

ACAP Preprocessing Work Flow

1) Run batchBIDSdcm.m

- Dependencies: BIDSdcm2nii.m, dcm2nii package
- Update batchBIDSdcm.m SOURCE_DIR, TAR_DIR and PAR_??? as required
- ???_LIST is an $n \times 4$ cell array with column number:
 - Regex expression to find dicom files
 - BIDS subfolder where the modality is stores (e.g. 'func', 'anat')
 - BIDS name of nifti file in printf format
 - BIDS name of nifti file in printf format if multiple runs exist

and the n rows correspond to n separate modalities e.g.:

(.*)ep2d(.*)Resting_State(.*)	func	sub-%s_ses-%s_task-resting_bold	sub-%s_ses-%s_task-resting_run-%02d_bold
(.*)_mprage_(.*)iso0.8\$	anat	sub-%s_ses-%s_T1w	sub-%s_ses-%s_run-%02d_T1w

- Switch to dcm2niix?

2) Run BOLDpreprocess.sh

- Dependencies: BOLDpreprocess.py, optiBET.sh
- Usage:

```
alexbarton$ bash BOLDpreprocess.sh [BIDS-ROOT-DIRECTORY] [#-OF-PARALLEL-PROCESSES] [optiBET.sh-LOCATION]
```

3) Manually Check T1's and make .csv list

- RepeatList.csv is an $n \times 3$.csv with column number:
 - Subject Name
 - Session Name
 - Run # to use

and the n rows being the number of subjects with repeat scans e.g.:

ACAP1215	B0	1
ACAP1217	B0	1
ACAP2136	M6	2
ACAP2137	M6	2
ACAP2140	M6	2
ACAP2142	M6	2
ACAP2143	M6	4

4) Run icaPrep.py

- a. Makes the .fsf files for each participant based on a template
- b. All fields that are replaced are strings in the form of
`XXXFIELDNAMEXXX` (e.g. XXXNVOLSXXX)
- c. Usage:

```
alexbarton$ python icaPrep.py [BIDS-ROOT-DIRECTORY] -t [T1-REPEATS.csv] -f [FSF-TEMPLATE-FILE.fsf]
```

5) Make list of .fsf files

- a. I use:

```
alexbarton$ ls -ld [/PATH/TO/FILES/.../sub*/ses*/*.fsf] > [FSF_LIST.txt]
```

6) Run:

```
alexbarton$ parallel -j[#-OF-PARALLEL-PROCESSES] feat :::: [FSF_LIST.txt]
```

Python packages:

- os
- sys
- argparse
- shutil
- csv
- nibabel
- re
- numpy
- glob
- nipytype (?unsure if needed?)