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SignalFlowEEG Book

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SignalFlowEEG Book

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Overview

Overview

This EEG signal analysis program is designed to provide a comprehensive and user-friendly solution for processing and analyzing EEG data in the context of clinical trials and studies. It leverages the power of open-source libraries, such as MNE, to offer a robust and flexible framework for clinical EEG analysis.

Features

- **Data Preprocessing:** Perform standard EEG preprocessing tasks, including filtering, artifact detection and removal, and channel selection.
- **Event Detection and Segmentation:** Accurately identify and segment EEG events of interest, such as evoked responses, oscillatory activity, and sleep stages.
- **Signal Analysis:** Apply a wide range of signal processing techniques, including time-domain analysis, frequency-domain analysis, and time-frequency analysis.
- **Statistical Analysis:** Conduct statistical tests to assess the significance of EEG findings, including group comparisons and correlations with clinical outcomes.
- **Visualization:** Generate informative visualizations, such as topographic maps, time-series plots, and spectrograms, to facilitate data exploration and presentation.
- **Integration with Clinical Data:** Seamlessly integrate EEG data with other clinical measures, such as cognitive assessments, behavioral observations, and medical records, to enable multimodal analysis.
- **Scalable and Automated Workflows:** Develop and deploy scalable and automated workflows to handle large-scale clinical datasets efficiently.
- **Reproducibility and Collaboration:** Ensure the reproducibility of analyses and enable collaborative work through the use of version control and documented workflows.

1 Set-up

This is a template for a simple Quarto book output to html, PDF or docx format. It includes a GitHub Action that will build the website automatically when you make changes to the files. The NOAA palette and fonts has been added to `theme.scss`. The webpage will be on the `gh-pages` branch. Serving the website files from this branch is a common way to keep all the website files from cluttering your main branch.

The GitHub Action installs R so you can have R code in your `qmd` or `Rmd` files. Note, you do not need to make changes to your `Rmd` files unless you need Quarto features like cross-references.

1.1 GitHub Set-up

- Click the green “use template” button to make a repository with this content. Make sure to make your repo public (since GitHub Pages doesn’t work on private repos unless you have a paid account) and check box to include all the branches (so that you get the `gh-pages` branch).
- Turn on GitHub Pages under Settings > Pages . You will set pages to be made from the `gh-pages` branch and root directory.
- Turn on GitHub Actions under Settings > Actions > General
- Edit the repo description and Readme to add a link to the webpage. When you edit the description, you will see the link url in the url box or you can click on the Actions tab or the Settings > Pages page to find the url.

2 Customize

2.1 Edit and add your pages

Edit the qmd or md files in the `content` folder. qmd files can include code (R, Python, Julia) and lots of Quarto markdown bells and whistles (like call-outs, cross-references, auto-citations and much more).

Each page should start with

```
---  
title: your title  
---
```

and the first header will be the 2nd level, so `##`. Note, there are situations where you leave off

```
---  
title: your title  
---
```

and start the qmd file with a level header `#`, but if using the default title yaml (in the `---` fence) is a good habit since it makes it easy for Quarto convert your qmd file to other formats (like into a presentation).

2.2 Add your pages the project

- Add the files to `_quarto.yml`

3 Customization

3.1 Quarto documentation

Quarto allow many bells and whistles to make nice output. Read the documentation [here](#) Quarto documentation.

3.2 Examples

Looking at other people's Quarto code is a great way to figure out how to do stuff. Most will have a link to a GitHub repo where you can see the raw code. Look for a link to edit page or see source code. This will usually be on the right. Or look for the GitHub icon somewhere.

- [Quarto gallery](#)
- [nmfs-openscapes](#)
- [Faye lab manual](#)
- [quarto-titlepages](#) Note the link to edit is broken. Go to repo and look in `documentation` directory.

4 Customization

4.1 Assessing Cortical Excitability Using Resting-State EEG

4.1.1 Introduction

Resting state EEG analysis can provide valuable insights into cortical excitability, which is often altered in neurodevelopmental disorders like Fragile X Syndrome (FXS). One powerful tool for assessing cortical excitability is the Fitting Oscillations & One-Over F (FOOOF) algorithm, which can decompose the resting state EEG power spectrum into its underlying oscillatory and aperiodic components.

In this analysis, we will use FOOOF to compare cortical excitability, as measured by the aperiodic component of the power spectrum, between a group of individuals with FXS and a matched control group. This will allow us to better understand the neurophysiological underpinnings of the altered sensory processing and hyperexcitability often observed in FXS.

4.1.2 Methods Overview

1. `p050_setup_environment`: Set up the analysis environment
2. `p100_load_data_catalog`: Retrieve file list and associated metadata
 - Preprocess the resting state EEG data, including filtering, artifact rejection, and channel selection.
 - Apply the FOOOF algorithm to the preprocessed, source localized data to extract the aperiodic and oscillatory components of the power spectrum for each participant.
 - Compare the aperiodic component between the FXS and control groups using appropriate statistical tests, such as t-tests or non-parametric tests, depending on the distribution of the data.

5 Rendering

The repo includes a GitHub Action that will render (build) the website automatically when you make changes to the files. It will be pushed to the `gh-pages` branch.

But when you are developing your content, you will want to render it locally.

5.1 Step 1. Make sure you have a recent RStudio

Have you updated RStudio since about August 2022? No? Then update to a newer version of RStudio. In general, you want to keep RStudio updated and it is required to have a recent version to use Quarto.

5.2 Step 2. Clone and create RStudio project

First, clone the repo onto your local computer. How? You can click `File > New Project` and then select “Version Control”. Paste in the url of the repository. That will clone the repo on to your local computer. When you make changes, you will need to push those up.

5.3 Step 3. Render within RStudio

RStudio will recognize that this is a Quarto project by the presence of the `_quarto.yml` file and will see the “Build” tab. Click the “Render website” button to render to the `_site` folder.

Previewing: You can either click `index.html` in the `_site` folder and specify “preview in browser” or set up RStudio to preview to the viewer panel. To do the latter, go to `Tools > Global Options > R Markdown`. Then select “Show output preview in: Viewer panel”.

6 R Markdown

You can include R Markdown files in your project.

6.1 R Markdown

This is an R Markdown document. Markdown is a simple formatting syntax for authoring HTML, PDF, and MS Word documents. For more details on using R Markdown see <http://rmarkdown.rstudio.com>.

When you click the **Knit** button a document will be generated that includes both content as well as the output of any embedded R code chunks within the document. You can embed an R code chunk like this:

```
summary(cars)
```

speed		dist	
Min.	: 4.0	Min.	: 2.00
1st Qu.	:12.0	1st Qu.	: 26.00
Median	:15.0	Median	: 36.00
Mean	:15.4	Mean	: 42.98
3rd Qu.	:19.0	3rd Qu.	: 56.00
Max.	:25.0	Max.	:120.00

6.2 Including Plots

You can also embed plots and reference them, like so Figure 6.1.

Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.

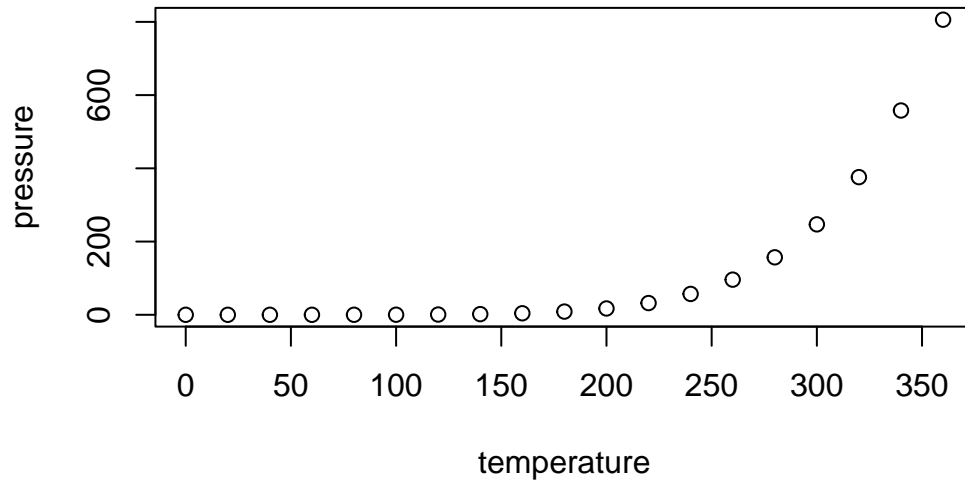


Figure 6.1: Plot of pressure

6.3 Including Tables

You can also embed tables and reference them with Table 6.1.

```
library(knitr)
kable(head(iris))
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
5.1	3.5	1.4	0.2	setosa
4.9	3.0	1.4	0.2	setosa
4.7	3.2	1.3	0.2	setosa
4.6	3.1	1.5	0.2	setosa
5.0	3.6	1.4	0.2	setosa
5.4	3.9	1.7	0.4	setosa

7 Rendering with Code

You can have code (R, Python or Julia) in your qmd file. You will need to have these installed on your local computer, but presumably you do already if you are adding code to your qmd files.

```
x <- c(5, 15, 25, 35, 45, 55)
y <- c(5, 20, 14, 32, 22, 38)
lm(x ~ y)
```

Call:

```
lm(formula = x ~ y)
```

Coefficients:

(Intercept)	y
1.056	1.326

7.1 Modify the GitHub Action

You will need to change the GitHub Action in `.github/workflows` to install these and any needed packages in order for GitHub to be able to render your webpage. The GitHub Action install R since I used that in `code.qmd`. If you use Python or Julia instead, then you will need to update the GitHub Action to install those.

If getting the GitHub Action to work is too much hassle (and that definitely happens), you can always render locally and publish to the `gh-pages` branch. If you do this, make sure to delete or rename the GitHub Action to something like

```
render-and-publish.old.yml
```

so GitHub does not keep trying to run it. Nothing bad will happen if you don't do this, but if you are not using the action (because it keeps failing), then you don't need GitHub to run it.

7.2 Render locally and publish to gh-pages branch

To render locally and push up to the `gh-pages` branch, open a terminal window and then `cd` to the directory with the Quarto project. Type this in the terminal:

```
quarto render gh-pages
```


8 References

Quarto has powerful references functionality. You can easily insert citations from Zotero libraries that you maintain in the cloud (on Zotero). This allows the whole team to update the library and you can sync up to that library. Read about this on the Quarto documentation on citations. Google youtube videos on this also to see it in action.

Add a `.bib` file in to your project or add a linked Zotero library via RStudio in Visual mode with Tools > Project Options... > R Markdown > select custom libraries from the Zotero dropdown.

Then you can type @ and you will see a dropdown of the references in your libraries. You can then select the ones to add. If you don't see the one you need, you can paste in the DOI and it will be added to your references file (with all the info). The references will be added to your references section of your book automatically.

See the `references.qmd` file for how to include the references.

- @ansley1981 will produce Ansley and Davis (1981)
- [@ansley1981] will produce (Ansley and Davis 1981).

References

Ansley, H. L. H., and C. D. Davis. 1981. "Migration and Standing Stock of Fishes Associated with Artificial and Natural Reefs on Georgia's Outer Continental Shelf." Brunswick, Georgia, USA.