Report from Brainhack Global 2020 Montreal

Load fMRIprep confounds in Python

Project URL: https://github.com/htwangtw/2021_08_Wang_load-confounds

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1 Introduction

fMRIprep [1] is a popular minimal preprocessing software for functional MRI data. 'Minimal preprocessing' refers to motion correction, field unwarping, normalization, bias field correction, and brain extraction. Confound regression and smoothing are exculded from the workflow. Instead, fMRIprep provides users with a large set of potential confound regressors that covers many denoising strategies. The users will have to select the confound regressors for denoising in the subsequent analysis. Loading a sensible subset of confounds is difficult and error prone for many strategies, such as ICA-AROMA [2] and CompCor [3]. load_confounds can access confound variables and provides preset strategies for confound selections. The loaded format is competible with nilearn analysis functions such as NiftiMasker and the GLM modules. The aim is to provide a easy and foolproof API for users to perform subsequent denoising of fMRIprep output.

2 Progress

At Brianhack Global Montreal 2020, the aim is to prepare the package ready for a potential Beta release. To prepare for the release, related issues involves completing the strategies missing and improve the user experience with better examples and error messages. Several issues has been identified before Brainhack and the full discussion can be found under load_confounds GitHub issue page. Participants at Brainhack were encouraged to pick up existing issues from the list. The following issues have been discussed and/or resolved:

2.1 Strategies

We worked on three strategies:

- Added ICA-AROMA [2] (contributed by Hao-Ting Wang)
- Added Scrubbing [4](contributed by Steven Meisler)
- Improved the anatomical mask selection for anatomical CompCor [3] (contributed by Steven Meisler)

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2.2 Demo

An executable demo using the nilearn developmental fMRI dataset (OpenNeuro ds000228) was added (contributed by Michael W. Weiss). The demo was adapted from an existing nilearn example on denoising 'Extracting signals from a brain parcellation'. This example notebook show how to extract signals from a brain parcellation and compute a correlation matrix, using different denoising strategies using the package.

2.3 Error message

New class NotFoundConfoundException was added for collecting all the missing parameter needed for the given noise component(s). A final error would be raised with the list of all parameters missing, rather than just the first encountered missing parameter. (Contributed by Michael W. Weiss and François Paugam)

2.4 Identify test dataset

OpenNeuro ds003 is adopted as the new test data, including all ICA-AROMA related components. However, non-steady-state volume and motion estmation mertic RMSD [5] are still missing. We consider to preprocess the nilearn demo dataset (OpenNeuro ds000228) with fMRIprep LTS release for getting all possible confounds for the future. (Discussions amongs Pierre Bellec, Hao-Ting Wang, Elizabeth DuPre, and Chris Markiewicz)

2.5 All contributor bot

allcontributors bot is added to track community contributions (Contributed by Pierre Bellec).

2.6 Add load_confounds to nixtract

In addition to the main package, there was a collaborative project with the developers of <code>nixtract</code>. <code>nixtract</code> is a tool that extract and process timeseries data from neuroimaging files. Annabelle Harvey and Dan Gale added <code>load_confounds</code> as a dependency of <code>nixtract</code> for reading fMRIPrep confound variables.

3 Results

load_confounds has now covered most of the noise components used in Ciric et al. [6]. The following noise components and a set of paramaters for dedicated approaches are supported.

- motion: the motion parameters including 6 translation/rotation (basic), and optionally derivatives, squares, and squared derivatives (full).
- high_pass: basis of discrete cosines covering slow time drift frequency band.
- wm_csf the average signal of white matter and cerebrospinal fluid masks (basic), and optionally derivatives, squares, and squared derivatives (full).
- **global**: the global signal (**basic**), and optionally derivatives, squares, and squared derivatives (**full**).
- compcor [3]: the results of a PCA applied on a mask based on either anatomy (anat), temporal variance (temp), or both (combined).

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• ica_aroma [2]: the results of an idependent component analysis (ICA) followed by identification of noise components. This can be implementing by incorporating ICA regressors (basic) or directly loading a denoised file generated by fMRIprep (full).

• scrub [4]: regressors coding for time frames with excessive motion, using threshold on frame displacement and standardized DVARS (basic) and suppressing short time windows using the (Power et al., 2014) appreach (full).

load_confounds can produce the following strategies from Ciric et al. [6]. The following table highlights the relevant options:

Strategy	high_pass	motion	wm_csf	global	compcor	ica_aroma	scrub
Params2	x		basic				
Params6	x	basic					
Params9	x	basic	basic	basic			full
Params9scrub	x	basic	basic				
Params24	x	full					
Params36	x	full	full	full			full
Params36scrub	x	full	full				
AnatCompCor	x	full			anat		
TempCompCor	x				temp		
ICAAROMA	x		basic			full	
AROMAGSR	x		basic	basic		full	
AggrICAAROMA	x		basic	basic		basic	

The future direction is to integrate <code>load_confounds</code> as part of <code>nilearn</code> for better reach to wider range of users that can be benifited from the package. To facilitate the nilearn inegration, we will add <code>sample_mask</code> to support volume-sensoring based scrubbing and improve the <code>nilearn</code> <code>NiftiMasker</code> related feature.

Availability of Supporting Data

Supplemental material has not been provided. More information about this project can be found at: https://github.com/htwangtw/2021_08_Wang_load-confounds. This report is generated at: https://github.com/htwangtw/2021_08_Wang_load-confounds.

Competing interests

None

Author's contributions

HTW wrote the software, HTW wrote the report.

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Reviewers

No reviewers has been added yet.

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