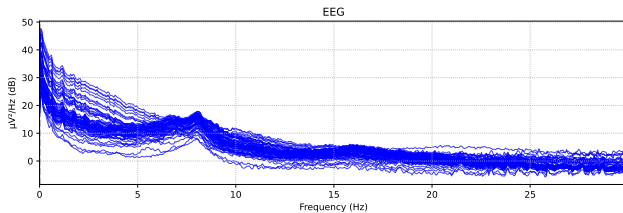
```
# Read raw data and downsample
from mne.io import read_raw_brainvision
raw = read_raw_brainvision('data/raw/05.vhdr')
raw = raw.resample(125.)

# Spectrum for a single channel
_ = raw.plot_psd(picks='Cz', color='b', spatial_colors=False)
```

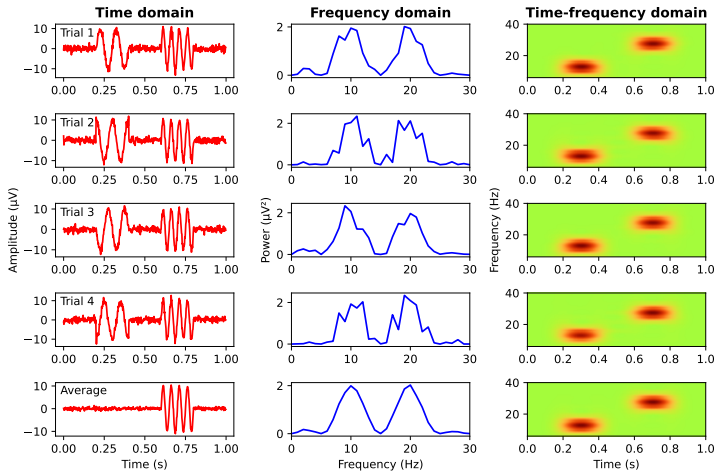


Frequency analysis in MNE

```
# Spectrum for all channels, restricted to 0--30 Hz  
_ = raw.plot_psd(fmax=30, color='b', spatial_colors=False)
```



Frequency analysis

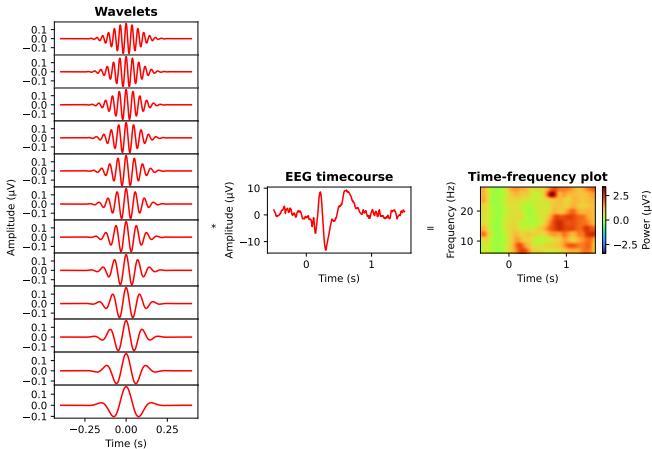


Time-frequency analysis

Goal: Estimate how power (or phase) at each frequency changes over time, e.g., in response to a stimulus

Approach: Apply the Fourier transform to a short time window and move this window over time points in the epoch → Short-term Fourier transform (STFT), Morlet wavelet convolution

Morlet wavelets



Baseline correction

- Must account for the $\frac{1}{f}$ scaling \rightarrow Divisive baseline
 - Subtract + divide: Percent signal change
 - Divide + take logarithm: Decibel
- Should end *before* stimulus onset to prevent smearing

Time-frequency analysis in MNE-Python

```
import numpy as np
from mne import events_from_annotations, Epochs
from mne.time_frequency import tfr_morlet

# Segmentation to epochs
events, _ = events_from_annotations(raw, regexp='Stimulus')
triggers = [201, 205]
epochs = Epochs(raw, events, triggers, tmin=-0.5, tmax=1.5, baseline=(-0.2, 0.))
print(epochs.get_data().shape)

## (240, 64, 251)

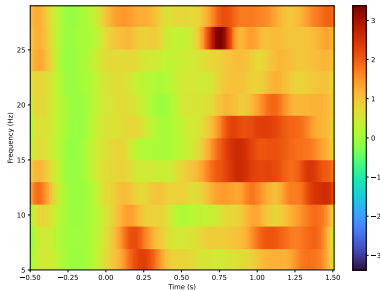
# Morlet wavelet decomposition
freqs = np.arange(6., 30., step=2.)
n_cycles = np.arange(3., 15., step=1.)
tfr = tfr_morlet(epochs, freqs, n_cycles, return_itc=False, average=False)
print(tfr.data.shape)

## (240, 64, 12, 251)
```


Time-frequency analysis in MNE-Python

```
# Divisive baseline correction to get percent signal change
tfr = tfr.apply_baseline((-0.3, -0.1), mode='percent')

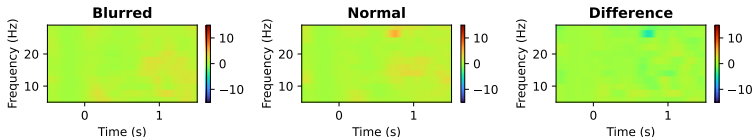
# Plot power averaged across all epochs and channels
tfr_ave = tfr.average()
tfr_ave.plot(picks='Cz', cmap='turbo')
```



Time-frequency analysis in MNE-Python

```
# Compute power per condition and the difference between conditions
tfr_aves = {'Blurred': tfr['201'].average(),
            'Normal': tfr['205'].average(),
            'Difference': tfr['201'].average() - tfr['205'].average()}

# Create multi-panel plot
import matplotlib.pyplot as plt
fig, axs = plt.subplots(nrows=1, ncols=len(tfr_aves), figsize=(9, 2))
for ix, (label, tfr_ave) in enumerate(tfr_aves.items()):
    axs[ix].set_title(label, weight='bold')
    tfr_ave.plot(picks='Cz', cmap='turbo', vmin=-15, vmax=15,
                axes=axs[ix], show=False)
```



Time-frequency analysis with the pipeline

```
# Import the Python package from R
pipeline <- reticulate::import("pipeline")

# Run the pipeline
res <- pipeline$group_pipeline(
  vhdr_files = "data/raw",
  log_files = "data/log",
  output_dir = "output",
  ocular_correction = "data/cali",
  triggers = c(201:208, 211:218),
  average_by = c("n_b", "DeviantPosRL", "n_b/DeviantPosRL"),
  perform_tfr = TRUE,
  tfr_freqs = seq(6, 30, 2),
  tfr_cycles = seq(3, 15, 1),
  tfr_baseline_tmin = -0.3,
  tfr_baseline_tmax = -0.1,
  tfr_components = list(
    "name":list("alpha"),
    "tmin":list(0.0),
    "tmax":list(0.2),
    "fmin":list(8.),
    "fmax":list(14.),
    "roi":list(
      c("P09", "P07", "P03", "P0z", "P04", "P08", "P010", "01", "0z", "02"),
    )
  )
)
```

Time-frequency analysis with the pipeline

- New column in the single trial data frame → Linear mixed model
- Permutation tests extended to the third (frequency) dimension

General considerations

- New parameters to be taken care of:
 - Design: Longer inter-trial interval
 - Wavelet frequencies + number of cycles (!)
 - Baseline correction window and method
 - Interpretation (narrow-band vs. broad-band, oscillation vs. rate of change)
- Everything matters