# Anomaly detection for dark matter direct detection

**Aim**: Use **event-by-event** prompt readout to train anomaly detector





# Semi-unsupervised pipeline: Combined anor events

Non-background events  $\rightarrow$  Higher TS

Semi-unsupervised pipeline: Combined anomaly function leverages discrete nature of detector

# What we get

VAE learns event energy...



# **Establish 1D two-sample test**

- No nuisance parameters
- Spectral info encoded
- No dependence on high level summary statistics (cS1/cS2)
- Trivially implement calibration data directly 0
- Likelihood free and model independent









# Clustering Considerations for Nested Sampling

# 5.3.1 Cluster recognition

Any cluster recognition algorithm can be substituted at this point. One must take care that this is not run too often, or one runs the risk of adding a large overhead to the calculation. In practice, checking for clustering every  $\sim O(n_{\text{live}})$  iterations is sufficient, since the prior will have only compressed by a factor *e*. We encourage users of POLYCHORD to experiment with their own preferred cluster recognition, in addition to that provided and described below.

"PolyChord: next-generation nested sampling" arXiv:1506.00171 Adam Ormondroyd University of Cambridge Kavli Institute for Cosmology





# Motive: primordial matter power spectrum flex-knot reconstruction



# Clustering choices are abundant!







# Clustering Considerations for Nested Sampling Adam Ormondroyd ano23@cam.ac.uk

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Clustering algorithms are integral to multi-modal nested sampling. for both region-based samplers such as MultiNest, and chain-based samplers such as PolyChord. Robust identification clusters of live points is crucial for effective spawning of new live points, prior volume estimation and therefore the total evidence calculation. Reliable cluster detection also allows the calculation of the sub-evidences of each cluster, which may correspond to different physical phenomena. We have explored extensions to the clustering approach within PolyChord, and found that including correlation between the volume estimates of clusters increases the accuracy of evidence calculations. We show how different clustering methods affect a reconstruction of the cosmological primordial matter power spectrum  $P_R(k)$ .

### **Clustering choice**



The nested sampling [1, 2] algorithm Poly-Chord was originally advertised suggesting that the user should experiment with their favourite cluster-identifying methods, but provides no guidance on how to do so [3, 4]. We have added an interface which allows the user to substitute any clustering strategy at the Python level, allowing for easier experimentation with alternatives such as the selection provided by scikit-learn and pyclustering [5,6]. Some algorithms are better suited than others

to identifying posterior modes of nested sampling live points, for example K-means and spectral clustering need to be told the number of clusters to look for, others may not assign every point to a cluster. Some approaches find clusters where there are none!

Examples of  $\mathcal{P}_{\mathcal{R}}(k)$  flexknots

with three knots each.

Prototype *L*-means

algorithm

-means with X+2 the likelihood at the centres -

## Application to cosmology $\mathcal{P}_{\mathcal{R}}(k)$

This investigation was initially motivated by a reconstruction of the primordial matter power spectrum  $\mathcal{P}_{\mathcal{R}}(k)$  using flexknots [7, 8], and the Planck 2018 likelihoods [9, 10]. Planck measured the  $C_{\ell}$  multipole range  $1 \leq \ell \leq 7000$ , corresponding to  $10^{-4} \le k/\text{Mpc}^{-1} \le 10^{-0.3}$  [11]. Flexknots are parameterisations of 1D functions, consisting of a series of splines (in this case linear) joined at knots. The number and positions of knots are determined by the data, which can be performed by either combining several runs with fixed number of knots, or the number of knots being itself a parameter. In the former case, we noticed that with three knots only the mode with the central knot towards the left was being fully explored. PolyChord's native K-nearest-neighbours clustering was unable to separate the positions of the central knot into two distinct clusters, so we explored both off-the-shelf clustering algorithms and an approach which includes likelihood information, L-means.







K-nearest-neighbours and DBSCAN fail to separate the central knot into two clusters, while X-means does so reliably. L-means is able to separate the central knot into multiple clusters, but also tends to over-cluster. Functional posterior plots were created using fgivenx [12] and scatterplots showing clusters were created using a development branch of anesthetic

### **Classes of Nested Sampler**

Region-based

+1+1



MultiNest PolyChord first uses the covariance maconstructs a series of ellipsoids [15-17] trix of the live points to whiten the space. These regions are usually expanded by a then performs Neal's slice sampling along numerical factor to improve their chances orthogonal directions in that space [3, 4] of fully enclosing the likelihood contour, GGNS (gradient-guided nested sampling) then a point is rejection-sampled from also implements both Neal's and Hamilthem. The curse of dimensionality means tonian slice sampling, along with uniform that these techniques are only effective up sampling and random walks [14]. to O(10) dimensions, as either rejection Multiple-modal problems render contour sampling becomes inefficient, or the expanwhitening ineffective, and a strategy is resion factor would have to be so low that quired to decide from which mode to sample significant regions of the likelihood contour since a random walk cannot pass through would be missed. Clustering can be used the likelihood contour to separate the live points into discrete regions, rather than a single sparse region.

Hybrid methods combine the two approaches in an attempt to alleviate the dimensionality scaling of region-based methods, while reducing the number of likelihood evaluations made outside the contour [14, 18-20].

### Correlated cluster volumes (in progress!)

When a cluster p is divided, the remaining prior volume  $X_p$  is divided among its subclusters X<sub>i</sub>. However, in nested sampling, we do not know the precise prior volumes, only expectation values and errors. This is divided according to the proportion of live points in each cluster n:



and the second second

Since  $\overline{X_iX_i} \neq \overline{X_i}$ , the error on the prior volumes estimates of each cluster are correlated. This is important when deciding from which cluster to sample: currently PolyChord chooses a cluster propotionally to its prior volume  $\overline{X_i}$ , but neglects their correlation. We are experimenting with drawing a set of  $X_i$  from their joint distribution before each live point is generated, which has shown promise with symmetric multi-modal likelihoods.

	References
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X-means and scikit-learn's

(H)DBSCAN

(ICC

# Enhancing Robustness

BSM Parameter Inference with n1D-CNN and Novel Data Augmentation

# Koay Yong Sheng

Stefano Moretti, Prashant Singh, Harri Waltari

<u>Keywords:</u>

- Parameter Inference
- Improving CNNs
- Data Augmentation
- Signal Region Evaluation



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# Given the following images, what is x<sub>signal</sub>?



# Impossible! No scale of reference! Recall CNN can only see images/pixels

Koay Yong Sheng, Stefano Moretti, Prashant Singh, Harri Waltari





# Auxiliary Information Important!





# CNNs need Aux Info to infer from <u>Different</u> Signal Regions

Koay Yong Sheng, Stefano Moretti, Prashant Singh, Harri Waltari

# n-1DCNN-wA

(Extending 1DCNN with Auxiliary Inputs)

# BSM Inference

# ANY SIGNAL REGION ANY FIXED NO. OF OBSERVABLES

# Poster A:14 Enhancing Robustness

BSM Parameter Inference with n1D-CNN and Novel Data Augmentation

Koay Yong Sheng





**Problem - Data Marginalized Forecasts for Bayesian** Model Comparison are Typically Infeasible

**Nested Sampling:** Generally applicable but too slow to explore data space **Savage–Dickey forecasts:** Restrictive assumptions and require nested models Both require explicit likelihoods.

Fully Bayesian Forecasts with Neural Bayes Ratio Estimation (Poster #9, arXiv:2309.06942) Thomas Gessey-Jones (tg400@cam.ac.uk) and Will Handley (wh260@cam.ac.uk)







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the network output f(d) converges to an invertible function  ${\cal G}$ of the Bayes ratio  $\mathcal{K}(d)$ between the competing models







Both require explicit likelihoods.





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