

Week 6 – First-level fMRI data analysis

L06-06. Making Custom Regressors (onsets2fmridesign.m)

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❖ Contents


06-08. Making custom regressors (onsets2fmridesign.m)

06-09. Residualize (canlab_connectivity_preproc.m)

06-10. Doing GLM with custom regressors (regress.m)




❖ onsets2fmridesign.m function in CanlabCore


 978da1fdc4 ▾




CanlabCore / CanlabCore / Model_building_tools / onsets2fmridesign.m

Go to file




...

 **torwager** plotDesign cosmetic update for events; cosmetic updates to getPredictors ✓

Latest commit 978da1f on 15 Jan  History

3 contributors   

549 lines (431 sloc) | 19.3 KB

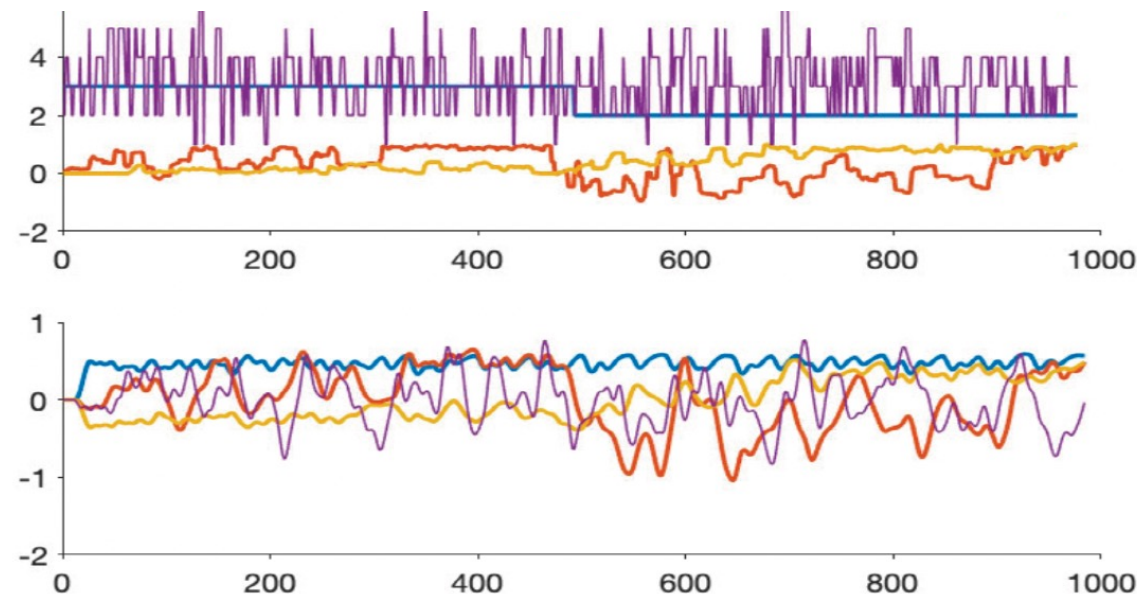
Raw Blame   

```
1 function [X, delta, delta_hires, hrf] = onsets2fmridesign(ons, TR, varargin)
2 % Builds design matrix X and delta function, given cell array of onset times in s
3 % One cell per condition per session, e.g., ons{1} = [24 27 29 44]';
```



❖ Why do we need custom regressors?

- When single-trial model is not appropriate, and you want to make “customized” regressor in a more flexible manner
- When you have **multiple experimental conditions** and want to identify brain regions that preferentially track each variable **controlling for** other variables, you need to make custom regressors for each variable



❖ How do we make custom regressors?

- Use onsets2fmridesign.m to apply HRF to the design matrix

```
X{run_i} = onsets2fmridesign([onsets{run_i}, durations{run_i}], 0.46, m*0.46, spm_hrf(1));
```

Onsets and durations
(in two columns)

TR (in seconds)

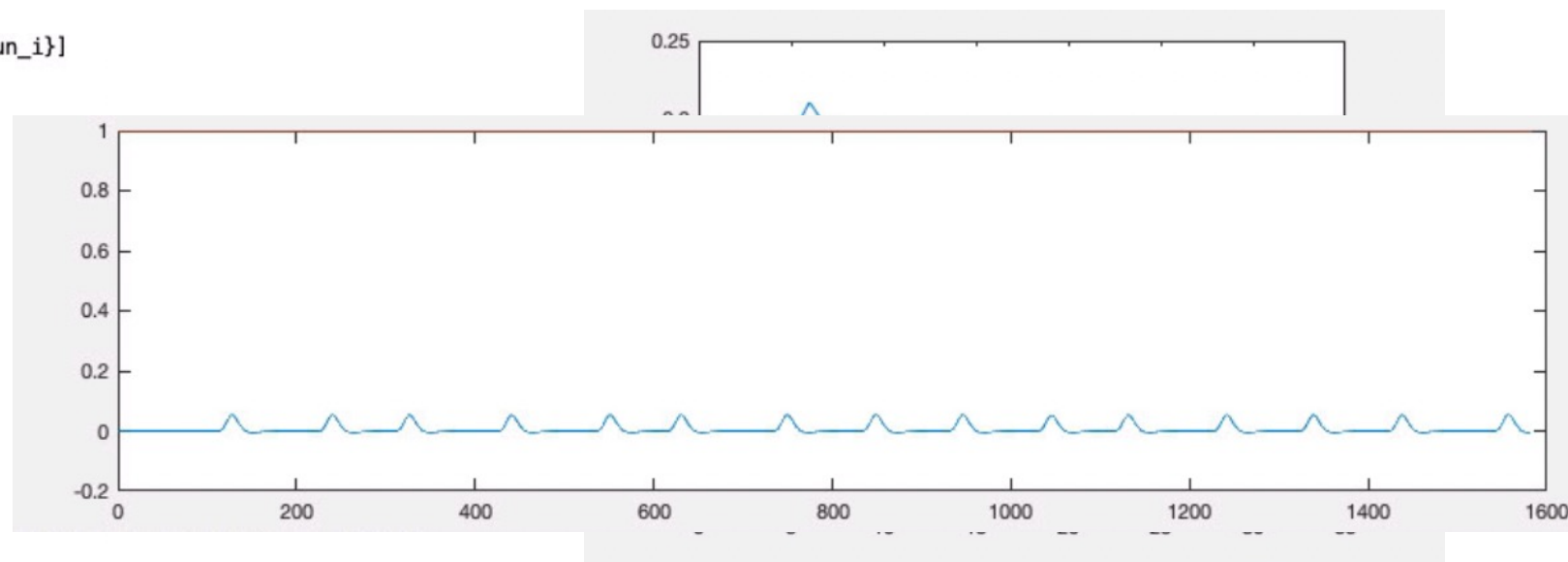
m = number of measurements in the run

spm_hrf(1)

```
>> [onsets{run_i}, durations{run_i}]
```

```
ans =
```

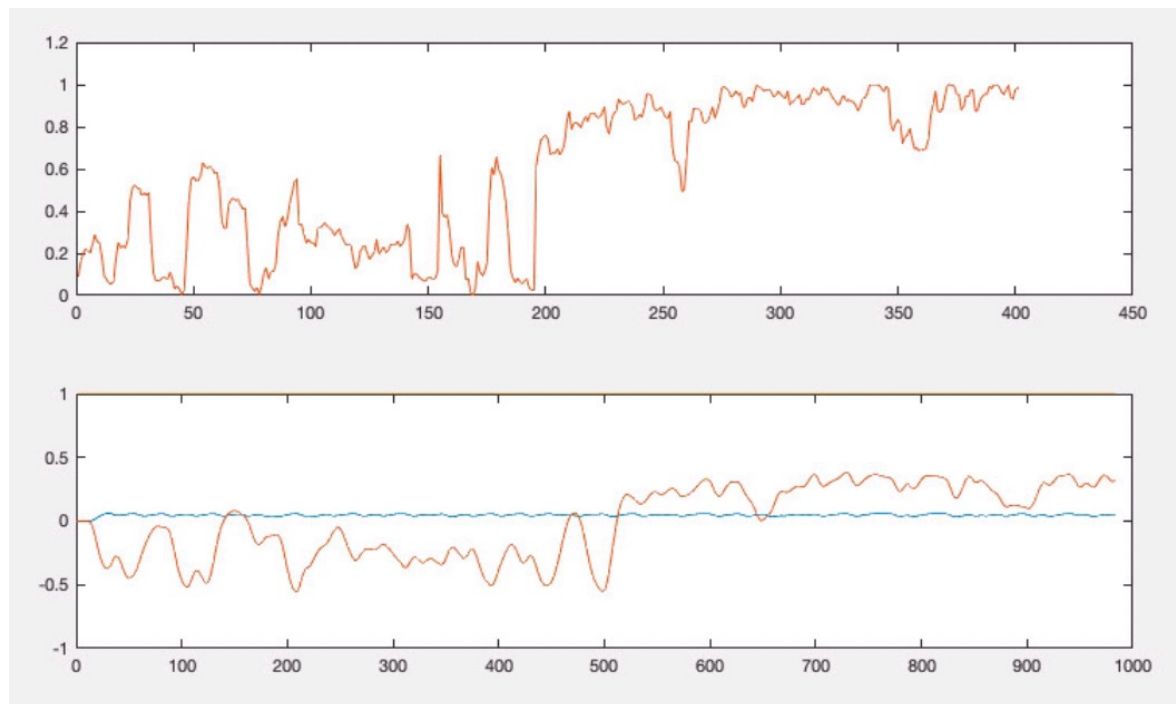
51.5173	4.8831
95.2491	4.7993
147.8637	4.8545
185.5474	4.8480
238.1989	4.8658
282.9421	4.7877
335.6083	4.7865
377.0859	4.9743
429.9263	4.7983
476.5022	4.8878
513.1364	4.9307
565.8369	4.8828
616.6137	4.7663
661.2807	4.7860
707.9787	4.9814



❖ How do we make custom regressors?

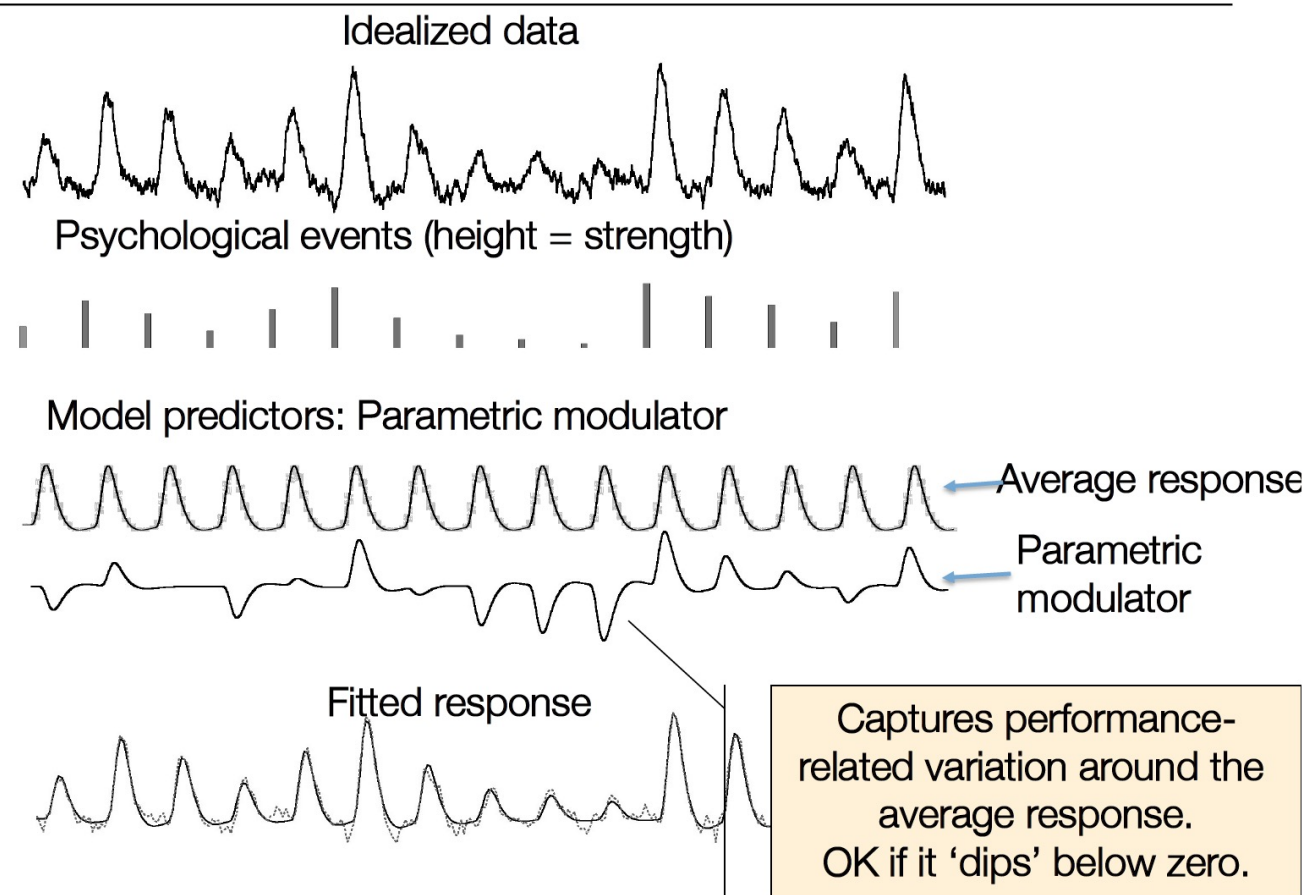
1. If you want to parametrically modulate onsets by modulator values, you can use 'parametric_standard' option and put two regressors per event type. (One to model the average response, and one for the mean-centered modulator values of modulator values in each cell).

```
X = onsets2fmridesign({[temp_onset', (temp_dur-minus_onset)']}, 0.46, m*0.46, spm_hrf(1), 'parametric_standard', {temp_wordnum'});
```



- ❖ For more details about parametric modulation, please watch:
Coursera (Principles of fMRI) - Part2:
module10 – parametric modulation
(<https://www.youtube.com/watch?v=iqwZmGOKqfM>)

Parametric modulation



Cocoan 101

<https://cocoanlab.github.io>

