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Clustering Considerations for Nested Sampling

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PolyChord was originally advertised encouraging users to experiment with their own clustering algorithms. Identifying clusters of nested sampling live points is critical for PolyChord to perform nested sampling correctly. We have updated the Python interface of PolyChordLite to allow straightforward substitution of different clustering methods.

Recent reconstructions of the primordial matter power spectrum $\mathcal{P}_{\mathcal{R}}(k)$ with a *flex-knot* revealed that the K-Nearest-Neighbours algorithm used by PolyChord cannot reliably detect the two posterior modes caused by cosmic variance and detector resolution. After exploring a number of different algorithms, we have found the X-means algorithm to be a reliable substitute for the power spectrum reconstruction.

This work prompted the development of additions to the post-processing tool *anesthetic*, allowing posterior modes corresponding to different live point clusters to be analysed and plotted independently.

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