

¹ PatchBatch: A Python Tool for Batch Analysis of ² Electrophysiology Data

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Software

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⁵ Summary

⁶ Electrophysiology methods such as patch clamp and two-electrode voltage clamp (TEVC)
⁷ have contributed greatly to our understanding of physiological determinants of disease,
⁸ such as the role of cardiac sodium channel dysfunction in common arrhythmias that can
⁹ cause sudden cardiac events. While traditional tools for data acquisition remain valuable to
¹⁰ electrophysiology researchers, data analysis throughput is limited by workflows that rely on
¹¹ outdated computational frameworks that can be drastically improved through purpose-built
¹² Python automation scripts. PatchBatch was built to standardize and streamline these workflows,
¹³ greatly reducing time expenditure needed for data analysis. This frees up resources for more
¹⁴ important tasks, facilitates greater data output, and makes feasible more exhaustive analysis
¹⁵ strategies for electrophysiology researchers specializing in voltage-clamped experiments.

¹⁶ Statement of Need

¹⁷ PatchBatch is a Python GUI program that provides a standardized data analysis pathway
¹⁸ from data acquisition to data interpretation for patch clamp electrophysiology data. While
¹⁹ programs such as PatchView ([Huang, 2024](#)) and SanPy ([Cudmore, 2024](#)) have been developed
²⁰ to streamline neuroscience and current-clamp data analysis, other electrophysiology domains
²¹ are still lacking such innovations. As a result, many scientists studying current-voltage
²² relationships, concentration-response curves, and ion channel characterizations via patch clamp
²³ or TEVC are frequently using tedious, outdated data processing schemas. Typical analysis of
²⁴ electrophysiology data through existing platforms is often repetitive, yet algorithmic, opening
²⁵ the opportunity for automation. It requires manually extracting the desired data by configuring
²⁶ analysis parameters and axis selection one file at a time (which are often the same for each
²⁷ file). In addition to having a high time cost, this process also invites user error resulting from
²⁸ tedious actions that must be repeated for each file. For one scientist, a day's patch clamp
²⁹ data can take an hour or more to process into a meaningful form. PatchBatch automates
³⁰ these repetitive data processing steps by enabling batch analysis of multiple data files at once
³¹ with the same parameters, cutting analysis time to minutes rather than hours. It can also
³² extract individual data sweeps from one or several files at a time, expediting sweep analyses
³³ that similarly are very tedious to perform manually. For example, one can extract the +100mV
³⁴ sweep of a full dataset from I-V relationship experiments in order to compare current waveforms
³⁵ measured at +100mV from multiple conditions, enabling rapid quantification of waveform
³⁶ changes associated with different conditions. Batch sweep export facilitates quick and easy
³⁷ plotting of multiple conditions for comparison.

³⁸ PatchBatch was built primarily for .wcp files output by WinWCP ([Dempster, 2024](#)), and it
³⁹ seeks to replicate many of its data analysis functionalities, leveraging Python to automate
⁴⁰ a subset of analysis workflows performed by many electrophysiologists. It is also compatible
⁴¹ with .abf formatted data files via pyABF ([Harden, 2022](#)). This program is intended to be

42 usable by anyone who needs to analyze patch clamp or TEVC data, while not requiring any
43 coding/programming knowledge. Analysis results can be copied or exported to a CSV file for
44 downstream analysis in a dedicated data visualization tool such as Excel or Graphpad Prism.
45 PatchBatch is intended to be an intermediate tool that serves as a quick pipeline from raw to
46 processed data, ready for drop-in integration into a separate application for publication-ready
47 figures.

48 The aforementioned WinWCP, as well as Clampfit([Molecular Devices, 2024](#)), remain valuable
49 tools capable of extended data analysis methods, many of which (such as action potential
50 detection) are not currently replicated in this program. PatchBatch provides a streamlined
51 analysis pathway for dual-channel (Current channel and Voltage channel) data containing
52 repeating Voltage/Current “sweeps”, from which scientists extract average or peak signals
53 from a specified temporal region in each sweep. WinWCP and Clampfit lack such a streamlined
54 analysis pathway, necessitating long, manual analysis processes that this program automates.
55 This program also adds further data processing capabilities in the form of Current Density
56 analysis, which simplifies the process of converting raw current values to current densities for
57 whole-cell recordings. By facilitating the rapid processing of raw data, PatchBatch builds on
58 previous software to provide a convenient platform for electrophysiologists to standardize and
59 streamline their data analysis protocols. This not only saves time, but enhances reproducibility
60 of analyses and supports more exhaustive data processing workflows.

61 **AI Usage Disclosure**

62 This software was developed with the aid of large language model (LLM) tools. ChatGPT-4o
63 and Gemini 2.5 were used during early development to produce a basic working prototype.
64 Claude Sonnet 4.0–4.5 and Opus 4.0–4.5 (Anthropic) were used in later development to
65 generate code and refactor the prototype into a user-friendly, fully functional software. All user-
66 facing design decisions were made by the author based on domain expertise in electrophysiology.
67 LLMs aided in the design of the data processing architecture, fine-tuning of the desired analysis
68 modes and GUI features, and troubleshooting coding errors. All code outputs were reviewed
69 and refined by the author for data operations integrity and overall functionality. All analysis
70 scripts were extensively validated by the author by comparing analysis outputs against known
71 reference values (Figure 3).

72 LLMs were used to develop pytest scripts to validate all data workflows. Pytests were validated
73 for accurate functionality by the author. Likewise, LLMs were used to script Continuous
74 Integration to run all pytests, and later cross-platform builds, upon each push to GitHub. All
75 build scripts (build_windows.py, build_macos.py, and pyproject.toml) were developed with
76 LLM assistance. Boilerplate documentation (BUILD-INSTRUCTIONS.md, CITATION.cff,
77 CODE_OF_CONDUCT.md, LICENSE.md) were generated with LLM assistance with light
78 refinement by the author. The design principles of ARCHITECTURE.md were written by the
79 author, while the service layer overview was guided by LLM assistance to comprehensively
80 describe the architecture that performs the core analysis workflow. All text of the present
81 document and the vast majority of README.md was written by the author and formatted
82 with LLM assistance.

83 Figures

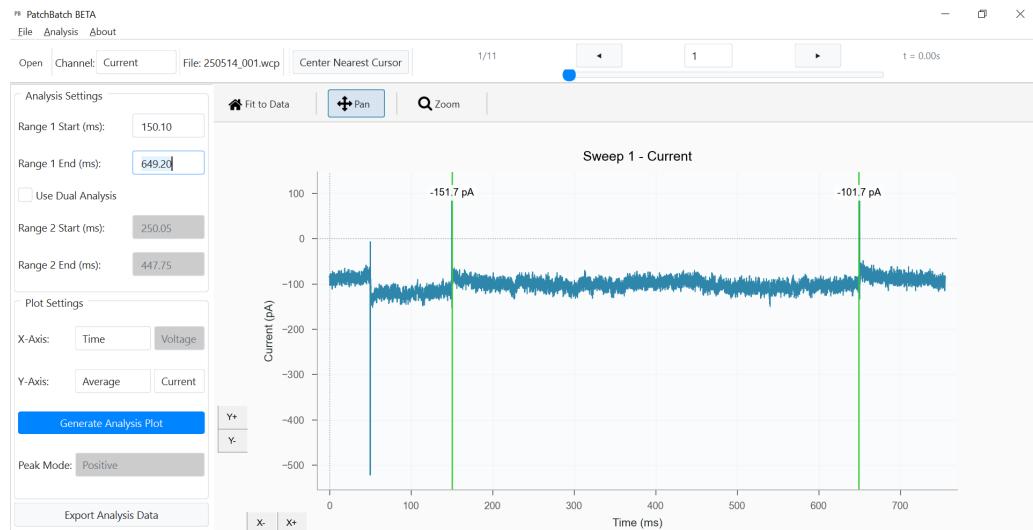


Figure 1: Entry point of PatchBatch.

84 Initial window where data sweeps are displayed and analysis ranges defined. Analysis parameters
 85 are entered on left-side control panel. The main plot features dragable cursors whose positions
 86 are coupled to the input fields in the control panel, enabling fine-tuning of analysis parameters.
 87 Users may select one or two time ranges per analysis as needed.

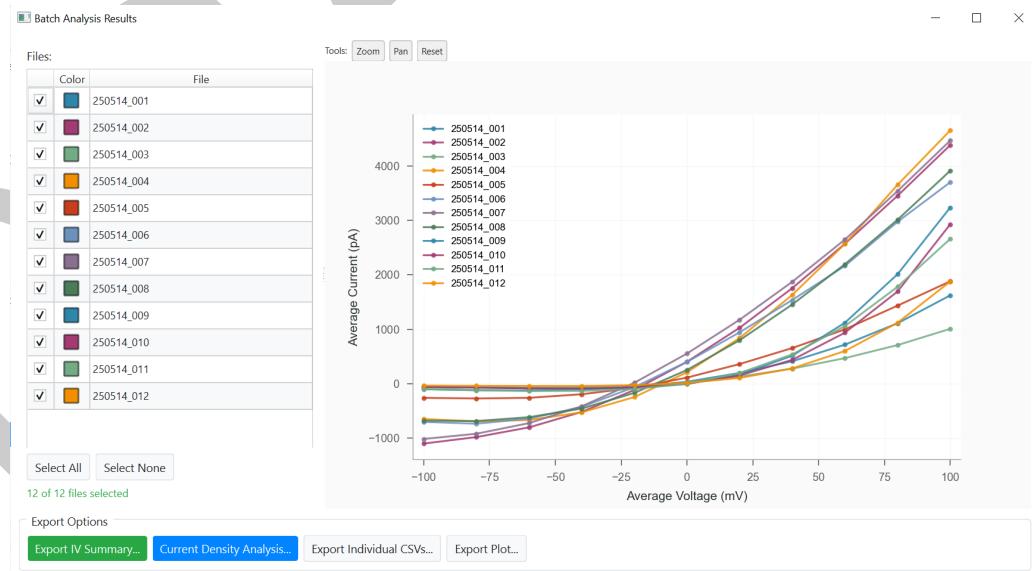


Figure 2: Batch Analysis window.

88 Batch analysis enables rapid output of full dataset, each analyzed by identical parameters.
 89 Batch-analyzed data may be exported as a single summary CSV file or several individual CSV
 90 files. Data may also be copied to clipboard directly from the program.

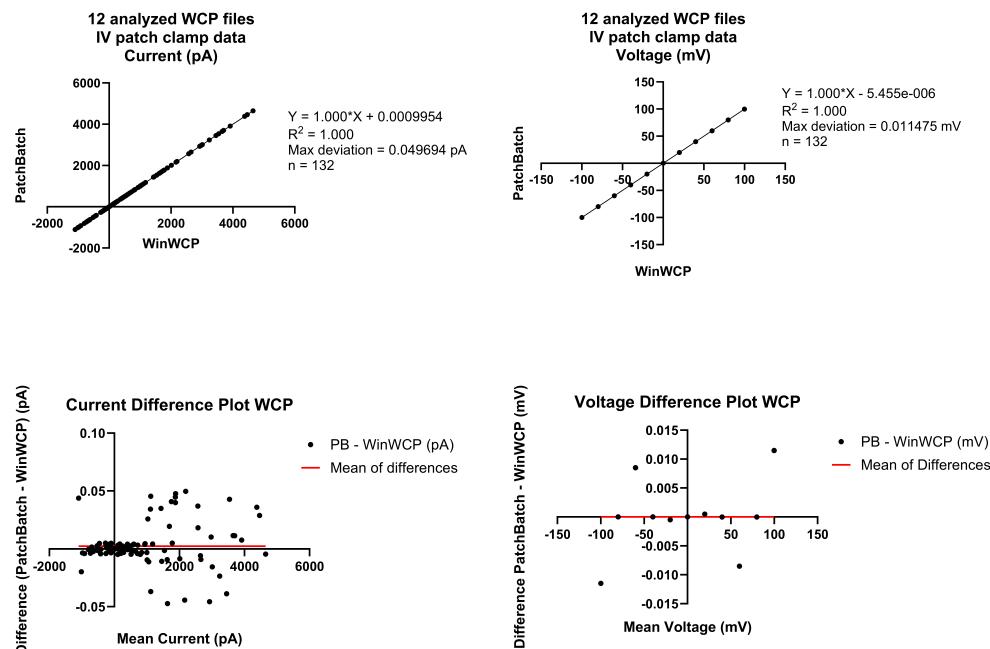


Figure 3: Validation of data operations.

91 Analysis outputs match those of WinWCP. A set of 12 files was analyzed in PatchBatch and
92 in WinWCP, each using identical analysis parameters. Largest discrepancy was 0.05 pA and
93 0.01 mV for Average Current and Average Voltage measurements respectively.

94 References

- 95 Cudmore, R. H. (2024). SanPy: Software for the analysis of electrophysiology data. In *GitHub*
96 *repository*. GitHub. <https://github.com/cudmore/SanPy>
- 97 Dempster, J. (2024). WinWCP: Whole cell program for patch clamp electrophysiology. In
98 *GitHub repository*. GitHub. <https://github.com/johndempster/WinWCPX>
- 99 Harden, S. W. (2022). pyABF (Version 2.3.5). <https://pypi.org/project/pyabf>
- 100 Huang, Z. (2024). PatchView: A python-based GUI for analysis of whole-cell patch clamp
101 recording data. In *GitHub repository*. GitHub. <https://github.com/ZeitgeberH/patchview>
- 102 Molecular Devices. (2024). Clampfit 11 software: pCLAMP software suite. Molecular Devices.
103 [https://www.moleculardevices.com/products/axon-patch-clamp-system/acquisition-and-](https://www.moleculardevices.com/products/axon-patch-clamp-system/acquisition-and-analysis-software/pclamp-software-suite)
104 [analysis-software/pclamp-software-suite](https://www.moleculardevices.com/products/axon-patch-clamp-system/acquisition-and-analysis-software/pclamp-software-suite)