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Initializing spm8 Batch
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A command line command to drive batch processing to segment the 'SPGR' to the standard template. This code will use SPM8 batch mode.

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

matlabScripts/spm8Batch/segHiRes/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 :

segHiRes_yymmdd_HH_MM_[USER]_[COMPUTER].m and
segHiRes_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:

segHiRes [OPTIONS] [subject_1] [subject_2] ... [subject_N]

Options

]	Key Parameter	Description
•		
+s/,	-d	enable debug flag.
	-D	enable super debug flag.
	-f sub-directory	<pre>set relative path functional directory, default=func</pre>
	-h HiResName	<pre>name of high resolution file, default is 'ehtlspgr'</pre>
	-M SUBJECTMASTER	set the directory to scan for subjects, default is Subjects/
	-t	enable testing flag, script will be built but not run.
	-T TEMPLATENAME	<pre>set the template to use for normalization, you can specify relative to /net/dysthymia//spm8/template</pre>
	.	<pre>specify an absolute directory path. default is /net/dysthymia//spm8/templates//T1.nii</pre>
	-U user-email	set the user name for email notification, default is "rcwelsh" of present shell.
	-w sub-directory	<pre>set coregistration path under func, default=coReg</pre>

e.g.

To segment the 'whet1spgr' :

segHiRes -f func/MSIT 050126zz

Current command default configuration is:

directory to coreg
name of the high res
: func//coReg/
: ehtlspgr

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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