

NWB File Format Specification

Version 1.0.3

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Introduction

Neurodata Without Borders: Neurophysiology is a project to develop a unified data format for cellular-based neurophysiology data, focused on the dynamics of groups of neurons measured under a large range of experimental conditions. Participating labs provided use cases and critical feedback to the effort. The design goals for the NWB format included:

Compatibility

- Cross-platform
- Support for tool makers

Usability

- Quickly develop a basic understanding of an experiment and its data
- Review an experiment's details without programming knowledge

Flexibility

- Accommodate an experiment's raw and processed data
- Encapsulate all of an experiment's data, or link to external data source when necessary

Extensibility

- Accommodate future experimental paradigms without sacrificing backwards compatibility.
- Support custom extensions when the standard is lacking

Longevity

- Data published in the format should be accessible for decades

[Hierarchical Data Format \(HDF\)](#) was selected for the NWB format because it met several of the project's requirements. First, it is a mature data format standard with libraries available in multiple programming languages. Second, the format's hierarchical structure allows data to be grouped into logical self-documenting sections. Its structure is analogous to a file system in which its "groups" and "datasets" correspond to directories and files. Groups and datasets can have attributes that provide additional details, such as authorities' identifiers. Third, its linking feature enables data stored in one location to be transparently accessed from multiple locations in the hierarchy. The linked data can be external to the file. Fourth, [HDFView](#), a free, cross-platform application, can be used to open a file and browse data. Finally, ensuring the ongoing accessibility of HDF-stored data is the mission of The HDF Group, the nonprofit that is the steward of the technology.

The NWB format standard is codified in a schema file written in a specification language created for this

project. The specification language describes the schema, including data types and associations. A new schema file will be published for each revision of the NWB format standard. Tool developers are expected to take advantage of this explicit description of the format. However, data publishers are not expected to do so. They can add custom datasets to existing TimeSeries, Epochs, Modules and Interfaces.

Top Level Groups

| Group | Description | Comment | Required |
|--------------|---|---|----------|
| /general | Experimental metadata, including protocol, notes and description of hardware device(s) | The metadata stored in this section should be used to describe the experiment. Metadata necessary for interpreting the data is stored with the data. | Yes |
| /acquisition | Data streams recorded from the system, including ephys, ophys, tracking, etc. | This group is read-only after the experiment is completed and timestamps are corrected to a common timebase. The data stored here may be links to raw data stored in external HDF5 files. This will allow keeping bulky raw data out of the file while preserving the option of keeping some/all in the file. | Yes |
| /stimulus | Data pushed into the system (eg, video stimulus, sound, voltage, etc) and secondary representations of that data (eg, measurements of something used as a stimulus) | This group is read-only after experiment complete and timestamps are corrected to common timebase. Stores both presented stimuli and stimulus templates, the latter in case the same stimulus is presented multiple times, or is pulled from an external stimulus library. | Yes |
| /epochs | Experimental intervals, whether that be logically distinct sub-experiments having a particular scientific goal, trials during an experiment, or epochs deriving from analysis of data. | Epochs provide pointers to time series that are relevant to the epoch, and windows into the data in those time series (i.e., the start and end indices of <i>TimeSeries::data[]</i> that overlap with the epoch). This allows easy access to a range of data in specific experimental intervals. | Yes |
| /processing | The home for processing <i>Modules</i> . These modules perform intermediate analysis of data that is necessary to perform before scientific analysis. Examples include spike clustering, extracting position from tracking data, stitching together image slices. | <i>Modules</i> are defined below. They can be large and express many data sets from relatively complex analysis (e.g., spike detection and clustering) or small, representing extraction of position information from tracking video, or even binary “lick/no-lick” decisions. Common software tools (e.g., klustakwik, MClust) are expected to read/write data here. | Yes |
| /analysis | Lab-specific and custom scientific analysis of data. There is no defined format for the content of this group – the format is up to the individual user/lab. | To facilitate sharing analysis data between labs, the contents here should be stored in standard types (eg, INCF types) and appropriately documented. | Yes |

The content of these organizational groups is more fully described in the section titled, [File Organization](#). The NWB format is based on *TimeSeries* and *Modules* and these are defined first.

NWB stores general optical and electrical physiology data in a way that should be understandable to a naive user after a few minutes using looking at the file in an HDF5 browser, such as HDFView. The format is designed to be friendly to and usable by software tools and analysis scripts, and to impose few a priori assumptions about data representation and analysis. Metadata required to understand the data itself (core metadata) is generally stored with the data. Information required to interpret the experiment (general metadata) is stored in the group 'general'. Most general metadata is stored in free-form text fields. Machine-readable metadata is stored as attributes on these free-form text fields.

The only API assumed necessary to read a NWB file is an HDF5 library (e.g., h5py in python, libhdf5 in C, JHI5 in Java).

Top Level datasets

Top-level datasets are for file identification and version information.

| Dataset | Type | Description | Comments | Required |
|----------------------|------------|---|--|----------|
| /nwb_version | text | File version string | Eg, "NWB-1.0.0". This is the identifying string (e.g., "NWB") with trailing major, minor and patch numbers. | Yes |
| /identifier | text | Unique file ID | Eg, concatenated lab name, file creation date/time and experimentalist, or a hash of these and/or other values. The goal is that the string should be unique to all other files. | Yes |
| /file_create_date | text array | Time file was created, UTC, and subsequent modifications to file. | Date + time, Use ISO format (eg, ISO 8601). File can be created after the experiment was run, so this may differ from experiment start time. Each modification to file adds new entry to array | Yes |
| /session_start_time | text | Time of experiment/session start, UTC | Date + time, Use ISO format (eg, ISO 8601). All times stored in the file use this time as reference (ie, time zero) | Yes |
| /session_description | text | One or two sentences describing the experiment and data in the file | | Yes |

Throughout the document, sizes are provided for the different datasets. These are both recommended sizes (for integers) and minimum sizes (for floats). The size (number of bits) defined for integer fields in the specification is the recommended size. Larger or smaller sizes can be used, both signed and unsigned, so long as the selected integer size encompasses the full range of data without over or underflow. For floating-point values, the size defined in the specification is the minimum size. A double-precision (float64) or larger can be used when a single-precision float is specified, but the reverse is not true.

TimeSeries

The file format is designed around a data structure called a *TimeSeries* which stores time-varying data. A *TimeSeries* is a superset of several INCF types, including signal events, image stacks and experimental events. To account for different storage requirements and different modalities, a *TimeSeries* is defined in a minimal form and it can be extended, or subclassed, to account for different modalities and data storage requirements. When a *TimeSeries* is extended, it means that the 'subclassed' instance maintains all of the data interface (eg, groups and datasets) as its parent and it has new groups and/or datasets of its own. The *TimeSeries* makes this process of defining such pairs more hierarchical.

Each *TimeSeries* has its own HDF5 group, and all datasets belonging to a *TimeSeries* are in that group. The group contains time and data components and users are free to add additional fields as necessary. There are two time objects represented. The first, *timestamps*, stores time information that is corrected to the experiment's time base (i.e., aligned to a master clock, with time-zero aligned to the starting time of the experiment). This field is used for data processing and subsequent scientific analysis. The second, *sync*, is an optional group that can be used to store the sample times as reported by the acquisition/stimulus hardware,

before samples are converted to a common timebase and corrected relative to the master clock. This approach allows the NWB format to support streaming of data directly from hardware sources.

| Dataset | Type | Description | Comment | Required |
|---------------------|---------------|---|--|---|
| data | polymorphic | Data values. Can also store binary data (eg, image frames) | This field may be a link to data stored in an external file, especially in the case of raw data. | Yes |
| _ unit (attr) | text | The base unit of measure used to store data. This should be in the SI unit. | This is the SI unit (when appropriate) of the stored data, such as Volts. If the actual data is stored in millivolts, the field 'conversion' below describes how to convert the data to the specified SI unit | Yes |
| _ conversion (attr) | float32 | Scalar to multiply each element in data to convert it to the specified <i>unit</i> | | Yes |
| _ resolution (attr) | float32 | Smallest meaningful difference between values in data, stored in the specified by <i>unit</i> | E.g., the change in value of the least significant bit, or a larger number if signal noise is known to be present. If unknown, use NaN | Yes |
| timestamps | float64 array | Timestamps for the samples stored in <i>data</i> . | Timestamps here have all been corrected to the common experiment master-clock. Time is stored as seconds and all timestamps are relative to experiment start time. Only one of <i>timestamps</i> and <i>starting_time</i> should be present | Yes -- unless <i>starting_time</i> is present |
| _ interval (attr) | int32 | The number of samples between each timestamp | Presently this value is restricted to 1 (ie, a timestamp for each sample) | Yes |
| _ unit (attr) | text | The string “Seconds” | All timestamps in the file are stored in seconds. Specifically, this is the number of seconds since the start of the experiment (i.e., since <i>session_start_time</i>) | Yes |
| starting_time | float64 | The timestamp of the first sample | When timestamps are uniformly spaced, the timestamp of the first sample can be specified and all subsequent ones calculated from the sampling rate. Only one of <i>timestamps</i> and <i>starting_time</i> should be present | Yes -- unless <i>timestamps</i> is present |
| _ rate (attr) | float32 | Sampling rate, in Hz | Rate information is stored in Hz | Yes |
| _ unit (attr) | text | The string “Seconds” | All timestamps in the file are stored in seconds. Specifically, this is the number of seconds since the start of the experiment (i.e., since <i>session_start_time</i>) | Yes |
| num_samples | int32 | Number of samples in data, or number of image frames. | This is important if the length of <i>timestamp</i> and <i>data</i> are different, such as for externally stored stimulus image stacks | Yes |
| control | uint8 array | Numerical labels that apply to each element in <i>data</i> [] | Optional field. If present, the control array should have the same number of elements as <i>data</i> [] | No |
| control_description | text array | Description of each control value | Array length should be as long as the highest number in control, generating an indexed array | No |

| | | | | |
|--------------|------------------|---|--|----|
| | | | for control values | |
| sync/ | group (optional) | Lab specific time and sync information as provided directly from hardware devices and that is necessary for aligning all acquired time information to a common timebase. The <i>timestamp</i> array stores time in the common timebase. | This group will usually only be populated in <i>TimeSeries</i> that are stored external to the NWB file, in files storing raw data. Once <i>timestamp</i> data is calculated, the contents of 'sync' are mostly for archival purposes. | No |

When data is streamed from experiment hardware it should be stored in an HDF5 dataset having the same attributes as *data*, with time information stored as necessary. This allows the raw data files to be separate file-system objects that can be set as read-only once the experiment is complete. *TimeSeries* objects in /acquisition will link to the *data* field in the raw time series. Hardware-recorded time data must be corrected to a common time base (e.g., timestamps from all hardware sources aligned) before it can be included in *timestamps*. The uncorrected time can be stored in the *sync* group.

The group holding the *TimeSeries* has the following attributes:

| Attribute | Type | Description | Comment | Required |
|----------------|------------|---|---|-----------|
| description | text | Description of <i>TimeSeries</i> | | Yes |
| source | text array | Name of <i>TimeSeries</i> or <i>Modules</i> that serve as the source for the data contained here. It can also be the name of a device, for stimulus or acquisition data | | Yes |
| comments | | Human-readable comments about the <i>TimeSeries</i> . This second descriptive field can be used to store additional information, or descriptive information if the primary description field is populated with a computer-readable string. | | Yes |
| ancestry | text array | The class-hierarchy of this <i>TimeSeries</i> , with one entry in the array for each ancestor. An alternative and equivalent description is that this <i>TimeSeries</i> object contains the datasets defined for all of the <i>TimeSeries</i> classes listed. The class hierarchy is described more fully below | For example: [0]="TimeSeries", [1]="ElectricalSeries" [1]="PatchClampSeries". The hierarchical order should be preserved in the array -- i.e., the parent object of subclassed element N in the array should be element N-1 | Yes |
| neurodata_type | text | The string "TimeSeries" | | Yes |
| data_link | text array | List of the paths of all <i>TimeSeries</i> that share a hard link to the same <i>data</i> field | Field is only present if links are present. List should include the path to this <i>TimeSeries</i> also | Yes |
| timestamp_link | | List of the paths of all <i>TimeSeries</i> that share a hard link to the same <i>timestamps</i> field | Field is only present if links are present. List should include the path to this <i>TimeSeries</i> also | Yes |
| missing_fields | text array | (Optional) This is a list of fields defined as 'required' parts of the | Only present if one or more recommended fields are missing. Note that a missing required | Recommend |

| | | | | |
|------|------|--|---|----|
| | | <i>TimeSeries</i> that are missing | field (such as <i>data</i> or <i>timestamps</i>) should generate an error by the API | d |
| help | text | (Optional) Displays the 'description' field of the time series stored in the specification file/language | | No |

The group holding the *TimeSeries* can be used to store additional information (HDF5 datasets) beyond what is required by the specification. I.e., an end user is free to add additional key/value pairs as necessary for their needs. It should be noted that such lab-specific extensions may not be recognized by analysis tools/scripts existing outside the lab. All datasets that are added and that are not defined by the schema should have an attribute 'neurodata_type' storing the string 'Custom'.

The *data* element in the *TimeSeries* will typically be an array of any valid HDF5 data type (e.g., a multi-dimensional floating point array). The data stored can be in any unit. The attributes of the data field must indicate the SI unit that the data relates to (or appropriate counterpart, such as color-space) and the multiplier necessary to convert stored values to the specified SI unit.

TimeSeries Class Hierarchy

The *TimeSeries* is a data structure/object. It can be “subclassed” to represent more narrowly focused modalities (e.g., electrical versus optical physiology) as well as new modalities (eg, video tracking of whisker positions). When it a *TimeSeries* is subclassed, new datasets can be added while all datasets of parent classes are preserved. An initial set of subclasses are described here. Users are free to define subclasses for their particular requirements. This can be done by adding a new *TimeSeries* entry to the specification language file. All datasets that are defined to be part of *TimeSeries* have the text attribute 'unit' that stores the unit specified in the documentation.

AbstractFeatureSeries extends TimeSeries

Abstract features, such as quantitative descriptions of sensory stimuli. The *TimeSeries::data* field is a 2D array, storing those features (e.g., for visual grating stimulus this might be orientation, spatial frequency and contrast). **Array structure:** [num frames] [num features]. Null stimuli (eg, uniform gray) can be marked as being an independent feature (eg, 1.0 for gray, 0.0 for actual stimulus) or by storing NaNs for feature values, or through use of the *TimeSeries::control* fields. A set of features is considered to persist until the next set of features is defined. The final set of features stored should be the null set.

| Dataset | Type | Description | Required |
|---------------|------------|--|-------------|
| feature | text array | Description of the features represented in <i>TimeSeries::data</i> | Yes |
| feature_units | text array | Array of features' units | Recommended |

AnnotationSeries extends TimeSeries

Stores, eg, user annotations made during an experiment. The *TimeSeries::data[]* field stores a text array, and timestamps are stored for each annotation (ie, interval=1). This is largely an alias to a standard *TimeSeries* storing a text array but that is identifiable as storing annotations in a machine-readable way.

IndexSeries extends TimeSeries

Stores indices to image frames stored in an ImageSeries. The purpose of the ImageIndexSeries is to allow a static image stack to be stored somewhere, and the images in the stack to be referenced out-of-order. This can be for the display of individual images, or of movie segments (as a movie is simply a series of images). The *data* field stores the index of the frame in the referenced ImageSeries, and the *timestamps* array indicates when that image was displayed. The ImageIndexSeries contains all datasets of *TimeSeries* plus the following:

| Dataset | Type | Description | Required |
|-------------------------|----------------------------|---|----------|
| indexed_timeseries | link: <i>TimeSeries</i> | HDF5 link to <i>TimeSeries</i> containing images that are indexed | Yes |
| indexed_timeseries_path | text | Path to linked <i>TimeSeries</i> | Yes |

IntervalSeries extends TimeSeries

Stores intervals of data. The *timestamps* field stores the beginning and end of intervals. The *data* field stores whether the interval just started (>0 value) or ended (<0 value). Different interval types can be represented in the same series by using multiple key values (eg, 1 for feature A, 2 for feature B, 3 for feature C, etc). The field *data* stores an 8-bit integer. This is largely an alias of a standard TimeSeries but that is identifiable as representing time intervals in a machine-readable way.

OptogeneticSeries extends TimeSeries

Optogenetic stimulus. The *data[]* field is in unit of watts.

| Dataset | Type | Description | Required |
|---------|------|--|----------|
| site | text | Name of site description in general/optogenetics | Yes |

RoiResponseSeries extends TimeSeries

ROI responses over an imaging plane. Each row in *data[]* should correspond to the signal from one ROI.

| Dataset | Type | Description | Required |
|-----------------------------|---------------------------------|--|----------|
| segmentation_interface | link: Segmentation interface | HDF5 link to image segmentation module defining ROIs | Yes |
| segmentation_interface_path | text | Path to segmentation module | Yes |
| roi_names | text array | List of ROIs represented, one name for each row of <i>data[]</i> | Yes |

SpatialSeries extends TimeSeries

Direction, e.g., of gaze or travel, or position. The *TimeSeries::data* field is a 2D array storing position or direction relative to some reference frame. [Array structure: \[num measurements\] \[num dimensions\]](#). Each SpatialSeries has a text dataset *reference_frame* that indicates the zero-position, or the zero-axes for

direction. For example, if representing gaze direction, “straight-ahead” might be a specific pixel on the monitor, or some other point in space. For position data, the 0,0 point might be the top-left corner of an enclosure, as viewed from the tracking camera. The unit of *data* will indicate how to interpret *SpatialSeries* values. A *SpatialSeries* has all the datasets of a *TimeSeries* plus the following:

| Dataset | Type | Description | Required |
|-----------------|------|--|-------------|
| reference_frame | text | Description defining what exactly “straight-ahead” means | Recommended |

ElectricalSeries extends TimeSeries

Stores acquired voltage data from extracellular recordings. The *data* field of an *ElectricalSeries* is a float array storing data in Volts. [Array structure: \[num time samples\] \[num channels\]](#). It contains all of the datasets of the basic *TimeSeries* as well as the following:

| Dataset | Type | Description | Required |
|---------------|-----------|--|----------|
| electrode_idx | int array | Indices to electrodes described in the experiment's electrode map array (under /general/extracellular_ephys) | Yes |

SpikeEventSeries extends ElectricalSeries

Stores “snapshots” of spike events (i.e., threshold crossings) in *data*. This may also be raw data, as reported by ephys hardware. If so, the *TimeSeries::description* field should describing how events were detected. All *SpikeEventSeries* should reside in a module (under EventWaveform interface) even if the spikes were reported and stored by hardware. All events span the same recording channels and store snapshots of equal duration. [TimeSeries::data array structure: \[num events\] \[num channels\] \[num samples\] \(or \[num events\] \[num samples\] for single electrode\)](#).

PatchClampSeries extends TimeSeries

Stores stimulus or response current or voltage. These are regular *TimeSeries* except for the addition of the following field:

| Dataset | Type | Description | Required |
|----------------|-------|---|-------------|
| electrode_name | text | Name of electrode entry in /general/intracellular_ephys | Yes |
| gain | float | Unit: Volt/Amp | Recommended |

VoltageClampStimulusSeries extends PatchClampSeries

CurrentClampStimulusSeries extends PatchClampSeries

These are aliases to standard *PatchClampSeries*. Their functionality is to better tag *PatchClampSeries* for machine (and human) readability of the file.

VoltageClampSeries extends PatchClampSeries

Stores current data recorded from intracellular voltage-clamp recordings. A corresponding *VoltageClampStimulusSeries* (stored separately as a stimulus) is used to store the voltage injected. The *VoltageClampSeries* has all of the datasets of an *PatchClampSeries* as well as the following:

| Dataset | Type | Description | Required |
|-----------------------------------|-------|--------------|-------------|
| capacitance_fast | float | Unit: Farads | Recommended |
| capacitance_slow | float | Unit: Farads | Recommended |
| resistance_comp_bandwidth | float | Unit: Hz | Recommended |
| resistance_comp_correction | float | Unit: % | Recommended |
| resistance_comp_prediction | float | Unit: % | Recommended |
| whole_cell_capacitance_comp | float | Unit: Farads | Recommended |
| whole_cell_series_resistance_comp | float | Unit: Ohms | Recommended |

CurrentClampSeries extends PatchClampSeries

Stores voltage data recorded from intracellular current-clamp recordings. A corresponding *CurrentClampStimulusSeries* (stored separately as a stimulus) is used to store the current injected. The *CurrentClampSeries* has all of the datasets of an *PatchClampSeries* as well as the following:

| Dataset | Type | Description | Required |
|--------------------------|-------|--------------|-------------|
| bias_current | float | Unit: Amps | Recommended |
| bridge_balance | float | Unit: Ohms | Recommended |
| capacitance_compensation | float | Unit: Farads | Recommended |

IZeroClampSeries extends CurrentClampSeries

Stores recorded voltage data from intracellular recordings when all current and amplifier settings are off (i.e., *CurrentClampSeries* fields will be zero). There is no *CurrentClampStimulusSeries* associated with an *IZero* series because the amplifier is disconnected and no stimulus can reach the cell.

ImageSeries extends TimeSeries

General image data that is common between acquisition and stimulus time series. Sometimes the image data

is stored in the HDF5 file in a raw format while other times it will be stored as an external image file in the host file system. The *data* field will either be binary data or empty. [TimeSeries::data array structure: \[frame\]\[y\]\[x\] or \[frame\]\[z\]\[y\]\[x\]](#). The *ImageSeries* contains all of the datasets of the *TimeSeries* as well as the following:

| Dataset | Type | Description | Required |
|----------------|-----------|---|--|
| format | text | Format of image. If this is “external” then the field <code>external_file</code> contains the path or URL information to that file. For tiff, png, jpg, etc, the binary representation of the image is stored in data. If the format is “raw” then the fields <code>bit_per_pixel</code> and <code>dimension</code> are used. For raw images, only a single channel is stored (eg, red) | Recommended |
| external_file | text | Path or URL to external file, if format = “external”. NOTE: this is only relevant if the image is stored in the file system as an image file. This field should NOT be used if the image is stored in another HDF5 file and that file is HDF5 linked to this file. | No -- only required if image is stored in an external image file |
| bits_per_pixel | int | Number of bits per image pixel | Recommended |
| dimension | int array | Number of pixels on x, y and z axes | Recommended |

ImageMaskSeries extends ImageSeries

An alpha mask that is applied to a presented visual stimulus. The `data[]` array contains an array of mask values that are applied to the displayed image. Mask values are stored as RGBA. Mask can vary with time. The timestamps array indicates the starting time of a mask, and that mask pattern continues until it's explicitly changed.

masked_imageseries (Link:ImageSeries) Link to ImageSeries that mask is applied to

masked_imageseries_path (text) Path to linked ImageSeries

| Dataset | Type | Description | Required |
|-------------------------|-------------------|--|----------|
| masked_imageseries | link: ImageSeries | Link to <i>ImageSeries</i> that mask is applied to | Yes |
| masked_imageseries_path | text | Path to linked <i>ImageSeries</i> | Yes |

TwoPhotonSeries extends ImageSeries

A special case of optical imaging. The *TwoPhotonSeries* has all the datasets of the *ImageSeries* as well as the following:

| Dataset | Type | Description | Required |
|---------------|-------------|---|-------------|
| pmt_gain | float | Photomultiplier gain | Recommended |
| field_of_view | float array | Width, height and depth of image, or imaged area (meters) | Recommended |

| | | | |
|----------------|-------|--|-------------|
| | | | d |
| imaging_plane | text | Name of imaging plane description in /general/optophysiology | Yes |
| scan_line_rate | float | Lines imaged per second. This is also stored in /general/optophysiology but is kept here as it is useful information for analysis, and so good to be stored w/ the actual data | Recommended |

OpticalSeries extends ImageSeries

Image data that is presented or recorded. A stimulus template movie will be stored only as an image. When the image is presented as stimulus, additional data is required, such as field of view (eg, how much of the visual field the image covers, or how what is the area of the target being imaged). If the *OpticalSeries* represents acquired imaging data, orientation is also important. The *OpticalSeries* has all datasets of the *ImageSeries* as well as the following

| Dataset | Type | Description | Required |
|---------------|-------------|---|-------------|
| field_of_view | float array | Width, height and depth of image, or imaged area (meters) | Recommended |
| distance | float | Distance of camera/monitor from target/eye | Recommended |
| orientation | text | Description of image relative to some reference frame (e.g., which way is 'up'). Must also specify frame of reference | Recommended |

Modules

NWB uses modules to store data for and represent the results of common data processing steps, such as spike sorting and image segmentation, that occur before scientific analysis of the data. Modules store the data used by software tools to calculate these intermediate results. Each module provides a list of the data it makes available, and it is free to provide whatever additional data that the module generates. Additional documentation is required for data that goes beyond standard definitions. All modules are stored in the /processing group.

Like *TimeSeries*, *Modules* are each have their own group where they store their data. Each module has the following attributes:

| Attribute | Type | Description | Comments | Required |
|----------------|------------|---|---|----------|
| interfaces | text array | Names of the data interfaces offered by this module. Interface descriptions are below | E.g., [0]="EventDetection", [1]= "Clusters", [2]= "FeatureExtraction" | Yes |
| neurodata_type | text | The string "Module" | | Yes |

Modules have subgroups for each interface that is a part of the module, as well as any additional custom datasets and subgroups that are stored in the module. The *interface* list provides a machine-readable way to identify what information a module provides. Interfaces are not required to publish data and included interfaces can extend beyond what is listed here, so long as the data made available through the new interface is documented.

The *source* field provides information that allows moving backward through the processing stream to find the source of data. Storage of additional provenance information (e.g., parameters, user decisions, etc.) is left to the module's developer, as the types of provenance information can vary greatly between modules.

Module Interfaces

Several module interfaces are listed. When a module provides an interface name in its *interfaces* list, it is required to have the data fields listed with that interface. The data published by each interface should reside in a group in the module having the same name as the interface.

All interfaces have the following attributes:

| Dataset | Type | Description | Required |
|----------------|------|--|----------|
| source | text | Path or description to the origin of the data represented in this module | Yes |
| neurodata_type | text | The string, “interface” | Yes |

BehavioralEvents

BehavioralEpochs

BehavioralTimeSeries

The objective of these interfaces is to provide generic hooks for software tools/scripts. This allows a tool/script to take the output one specific interface (e.g., UnitTimes) and plot that data relative to another data modality (e.g., behavioral events) without having to define all possible modalities in advance. Declaring one of these interfaces means that one or more *TimeSeries* of the specified type is published. These *TimeSeries* should reside in a group having the same name as the interface. For example, if a BehavioralTimeSeries interface is declared, the module will have one or more *TimeSeries* defined in the module sub-group “BehavioralTimeSeries”. BehavioralEpochs should use *IntervalSeries*. BehavioralEvents is used for irregular events. BehavioralTimeSeries is for continuous data.

Clustering

Clustered spike data, whether from automatic clustering tools (e.g., klustakwik) or as a result of manual sorting. A Clustering module publishes the following datasets:

| Dataset/group | Type | Description | Required |
|----------------|--------------|---|----------|
| Clustering/ | group | | Yes |
| _times | double array | Times of clustered events, in seconds. This may be a link to <i>times</i> field in associated FeatureExtraction module. Array structure: [num events] | Yes |
| _num | int array | Cluster number for each event Array structure: [num events] | Yes |
| _description | text | Description of clusters or clustering (e.g., cluster 0 is electrical noise, clusters curated using Klusters, etc) | Yes |
| _cluster_nums | int array | List of cluster numbers that are a part of this set (cluster numbers can be non-continuous) | Yes |
| _peak_over_rms | float array | Maximum ratio of waveform peak to RMS on any channel in the cluster (provides a | Yes |

| | | | |
|--|--|---|--|
| | | basic clustering metric). Array structure: [num clusters] | |
|--|--|---|--|

ClusterWaveforms

The mean waveform shape, including standard deviation, of the different clusters. Ideally, the waveform analysis should be performed on data that is only high-pass filtered. This is a separate module because it is expected to require updating. For example, IMEC probes may require different storage requirements to store/display mean waveforms, requiring a new interface or an extension of this one. A ClusterWaveform module publishes the following datasets:

| Dataset/group | Type | Description | Required |
|-----------------------------|---|--|----------|
| ClusterWaveforms/ | group | | Yes |
| _ waveform_mean | 2D float array | The mean waveform for each cluster, using the same indices for each wave as cluster numbers in the associated Clustering module (i.e, cluster 3 is in array slot [3]). Waveforms corresponding to gaps in cluster sequence should be empty (e.g., zero-filled) | Yes |
| _ waveform_sd | 2D float array | Stdev of waveforms for each cluster, using the same indices as in mean | Yes |
| _ waveform_filtering | text | Filtering applied to data before generating mean/sd | Yes |
| _ clustering_interface | link: <i>Clustering</i> interface | HDF5 link to Clustering interface that was the source of the clustered data | Yes |
| _ clustering_interface_path | text | Path to linked clustering interface | Yes |

CompassDirection

With a CompassDirection interface, a module publishes a *SpatialSeries* object representing a floating point value for theta. The *SpatialSeries::reference_frame* field should indicate what direction corresponds to “0” and which is the direction of rotation (this should be “clockwise”). The *si_unit* for the *SpatialSeries* should be “radians” or “degrees”.

| Dataset/group | Type | Description | Required |
|----------------------------|----------------------|---|----------|
| CompassDirection/ | group | | Yes |
| _ < <i>SpatialSeries</i> > | <i>SpatialSeries</i> | One of possibly many <i>SpatialSeries</i> storing direction. Name should be informative | Yes |

DfOverF

dF/F information about a region of interest (ROI). Storage hierarchy of dF/F should be the same as for segmentation (ie, same names for ROIs and for image planes).

| Dataset/group | Type | Description | Required |
|---------------|------|-------------|----------|
|---------------|------|-------------|----------|

| | | | |
|-----------------------|--------------------------|--|-----|
| DfOverF/ | group | | Yes |
| _ <RoiResponseSeries> | <i>RoiResponseSeries</i> | One of possibly many RoiResponseSeries, one for each imaging plane. Name should match entry in /general/optophysiology | Yes |

EventDetection

Detected spike events from voltage trace(s).

| Dataset/group | Type | Description | Required |
|--------------------------------|-------------------------------|---|----------|
| EventDetection/ | group | | Yes |
| _ times | double array | Timestamps of events, in Seconds | Yes |
| _ detection_method | text | Description of how events were detected, such as voltage threshold, or dV/dT threshold, as well as relevant values. | Yes |
| _ source_idx | int array | Indices into source <i>ElectricalSeries::data</i> array corresponding to time of event. Module description should define what is meant by time of event (e.g., .25msec before action potential peak, zero-crossing time, etc). The index points to each event from the raw data | Yes |
| _ source_electricalseries | link: <i>ElectricalSeries</i> | HDF5 link to <i>ElectricalSeries</i> that this data was calculated from. Metadata about electrodes and their position can be read from that <i>ElectricalSeries</i> so it's not necessary to mandate that information be stored here | Yes |
| _ source_electricalseries_path | text | Path to linked <i>ElectricalSeries</i> | Yes |

EventWaveform

Represents either the waveforms of detected events, as extracted from a raw data trace in /acquisition, or the event waveforms that were stored during experiment acquisition.

| Dataset/group | Type | Description | Required |
|----------------------|-------------------------|--|----------|
| EventWaveform/ | group | | Yes |
| _ <SpikeEventSeries> | <i>SpikeEventSeries</i> | One of possibly many. Name should be informative | Yes |

EyeTracking

Eye-tracking data, representing direction of gaze.

| Dataset/group | Type | Description | Required |
|---------------|------|-------------|----------|
|---------------|------|-------------|----------|

| | | | |
|-------------------|----------------------|--|-----|
| EyeTracking/ | group | | Yes |
| _ <SpatialSeries> | <i>SpatialSeries</i> | One of possibly many SpatialSeries storing direction of gaze | Yes |

FeatureExtraction

Features, such as PC1 and PC2, that are extracted from signals stored in a SpikeEvent *TimeSeries* or other source.

| Dataset/group | Type | Description | Required |
|--------------------|--------------|---|----------|
| FeatureExtraction/ | group | | Yes |
| _ features | float array | Multi-dimensional array of features extracted for each event. Array structure: [num events] [num channels] [num features] | Yes |
| _ times | double array | Times of events that features correspond to (can be a link). Array structure: [num events] | Yes |
| _ description | text array | Description of features (eg, "PC1") for each of the extracted features. Array structure: [num features] | Yes |
| _ electrode_idx | int array | Indices to electrodes described in the experiment's electrode map array (under /general/extracellular_ephys). Array structure: [num channels] | Yes |

FilteredEphys

Ephys data from one or more channels that has been subjected to filtering. Examples of filtered data include Theta and Gamma (LFP has its own interface). FilteredEphys modules publish an *ElectricalSeries* for each filtered channel or set of channels. The name of each *ElectricalSeries* is arbitrary but should be informative. The source of the filtered data, whether this is from analysis of another time series or as acquired by hardware, should be noted in each's *TimeSeries::description* field. There is no assumed 1::1 correspondence between filtered ephys signals and electrodes, as a single signal can apply to many nearby electrodes, and one electrode may have different filtered (e.g., theta and/or gamma) signals represented.

| Dataset/group | Type | Description | Required |
|----------------------|---------------------------|---|----------|
| FilteredEphys/ | group | | Yes |
| _ <ElectricalSeries> | <i>ElectricalSeries</i> s | One of possibly many ElectricalSeries storing DSP filtered data | Yes |

Fluorescence

Fluorescence information about a region of interest (ROI). Storage hierarchy of fluorescence should be the same as for segmentation (ie, same names for ROIs and for image planes).

| Dataset/group | Type | Description | Required |
|---------------|------|-------------|----------|
|---------------|------|-------------|----------|

| | | | |
|-----------------------|-------------------|--|-----|
| Fluorescence/ | group | | Yes |
| _ <RoiResponseSeries> | RoiResponseSeries | One of possibly many RoiResponseSeries, one for each imaging plane. Name should match entry in /general/optophysiology | Yes |

ImageSegmentation

Stores pixels in an image that represent different regions of interest (ROIs) or masks. All segmentation for a given imaging plane is stored together, with storage for multiple imaging planes (masks) supported. Each ROI is stored in its own subgroup, with the ROI group containing both a 2D mask and a list of pixels that make up this mask. Segments can also be used for masking neuropil. If segmentation is allowed to change with time, a new imaging plane (or module) is required and ROI names should remain consistent between them.

| Dataset/group | Type | Description | Required |
|------------------------|----------------|---|-------------|
| ImageSegmentation/ | group | | No |
| _ <imaging_plane> | group | Group name is human-readable description of imaging plane | Yes |
| _ _ description | text | Description of image plane, recording wavelength, depth, etc | Recommended |
| _ _ imaging_plane_name | text | Name of imaging plane under general/optophysiology | Yes |
| _ _ roi_list | text array | List of ROIs in this imaging plane | Yes |
| _ _ <roi_name> | group | Name of ROI | No |
| _ _ _ img_mask | 2D float array | Mask, represented in 2D ([y][x]) intensity image | Yes |
| _ _ _ pix_mask | 2D int array | List of pixels (x,y) that compose the mask | Yes |
| _ _ _ pix_mask_weight | float array | Weight of each pixel listed in pix_mask | Yes |
| _ _ _ roi_description | text | Description of the ROI | Yes |
| _ _ reference_images | group | | Yes |
| _ _ _ <image_name> | ImageSeries | One or more image stacks that the masks apply to (can be one-element stack) | Yes |

Imaging_Retinotopy

Intrinsic signal optical imaging or widefield imaging for measuring retinotopy. Stores orthogonal maps (e.g., altitude/azimuth; radius/theta) of responses to specific stimuli and a combined polarity map from which to identify visual areas.

Note: for data consistency, all images and arrays are stored in the format [row][column] and [row, col], which equates to [y][x]. Field of view and dimension arrays may appear backward (i.e., y before x).

| Dataset/group | Type | Description | Required |
|---------------|------|-------------|----------|
|---------------|------|-------------|----------|

| Imaging_Retinotopy/ | group | | No |
|------------------------|----------------|--|-------------|
| _axis_descriptions | text array | Two-element array describing the contents of the two response axis fields. Description should be something like ['altitude', 'azimuth'] or ['radius', 'theta'] | Yes |
| _axis_1_phase_map | 2D float array | Phase response to stimulus on the first measured axis | Yes |
| __unit (attr) | Text | Unit that axis data is stored in (e.g., degrees) | Yes |
| __dimension (attr) | int array | Number of rows and columns in the image. NOTE: row, column representation is equivalent to height,width | Yes |
| __field_of_view (attr) | float array | Size of viewing area, in meters | Yes |
| _axis_1_power_map | 2D float array | Power response on the first measured axis. Response is scaled so 0.0 is no power in the response and 1.0 is maximum relative power | Recommended |
| __dimension (attr) | int array | Number of rows and columns in the image. NOTE: row, column representation is equivalent to height,width | Yes |
| __field_of_view (attr) | float array | Size of viewing area, in meters | Recommended |
| _axis_2_phase_map | 2D float array | Phase response to stimulus on the second measured axis | Yes |
| __unit (attr) | Text | Unit that axis data is stored in (e.g., degrees) | Yes |
| __dimension (attr) | int array | Number of rows and columns in the image. NOTE: row, column representation is equivalent to height,width | Yes |
| __field_of_view (attr) | float array | Size of viewing area, in meters | Yes |
| _axis_2_power_map | 2D float array | Power response on the second measured axis. Response is scaled so 0.0 is no power in the response and 1.0 is maximum relative power | Recommended |
| __dimension (attr) | int array | Number of rows and columns in the image. NOTE: row, column representation is equivalent to height,width | Yes |
| __field_of_view (attr) | float array | Size of viewing area, in meters | Recommended |
| _sign_map | 2D float array | sine of the angle between the direction of the gradient in axis_1 and axis_2 | No |
| __dimension (attr) | int array | Number of rows and columns in the image. NOTE: row, column representation is equivalent to height,width | Yes |
| _vasculature_image | 2D int array | Gray-scale anatomical image of cortical surface. Array structure: [rows][columns] | Recommended |
| __format (attr) | text | Format of image. Right now only 'raw' supported | Yes |
| __dimension (attr) | int array | Number of rows and columns in the image. NOTE: row, column representation is equivalent to height,width | Yes |

| | | | |
|-------------------------------------|--------------------|--|-------------|
| <code>--bits_per_pixel</code> | <i>int</i> | Number of bits used to represent each value. This is necessary to determine maximum (white) pixel value | Yes |
| <code>--field_of_view (attr)</code> | <i>float array</i> | Size of viewing area, in meters | Recommended |
| <code>_focal_depth_image</code> | 2D int array | Gray-scale image taken with same settings/parameters (e.g., focal depth, wavelength) as data collection. Array format: [rows][columns] | Recommended |
| <code>--format (attr)</code> | <i>text</i> | Format of image. Right now only 'raw' supported. Array format: [rows][columns]. | Yes |
| <code>--dimension (attr)</code> | <i>int array</i> | Number of rows and columns in the image. NOTE: row, column representation is equivalent to height,width | Yes |
| <code>--bits_per_pixel</code> | <i>int</i> | Number of bits used to represent each value. This is necessary to determine maximum (white) pixel value | Yes |
| <code>--focal_depth</code> | <i>float</i> | Focal depth offset, in meters | Recommended |
| <code>--field_of_view (attr)</code> | <i>float array</i> | Size of viewing area, in meters | Recommended |

LFP

LFP data from one or more channels. The electrode map in each published *ElectricalSeries* will identify which channels are providing LFP data. Filter properties should be noted in the *ElectricalSeries* description or comments field.

| Dataset/group | Type | Description | Required |
|---|-------------------------|---|----------|
| LFP/ | group | | No |
| <code>_ <ElectricalSeries></code> | <i>ElectricalSeries</i> | One of possibly many <i>ElectricalSeries</i> storing LFP data | Yes |

MotionCorrection

Publishes an image stack where all frames are shifted (registered) to a common coordinate system, to account for movement and drift between frames.

| Dataset/group | Type | Description | Required |
|---|-----------------------------|--|----------|
| MotionCorrection/ | group | | No |
| <code>_ <image stack name></code> | group | One of possibly many. Name should be informative | Yes |
| <code>--original</code> | link: <i>ImageSeries</i> | Link to image series that is being registered | Yes |
| <code>--xy_translation</code> | <i>TimeSeries</i> | Stores the x,y delta necessary to align each frame to the common coordinates | Yes |

| | | | |
|-------------|--------------------|---|-----|
| __corrected | <i>ImageSeries</i> | Image stack with frames shifted to the common coordinates | Yes |
|-------------|--------------------|---|-----|

Position

Position data, whether along the x, x/y or x/y/z axis.

| Dataset/group | Type | Description | Required |
|----------------------------|----------------------|--|----------|
| Position/ | group | | No |
| _ < <i>SpatialSeries</i> > | <i>SpatialSeries</i> | One of possibly many <i>SpatialSeries</i> storing position | Yes |

PupilTracking

Eye-tracking data, representing pupil size.

| Dataset/group | Type | Description | Required |
|-------------------------|-------------------|---|----------|
| PupilTracking/ | group | | No |
| _ < <i>TimeSeries</i> > | <i>TimeSeries</i> | One of possibly many <i>TimeSeries</i> storing pupil size | Yes |

UnitTimes

Event times in the observed units (eg, cell, synapse, etc). The UnitTimes group contains a group for each unit. The name of the group should match the value in the source module, if that is possible/relevant (e.g., name of ROIs from Segmentation module).

| Dataset/group | Type | Description | Required |
|---------------------|--------------|--|-------------|
| UnitTimes/ | group | | No |
| _ unit_list | text array | List of units present | Yes |
| _ <unit_N>/ | group | | Yes |
| __ times | double array | Spike times for the unit (exact or estimated) | Yes |
| __ source | text | Name, path or description of where unit times originated. This is necessary only if the info here differs from or is more fine-grained than the interface's source field | Recommended |
| __ unit_description | text | Description of the unit (eg, cell type) | Yes |

File Organization

Top-level group: general

General experimental metadata, including animal strain, experimental protocols, experimenter, devices, etc, are stored under 'general'. Core metadata (e.g., that required to interpret data fields) is stored with the data itself, and implicitly defined by the file specification (eg, time is in seconds). The strategy used here for storing non-core metadata is to use free-form text fields, such as would appear in sentences or paragraphs from a Methods section. Metadata fields are text to enable them to be more general, for example to represent ranges instead of numerical values. Machine-readable metadata is stored as attributes to these free-form datasets.

All entries in the below table are to be included when data is present. Unused groups (e.g., intracellular_ephys in an optophysiology experiment) should not be created unless there is data to store within them.

| Datasets & subgroups | Type | Description | Comments |
|------------------------|-------|--|---|
| session_id | text | Lab-specific ID for | Only 1 session_id per file, with all time aligned to experiment start time. |
| experimenter | text | Who performed the experiment | More than one person OK. Can specify roles of different people involved. |
| institution | text | Institution where experiment was performed | |
| lab | text | Lab where experiment was performed | |
| related_publications | text | Publication information. | PMID, DOI, URL, etc. If multiple, concatenate together and describe which is which. |
| notes | text | Notes about the experiment | Things particular to this experiment |
| experiment_description | text | General description of the experiment | Can be from Methods |
| data_collection | text | Notes about data collection and analysis | Can be from Methods |
| stimulus | text | Notes about stimuli, such as how and where presented | Can be from Methods |
| pharmacology | text | Description of drugs used, including how and when they were administered | Anesthesia(s), painkiller(s), etc., plus dosage, concentration, etc. |
| surgery | text | Narrative description of surgery/surgeries, including date(s) and who performed surgery | Much can be copied from Methods |
| virus | text | Information about virus(es) used in experiment, including virus ID, source, date made, injection location and volume, etc. | Content can be from Methods section of paper. |
| slices | text | Description of slice(s) in free-form text, including information about preparation, thickness, orientation, temperature and bath solution. | Content can be from Methods section of paper. |
| protocol | text | Experimental protocol, if applicable | E.g., include IACUC protocol |
| subject | group | | |
| _ subject_id | text | ID of animal/person used/participating in experiment (lab convention) | |
| _ description | text | Description of subject and where subject came from (eg, breeder, if animal) | |
| _ species | text | Species | |
| _ genotype | text | Genetic strain | If absent, assume WT |

| | | | |
|----------|------|--|--|
| _ sex | text | | |
| _ age | text | Age of person, animal, embryo | |
| _ weight | text | Weight at time of experiment, at time of surgery, and at other important times | |

| Datasets & subgroups | Type | Description | Comments |
|-------------------------------|-------------|---|---|
| intracellular_ephys | group | | |
| _ filtering | text | Description of filtering used | Includes filtering type and parameters, frequency fall-off, etc. If this changes between <i>TimeSeries</i> , filter description should be stored as a text attribute for each <i>TimeSeries</i> . |
| _ electrode_X | group | One of possibly many. | Name should be informative. This can optionally be a free-form text field that includes relevant description and metadata. |
| _ _ description (attr) | text | Recording description, description of electrode (eg, whole-cell, sharp) | Free-form text (can be from Methods) |
| _ _ location | text | Area, layer, comments on estimation, stereotaxic coords (if in vivo), etc. | |
| _ _ device | text | Name(s) of device(s) in /general/devices | |
| _ _ slice | text | Name of entry in /general/slices, if relevant | |
| _ _ initial_access_resistance | text | | |
| _ _ seal | text | | |
| _ _ resistance | text | Electrode resistance | unit: Ohm |
| extracellular_ephys | group | | |
| _ electrode_map | float array | Physical location of electrodes (x,y,z, in meters) | Location of electrodes relative to one another. This records the points in space. If an electrode is moved, it needs a new entry in the electrode map for its new location. Otherwise format doesn't support using the same electrode in a new location, or processing spikes pre/post drift. |
| _ electrode_group | text array | Identification string for probe, shank or tetrode each electrode resides on. Name should correspond to one of electrode_group_N groups below. | There's one entry here for each element in electrode_map. All elements in an electrode group should have a functional association, for example all being on the same planar electrode array, or on the same shank. |
| _ impedance | text array | Impedance of electrodes listed in electrode_map | Text, in the event that impedance is stored as range and not a fixed value |
| _ filtering | text | Description of filtering used | Includes filtering type and parameters, frequency fall-off, etc. If this changes between <i>TimeSeries</i> , filter description should be stored as a text attribute for each <i>TimeSeries</i> . |

| Datasets & subgroups | Type | Description | Comments |
|---------------------------|----------------|--|--|
| _electrode_group_X | group | One of possibly many groups, one for each electrode group. If the groups have a hierarchy, such as multiple probes each having multiple shanks, that hierarchy can be mirrored here, using groups for electrode_probe_N and subgroups for electrode_group_N. | Name is arbitrary but should be meaningful. |
| __description | text | Description of probe/shank | |
| __location | text | Description of probe location | E.g., stereotaxic coordinates and other data, e.g., drive placement, angle and orientation and tetrode location in drive and tetrode depth |
| __device | link | Name(s) of device(s) in /general/devices | |
| optophysiology | group | | |
| _imaging_plane_X | group | One of possibly many groups describing an imaging plane | Name is arbitrary but should be meaningful. It is referenced by TwoPhotonSeries and also ImageSegmentation and DfOverF interfaces |
| __manifold | 3D float array | Physical position of each pixel. Array structure: [column][row][xyz] | “xyz” represents the position of the pixel relative to the defined coordinate space |
| ___unit (attr) | text | Base unit that coordinates are stored in (e.g., Meters) | |
| ___conversion (attr) | float | Multiplier to get from stored values to specified unit (e.g., 1e-3 for millimeters) | |
| __reference_frame | text | Describes position and reference frame of manifold based on position of first element in manifold. For example, text description of anatomical location or vectors needed to rotate to common anatomical axis (eg, AP/DV/ML) | This field is necessary to interpret manifold. If manifold is not present then this field is not required |
| __channel_X | group | One of possibly many groups storing channel-specific data | Name is arbitrary but should be meaningful |
| ___emission_lambda | text | Emission lambda for channel | |
| ___description | text | Any notes or comments about the channel | |
| __indicator | text | Calcium indicator | |
| __excitation_lambda | text | Excitation wavelength | |
| __imaging_rate | text | | |

| | | | |
|------------|------|------------------------------------|--|
| __location | text | Location of image plane | |
| __device | text | Name of device in /general/devices | |

| Datasets & subgroups | Type | Description | Comments |
|----------------------|-------|--|---|
| optogenetics | group | | |
| _site_X | group | One of possibly many groups describing an optogenetic stimulation site | Name is arbitrary but should be meaningful. Name is referenced by OptogeneticSeries |
| __description | text | | |
| __device | text | Name of device in /general/devices | |
| __excitation_lambda | text | Excitation wavelength | |
| __location | text | Location of stimulation site | |
| devices | group | Description of hardware devices used during experiment. | Eg, monitors, ADC boards, microscopes, etc |
| _device_X | text | One of possibly many. Information about device and device description. | Name should be informative. Contents can be from Methods. |

Top-level group: acquisition

Acquired data includes tracking and experimental data streams (ie, everything measured from the system)

| Datasets & subgroups | Type | Description | Comments | Required |
|----------------------|--------------------|--|---|-------------|
| timeseries | group | Acquired TimeSeries | When importing acquisition data to an NWB file, all acquisition/tracking/stimulus data must already be aligned to a common time frame. It is assumed that this task has already been performed. | Yes |
| _timeseries_X | generic TimeSeries | One of possibly many acquisition TimeSeries objects. | Name is arbitrary. The TimeSeries::data[] dataset may be a link to raw acquired data in an external file. | No |
| images | group | | | Yes |
| _image_X | binary | Photograph of experiment or experimental setup (video also OK) | Name is arbitrary | No |
| __format (attr) | text | format of image | eg, jpg, png, mpeg | Yes |
| __description (attr) | text | human description of image | If image is of slice data, include slice thickness and orientation, and reference to appropriate entry in /general/slices | Recommended |

If bulky data is stored in the /acquisition group, the data can exist in a separate HDF5 file that is linked to by the file being used for processing and analysis.

When converting data from another format into NWB, there will be times that some data, particularly the raw

data in *acquisition* and *stimulus*, is not included as part of the conversion. In such cases, a *TimeSeries* should be created that represents the missing data, even if the contents of that *TimeSeries* are empty. This helps to interpret the data in the file.

Top-level group: stimulus

Stimuli are here defined as any signal that is pushed into the system as part of the experiment (eg, sound, video, voltage, etc). Many different experiments can use the same stimuli, and stimuli can be re-used during an experiment. The stimulus group is organized so that one version of template stimuli can be stored and these be used multiple times. These templates can exist in the present file or can be HDF5-linked to a remote library file.

| Datasets and subgroups | Type | Description | Comments | Required |
|------------------------|--------------------|---|--|----------|
| templates | group | Template stimuli | Time stamps in templates are based on stimulus design and are relative to the beginning of the stimulus. When templates are used, the stimulus instances must convert presentation times to the experiment's time reference frame. | Yes |
| _timeseries_X | generic TimeSeries | One of possibly many templates, stored as a time series | Name is arbitrary. | No |
| presentation | group | Stimuli presented during the experiment | | Yes |
| _timeseries_X | generic TimeSeries | One of possibly many stimuli, stored as a time series | Name is arbitrary. The TimeSeries::data[] field for the stimulus may be a link to data a stimulus template. | No |

See above, in *acquisition*, about missing *TimeSeries*.

Top-level group: epochs

An experiment can be separated into one or many logical intervals, with the order and duration of these intervals often definable before the experiment starts. In this document, and in the context of NWB, these intervals are called 'epochs'. Epochs have acquisition and stimulus data associated with them, and different epochs can overlap. Examples of epochs are the time when a rat runs around an enclosure or maze as well as intervening sleep sessions; the presentation of a set of visual stimuli to a mouse running on a wheel; or the uninterrupted presentation of current to a patch-clamped cell. Epochs can be limited to the interval of a particular stimulus, or they can span multiple stimuli. Different windows into the same time series can be achieved by including multiple instances of that time series, each with different start/stop times.

| Datasets and subgroups | Type | Description | Comments | Required |
|------------------------|------|---|---|----------|
| epoch_X | | One of possibly many different experimental epoch | Name is arbitrary but must be unique within the experiment. | No |
| _neurodata_type (attr) | text | The string "Epoch" | | Yes |
| _description | text | | | Yes |

| | | | | |
|---------------|------------|--|--|-----|
| _start_time | double | | | Yes |
| _stop_time | double | | | Yes |
| _tags | text array | User-defined tags used throughout the epochs. Tags are to help identify or categorize epochs | E.g., can describe stimulus (if template) or behavioral characteristic (e.g., “lick left”) | Yes |
| _timeseries_X | | One of possibly many input or output streams recorded during epoch | Name is arbitrary and does not have to match the TimeSeries that it refers to | No |
| __idx_start | int | Epoch's start index in TimeSeries data[] field | This can be used to calculate location in TimeSeries timestamp[] field | Yes |
| __count | int | Number of data samples available in this time series, during this epoch | Changed from stop_idx to lessen ambiguity | Yes |
| __timeseries | | link to TimeSeries | | Yes |

The top-level group *epochs* has the following attributes:

| Attribute | Type | Description | Comments | Required |
|-----------|------|---|----------|---------------------------------------|
| links | text | A human-readable list mapping <i>TimeSeries</i> entries in the epoch to the path of the <i>TimeSeries</i> within the file | | Yes, if <i>TimeSeries</i> are present |

Top-level group: processing

'Processing' refers to intermediate analysis of the acquired data to make it more amenable to scientific analysis. These are performed using *Modules*, as defined above. All modules reside in the processing group.

Top-level group: analysis

The file can store lab-specific and custom data analysis without restriction on its form or schema, reducing data formatting restrictions on end users. Such data should be placed in the analysis group. The analysis data should be documented so that it is sharable with other labs.

Extending the format

Extensibility is handled by allowing users to add new datasets to existing *TimeSeries*, *Epochs*, *Modules* and *Interfaces*. What is described in this document is the minimum data that these objects store. The user is free to store additional data as necessary. Documentation should be provided to describe extra data if it is not clear from the context what the data represent.

More extensive modifications, such as defining new *TimeSeries* or *Interfaces*, can be achieved by extending the Specification Language. Generally speaking, language-level extensions will only be necessary for changes that need to be recognized by new software tools. Popular extensions can be proposed and added to the official format specification. Documentation on how to extend the specification itself will be forthcoming.

This approach allows extensibility without breaking backward compatibility.

Acknowledgements

The Neurodata Without Borders: Neurophysiology Initiative is funded by GE, the Allen Institute for Brain Science, the Howard Hughes Medical Institute (HHMI), The Kavli Foundation and the International Neuroinformatics Coordinating Facility. Our founding scientific partners are the Allen Institute, the Svoboda Lab at the Janelia Research Campus of HHMI, the Meister Lab at the California Institute of Technology, the Buzsáki Lab at New York University School of Medicine, and the University of California, Berkeley. Ovation.io is our founding development partner. Ken Harris at University College London provided invaluable input and advice.