

# epyc: Computational experiment management in Python

## Simon Dobson<sup>1</sup>

1 School of Computer Science, University of St Andrews, Scotland UK

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#### **Software**

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# Summary

epyc is a Python module for designing, executing, storing, and analysing the results of large sets of (possibly long-running) computational experiments, as are often found when writing simulations of complex networks and other domains. It allows the same experimental code to be run on single machines, multicore machines, and computational clusters without modification, and automatically manages the execution of an experiment for different parameter values and for multiple repetitions.

## Statement of need

Computation and simulation have joined theory and experiment as a third component of many scientific fields. Computational experiments share many features with traditional experiments: they need to be structured, repeated, repeatable, reproducible, and interpretable (Software Sustainability Institute, 2020). There is often a need to perform a computation at a number of points across a multi-dimensional parameter space, aggregating multiple repetitions and wrangling results for analysis and presentation. Often the experiments being performed are on such a scale as to require the use of multicore machines and/or computing clusters to perform multiple experiments simultaneously. The logistics of managing this process can become quite complicated, and benefits from automation.

epyc defines a conceptual framework for performing series of computational experiments, and implements this in pure Python. Computational "laboratories" perform a collection of "experiments" whose parameters and results are recorded in a "lab notebook" for later retrieval. Laboratories take a parameter space consisting of any number of individual parameters having either single values or ranges, and use a computational experimental "design" to determine which combinations are used in running experiments.

### Main features

Modern research software has to operate in a range of computational environments and at a range of scales. epyc is fully portable to any Python environment, and can operate across the scales. Laboratories can be sequential (a single core on a machine), parallel (multiple cores on a machine), or clustered (multiple cores on multiple machines). A key design goal of epyc is to allow the same experimental code to run under all these configurations, to maximise the re-use of experiments. Cluster-based laboratories can be accessed asynchronously, allowing a set of experiments to be started and their results retrieved later without remaining connected to the cluster: useful for laptop users. epyc also integrates fully into the Jupyter ecosystem, allowing notebook-based experimental protocols to still use compute clusters.

Experimental design is a neglected feature of computational science, despite its strong presence in traditional settings (Bailey, 2009). The choice of design translates an experiment and a space of possible parameters into a collection of experimental runs at selected points in that



parameter space. Typically choices are to use every combination of parameters (a factorial design) or to take corresponding values from each parameter range (a pointwise design): other designs can easily be added.

epyc defines a lifecycle for experiments through which they are configured to their parameters, set-up, run, and torn-down, before returning the experimental results alongside the experimental parameters and some metadata on the execution of the run, to be inserted into a notebook. A small set of experiment combinators separate the logic of a single experiment from structuring and other tasks. This means that any experiment can be repeated and statistically summarised as a single unit, and also encourages the development of re-usable computational analytics rather than manual handling that can be harder to reproduce.

Lab notebooks can be persistent to store the results of experiments in a non-destructive form, structured as individual result sets with notes and metadata. Notebooks store all the data and metadata in a portable format to improve the reproducibility of computational experiments. Results can be accessed through pandas DataFrames for easy wrangling and visualisation. Lab notebooks store their data in HDF5, a standard format for large-scale archival storage that augments raw data with annotations and metadata, as well as making results obtained in epyc easily portable to other tools.

# Differences with other tools

epyc is designed to manage experiments rather than provide sophisticated workflows, since for many application such workflows are largely unnecessary. Instead, epyc defines a conceptual model of computational experiments (laboratories, designs, experiments, and notebooks) and supports each component as flexibly and orthogonally as possible.

Unlike Dask, epyc works independently of libraries such as numpy or scikit-learn (although it can run experiments using these tools). It also adopts a control- rather than data-driven approach (also unlike Dask, and unlike more traditional workflow engines such as Taverna or SnakeMake) which leads to more predictable computational requirements and fits better with the approaches of certain disciplines.

epyc is perhaps closest in spirit to Sacred, which also emphasises the management of experimental configurations for reproducibility. epyc provides more support for executing experiments uniformly across entire spaces of parameters, with integrated use of parallelism, and under different experimental designs where required.

# Main applications

epyc arose as part of an on-going research programme in network science and complex systems, as a way to manage the thousands of repeated simulations needing to be performed across large parameter spaces. It provides experiment management for the epydemic simulation framework for complex networks, from which it has contributed to a series of papers exploring networks and their processes (for example Mann et al. (2021a), and a book on network-based disease modelling (Dobson, 2020). While originally closely integrated into this system, epyc is now a stand-alone domain-independent experimental management framework.

# Compatibility and availability

epyc works with versions of Python from Python 3.6, and can be installed directly from PyPy using pip. API and tutorial documentation can be found on ReadTheDocs. Source code is available on GitHub, where issues can also be reported.



## References

- Bailey, R. A. (2009). *Design of comparative experiments*. Cambridge University Press. https://doi.org/10.1017/CBO9780511611483
- Dobson, S. (2020). *Epidemic modelling some notes, maths, and code*. Independent Publishing Network. ISBN: 978-183853-565-0
- Mann, P., Smith, V. A., Mitchell, J., & Dobson, S. (2021a). Symbiotic and antagonistic disease dynamics on clustered networks using bond percolation. *Physical Review E*, 104(2). https://doi.org/10.1103/PhysRevE.104.024303
- Mann, P., Smith, V. A., Mitchell, J., & Dobson, S. (2021b). Two-pathogen model with competition on clustered networks. *Physical Review E*, 103(6). https://doi.org/10.1103/PhysRevE.103.062308
- Pitcher, M., Bowness, R., Dobson, S., Eftimie, R., & Gillespie, S. (2020). Modelling the effects of environmental heterogeneity within the lung on the tuberculosis lifecycle. *Journal of Theoretical Biology*, 506. https://doi.org/10.1101/2019.12.12.871269
- Software Sustainability Institute. (2020). *Guides for researchers*. https://www.software.ac.uk/resources/guides/guides-researchers