

- MultiSim: A Python Toolbox for Simulating Datasets
 with time resolved multivariate effects
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Summary

In MEG/EEG research, validating analysis pipelines is hampered by the lack of ground-truth neural signals in real data. SimMEG fills this gap by generating realistic, time-locked multivariate effects of known magnitude that you can inject into simulated sensor data. You can then run any pipeline—e.g. decoding, sensor-level statistics, or source estimation—against these datasets to benchmark sensitivity and specificity.

Key benefits include:

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- Testing whether your pipeline reliably detects effects of a chosen size.
- Providing demonstrable, reproducible benchmarks for reviewers or collaborators.
- Offering a controlled teaching environment for newcomers.

Below, we describe the rationale (Statement of needs), and the data-generation method (Methods), a hands-on example (Results), and potential extensions (Discussion).

Statement of needs

Multivariate pattern analysis (MVPA) is now routine in cognitive neuroscience for probing how the brain represents information (Haxby et al., 2001; Haynes & Rees, 2006; Kriegeskorte et al., 2008; Poldrack et al., 2009; Ritchie et al., 2019). Applied to high-temporal-resolution electrophysiology signals such as electro and magneto-encephalography (EEG and MEG respectively), decoding techniques reveal the millisecond-by-millisecond unfolding of mental representations (Cichy et al., 2014; Consortium et al., 2025; King et al., 2016; King & Dehaene, 2014; Kok et al., 2017). Strikingly, despite the ubiquity of MVPA techniques, to our knowledge, no method exists to test the sensitivity and specificity of decoding analysis pipelines, nor to estimate, before data collection, how many trials and how many participants are required to detect an effect of a given size.

40 MultiSim addresses this gap by letting investigators simulate time-resolved multi-channel signals



with paramaters matching that of their recording setups, and specify multivariate effects with known timing, spatialization and strength, while controlling channel covariance, sensory noise and between subjects variability (see code block below for a minimal working example and Figure 1 for a visualization of the pipeline). Our algorithm produces multi-subject data sets in which ground truth effects are known. By running their pipeline on these data, researchers obtain a direct read-out of its true-positive rate (can it recover the injected effects?) and false-positive rate (does it raise alarms when nothing is present). In addition, our simulator can be used to perform computational power analysis, to determine the number of trials and subjects, by iterating over these parameters.

This toolbox promotes best-practice MVPA by giving researchers a tailored benchmark for their specific experimental designs, a testbed for developing new decoding methods, and a principled way to check that planned studies are properly powered—ultimately enabling more reliable and efficient investigations of brain function.



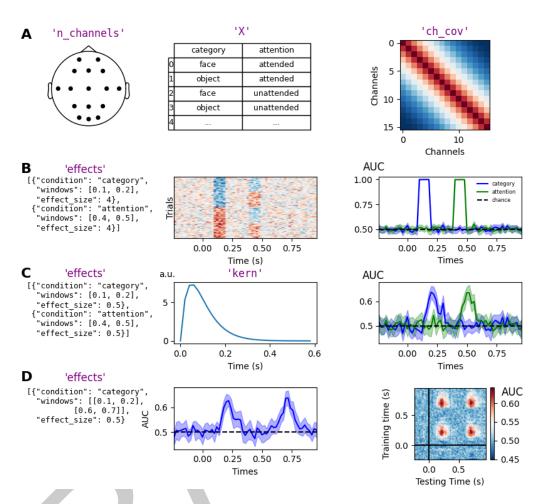


Figure 1: Figure 1. Overview of the simulation and decoding framework. A. General data parameters for the simulation. Left: n_channels corresponds to the number of channels in the montage, with 2 experimental conditions. Middle: X is the design matrix (each column is an experimental condition and each row a trial). Right: ch_cov is the channel by channel covariance matrix of the data to be simulated. B. Minimal simulation example with effects for each experimental condition with large effect size. Left: effects is a dictionary that specifies the "condition", time window ("windows") and effect size ("effect_size") of each effect to simulate. The example specifies an effect of category from 0.1 and 0.2 s with an effect size of 4, and an effect of the attention condition from 0.4 to 0.5 with an effect size of 4. Middle: example of the activation of a single channel. Right: resulting decoding accuracy (using a Support vector machine classifier). C. Example of simulated effects with a an added gamma kernel to simulate effects with biologically plausible temporal dynamcs. Left: effects similar to that of B but with effect size of 0.5 for each condition. Middle: gamma kernel, specifying the temporal dynamics of the multivariate effect. Right: Resulting decoding accuracy. D. example of simulated data with cross temporal generalization of the category effect. Left: effects dictionary specifies two different time windows for the effect of category as a list. Middle: resulting decoding accuracy. Right: Temporal generalization of the decoding.

Code Quality and Documentation

SimMEG is hosted on GitHub. Examples and API documentation are available on the platform here. We provide installation guides, algorithm introductions, and examples of using the package with Jupyter Notebook. We further provide the full mathematical details of our simulation here. The package is available on Linux, macOS and Windows for Python >=3.12 It can be installed with pip install simMEG. To ensure high code quality, all implementations adhere to the PEP8 code style [REF], enforced by ruff [REF], the code formatter black and



- 61 the static analyzer prospector. The documentation is provided through docstrings using the
- 62 NumPy conventions and build using Sphinx.

Acknowledgements

64 References

65 {bibliography}

5 Supplementary

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