# **BIDScoin**

Release 3.0.6

Feb 04, 2021

# Contents

3

## 1 BIDScoin functionality

2	Note:		5
	2.1	Installati	on
		2.1.1	Dcm2niix installation
		2.1.2	Python 3 installation
		2.1.3	BIDScoin installation
	2.2	Data pre	paration
		2.2.1	Required source data structure
		2.2.2	Data management utilities
	2.3	The BID	Scoin workflow
		2.3.1	Step 1a: Running the bidsmapper
		2.3.2	Step 1b: Running the bidseditor 12
		2.3.3	Step 2: Running the bidscoiner
	2.4	Finishin	g up
		2.4.1	Adding meta-data
		2.4.2	Data sharing utilities
		2.4.3	BIDS validation
	2.5	Options	
		2.5.1	BIDScoin
		2.5.2	dcm2niix
		2.5.3	Plugins
	2.6 Advanced usage		d usage
		2.6.1	Site specific / customized template
		2.6.2	Plugins
	2.7	7 Screenshots	
		2.7.1	The bidseditor
	2.8	Demo ar	nd tutorial
		2.8.1	BIDS introduction and BIDScoin demo 24
		2.8.2	Hands-on futorial 24



BIDScoin is a user friendly open-source python toolkit that converts ("coins") source-level (raw) neuroimaging datasets to nifti / json / tsv data-sets that are organized following the Brain Imaging Data Structure, a.k.a. the BIDS standard. Rather then depending on complex or ambiguous programmatic logic for the identification of imaging modalities, BIDScoin uses a direct mapping approach to identify and convert the raw source data into BIDS data. Different runs of source data are identified by reading information from MRI header files (DICOM or PAR/REC; e.g. 'ProtocolName') and the mapping information about how these different runs should be named in BIDS can be specified a priori as well as interactively by the researcher – bringing in the missing knowledge that often exists only in his or her head!

Because all the mapping information can be easily edited with a Graphical User Interface (GUI), BIDScoin requires no programming knowledge in order to use it.

BIDScoin is developed at the Donders Institute of the Radboud University.

# CHAPTER 1

# **BIDScoin functionality**

- [x] DICOM source data
- [x] PAR / REC source data (Philips)
- [] P7 source data (GE)
- [] Nifti source data
- [x] Fieldmaps\*
- [x] Multi-echo data\*
- [x] Multi-coil data\*
- [x] PET data\*
- [] Stimulus / behavioural logfiles
- [] Physiological data
- [x] Plug-ins
- [x] Defacing
- [x] Multi-echo combination
  - \* = Experimental support for PAR / REC source data

```
Are you a python programmer with an interest in BIDS who knows all about GE and / or_

→Philips data?

Are you experienced with parsing stimulus presentation log-files? Or do you have_

→ideas to improve

the this toolkit or its documentation? Have you come across bugs? Then you are highly_

→encouraged to

provide feedback or contribute to this project on https://github.com/Donders-

→Institute/bidscoin.
```

# CHAPTER 2

Note:

#### The full BIDScoin documentation is hosted at Read the Docs

Issues can be reported at Github

## 2.1 Installation

BIDScoin can be installed and should work on Linux, MS Windows and on OS-X computers (this latter option has not been tested) that satisfy the system requirements:

- dcm2niix
- python 3.6 or higher

#### 2.1.1 Dcm2niix installation

BIDScoin relies on dcm2niix to convert the source imaging data to nifti. Please download and install dcm2niix yourself according to the instructions. When done, make sure that the path to the dcm2niix binary / executable is set correctly in the BIDScoin Options in the [path\_to\_bidscoin]/heuristics/bidsmap\_template.yaml file or in the Site specific / customized template file.

#### 2.1.2 Python 3 installation

BIDScoin is a python package and therefore a python interpreter needs to be present on the system. On Linux this is usually already the case, but MS Windows users may need to install python themselves. See e.g. this python 3 distribution for instructions. They may also need to install the MS Visual C++ build tools (sorry for this pain).

#### 2.1.3 BIDScoin installation

To install BIDScoin run the following command in your command-shell (institute users may want to activate a virtual / conda python environment first):

```
$ pip install bidscoin
```

This will give you the latest stable release of the software. To get the very latest (development) version of the software you can install the package directly from the github source code repository:

\$ pip install git+https://github.com/Donders-Institute/bidscoin

If you want to edit the code or want to contribute back to the project, you can use the -e option:

\$ pip install -e git+https://github.com/Donders-Institute/bidscoin#egg=bidscoin

If you do not have git (or any other version control system) installed you can download and unzip the code yourself in a directory named e.g. bidscoin and run (again, with or without the -e option):

\$ pip install -e bidscoin

#### **Updating BIDScoin**

Run the pip command as before with the additional --upgrade option:

\$ pip install --upgrade bidscoin

#### **Caution:**

- The bidsmaps are not garanteed to be compatible between different BIDScoin versions.
- After a succesful BIDScoin installation or upgrade, it may be needed to (re)do any adjustments that were done on the Site specific / customized template file(s) (so make a back-up of these before you upgrade).

## 2.2 Data preparation

#### 2.2.1 Required source data structure

BIDScoin requires that the source data input folder is organized according to a sub-identifier/ [ses-identifier]/sessiondata structure (NB: the ses-identifier subfolder is optional). The sessiondata can have various formats, as shown in the following examples:

1. A 'seriesfolder' organization. A series folder contains a single data type and are typically acquired in a single run – a.k.a 'Series' in DICOM speak. This is how users receive their data from the (Siemens) scanners at the DCCN:

```
sourcedata
|-- sub-001
   |-- ses-mri01
      |-- 001-localizer
   |-- 00001_1.3.12.2.1107.5.2.19.45416.2017121914582956872274162.IMA
   |-- 00002_1.3.12.2.1107.5.2.19.45416.2017121914583757650874172.IMA
`-- 00003_1.3.12.2.1107.5.2.19.45416.2017121914583358068374167.IMA
|-- 002-t1_mprage_sag_p2_iso_1.0
|-- 00002_1.3.12.2.1107.5.2.19.45416.2017121915051526005675150.IMA
```

```
|-- 00003_1.3.12.2.1107.5.2.19.45416.2017121915051520026075138.IMA
       |-- 00004_1.3.12.2.1107.5.2.19.45416.2017121915051515689275130.IMA
   [..]
[..]
-- ses-mri02
|-- 001-localizer
| |-- 00001_1.3.12.2.1107.5.2.19.45416.2017121914582956872274162.IMA
|-- 00002_1.3.12.2.1107.5.2.19.45416.2017121914583757650874172.IMA
       1
`-- 00003_1.3.12.2.1107.5.2.19.45416.2017121914583358068374167.IMA
       1
[..]
|-- sub-002
   `-- ses-mri01
|-- 001-localizer
| |-- 00001_1.3.12.2.1107.5.2.19.45416.2017121914582956872274162.IMA
|-- 00002_1.3.12.2.1107.5.2.19.45416.2017121914583757650874172.IMA
`-- 00003_1.3.12.2.1107.5.2.19.45416.2017121914583358068374167.IMA
       [..]
[..]
```

2. A 'DICOMDIR' organization. A DICOMDIR is dictionary-file that indicates the various places where all the DICOM files are stored of each DICOM Series. This is how data is often exported in clinical settings:

```
sourcedata
|-- sub-001
  |-- DICOM
`-- 00001EE9
  `-- AAFC99B8
  `-- AA547EAB
   |-- 00000025
  | |-- EE008C45
| |-- EE027F55
| |-- EE03D17C
                | [..]
T
   |-- 000000B4
| |-- EE07CCDA
|-- EE0E0701
|-- EE0E200A
   [..]
[..]
`-- DICOMDIR
|-- sub-002
   [..]
[..]
```

3. A flat DICOM organization. In a flat DICOM organization all the DICOM files of all the different Series are simply put in one large directory. This is how data is sometimes exported in clinical settings:

```
sourcedata
|-- sub-001
| `-- ses-mri01
| |-- IM_0001.dcm
| |-- IM_0002.dcm
```

```
| |-- IM_0003.dcm
| [..]
|
|-- sub-002
| `-- ses-mri01
| |-- IM_0001.dcm
| |-- IM_0002.dcm
| |-- IM_0003.dcm
| [..]
[..]
```

4. A PAR/REC organization. All PAR/REC(/XML) files of all the different Series are put in one directory. This is how users often export their data from Philips scanners in research settings:

```
sourcedata
|-- sub-001
   `-- ses-mri01
       |-- TCHC_066_1_WIP_Hanneke_Block_2_SENSE_4_1.PAR
|-- TCHC_066_1_WIP_Hanneke_Block_2_SENSE_4_1.REC
|-- TCHC_066_1_WIP_IDED_SENSE_6_1.PAR
|-- TCHC_066_1_WIP_IDED_SENSE_6_1.REC
|-- TCHC_066_1_WIP_Localizer_CLEAR_1_1.PAR
        |-- TCHC_066_1_WIP_Localizer_CLEAR_1_1.REC
[..]
|-- sub-002
   `-- ses-mri01
|-- TCHC_066_1_WIP_Hanneke_Block_2_SENSE_4_1.PAR
       |-- TCHC_066_1_WIP_Hanneke_Block_2_SENSE_4_1.REC
       |-- TCHC_066_1_WIP_IDED_SENSE_6_1.PAR
       |-- TCHC_066_1_WIP_IDED_SENSE_6_1.REC
|-- TCHC_066_1_WIP_Localizer_CLEAR_1_1.PAR
|-- TCHC_066_1_WIP_Localizer_CLEAR_1_1.REC
[..]
[..]
```

**Note:** You can store the sessiondata in any of the above data organizations as zipped (.zip) or tarzipped (e.g. .tar.gz) archive files. BIDScoin workflow tools will unpack/unzip those archive files in a temporary folder and will process the sessiondata from there. The BIDScoin tools will run *dicomsort* in a temporary folder for flat/DICOMDIR data to sort them in seriesfolders. BIDScoin tools that work from a temporary folder has the downsde of getting a speed penalty. Also note that privacy-sensitive data samples will then be stored in [bidsfolder]/ code/bidscoin/provenance.

## 2.2.2 Data management utilities

#### dicomsort

The dicomsort command-line tool is a utility to move your flat- or DICOMDIR-organized files (see *above*) into a 'seriesfolder' organization (see above). This can be useful to organise your source data in a more convenient and human readable way, as DICOMDIR or flat DICOM directories can often be hard to comprehend. The BIDScoin tools will run dicomsort in a temporary folder if your data is not already organised in series-folders, so in principle you don't really need to run it yourself. Running dicomsort beforehand does, however, give you more flexibility in handling special cases that are not handled properly and it can also give you a speed benefit.

```
usage: dicomsort [-h] [-i SUBPREFIX] [-j SESPREFIX] [-f FIELDNAME] [-r]
                 [-e EXT] [-n] [-p PATTERN] [-d]
                 dicomsource
Sorts and / or renames DICOM files into local subdirectories with a (3-digit)
SeriesNumber-SeriesDescription directory name (i.e. following the same listing
as on the scanner console)
positional arguments:
 dicomsource
                        The name of the root folder containing the
                        dicomsource/[sub/][ses/]dicomfiles and / or the
                        (single session/study) DICOMDIR file
optional arguments:
                        show this help message and exit
 -h, --help
 -i SUBPREFIX, --subprefix SUBPREFIX
                        Provide a prefix string for recursive searching in
                        dicomsource/subject subfolders (e.g. "sub") (default:
                        None)
 -j SESPREFIX, --sesprefix SESPREFIX
                        Provide a prefix string for recursive searching in
                        dicomsource/subject/session subfolders (e.g. "ses")
                        (default: None)
 -f FIELDNAME, --fieldname FIELDNAME
                        The dicomfield that is used to construct the series
                        folder name ("SeriesDescription" and "ProtocolName"
                        are both used as fallback) (default:
                        SeriesDescription)
 -r, --rename
                        Flag to rename the DICOM files to a PatientName_Series
                        Number_SeriesDescription_AcquisitionNumber_InstanceNum
                        ber scheme (recommended for DICOMDIR data) (default:
                        False)
 -e EXT, --ext EXT
                       The file extension after sorting (empty value keeps
                        the original file extension), e.g. ".dcm" (default: )
 -n, --nosort
                        Flag to skip sorting of DICOM files into SeriesNumber-
                        SeriesDescription directories (useful in combination
                        with -r for renaming only) (default: False)
 -p PATTERN, --pattern PATTERN
                        The regular expression pattern used in
                        re.match(pattern, dicomfile) to select the dicom files
                        (default: .*\.(IMA|dcm)$)
 -d, --dryrun
                        Add this flag to just print the dicomsort commands
                        without actually doing anything (default: False)
examples:
 dicomsort /project/3022026.01/raw
 dicomsort /project/3022026.01/raw --subprefix sub
 dicomsort /project/3022026.01/raw --subprefix sub-01 --sesprefix ses
 dicomsort /project/3022026.01/raw/sub-011/ses-mri01/DICOMDIR -r -e .dcm
```

#### rawmapper

Another command-line utility that can be helpful in organizing your source data is rawmapper. This utility can show you the overview (map) of all the values of DICOM-fields of interest in your data-set and, optionally, use these fields to rename your source data sub-folders (this can be handy e.g. if you manually entered subject-identifiers as [Additional info] at the scanner console and you want to use these to rename your subject folders).

```
usage: rawmapper [-h] [-s SESSIONS [SESSIONS ...]]
                 [-d DICOMFIELD [DICOMFIELD ...]] [-w WILDCARD]
                 [-o OUTFOLDER] [-r] [-n SUBPREFIX] [-m SESPREFIX]
                 [--dryrun]
                 sourcefolder
Maps out the values of a dicom field of all subjects in the sourcefolder, saves
the result in a mapper-file and, optionally, uses the dicom values to rename
the sub-/ses-id's of the subfolders. This latter option can be used, e.g.
when an alternative subject id was entered in the [Additional info] field
during subject registration (i.e. stored in the PatientComments dicom field)
positional arguments:
 sourcefolder
                        The source folder with the raw data in
                   sub-#/ses-#/series organisation
optional arguments:
                        show this help message and exit
 -h, --help
 -s SESSIONS [SESSIONS ...], --sessions SESSIONS [SESSIONS ...]
                    Space separated list of selected sub-#/ses-# names /
                    folders to be processed. Otherwise all sessions in the
                   bidsfolder will be selected (default: None)
 -d DICOMFIELD [DICOMFIELD ...], --dicomfield DICOMFIELD [DICOMFIELD ...]
                    The name of the dicomfield that is mapped / used to
                    rename the subid/sesid foldernames (default:
                    ['PatientComments'])
 -w WILDCARD, --wildcard WILDCARD
                    The Unix style pathname pattern expansion that is used
                    to select the series from which the dicomfield is
                   being mapped (can contain wildcards) (default: *)
 -o OUTFOLDER, --outfolder OUTFOLDER
                    The mapper-file is normally saved in sourcefolder or,
                    when using this option, in outfolder (default: None)
                       If this flag is given sub-subid/ses-sesid directories
 -r, --rename
                    in the sourcefolder will be renamed to sub-dcmval/ses-
                    dcmval (default: False)
 -n SUBPREFIX, --subprefix SUBPREFIX
                    The prefix common for all the source subject-folders
                    (default: sub-)
 -m SESPREFIX, --sesprefix SESPREFIX
                    The prefix common for all the source session-folders
                    (default: ses-)
                        Add this flag to dryrun (test) the mapping or renaming
 --dryrun
                    of the sub-subid/ses-sesid directories (i.e. nothing
                    is stored on disk and directory names are not actually
                    changed)) (default: False)
examples:
 rawmapper /project/3022026.01/raw/
 rawmapper /project/3022026.01/raw -d AcquisitionDate
 rawmapper /project/3022026.01/raw -s sub-100/ses-mri01 sub-126/ses-mri01
 rawmapper /project/3022026.01/raw -r -d ManufacturerModelName AcquisitionDate --
→dryrun
 rawmapper raw/ -r -s sub-1*/* sub-2*/ses-mri01 --dryrun
 rawmapper -d EchoTime -w *fMRI* /project/3022026.01/raw
```

Note: If these data management utilities do not satisfy your needs, then have a look at this reorganize\_dicom\_files

tool.

# 2.3 The BIDScoin workflow

Having an organized source data folder, the actual data-set conversion to BIDS is performed by the (1a) the bidsmapper, (1b) the bidseditor and (2) the bidscoiner command-line tools. The bidsmapper makes a map of the different kind of datatypes in your source dataset, with the bidseditor you can edit this map, and the bidscoiner does the actual work to convert the source data into BIDS. By default (but see the -i option of the bidsmapper below), step 1a automatically launches step 1b, so in it's simplest form, all you need to do to convert your raw source data into BIDS is to run two simple commands, e.g.:

```
$ bidsmapper sourcefolder bidsfolder
$ bidscoiner sourcefolder bidsfolder
```

## 2.3.1 Step 1a: Running the bidsmapper

```
usage: bidsmapper [-h] [-b BIDSMAP] [-t TEMPLATE] [-n SUBPREFIX]
                  [-m SESPREFIX] [-i {0,1,2}] [-v]
                  sourcefolder bidsfolder
Creates a bidsmap.yaml YAML file in the bidsfolder/code/bidscoin that maps the_
→information
from all raw source data to the BIDS labels. You can check and edit the bidsmap file,
⇔with
the bidseditor (but also with any text-editor) before passing it to the bidscoiner...
\rightarrowSee the
bidseditor help for more information and useful tips for running the bidsmapper in,
⇒ interactive
mode (which is the default).
N.B.: Institute users may want to use a site-customized template bidsmap (see the
--template option). The bidsmap_dccn template from the Donders Institute can serve as
an example (or may even mostly work for other institutes out of the box).
positional arguments:
 sourcefolder
                        The study root folder containing the raw data in
                        sub-#/[ses-#/]data subfolders (or specify --subprefix
                        and --sesprefix for different prefixes)
 bidsfolder
                        The destination folder with the (future) bids data and
                        the bidsfolder/code/bidscoin/bidsmap.yaml output file
optional arguments:
                        show this help message and exit
 -h, --help
 -b BIDSMAP, --bidsmap BIDSMAP
                        The bidsmap YAML-file with the study heuristics. If
                        the bidsmap filename is relative (i.e. no "/" in the
                        name) then it is assumed to be located in
                        bidsfolder/code/bidscoin. Default: bidsmap.yaml
 -t TEMPLATE, --template TEMPLATE
                        The bidsmap template with the default heuristics (this
                        could be provided by your institute). If the bidsmap
                        filename is relative (i.e. no "/" in the name) then it
```

```
is assumed to be located in bidsfolder/code/bidscoin.
                        Default: bidsmap_template.yaml
 -n SUBPREFIX, --subprefix SUBPREFIX
                        The prefix common for all the source subject-folders.
                        Default: 'sub-'
 -m SESPREFIX, --sesprefix SESPREFIX
                        The prefix common for all the source session-folders.
                        Default: 'ses-'
 -s, --store
                        Flag to store the provenance data samples in the
                        bidsfolder/'code'/'provenance' folder
 -i {0,1,2}, --interactive {0,1,2}
                        {0}: The sourcefolder is scanned for different kinds
                        of scans without any user interaction. {1}: The
                        sourcefolder is scanned for different kinds of scans
                        and, when finished, the resulting bidsmap is opened
                        using the bidseditor. {2}: As {1}, except that already
                        during scanning the user is asked for help if a new
                        and unknown run is encountered. This option is most
                        useful when re-running the bidsmapper (e.g. when the
                        scan protocol was changed since last running the
                        bidsmapper). Default: 1
 -v, --version
                        Show the BIDS and BIDScoin version
examples:
 bidsmapper /project/foo/raw /project/foo/bids
 bidsmapper /project/foo/raw /project/foo/bids -t bidsmap_dccn
```

The bidsmapper will scan your sourcefolder to look for different runs (scan-types) to create a mapping for each run to a bids output name (a.k.a. the 'bidsmap'). By default (but see the -i option above), when finished the bidsmapper will automatically launch *step 1b*, as described in the next section (but step 1b can also always be run separately by directly running the bidseditor).

Tip: Use the -t bidsmap\_dccn option and see if it works for you. If not, consider adapting it to your needs.

### 2.3.2 Step 1b: Running the bidseditor

```
usage: bidseditor [-h] [-b BIDSMAP] [-t TEMPLATE] [-d DATAFORMAT]
                  [-n SUBPREFIX] [-m SESPREFIX]
                  bidsfolder
This tool launches a graphical user interface for editing the bidsmap.yaml file
that is e.g. produced by the bidsmapper or by this bidseditor itself. The user can
fill in or change the BIDS labels for entries that are unidentified or sub-optimal,
such that meaningful BIDS output names will be generated from these labels. The saved
bidsmap.yaml output file can be used for converting the source data to BIDS using
the bidscoiner.
positional arguments:
 bidsfolder
                   The destination folder with the (future) bids data
optional arguments:
 -h, --help
                    show this help message and exit
 -b BIDSMAP, --bidsmap BIDSMAP
```

```
The bidsmap YAML-file with the study heuristics. If
                    the bidsmap filename is relative (i.e. no "/" in the
                    name) then it is assumed to be located in
                   bidsfolder/code/bidscoin. Default: bidsmap.yaml
 -t TEMPLATE, --template TEMPLATE
                    The bidsmap template with the default heuristics (this
                    could be provided by your institute). If the bidsmap
                    filename is relative (i.e. no "/" in the name) then it
                    is assumed to be located in bidsfolder/code/bidscoin.
                   Default: bidsmap_template.yaml
 -d DATAFORMAT, --dataformat DATAFORMAT
                   The format of the source data, e.g. DICOM or PAR.
                   Default: DICOM
 -n SUBPREFIX, --subprefix SUBPREFIX
                   The prefix common for all the source subject-folders.
                   Default: 'sub-'
 -m SESPREFIX, --sesprefix SESPREFIX
                    The prefix common for all the source session-folders.
                    Default: 'ses-'
examples:
 bidseditor /project/foo/bids
 bidseditor /project/foo/bids -t bidsmap_dccn.yaml
 bidseditor /project/foo/bids -b my/custom/bidsmap.yaml
Here are a few tips & tricks:
_____
DICOM Attributes
 An (DICOM) attribute label can also be a list, in which case the BIDS labels /_
→mapping
 are applied if a (DICOM) attribute value is in this list. If the attribute value is
 empty it is not used to identify the run. Wildcards can also be given, either as a
⇔single
  '*', or enclosed by '*'. Examples:
      SequenceName: '*'
       SequenceName: '*epfid*'
       SequenceName: ['epfid2d1rs', 'fm2d2r']
       SequenceName: ['*epfid*', 'fm2d2r']
  NB: Editing the DICOM attributes is normally not necessary and adviced against
Dvnamic BIDS labels
 The BIDS labels can be static, in which case the label is just a normal string, or
\rightarrow dvnamic.
 when the string is enclosed with pointy brackets like `<attribute name>` or
  `<<argument1><argument2>>`. In case of single pointy brackets the label will be,
⇔replaced
 during bidsmapper, bidseditor and bidscoiner runtime by the value of the (DICOM).
→attribute
 with that name. In case of double pointy brackets, the label will be updated for_
→each
 subject/session during bidscoiner runtime. For instance, then the `run` label `<<1>>
→` in
 the bids name will be replaced with `1` or increased to `2` if a file with runindex.
\rightarrow 1
 already exists in that directory.
```

Fieldmaps: suffix Select 'magnitude1' if you have 'magnitude1' and 'magnitude2' data in one series-∽folder (this is what Siemens does) -- the bidscoiner will automatically pick up the → 'magnitude2' data during runtime. The same holds for 'phase1' and 'phase2' data. See the BIDS specification for more details on fieldmap suffixes Fieldmaps: IntendedFor You can use the `IntendedFor` field to indicate for which runs (DICOM series) a\_ → fieldmap was intended. The dynamic label of the `IntendedFor` field can be a list of string\_ →patterns that is used to include all runs in a session that have that string pattern in. →their BIDS file name. Example: use `<<task>>` to include all functional runs or `<<Stop\*Go>  $\leftrightarrow$  < Reward >>` to include "Stop1Go"-, "Stop2Go"- and "Reward"-runs. NB: The fieldmap might not be used at all if this field is left empty! Manual editing / inspection of the bidsmap You `can of course also directly edit or inspect the `bidsmap.yaml` file yourself\_ ⇔with any text editor. For instance to merge a set of runs that by adding a '\*' wildcard to a\_ →DICOM attribute in one run item and then remove the other runs in the set. See ./docs/ →bidsmap.md and ./heuristics/bidsmap\_dccn.yaml for more information.

As shown below, the main window of the bidseditor opens with the BIDS map tab that contains a list of input samples that uniquely represents all the different files that are present in the source folder, together with the associated BIDS output name. The path in the BIDS output name is shown in red if the modality is not part of the BIDS standard, striked-out gray when the runs will be ignored in the conversion to BIDS, otherwise it is colored green. Double clicking the sample (DICOM) filename opens an inspection window with the full header information (double clicking sample filenames works throughout the GUI).

	ant labels			
	ant labels			
	n < <sourcefilepath>&gt;</sourcefilepath>			
	mples			_
^	DICOM input	BIDS output	Action	
01	00002_1.3.12.2.1107.5.2.19.45416.2017121915051526005675150.IMA	anat\sub-001_ses-mri01_acq-t1mpragesagp2iso10_run-1_T1w.*	Edit	
02	00001_1.3.12.2.1107.5.2.43.66068.2019040818580312385121448.IMA	anat\sub-POM1FM0031237_ses-mri01_acq-T1p21mmfov256sagTl880ukbiobank_run-1_T1w.*	Edit	
03	00001_1.3.12.2.1107.5.2.43.66068.2019040818580311755321445.IMA	anat\sub-POM1FM0031237_ses-mri01_acq-T1p21mmfov256sagTl880ukbiobank_run-1_T1w.*	Edit	
04	00001_1.3.12.2.1107.5.2.43.66068.2019040819361792477841719.IMA	extra_data\sub-POM1FM0031237_ses-mri01_acq-3dflair1mmserialimagingadaptedND_run-1.*	Edit	
005	00001_1.3.12.2.1107.5.2.43.66068.2019040819361797231241725.IMA	anat\sub-POM1FM0031237_ses-mri01_acq-3dflair1mmserialimagingadapted_run-1_FLAIR.*	Edit	
006	00002_1.3.12.2.1107.5.2.19.45416.2017121915074837751578028.IMA	func\sub-001_ses-mri01_task-cmrr25isombme3TR1500TaskPart1_acq-epfid2d384_run-1_echo-2_sbref.*	Edit	
007	00002_1.3.12.2.1107.5.2.19.45416.2017121915074990016378598.IMA	func\sub-001_ses-mri01_task-cmrr25isombme3TR1500TaskPart1_acq-epfid2d384_run-1_echo-2_bold.*	Edit	
800	00003_1.3.12.2.1107.5.2.19.45416.2017121916140779237785804.IMA	func\sub-001_ses-mri01_task-cmrr25isombme3TR1500TaskPart1_acq-epfid2d384_run-1_echo-3_bold.*	Edit	
09	00001_1.3.12.2.1107.5.2.43.66068.2019040819002138085426936.IMA	leave_out\sub-POM1FM0031237_ses-mri01_acq-MB8fMRIfov21024mmukbiobankSBRef_run-1.*	Edit	
010	00001_1.3.12.2.1107.5.2.43.66068.2019040819002162138127114.IMA	leave_out\sub-POM1FM0031237_ses-mri01_acq-MB8fMRIfov21024mmukbiobank_run-1.*	Edit	
)11	00001_1.3.12.2.1107.5.2.43.66068.2019040819135564837635013.IMA	func\sub-POM1FM0031237_ses-mri01_task-MB6fMRI20isoTR1000TE34_acq-epfid2d1104_run-1_echo-1_bold.*	Edit	
)12	00001_1.3.12.2.1107.5.2.19.45416.2017121916090972370383869.IMA	fmap\sub-001_ses-mri01_acq-grefieldmappingMB3ME_run-1_magnitude1.*	Edit	
)13	00001_1.3.12.2.1107.5.2.19.45416.2017121916091054452284165.IMA	fmap\sub-001_ses-mri01_acq-grefieldmappingMB3ME_run-1_phasediff.*	Edit	
)14	00073_1.3.12.2.1107.5.2.43.66068.2019040819301282781840881.IMA	fmap\sub-POM1FM0031237_ses-mri01_acq-diffUKBioBankAdaptedMB3invertedSBRef_dir-COL_run-1_epi.*	Edit	
)15	00001_1.3.12.2.1107.5.2.43.66068.2019040819301443079241259.IMA	fmap\sub-POM1FM0031237_ses-mri01_acq-diffUKBioBankAdaptedMB3inverted_dir-COL_run-1_epi.*	Edit	
016	PET.IMA	pet\sub-001_ses-mri01_task-PETdynLM20170420191301000000_acq-FluoroLdopa_run-1_pet.*	Edit	
)17	sub-034_ses-drug3_task-RL_file-1_dicom.IMA	extra_data\sub-034_ses-drug3_acq-SeriesDescription_run-1.*	Edit	

The user can click the Edit button for each list item to open a new edit window, as show below. In this interface, the right BIDS Modality (drop down menu) and the suffix label (drop down menu) can set correctly, after which the associated BIDS Labels can be edited (double click black items). As a result, the new BIDS Output name is then shown in the bottom text field. This is how the BIDS output data will look like and, if this looks all fine, the user can store this mapping to the bidsmap and return to the main window by clicking the OK button.

DICOM input		BIDS output		
Provenance		Modality		
filename 00001_1.3.12.2.1107.5.2.43.67027.2020042808484341531102371.IMA		anat		
		Labels		
		acq t1mpragesagipat21p0iso		
Attributes		ce		
Modality	MR	rec		
ProtocolName	t1_mprage_sag_ipat2_1p0iso	run <<1>>		
SeriesDescription	t1_mprage_sag_ipat2_1p0iso	mod		
ImageType	['ORIGINAL', 'PRIMARY', 'M', 'ND', 'NORM']	suffix T1w		
SequenceName	*tfl3d1_16ns			
SequenceVariant	['SK', 'SP', 'MP']			
ScanningSequence	['GR', 'IR']			
MRAcquisitionType	3D			
SliceThickness	1			
FlipAngle	8			
EchoNumbers	1	Output name		
EchoTime	3.03	anat\sub-PrismaFitDCCNsequences_acq-t1mpragesagipat21p0iso_run-1_T1w.*		
RepetitionTime	2300			
PhaseEncodingDirecti	on			

Tip: Use the Export button to append new or unknown run items to your template bidsmap for usage in other studies

Finally, if all BIDS output names in the main window are fine, the user can click on the Save button and proceed with

running the bidscoiner tool.

### 2.3.3 Step 2: Running the bidscoiner

```
usage: bidscoiner [-h] [-p PARTICIPANT_LABEL [PARTICIPANT_LABEL ...]] [-f]
                  [-s] [-b BIDSMAP] [-n SUBPREFIX] [-m SESPREFIX] [-v]
                  sourcefolder bidsfolder
Converts ("coins") datasets in the sourcefolder to nifti / json / tsv datasets in the
bidsfolder according to the BIDS standard. Check and edit the bidsmap.yaml file to
your needs using the bidseditor tool before running this function. You can run
bidscoiner after all data is collected, or run / re-run it whenever new data has
been added to the source folder (presuming the scan protocol hasn't changed). If you
delete a (subject/) session folder from the bidsfolder, it will be re-created from the
sourcefolder the next time you run the bidscoiner.
Provenance information, warnings and error messages are stored in the
bidsfolder/code/bidscoin/bidscoiner.log file.
positional arguments:
                        The source folder containing the raw data in
 sourcefolder
                        sub-#/[ses-#]/data format (or specify --subprefix and
                        --sesprefix for different prefixes)
 bidsfolder
                        The destination / output folder with the bids data
optional arguments:
 -h, --help
                       show this help message and exit
 -p PARTICIPANT_LABEL [PARTICIPANT_LABEL ...], --participant_label PARTICIPANT_LABEL_
→ [PARTICIPANT_LABEL ...]
                        Space seperated list of selected sub-# names / folders
                        to be processed (the sub- prefix can be removed).
                        Otherwise all subjects in the sourcefolder will be
                        selected
 -f, --force
                        If this flag is given subjects will be processed,
                        regardless of existing folders in the bidsfolder.
                        Otherwise existing folders will be skipped
 -s, --skip_participants
                        If this flag is given those subjects that are in
                        particpants.tsv will not be processed (also when the
                        --force flag is given). Otherwise the participants.tsv
                        table is ignored
 -b BIDSMAP, --bidsmap BIDSMAP
                        The bidsmap YAML-file with the study heuristics. If
                        the bidsmap filename is relative (i.e. no "/" in the
                        name) then it is assumed to be located in
                        bidsfolder/code/bidscoin. Default: bidsmap.yaml
 -n SUBPREFIX, --subprefix SUBPREFIX
                        The prefix common for all the source subject-folders.
                        Default: 'sub-'
 -m SESPREFIX, --sesprefix SESPREFIX
                        The prefix common for all the source session-folders.
                        Default: 'ses-'
 -v, --version
                        Show the BIDS and BIDScoin version
examples:
 bidscoiner /project/foo/raw /project/foo/bids
 bidscoiner -f /project/foo/raw /project/foo/bids -p sub-009 sub-030
```

Tip: Check your json sidecar files of your fieldmaps, in particular see if they have the expected IntendedFor values.

**Note:** The provenance of the produced BIDS data-sets is stored in the [bidsfolder]/code/bidscoin/bidscoiner.log file. This file is also very useful for debugging / tracking down bidscoin issues.

# 2.4 Finishing up

After a successful run of bidscoiner, the work to convert your data in a fully compliant BIDS dataset is unfortunately not yet fully over and, depending on the complexity of your data-set, additional tools may need to be run and meta-data may need to be entered manually (not everything can be automated).

#### 2.4.1 Adding meta-data

For instance, you should update the content of the dataset\_description.json and README files in your bids folder and you may need to provide e.g. additional \*\_scans.tsv,\*\_sessions.tsv or participants.json files (see the BIDS specification for more information). Moreover, if you have behavioural log-files you will find that BIDScoin does not (yet) support converting these into BIDS compliant \*\_events.tsv/json files (advanced users are encouraged to use the bidscoiner plug-in possibility and write their own log-file parser).

#### 2.4.2 Data sharing utilities

#### **Multi-echo combination**

Before sharing or pre-processing their images, users may want to combine the separate the individual echos of multiecho MRI acquisitions. The echcombine-tool is a wrapper around mecombine that writes BIDS valid output.

```
usage: echocombine [-h] [-p PARTICIPANT_LABEL [PARTICIPANT_LABEL ...]]
                   [-o {fmap, anat, func, dwi, beh, pet, extra_data, derivatives}]
                   [-a {PAID, TE, average}] [-w [WEIGHTS [WEIGHTS ...]]]
                   bidsfolder pattern
A wrapper around the 'mecombine' multi-echo combination tool (https://github.com/
→Donders-Institute/multiecho).
This wrapper is fully BIDS-aware (a 'bidsapp') and writes BIDS compliant output
positional arguments:
 bidsfolder
                        The bids-directory with the (multi-echo) subject data
                        Globlike recursive search pattern (relative to the
 pattern
                        subject/session folder) to select the first echo of
                        the images that need to be combined, e.g.
                        '*task-*echo-1*'
optional arguments:
 -h, --help
                        show this help message and exit
  -p PARTICIPANT_LABEL [PARTICIPANT_LABEL ...], --participant_label PARTICIPANT_LABEL_
→ [PARTICIPANT_LABEL ...]
```

```
Space separated list of sub-# identifiers to be
                        processed (the sub- prefix can be left out). If not
                        specified then all sub-folders in the bidsfolder will
                        be processed (default: None)
 -o {fmap,anat,func,dwi,beh,pet,extra_data,derivatives}, --output {fmap,anat,func,
→dwi,beh,pet,extra_data,derivatives}
                        A string that determines where the output is saved. It
                        can be the name of a BIDS modality folder, such as
                        'func', or of the derivatives folder, i.e.
                        'derivatives'. If output = [the name of the input
                        modality folder] then the original echo images are
                        replaced by one combined image. If output is left
                        empty then the combined image is saved in the input
                        modality folder and the original echo images are moved
                        to the extra_data folder (default: None)
 -a {PAID,TE,average}, --algorithm {PAID,TE,average}
                        Combination algorithm (default: TE)
 -w [WEIGHTS [WEIGHTS ...]], --weights [WEIGHTS [WEIGHTS ...]]
                        Weights for each echo (default: None)
examples:
 echocombine /project/3017065.01/bids func/*task-stroop*echo-1*
 echocombine /project/3017065.01/bids *task-stroop*echo-1* -p 001 003
 echocombine /project/3017065.01/bids func/*task-*echo-1* -o func
 echocombine /project/3017065.01/bids func/*task-*echo-1* -o derivatives -w 13 26 39_
\rightarrow 52
 echocombine /project/3017065.01/bids func/*task-*echo-1* -a PAID
```

#### Defacing

Before sharing or pre-processing their images, users may want to deface their anatomical MRI acquisitions as to protect the privacy of their subjects. The deface-tool is a wrapper around pydeface that writes BIDS valid output.

```
usage: deface [-h] [-p PARTICIPANT_LABEL [PARTICIPANT_LABEL ...]]
              [-o {fmap,anat,func,dwi,beh,pet,extra_data,derivatives}] [-c]
              [-n NATIVESPEC] [-a ARGS]
             bidsfolder pattern
A wrapper around the 'pydeface' defacing tool (https://github.com/poldracklab/
→pydeface).
This wrapper is fully BIDS-aware (a 'bidsapp') and writes BIDS compliant output
positional arguments:
 bidsfolder
                        The bids-directory with the (multi-echo) subject data
 pattern
                        Globlike search pattern (relative to the
                        subject/session folder) to select the images that need
                        to be defaced, e.g. 'anat/*_T1w*'
optional arguments:
 -h, --help
                        show this help message and exit
  -p PARTICIPANT_LABEL [PARTICIPANT_LABEL ...], --participant_label PARTICIPANT_LABEL_
↔ [PARTICIPANT_LABEL ...]
                        Space separated list of sub-# identifiers to be
                        processed (the sub- prefix can be left out). If not
```

```
specified then all sub-folders in the bidsfolder will
                        be processed (default: None)
 -o {fmap,anat,func,dwi,beh,pet,extra_data,derivatives}, --output {fmap,anat,func,
→dwi,beh,pet,extra_data,derivatives}
                        A string that determines where the defaced images are
                        saved. It can be the name of a BIDS modality folder,
                        such as 'anat', or of the derivatives folder, i.e.
                        'derivatives'. If output is left empty then the
                        original images are replaced by the defaced images
                        (default: None)
 -c, --cluster
                        Flag to submit the deface jobs to the high-performance
                        compute (HPC) cluster (default: False)
 -n NATIVESPEC, --nativespec NATIVESPEC
                        DRMAA native specifications for submitting deface jobs
                        to the HPC cluster (default: -1
                        walltime=00:30:00,mem=1gb)
 -a ARGS, --args ARGS Additional arguments (in dict/json-style) that are
                        passed to pydeface. See examples for usage (default:
                        {})
examples:
 deface /project/3017065.01/bids anat/*_T1w*
 deface /project/3017065.01/bids anat/*_T1w* -p 001 003 -o derivatives
 deface /project/3017065.01/bids anat/*_T1w* -n "-1 walltime=00:60:00,mem=2gb"
 deface /project/3017065.01/bids anat/*_T1w* -a '{"cost": "corratio", "verbose": ""}'
```

#### 2.4.3 BIDS validation

If all of the above work is done, you can (and should) run the web-based bidsvalidator to check for inconsistencies or missing files in your bids data-set (NB: the bidsvalidator also exists as a command-line tool).

## 2.5 Options

BIDScoin has different options and settings (see below) that can be adjusted per study bidsmap. You can use a text editor to edit the bidsmap template [path\_to\_bidscoin]/heuristics/bidsmap\_template.yaml if you want to adjust the default)

#### 2.5.1 BIDScoin

- version: should correspond with the version in ../bidscoin/version.txt
- bidsignore: Semicolon-separated list of entries that are added to the .bidsignore file (for more info, see BIDS specifications), e.g.:

```
- extra_data/;pet/;myfile.txt;yourfile.csv
```

#### 2.5.2 dcm2niix

The nifti- and json-files are generated with dcm2niix. Here you can adjust how dcm2niix is used:

• path: Command to set the path to dcm2niix, e.g.:

I bids\code\bidscoin\bidsmap.yaml - BIDS editor -	
ile Help	
File browser Options BIDS map	
bidscoin	
version 2.3.1 bidsignore extra_data/	Test
dcm2niix	
path module add dcm2niix; args -b y -z y -i n	Test
Plugins	
path //fileserver.dccn.nl/marzwi\$/python/bidscoin/bidscoin/plugins/README.py path	Test Select
Reset Save	Help

Fig. 1: The bidseditor options window with the different BIDScoin settings

- module add dcm2niix/1.0.20180622; (note the semi-colon at the end)
- PATH=/opt/dcm2niix/bin:\$PATH; (note the semi-colon at the end)
- /opt/dcm2niix/bin/ (note the slash at the end)
- '\"C:\\Program Files\\dcm2niix\"' (note the quotes to deal with the whitespace)
- args: Argument string that is passed to dcm2niix. Click [Test] and see the terminal output for usage

Tip: SPM users may want to use '-z n', which produces unzipped nifti's

#### 2.5.3 Plugins

BIDScoin provides the possibility for researchers to write custom python functions that will be executed at bidsmapper and bidscoiner runtime. To use this functionality, enter the name of the module (default location is the plugins-folder; otherwise the full path must be provided) in the bidsmap dictionary file to import the plugin functions. See advanced usage for more details.

# 2.6 Advanced usage

#### 2.6.1 Site specific / customized template

If you want to convert many studies with similar acquisition protocols then you **may** consider (NB: this is in no way necessary) creating your own customized bidsmap template. This template can then be passed to the bidsmapper tool (instead of the default [path\_to\_bidscoin]/heuristics/bidsmap\_template.yaml template) to automatically identify the different scans in your (future) studies and map these to the correct BIDS modalities.

Generally speaking, a bidsmap file contains a collection of key-value dictionaries that define unique mappings between different types (runs) of source data onto BIDS outcome data. As illustrated in the figure below, each run item in the bidsmap has a provenance key-value pair to store the pathname of a representative data sample of that run. Each run item also contains a source data attributes object, i.e. a key-value dictionary with keys and values that are extracted from the provenance data sample, as well as a bids object, i.e. a key-value dictionary that determines the filename of the BIDS output file. The different key-value pairs in the attributes dictionary represent properties of the source data and should uniquely identify the different runs in a session. But these attrribute-values should not vary between sessions, making the length of the bidsmap only dependent on the acquisition protocol and not on the number of subjects and sessions in the data collection. The difference between a bidsmap template and the study bidsmap that comes out of the bidsmapper is that the template contains / defines the key-values that will be used by the bidsmapper and that the template contains all possible runs. The study bidsmap contains only runs that were encountered in the study, with key-values that are specific for that study. A bidsmap has different sections for different source data modalities, i.e. DICOM, PAR, P7, Nifti, FileSystem, as well as a section for the BIDScoin Options. Within each source data section there sub-sections for the different BIDS modalities, i.e. for anat, func, dwi, fmap, pet, beh and extra\_data, and for the participant\_label and session\_label. It is important to note that BIDScoin tools, given a data sample, will go through the bidsmap (from top to bottom) until they come across a run with attribute values that match the attribute values of the data sample (NB: empty values are ignored). At that point a bidsmapping is made, i.e. the bids values will be taken to contruct a BIDS output filename.

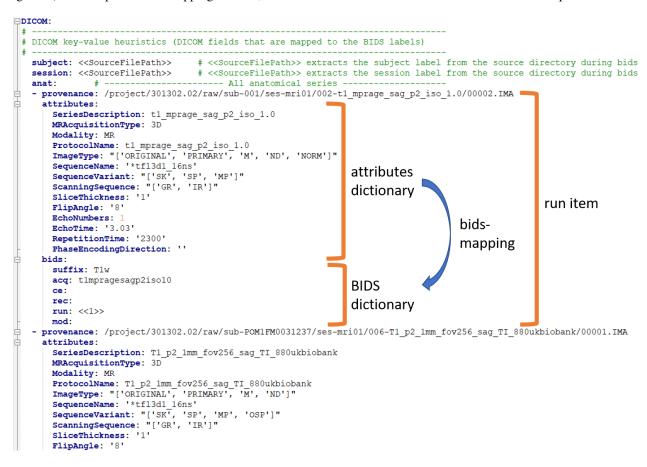


Fig. 2: A snippet of a study bidsmap, showing a DICOM section with a few run items in the anat subsection

To create your own template bidsmap you can best first make a copy of the default template ([path\_to\_bidscoin]/heuristics/bidsmap\_template.yaml) or of the DCCN example template ([path\_to\_bidscoin]/heuristics/bidsmap\_dccn.yaml) and customize that bidsmap to your needs:

1. Using the bidseditor. This is the easiest way to create a bidsmap template since it uses only a GUI and doesn't

require in-depth knowledge of bidsmaps and YAML files. If you have a run item in your study that you would like to be automatically mapped in other / future studies you can simply append that run to the standard or to a custom template bidsmap by editting it to your needs and click the Export button (see below). With the GUI you can still use advanced features, such as Unix shell-style wildcards in the values of the source attributes (see left panel), or such as using lists of attribute values (of which either one can match), or simply empty field to ignore the item. The main limitation of using the GUI is that the run items are always appended to a bidsmap template, meaning that they are last in line and will be used only if no other item in the template matches. It also means that like this you cannot edit the already existing run items in the bidsmap. Another (smaller) limitation is that with the GUI you cannot make usage of YAML anchors and references. Both limitations are overcome when directly editting the template bidsmap yourself using a text editor (see next point).

Edit BIDS mapping		- D X
DICOM input		BIDS output
Provenance		Modality
path C:\Users\Ma	rcel\Data\bidscoin\raw\sub-PrismaFit_DCCN_sequences\007	anat $\checkmark$
filename 00001_1.3.12.2.1107.5.2.43.67027.2020042808484341531102371.IMA		Labels
		acq t1mpragesagipat21p0iso
Attributes		ce
Modality	MR	rec
ProtocolName	t1_mprage_sag_ipat2_1p0iso	run <<1>>
SeriesDescription	t1_mprage_sag_ipat2_1p0iso	mod
ImageType	['ORIGINAL', 'PRIMARY', 'M', 'ND', 'NORM']	suffix T1w ~
SequenceName	*tfl3d1_16ns	
SequenceVariant	['SK', 'SP', 'MP']	
ScanningSequence	['GR', 'IR']	
MRAcquisitionType	3D	
SliceThickness	1	
FlipAngle	8	
EchoNumbers	1	Output name
EchoTime	3.03	anat\sub-PrismaFitDCCNsequences_acq-t1mpragesagipat21p0iso_run-1_T1w.*
RepetitionTime	2300	
PhaseEncodingDirecti	on	
Reset		OK Export Cancel Help

Fig. 3: The edit window with the option to export the customized mapping of run a item

2. Using a text editor. This is the most powerful way to create or modify a bidsmap template but requires more indepth knowledge of YAML and of how BIDScoin identifies different acquisitions in a protocol given a bidsmap. How you can customize your template is well illustrated by the DCCN template bidsmap ([path\_to\_bidscoin]/heuristics/bidsmap\_dccn.yaml). If you open that template, there are a few things to take notice of (as shown in the template snippet below). First, you can see that the DCCN template makes use of YAML anchors and aliases (to make maintanance more sustainable). The second thing to notice is that, of the first run, all values of the attribute dictionary are empty, meaning that it won't match any run / will be ignored. In that way, however, the subsequent runs that alias (<<: \*anatattributes\_dicom) this anchor (&anatattributes\_dicom) will inherit only the keys and can inject their own values, as shown in the second run. The first run of each modality sub-section (like anat) also serves as the default bidsmapping when users manually overrule / change the bids modality using the bidsmapper GUI. Finally, it is important to take notice of the usage of the lists of values (any of which can match) and Unix shell-style wildcards (see DICOM Attributes).

```
anat: # ------ All anatomical runs ------
- provenance: ~ # The first run item with empty attributes will not_
→match anything but will be used when changing modality in the bidseditor GUI ->_
→suffix = Tlw
attributes: &anatattributes_dicom
Modality: ~
```

```
ProtocolName: ~
   SeriesDescription: ~
   ImageType: ~
   SequenceName: ~
   SequenceVariant: ~
   ScanningSequence: ~
   MRAcquisitionType: ~
   SliceThickness:
   FlipAngle: ~
   EchoNumbers: ~
   EchoTime: ~
   RepetitionTime: ~
   PhaseEncodingDirection: ~
 bids: &anatbids dicom
   acq: <SeriesDescription> # A dynamic label which will be replaced during.
⇔bidscoiner runtime with the DICOM attribute value
   ce: ~
   rec: ~
   run: <<1>>
                                # A dynamic label that will be increased during_
→ bidscoiner runtime. NB: changing this value may lead to collisions / overwriting of.
→BIDS data
   mod: ~
   suffix: T1w
                                # The second run item with non-empty attributes (
- provenance: ~
→ 'SeriesDescription' and 'MRAcquisitionType') will match any run with these.
→attribute values
 attributes:
   <<: *anatattributes dicom
   SeriesDescription: ['*mprage*', '*MPRAGE*', '*MPRage*', '*t1w*', '*T1W*']
   MRAcquisitionType: 3D
 bids:
    <<: *anatbids_dicom
   suffix: T1w
```

Snippet from the ''bidsmap\_dccn.yaml'' template, showing a DICOM section with the first two run items in the anat subsection

### 2.6.2 Plugins

BIDScoin has the option to import plugins to further automate / complete the conversion from source data to BIDS. The plugin takes is called each time the BIDScoin tool has finished processing a run or session, with arguments containing information about the run or session, as shown in the plugin example code below. The functions in the plugin module should be named bidsmapper\_plugin to be called by bidsmapper and bidscoiner\_plugin to be called by bidscoiner.

```
The full-path name of the raw-data series folder
    :param seriesfolder:
                              The study bidsmap
    :param bidsmap:
    :param bidsmap_template: Full BIDS heuristics data structure, with all options,
→ BIDS labels and attributes, etc
    :return:
                                The study bidsmap with new entries in it
    .....
   LOGGER.debug(f'This is a bidsmapper demo-plugin working on: {seriesfolder}')
   return bidsmap
def bidscoiner_plugin(session: Path, bidsmap: dict, bidsfolder: Path, personals:
→dict) -> None:
    ......
    The plugin to cast the series into the bids folder
    :param session: The full-path name of the subject/session raw data source.
⇔folder
   :param bidsmap: The full mapping heuristics from the bidsmap YAML-file
    :param bidsfolder: The full-path name of the BIDS root-folder
   :param personals: The dictionary with the personal information
    :return:
                       Nothing
    .....
   LOGGER.debug(f'This is a bidscoiner demo-plugin working on: {session} ->
\leftrightarrow {bidsfolder}')
```

Plugin example code

# 2.7 Screenshots

### 2.7.1 The bidseditor

# 2.8 Demo and tutorial

### 2.8.1 BIDS introduction and BIDScoin demo

A good starting point to learn more about BIDS and BIDScoin is to watch this presentation from the OpenMR Benelux 2020 meeting (slides). The first 14 minutes Robert Oostenveld provides a general overview of the BIDS standard, after which Marcel Zwiers presents the design of BIDScoin and demonstrates hands-on how you can use it to convert a dataset to BIDS.

### 2.8.2 Hands-on tutorial

The following tutorial is somewhat tailored to the dataflow in the DCCN (e.g. using the module system to set the BID-Scoin shell environment), but should nevertheless make the basic parts of the BIDScoin workflow clear for everyone.

1. **Preparation.** Activate the bidscoin environment and create a tutorial playground folder in your home directory by executing these bash commands (see also module help bidscoin):

	owser Options BIDS map			
srticip	ant labels			
ubje	ct < <sourcefilepath>&gt;</sourcefilepath>			
essio	on < <sourcefilepath>&gt;</sourcefilepath>			
ata sa	Imples			
^	DICOM input	BIDS output	Action	1
01	00002_1.3.12.2.1107.5.2.19.45416.2017121915051526005675150.IMA	anat\sub-001_ses-mri01_acq-t1mpragesagp2iso10_run-1_T1w.*	Edit	
)2	00001_1.3.12.2.1107.5.2.43.66068.2019040818580312385121448.IMA	anat\sub-POM1FM0031237_ses-mri01_acq-T1p21mmfov256sagTl880ukbiobank_run-1_T1w.*	Edit	
03	00001_1.3.12.2.1107.5.2.43.66068.2019040818580311755321445.IMA	anat\sub-POM1FM0031237_ses-mri01_acq-T1p21mmfov256sagTl880ukbiobank_run-1_T1w.*	Edit	
)4	00001_1.3.12.2.1107.5.2.43.66068.2019040819361792477841719.IMA	extra_data\sub-POM1FM0031237_ses-mri01_acq-3dflair1mmserialimagingadaptedND_run-1.*	Edit	
)5	00001_1.3.12.2.1107.5.2.43.66068.2019040819361797231241725.IMA	anat\sub-POM1FM0031237_ses-mri01_acq-3dflair1mmserialimagingadapted_run-1_FLAIR.*	Edit	
)6	00002_1.3.12.2.1107.5.2.19.45416.2017121915074837751578028.IMA	func\sub-001_ses-mri01_task-cmrr25isombme3TR1500TaskPart1_acq-epfid2d384_run-1_echo-2_sbref.*	Edit	
07	00002_1.3.12.2.1107.5.2.19.45416.2017121915074990016378598.IMA	func\sub-001_ses-mri01_task-cmrr25isombme3TR1500TaskPart1_acq-epfid2d384_run-1_echo-2_bold.*	Edit	
08	00003_1.3.12.2.1107.5.2.19.45416.2017121916140779237785804.IMA	func\sub-001_ses-mri01_task-cmrr25isombme3TR1500TaskPart1_acq-epfid2d384_run-1_echo-3_bold.*	Edit	
09	00001_1.3.12.2.1107.5.2.43.66068.2019040819002138085426936.IMA	leave_out\sub-POM1FM0031237_ses-mri01_acq-MB0fMRIfov21024mmukbiobankSBRef_run-1.*	Edit	
10	00001_1.3.12.2.1107.5.2.43.66068.2019040819002162138127114.IMA	leave_out\sub-POM1FM0031237_ses-mri01_acq-MB8fMRIfov21024mmukbiobank_run-1.*	Edit	
11	00001_1.3.12.2.1107.5.2.43.66068.2019040819135564837635013.IMA	func\sub-POM1FM0031237_ses-mri01_task-MB6fMRI20isoTR1000TE34_acq-epfid2d1104_run-1_echo-1_bold.*	Edit	
12	00001_1.3.12.2.1107.5.2.19.45416.2017121916090972370383869.IMA	fmap\sub-001_ses-mri01_acq-grefieldmappingMB3ME_run-1_magnitude1.*	Edit	
13	00001_1.3.12.2.1107.5.2.19.45416.2017121916091054452284165.IMA	fmap\sub-001_ses-mri01_acq-grefieldmappingMB3ME_run-1_phasediff.*	Edit	
14	00073_1.3.12.2.1107.5.2.43.66068.2019040819301282781840881.IMA	fmap\sub-POM1FM0031237_ses-mri01_acq-diffUKBioBankAdaptedMB3invertedSBRef_dir-COL_run-1_epi.*	Edit	
15	00001_1.3.12.2.1107.5.2.43.66068.2019040819301443079241259.IMA	fmap\sub-POM1FM0031237_ses-mri01_acq-diffUKBioBankAdaptedMB3inverted_dir-COL_run-1_epi.*	Edit	
16	PET.IMA	pet\sub-001_ses-mri01_task-PETdynLM20170420191301000000_acq-FluoroLdopa_run-1_pet.*	Edit	
17	sub-034_ses-drug3_task-RL_file-1_dicom.IMA	extra_data\sub-034_ses-drug3_acq-SeriesDescription_run-1.*	Edit	

Fig. 4: The main window with the bidsmap overview

Edit BIDS mapping		- 0	×
DICOM input		BIDS output	
Provenance		Modality	
path C:\Users\Ma	rcel\Data\bidscoin\raw\sub-PrismaFit_DCCN_sequences\007	anat	$\sim$
		Labels	
Attributes		acq t1mpragesagipat21p0iso ce	
Modality	MR	rec	
ProtocolName	t1_mprage_sag_ipat2_1p0iso	run <<1>>	
SeriesDescription	t1_mprage_sag_ipat2_1p0iso	mod	
ImageType	['ORIGINAL', 'PRIMARY', 'M', 'ND', 'NORM']	suffix T1w	$\sim$
SequenceName	*tfl3d1_16ns		
SequenceVariant	['SK', 'SP', 'MP']		
ScanningSequence	['GR', 'IR']		
MRAcquisitionType	3D		
SliceThickness	1		
FlipAngle	8		
EchoNumbers	1	Output name	
EchoTime	3.03	anat\sub-PrismaFitDCCNsequences_acq-t1mpragesagipat21p0iso_run-1_T1w.*	
RepetitionTime	2300		
PhaseEncodingDirecti	ion		
Reset		OK Export Cancel	Help

Fig. 5: The edit window for customizing the individual bidsmap entries

■ bids\code\bidscoin\bidsmap.yaml - BIDS editor —	
ile Help	
File browser Options BIDS map	
bidscoin	
version 2.3.1 bidsignore extra_data/	Test
dcm2niix	
path module add dcm2niix; args -b y -z y -i n	Test
Plugins	
path //fileserver.dccn.nl/marzwi\$/python/bidscoin/bidscoin/plugins/README.py path	Test Select
Reset	Help

Fig. 6: The options window with BIDScoin settings

```
$ module add bidscoin
$ source activate /opt/bidscoin
$ pulltutorialdata
$ cd bidscointutorial
```

The new bidscointutorial folder contains a raw source-data folder and a bids\_ref reference BIDS folder, i.e. the intended end product of this tutorial. In the raw folder you will find these DICOM series:

```
001-localizer_32ch-head
                                         A localizer scan that is not scientifically
→relevant and can be left out of the BIDS dataset
002-AAHead_Scout_32ch-head
                                         A localizer scan that is not scientifically,
\hookrightarrowrelevant and can be left out of the BIDS dataset
007-t1_mprage_sag_ipat2_1p0iso An anatomical T1-weighted scan
047-cmrr_2p4iso_mb8_TR0700_SBRef A single-band reference scan of the
→subsequent multi-band functional MRI scan
048-cmrr_2p4iso_mb8_TR0700
                                          A multi-band functional MRI scan
049-field_map_2p4iso
                                          The fieldmap magnitude images of the first.
\hookrightarrowand second echo. Set as "magnitude1", bidscoiner will recognize the format. This
\leftrightarrow fieldmap is intended for the previous functional MRI scan
050-field_map_2p4iso
                                          The fieldmap phase difference image of the
\rightarrow first and second echo
059-cmrr_2p5iso_mb3me3_TR1500_SBRef
                                         A single-band reference scan of the
→subsequent multi-echo functional MRI scan
060-cmrr_2p5iso_mb3me3_TR1500
                                          A multi-band multi-echo functional MRI scan
061-field_map_2p5iso
                                          Idem, the fieldmap magnitude images of the
\rightarrow first and second echo, intended for the previous functional MRI scan
062-field_map_2p5iso
                                          Idem, the fieldmap phase difference image of
→the first and second echo
```

Let's begin with inspecting this new raw data collection:

- Are the DICOM files for all the bids/sub-\* folders organised in series-subfolders (e.g. sub-001/ ses-01/003-T1MPRAGE/0001.dcm etc)? Use dicomsort if this is not the case. A help text for all BID-Scoin tools is available by running the tool with the -h flag (e.g. rawmapper -h)
- Use the rawmapper command to print out the DICOM values of the "EchoTime", "Sex" and "AcquisitionDate"

of the fMRI series in the raw folder

2. **BIDS mapping.** Now we can make a bidsmap, i.e. the mapping from DICOM source-files to BIDS target-files. To that end, scan all folders in the raw data collection by running the bidsmapper command:

\$ bidsmapper raw bids

- In the GUI that appears, rename the task and acquisition label of the functional scans into something more readable, e.g. task-Reward for the acq-mb8 scans and "task-Stop" for the acq-mb3me3 scans. Also make the name of the T1 scan more pleasant, e.g. by naming the aquisition label simply acq-mprage.
- Add a search pattern to the IntendedFor field such that it will select your fMRI runs correctly (see the bidseditor fieldmap section for more details)
- Since for this dataset we only have one session per subject, remove the session label (and note how the output names simplify, omitting the session subfolders and labels)
- When all done, (re)open the bidsmap.yaml file and change the options such that you will get non-zipped nifti data (i.e. \*.nii instead of \*.nii.gz) in your BIDS data collection. You can use a text editor or, much better, run the bidseditor command line tool.
- 3. **BIDS coining.** The next step is very simple (and can be repeated when new data comes in) and just applies the previously made bidsmap when converting the source data into a BIDS collection. To do this run the bidscoiner commandline tool (note that the input is the same as for the bidsmapper):

\$ bidscoiner raw bids

- Check your bids/code/bidscoin/bidscoiner.log (the complete terminal output) and bids/ code/bidscoin/bidscoiner.errors (the summary that is also printed at the end) files for any errors or warnings. You shouldn't have any :-)
- Compare the results in your bids/sub-\* subject folders with the in bids\_ref reference result. Are the file and foldernames the same (don't worry about the multi-echo images and the extra\_data images, they are combined/generated as described below)? Also check the json sidecar files of the fieldmaps. Do they have the right "EchoTime" and "IntendedFor" fields?
- What happens if you re-run the bidscoiner command? Are the same subjects processed again? Re-run "sub-001".
- 4. **Finishing up.** Now that you have converted the data to BIDS, you still need to do some manual work to make it fully ready for data analysis and sharing
- Combine the echos using the echocombine tool, such that the individual echo images are replaced by the echcombined image
- Deface the anatomical scans using the echocombine tool. This will take a while, but will obviously not work well for our phantom dataset. Therefore store the 'defaced' output in the extra\_data folder (instead of e.g. overwriting the existing images)
- Inspect the bids/participants.tsv file and decide if it is ok.
- Update the dataset\_description.json and README files in your bids folder
- As a final step, run the bids-validator on your ~/bids\_tutorial folder. Are you completely ready now to share this dataset?