

A tutorial of:

1) downloading needed files from brain observatory api and converting them to matlab tables

2) building a `brain_observatory_cache` object to a) get general information of the whole brain observatory dataset

b) select sessions by

conditions such as brain

areas, imaging depth and

stimuli

c) download nwb files of selected sessions

3) importing imaging data from nwb files to a) get fluorescence traces

b) plot fluorescence traces

c) transform the interested

fluorescence trace data of

the interested subexperiment

into raster formats for decoding

important information about this dataset:

An "experiment container" (named by allen institute) contains a set of "subexperiments" (defined by us as one subexperiment only adopted one kind of stimuli, allen institute doesn't seem to have a name for this concept) that were operated on a single mouse, recorded in a single brain space (same targeted_structure and same imaging depth), performed during one out of three sessions (allen institute equates "session" with "ophys_experiment", in allen_sdk they use "ophys_experiment", in whitepapers they use "session").

Different sessions in the same experiment container may have adopted the same stimulus, which means they may share the same type of "subexperiment":

```
session_by_stimuli.three_session_A =  
{'drifting_gratings','natural_movie_one','natural_movie_three','spontaneous_activity'};  
session_by_stimuli.three_session_B =  
{'static_gratings','natural_scene','natural_movie_one','spontaneous_activity'};  
session_by_stimuli.three_session_C =  
{'locally_sparse_noise_four_degree','natural_movie_one','natural_movie_two','spontaneous_activity'};  
session_by_stimuli.three_session_C2 =  
{'locally_sparse_noise_four_degree','locally_sparse_noise_eight_degree', ...  
'natural_movie_one','natural_movie_two','spontaneous_activity'};
```

our shorthand: `container = experiment_container`

the three names we use: `container > session > subexperiment`

0)

```
% set your bot_dir
```

```
bot_dir_name = '/om/user/xf15/Brain-Observatory-Toolbox/';

% add path to bot_dir
addpath(bot_dir_name)
```

1) download needed files from brain observatory api and converting them to matlab tables

```
get_files_from_brain_obs_api()

load('references')
```

2) build a brain_observatory_cache object

```
% building a brain_observatory_cache object
boc = brain_observatory_cache (references)
```

```
boc =
  brain_observatory_cache with properties:
    session_table: [597×14 table]
    container_table: [199×13 table]
    selected_session_table: [597×14 table]
    stimuli: []
    targeted_structure: []
    imaging_depth: []
    container_id: []
    session_id: []
    session_type: []
    references: [1×1 struct]
```

2a) get general information of the whole brain observatory dataset

```
boc.get_total_of_containers()
```

```
ans = 199
```

```
boc.get_all_imaing_depths()
```

```
ans = 11×1 cell array
    '175'
    '265'
    '275'
    '300'
    '320'
    '325'
    '335'
    '350'
    '365'
    '375'
    '435'
```

```
boc.get_all_cre_lines()
```

```
ans = 6×1 cell array
```

```

'Cux2-CreERT2'
'Emx1-IRES-Cre'
'Nr5a1-Cre'
'Rbp4-Cre_KL100'
'Rorb-IRES2-Cre'
'Scnn1a-Tg3-Cre'

```

```
boc.get_all_targeted_structures()
```

```

ans = 6x1 cell array
    'VISal'
    'VISam'
    'VISl'
    'VISp'
    'VISpm'
    'VISrl'

```

```
boc.get_all_session_types()
```

```

ans = 4x1 cell array
    'three_session_A'
    'three_session_B'
    'three_session_C'
    'three_session_C2'

```

```
boc.get_all_stimuli()
```

```

ans = 9x1 cell array
    'drifting_gratings'
    'locally_sparse_noise_eight_degree'
    'locally_sparse_noise_four_degree'
    'natural_movie_one'
    'natural_movie_three'
    'natural_movie_two'
    'natural_scene'
    'spontaneous_activity'
    'static_gratings'

```

```
boc.get_summary_of_container_along_targeted_structures()
```

VISal	33
VISam	25
VISl	36
VISp	54
VISpm	35
VISrl	16

```
boc.get_summary_of_containers_along_imaging_depths()
```

175	50
265	1
275	72
300	4
320	1
325	3
335	3
350	33
365	1
375	30
435	1

```
boc.get_summary_of_containers_along_depths_and_structures()
```

```
ans = 12x7 table
```

	VISal	VISam	VISl	VISp	VISpm	VISrl	total
	-----	-----	-----	-----	-----	-----	-----
175	8	5	10	11	11	5	50
265	1	0	0	0	0	0	1
275	12	9	15	20	11	5	72
300	1	0	1	1	1	0	4
320	0	1	0	0	0	0	1
325	0	1	1	0	1	0	3
335	0	0	0	2	1	0	3
350	4	4	4	13	4	4	33
365	1	0	0	0	0	0	1
375	6	4	5	7	6	2	30
435	0	1	0	0	0	0	1
total	33	25	36	54	35	16	199

2b) select sessions by conditions such as brain areas, imaging depth and stimuli

```
% Example: search for experiments that primary visual cortex was  
% recorded at 275 mm deep as drifting gratings were shown
```

```
% reinitialize to have a "clean start"  
boc = brain_observatory_cache (references)
```

```
boc =  
brain_observatory_cache with properties:
```

```
    session_table: [597x14 table]  
    container_table: [199x13 table]  
selected_session_table: [597x14 table]  
      stimuli: []  
targeted_structure: []  
    imaging_depth: []  
    container_id: []  
    session_id: []  
    session_type: []  
    references: [1x1 struct]
```

```
% set conditions  
boc.get_sessions_by_stimuli('drifting_gratings')
```

```
ans =  
brain_observatory_cache with properties:
```

```
    session_table: [597x14 table]  
    container_table: [199x13 table]  
selected_session_table: [199x14 table]  
      stimuli: 'drifting_gratings'  
targeted_structure: []  
    imaging_depth: []  
    container_id: []  
    session_id: []  
    session_type: []  
    references: [1x1 struct]
```

```
boc.get_sessions_by_targeted_structure('VISp')
```

```
ans =
  brain_observatory_cache with properties:
    session_table: [597x14 table]
    container_table: [199x13 table]
    selected_session_table: [54x14 table]
    stimuli: 'drifting_gratings'
    targeted_structure: 'VISp'
    imaging_depth: []
    container_id: []
    session_id: []
    session_type: []
    references: [1x1 struct]
```

```
boc.get_sessions_by_imaging_depth(275)
```

```
ans =
  brain_observatory_cache with properties:
    session_table: [597x14 table]
    container_table: [199x13 table]
    selected_session_table: [20x14 table]
    stimuli: 'drifting_gratings'
    targeted_structure: 'VISp'
    imaging_depth: 275
    container_id: []
    session_id: []
    session_type: []
    references: [1x1 struct]
```

```
boc.get_sessions_by_container_id(527550471)
```

```
ans =
  brain_observatory_cache with properties:
    session_table: [597x14 table]
    container_table: [199x13 table]
    selected_session_table: [1x14 table]
    stimuli: {4x1 cell}
    targeted_structure: 'VISp'
    imaging_depth: 275
    container_id: 527550471
    session_id: 527745328
    session_type: {'three_session_A'}
    references: [1x1 struct]
```

```
% get more detailed metadata of selected session(s)
boc.selected_session_table
```

```
ans = 1x14 table
    date_of_acquisition    experiment_container_id    fail_eye_tracking    id    imaging_depth
    -----
    '2016-07-08T15:59:05Z'    5.2755e+08    true    5.2775e+08    275
```

```
% % you can also use brain_observatory_cache to look up metadata of
% % selected session(s) by container_id or session_id
% %
% boc = brain_observatory_cache(references)
```

```
% boc.get_sessions_by_container_id (506823562)
% boc.get_sessions_by_session_id (527676626)
```

2c) download nwb files of selected sessions

```
% download nwb file of the first session in selected sessions into a directory called nwb_files
nwb_dir_name = [bot_dir_name, 'nwb_files/'];

% the size of a nwb file is at the scale of 100 MB
boc.get_session_data(nwb_dir_name);
```

desired nwb file already exists

3 import imaging data from nwb files

```
% add path to nwb files
addpath ([bot_dir_name, 'nwb_files/'])
```

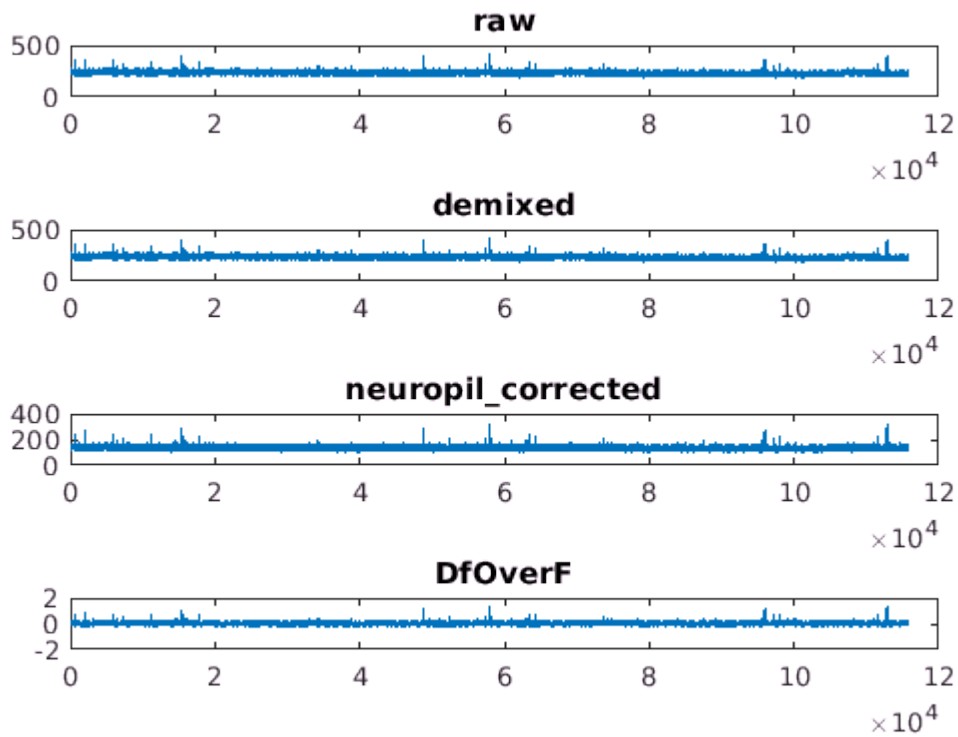
3a) get_fluorescence_traces_of_selected_session

```
[raw,demixed,neuropil_corrected,Df0verF] = get_fluorescence_traces_of_selected_session (boc.session_id, cell_specimen_id);
```

Elapsed time is 5.519662 seconds.

3b) plot the f traces of one cell in session 517745328

```
cell_specimen_id = 529022196;
plot_fluorecence_traces_of_selected_cell_in_selected_session(boc.session_id, cell_specimen_id);
```



3c) transform data of the selected fluorescence trace of the selected subexperiment into raster format

```
raster_dir_name = [bot_dir_name, 'raster/'];

stimuli = 'drifting_gratings';
fluorescence_trace = DfOverF;

current_raster_dir_name = transform_fluorescence_trace_into_raster_format(fluorescence_trace,
    boc.session_id, stimuli, raster_dir_name);
```

/om/user/xf15/Brain-Observatory-Toolbox/raster/drifting_gratings_527745328/ already exists