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# In-Vivo-Imaging-Pipeline

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## MANAGEMENT

### 1.1 ConfigHandling module

`ConfigHandling.config_generator(Config: Optional[dict] = None) → None`

Generates a configuration file

**Parameters**

**Config** (*dict*) – Configuration to write to file (Optional)

**Return type**

None

`ConfigHandling.config_reader(File: Optional[str]) → dict`

Reads config into a dictionary

**Parameters**

**File** (*str*) – Configuration File (Optional)

**Returns**

Configuration

**Return type**

dict

### 1.2 Organization module

`class Organization.BehavioralExperiment(Meta: Tuple[str, str], ExperimentName: str)`

Bases: *Experiment*

Experiment class for a generic day of a behavioral task

**Required Inputs**

*Meta* : Passed meta from experimental hierarchy (directory, mouse\_id)

*ExperimentName* : Title of ExperimentName

**Properties**

*mouse\_id* : Identifies which mouse this data belongs to

*instance\_data* : Identifies when this behavioral ExperimentName was created

### Attributes

*data* : Pandas dataframe of synced data  
*folder\_dictionary* : A dictionary of relevant folders for this behavioral ExperimentName  
*modifications* : List of modifications made to this behavioral ExperimentName  
*multi\_index*: Pandas multi-index of behavioral components  
*state\_index* : look-up table / index relating states to integers  
*trial\_parameters* : behavioral parameters

### Methods

*copy\_raw\_behavioral\_data* : Interactive tool for copying raw behavioral data  
*load\_data* : Loads all data  
*record\_mod* : Records a modification made to the behavioral ExperimentName (Date & Time)  
*update\_folder\_dictionary* : This function re-indexes all folders in the folder dictionary

**copy\_raw\_behavioral\_data()** → Self

Interactive tool for copying raw behavioral data

#### Return type

Any

**load\_data()** → Self

Loads behavioral data

#### Return type

Any

**class Organization.Data(Path: str)**

Bases: object

This is a class for managing a folder of unorganized data files

### Required Inputs

*Path* : path to folder

### Self Methods

*find\_matching\_files* : Finds all matching files  
*reindex* : Function that indexed the files within folder again  
*find\_all\_ext* : Finds all files with specific extension

### Properties

*instance\_data* : Data created  
*path* : path to folder  
*files* : List of files in folder

**property files:** List[str]

**find\_all\_ext**(*Ext: str*) → Optional[List[str]]

Finds all files with specific extension

**Parameters**

**Ext** (*str*) – Filename extension

**Returns**

List of files

**Return type**

List[str]

**find\_matching\_files**(*Filename: str, Folder: Optional[str] = None*) → Optional[Tuple[str]]

Finds all matching files

**Parameters**

- **Filename** (*str*) – Filename or ID to search for
- **Folder** (*Any*) – Specify folder filename in

**Returns**

Matching file/s

**Return type**

Any

**property folders: dict**

Dictionary of folders in path

**Return type**

dict

**property instance\_date: str**

Date Created

**Return type**

str

**property path: str**

Path to folder

**Return type**

str

**reindex()** → Self

Function that indexes the files within folder again

**class** Organization.**Experiment**(*Meta: Tuple[str, str], ExperimentName: str*)

Bases: object

Experiment class for a generic experiment

**Required Inputs**

*Meta* : Passed meta from mouse (directory, mouse\_id)

*ExperimentName* : Title of experiment

**Properties**

*instance\_data* : Identifies when this behavioral experiment was created

*mouse\_id* : Identifies which mouse this data belongs to

### Attributes

*data* : a pandas dataframe containing synchronized data

*folder\_dictionary* : A dictionary of relevant folders for this experiment

*modifications* : List of modifications made to this experiment

### Public Methods

*copy\_data* : Interactive tool to copy data to directory (Intended to be overwritten during inheritance)

*load\_data* : Loads all data (Intended to be overwritten during inheritance)

*record\_mod* : Records a modification made to the experiment (Date & Time)

*update\_folder\_dictionary* : This function re-indexes all folders in the folder dictionary

**copy\_data()** → Self

Interactive tool to copy data to directory

**Return type**

Any

**property experiment\_id:** str

**property instance\_date:** str

Date created

**Return type**

str

**property mouse\_id:** str

ID of mouse

**Return type**

str

**record\_mod()** → Self

Records a modification made to the behavioral ExperimentName (Date & Time)

**Return type**

Any

**update\_folder\_dictionary()** → Self

This function re-indexes all folders in the folder dictionary

**Return type**

Any

**class Organization.Figures**(Path: str)

Bases: *Data*

Data Folder specifically for storing figures.

**view\_figure**(Name: str) → plt.Figure

Function identifies and views a figure based on supplied name

**Parameters**

**Name** (str) – Name of figure (can be partial)



**Returns**

the plotted figure

**Return type**

Any

**class** Organization.**Images**(Path: str)Bases: *Data*

Data Folder specifically for folders containing raw images.

**property** channels**property** file\_format**property** frames**property** height**property** imaging\_files**property** meta\_files**property** num\_imaging\_files**property** num\_meta\_files**property** planes**reorganize\_bruker\_files**() → None

This function extracts out the meta files and saves in a new directory

**Return type**

None

**property** width**class** Organization.**ImagingAnalysis**(Path: str)Bases: *Data*

Data Folder specifically for imaging analysis folders.

**Self Methods***load\_fissa\_exports* : loads fissa exported files*load\_cascade\_exports* : loads cascade exported files*load\_suite2p* : loads suite2p exported files*export\_registration\_to\_denoised* : moves registration to new folder for namespace compatibility when skipping denoising step*clean\_up\_motion\_correction* : This function removes the reg\_tif folder and registered.bin generated during motion correction.*clean\_up\_compilation* : This function removes the compiled tif files*add\_notes* : Function adds notes**add\_notes**(Step: str, KeyOrDict: Union[str, dict], Notes: Optional[Any] = None) → Self

Function adds notes indicating steps

**Parameters**

- **Step** – Step of Analysis

- **Step** – str
- **KeyOrDict** (*Union[str, dict]*) – Either a Key or a dictionary containing multiple key-value (note) pairs
- **Notes** (*Optional[Any]*) – If using key, then notes is the paired value

**Return type**

Any

**clean\_up\_compilation()** → Self

This function removes the compiled tif files generated inside CompiledImagingData (You can avoid the creation of these in the first place by changing suite2p parameters)

**Return type**

Any

**clean\_up\_motion\_correction()** → Self

**This function removes the reg\_tif folder and registered.bin generated during motion correction.**  
(You can avoid the creation of these in the first place by changing suite2p parameters)

**Return type**

Any

**property current\_ExperimentName:** str

ExperimentName of Analysis

**Return type**

str

**default\_folders()**

**export\_registration\_to\_denoised()**

moves registration to new folder for namespace compatibility

**Returns**

**load\_cascade\_exports()** → Tuple[ndarray, ndarray, ndarray, dict]

This function loads the Spike Times, Spike Prob, Discrete Approximation and ProcessedInferences files exported from Cascade

**Returns**

SpikeTimes, SpikeProb, DiscreteApproximation, Processed Inferences

**Return type**

tuple[Any, Any, Any, dict]

**load\_fissa\_exports()** → Tuple[dict, dict, dict]

This function loads the prepared and separated files exported from Fissa

**Returns**

Prepared, Separated, ProcessedTraces

**Return type**

tuple[dict, dict, dict]

**load\_suite2p(\*args: str)**

**class** Organization.**ImagingBehaviorExperiment**(*Meta: Tuple[str, str], ExperimentName: str*)

Bases: *ImagingExperiment, BehavioralExperiment*

Experiment class for a generic day of an Imaging / Behavioral experiment

### Required Inputs

*Meta* : Passed meta from experimental hierarchy (directory, mouse\_id)

*ExperimentName* : Title of ExperimentName

### Properties

*mouse\_id* : Identifies which mouse this data belongs to

*instance\_data* : Identifies when this behavioral ExperimentName was created

### Attributes

*data* : Pandas dataframe of synced data

*folder\_dictionary* : A dictionary of relevant folders for this behavioral ExperimentName

*modifications* : List of modifications made to this behavioral ExperimentName

*meta* : bruker metadata

*multi\_index*: Pandas multi-index of behavioral components

*state\_index* : look-up table / index relating states to integers

*trial\_parameters* : behavioral parameters

### Methods

*copy\_raw\_imaging\_data* : Interactive tool for copying raw imaging data

*copy\_raw\_behavioral\_data* : Interactive tool for copying raw behavioral data

*load\_data* : Loads all data

*record\_mod* : Records a modification made to the experiment (Date & Time)

*update\_folder\_dictionary* : This function re-indexes all folders in the folder dictionary

**copy\_data()** → Self

Interactive tool to copy data to directory

#### Return type

Any

**load\_data**(*ImagingParameters: Optional[Union[dict, list[dict]]] = None, \*args: Optional[Tuple[str, str]], \*\*kwargs*) → Self

Loads all data

#### Parameters

**ImagingParameters** – Parameters for some imaging dataset or list of datasets

(e.g., for two different sampling rates) :type ImagingParameters: Optional[dict] :param args: Optionally pass Sync Key to synchronize bruker recordings :type args: Tuple[str, str] :param kwargs: passed to internal functions taking kwargs :rtype: Any

**class** Organization.**ImagingExperiment**(*Meta: Tuple[str, str], ExperimentName: str*)

Bases: *Experiment*

Experiment class for a generic imaging experiment

### Required Inputs

*Meta* : Passed meta from mouse (directory, mouse\_id)

*ExperimentName* : Title of experiment

### Properties

*mouse\_id* : Identifies which mouse this data belongs to

*instance\_data* : Identifies when this experiment was created

### Attributes

*data* : a pandas dataframe containing synchronized data

*folder\_dictionary* : A dictionary of relevant folders for this experiment

*meta* : bruker metadata

*modifications* : List of modifications made to this experiment

### Public Methods

*add\_image\_sampling\_folder* : Generates a folder for containing imaging data of a specific sampling rate

*copy\_raw\_imaging\_data* : Interactive tool for copying raw imaging data

*load\_data* : Loads all data

*record\_mod* : Records a modification made to the experiment (Date & Time)

*update\_folder\_dictionary* : This function re-indexes all folders in the folder dictionary

**add\_image\_sampling\_folder**(*SamplingRate: int*) → Self

Generates a folder for containing imaging data of a specific sampling rate

#### Parameters

**SamplingRate** (*int*) – Sampling Rate of Dataset in Hz

#### Return type

Any

**copy\_data**() → Self

Interactive tool to copy data to directory

#### Return type

Any

**copy\_raw\_imaging\_data**() → Self

This function copies raw imaging data to the appropriate folder

#### Return type

Any

**load\_data**(*ImagingParameters: Optional[Union[dict, list[dict]]] = None*) → Self

Loads all data

#### Parameters

**ImagingParameters** (*Optional[dict]*) – Parameters for some imaging dataset or list of datasets (e.g., for two different sampling rates)

#### Return type

Any

**class** Organization.**Mouse**(\*\*kwargs)

Bases: object

Class for Organizing & Managing Experimental Data Across Sessions

### Keyword Arguments

*Logfile* : Path to existing log file (str, default None)  
*Mouse* : Mouse ID (str, default None)  
*Condition* : Experimental Condition (str, default None)  
*Directory* : Directory for hierarchy (str, default None)  
*Study* : Study (str, default None)  
*StudyMouse* : Study ID (str, default None)

### Properties

*mouse\_id* : ID of Mouse  
*log\_file* : Log Filename Path  
*experimental\_condition* : Experiment condition of the mouse  
*instance\_data* : Date when this experimental hierarchy was created

### Attributes

*directory* : Experimental Hierarchy Directory  
*experiments* : Names of included experiments  
*study* : Study  
*study\_mouse* : ID of mouse in study  
*modifications* : modifications made to this file

### Public Class Methods

*load* : Function that loads the entire mouse

### Public Methods

*create* : This function creates the directory/logs/organization.json if it doesn't exist  
*check\_log* : Checks Log Status  
*create\_log\_file* : Creates log file  
*pass\_meta* : Passes directory/mouse id  
*record\_mod* : Record modification of experiment  
*record\_experiments\_mod* : Record modification of experiments  
*save* : Saves mouse to organization.json  
*start\_log* : Starts Log

### Private Class Methods

*\_generate\_analysis\_subdirectory* : Generate Analysis  
*\_generate\_analysis\_technique\_subdirectory* : Generate Analysis Technique  
*\_generate\_behavior\_subdirectory* : Generate Behavioral Folder  
*\_generate\_directory\_structure* : Generates the Directory Structure (The structured folders where data stored)

*\_generate\_experiment\_folders* : Generate Behavioral ExperimentName Folder  
*\_generate\_histology\_directory* : Generates Histology Folder  
*\_generate\_imaging\_subdirectory* : Generate Imaging Folder  
*\_generate\_roi\_matching\_index\_directory* : Generate ROI Matching Folder

**check\_log()** → Self

Checks log status

**Return type**

Any

**create()** → Self

This function generates the directory hierarchy in one step

**Return type**

Any

**create\_experiment**(*ExperimentName: str, Type: Optional[str, Experiment]* = 'Experiment', *\*\*kwargs*) → Self

Generates an experiment ExperimentName folder and attribute

Kwargs are passed to underlying functions

**Parameters**

- **ExperimentName** (*str*) – Name of experimental ExperimentName
- **Type** (*Optional[str, Experiment]*) – Type of experiment (Optional, default = Experiment)

**Return type**

Any

**create\_log\_file()** → Self

Creates log file

**Return type**

Any

**end\_log()** → Self

Ends Logging

**Return type**

Any

**property experimental\_condition:** str

Experiment condition of the mouse

**Return type**

str

**property instance\_date:** str

Date when this experimental hierarchy was created

**Return type**

str

**classmethod load**(*Directory: Optional[str] = None*) → *Mouse*

Function that loads the entire mouse

**Parameters**

**Directory** (*Optional[str]*) – Directory containing the organization.json file and associated data

**Returns**

Mouse

**Return type**

ExperimentManagement.Organization.Mouse

**property log\_file:** **str**

Log Filename Path

**Return type**

str

**property mouse\_id:** **str**

ID of Mouse

**Return type**

str

**pass\_meta()** → Tuple[str, str]

Passes directory/mouse id

**Returns**

directory/mouse id

**Return type**

tuple[str, str]

**record\_experiment\_mod**(*ExperimentNameKey: str, \*args*) → Self

Record modification of experiment (Data, Time, *\*args*)

**Parameters**

- **ExperimentNameKey** (*str*) – The key name for the ExperimentName
- **args** (*str*) – A string explaining the modification

**Return type**

Any

**record\_mod**(*\*args: str*) → Self

Record modification of experiment (Data, Time, *\*args*)

**Parameters**

**args** (*str*) – A string explaining the modification

**Return type**

Any

**save()** → Self

Saves Mouse to json

**Return type**

Any

**start\_log()** → Self

Starts Log

**Return type**

Any

**update\_all\_folder\_dictionaries()** → Self

This function iterates through all behavioral ExperimentNames to update their folder dictionaries

**Return type**

Any

**class** Organization.Study

Bases: object

Organization.**generate\_read\_me**(*AbsoluteFilePath: str, Text: str*) → None

Generate a read me file

**Parameters**

- **AbsoluteFilePath** (*str*) – Filename path
- **Text** (*str*) – Text inside

**Return type**

None

Organization.**get\_date**()

Organization.**get\_time**()

## 1.3 UserInterfaces module

UserInterfaces.**select\_directory**(*\*\*kwargs*) → str

UserInterfaces.**verbose\_copying**(*src, dst*) → None



## INDICES AND TABLES

- `genindex`
- `modindex`
- `search`



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