# Beyond Generalised Bayes: Prediction-Centric Alternatives





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#### Chris. J. Oates

#### https://postbayes.github.io/seminar/

March 2025

Different communities use different conventions and standards in defining a "model":

discrete/continuous;

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- deterministic/stochastic;
- based on mathematical equations/computer simulation;





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SimBoal What Is FEA | Finite Element Analysis ...



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Understanding the Finite Element Method ...

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Finite Element Method ...





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 MathiWorks What is Finite Element Analysis ...





The Finite Element Method (FEM) - A ....





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A Blog - Spatial Corp. What is Finite Element Analysis ... An Introduction to Finite Element Modeling

The Finite Element Method (FEM) - ...

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CAD/CAM Services 3D CAD Models used in Design Proce...



The example of GAD model [5 ...

Professional 3D CAD Modeling Softw...



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Fast CAD Model Comparison tool





SD-Ace Difference Between GAD and 3D Mod ....



Siemens Digital Industries Software Blogs

CAD preparation for CFD simulation ....



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Focus on deterministic models

$$M_{\theta}: \mathcal{X} \to \mathcal{Y}$$

with parameters denoted  $\theta \in \Theta$ .

**Challenge:** How to use such a "model" for statistical inference and (causal) prediction?

**Usual Solution:** Turn the "model"  $M_{\theta}$  into a "statistical model"

$$P_{\theta}$$
:  $y_i = M_{\theta}(x_i) + \epsilon_i, \qquad \epsilon_i \stackrel{\text{iid}}{\sim} \mathcal{N}(0, \sigma^2 I)$ 

using knowledge of the equipment used to make the measurement.

Unfortunately a good "model" can lead to a misspecified "statistical model"..

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Figure: ERK signalling model.

Systems biology has invested decades of effort into the design of detailed ODE descriptions of cellular signalling pathways, with thousands of models hosted on repositories such as BioModels [Malik-Sheriff et al., 2020].

#### e.g. ERK signalling is modelled as

 $\frac{\mathrm{d}u}{\mathrm{d}x} = f_{\theta}(x, u), \qquad \theta \in \mathbb{R}^{11}$ 

Data are (reasonably, as far as this talk is concerned) treated as noisy observations of molecular concentrations u(x) at discrete times  $x_{1:n}$ .

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#### Other Choices of $L_n$ and $\lambda_n$

For various other choices of L<sub>n</sub> and  $\lambda_n$ , generalised Bayesian methods can produce robust posteriors suitable for dealing with certain forms of statistical model misspecification [see Hooker and Vidyashankar, 2014, Ghosh and Basu, 2016, Knoblauch et al., 2018, Schmon et al., 2020, Chérief-Abdellatif and Alquier, 2020, Dellaporta et al., 2022, Husain and Knoblauch, 2022, Altamirano et al., 2023, 2024, Duran-Martin et al., 2024].
# TL/DR: Generalised Bayesian Inference is a Bit Complicated

Generalisations of Bayesian inference have been proposed for when the model is misspecified.

- e.g. generalised Bayesian inference [Bissiri et al., 2016, Knoblauch et al., 2022]:
  - **>** parameter  $\theta \in \Theta$
  - ▶ IID data  $y_{1:n} = (y_1, \ldots, y_n) \in \mathcal{Y}^n$
  - ▶ loss function  $L_n : \Theta \times \mathcal{Y}^n \to \mathbb{R}$
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#### **Concentration of Generalised Posterior**

For convenient choices of  $L_n$  and  $\lambda_n$ , the magnitude of the data-fit term generally increases with *n*. As a result,  $Q_n^{\dagger} \rightarrow \delta_{\theta^{\dagger}}$  [Miller, 2021].

So need to tune the learning rate...

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#### **Tuning the Learning Rate**

Several ideas have been proposed to select the learning rate  $\lambda_n$  - c.f. Ryan Martin's talk. But these involve approximations and/or can be computationally demanding.

Tuning the learning rate is complicated - seek alternative to generalised Bayes...?

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### Setting Given a model $P_{\theta}$ that is useful (e.g. for causal prediction) but misspecified.

Step 1: Mitigate Misspecification Form a mixture model

$${\sf P}_Q = \int {\sf P}_ heta \, \mathrm{d} {\sf Q}( heta) \in {\mathcal P}({\mathcal Y}).$$

**Step 2: Learn** Q For example, by matching the predictive distribution of  $P_Q$  to the dataset.

#### Example: Nonparametric Maximum Likelihood

This approach solves

$$\underset{Q \in \mathcal{P}(\Theta)}{\operatorname{arg\,min}} - \frac{1}{n} \sum_{i=1}^{n} \log p_Q(y_i)$$

where  $p_Q$  is a density for the mixture model  $P_Q$  [see Chapter 5 of Lindsay, 1995].

- ▶ approximates  $KL(P_*, P_Q)$  when  $y_{1:n}$  is a collection of *n* independent samples from  $P_* \in \mathcal{P}(\mathcal{Y})$
- lack of regularisation causes computational difficulties and non-identifiability [see e.g. Laird, 1978], as the minimising measure will generally be fully atomic, see Lindsay [1995, e.g. Theorem 21 in Chapter 5] and Jordan-Squire [2015].

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predictive fit assessed using log-predictive density (or any proper scoring rule)

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Joint work with Zheyang Shen (Newcastle), Jeremias Knoblauch (UCL), and Sam Power (Bristol)

• (for now) IID data 
$$y_{1:n} = (y_1, \dots, y_n) \in \mathcal{Y}^n$$

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### **Regularisation Target**

 $Q_0$  acts on  $Q_n$  in essentially the same way that  $Q_0$  acts on Gibbs measures like  $Q_n^{\dagger}$ , as a reference measure in a Radon–Nikodym derivative [Bissiri et al., 2016, Knoblauch et al., 2022]. That is, once can reason about 'updating belief distributions' using PCUQ.

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### Learning Rate

Compared to generalised Bayes, PCUQ depends less critically on the learning rate  $\lambda_n$ . i.e. support of  $Q_n$  is not a singleton set when  $\lambda_n \to 0$ .

# Model Misspecification in Cell Signalling



Figure: PCUQ predictive for the ERK signalling model.

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# Model Misspecification in Cell Signalling



Figure: PCUQ predictive for the ERK signalling model.

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# A Bit More Detail

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#### To define the *predictive fit* for PCUQ, we need:

▶ kernel  $k : \mathcal{Y} \times \mathcal{Y} \to \mathbb{R}$ , e.g.  $k(y, y') = (yy') + (yy')^2$ 

- reproducing kernel Hilbert space (RKHS) H(k) [see Berlinet and Thomas-Agnan, 2011, for background]
- kernel mean embedding

$$\mu_k(P) := \int k(\cdot, y) \, \mathrm{d}P(y) \in \mathcal{H}(k).$$

The divergence of a candidate  $P \in \mathcal{P}(\mathcal{Y})$  from the data-generating distribution  $P_*$  can be quantified using *maximum mean discrepancy* (MMD):

$$\mathsf{MMD}(P_{\star}, P) = \|\mu_{k}(P_{\star}) - \mu_{k}(P)\|_{\mathcal{H}(k)}$$
  
e.g.= 
$$\left\| \begin{bmatrix} \mathbb{E}_{Y \sim P_{\star}}[Y] \\ \mathbb{E}_{Y \sim P_{\star}}[Y^{2}] \end{bmatrix} - \begin{bmatrix} \mathbb{E}_{Y \sim P}[Y] \\ \mathbb{E}_{Y \sim P}[Y^{2}] \end{bmatrix} \right\|$$

The MMD is a proper metric if k is a *characteristic* kernel [Sriperumbudur et al., 2011]; our use of MMD is justified by its interpretation as a statistical divergence induced by a *proper scoring rule* [Dawid, 1986].

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e.g.= 
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Interpretation as Kernel Stein Discrepancy

This reveals one possible interpretation of (1) as a *kernel Stein discrepancy* [Chwialkowski et al., 2016, Liu et al., 2016, Gorham and Mackey, 2017] corresponding to the *Stein kernel*  $\kappa_{P_*}$  [Oates et al., 2017].

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#### Sensible in the Well-Specified Context

If  $P_{\star} = P_{\theta_{\star}}$  for some unique  $\theta_{\star} \in \Theta$ , then (1) is uniquely minimised by  $Q = \delta_{\theta_{\star}}$  provided k is a characteristic kernel.

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#### Estimation

Of course, the true data-generating distribution  $P_{\star}$  in (1) is unknown and must be approximated. In PCUQ we use the empirical distribution  $P_n$  in lieu of  $P_{\star}$ .

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#### Regularisation

A plug-in approximation necessitates additional regularisation, since otherwise minimisation of  $Q \mapsto MMD(P_n, P_Q)$  would result in a discrete distribution where each atom corresponds to a value of  $\theta$  that explains one of the data points well.

The output  $Q_n$  of PCUQ is a minimiser of the entropy-regularised objective

$$\mathcal{F}_n(Q) = \mathcal{E}_n(Q) + \lambda_n \int \log q(\theta) \, \mathrm{d}Q(\theta), \tag{2}$$

where the free energy  $\mathcal{E}_n(Q)$  is identical, after algebraic manipulation, to

$$\mathcal{E}_n(Q) \stackrel{+C}{=} \int v(\theta) \, \mathrm{d}Q(\theta) + \frac{1}{2} \iint \kappa_{P_n}(\theta, \vartheta) \, \mathrm{d}Q(\theta) \mathrm{d}Q(\vartheta),$$

#### and where q and $q_0$ are respectively densities for Q and $Q_0$ .

Wasserstein Gradient Flow

For the entropy-regularised functional  $\mathcal{F}_n$  (2), we can simulate a Wasserstein gradient flow via a McKean–Vlasov process [Ambrosio et al., 2008]

$$d\theta_t = -\nabla_W \mathcal{E}_n(Q^t)(\theta_t) + \sqrt{2\lambda_n} dW_t,$$
(3)  
$$\nabla_W \mathcal{E}_n(Q^t)(\theta_t) = \nabla v(\theta_t) + \int \nabla_1 \kappa_{P_n}(\theta_t, \vartheta) dQ^t(\vartheta)$$

where  $Q^t = \text{law}(\theta_t)$ ,  $\nabla_W$  denotes the Wasserstein gradient,  $(W_t)_{t\geq 0}$  is a Wiener process on  $\mathbb{R}^p$  and, for the bivariate function  $\kappa_{P_n}$ , the notation  $\nabla_1 \kappa_{P_n}$  denotes differentiation with respect to the first argument.

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#### Simulation as an Interacting Particle System

Discretise  $Q^t$  into a system of N evolving particles  $\theta_t^1, \theta_t^2, \dots, \theta_t^N$ , whose evolution is governed by the following system of *stochastic differential equations* (SDEs):

$$\mathrm{d}\theta_t^i = -\left(\nabla v(\theta_t^i) + \frac{1}{N-1}\sum_{j\neq i} \nabla_1 \kappa_{P_n}(\theta_t^i, \theta_t^j)\right) \mathrm{d}t + \sqrt{2\lambda_n} \mathrm{d}W_t^i$$

where  $(W_t^i)_{t\geq 0}$  are N independent Wiener processes on  $\mathbb{R}^p$ . An Euler–Maruyama discretisation incurs per-iteration computational complexity  $O(nN^2)$  and storage complexity (with caching) of O(n + N).



Figure: Interacting particle system for approximation of  $Q_n$ . (N = number of particles used)

Convergence of the Gradient Flow

Though theoretically convex, in practice gradients are small in low probability region; we mitigated this by initialising close to the Bayesian MAP  $\theta^{\dagger}$ .

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## Extension to Dependent Data

- each  $y_i$  is associated with a covariate  $x_i \in \mathcal{X}$  and generated according to an (unknown) conditional distribution  $P_*(\cdot|x_i)$
- ▶ have a conditional model  $\{P_{\theta}(\cdot|x)\}_{\theta\in\Theta}$  for each  $x \in \mathcal{X}$

Idea: Suppose that [Alquier and Gerber, 2024]

$$\{(x_i, y_i)\}_{i=1}^n \stackrel{\text{iid}}{\sim} \bar{P}_{\star}(\mathrm{d} x, \mathrm{d} y) := \frac{1}{n} \sum_{i=1}^n \delta_{x_i}(\mathrm{d} x) P_0(\mathrm{d} y | x_i)$$

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Choice of Kernel

For example, if  $\mathcal{X} \subset \mathbb{R}^{d_{\mathcal{X}}}$  and  $\mathcal{Y} \subset \mathbb{R}^{d_{\mathcal{Y}}}$ , we may consider the Gaussian kernel

$$k((x,y),(x',y')) = \exp\left(-\frac{\|x-x'\|^2}{\ell_{\mathcal{X}}^2} - \frac{\|y-y'\|^2}{\ell_{\mathcal{Y}}^2}\right)$$

with bandwidths  $\ell_X$  and  $\ell_Y$  to be specified. (We take  $\ell_X o 0$ , as recommended in Alquier and Gerber [2024].)

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For example, if  $\mathcal{X} \subset \mathbb{R}^{d_{\mathcal{X}}}$  and  $\mathcal{Y} \subset \mathbb{R}^{d_{\mathcal{Y}}}$ , we may consider the Gaussian kernel

$$k((x,y),(x',y')) = \exp\left(-\frac{\|x-x'\|^2}{\ell_{\mathcal{X}}^2} - \frac{\|y-y'\|^2}{\ell_{\mathcal{Y}}^2}\right)$$

with bandwidths  $\ell_{\mathcal{X}}$  and  $\ell_{\mathcal{Y}}$  to be specified. (We take  $\ell_{\mathcal{X}}\to 0$ , as recommended in Alquier and Gerber [2024].)

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Notable omissions:

Bayesian exponentially tilted empirical likelihood, conformal prediction, martingale posteriors (next chapter!), ...

If you would like to read more about our approach:

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