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# **CalSciPy**

***Release 0.0.5***

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**CalSciPy** contains a variety of useful methods for handling, processing, and visualizing calcium imaging data. It's intended to be a collection of useful, well-documented functions often used in boilerplate code alongside software packages such as [Caiman](#), [SIMA](#), and [Suite2P](#).

## 1.1 Motivation

I noticed I was often re-writing or copy/pasting a lot of code between environments when working with calcium imaging data. I started this package so you don't have to. No more wasting time writing 6 lines to simply preview your tiff stack, extract a particular channel, or bin some spikes. No more vague exceptions or incomplete documentation when re-using a hastily-made function from 2 months ago. Alongside these time-savers, I've also included some more non-trivial methods that are particularly useful.

## 1.2 Limitations

The current distribution for the package is incomplete. When each module has its associated unit tests complete, it will be pushed.

## 1.3 Installation

Enter `pip install CalSciPy` in your terminal.



- *Coloring*
- *Event Processing*
- *Input/Output (I/O)*
- *Image Processing*
- *Interactive Visuals*
- *Reorganization*
- *Signal Processing*
- *Static Visuals*

## 2.1 Coloring

### 2.1.1 Description

Write me

Write me

Write me

Write me

## 2.1.2 Methods

Import me

## 2.2 Event Processing

### 2.2.1 Description

Write me

Write me

Write me

Write me

### 2.2.2 Methods

Import me

## 2.3 Input/Output (I/O)

### 2.3.1 Description

Write me

Write me

Write me

Write me

### 2.3.2 Methods

Import me

## 2.4 Image Processing

### 2.4.1 Description

Write me

Write me

Write me

Write me



## 2.4.2 Methods

Import me

## 2.5 Interactive Visuals

### 2.5.1 Description

Write me

Write me

Write me

Write me

### 2.5.2 Methods

Import me

## 2.6 Reorganization

### 2.6.1 Description

Write me

Write me

Write me

Write me

### 2.6.2 Methods

Import me

## 2.7 Signal Processing

### 2.7.1 Description

Write me

Write me

Write me

Write me

## 2.7.2 Methods

Import me

## 2.8 Static Visuals

### 2.8.1 Description

Write me

Write me

Write me

Write me

### 2.8.2 Methods

Import me

### 3.1 Submodules

#### 3.1.1 CalSciPy.coloring module

#### 3.1.2 CalSciPy.event\_processing module

#### 3.1.3 CalSciPy.image\_processing module

#### 3.1.4 CalSciPy.interactive\_visuals module

#### 3.1.5 CalSciPy.io module

CalSciPy.io.**determine\_bruker\_folder\_contents**(*folder*)

Function determine contents of the bruker folder

**Parameters**

**folder** (*Union[str, pathlib.Path]*) -- Folder containing bruker imaging data

**Returns**

channels, planes, frames, Height, Width

**Return type**

tuple

CalSciPy.io.**load\_all\_tiffs**(*folder*)

Load a sequence of tiff stacks

**Parameters**

**folder** (*Union[str, pathlib.Path]*) -- Folder containing a sequence of tiff stacks

**Returns**

complete\_image numpy array [Z x Y x X] as uint16

**Return type**

np.ndarray

CalSciPy.io.load\_binary\_meta(*path*)

Loads meta file for binary video

**Parameters****path** (*Union[str, pathlib.Path]*) -- The meta file (.txt ext) or directory containing metafile**Returns**

A tuple containing the number of frames, y pixels, and x pixels [Z x Y x X]

**Return type**

tuple[int, int, int, str]

CalSciPy.io.loadbruker\_tiffs(*folder, channels=None, planes=None*)

Load a sequence of tiff files from a directory.

Designed to compile the outputs of a certain imaging utility that exports recordings such that each frame is saved as a single tiff.

**Parameters**

- **folder** (*Union[str, pathlib.Path]*) -- Folder containing a sequence of single frame tiff files
- **channels** (*Optional[int]*) -- channel to load
- **planes** (*Optional[int]*) -- plane to load

**Returns**

complete\_image: All tiff files in the directory compiled into a single array (Z x Y x X, uint16)

**Return type**

Tuple[np.ndarray]

CalSciPy.io.load\_mapped\_binary(*filename, meta\_filename, \*\*kwargs*)

Loads a raw binary file in the workspace without loading into memory

Enter the path to autofill (assumes Filename &amp; meta are path + binary\_video, video\_meta.txt)

**Parameters**

- **filename** (*str*) -- filename for binary video
- **meta\_filename** (*str*) -- filename for meta file
- **mode** -- pass mode to numpy.memmap (str, default = "r")

**Returns**

memmap(numpy) array [Z x Y x X]

**Return type**

np.memmap

CalSciPy.io.load\_raw\_binary(*path, meta\_filename*)

Loads a raw binary file

Enter the path to autofill (assumes Filename &amp; meta are path + binary\_video, video\_meta.txt)

**Parameters**

- **path** (*str*) -- absolute filepath for binary video or directory containing a file named binary video

- **meta\_filename** (*Optional[str]*) -- absolute path to meta file

**Returns**

numpy array [Z x Y x X]

**Return type**

Any

`CalSciPy.io.load_single_tiff(filename, num_frames)`

Load a single tiff file

**Parameters**

- **filename** (*Union[str, pathlib.Path]*) -- absolute filename
- **num\_frames** (*int*) -- number of frames

**Returns**

numpy array [Z x Y x X]

**Return type**

np.ndarray

`CalSciPy.io.pretty_printbruker_command(channels, planes, frames, height, width)`

Function simply prints the bruker folder contents detected

**Parameters**

- **channels** (*int*) -- Number of channels
- **planes** (*int*) -- Number of planes
- **frames** (*int*) -- Number of frames
- **height** (*int*) -- Height of Image (Y Pixels)
- **width** -- Width of Image (X Pixels)

**Return type**

None

`CalSciPy.io.repackagebruker_tiffs(input_folder, output_folder, *args)`

Repackages a sequence of tiff files within a directory to a smaller sequence of tiff stacks. Designed to compile the outputs of a certain imaging utility that exports recordings such that each frame is saved as a single tiff.

**Parameters**

- **input\_folder** (*Union[str, pathlib.Path]*) -- Directory containing a sequence of single frame tiff files
- **output\_folder** (*Union[str, pathlib.Path]*) -- Empty directory where tiff stacks will be saved
- **args** (*int*) -- optional argument to indicate the repackaging of a specific channel and/or plane

**Return type**

None

`CalSciPy.io.save_raw_binary(images, path, meta_filename)`

This function saves a tiff stack as a binary file

**Parameters**

- **images** (*np.ndarray*) -- Images to be saved [Z x Y x X]

- **path** (*str*) -- absolute filepath for saving binary video or directory containing a file named binary video
- **meta\_filename** (*str*) -- absolute filepath for saving meta

**Return type**

None

`CalSciPy.io.save_single_tiff(images, path, type_=<class 'numpy.uint16'>)`

Save a numpy array to a single tiff file as type uint16

**Parameters**

- **images** (*Any*) -- numpy array [frames, y pixels, x pixels]
- **path** (*Union[str, pathlib.Path]*) -- filename or absolute path
- **type** (*Optional[Any]*) -- type for saving

**Return type**

None

`CalSciPy.io.save_tiff_stack(images, output_folder, type_=<class 'numpy.uint16'>)`

Save a numpy array to a sequence of tiff stacks

**Parameters**

- **images** (*Any*) -- A numpy array containing a tiff stack [Z x Y x X]
- **output\_folder** (*Union[str, pathlib.Path]*) -- A directory to save the sequence of tiff stacks in uint16
- **type** (*Optional[Any]*) -- type for saving

**Return type**

None

`CalSciPy.io.save_video(images, path, fps=30)`

Function writes video to .mp4

**Parameters**

- **images** (*Any*) -- Images to be written
- **path** (*Union[str, pathlib.Path]*) -- Filename (Or Complete Path)
- **fps** (*Union[float, int]*) -- frame rate

**Return type**

None

### 3.1.6 CalSciPy.reorganization module

`CalSciPy.reorganization.merge_traces(traces_as_tensor)`

Concatenate multiple trials or tiffs into single matrix:

**Parameters****traces\_as\_tensor** (*ndarray*) --**Returns****Return type**

np.ndarray

### **3.1.7 CalSciPy.static\_visuals module**

### **3.1.8 CalSciPy.trace\_processing module**

## **3.2 Module contents**





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#### 4.1 Parameterized Decorators

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### 4.1.1 Parsing Decorators

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### 4.1.2 Validation Decorators

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### 4.1.3 Terminal Style

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## 4.2 PyTest

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### 4.2.1 Sample Datasets

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### 4.2.2 Tests

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## CHAPTER 5

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