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Initializing spm8 Batch
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A command line interface to slice time correction procedures in FSL. This uses the FSL routines : slicetimer.

The 'sliceTime' command builds a script in the directory:

matlabScripts/spm8Batch/sliceTime/YYYY_MM directory

directly below the current directory and will launch this script in background. The names of the matlab and shell scripts will be in the format of \cdot

sliceTime_yymmdd_HH_MM_[USER]_[COMPUTER].m and sliceTime_yymmdd_HH_MM_[USER]_[COMPUTER].sh

where yymmdd is the date and HH_MM is the time.

This script is expecting to find the necessary components in the directory scruture of $\boldsymbol{:}$

Usage:

sliceTime [OPTIONS] [subject_1] [subject_2] ... [subject_N]

Options

Ke	y Parameter	Description
_	A	process all runs found in 'func' directory, default is unspecified.
_	d	enable debug flag.
_	D	enable super debug flag.
-	f sub-directory	set relative path functional directory, default=func
	i #1 - #2	set the start and end run numbers to scan for inclusively
+,		default is none.
-	M SUBJECTMASTER	set the directory to scan for subjects, default is Subjects/
-	t	enable testing flag, script will be built but not run.
-	U user-email	set the user name for email notification, default is "rcwelsh" of present shell.
-	v volume-wild	set the volume name wild-card, default "prun".
_	# RUN-NUMBER	add run # to list to process, default is none.

e.g.

To slice-time correct fmri volumes in runs 1,2,4 you would:

sliceTime -v prun -f func/MSIT -s img/ -n 10 -r 1 -r 2 -r 4 060402rw To realign fmri volumes in inclusive range of 1-10, do: sliceTime -v prun -f func/MSIT -s img/ -n 10 -i 1-10 060402rw Current command default configuration is:

Functional Path : func/ Sub-directory Volume Wildcard : prun fMRI TR FSLOUTPUTTYPE : 2 : NIFTI Number of runs to realign : 0 functional images path : func/ Subject directory : Subjects/

User : rcwelsh

MATLAB : /net/misc/matlab2007b/bin/matlab

SANDBOXHOST : dysthymia

SANDBOX : /dysthymia/sandbox/

SANDBOXPID

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