

anesthetic: nested sampling visualisation

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Software

■ Review 🖸

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Summary

anesthetic is a Python package for processing nested sampling runs, and will be useful for any scientist or statistician who uses nested sampling software. anesthetic unifies many existing tools and techniques in an extensible framework which is intuitive for users familiar with the standard Python packages, namely NumPy, SciPy, Matplotlib and pandas. It has been extensively used in the recent cosmological papers (W. Handley & Lemos, 2019a, 2019b).

Nested sampling

Nested sampling (Skilling, 2006) is an alternative to Markov-Chain-Monte-Carlo techniques (Hastings, 1970). Given some data D, for a scientific model M with free parameters θ , Bayes theorem states:

$$P(\theta|D) = \frac{P(D|\theta)P(\theta)}{P(D)}.$$

Traditional MCMC approaches ignore the Bayesian evidence P(D) and instead focus on the problem of generating samples from the posterior $P(\theta|D)$ using knowledge of the prior $P(\theta)$ and likelihood $P(D|\theta)$. Nested sampling reverses this priority, and instead computes the evidence P(D) (the critical quantity in Bayesian model comparison (Trotta, 2008)), producing posterior samples as a by-product. Nested sampling does this by evolving a set of live points drawn from the prior under a hard likelihood constraint which steadily increases, causing the live points to contract around the peak(s) of the likelihood. The history of the live-point evolution can be used to reconstruct both the evidence and posterior samples, as well as the density of states and consequently the full partition function.

Current publicly available implementations of nested sampling include MultiNest (Feroz, Hobson, & Bridges, 2009), PolyChord (W. J. Handley, Hobson, & Lasenby, 2015a, 2015b)(Higson, 2018), DNest (Brewer & Foreman-Mackey, 2018) and dynesty (Speagle, 2019), all of which have been incorporated into a wide range of cosmological (Brinckmann & Lesgourgues, 2019; Antony Lewis & Bridle, 2002; Zuntz et al., 2015) and particle physics (The GAMBIT Scanner Workgroup:, 2017) codes.



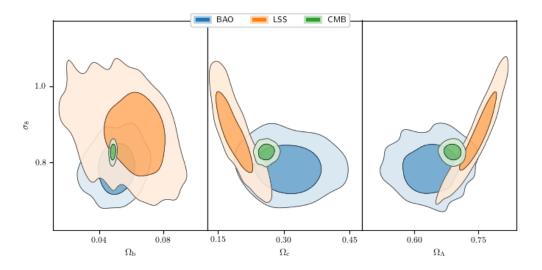


Figure 1: Marginalised posterior plots produced by anesthetic. The x axes indicate the fraction of normal matter, dark matter and dark energy respectively, whilst the y-axis is the amplitude of mass fluctuation in our late-time universe. The three measurements were performed using measurements of baryonic acoustic oscillations, large scale structure and the cosmic microwave background (W. Handley & Lemos, 2019a). It is an open cosmological and statistical questions whether the LSS and CMB are consistent with one another.

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anesthetic acts on outputs of nested sampling software packages. It can:

- 1. Compute inferences of the Bayesian evidence (Trotta, 2008), the Kullback-Leibler divergence (Kullback & Leibler, 1951) of the distribution, the Bayesian model dimensionality (W. Handley & Lemos, 2019b) and the full partition function.
- 2. Dynamically replay nested sampling runs.
- 3. Produce one- and two-dimensional marginalised posterior plots (Figure 1).

A subset of computations from item 1 is provided by many of the nested sampling software packages. anesthetic allows you to compute these independently and more accurately, providing a unified set of outputs and separating these computations from the generation of nested samples.

Item 2 is useful for users that have experienced the phenomenon of 'live point watching' – the process of continually examining the evolution of the live points as the run progresses in an attempt to diagnose problems in likelihood and/or sampling implementations. The GUI provided allows users to fully reconstruct the run at any iteration, and examine the effect of dynamically adjusting the thermodynamic temperature.

Finally, it is important to recognise that the functionality from item 3 is also provided by many other high-quality software packages, such as getdist (Anthony Lewis, 2015), corner (Foreman-Mackey, 2016), pygtc (Bocquet & Carter, 2016), dynesty (Speagle, 2019) and MontePython (Brinckmann & Lesgourgues, 2019). anesthetic adds to this functionality by:

• Performing kernel density estimation using the state-of-the-art fastkde (T. A. O'Brien, Kashinath, Cavanaugh, Collins, & O'Brien, 2016) algorithm.



- Storing samples and plotting grids as a weighted pandas. DataFrame, which is more consistent with the scientific Python canon, allows for unambiguous access to samples and plots via their reference names, and easy definition of new parameters.
- Using a contour colour scheme that is better suited to plotting distributions with uniform probability, which is important if one wishes to plot priors along with posteriors.

The source code for anesthetic is available on GitHub, with its automatically generated documentation at ReadTheDocs and a pip-installable package on PyPi. An example interactive Jupyter notebook is given using Binder (Jupyter et al., 2018). Continuous integration is implemented with Travis and Circle.

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