

Tutorial: Basic introduction to SPM

Methods and Models in fMRI, 20.09.2016

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Translational Neuromodeling Unit

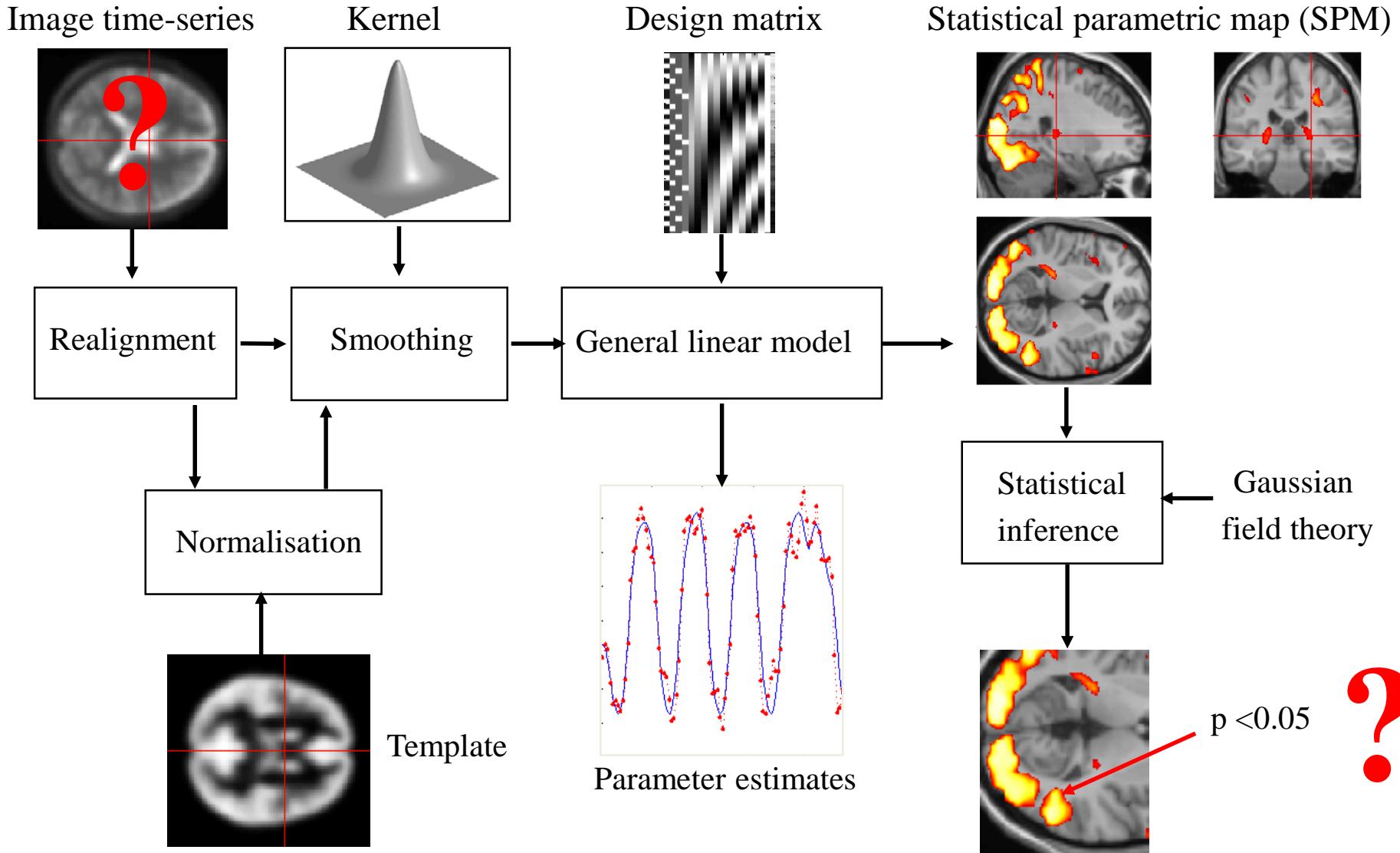


University of
Zurich^{UZH}

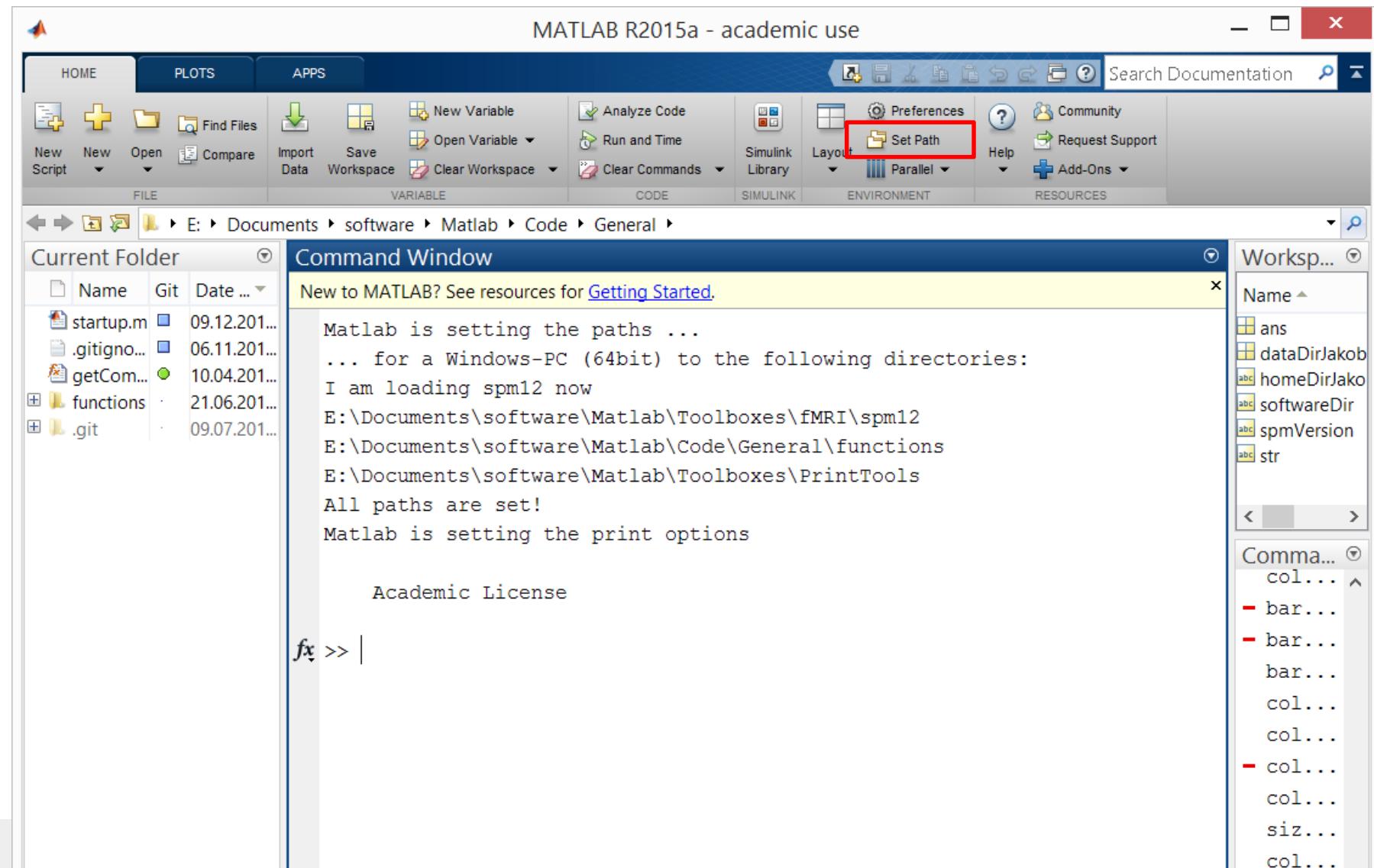
ETH

Eidgenössische Technische Hochschule Zürich
Swiss Federal Institute of Technology Zurich

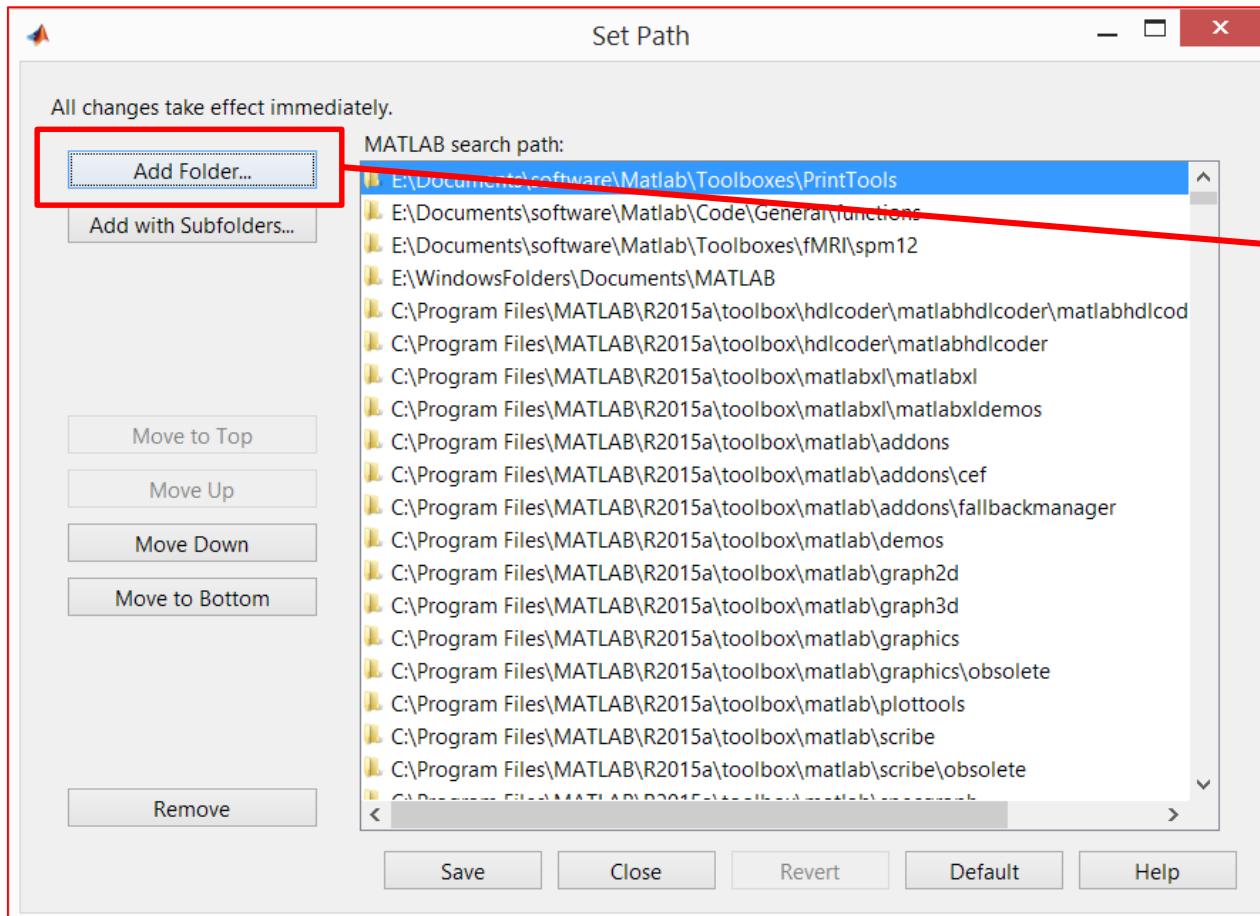
Overview of SPM



Set path for SPM

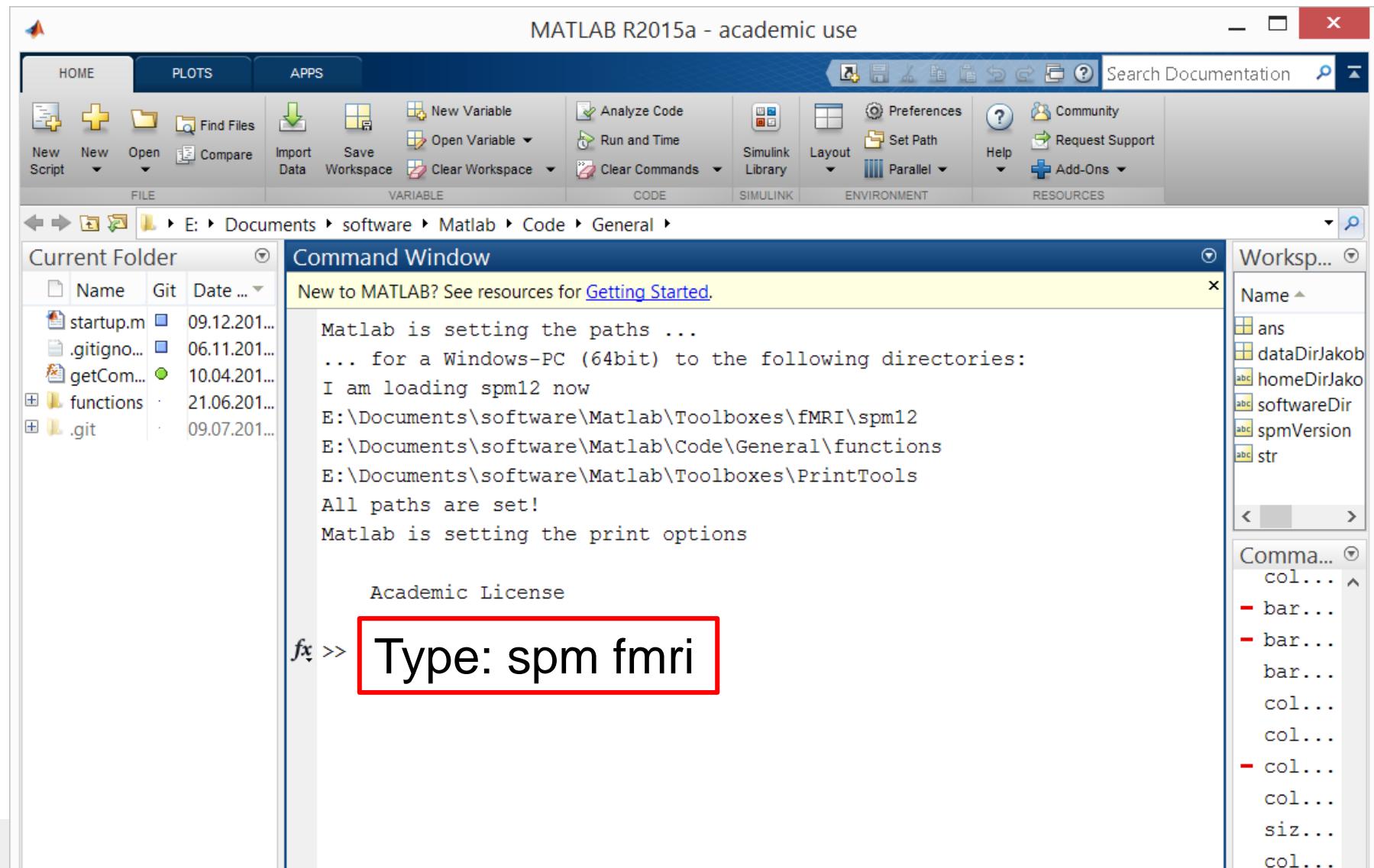


Set path for SPM

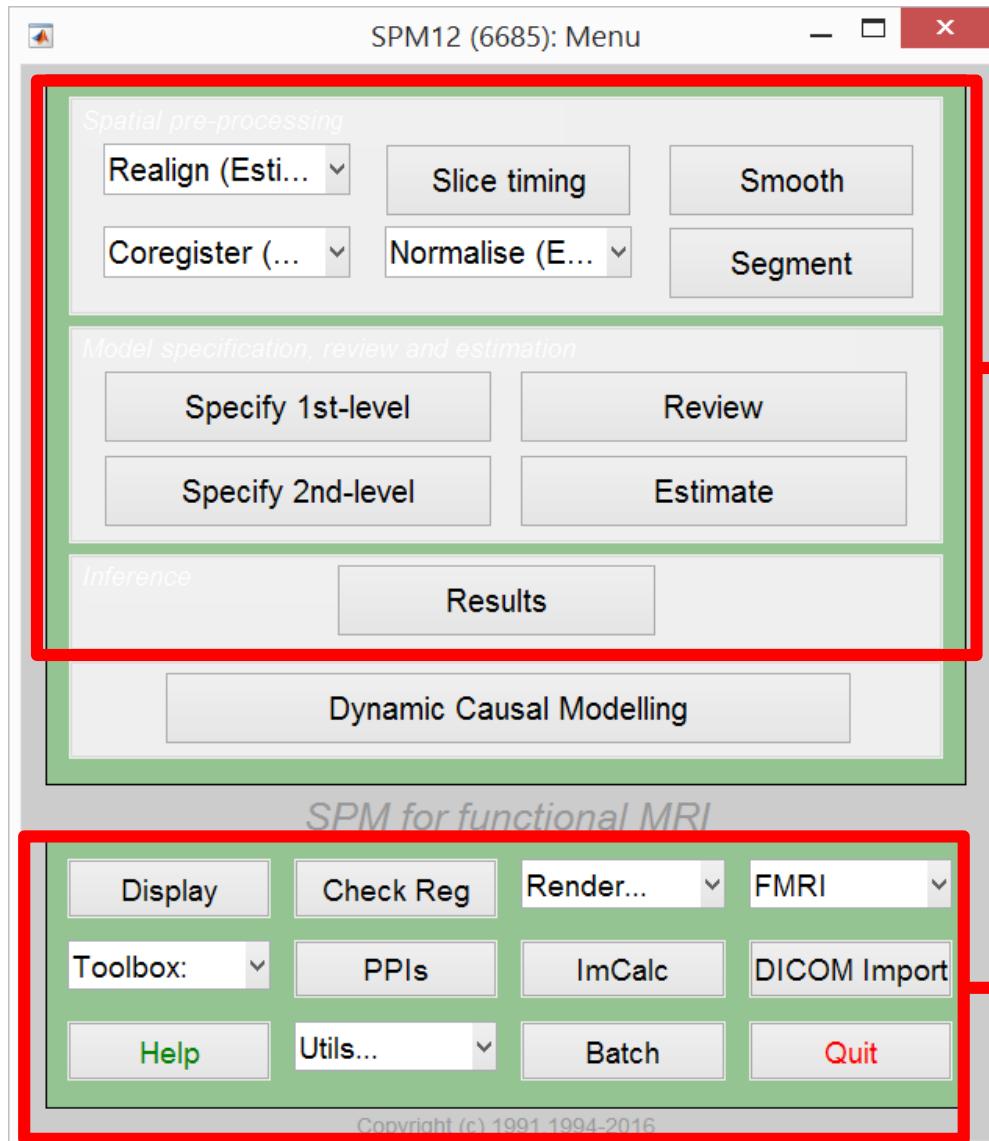


Select the
SPM folder.

Starting SPM



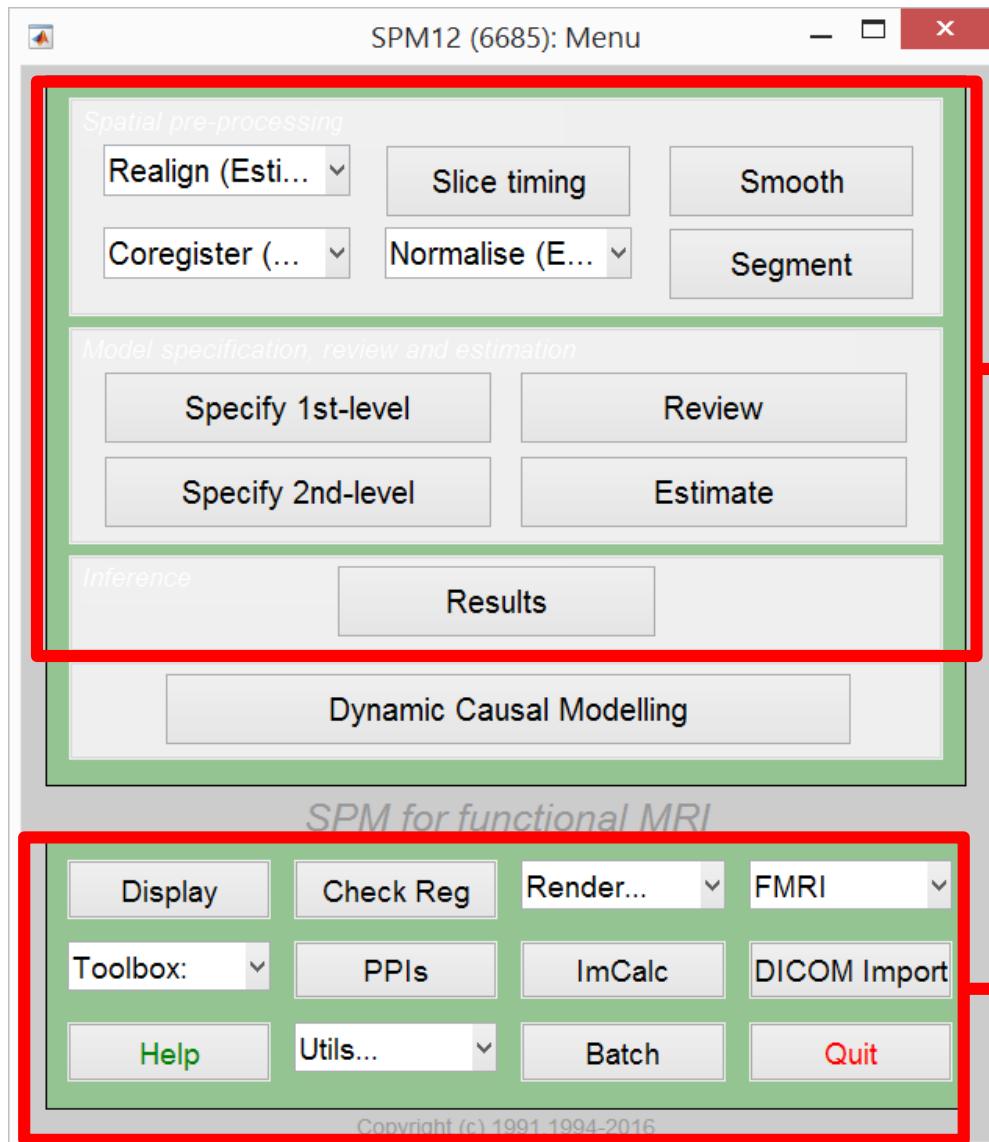
The SPM main menu



Parts needed for
basic fMRI analysis.

Additional useful
functions.

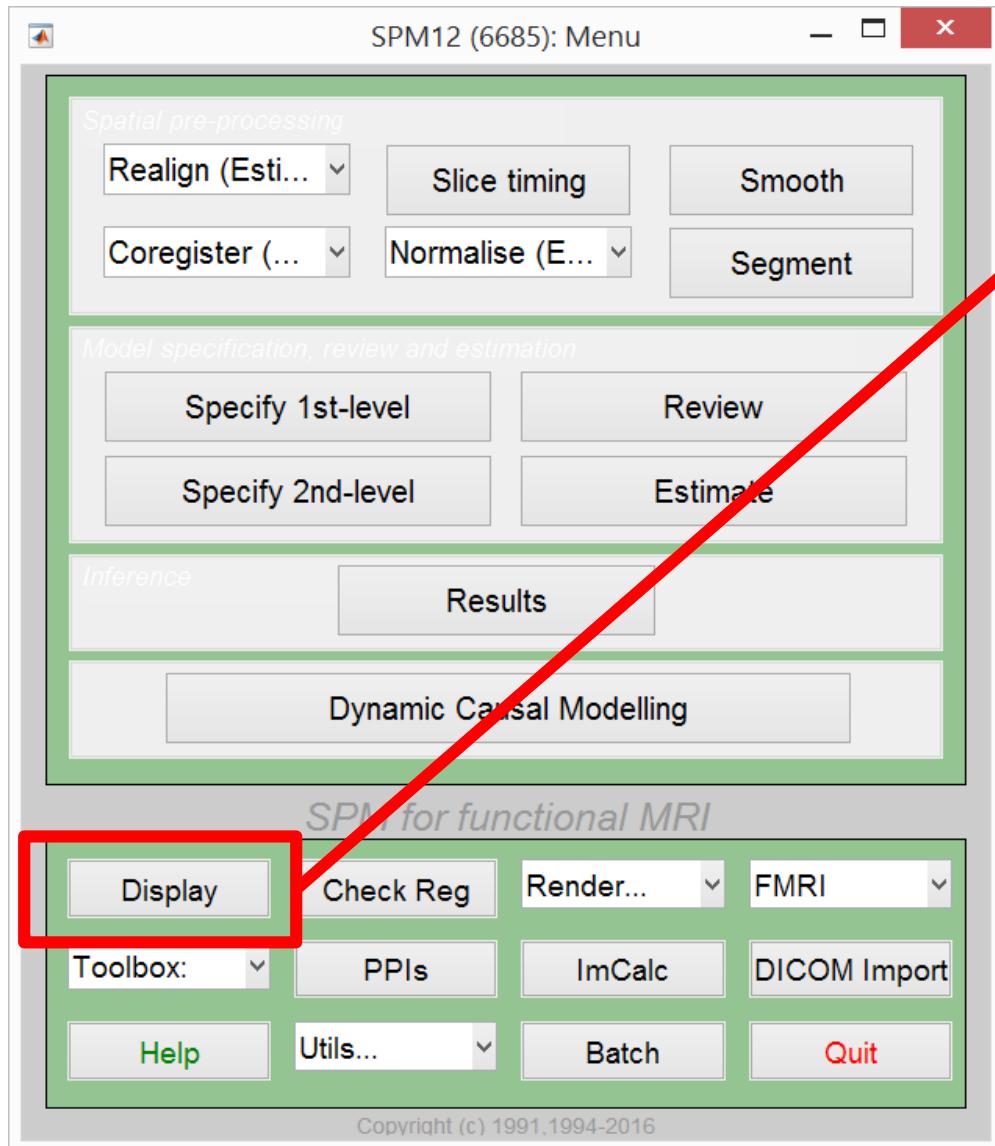
The SPM main menu



Parts needed for
basic fMRI analysis.

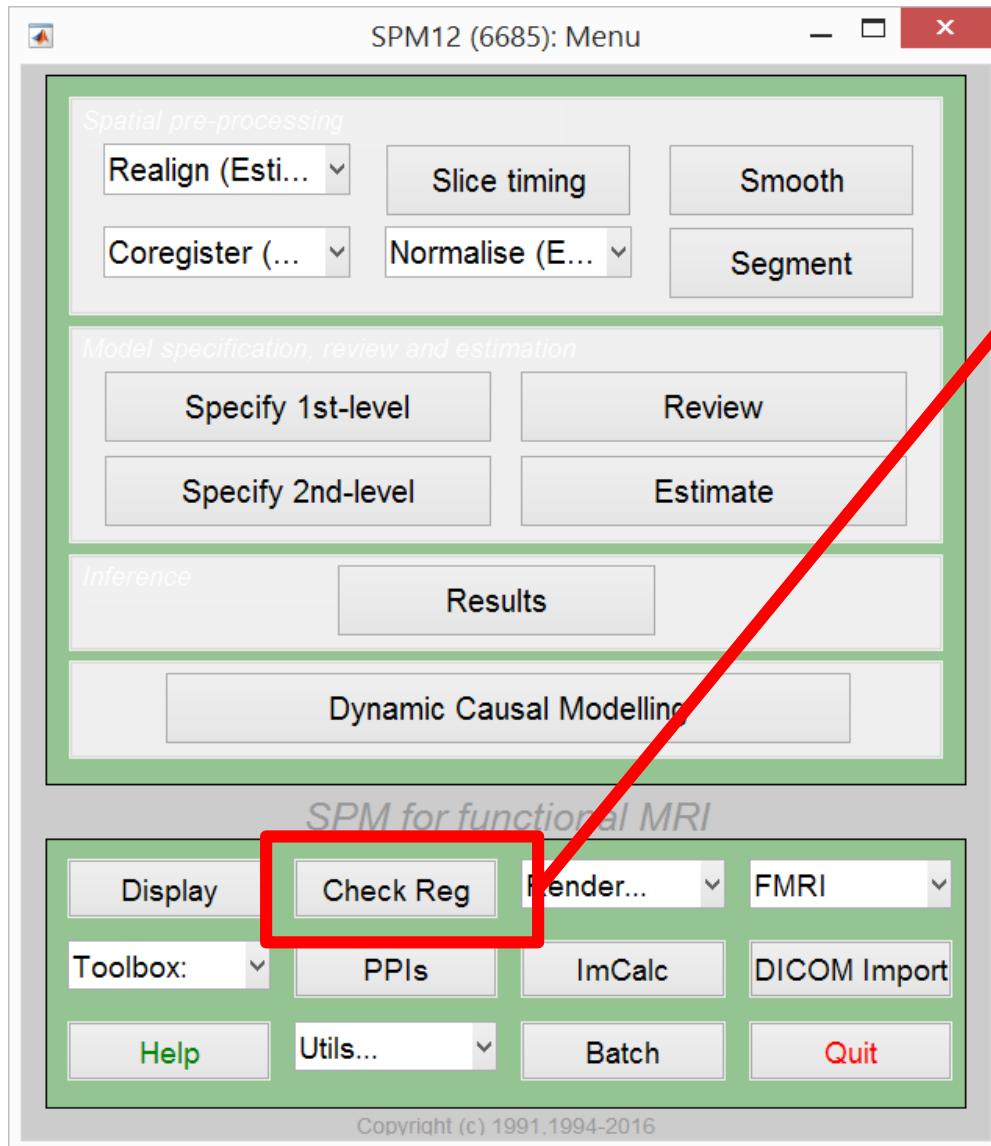
Additional useful
functions.

Display



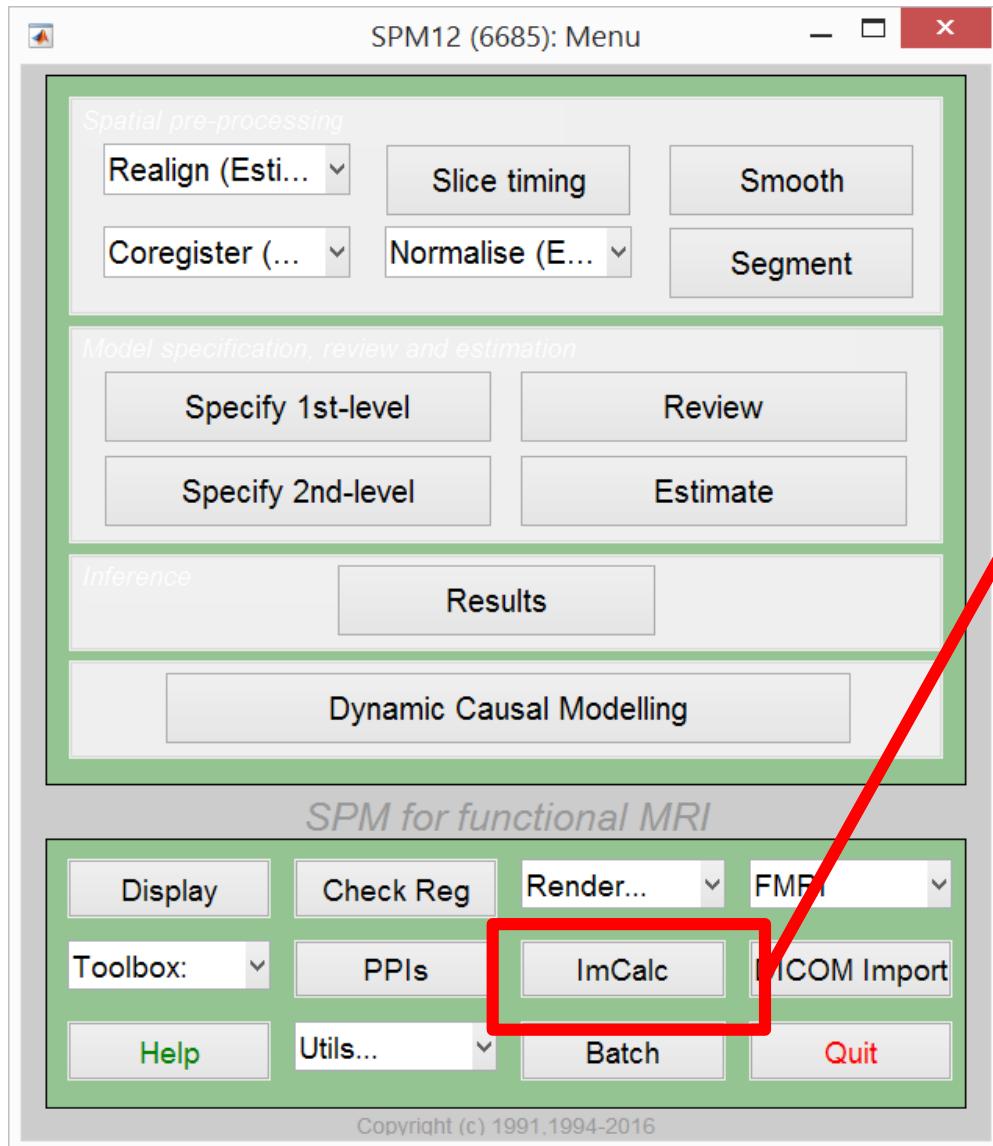
Display one image.

Check Reg



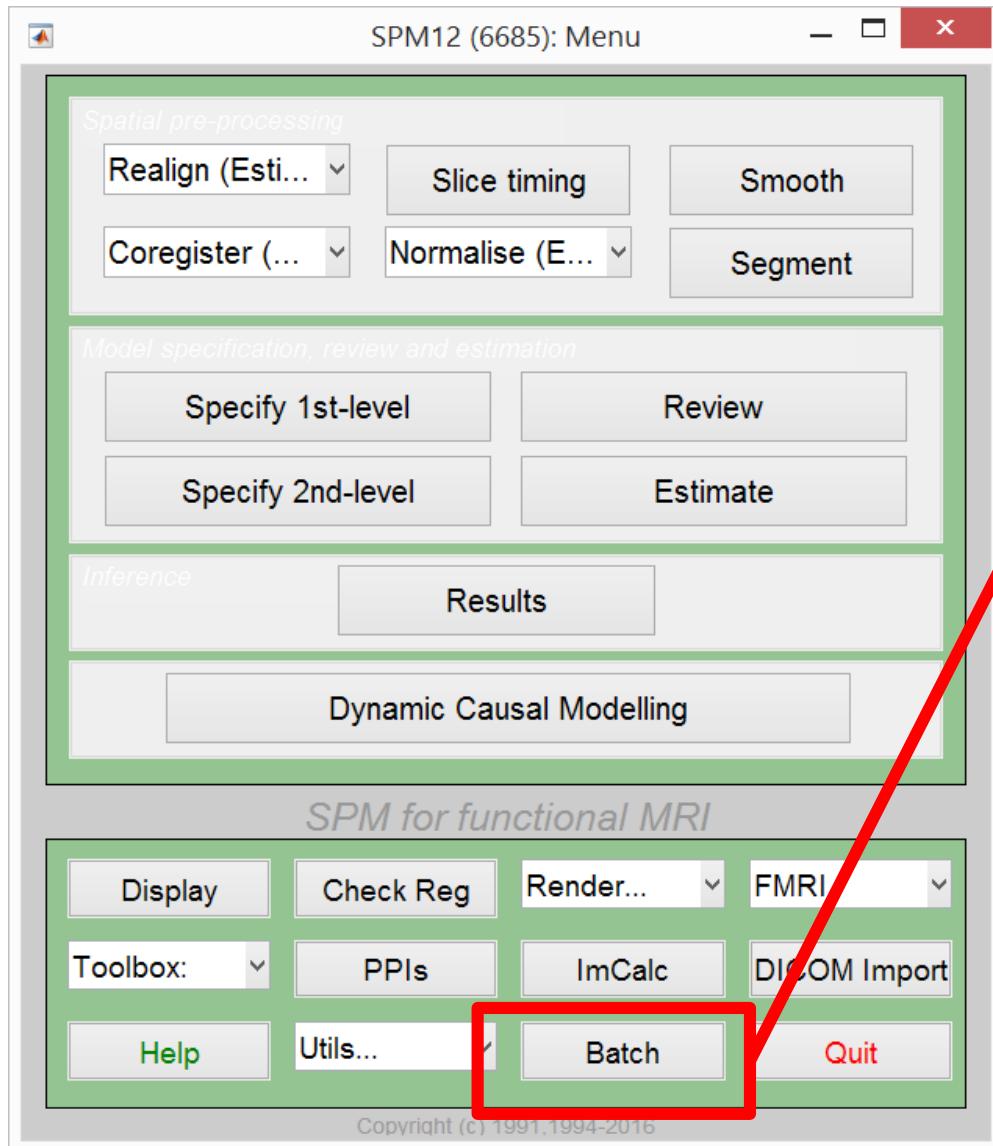
Display several images for comparison.

ImCalc



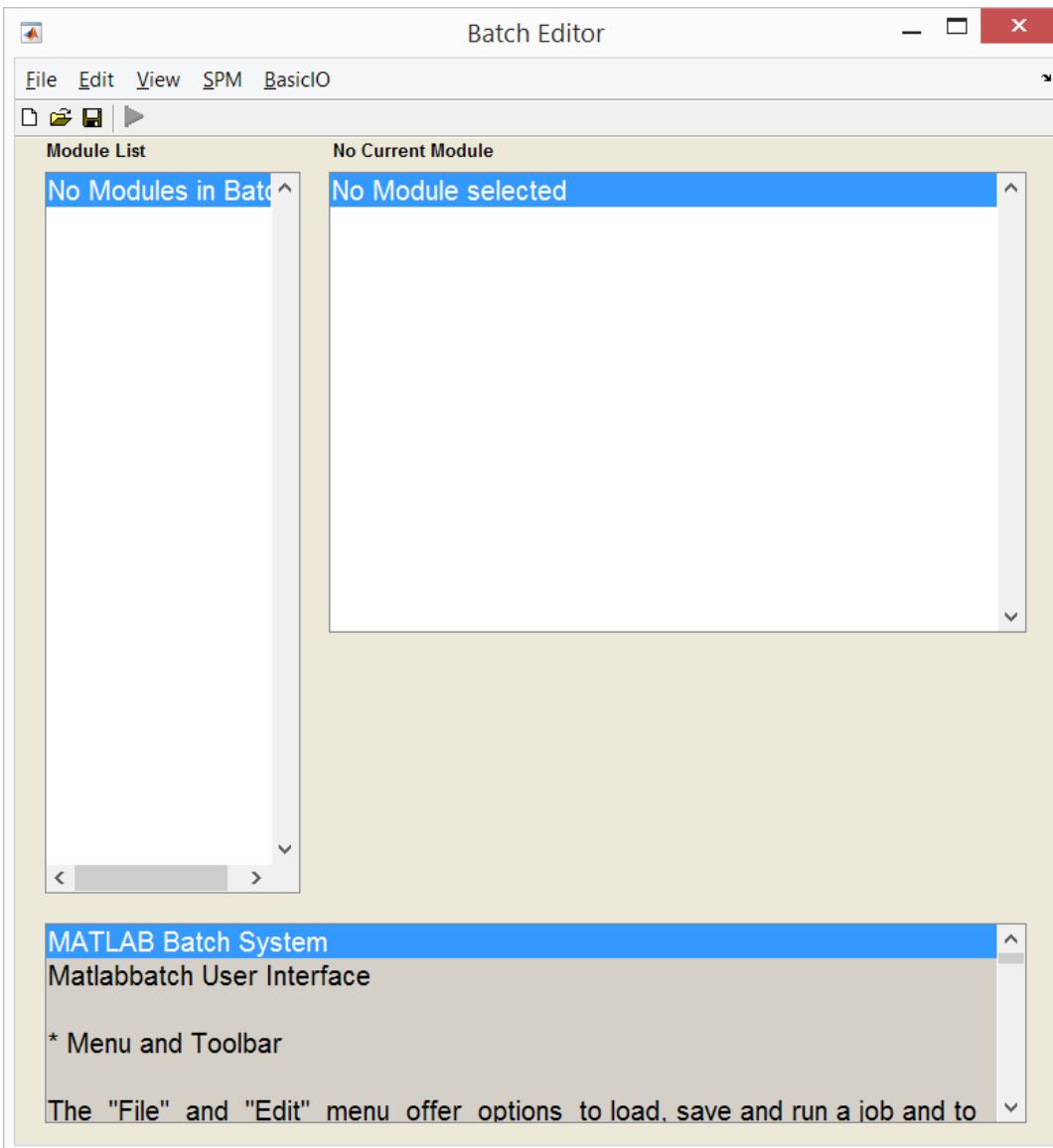
Do simple calculations with one or several images.
Example: Threshold anatomy ...

ImCalc

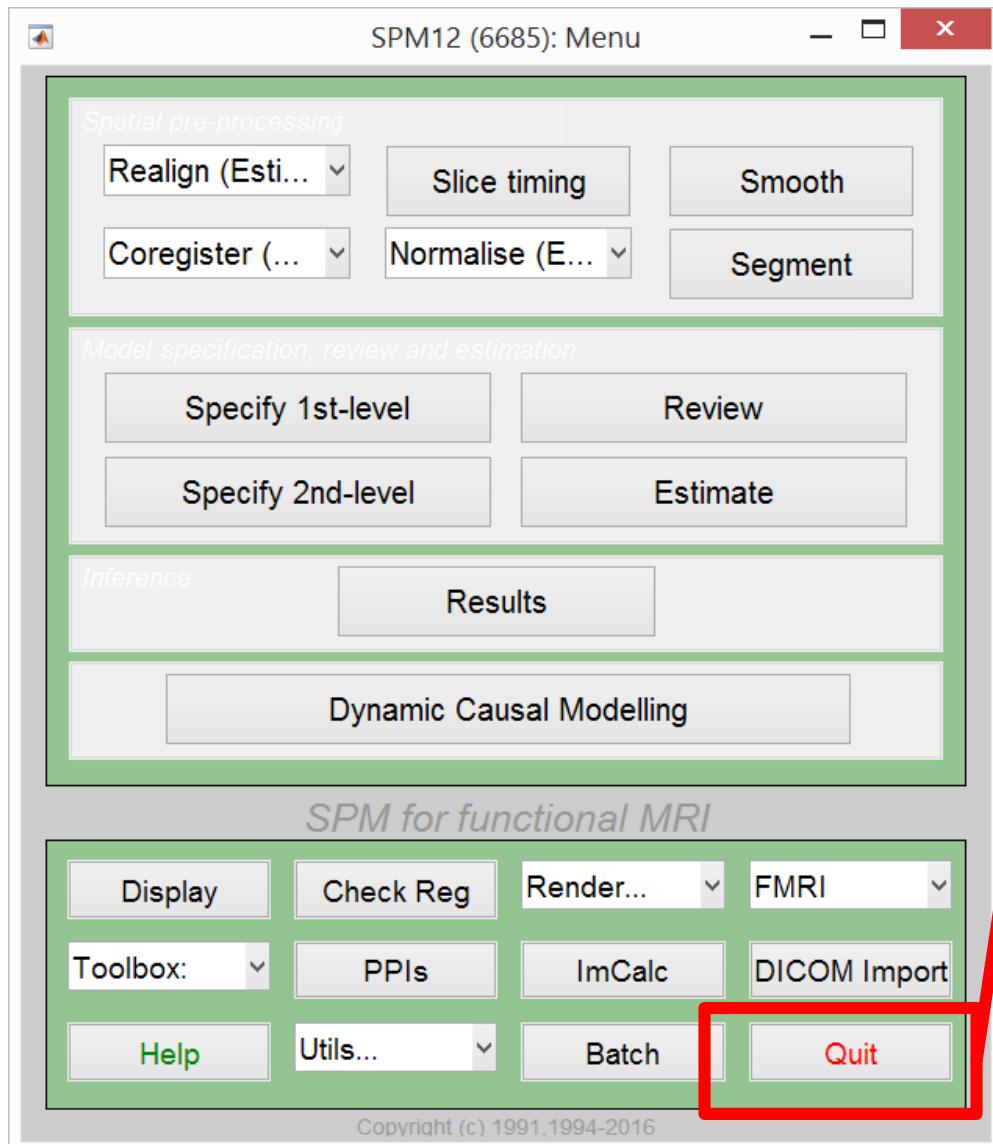


Start the batch editor. Often the batch editor also starts, if one clicks on one of the other buttons.

The batch editor



Quitting spm



Exit spm.

A simple example – thresholding anatomy

The screenshot shows the SPM Batch Editor interface with the 'Image Calculator' module selected in the 'Module List'. The 'Current Module: Image Calculator' panel displays the following settings:

- Help on: Image Calculator
- Input Images: ...2016\face_rep\Structural\sM03953_0007.img,1
- Output Filename: GreyFromThreshold
- Expression: $(i1 > 75) \& (i1 < 105)$
- Additional Variables: None
- Options:
 - Data Matrix: No - don't read images into data matrix
 - Masking: No implicit zero mask
 - Interpolation: Trilinear
 - Data Type: INT16 - signed short

The 'Current Item: Expression' panel shows the expression $(i1 > 75) \& (i1 < 105)$. Below it is a 'Specify...' button.

Expression
Example expressions (f):

- * Mean of six images (select six images)
 $f = '(i1+i2+i3+i4+i5+i6)/6'$
- * Make a binary mask image at threshold of 100
 $f = 'i1 < 100'$

Preparation for tutorials (if you want to follow on your own laptop, optional)

- Install matlab via your university.
- Download spm (<http://www.fil.ion.ucl.ac.uk/spm/software/spm12/>) and add to Matlab path.
- Download dataset face_rep.zip from http://www.fil.ion.ucl.ac.uk/spm/data/face_rep/
- Download face_rep_spm12_batch.m from the same source.
- Extract the data in a folder on your computer. Within the face_rep folder create a folder called «batches».