

Week 6 – First-level fMRI data analysis

L06-07. Residualize (canlab_connectivity_preproc.m)

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❖ Canlab_connectivity_preproc.m


 master ▾




CanlabCore / CanlabCore / @fmri_data / canlab_connectivity_preproc.m

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 **wanirepo** make the residualize more efficient by reordering the matrix product ✓

Latest commit 9746178 on 2 Aug 2019  History

👤 3 contributors   

681 lines (542 sloc) | 23.5 KB

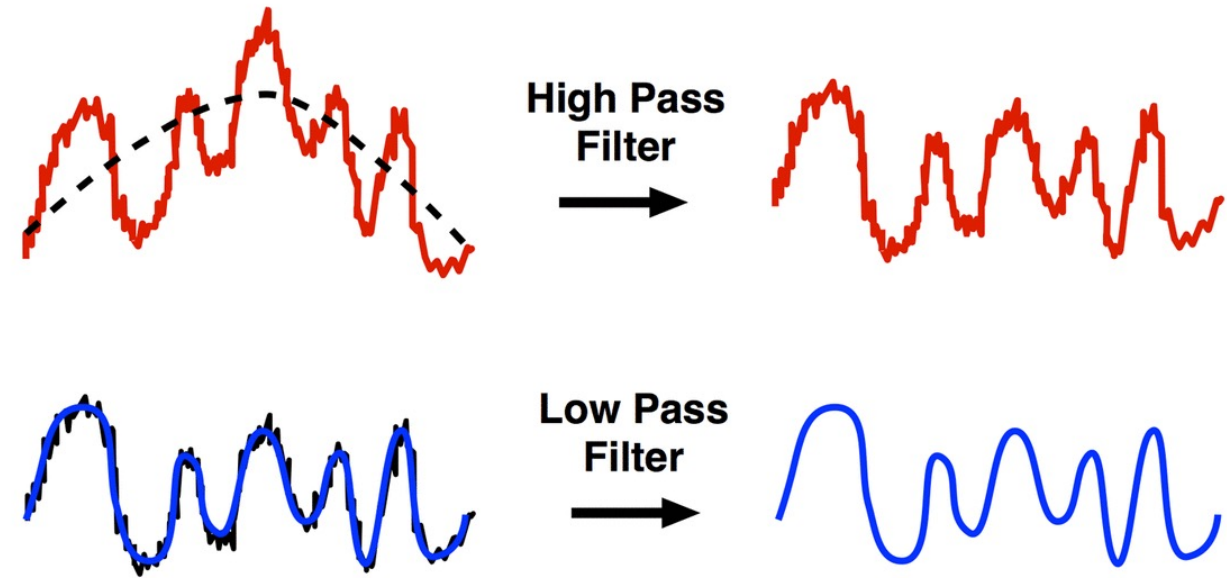
Raw Blame   

```
1 function [dat, roi_val, maskdat, beta_dat, beta_roi_val] = canlab_connectivity_preproc(dat, varargin)
2
3 % This function prepares data for connectivity analysis by removing nuisance
4 % variables and temporal filtering (high, low, or bandpass filter). This also
5 % can extract values from given masks and return averaged activity or pattern
6 % expression values.
7 %
```



❖ Additional preprocessing

- **'canlab_connectivity_preproc.m'** function
 - additional_nuisance
 - temporal filtering (e.g., bpf, lpf, hpf)
 - winsorize



❖ Why do you use this instead of spm batch?

- When trial structure is not clear but you need to residualize using nuisance and apply pass filtering
 - Continuous design (naturalistic design)
 - Connectivity analysis



❖ Features

```
roi_masks = which('weights_NSF_grouppred_cvpcr.img');  
[preprocessed_dat, roi_val] = canlab_connectivity_preproc(dat, 'vw', 'datdir',  
    subject_dir, 'bpf', [.008 .25], TR, 'extract_roi', roi_masks, 'pattern_expression');
```

- can regress **out nuisance variables with any additional nuisance matrix**
- can remove signal from ventricle and white matter
- can do temporal filtering, including high-pass, low-pass, or bandpass filtering (it uses conn_filter.m from conn toolbox, which is using fast fourier transform)
- can extract data from given ROIs and return averaged value or pattern expression value (dot-product)
- It is possible to **extract ROI averages** or linear patterns, save them, and run this function afterwards. This will give the same result as filtering each voxel and then calculating the ROI average/pattern response.
- (optional) can run additional GLM model with additional regressors if they are specified. This runs the GLM *additionally*, therefore do not change residualized image data or roi_vals. This GLM uses the same nuisance matrix from above. output: beta_dat



❖ Steps in order

1. Remove nuisance covariates (and linear trend if requested)
2. Remove ventricle and white matter - default: uses canonical images
3. High/low/bandpass filter on BOTH data and covariates before regression (plot)
4. Residualization using regression (plot)
5. (optional) Run additional GLM using the same additional preprocessing -> beta_dat
6. Winsorize based on distribution of full data matrix (plot)
7. Extract region-by-region average ROI or pattern expression data

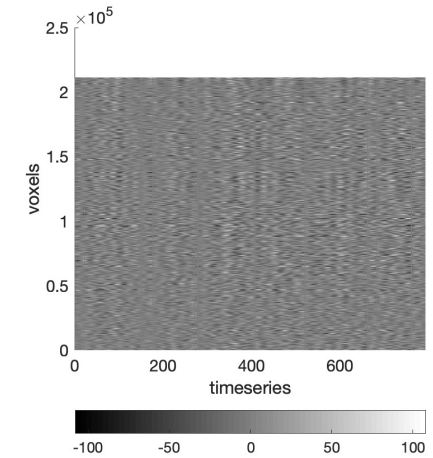
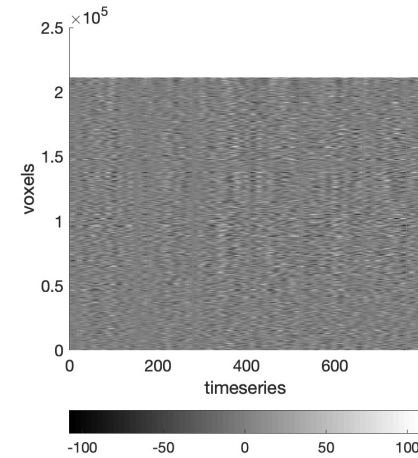
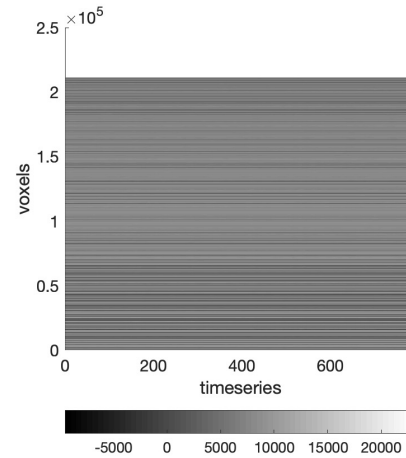
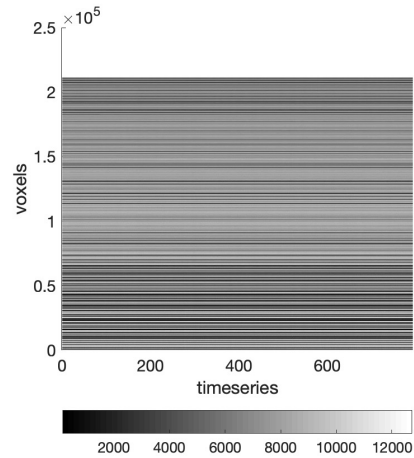
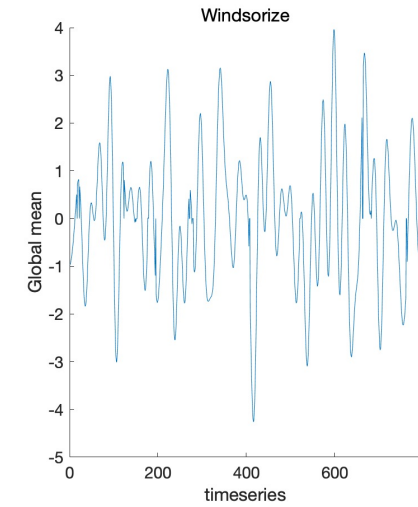
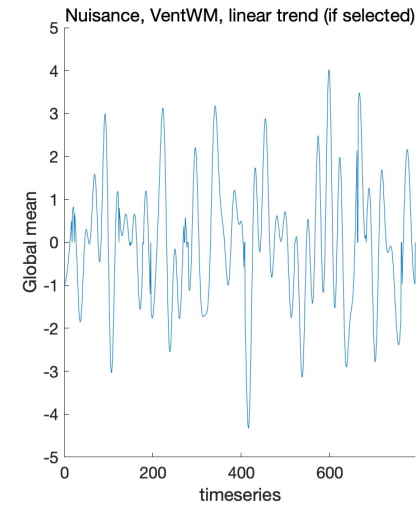
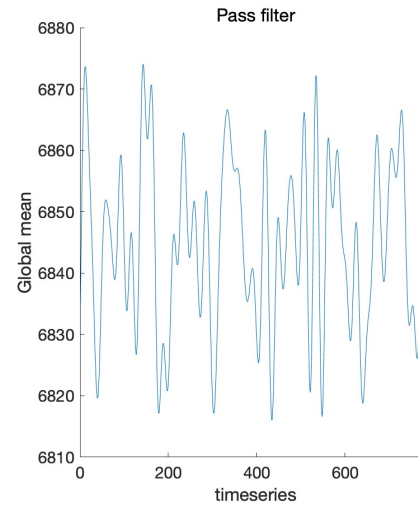
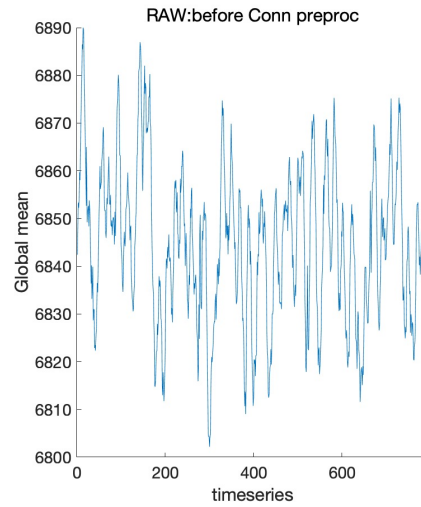


❖ Codes and plots

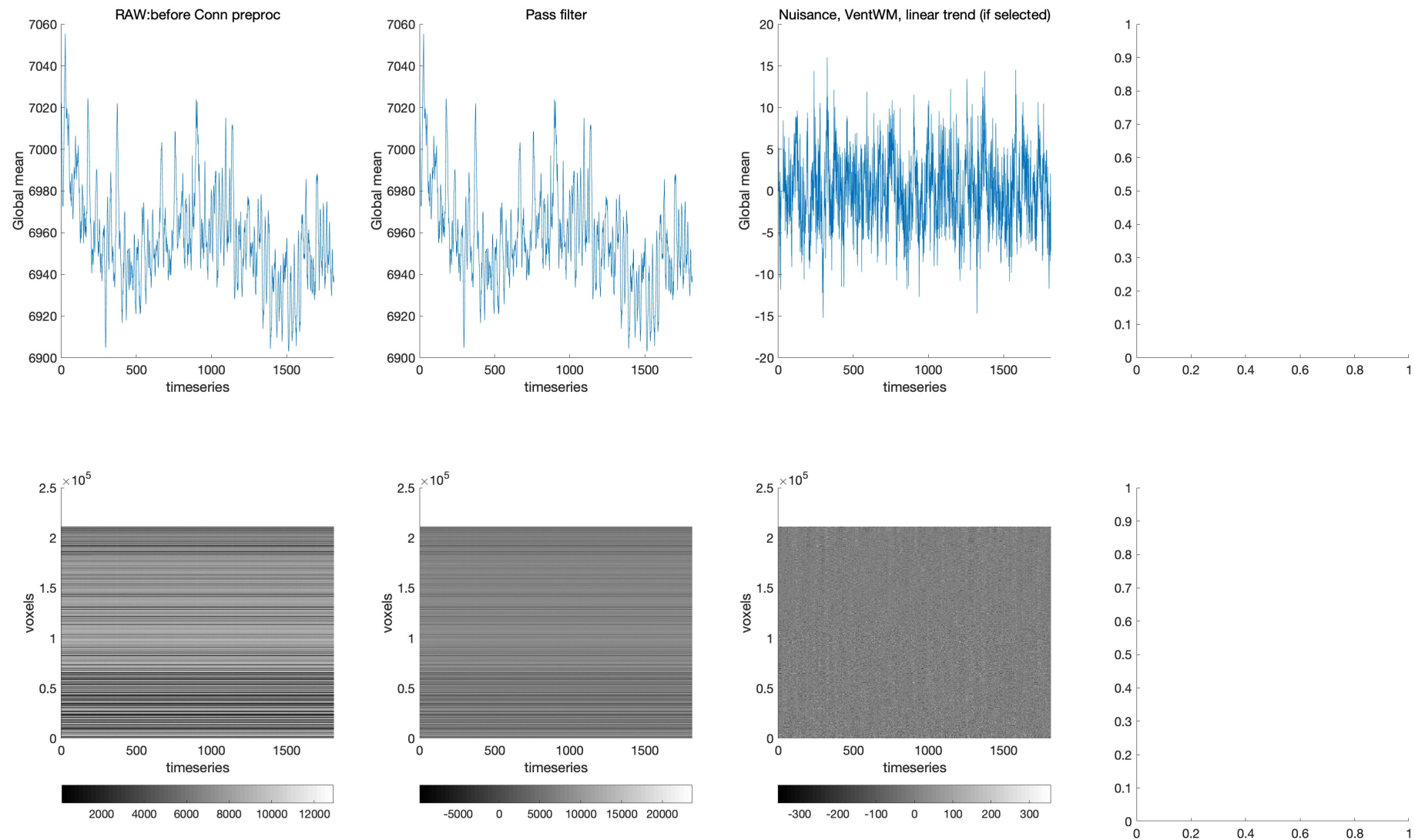
```
if run_i == 1 || run_i == 6
    preprocessed_dat = canlab_connectivity_preproc_new(dat, 'bpf', [0.008 0.1], tr, 'windsorize', 5); % no design matrix for resting runs
else
    preprocessed_dat = canlab_connectivity_preproc_new(dat, 'bpf', [0.008 0.1], tr, 'additional_nuisance', additional_reg.X{run_i-1}, 'windsorize', 5);
end
```



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Cocoan 101

<https://cocoanlab.github.io>

