

# Getting Started with FSL

Richard Yaxley

Brain Imaging and Analysis Center  
Duke University

December 9, 2008

## Introduction

Tools

Accessing Data

## Preprocessing

Preparing Images for Analysis

Specifying Events for Analysis

## FEAT

Prestats

# FMRIB Software Library (FSL)



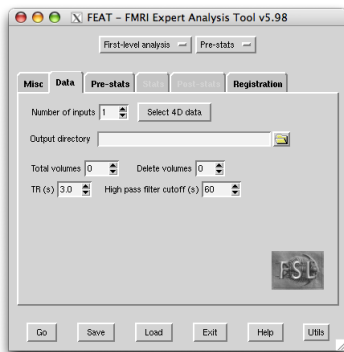
▶ FSL

- ▶ <http://www.fmrib.ox.ac.uk/fsl/>
- ▶ <http://www.fmrib.ox.ac.uk/fsl/feat5/>

▶ BIAC Wiki

- ▶ <http://fourier.biac.duke.edu/wiki/doku.php/biac:fsl:guide>

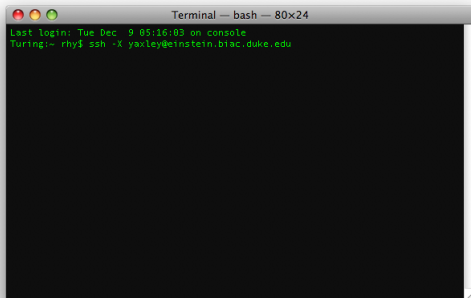
# FMRI Expert Analysis Tool (FEAT)



*“a software tool for high quality model-based FMRI data analysis, with an easy-to-use graphical user interface (GUI).”*

## Connect to Cluster

- ▶ Secure Shell (SSH) via Terminal in OS X, F-Secure in XP



A screenshot of a macOS Terminal window. The title bar reads "Terminal - bash - 80x24". The terminal content shows a successful SSH connection to a remote host. The first line is "Last login: Tue Dec 9 05:16:03 on console". The second line shows the prompt "Turing:~ rhy\$" followed by the command "ssh -X yaxley@beinstein.biac.duke.edu" which has been executed.

```
Terminal - bash - 80x24
Last login: Tue Dec 9 05:16:03 on console
Turing:~ rhy$ ssh -X yaxley@beinstein.biac.duke.edu
```

## Access Data

- ▶ Login to Einstein:  
`$ ssh -X Username@einstein.biac.duke.edu`
- ▶ Login to Node4:  
`$ qinteract`
- ▶ Mount experiment directories:  
`$ lnexp Exp.01`
- ▶ Navigate to experiment:  
`$ cd ~/experiments/Exp.01`

## Reorient Raw Data

- ▶ BIAC images must be reoriented from LPS (Left,Posterior,Superior) to LAS (Left,Anterior,Superior)
- ▶ Functional:  

```
$ bxhreorient --orientation LAS run004.bxh  
reoriented_run01.bxh
```
- ▶ Anatomical:  

```
$ bxhreorient --orientation LAS series002.bxh  
reoriented_anat.bxh
```
- ▶ Repeat for every functional image

## Generate NIFTI-formatted Images

▶ Functional:

```
$ bxh2analyze --nifti -s -v run001_01.bxh run01
```

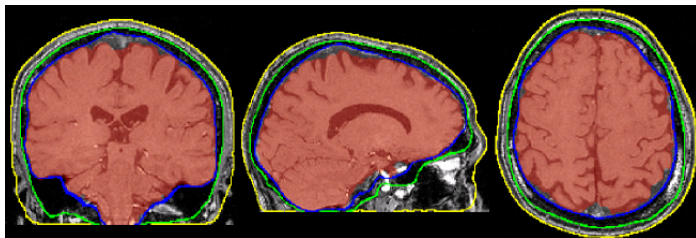
- ▶ This will create one large file (run01.img) that contains the functional data in 4D format and a NIFTI-formatted header (run01.hdr)

▶ Anatomical:

```
$ bxh2analyze --niftihdr -s -v series003.bxh anat
```

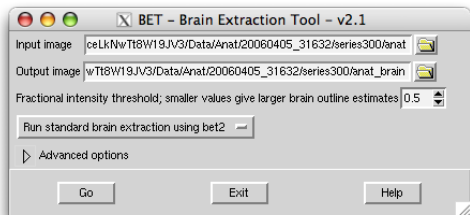


## Brain Extraction Tool



- ▶ BET deletes non-brain tissue from an image of the whole head
- ▶ <http://www.fmrib.ox.ac.uk/analysis/research/bet/>

# Generate Anatomical Brain Mask



- ▶ Use GUI or CLI to generate 'anat\_brain.nii.gz'
- ▶ Anatomical:
 

```
$ bet anat anat_brain -f 0.5 -g 0
```
- ▶ Functional: Perform later in 'Prestats'

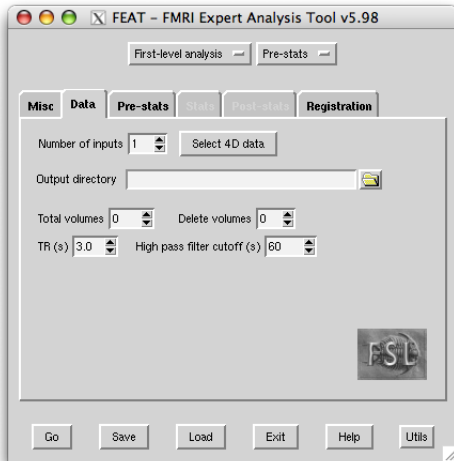
# Time-series of Events

- ▶ 3-column tab-delimited text files

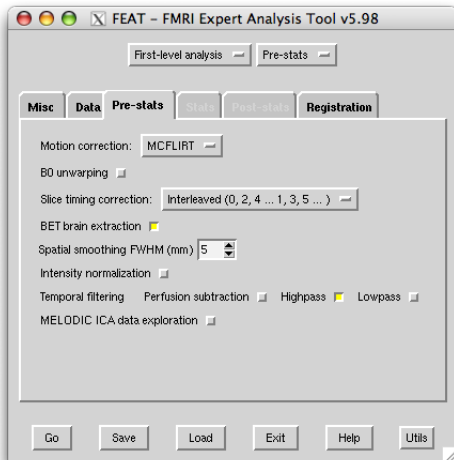
Onset	Duration	Weight
30	1	1
60	1	1
90	1	1
120	1	1

- ▶ One file for each condition of each run for each subject
- ▶ Example: 3 conditions X 6 runs X 20 subjects = 360 text files

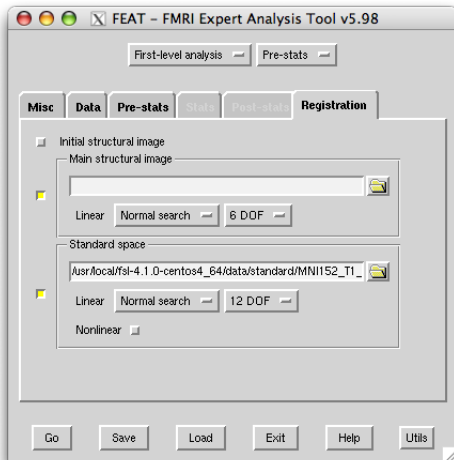
# Data



# Prestats



# Registration



# Output

- ▶ Output placed in `run01.feats/`

The screenshot shows a web browser window titled 'FSL' with the address bar containing the file path: `file:///Users/rhy/Documents/Demo.01/31700/run01/run01.feats/report_log.html`. The page content includes:

## FEAT Report

`/Users/rhy/Documents/Demo.01/31700/run01/run01.feats`  
 Started at Tue Dec 9 06:50:22 EST 2008 **STILL RUNNING.**

[Pre-stats](#) - [Stats](#) - [Post-stats](#) - [Registration](#) - [Log](#)

---

### Progress Report / Log

Started at Tue Dec 9 06:50:22 EST 2008

Feat main script

```

/bin/cp /tmp/feat_zY8T2p.fsf design.fsf
/usr/local/fsl/bin/feat_model design
mkdir .files;cp /usr/local/fsl/doc/fsl.css .files;ln -s /usr/local/fsl/doc/images .files/images
/usr/local/fsl/bin/fsl_sub -T 18 -l logs -N feat2_pre /usr/local/fsl/bin/feat /Users/rhy/Documents/Demo.01/31700/run01/ru
  
```

Prestats

## Conclusion

- ▶ Prepared the functional and anatomical images (reoriented, converted to NIFTI, and created brain mask from hi-res image)
- ▶ Specified event time-series for conditions
- ▶ Preprocessed and checked
- ▶ Now ready to setup model...