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# Meta-info

Meta-info is information not stored in the respective data files directly, like electrode location, unit\_IDs, SNR/Stability/single rating and procedures

# electrode\_depths file

Find/create the electrode depth file corresponding to your monkey in the following location: C:\Users\username\Dropbox\DAG\phys\Monkey\_phys\_dpz\Electrode\_depths\_Mon.m (replace username with your username, Monkey with the monkey's name and Mon with the first three letters of the monkey's name).

It should look like this:

```
 k=0; \\ k=k+1; \; Session\{k\}=20150508; \; block\{k\}=1; \; channels\{k\}=[3]; \; z\{k\}=[15000]; \\ k=k+1; \; Session\{k\}=20150508; \; block\{k\}=2; \; channels\{k\}=[3]; \; z\{k\}=[15000]; \\ k=k+1; \; Session\{k\}=20150508; \; block\{k\}=3; \; channels\{k\}=[1 \; 3]; \; z\{k\}=[15735 \; 14879];
```

Note that here each BLOCK has one line, channels contains only useful channels (there was an electrode which was not broken) and the depths (z) correspond to the channels. Electrode depths within certain range will be taken as the same LFP site and combined for waveclus sorting.

# same\_cells file

Find/create the same cells file corresponding to your monkey in the following location: C:\Users\username\Dropbox\DAG\phys\Monkey\_phys\_dpz\Same\_cells\_Mon.m (replace username with your username, Monkey with the monkey's name and Mon with the first three letters of the monkey).

It should look like this:

```
 k=0; \\ k=k+1; \; Session\{k\}=20150508; \; channel\{k\}=3; \; blocks\{k\}=[1 \quad 2]; \; sortcodes\{k\}=[1 \quad 1]; \\ k=k+1; \; Session\{k\}=20150513; \; channel\{k\}=1; \; blocks\{k\}=[1 \quad 2]; \; sortcodes\{k\}=[1 \quad 1]; \\ k=k+1; \; Session\{k\}=20150514; \; channel\{k\}=3; \; blocks\{k\}=[1 \quad 2]; \; sortcodes\{k\}=[1 \quad 1]; \\ k=k+1; \; Session\{k\}=20150514; \; channel\{k\}=3; \; blocks\{k\}=[1 \quad 2]; \; sortcodes\{k\}=[1 \quad 1]; \\ k=k+1; \; Session\{k\}=20150514; \; channel\{k\}=3; \; blocks\{k\}=[1 \quad 2]; \; sortcodes\{k\}=[1 \quad 2]; \; sortcodes\{k
```

Note that here each CELL has one line, channel is unique per line, sortcodes correspond to blocks. The same cell might be assigned to different sortcodes in different blocks. **Importantly, you don't need to fill in each cell here, only the cells that were recorded across several blocks.** 

### sorted neurons table

Find/create the Sorted neurons excel table corresponding to your monkey in the following location: C:\Users\username\Dropbox\DAG\phys\Monkey\_phys\_dpz\Mon\_sorted\_neurons.xlsx (replace username with your username, Monkey with the monkey's name and Mon with the first three letters of the monkey's name).

There are two main sheets in this excel file:

- automatic sorting
- final sorting

The idea is, that a large part of the information can be updated automatically (Create excel entries with phys\_gui), while some information has to be added manually in the final\_sorting sheet. For both sheets, each line represents one unit recorded in one block (and run) AND/OR the corresponding LFP site

### **Automatically created entries**

Monkey	Session	Date	Run	Block	Chan	z	Unit	N_spk	Neur	ron_ID	Site	_ID		Times_same_unit
Lin	1	2010508	3	1	1	15000	a	15251	Lin_2	0150508_01	Lin_2	20150508_9	Site_01	1
Lin	1	2010508	4	2	1	15000	a	15251	Lin_2	0150508_01	Lin_2	20150508_9	Site_01	1

- Monkey: monkey initials (first 3 letters)
- Session: A session counter (irrelevant, but helpful for browsing)
- · Date: Date of recording
- Run: Corresponding behavioral run
- Block: ephys recording block
- Chan: electrode number (channel)
- z: Electrode depth (from guide tube tip) taken from Electrode depths file
- Unit: sortcode assigned to this cell (a=first,b=second,...)
- N spk: Total spike count
- Neuron ID: Cell identifier (the Same cells file can be recorded in several blocks!)
- Site\_ID: LFP site identifier (Based on electrode depth, same site can be recorded in different blocks and at the same time with different units)
- Times\_same\_unit: For identifying unique units

### Manually added entries

SNR_rank	Single_rank	Stability_rank	Set	Target	Hemisphere	Grid	x	у
1	1	0	1	dPul	left	22.5	-5	8
3	2	1	1	dPul	left	22.5	-5	8

The first three entries should be taken from your notes during sorting.

- SNR rank: Signal to noise ratio assessment (1 best SNR to 3 marginal SNR)
- Single rank: Multiunit or single unit (1 Single unit (SU) to 3 Multiunit (MU) )
- Stability\_rank: How stable were the features across time (0 unit appeared or disappeared during the course of the run; 1 stable)

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The other entries typically don't change within a session and can be taken from the handwritten log (notes taken during recording, to be found in Y:\Logs\Phys

• Set: Dataset, relevant for assigning to specific project (see dataset convention

• Target: Area

• Hemisphere: left/right

• Grid: grid identifier (diameter.number)

• x: grid hole coordinate x

• y: grid hole coordinate y

#### **Electrode travel**

Last, but not least, there is another entry which is typically calculated inside the table using different approaches per experimenter: Electrode\_travel. Several columns serve the purpose of calculating the electrode travel (from top of the grid).

#### **Dataset convention**

Numeric code	Meaning
General notes	Neuron set IDs with no decimals correspond to the ones in which the task had the desired characteristics for a given dataset, e.g. set =2 gaze modulation with 3 initial starting locations 15° horizontal eccentricity and 10° vertical eccentricity; set=2.1 even when recorded before than 2, it contains a different eccentricity that was not used for the final dataset
Additional notes	when unit ID has a termination "a" e.g Lin_20160506_05a it means for that neuron the saccade target size matched the reach target size
1	Direct-visually- and memory-guided saccade dataset
2	Gaze modulation dataset, i.e. memory saccades, rectangular grid of targets starting at -15°, 0°, +15°
2.1	Memory saccades center-out / out-center 12° and 24° eccentricity (blocked per run)
2.2	Memory saccades center-out & out-center 12° and 24° eccentricity (interleaved in the same run)
2.3	Memory saccades, rectangular grid of targets starting at -12°, 0°, +12°, saccading 12° peripherally. In this dataset there were trials in which fixation was required instead of a saccade. From this dataset onwards ITI was recorded as well
2.4	Memory saccades, rectangular grid of targets starting at -12°, 0°, +12°, or -15°, 0°, +15° In this dataset there were trials in which fixation was required instead of a saccade
3	Reach dataset
3.1	Reach dataset, joint reach task

To differentiate inactivation from control runs, we also indicate this by assigning a different data(sub-)set. Ideally, following the logic above, this would mean something like 3.5 for control and 3.6 for inactivation. Currently 31 and 32 are used respectively. For convention, Inactivation data(sub-)set value should always be higher than the respective control.

#### Other entries ???

Difficult\_sorting\_or\_unstable\_Other\_potential\_reasons\_to\_ignore\_or\_check Runcomplete Usable Comments task

### plx files table

The plx file tables contain two sheets:

- to\_use determines which plx file will be used next time the TDT and monkeypsych data is combined
- in\_use stores information about which plx file was used last time data was combined

In most cases, you won't have to worry about this, becuase you can automatically update the plx file table using phys gui. However, you CAN in principle also update this manually.

Both sheets look like this:

Monkey	Date	Block	Sorttype	Plx_file_extension
Lin	20150506	8	Snippets	1
Lin	20150508	1	realigned	1
Lin	20150508	2	realigned	2

# automatic processing logs

Every time you run phys\_gui\_execute, two new files will be created in Y:\Data\All\_phys\_preprocessing\_log\Monkey\_phys

• attempted <u>YYYYMMDD-hhmm - stored settings before any execution \* executed YYYYMMDD-hhmm - stored settings after running the entire execution (successfully)</u>

This allows to track what processing has been performed, and in case of unsuccessful completion recover the itnended settings

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