

Introduction to Nested Sampling



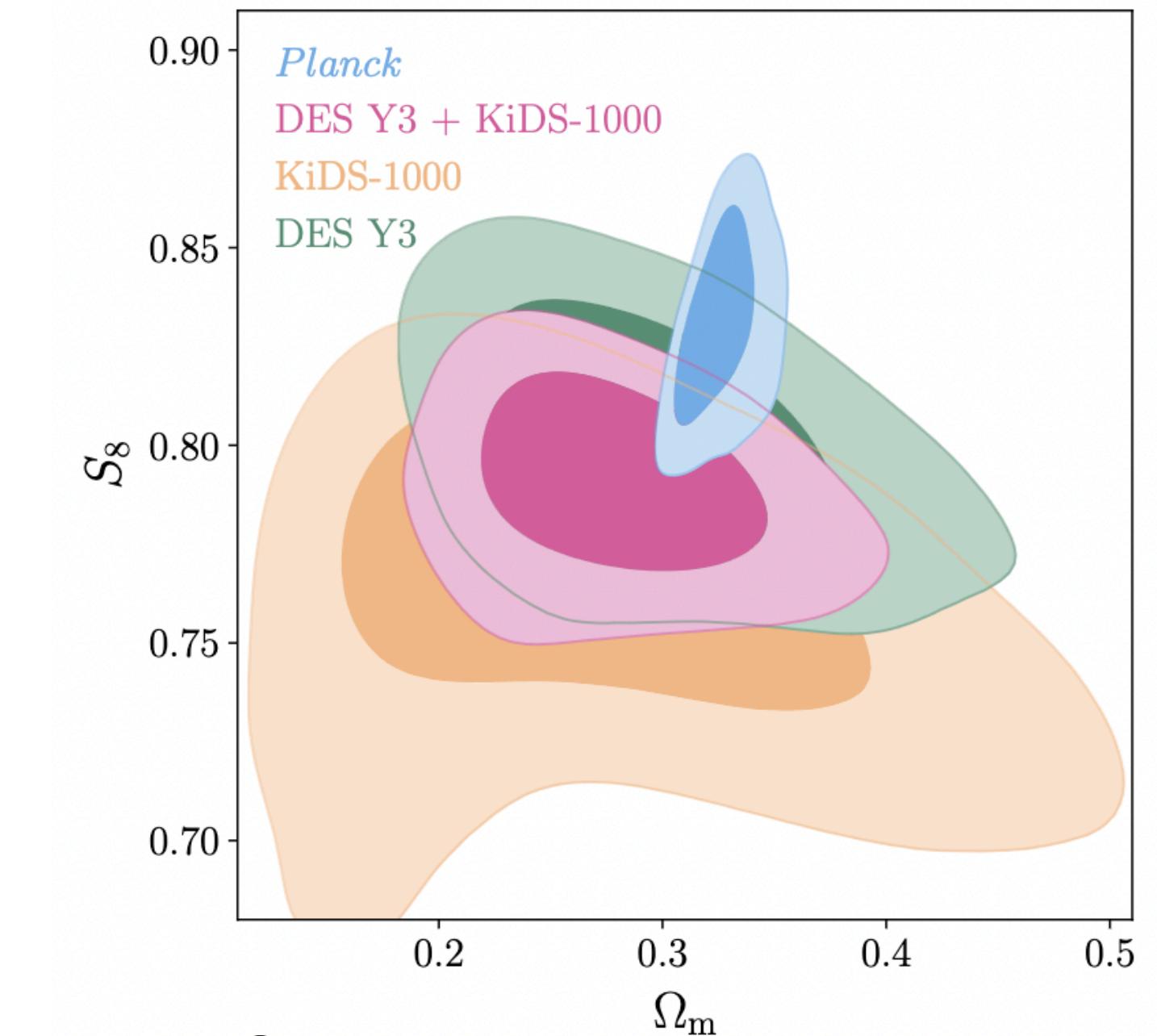
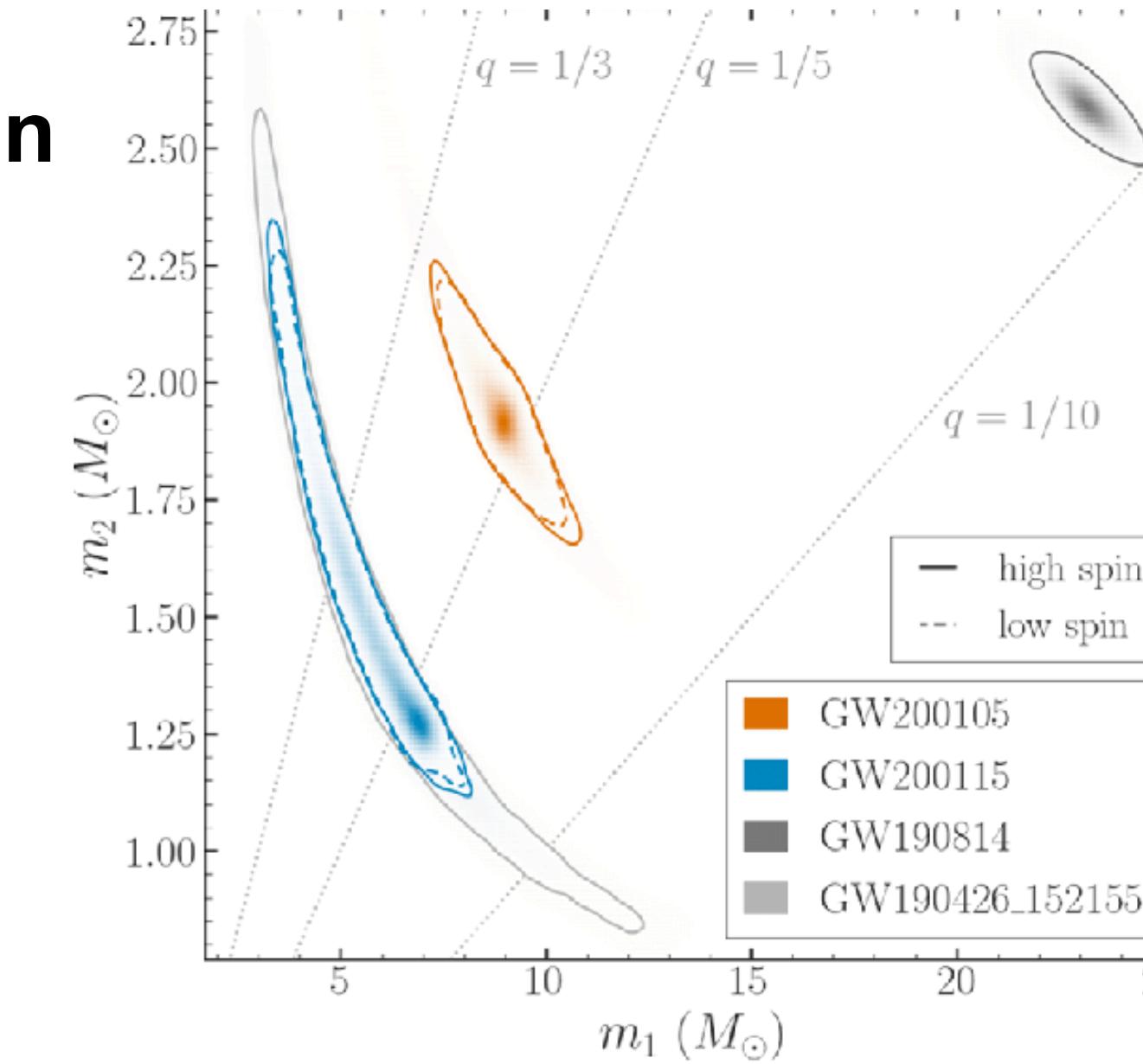
Dynesty is one of the package for Nested Sampling

What I've understood so far ...

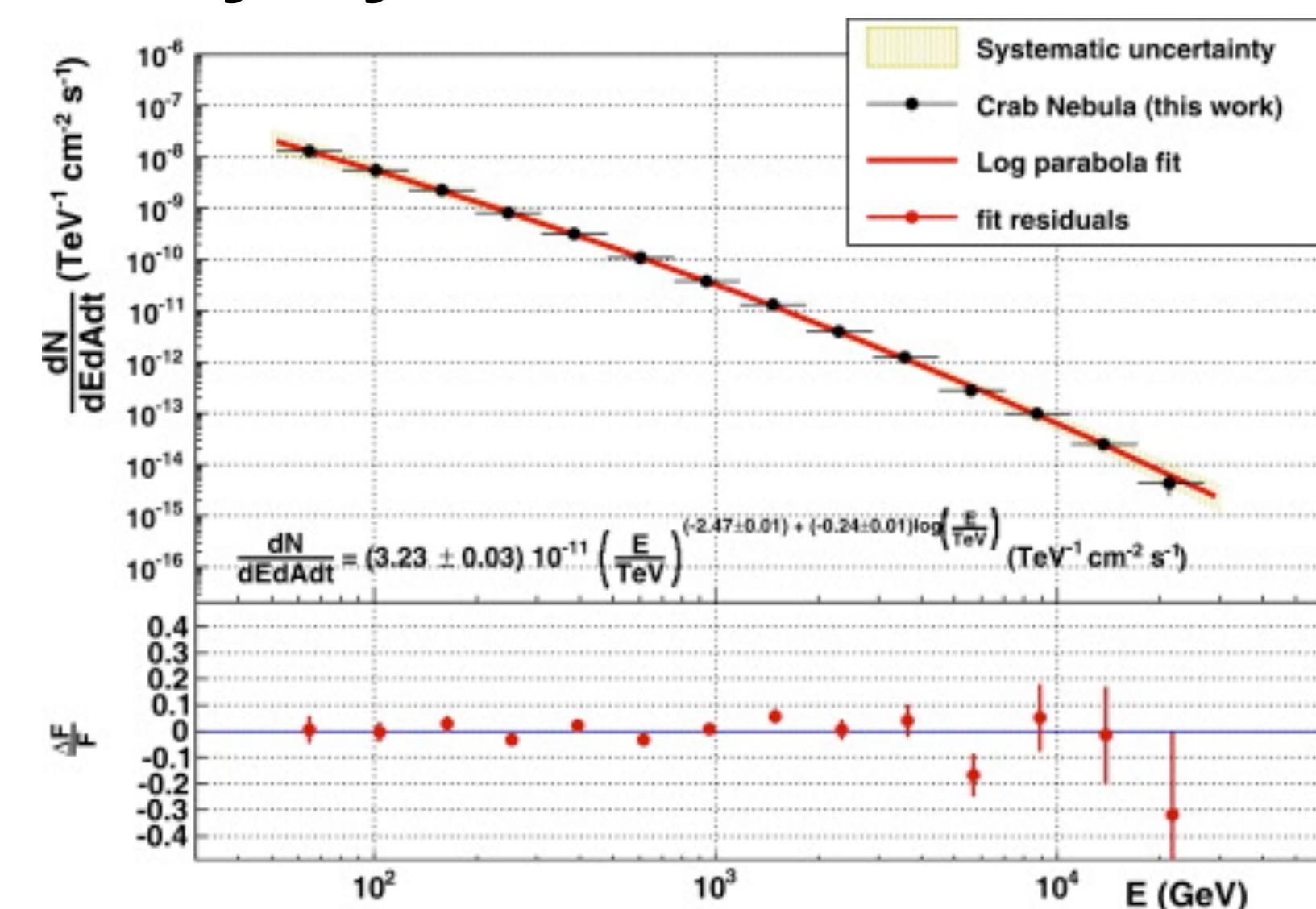
Fabio Acero,

A global shift towards Bayesian frameworks

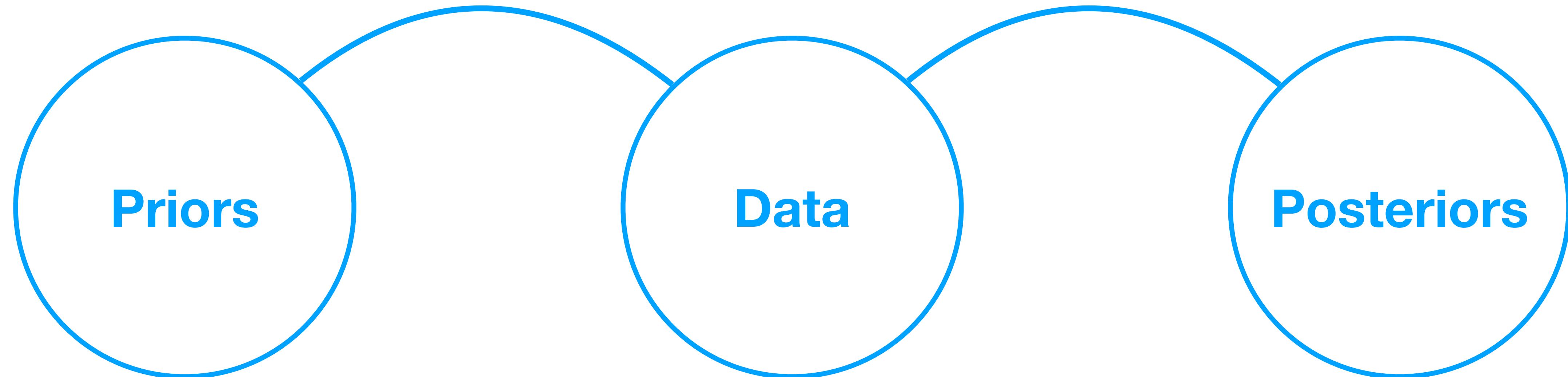
- Bayesian analyses have been dominant in cosmology for 1 or 2 decades
 - Complex models
 - Complex dataset.
 - Need to combine multiple datasets



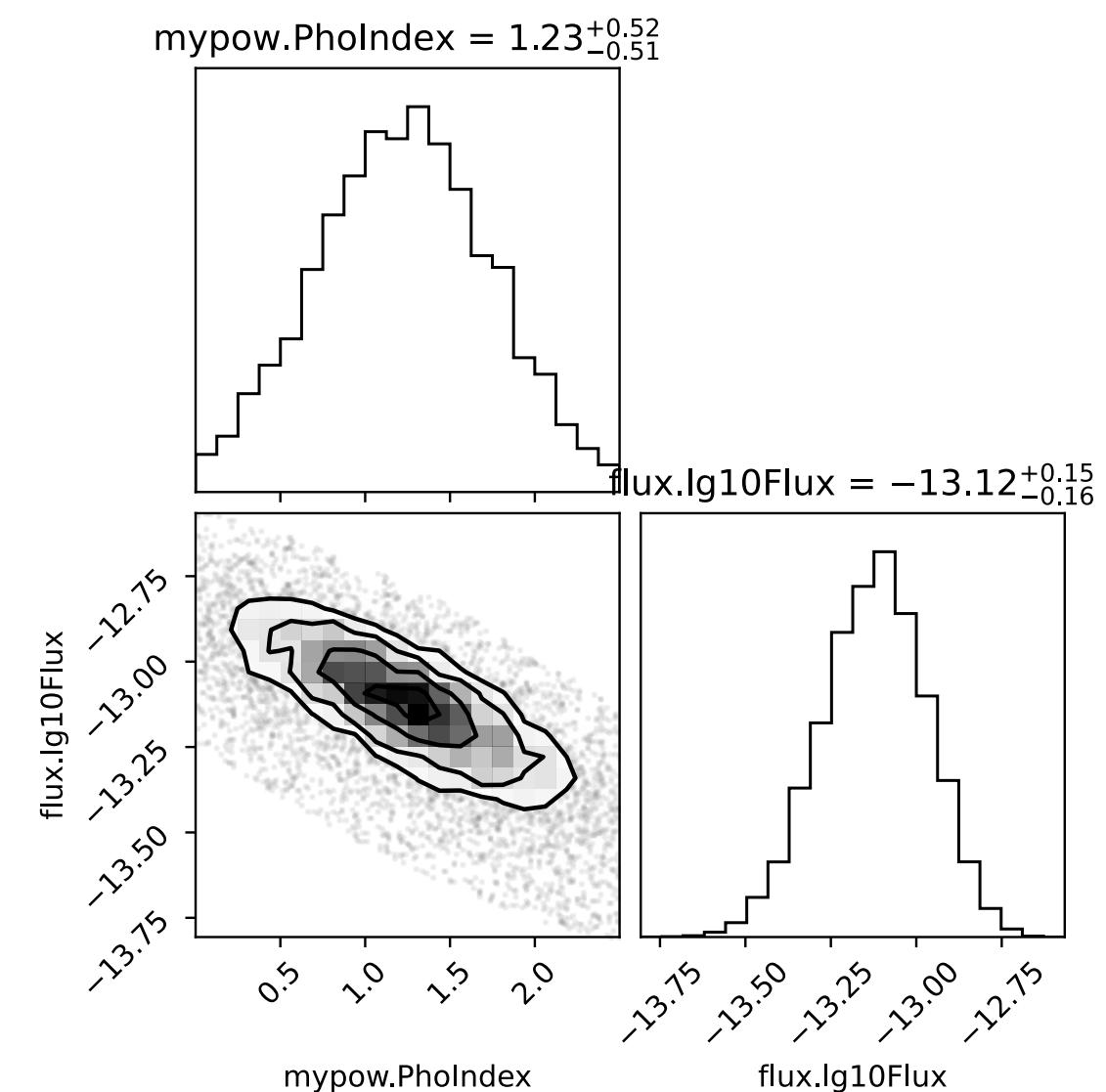
- In High-Energy astrophysics we're mostly loyal to Minuit and old fashioned Chi2 and LogLikelihood
 - Complex sources & instruments
 - But simpler models (POW) ?
 - But moving towards Bayesian



What is a Bayesian analysis (user point of view)



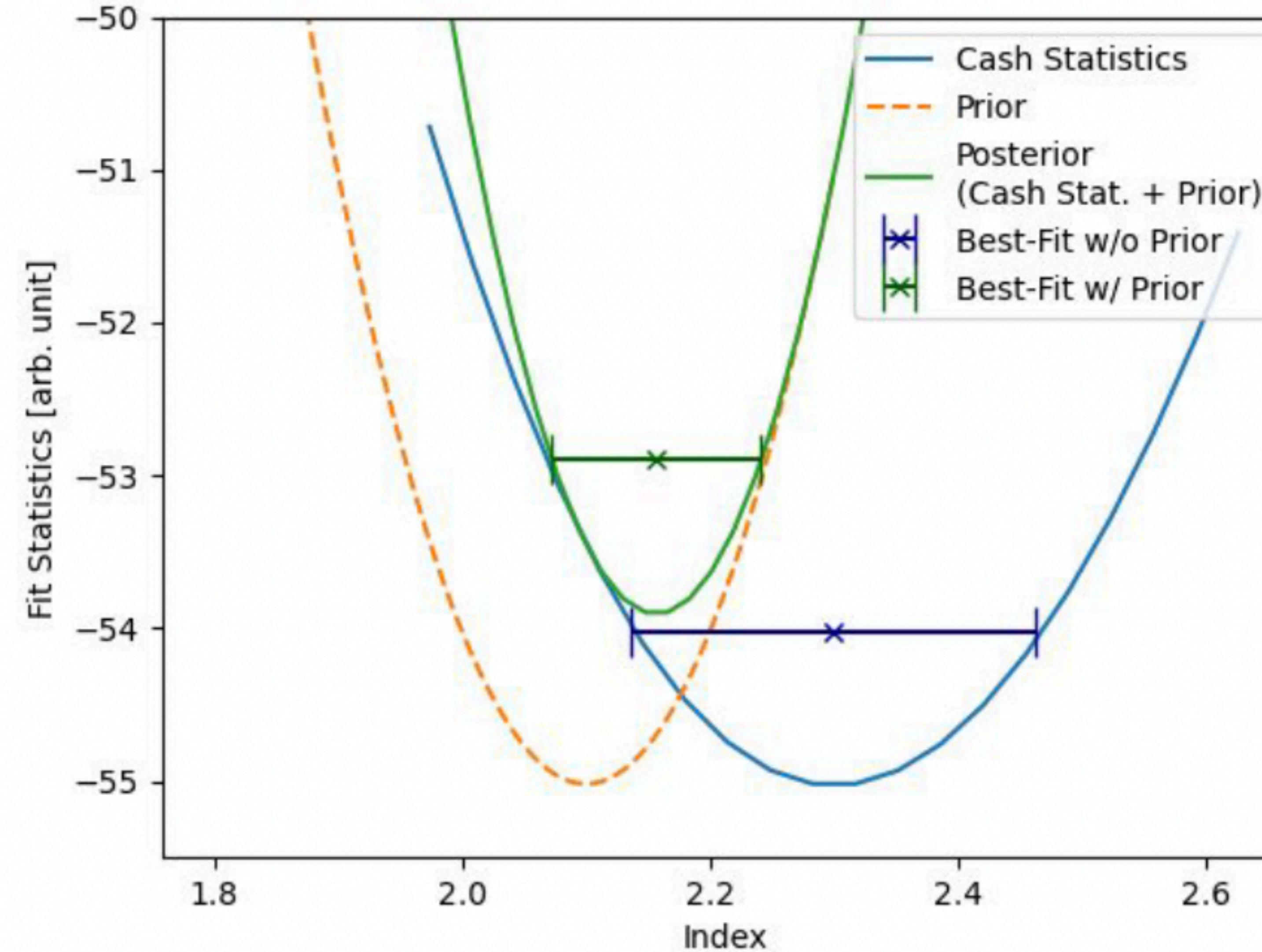
- Prior information on parameters
 - E.g. : $\text{Norm} > 0$
 - $P1 < \text{Param} < P2$
 - Norm_bkg Gaussian(1, sigma)
- Modified Likelihood
 - Likelihood = data_term + priors



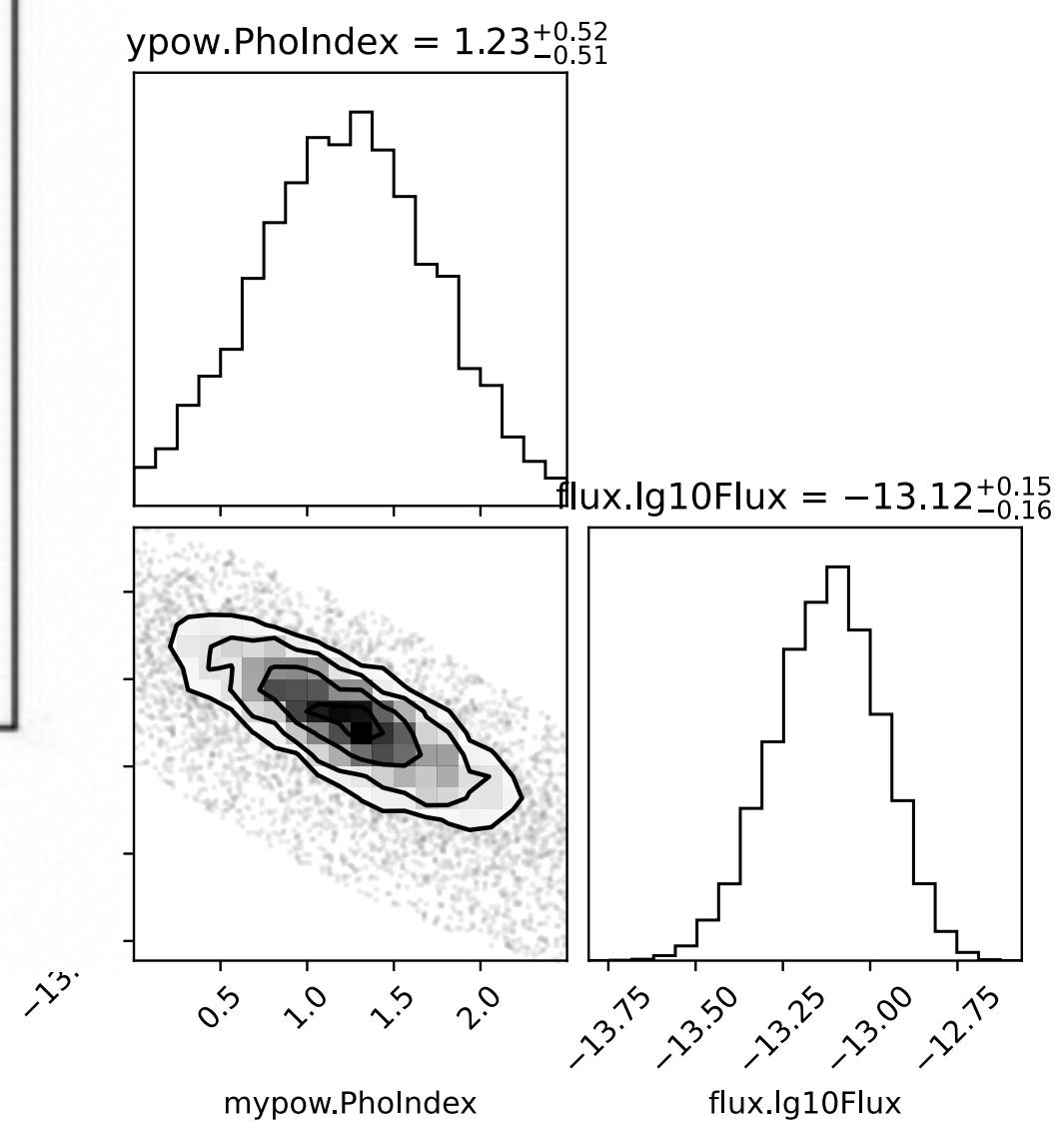
What is a Bayesian analysis (user point of view)

- Prior information
- E.g. : Norm > 0
- P1 < Param < F
- Norm_bkg Gal

Pri



osteriors

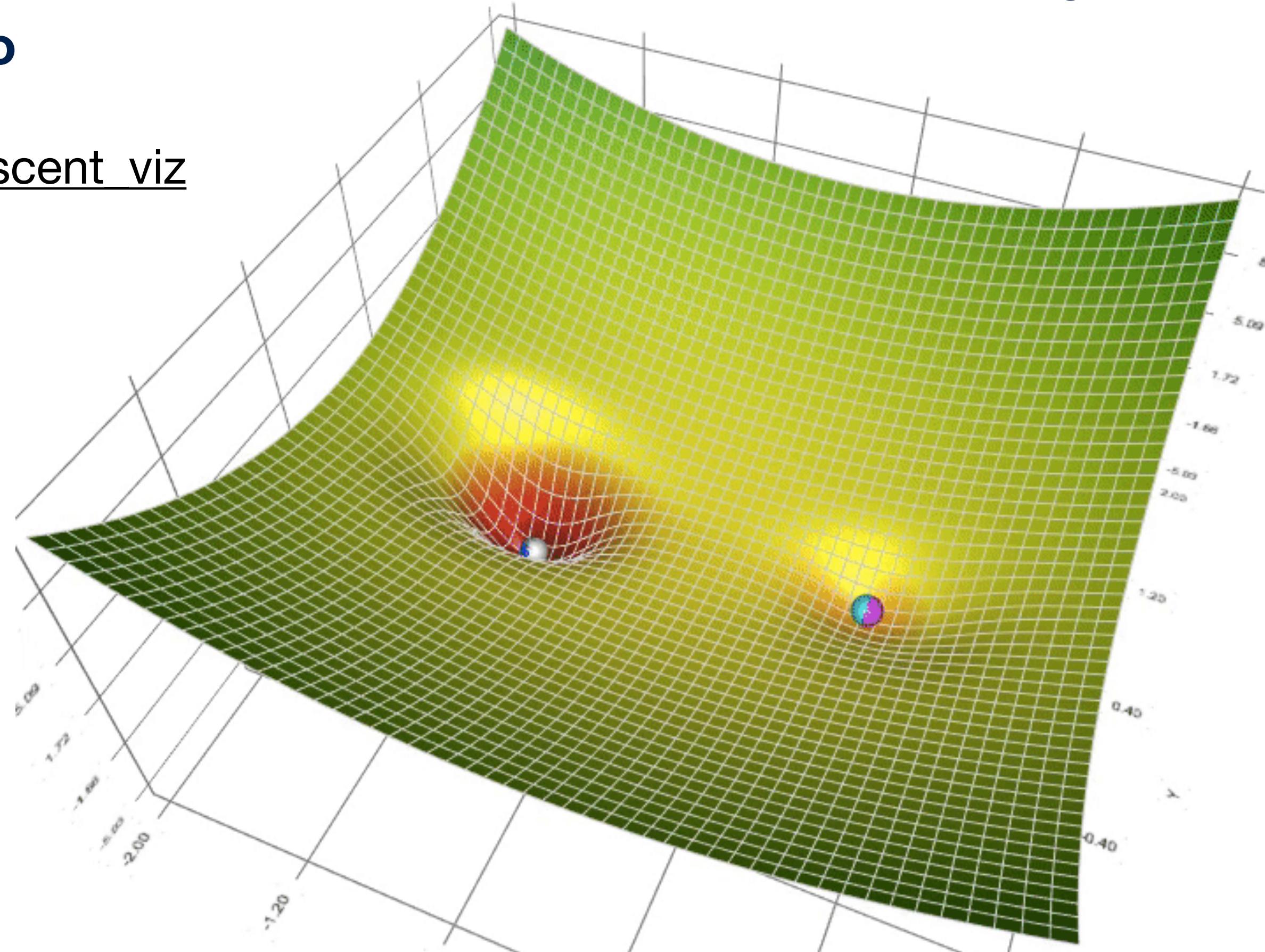


What is a fit ?

- Need a loss function + minimizer :
 - Gradient Descent (e.g. Scipy minimize, iMinuit, sherpa fit, etc)
 - Markov chain Monte Carlo
 - Nested Sampling metho

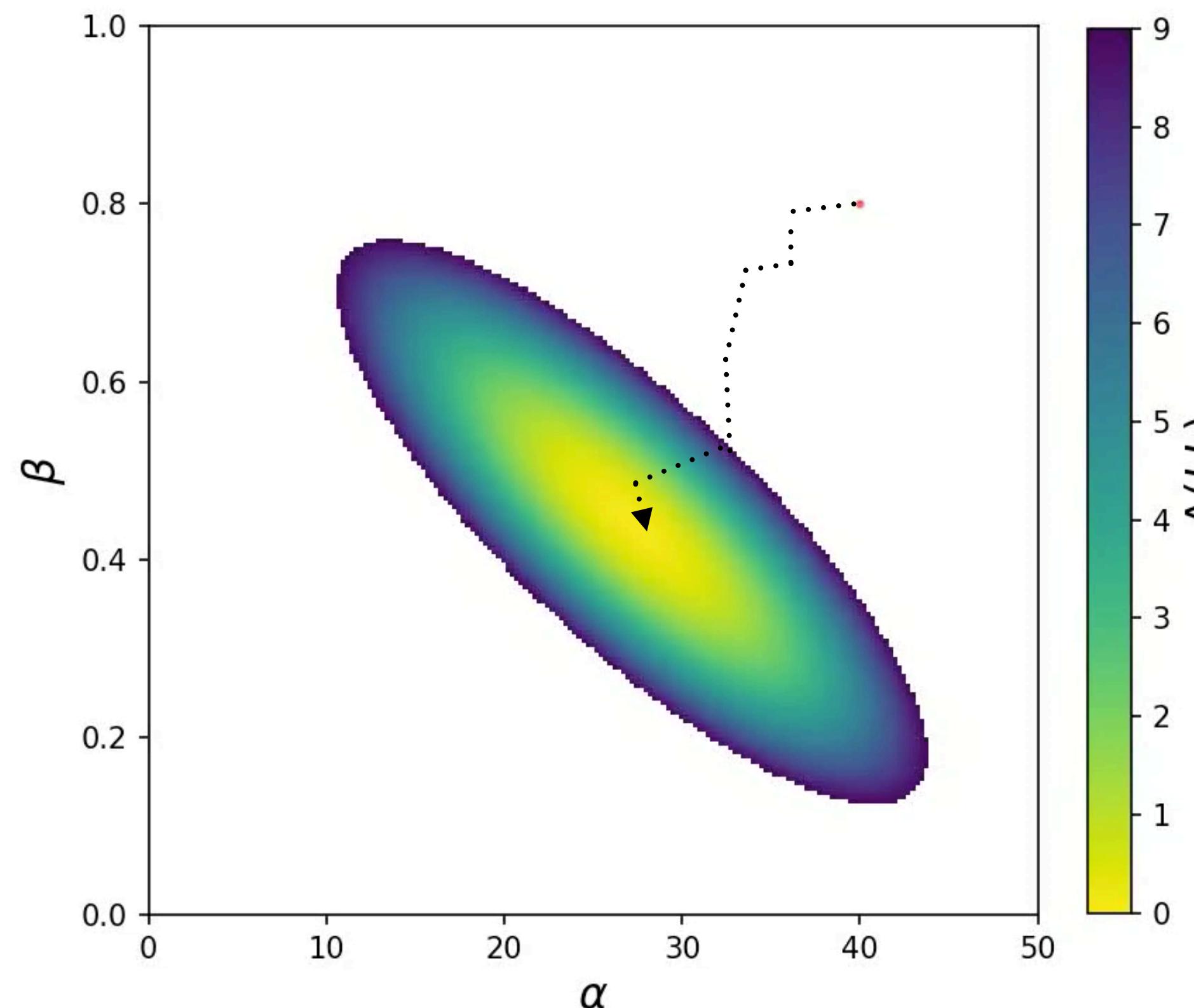
Never take a best-fit for granted

github.com/lilipads/gradient_descent_viz



Traditional fitting method

- Gradient descent based method
 - Levenberg Marquardt
 - Migrad in Minuit for example
 - Gradient is estimated numerically at each step

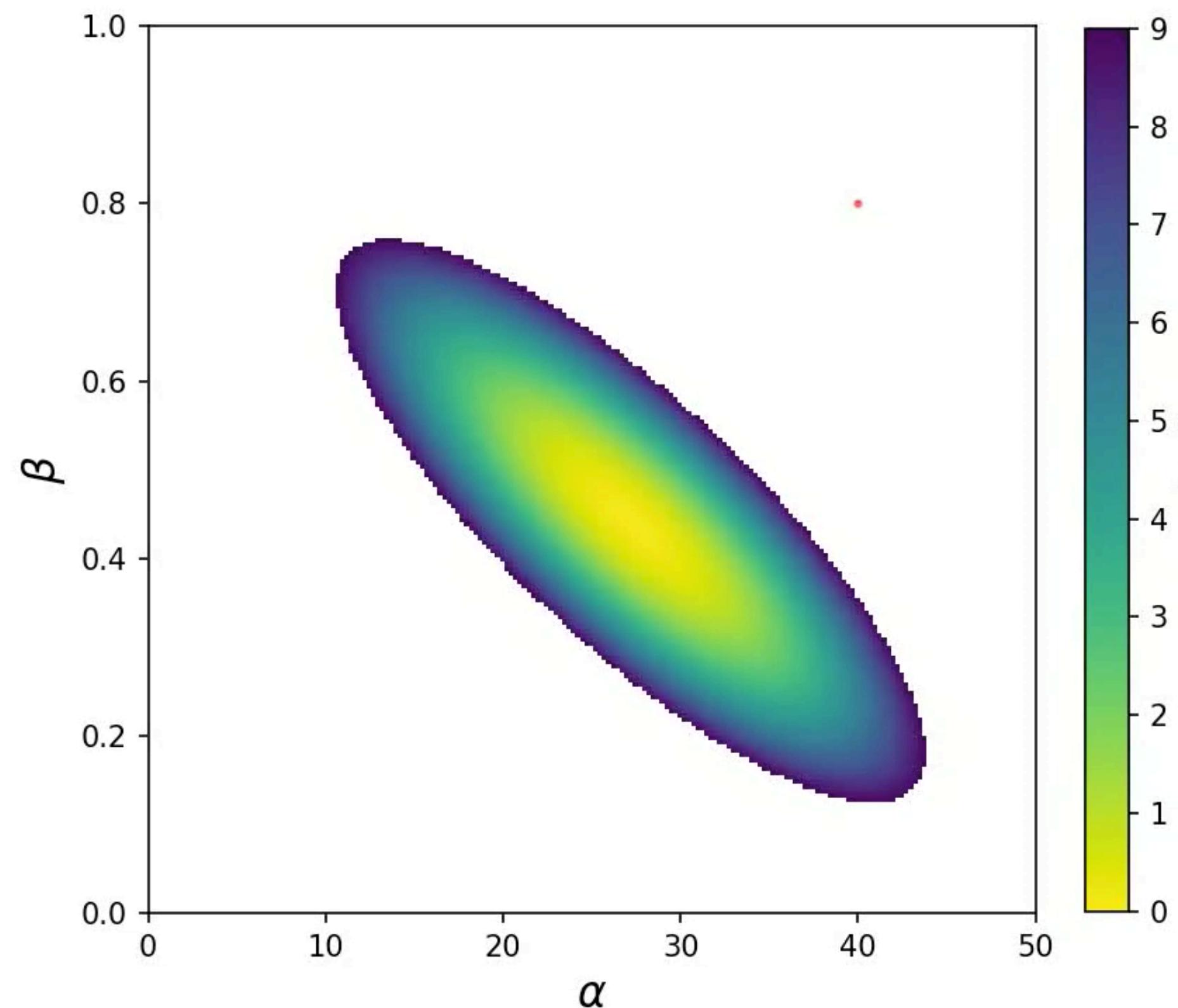


Once at best-fit stops
No information about local likelihood
Sometimes fit fails :

| Migrad | |
|---------------------------------|------------------------|
| FCN = -2.676e+07 | Nfcn = 378 |
| EDM = 0.00356 (Goal: 2e-06) | time = 2.3 sec |
| INVALID Minimum | No Parameters at limit |
| ABOVE EDM threshold (goal x 10) | Below call limit |
| Covariance | Hesse ok |
| APPROXIMATE | NOT pos. def. |
| FORCED | |

Markov Chain Monte Carlo (MCMC)

- What are they:
 - Monte Carlo: samples are used to approximate the probability distribution
 - Markov Chain: semi-random walk in potential
 - Walkers explore the local likelihood



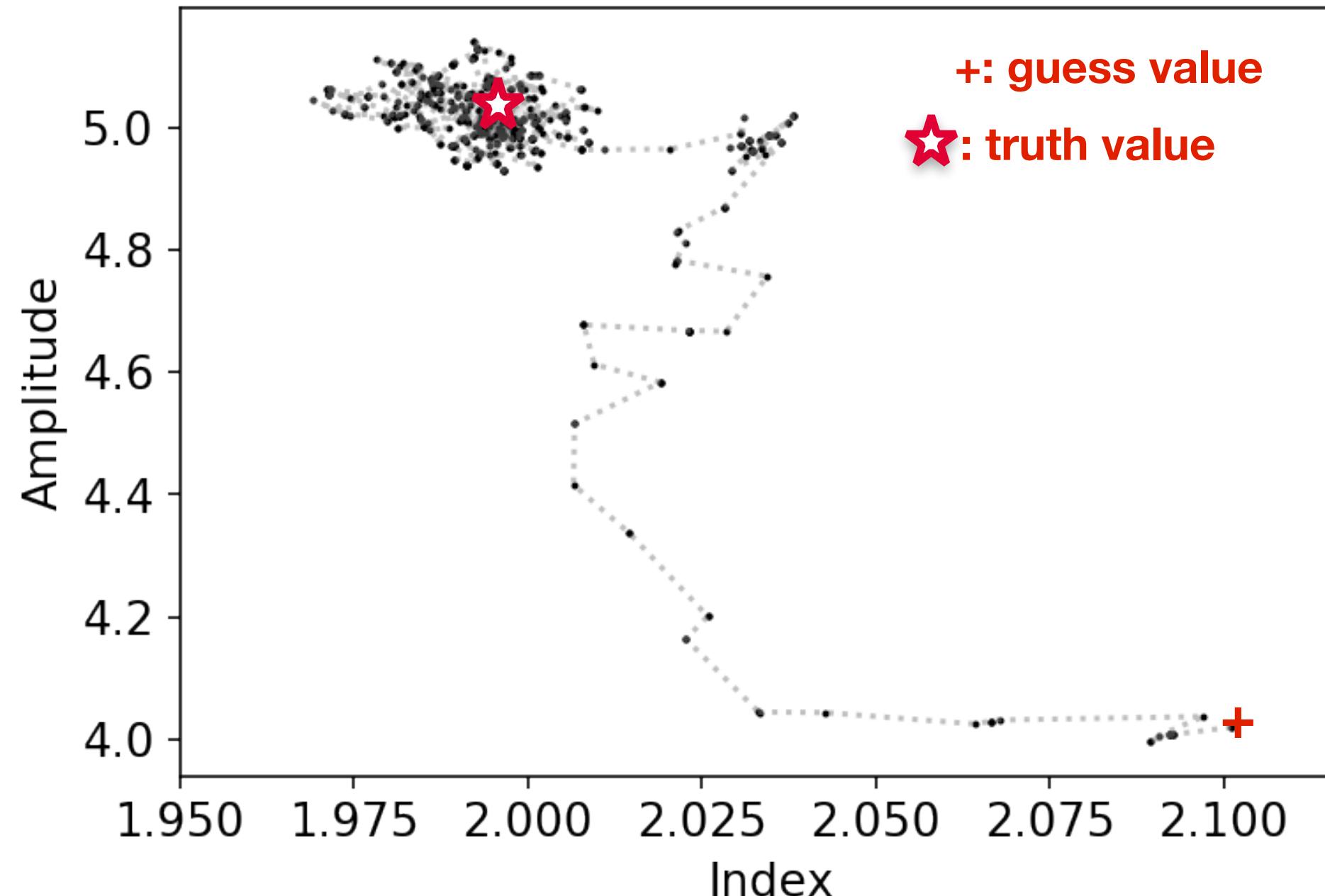
Random walk directed by potential (likelihood)
spend most of their time in interesting region

Technically not a fit (no convergence) it's
a phase space parameter exploration

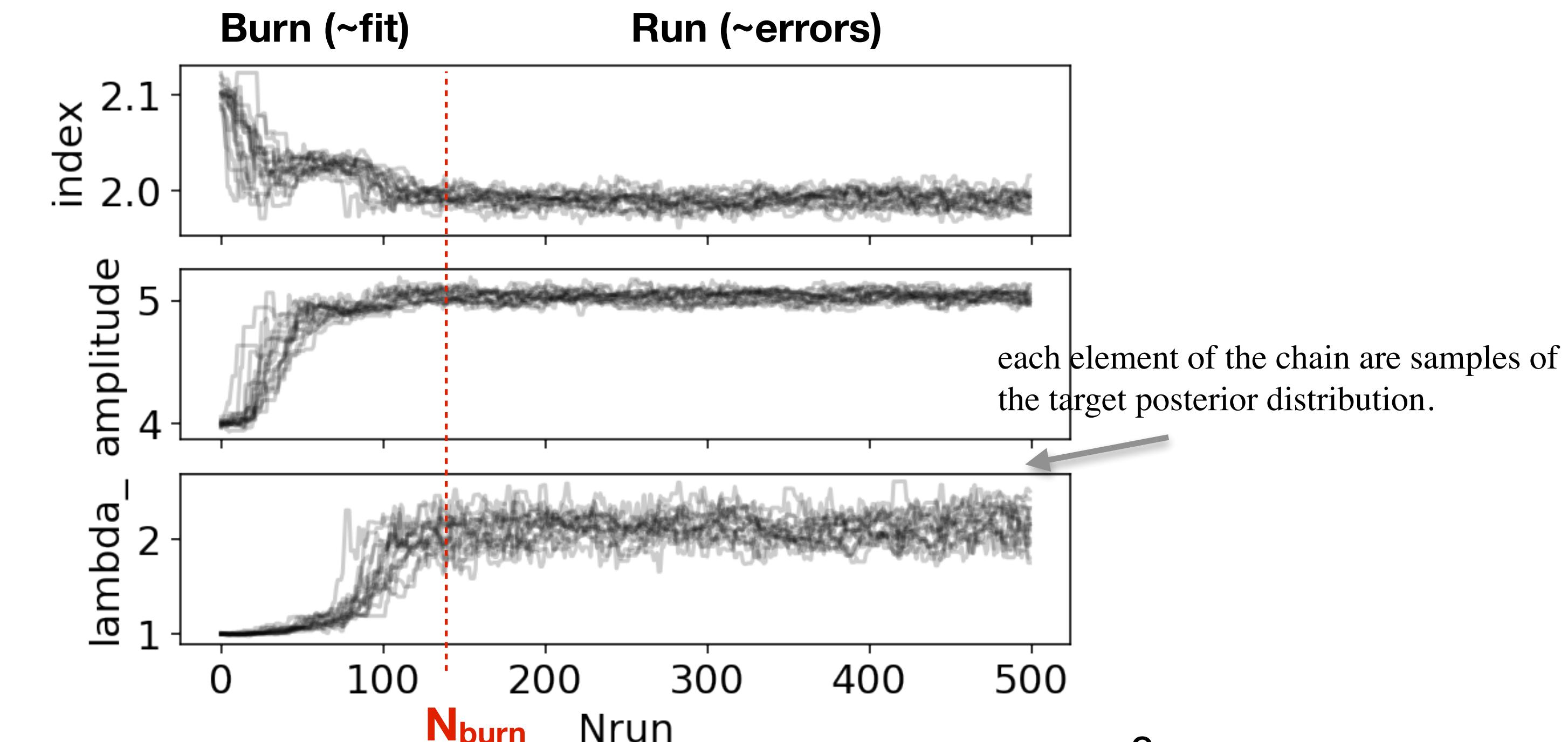
MCMC issues : inside out approach

- Need to define an initialization point
 - Might stay trapped in local minima. Could miss multimodal posterior
- Need to define "by eye" an burn period
 - No clear automated criteria
- Need to evolve walkers for "sufficiently long" to sample target posterior
 - How long is good enough ?

1 walker evolving for 500 steps



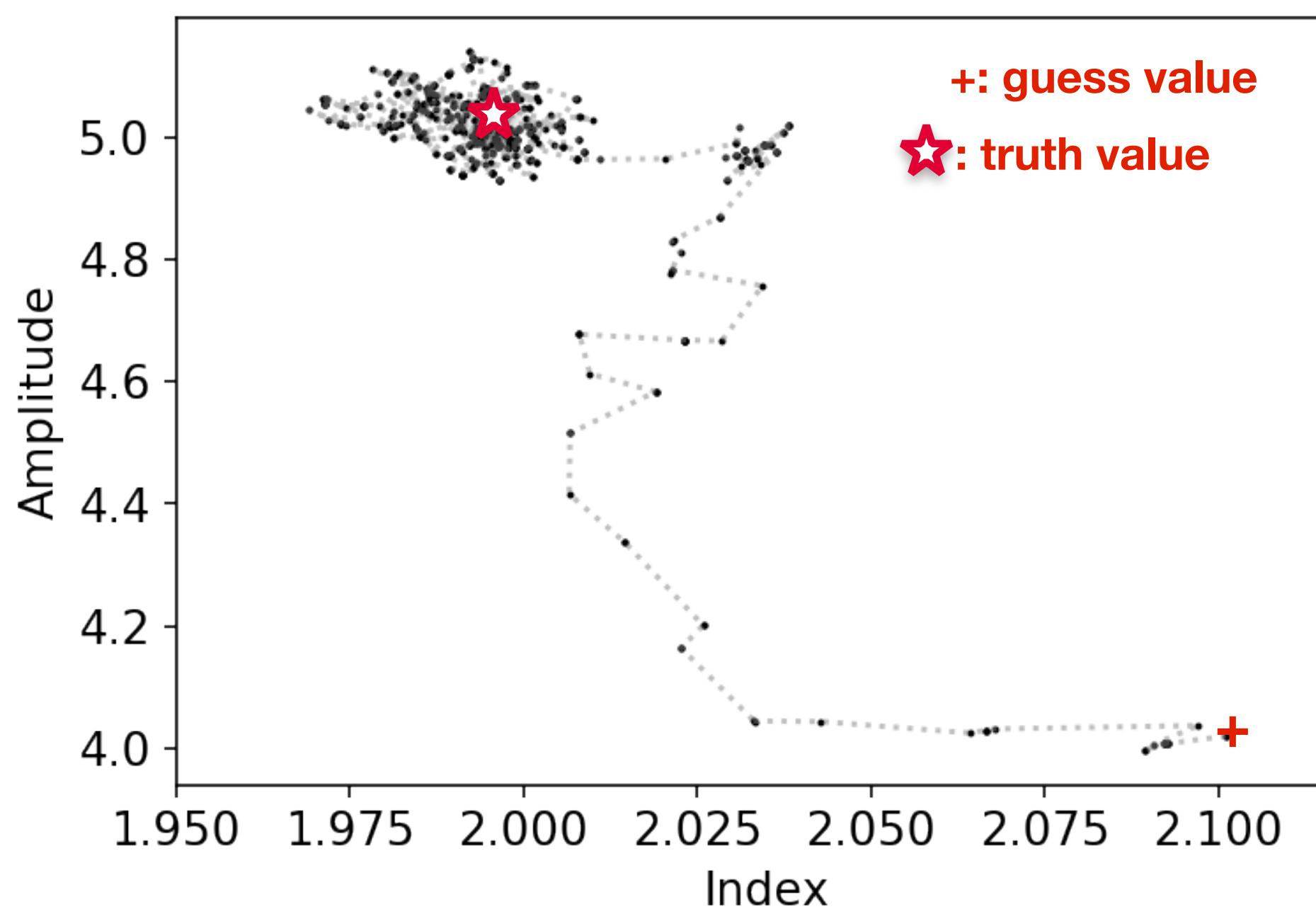
10 walkers evolving for 500 steps



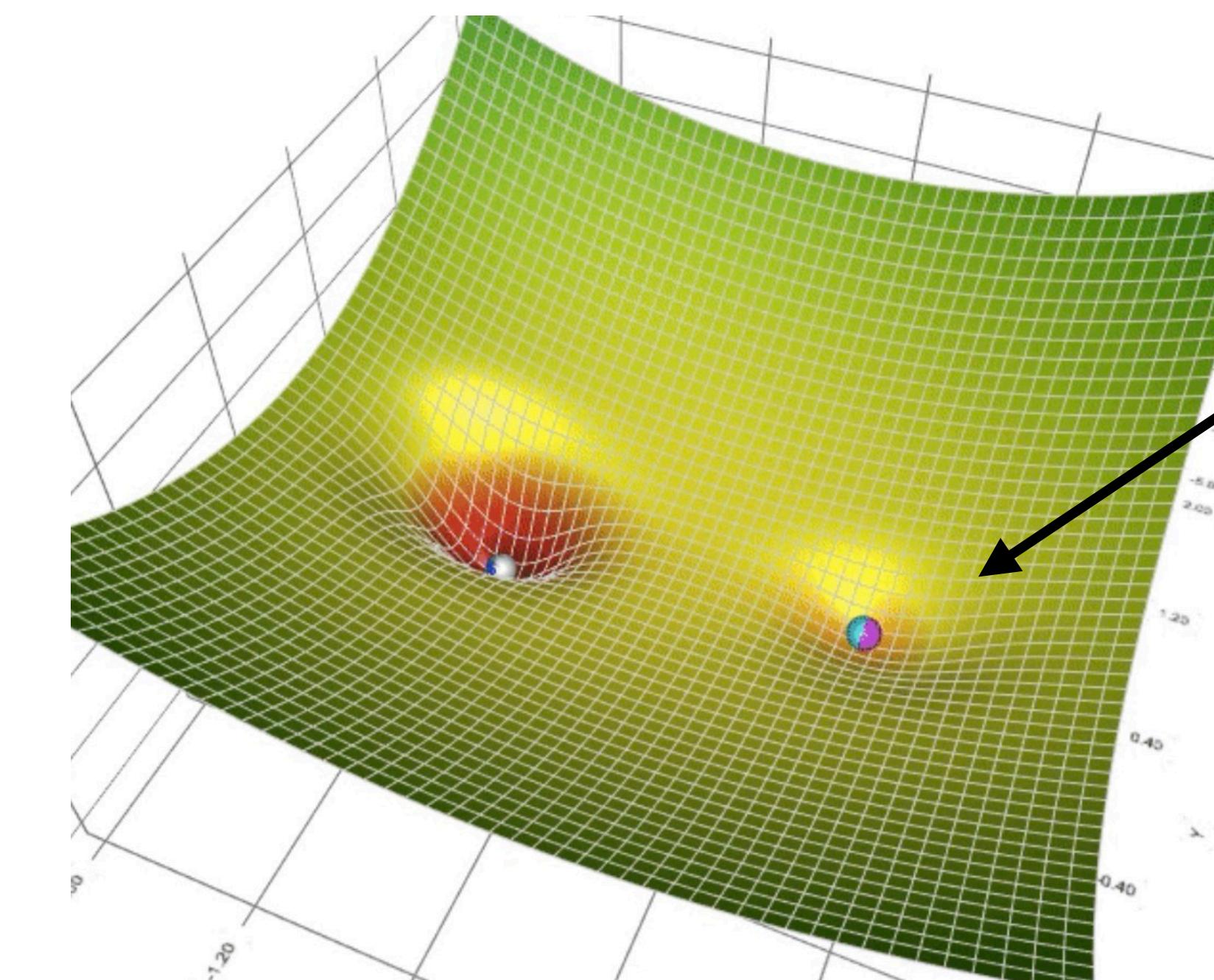
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1 walker evolving for 500 steps



10 walkers evolving for 500 steps



But what if all your walkers end up here ?

Nested Sampling methods

- Nested Sampling (Skilling, 2004) is a Monte Carlo algorithm for estimating an integral over a model parameter space Θ
 - Integral :
 - $\int \text{Like(Data}|\Theta) * \text{Prior}(\Theta) d\Theta = Z = \text{Bayesian evidence}$
 - Z can be used to compare models even if not nested (morphology : mwI template vs disk)
 - This integral is also what will provide the normalized posterior distributions
 - Main idea is integration is done by switching frame from many Θ to a volume variable

Unlike MCMC methods, which attempt to estimate the posterior $\mathcal{P}(\Theta)$ directly, Nested Sampling instead focuses on estimating the evidence

$$Z \equiv \int_{\Omega_\Theta} \mathcal{P}(\Theta) d\Theta = \int_{\Omega_\Theta} \mathcal{L}(\Theta) \pi(\Theta) d\Theta \quad (6)$$

As this integral is over the entire multi-dimensional domain of Θ , it is traditionally very challenging to estimate.

Nested Sampling approaches this problem by refactoring this integral as one taken over prior volume X of the enclosed parameter space

$$Z = \int_{\Omega_\Theta} \mathcal{L}(\Theta) \pi(\Theta) d\Theta = \int_0^1 \mathcal{L}(X) dX \quad (7)$$

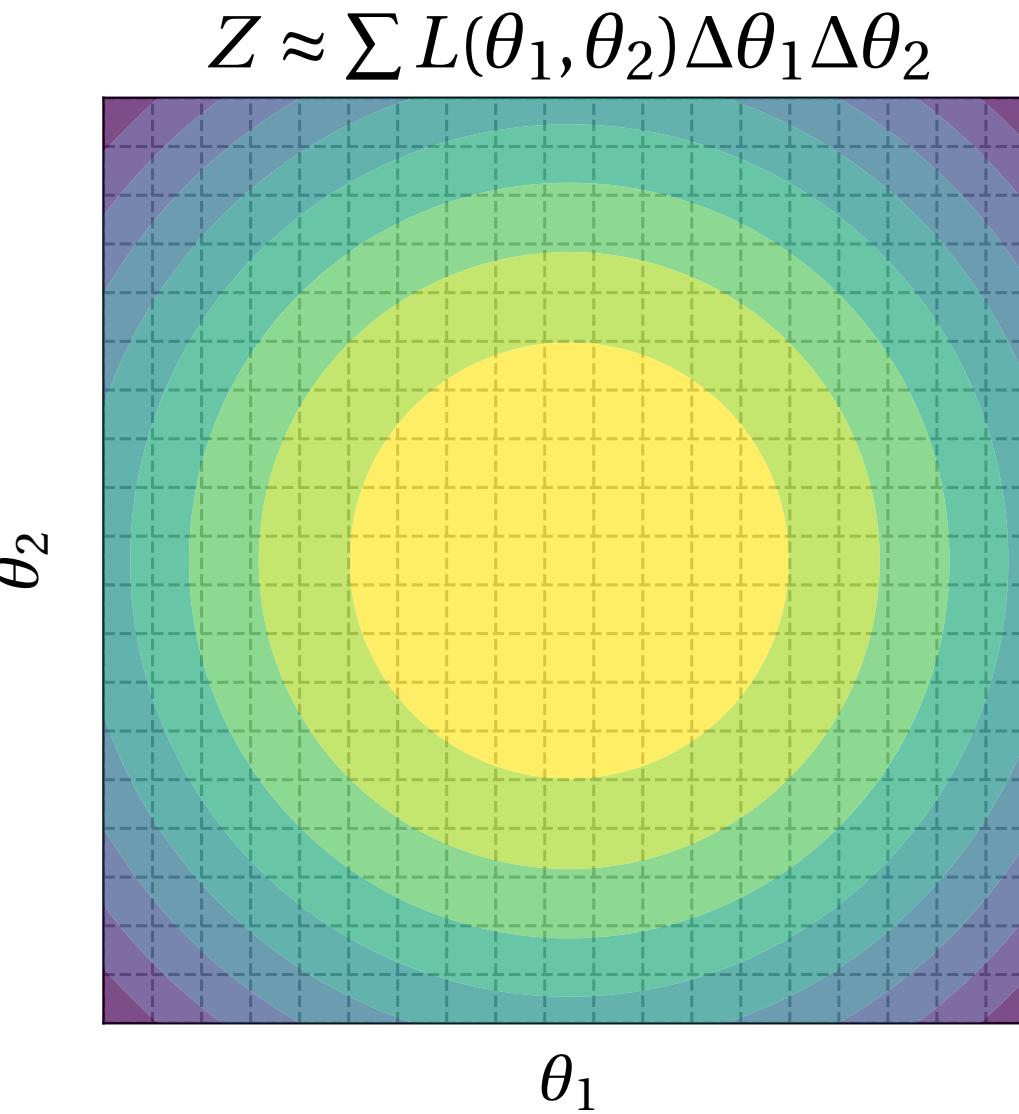
- **Analogy to spherical coordinates:**

$$\int \mathcal{P}(x, y, z) dx dy dz = \int \mathcal{P}(V(r)) dV(r) = \int \mathcal{P}(r) 4\pi r^2 dr$$

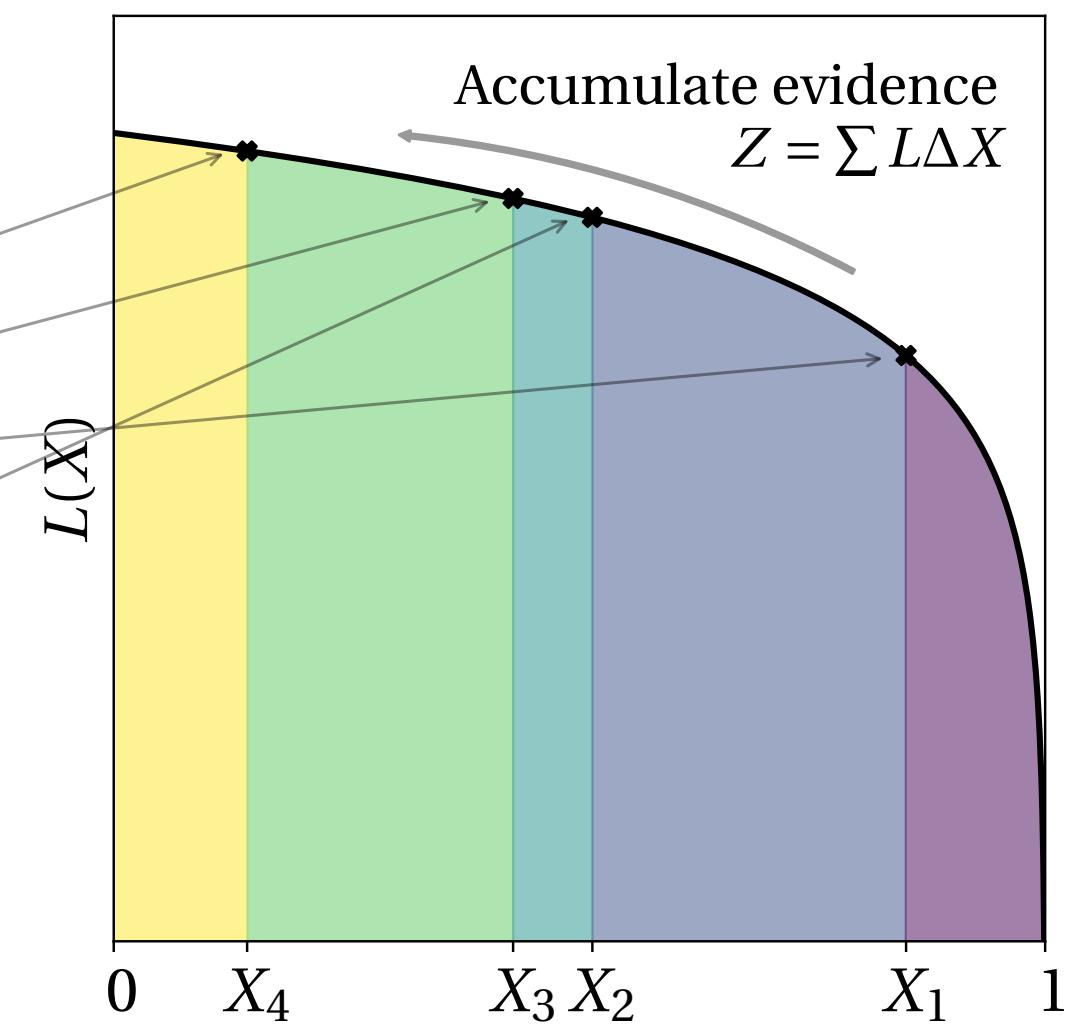
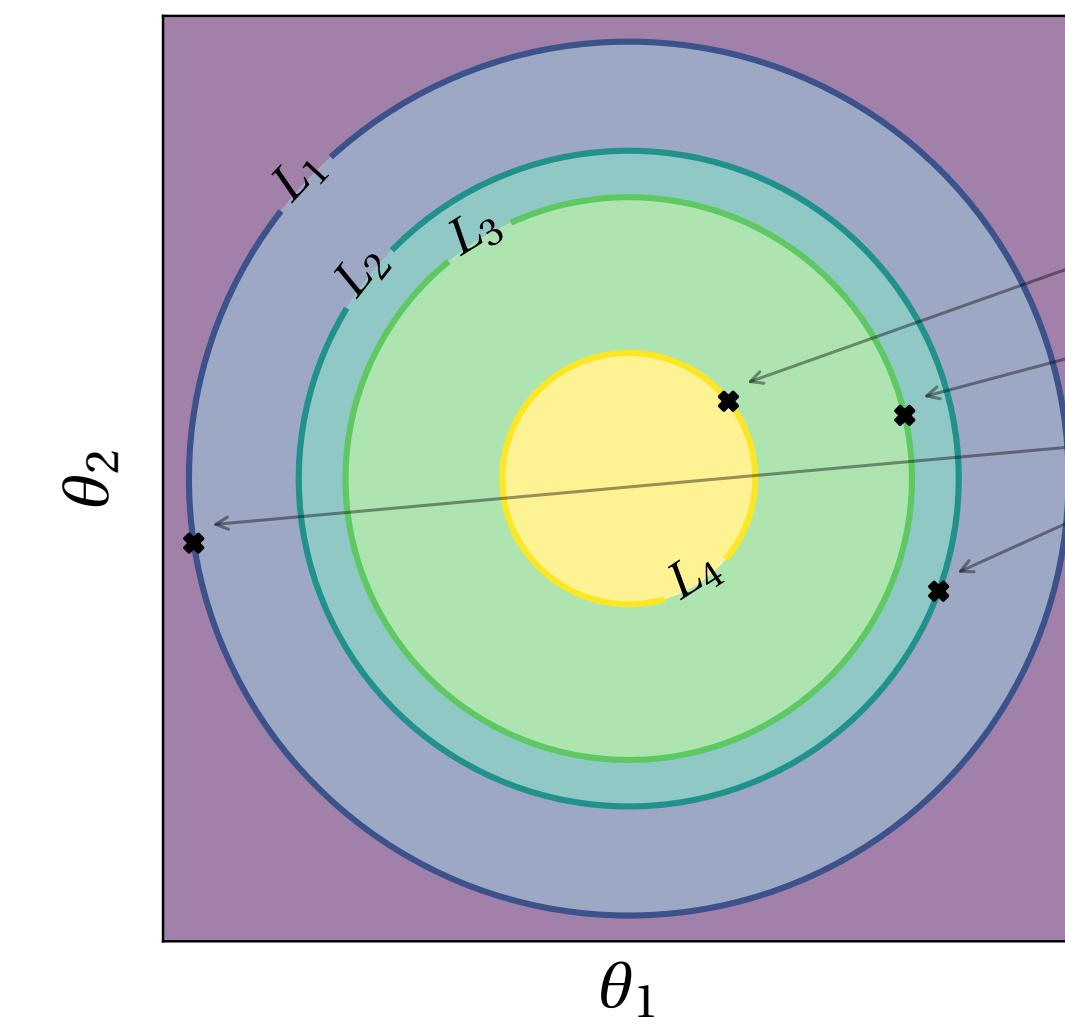
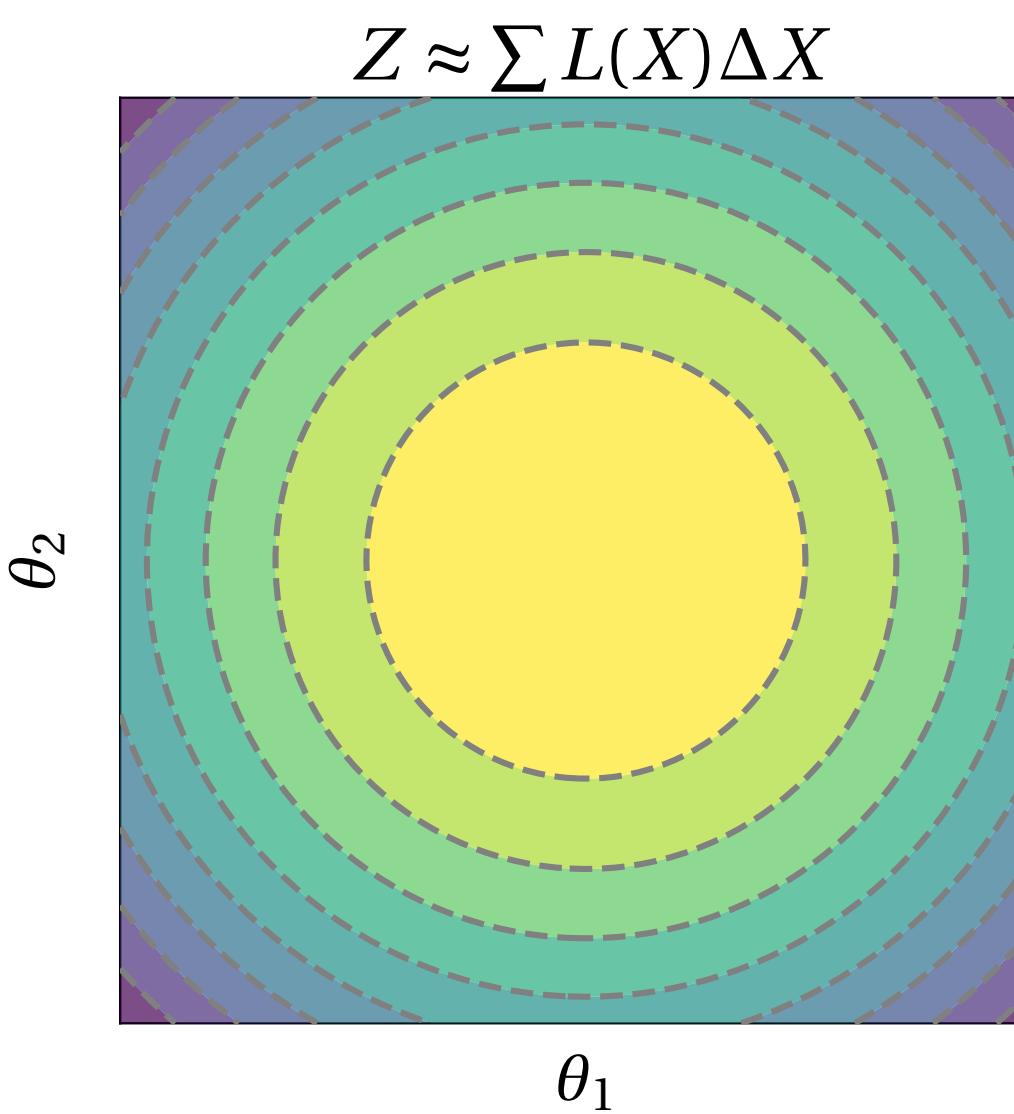
Nested means integrating in nested shells of iso-likelihood

- Goal of the method is to integrate over shells (like an onion) to approximate the integral

Direct space : more complex



Volume space : integration over shell of iso-LogLike

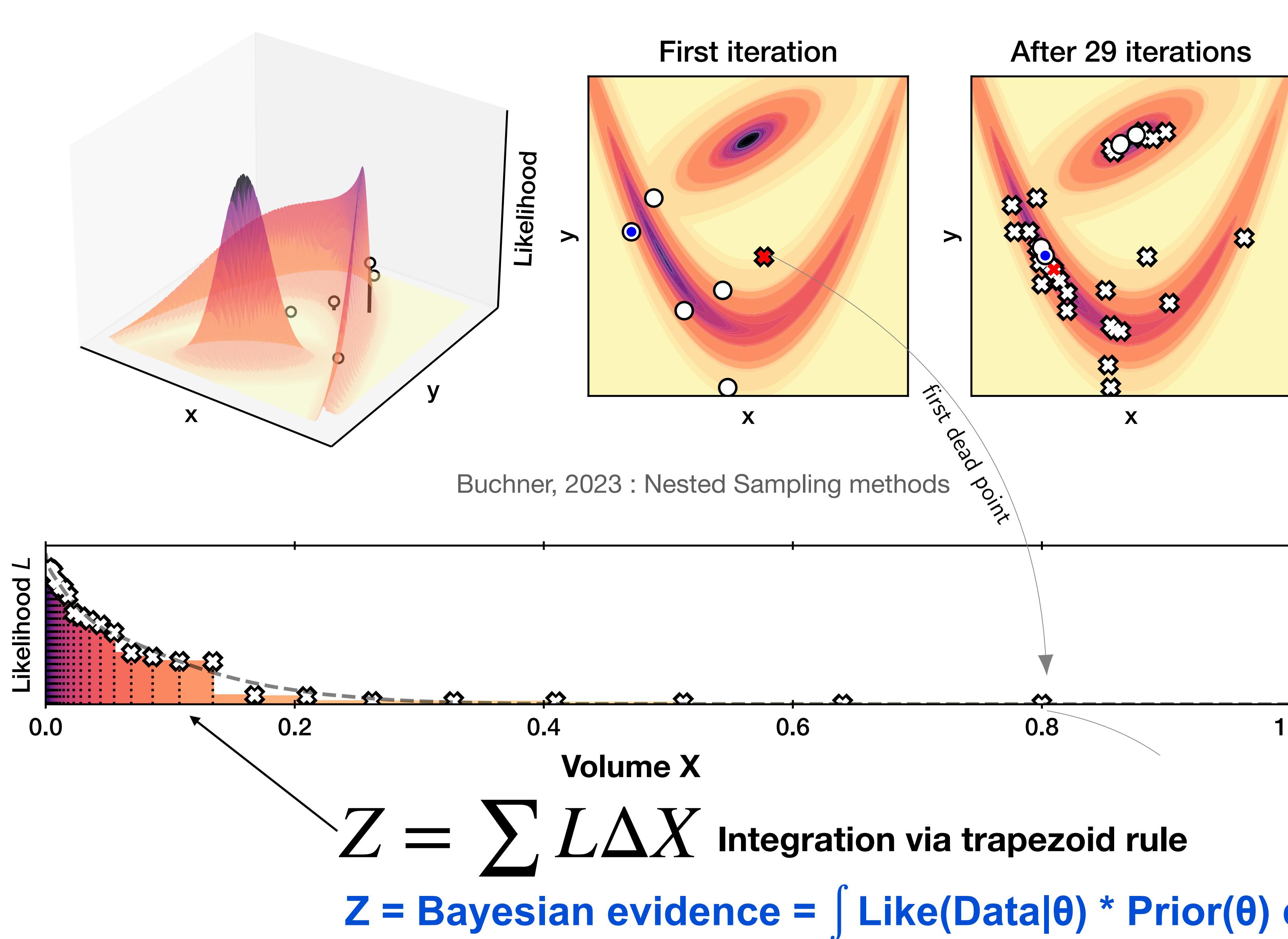


a | The NS **evidence identity**. The colours represent contours of a two-dimensional **likelihood** function. Rather than summing over little cubes (left), we combine cubes of similar **likelihood** together and sum over them (right).

b | NS on a two dimensional problem. We show the dead points and their **iso-likelihood contours** (left) and the corresponding contributions to the **evidence** integral (right). The volumes X_i are estimated statistically in NS.

Ashton, 2002 in Nature : Nested Sampling for physical scientists

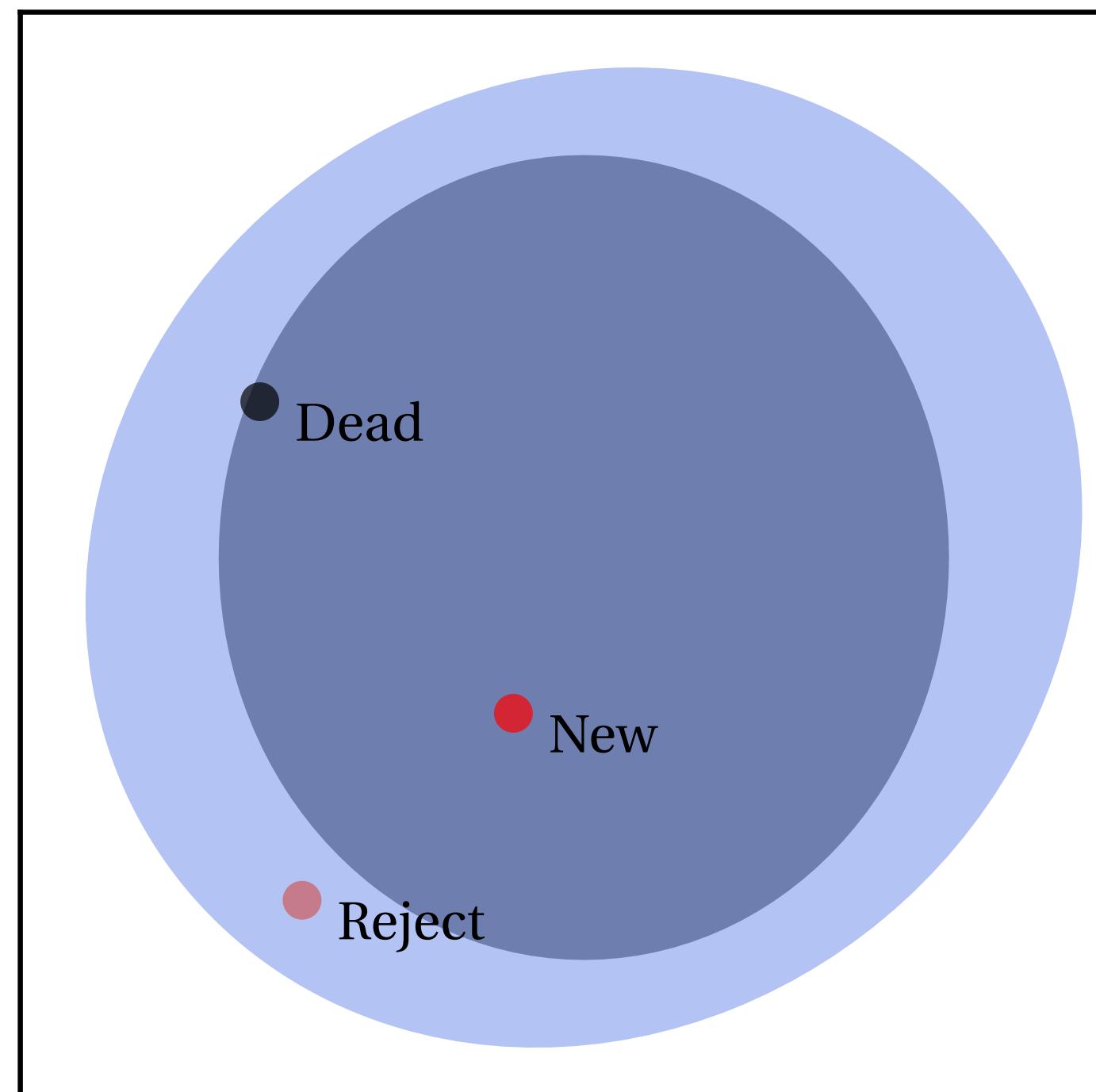
Basic steps of Nested Sampling



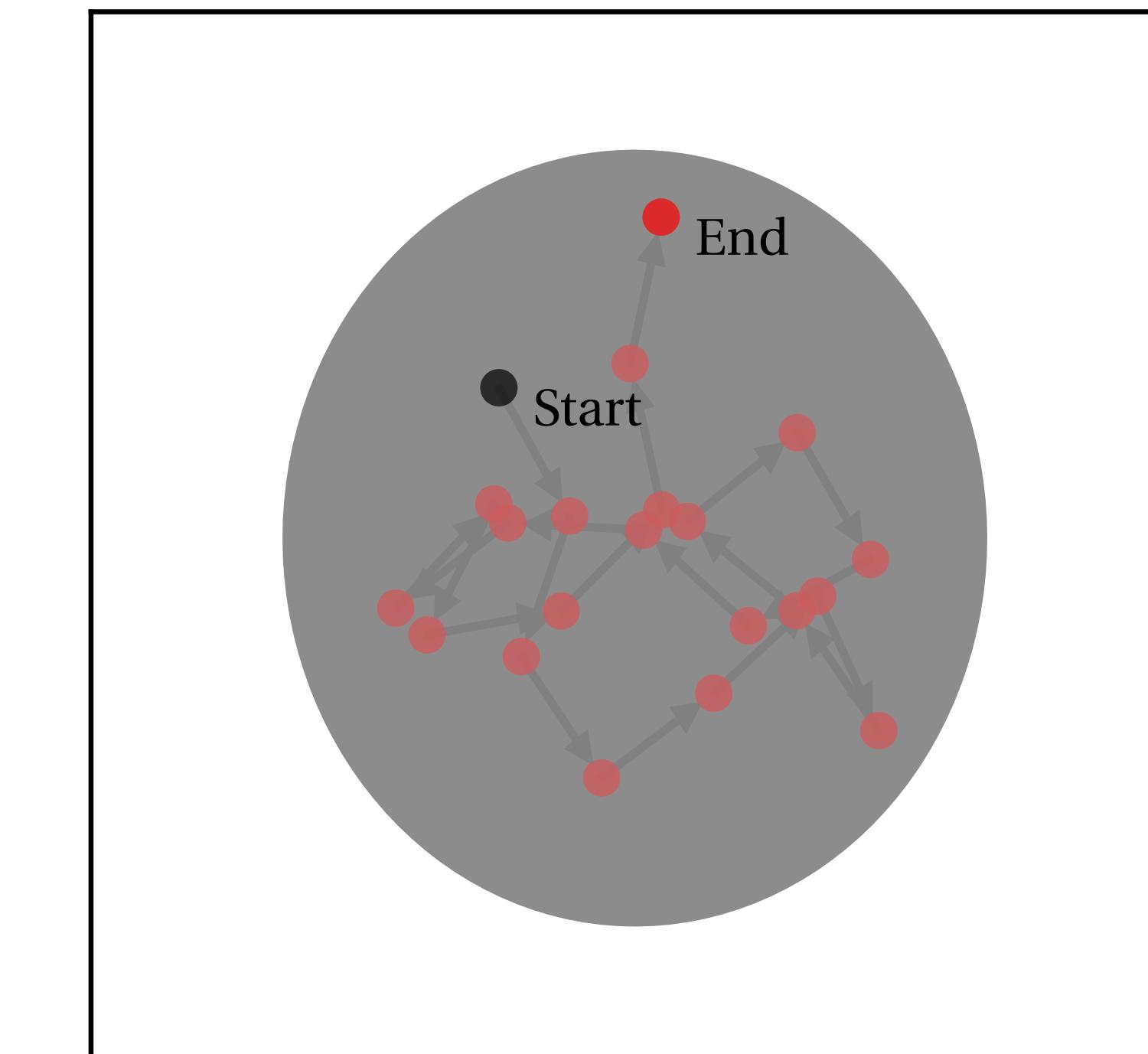
- Define priors on params
- Transform param space to volume unity cube X
- Draw random uniform N_{live} ($\sim 400-1000$)
- Iterate until stop criteria:
 - Remove worst LogLike point
 - Draw new point with a better LL
 - Likelihood-restricted prior sampling
 - (The tricky part)
 - Increment $Z = Z + L^* \Delta X$
 - Stop criteria : $\Delta Z/Z < \text{tolerance}$

Sampling from a restricted prior ($\text{Like}_{\text{new}} > \text{Like}_{\text{removed}}$)

- The tricky part : how to sample points with a better LogLike
 - This means drawing point inside the iso LogLike contours that we don't know
- Take advantage from the fact that if NLive is large enough. Surviving points already provide trace the landscape
- So drawing an encapsulating ellipsoid and try to sample from this ellipsoid. Reject if needed
- Or if landscape too complex or high dimension use a step sampler



Ellipsoid sampling

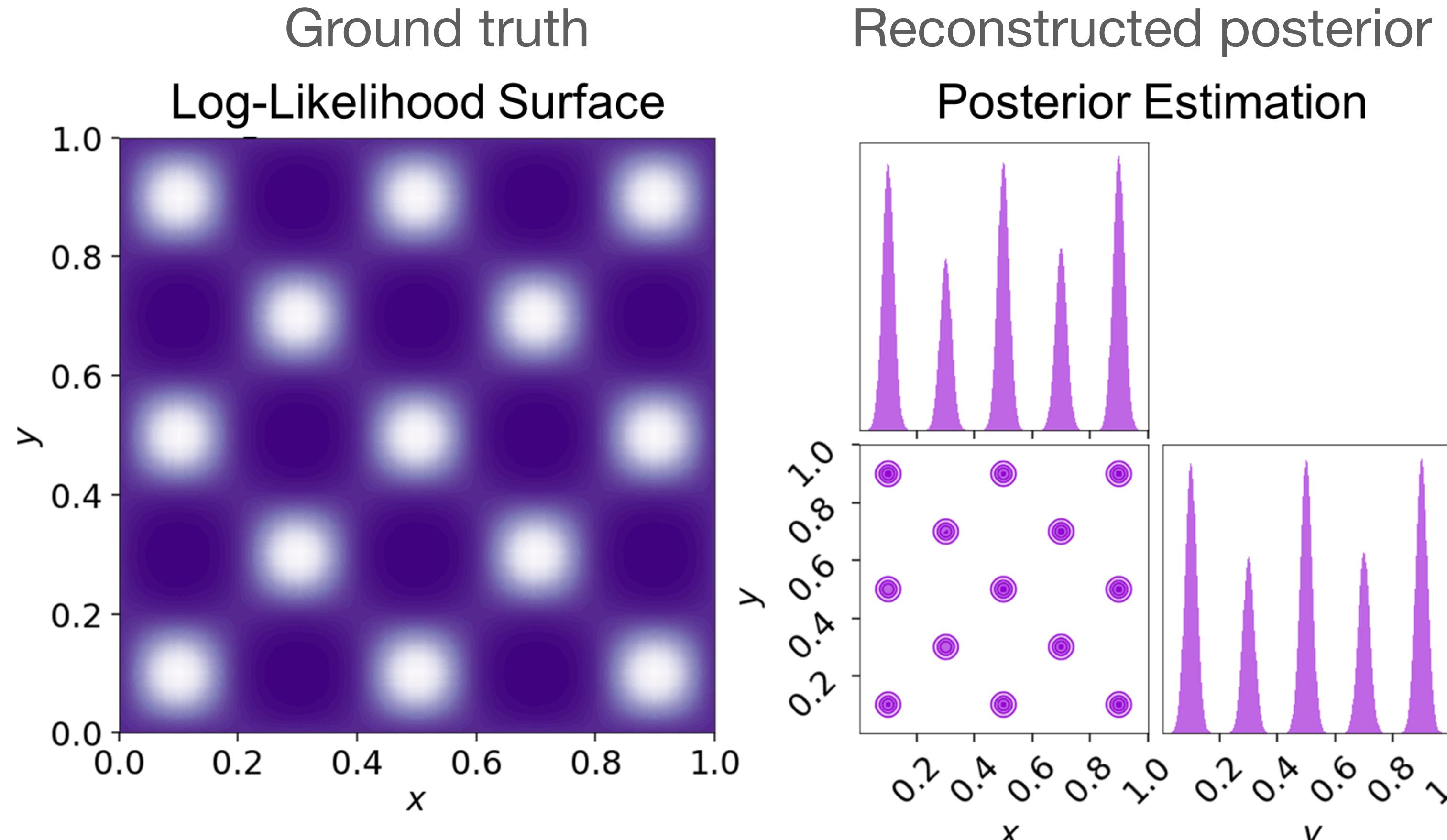


Random walk

Ashton, 2022

Complex landscapes

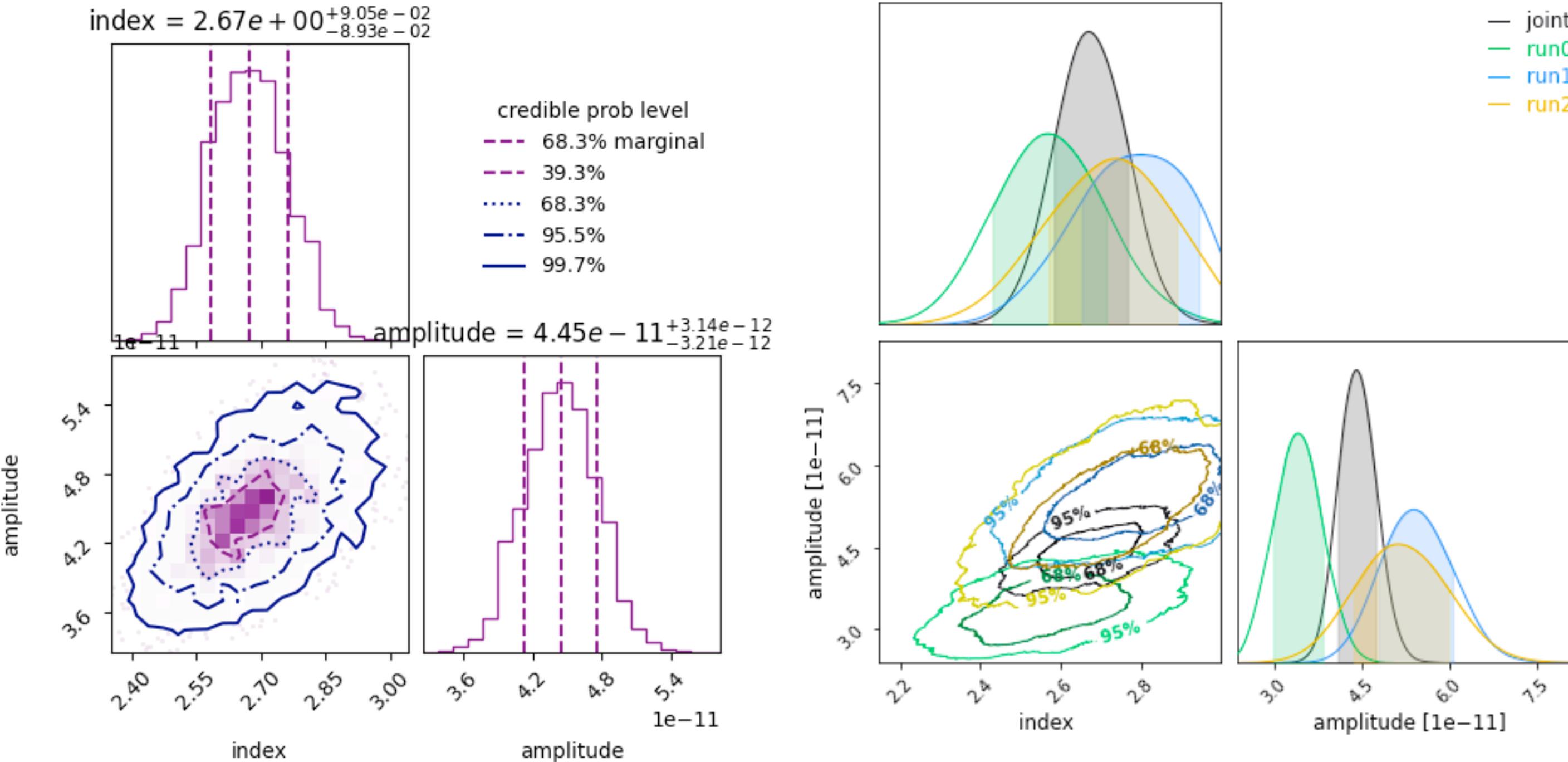
- Works perfectly well in landscapes that would drive crazy any MCMC or gradient descent method



Speagle, 2020

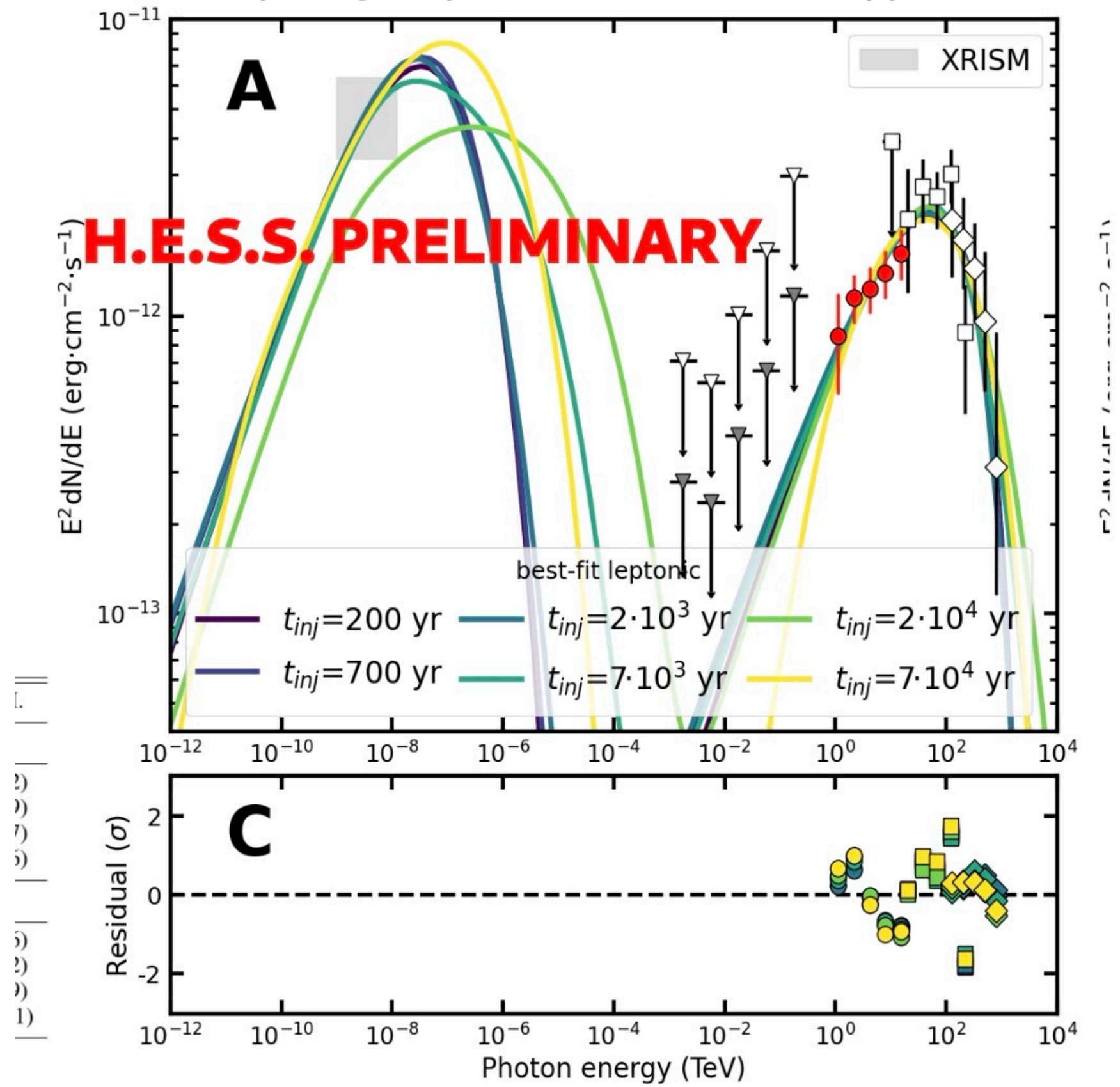
From simple to complex analyses

Crab spectral analysis

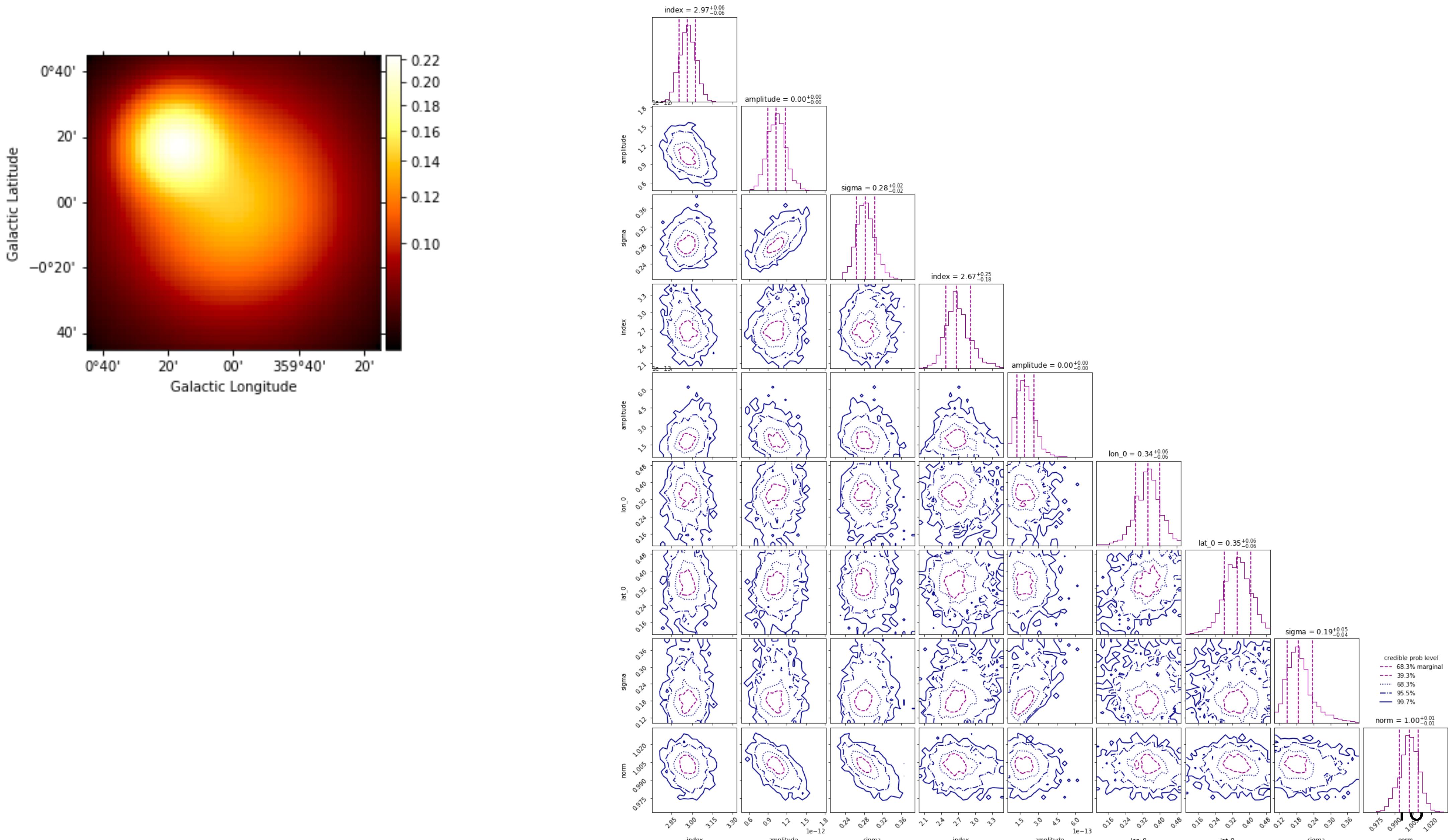


V4641 Sgr - Laura Olivera-Nieto - ICRC '25

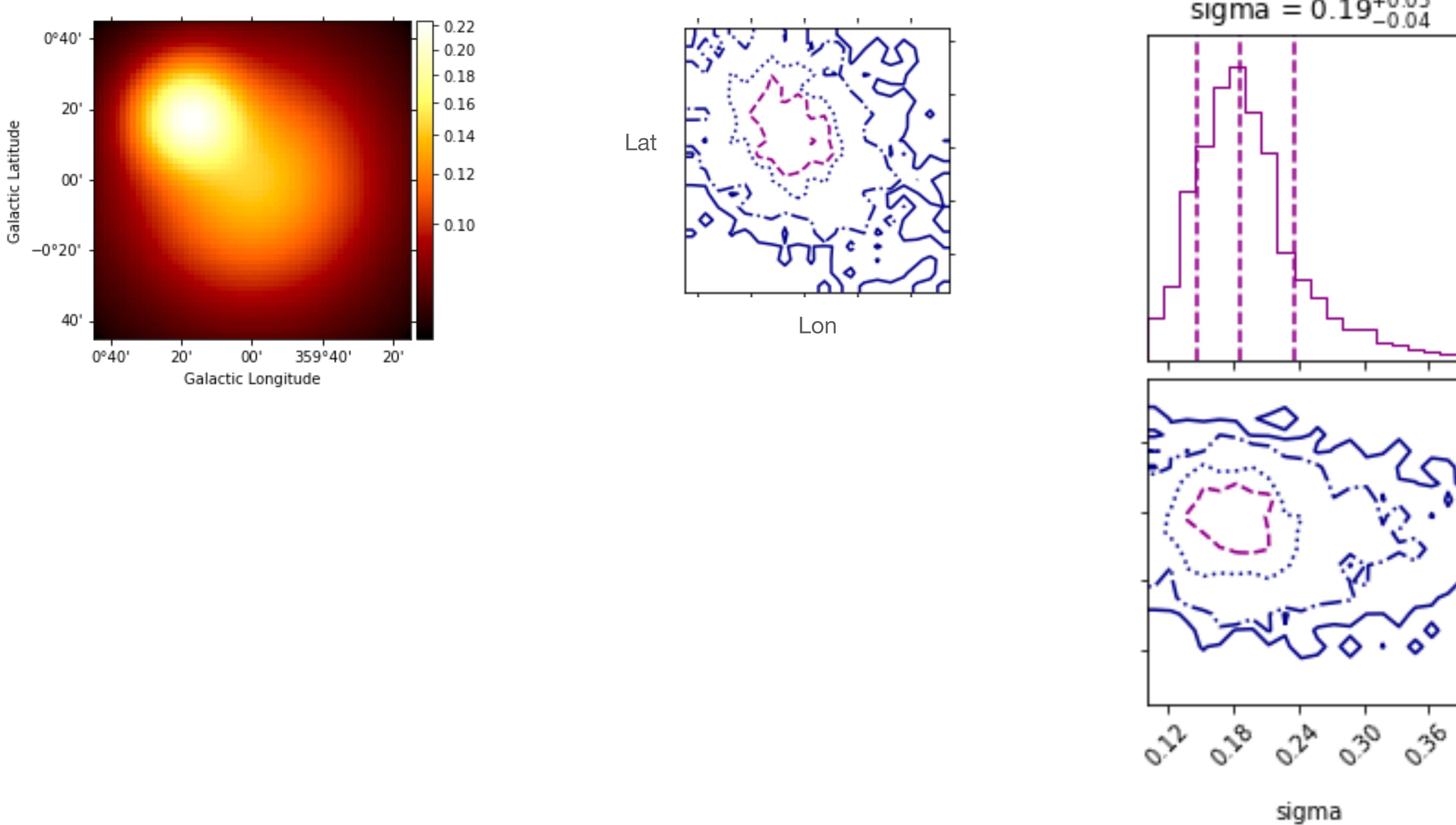
Fit using GAMERA and bayesian sampling implemented in Gammapy 1.3



Degeneracies in complex regions



Degeneracies in complex regions



Conclusion

- Nested Sampling Philosophy : Outside-Inside method, 500 independent live points init all over the parameter space (unlike 10-30 MCMC walkers init in small ball)
- Nested sample allows to lift several issues of MCMC techniques:
 - No need for init point. No burn-in period. No issue with multi-modal distribution.
 - NS has a well defined stopping criteria unlike MCMC
- In addition to estimating posterior distributions, calculate the Bayesian evidence Z :
 - This enables to compare different models that are not nested by comparing ΔZ between models
 - This is not possible with MCMC as only a local part of the likelihood is sampled
- More robust than traditional gradient descent fitting. Little human intervention needed to debug "failed fit".
- CONS: Need to define priors & slower but can be // easily but you get much more information than a simple fit.

Existing Nested sampling frameworks

| Code | Methods | Dynamic | Languages | Field | Pub. Year | Ashton, 2022 |
|----------------------------|--|-----------|-------------------------------------|---------------------|-----------|--------------|
| CosmoNest [60, 61] | ellipsoid | fixed | Fortran | Cosmology | 2006 | |
| MultiNest [48, 84] | multi-ellipsoid | fixed | Fortran, C/C++, Python | Cosmology | 2008 | |
| DIAMONDS [249] | multi-ellipsoid | fixed | C++ | Astrophysics | 2015 | |
| nestle [250] | ellipsoid, multi-ellipsoid | fixed | Python | Astrophysics | 2015 | |
| nessai [90, 91] | normalising flow ellipsoid | fixed | Python | Gravitational waves | 2021 | |
| (dy)PolyChord [53, 65] | slice | dynamic | Fortran, C/C++, Python | Cosmology | 2015 | |
| LALInferenceNest [180] | random walk, ensemble, differential evolution | fixed | C | Gravitational waves | 2015 | |
| Nested_fit [104, 257, 258] | random walk | fixed | Fortran | Atomic physics | 2016 | |
| cpnest [259] | slice, differential evolution, Gauss, Hamiltonian, ensemble | fixed | Python | Gravitational waves | 2017 | |
| pymatnest [44] | random walk, Galilean, symplectic Hamiltonian | fixed | Python | Materials | 2017 | |
| NNest [261] | normalising flow random walk | fixed | Python | Cosmology | 2019 | |
| DNest5 [55] | user-defined, random walk | diffusive | C++ | Astrophysics | 2020 | |
| BayesicFitting [263] | random walk, slice, Galilean, Gibbs | fixed | Python | Astronomy | 2021 | |
| dynesty [52] | ellipsoid, multi-ellipsoid, MLFriends & Gauss, slice, Hamiltonian | dynamic | Python | Astrophysics | 2020 | |
| UltraNest [92] | MLFriends + ellipsoid & Gauss, hit-and-run, slice | reactive | Python, Julia, R, C/C++, Fortran | Astrophysics | 2020 | |
| jaxns [266] | multi-ellipsoid & slice | fixed | jax | Astronomy | 2021 | |