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   Initializing spm8 Batch


  A command line command to drive batch processing to warp the 'SPGR'
  to the standard template.


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

        matlabScripts/spm8Batch/warpHiRes/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 :
          warpHiRes_yymmdd_HH_MM_[USER]_[COMPUTER].m    and
          warpHiRes_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 :

      yymmddLF/                              (subject directory)
              anatomy/
              func/
                  coReg/
                  run_XX/

   Usage:

        warpHiRes [OPTIONS] [subject_1] [subject_2] ... [subject_N]


   Options

     Key Parameter            Description
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       -d                     enable debug flag.

       -D                     enable super debug flag.

       -f sub-directory       set relative path functional directory,
                              default=func

       -h HiResName           name of high resolution file,
                              default is 'eht1spgr'

       -M SUBJECTMASTER       set the directory to scan for subjects,
                              default is Subjects/

       -n PrependName         name to add to output file name,
                              default is none.

       -t                     enable testing flag, script will be built
                              but not run.

       -T TEMPLATENAME        set the template to use for normalization,
                              you can specify relative to /net/dysthymia//spm8/template
+s/, or.
                              specify an absolute directory path.
                              default is /net/dysthymia//spm8/templates//T1.nii

       -U user-email          set the user name for email notification,
                              default is "rcwelsh" of present shell.

       -w sub-directory       set coregistration path under func,
                              default=coReg

       -z VOXELSIZE           set the voxel size for resampling,
                              default is spm default.
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e.g.

   To warp the 'het1spgr.img' to the standard template:

      warpHiRes -p func/MSIT 050126zz


Current command default configuration is:


directory to coreg      : func//coReg//
name of the high res     : eht1spgr
template image dir/name   : /net/dysthymia//spm8/templates//T1.nii
prepending name        : w
Voxel dimension (0=spm)   : 0

functional images path    : func/
Subject directory       : Subjects/

spm8 is located in      : /net/dysthymia//spm8
spm8Batch is located in   : /net/dysthymia//spm8Batch
spm8 patch is located in : /net/dysthymia//spm8Batch/matlabScripts
auxillary matlab path    : /net/dysthymia//spm8Batch/spm8_patch

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

SANDBOXHOST        : dysthymia
SANDBOX          : /dysthymia/sandbox/
SANDBOXPID        :

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