

2025-2026 | Neurorobotics
Lab02 | GDF data and EEG manipulation

1. Installing and using the biosig toolbox

- a. Follow the instructions in the slides
- b. Download biosig toolbox from <http://biosig.sourceforge.net/download.html>
- c. Add biosig toolbox to your MATLAB path
- d. Verify that you can call the function `sload()`

2. EEG data

- a. Download EEG data from the repository in the Moodle
- b. For now, use only the first GDF file:
`ah7.20170613.161402.offline.mi.mi_bhbf.gdf`

3. First plots of EEG data

- a. Load the EEG data from the provided GDF file with `sload()` function
- b. Plot 5 seconds of an EEG channel (amplitude vs. time)
- c. Subplot 5 seconds of 3 EEG channels (amplitude vs. time)
- d. Adjust the plots to have the same scale for all plotted channels

4. Creation of the label vectors for the provided GDF file

- a. Create `Tk` [*trial vector*] (1, 2, 3, ... N)
- b. Create `Fk` [*fixation periods*] (0 or event value)
- c. Create `Ak` [*cue periods*] (0 or event value)
- d. Create `CFk` [*continuous feedback periods*] (0 or event value)
- e. Create `Xk` [*hit/miss periods*] (0 or event value)
- f. Plot the label vectors

5. Concatenate each offline GDF file provided

- a. Load each offline GDF file
- b. Concatenate the EEG
- c. Concatenate the events (POS, TYP, DUR).
Be aware of the positions of the events from the second GDF file on

6. Creation of the label vectors for the concatenated data

7. Trial extraction

- a. Exploiting the label vectors, extract trials from the concatenated data
- b. Store the trials in a matrix [`samples x channels x trials`]
- c. Create a vector `Ck` with cue information regarding each trial [`trials x 1`]

8. Visualization

- a. Select a trial for each cue and plot a channel of your choice
- b. Compute the grand averages for each cue and plot a channel of your choice