# Module Interface Specification for Live Neuro

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## 1 Revision History

Date	Version	Notes
March 17	1.0	Initial Draft

## 2 Symbols, Abbreviations and Acronyms

See SRS Documentation at SRS

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## 3 Introduction

This document specifies the interfaces for modules in the Live Neuro system, an interactive neural data visualization tool. It complements the SRS and MG, with full implementation details available at GitHub Repository.

## 4 Notation

The following table summarizes the primitive data types used by Live Neuro.

Data Type	Notation	Description	
String	S	a character string	
data type dictionary	$\mathbb{D}$	a storage type for KV structures	
bool	$\mathbb{B}$	Boolean data type, which can hold one of two values: either true or false	
integer	$\mathbb{Z}$	a number without a fractional component in $(-\infty, \infty)$	
natural number	N	a number without a fractional component in $[1, \infty)$	
real	$\mathbb{R}$	any number in $(-\infty, \infty)$	

The specification of Live Neuro uses some derived data types: sequences, strings, and tuples. Sequences are lists filled with elements of the same data type. Strings are sequences of characters. Tuples contain a list of values, potentially of different types. In addition, Live Neuro uses functions, which are defined by the data types of their inputs and outputs. Local functions are described by giving their type signature followed by their specification.

## 5 Module Decomposition

The following table is taken directly from the Module Guide document for this project.

Level 1	Level 2	Module ID
Hardware-Hiding Module	Hardware-Hiding Module	M1
Behaviour-Hiding Module	Input Format Module Data Processing Module Visualization Module	M2 M3 M4
Software Decision Module	TRF Calculation Module	M5

Table 1: Module Hierarchy

## 6 MIS of Hardware-Hiding Module(M1)

### 6.1 Module

M1: OS Abstraction Layer

#### 6.2 Uses

Directly interacts with OS APIs (e.g., file I/O, hardware drivers).

## 6.3 Syntax

### 6.3.1 Exported Access Programs

Name	In	Out	Exceptions
readFile	S (file path)	Sequence	${ m File Not Found Error}$
savePlot	plot data	$\mathrm{file}(.\mathrm{jpeg})$	SavePlotError

#### 6.4 Semantics

#### 6.4.1 Environment Variables

File system, display hardware.

### 6.4.2 Assumptions

OS compatibility (Linux/Windows/macOS)

#### 6.4.3 Access Routine Semantics

- readFile(): Reads neural data from disk, returns a sequence of  $\mathbb{R}$  numbers.
- savePlot(): Renders visualization output to screen or file.

## 7 MIS of Input Format Module

#### 7.1 Module

Multi-Format MEG/EEG Data Parser

#### 7.2 Uses

M1 (readFile for raw data loading).

## 7.3 Syntax

#### 7.3.1 Exported Constants

supported formats = [EDF, FIF, BrainVision](supported data formats). max channels = 256(maximum allowed channels per dataset).

#### 7.3.2 Exported Access Programs

Name	In	Out	Exceptions
load edf()	S (EDF file path)	$\mathbb{D}$ < $\mathbb{S}$ : $\mathbb{R}[]$ >	EDFFileError
load fif()	S (FIF file path)	$\mathbb{D} {<} \mathbb{S} : \mathbb{R}[] {>}$	FIFFileError
load brain-	S (.vhdr file path)	$\mathbb{D} < \mathbb{S}: \mathbb{R}[] >$	VHDRFileError
vision()			

#### 7.4 Semantics

#### 7.4.1 State Variables

- currentFormat: S(last detected data format, e.g., "FIF").
- metadataCache: D(cached metadata from parsed files).

#### 7.4.2 Environment Variables

#### 7.4.3 Access Routine Semantics

• load edf():

Output:

data: Time-series array of shape[channels  $\times$  samples].

Exception:

EDFHeaderError: Invalid EDF header structure.

Implementation: Uses eelbrain for EDF parsing.

• load fif():

Output:

data: MEG/EEG sensor data.

Exception:

FIFFileError: FIF file version mismatch or missing data tags.

Implementation: Relies on mne-python library.

• load brainvision():

Output:

data: EEG data segmented by markers

Exception:

VHDRParseError: Inconsistent header fields in.vhdr file.

#### 7.4.4 Local Functions

• parse edf header(): Extracts EDF header fields and validates integrity.

• align brainvision files(): Synchronizes.vhdr,.vmrk, and.eeg data.

## 8 MIS of Data Processing Module

### 8.1 Module

Statistical Preprocessing and Validation

Primary Function: Validates input data integrity and applies statistical preprocessing to neural signals.

#### 8.2 Uses

- M2(Input Format Module): Receives parsed raw data (e.g., MEG/EEG time-series).
- M5(TRF Calculation Module): Provides preprocessed data for dipole current computation.

## 8.3 Syntax

#### 8.3.1 Exported Access Programs

Name	In	Out	Exceptions
validate in-	$\mathbb{D}$ < $\mathbb{S}$ : $\mathbb{R}[]$ >	$\mathbb{B}$	InvalidDataError
put compute statistics	$\mathbb{R}[]$	D <s: r=""></s:>	NaNError

#### 8.4 Semantics

#### 8.4.1 State Variables

• validatedSignals:  $\mathbb{D} < \mathbb{S}$ :  $\mathbb{R}[] > (cached validated data)$ .

• baselineStats:  $\mathbb{D} < \mathbb{S}$ :  $\mathbb{R} >$ (mean and std of reference signals).

#### 8.4.2 Access Routine Semantics

• validate input(raw data:  $\mathbb{D}$ ):

Validation Steps:

Check if raw data data is of type  $\mathbb{R}[]$  and non-empty.

Ensure no NaN or infinite values in the signal.

Output:

Returns True if validation passes.

Exceptions:

InvalidDataError: Non-numeric values or mismatched channel counts.

• compute statistics( $\mathbb{R}[]$ ):

Output:

mean, std, min, max and other statistical information

Exceptions:

NaNError if signal contains invalid values after preprocessing

## 9 MIS of Visualization Module

#### 9.1 Module

Interactive Neuro-imaging Module

#### 9.2 Uses

M1 (writePlot), M3 (processed data), M5 (TRF results)

## 9.3 Syntax

#### 9.3.1 Exported Access Programs

$\mathbf{Name}$	${ m In}$	$\mathbf{Out}$	Exceptions
plot	$\text{List} < \mathbb{R}[] >$	void	Plotting
	(dipole curre	nts),	Error
	List <s>(plot ty</s>	vpe),	
	S(interaction type		
-			

### 9.4 Semantics

#### 9.4.1 State Variables

- activePlots:  $\mathbb{D}$ <PlotID, PlotData> (metadata for open plots).
- linkedViews: Set<PlotID>(plots synchronized via linkPlots).

#### 9.4.2 Environment Variables

GPU acceleration (enabled via plotly.graph\_objects and nilearn.plotting)

#### 9.4.3 Assumptions

#### 9.4.4 Access Routine Semantics

- renderCorticalMap():
  - Output: Generates brain activation map using plotly.graph\_objects and nilearn.plotting
  - Exceptions: Plotting Error

### 9.4.5 Local Functions

• \_updateLinkedAxes(): Propagates axis changes to all linked plots.

## 10 Appendix

## 10.1 Dependencies

- Plotly: Used for interactive HTML5 visualizations (zooming, panning, tooltips).
- Nilearn: Handles neuroimaging-specific rendering (glass brain plots).

## 10.2 Performance Notes

• GPU acceleration is optional but recommended for real-time interaction with large datasets (>10 samples).