



Simulation-based inference of Bayesian hierarchical models while checking for model misspecification



MaxEnt 2022 conference
Institut Henri Poincaré, Paris

Florent Leclercq

www.florent-leclercq.eu

Institut d'Astrophysique de Paris
CNRS & Sorbonne Université

In collaboration with the Aquila Consortium

www.aquila-consortium.org

19 July 2022

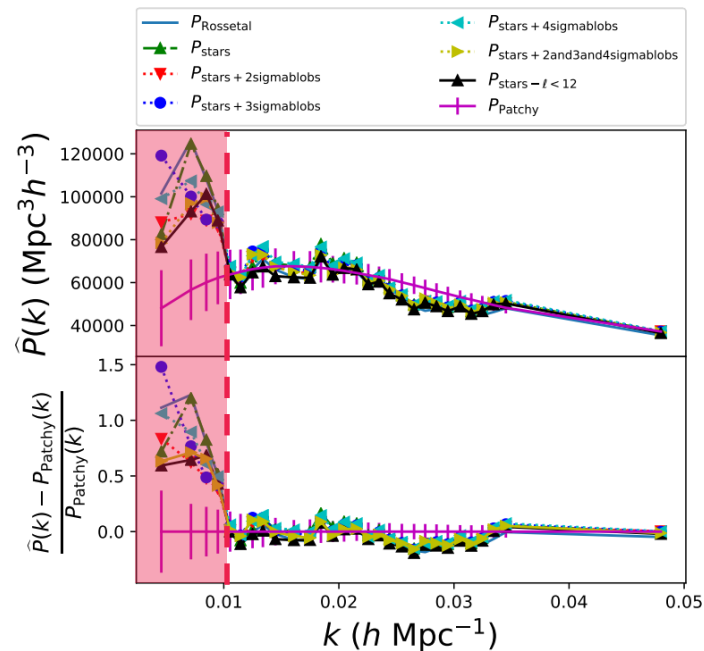
The issue of model misspecification in Bayesian inference and in simulation-based inference (SBI)

- [Model misspecification](#) 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.

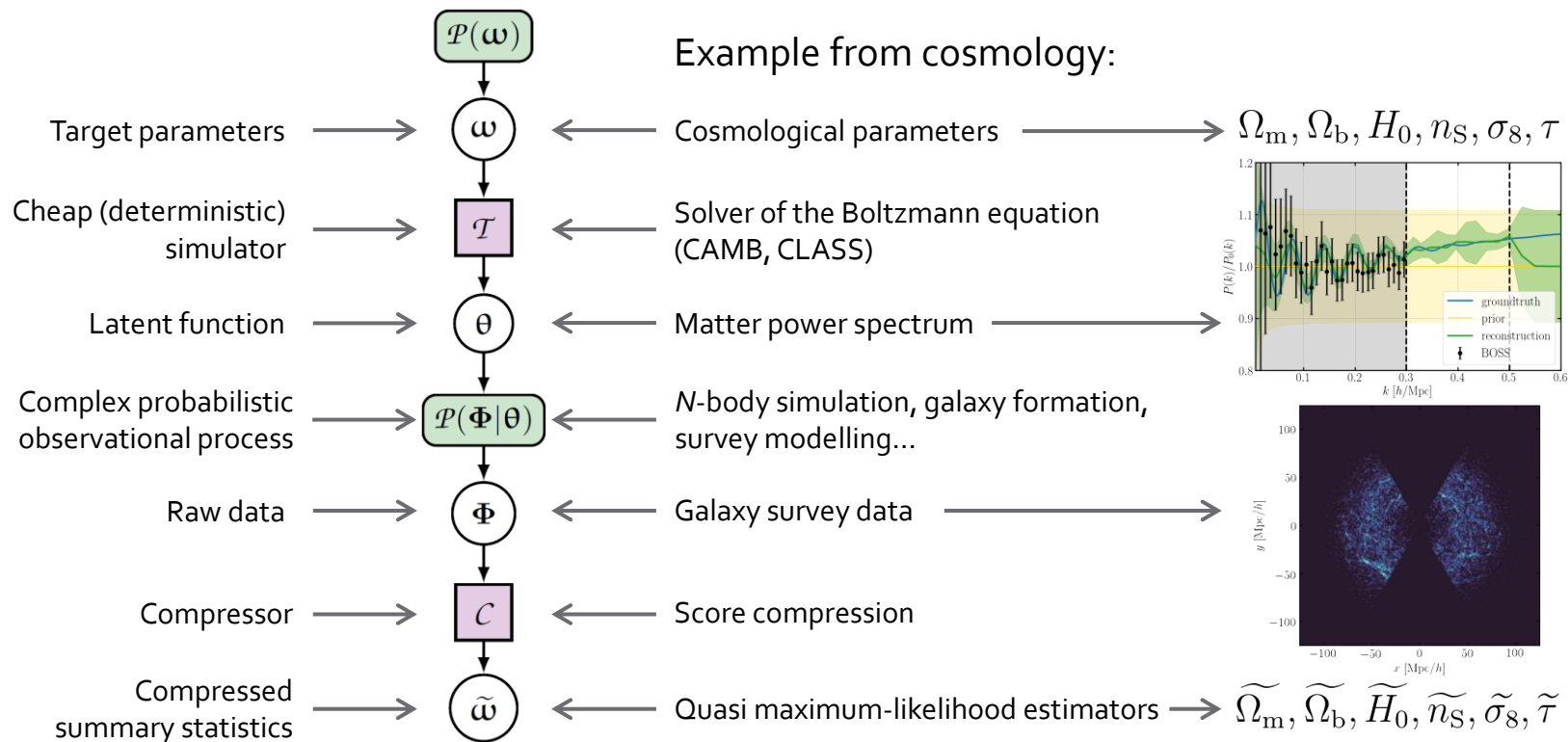
- Model misspecification: a major challenge particularly for approaches that marginalise over latent variables, such as [simulation-based inference](#) (SBI).
- Some recent work: [diagnosing such issues](#) and [performing conservative belief updates](#) in the presence of model misspecification, via 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

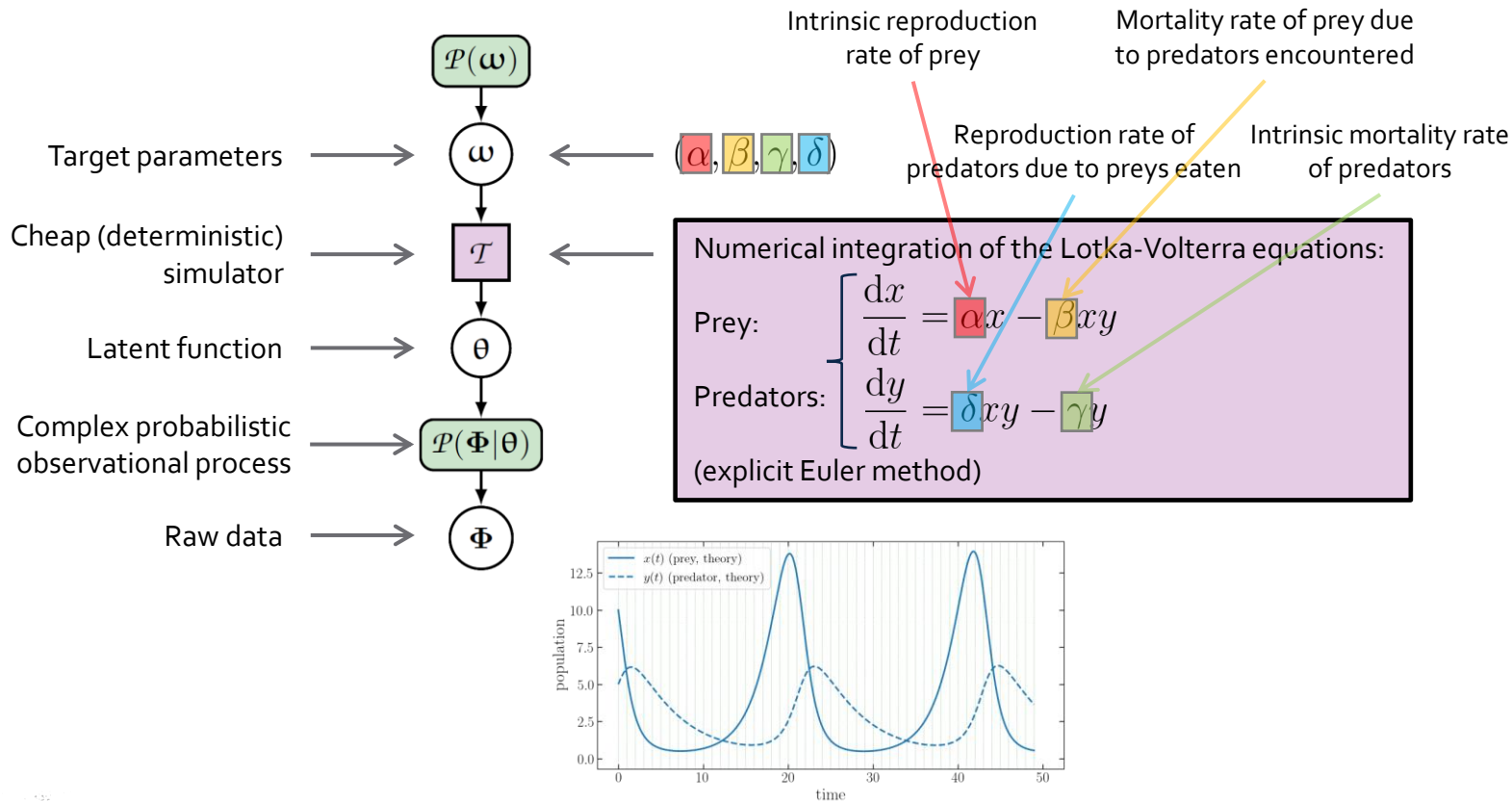


Kalus et al. 1806.02789

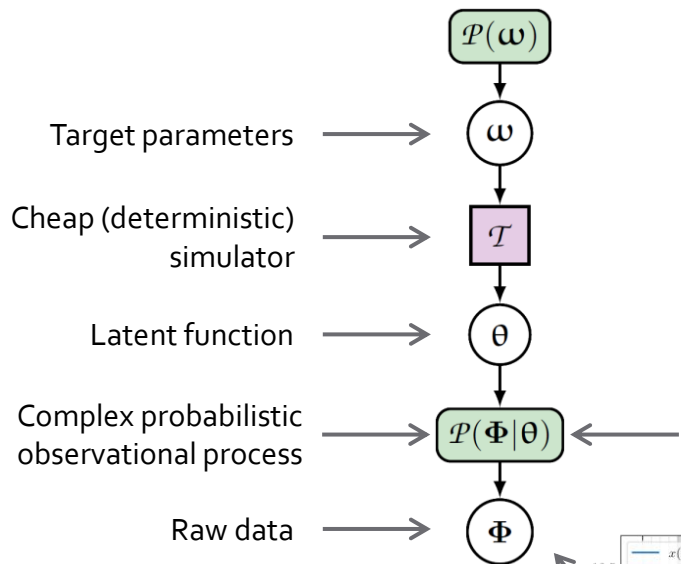
A general class of Bayesian hierarchical models (BHM): Complex observations of a latent function controlled by top-level parameters



A prey-predator model with observational effects



A prey-predator model with observational effects



Model A (correct):

- Signal:** a retarded a non-linear observation of the true functions

$$\begin{cases} s_x(t_{i+1}) = e_x(t_i) [x(t_i) - p x(t_i)y(t_i) + q x(t_i)^2] \\ s_y(t_{i+1}) = y(t_i) + p x(t_i)y(t_i) - q y(t_i)^2 \end{cases}$$

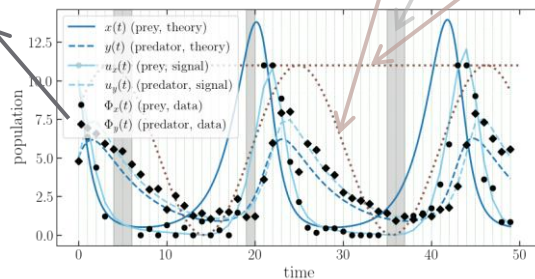
- Noise:** additive Gaussian noise with zero mean

$$\begin{cases} u_x(t) = s_x(t) + n_x^D(t) + n_x^O(t) \\ u_y(t) = s_y(t) + n_y^D(t) + n_y^O(t) \end{cases}$$

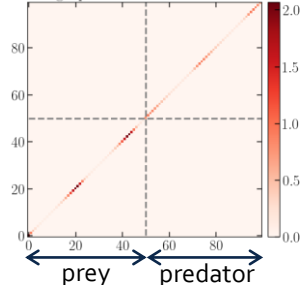
$$\begin{aligned} n_x^D(t) &\sim \mathcal{G}\left[0, \begin{pmatrix} r x(t) & t \sqrt{x(t)y(t)} \\ t \sqrt{x(t)y(t)} & x(t) \end{pmatrix}\right] \\ n_y^D(t) &\sim \mathcal{G}\left[0, \begin{pmatrix} r y(t) & t \sqrt{x(t)y(t)} \\ t \sqrt{x(t)y(t)} & y(t) \end{pmatrix}\right] \end{aligned}$$

- Censoring:** a mask and a threshold

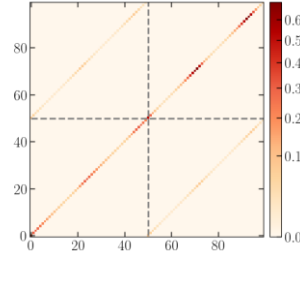
- Model B (misspecified):** assumes the true functions are directly observed, does not account for observational noise nor threshold.



Demographic noise covariance matrix

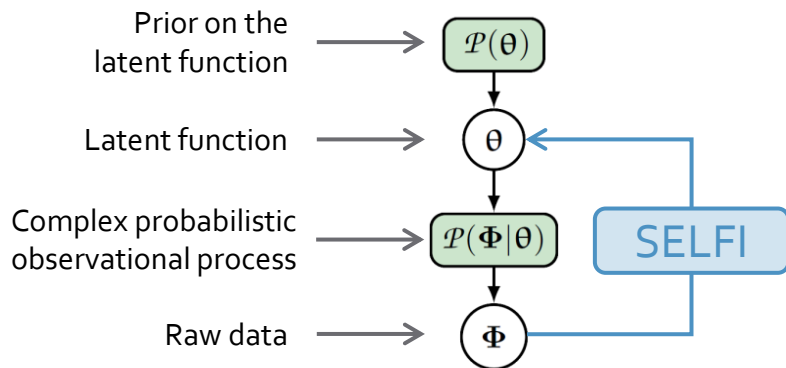


Observational noise covariance matrix

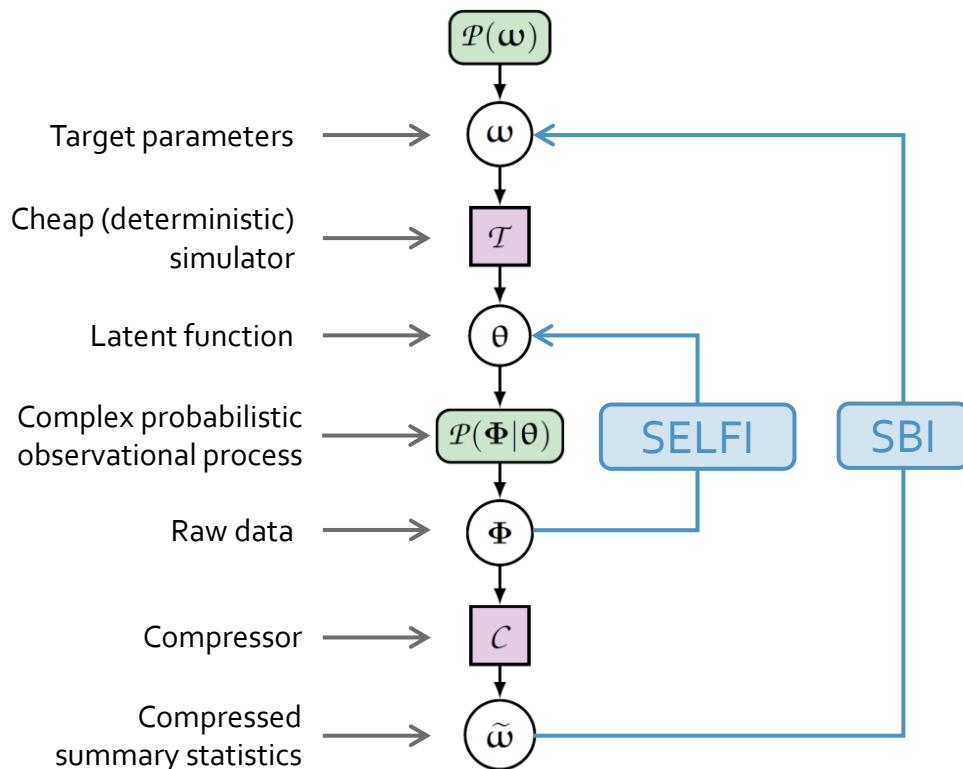


Key idea: a two-step SBI process that recycles simulations

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



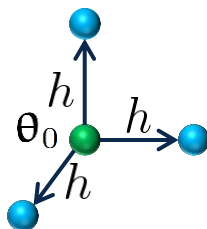
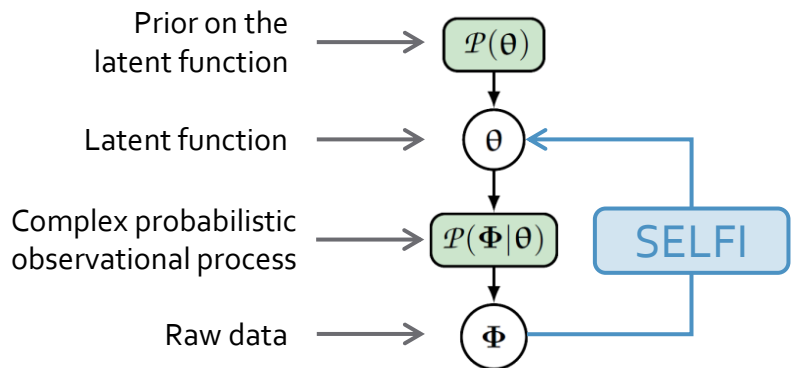
Key idea: a two-step SBI process that recycles simulations



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

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

SELFI-2 (second order): coming soon!
- Gradients, Hessian matrix, etc. of the black-box can be evaluated via finite differences in parameter space.

Leclercq et al. 1902.10149



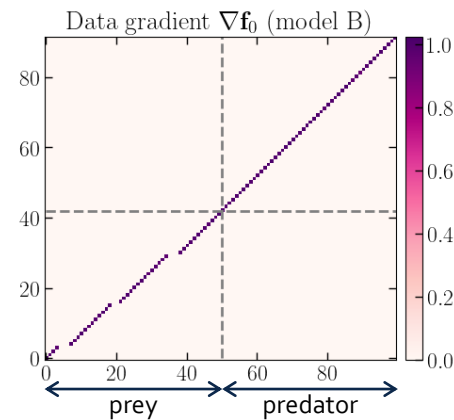
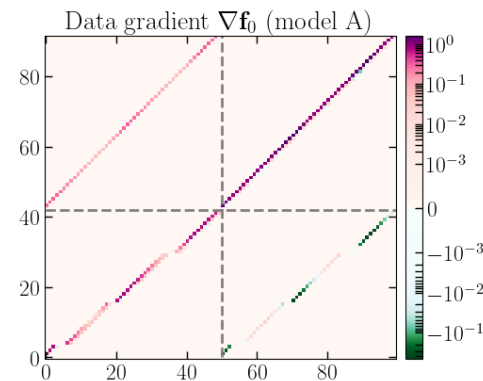
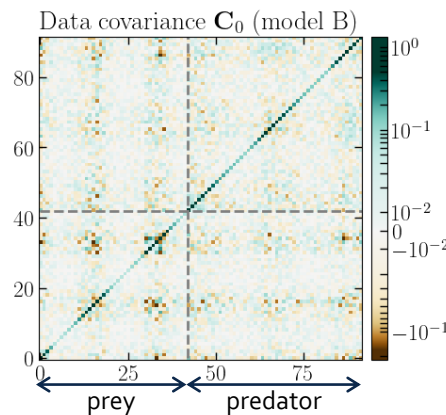
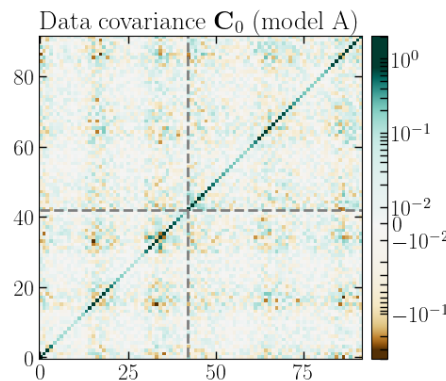
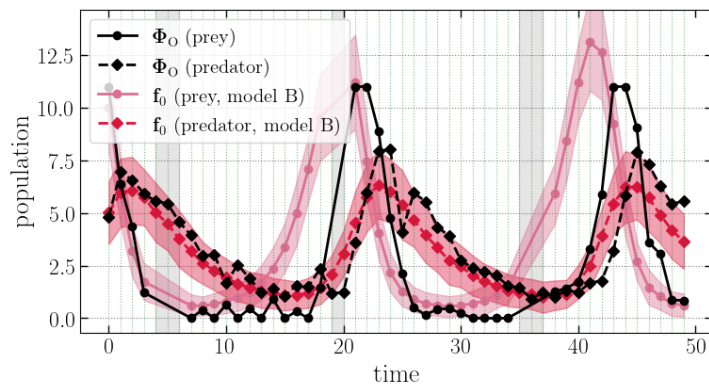
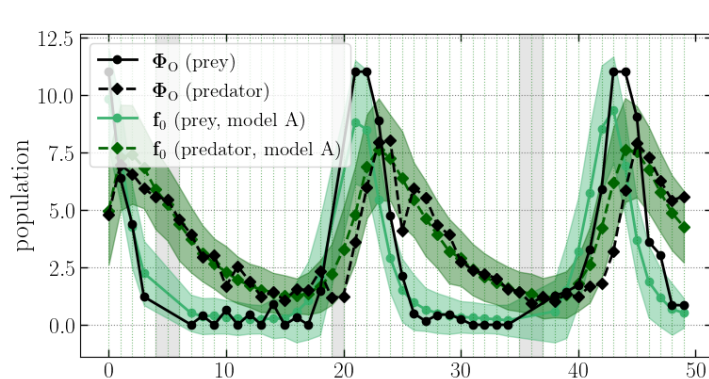
Florent Leclercq

SBI of BHM's while checking for model misspecification

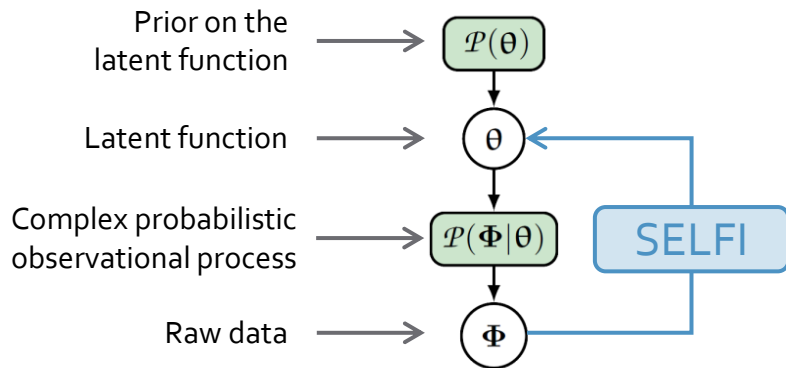
19/07/2022

8

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



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



- Linearisation of the black-box:

$$\hat{\Phi}_{\theta} \approx \mathbf{f}_0 + \nabla \mathbf{f}_0 \cdot (\theta - \theta_0)$$

- Further assume:

- Gaussian prior: $\mathcal{P}(\theta) = \mathcal{G}(\theta_0, \mathbf{S})$
- Gaussian effective likelihood: $\mathcal{P}(\Phi|\theta) = \mathcal{G}[\mathbf{f}(\theta), \mathbf{C}_0]$

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

expansion point observed summaries

$$\text{mean: } \gamma \equiv \theta_0 + \mathbf{\Gamma} (\nabla \mathbf{f}_0)^\top \mathbf{C}_0^{-1} (\Phi_O - \mathbf{f}_0)$$

$$\text{covariance: } \mathbf{\Gamma} \equiv [(\nabla \mathbf{f}_0)^\top \mathbf{C}_0^{-1} \nabla \mathbf{f}_0 + \mathbf{S}^{-1}]^{-1}$$

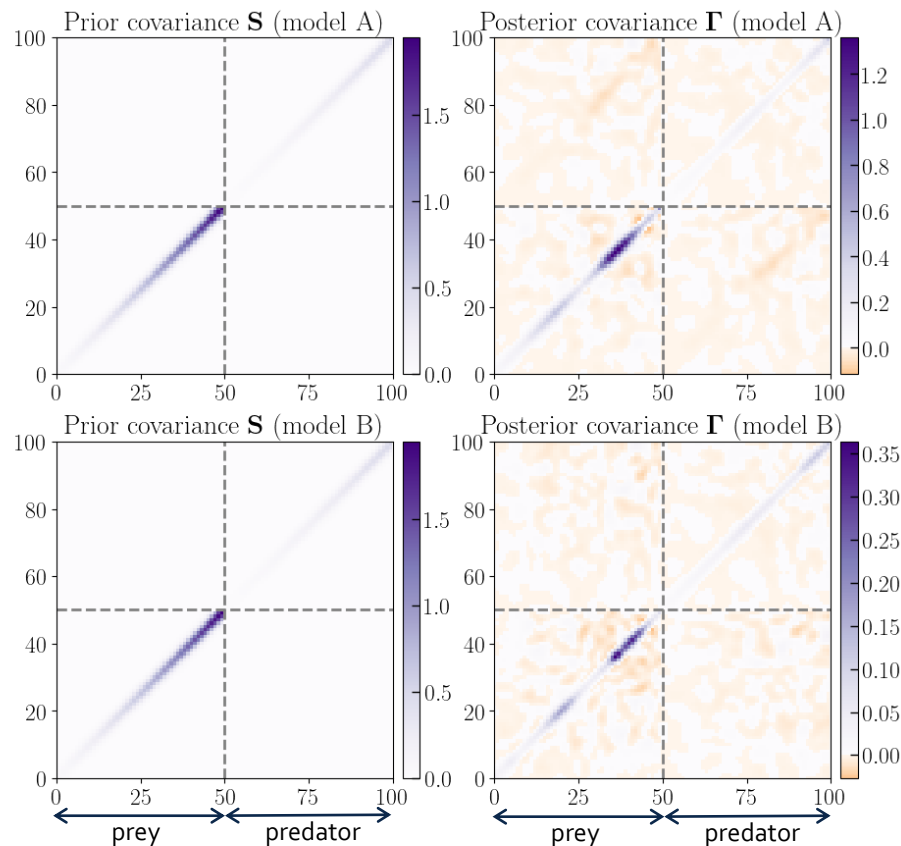
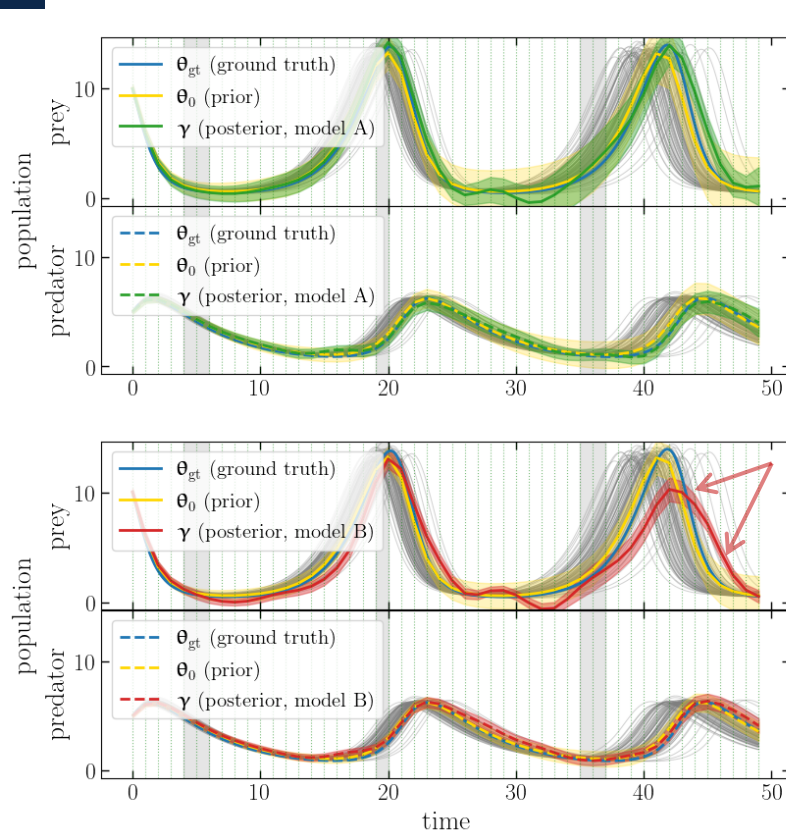
covariance of summaries gradient of the black-box prior covariance

- \mathbf{f}_0 , \mathbf{C}_0 and $\nabla \mathbf{f}_0$ can be evaluated through simulations only.
- The number of required simulations is fixed *a priori* (contrary to MCMC).
- The workload is perfectly parallel.

Leclercq *et al.* 1902.10149



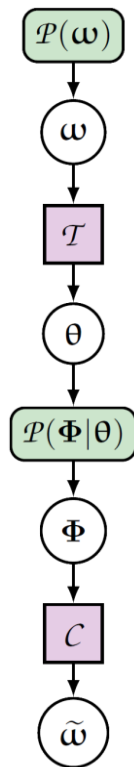
Prey-predator model: inference of the latent population functions



- Qualitatively: the shape of the reconstructed θ is useful as a [check for model misspecification](#) (independent theoretical understanding).
- Quantitatively: we can use the Mahalanobis distance between the reconstruction γ and the prior distribution $\mathcal{P}(\theta)$:

$$d_M(\gamma, \theta_0 | \mathbf{S}) \equiv \sqrt{(\gamma - \theta_0)^\top \mathbf{S}^{-1} (\gamma - \theta_0)}$$

- In the example:
 - $d_M(\gamma, \theta_0 | \mathbf{S}) \approx 5.35$ for model A
 - $d_M(\gamma, \theta_0 | \mathbf{S}) \approx 12.54$ for model B
 - $\langle d_M(\mathcal{T}(\omega), \theta_0 | \mathbf{S}) \rangle \approx 9.43$ in fiducial simulations



- The score function $\nabla_{\omega} \hat{\ell}_{\omega_0}$ is the gradient of the log-likelihood at fiducial point ω_0 in parameter space.
- A quasi maximum-likelihood estimator for the parameters is

$$\mathcal{C}(\Phi) = \tilde{\omega} \equiv \omega_0 + \mathbf{F}_0^{-1} [(\nabla_{\omega} \mathbf{f}_0)^\top \mathbf{C}_0^{-1} (\Phi - \mathbf{f}_0)]$$

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

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

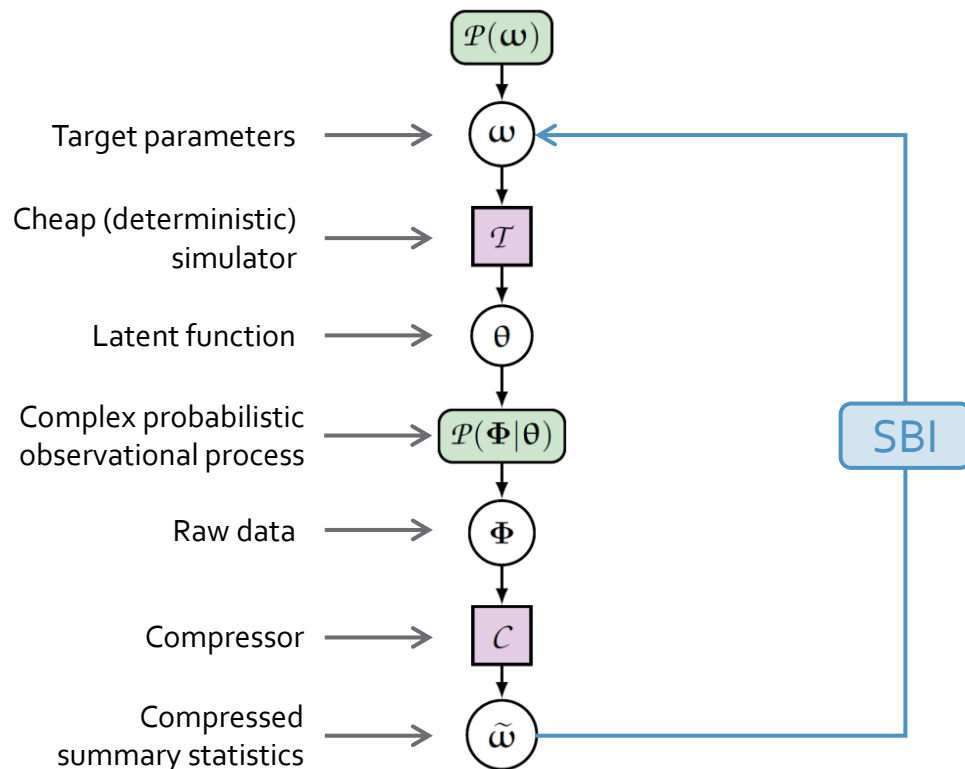
Already computed
for SELF1

Cheap via finite
differences

- Score compression is optimal in the sense that it [preserves the Fisher information content](#) of the data.

Alsing & Wandelt, 1712.00012

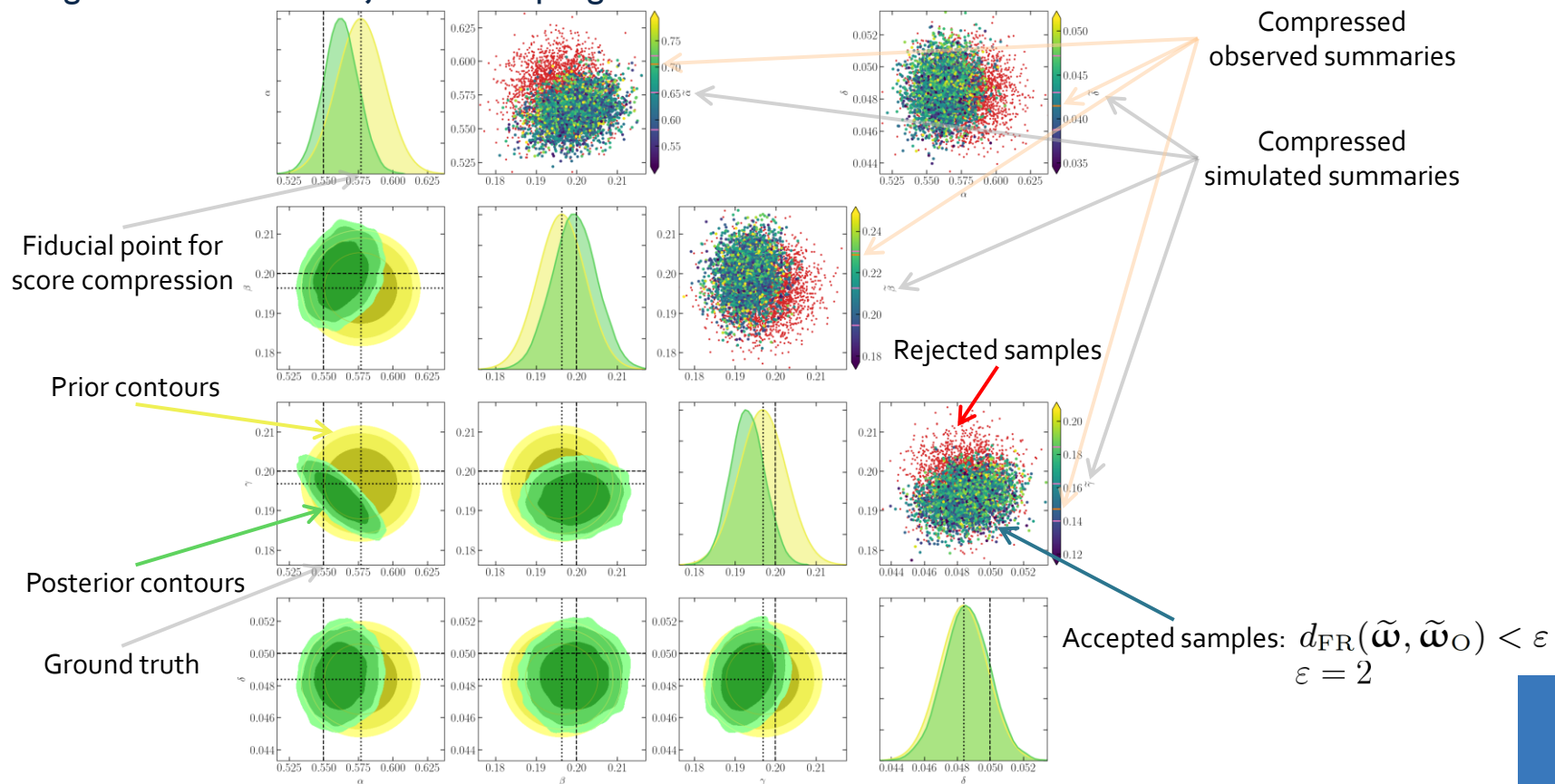
Simulation-based inference of top-level target parameters



- Any SBI algorithm can be used to obtain the posterior $\mathcal{P}(\omega|\tilde{\omega}_O)$.
- 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 [Fisher-Rao distance](#) between simulated summaries $\tilde{\omega}$ and observed summaries $\tilde{\omega}_O$:

$$d_{\text{FR}}(\tilde{\omega}, \tilde{\omega}_O) \equiv \sqrt{(\tilde{\omega} - \tilde{\omega}_O)^\top \mathbf{F}_0 (\tilde{\omega} - \tilde{\omega}_O)}$$

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



- A novel [two-step simulation based Bayesian approach](#), combining SELFI and SBI, to tackle the issue of model misspecification for a large class of BHM.
 - 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):
 - SELFI quantities provide a score compressor for free.
 - General advantages of SBI with respect to likelihood-based methods are preserved.
 - Inference does not depend on the assumptions made to check for model misspecification.
- A computationally efficient and easily applicable framework to perform [SBI of BHMs while checking for model misspecification](#).

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

Backup slides

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

Assumptions:

1. The population functions θ are Gaussian-distributed.
2. They are strongly constrained to live close to $\theta_0 = \mathcal{T}(\omega_0)$.
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:

mean: θ_0

covariance: $S \equiv \alpha_{\text{norm}}^2 \mathbf{K} \circ \mathbf{V}$

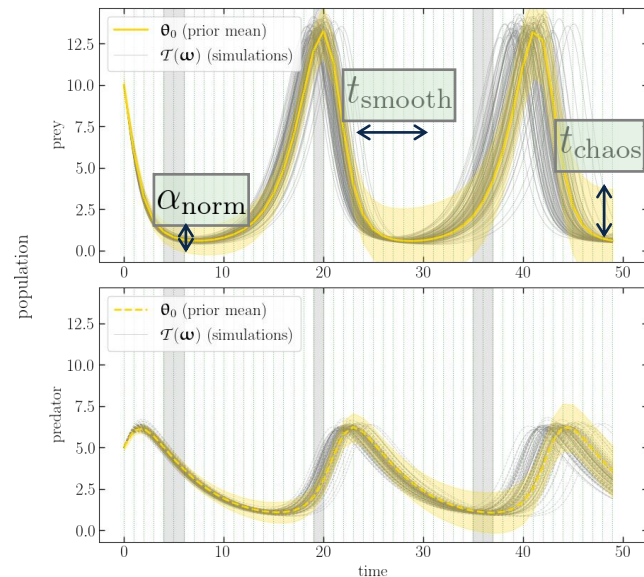
$$(\mathbf{K}_z)_{ij} \equiv \left[-\frac{1}{2} \left(\frac{(t_i - t_j)^2}{t_{\text{smooth}}^2} \right) \right]$$

Smoothness of the population function Chaotic behaviour of the system

$$\mathbf{K} \equiv \begin{pmatrix} \mathbf{K}_x & 0 \\ 0 & \mathbf{K}_y \end{pmatrix}$$

$$\mathbf{V} \equiv \begin{pmatrix} x_0 \mathbf{u} \mathbf{u}^\top & 0 \\ 0 & y_0 \mathbf{u} \mathbf{u}^\top \end{pmatrix}$$

$$(\mathbf{u})_i \equiv 1 + \frac{t_i}{t_{\text{chaos}}}$$



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

