

Simulation-based inference, Bayesian hierarchical models, and model misspecification



Learning the Universe Meeting

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Making inferences requires advanced Bayesian techniques

• Complex computer models are incorporated into Bayesian hierarchical models:



• The challenge: using new statistical methods is necessary. Two approaches are possible:



\bigcap	Simulation-based inference:	
	approximate statistical analysis	
	arbitrary data model	



Correlation functions versus field-level inference



$$f = \frac{1}{\alpha} \left[\exp\left(\alpha g - \frac{1}{2}\alpha^2\right) - 1 \right]$$





Gaussian field with 2PCF: $\xi_g(r) = \exp\left(-\frac{1}{4}\frac{r^2}{\beta^2}\right)$



- <u>2PCF likelihood-based analysis</u> is *imprecise* and *inaccurate*
- <u>2PCF simulation-based inference</u> is *imprecise* but *accurate*
- <u>Full-field data assimilation</u> is *precise* and *accurate*



Leclercq & Heavens, 2103.04158

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SBI, BHMs, and model misspecification: SELFI+BOLFI approach 15/09/2022

Companion repository:

https://github.com/florent-leclercg/correlations vs field

The issue of model misspecification in Bayesian inference and in simulation-based inference (SBI)

- <u>Model misspecification</u> arises when model differs from actual data-generating process.
- An example in cosmology: the galaxy *power spectrum*.

Due to observational systematics, we are unable to formulate *any* model that fits the data at large scales.

- Model misspecification: a major challenge particularly for approaches that marginalise over latent variables, such as <u>simulation-based inference</u> (SBI).
- Some recent work: diagnosing such issues and performing conservative belief updates in the presence of model misspecification, via e.g. tempering of the explicit likelihood (when it exists) or of a loss function.

Frazier, Robert & Rousseau 1708.01974, Thomas & Corander 1912.05810, Thomas *et al.* 2002.09377





A general class of Bayesian hierarchical models (BHMs): Complex observations of a latent function controlled by top-level parameters





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Key idea: a two-step SBI process that recycles simulations



- 1. Inference of the latent function θ , to check for model misspecification:
 - SELFI algorithm

Leclercq, MaxEnt 2022 proceedings (on arXiv next week)



Key idea: a two-step SBI process that recycles simulations



Leclercq, MaxEnt 2022 proceedings (on arXiv next week)

- 1. Inference of the latent function θ , to check for model misspecification:
 - SELFI algorithm
- 2. Simulation-based inference of ω :
 - Approximate Bayesian Computation (ABC), Likelihood-Free Rejection Sampling
 - Density/ratio estimation (DELFI / NRE)
 - Bayesian optimisation (BOLFI)
 - others...

Important: the simulations necessary for step **1**. are recycled for data compression, which is required for step **2**.



Latent function inference: the SELFI approach (Simulator Expansion for Likelihood-Free Inference)



We aim at inferring the latent function θ , which usually contains most/all of the information on ω .

(initial power spectrum in cosmology, prey/predator population functions in ecology)

- This requires doing SBI in $d = \mathcal{O}(100) \mathcal{O}(1,000)$
- If we trust the results of earlier experiments, we can Taylor-expand the black-box around an expansion point θ_0 :

$$\hat{\Phi}_{\theta} \approx \mathbf{f}_0 + \nabla \mathbf{f}_0 \cdot (\theta - \theta_0) + \frac{1}{2} (\theta - \theta_0)^{\mathsf{T}} \cdot \mathbf{H} \cdot (\theta - \theta_0) + \dots$$

SELFI-2 (second order): coming soon!

Gradients, Hessian matrix, etc. of the black-box can be evaluated via finite differences in parameter space.



Latent function inference: the SELFI approach (Simulator Expansion for Likelihood-Free Inference)



- Linearisation of the black-box: $\mathbf{\hat{\Phi}}_{\mathbf{\theta}} \approx \mathbf{f}_0 + \nabla \mathbf{f}_0 \cdot (\mathbf{\theta} - \mathbf{\theta}_0)$
- Further assume:
 - Gaussian prior: $\mathcal{P}(\mathbf{\theta}) = \mathcal{G}(\mathbf{\theta}_0, \mathbf{S})$
 - Gaussian effective likelihood: $\mathcal{P}(\mathbf{\Phi}|\mathbf{\theta}) = \mathcal{G}[\mathbf{f}(\mathbf{\theta}), \mathbf{C}_0]$
- Leclercq et al. 1902.10149



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• The posterior is Gaussian and analogous to a Wiener filter:

 $\begin{array}{ll} \mbox{expansion point} & \mbox{observed summaries} \\ \mbox{mean:} \ensuremath{\boldsymbol{\gamma}} \equiv \ensuremath{\boldsymbol{\theta}}_0 + \ensuremath{\boldsymbol{\Gamma}} (\nabla \mathbf{f}_0)^\intercal \ensuremath{\mathbf{C}}_0^{-1} (\ensuremath{\boldsymbol{\Phi}}_O - \ensuremath{\mathbf{f}}_0) \\ \mbox{covariance:} \ensuremath{\boldsymbol{\Gamma}} \equiv \left[(\nabla \mathbf{f}_0)^\intercal \ensuremath{\mathbf{C}}_0^{-1} \nabla \mathbf{f}_0 + \ensuremath{\mathbf{S}}_{-1}^{-1} \right]^{-1} \\ \mbox{covariance of summaries} \\ \ensuremath{\text{gradient of the black-box}} \end{array}$

- f_0, C_0 and ∇f_0 can be evaluated through simulations only.
- The number of required simulations is fixed *α priori* (contrary to MCMC).
- The workload is perfectly parallel.

Check for model misspecification and data compression for SBI

 $\mathcal{P}(\boldsymbol{\omega})$

ω

θ

 $\mathcal{P}(\mathbf{\Phi}|\mathbf{\theta})$

 $\mathbf{\Phi}$

 $\widetilde{\omega}$

- Qualitatively: the shape of the reconstructed θ is useful as a <u>check for</u> <u>model misspecification</u> (independent theoretical understanding).
- Quantitatively: we can use the Mahalanobis distance between the reconstruction γ and the prior distribution 𝒫(θ):

 $d_{\mathrm{M}}(\boldsymbol{\gamma}, \boldsymbol{\theta}_{0} | \mathbf{S}) \equiv \sqrt{(\boldsymbol{\gamma} - \boldsymbol{\theta}_{0})^{\mathsf{T}} \mathbf{S}^{-1} (\boldsymbol{\gamma} - \boldsymbol{\theta}_{0})}$

Leclercq, MaxEnt 2022 proceedings (on arXiv next week)



for the parameters is

The score function $\nabla_{\alpha}\ell_{\alpha,0}$ is the

gradient of the log-likelihood at

Fisher matrix: $\mathbf{F}_0 = (\nabla_{\boldsymbol{\omega}} \mathbf{f}_0)^{\mathsf{T}} \mathbf{C}_0^{-1} \nabla_{\boldsymbol{\omega}} \mathbf{f}_0$

fiducial point $\boldsymbol{\omega}_0$ in parameter space.

A guasi maximum-likelihood estimator

 $\mathcal{C}(\mathbf{\Phi}) = \widetilde{\mathbf{\omega}} \equiv \mathbf{\omega}_0 + \mathbf{F}_0^{-1} \left[(\nabla_{\mathbf{\omega}} \mathbf{f}_0)^{\mathsf{T}} \mathbf{C}_0^{-1} (\mathbf{\Phi} - \mathbf{f}_0) \right]$

 $\nabla_{\boldsymbol{\omega}} \mathbf{f}_0 = \nabla \mathbf{f}_0 \cdot \nabla_{\boldsymbol{\omega}} \mathcal{T}_0$

Already computed

for SELFI

Score compression is optimal in the sense that it <u>preserves the Fisher</u> information content of the data.

Cheap via finite

differences

Alsing & Wandelt, 1712.00012

Simulation-based inference of top-level target parameters



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• Any SBI algorithm can be used to obtain the posterior $\mathscr{P}(\boldsymbol{\omega}|\widetilde{\boldsymbol{\omega}}_{\mathrm{O}})$.

Final inference:

- does not depend on the assumptions made to check for model misspecification,
- is unbiased (only more conservative) in case data compression is lossy.
- Non-parametric approaches can use the <u>Fisher-Rao distance</u> between simulated summaries w and observed summaries w_O:

$$d_{\rm FR}(\widetilde{\boldsymbol{\omega}},\widetilde{\boldsymbol{\omega}}_{\rm O}) \equiv \sqrt{(\widetilde{\boldsymbol{\omega}}-\widetilde{\boldsymbol{\omega}}_{\rm O})^{\mathsf{T}} \mathbf{F}_0(\widetilde{\boldsymbol{\omega}}-\widetilde{\boldsymbol{\omega}}_{\rm O})}$$



Prey-predator model: inference of target population parameters using Likelihood-Free Rejection Sampling



Simulation-efficient inference of target parameters: BOLFI: Data acquisition

 Simulations are obtained from sampling an adaptively-constructed proposal distribution, using the regressed effective likelihood



F. Nogueira, https://github.com/fmfn/BayesianOptimization



Simulation-efficient inference of target parameters: BOLFI: Re-analysis of the JLA supernova sample



- The number of required simulations is reduced by:
 - 2 orders of magnitude with respect to likelihood-free rejection sampling (for a much better approximation of the posterior)
 - 3 orders of magnitude with respect to exact Markov Chain Monte Carlo sampling
- Bayesian optimisation can also be applied to the "true" likelihood (if known) or to build iteratively an emulator of the data model

Leclercq 2018, 1805.07152



Conclusions

- A novel <u>two-step simulation based Bayesian approach</u>, combining SELFI and SBI, to tackle the issue of model misspecification for a large class of BHMs.
- Advantages of the first step (SELFI):
 - Even if the inference is in high dimension, the simulator remains a black-box.
 - The number of simulations is fixed *a priori* by the user.
 - The computational workload is perfectly parallel.
 - The linearised data model is trained once and for all independently of the data vector (amortisation).
- Advantages of the second step (SBI/BOLFI):
 - SELFI quantities provide a score compressor for free.
 - Inference does not depend on the assumptions made to check for model misspecification.
 - General advantages of SBI with respect to likelihood-based methods are preserved.
 - BOLFI is particularly suitable for very expensive simulators.

A computationally efficient and easily applicable framework to perform <u>SBI of BHMs while</u> <u>checking for model misspecification</u>.

pySELFI is publicly available at <u>https://pyselfi.florent-leclercq.eu</u>.



Backup slides



A prey-predator model with observational effects





A prey-predator model with observational effects



Prey-predator model: diagnostics of the linearised black-box





Prey-predator model: inference of the latent population functions



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A family of priors for population functions in prey-predator systems

Assumptions:

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- 1. The population functions θ are Gaussiandistributed.
- 2. They are strongly constrained to live close to $\theta_0 = T(\boldsymbol{\omega}_0)$.
- 3. x(t) and y(t) are smooth functions of time.
- 4. The uncertainty on x(t) and y(t) grows with time.
- Gaussian prior: Overall prior uncertainty mean: θ_0 covariance: $\mathbf{S} \equiv \alpha_{norm}^2 \mathbf{K} \circ \mathbf{V}$

 $(\mathbf{K}_z)_{ij} \equiv \left[-\frac{1}{2} \left(\frac{t_i - t_j}{t_{\text{smooth}}} \right)^2 \right]$



• The 3 free hyperparameters $\{\alpha_{norm}, t_{smooth}, t_{chaos}\}$ can be optimised using simulations.

Smoothness of the population function Chaotic behaviour of the system

$$\begin{pmatrix} t_i - t_j \\ \hline t_{\text{smooth}} \end{pmatrix}^2 \qquad \mathbf{K} \equiv \begin{pmatrix} \mathbf{K}_x & 0 \\ 0 & \mathbf{K}_y \end{pmatrix} \qquad \mathbf{V} \equiv \begin{pmatrix} x_0 \, \mathbf{u} \mathbf{u}^{\mathsf{T}} & 0 \\ 0 & y_0 \, \mathbf{u} \mathbf{u}^{\mathsf{T}} \end{pmatrix} \qquad (\mathbf{u})_i \equiv 1 + \frac{t_i}{\overline{t_{\text{chaos}}}}$$

