Black Box Probabilistic Numerics

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The Alan Turing Institute

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Motivation

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State of the universe:
$$u = (u(t))_{t \in T}, u \in U$$

- ▶ Information: $A : U \to \mathbb{R}^n$, some $n \in \{1, 2, ...\}$
- **Quantity of interest**: $Q : U \to \mathbb{R}^m$, some $m \in \{1, 2, ...\} \cup \{\infty\}$
- e.g. for numerical integration we might have

$$Q(u) = \int_0^1 u(t) dt, \qquad A(u) = [u(0), u(x), u(2x), \dots, u(1)].$$

Linear information enables us to use a conjugate Gaussian framework:

- 1. Select a Gaussian process $(U(t))_{t \in T}$ to represent epistemic uncertainty in $(u(t))_{t \in T}$.
- 2. Compute the conditional

$$U|(A = a) \sim \mathcal{GP}(m_{U|a}, k_{U|a})$$

$$m_{U|a}(t) = A_{t'}k(t, t')[A_tA_{t'}k(t, t')]^{-1}a$$

$$k_{U|a}(t, t') = k(t, t') - A_{t'}k(t, t')[A_tA_{t'}k(t, t')]^{-1}A_tk(t, t')$$

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Using the same notation, consider instead

$$Mu = b,$$
 $u = (u_1, \ldots, u_d)^\top \in \mathbb{R}^d.$

The matrix-vector products computed in the popular conjugate gradient method are

$$\langle s^{(1)}, b \rangle, \qquad s^{(1)} = b$$

 $\langle s^{(2)}, b \rangle, \qquad s^{(2)} = \text{cubic in } b$
 $\langle s^{(3)}, b \rangle, \qquad s^{(3)} = \text{ninth powers of } b$

So it seems natural to let

$$\Lambda(u) = \begin{bmatrix} \langle s^{(1)}, Mu \rangle \\ \langle s^{(2)}, Mu \rangle \\ \vdots \end{bmatrix}$$

... but this is **nonlinear** information!

This problem is not easily fixed.

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Aim: A *black box* that enables state-of-the-art numerical algorithms to be immediately exploited in the context of probabilistic numerics (PN).

Key Idea: Predict the limit of a sequence of increasingly accurate approximations produced by a traditional numerical method.

Bonus: A statistical perspective on *extrapolation methods*.

GPs: For concreteness, we will predict using GPs, but other predictive models could be used.

Compared to standard PN:

- applicable to nonlinear information
- state-of-the-art performance and functionality (in principle, at least)
- (\checkmark) provably higher order of convergence relative to a single application of the numerical method
- (X) multiple realisations of a numerical method are required
- (X) a joint statistical model has to be built for not just the quantity of interest but also for the error associated with the output of a traditional numerical method.

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An extrapolation method is an estimate for the limit

$$\lim_{x\to 0} f(x)$$

which can be based on values $\{f(\mathbf{x}_1), \ldots, f(\mathbf{x}_n)\}$ with $\{\mathbf{x}_1, \ldots, \mathbf{x}_n\} \subset (0, \infty)^d$ such that the associated computational cost falls within a notional budget.

Applications:

- finite difference approximation to derivatives
- numerical integration
- numerical solution of differential equations
- modern computer codes

Solutions:

- Richardson [1911] (higher-order convergence guaranteed)
- other extrapolation methods (NA-informed, typically univariate quantity of interest)
- multi-fidelity modelling (flexible, data-driven, multivariate quantity of interest)

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- numerical solution of differential equations
- modern computer codes

Solutions:

- Richardson [1911] (higher-order convergence guaranteed)
- other extrapolation methods (NA-informed, typically univariate quantity of interest)
- multi-fidelity modelling (flexible, data-driven, multivariate quantity of interest)

An extrapolation method is an estimate for the limit

$$\lim_{x\to 0} f(x)$$

which can be based on values $\{f(\mathbf{x}_1), \ldots, f(\mathbf{x}_n)\}$ with $\{\mathbf{x}_1, \ldots, \mathbf{x}_n\} \subset (0, \infty)^d$ such that the associated computational cost falls within a notional budget.

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Example: Higher-order convergence with Richardson extrapolation

Figure: Lewis Fry Richardson (1881 - 1953; born in Newcastle upon Tyne)

Suppose that



so that f(x) is a *first-order* approximation to f(0).

Then

$$2f(x) - f(2x)$$

= 2[f(0) + f'(0)x + O(x²)]
- [f(0) + f'(0)(2x) + O(x²)]
= f(0) + O(x²)

is a *second-order* approximation to f(0)

In general

 $f^{(s)}$ cts \implies combine *s* evaluations of *f* to get order-*s* approximation to f(0).

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Figure: Christiaan Huygens (1629 -1695; born in the Hague)

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- extrapolation methods do not provide probabilistic UQ
- multi-fidelity modelling is not NA-informed and lacks higher-order convergence guarantees
- neither method enables experimental design for $\{x_1, \ldots, x_n\}.$

Seek a extrapolation method that:

- provides probabilistic UQ
- is NA-informed
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Black Box Probabilistic Numerics

Given an explicit error bound b such that $f(\mathbf{x}) = f(\mathbf{0}) + O(b(\mathbf{x}))$ and $b(\mathbf{0}) = 0$.

How to encode this knowledge into a probabilistic regression model?

Gauss–Richardson Extrapolation (GRE)

Samples $g \sim \mathcal{GP}(0, k)$ from a Gaussian process with covariance kernel

$$k(\mathbf{x},\mathbf{x}') = \sigma^2 \left\{ k_0^2 + b(\mathbf{x})b(\mathbf{x}')k_e(\mathbf{x},\mathbf{x}') \right\}, \qquad \mathbf{x},\mathbf{x}' \in (0,\infty)^d,$$

a.s. satisfy $g(\mathbf{x}) - g(\mathbf{0}) = O(b(\mathbf{x}))$ as well.

So we use this GP regression model, trained on $[f(X_n)]_i = f(\mathbf{x}_i)$, to predict $f(\mathbf{0})$.

Only need to work with *linear* information!

Since k_0^2 is proportional to the prior variance for $f(\mathbf{0})$, we seek to let $k_0^2 \to \infty$, representing the flat prior limit / universal kriging:

$$f(\mathbf{0})|f(X_n) \sim \mathcal{N}(m_n[f], v_n[f]), \qquad m_n[f] = \frac{1^\top \mathsf{K}_b^{-1} f(X_n)}{1^\top \mathsf{K}_b^{-1} 1}, \qquad v_n[f] = \frac{\sigma_n^2[f]}{1^\top \mathsf{K}_b^{-1} 1},$$

where $[K_b]_{i,j} = k_b(\mathbf{x}_i, \mathbf{x}_j) = b(\mathbf{x}_i)b(\mathbf{x}_j)k_e(\mathbf{x}_i, \mathbf{x}_j)$ and $\sigma_n^2[f]$ is a scale estimator, to be specified.

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▶ Let $A \in \mathbb{R}^{m \times m}$.

- An eigenvalue $\lambda \in \mathbb{R}$ satisfies $Av = \lambda v$ for some $v \in \mathbb{R}^m$.
- All such matrices have *n* (possibly complex or repeated) eigenvalues, say $\{\lambda_1, \ldots, \lambda_m\}$.
- Eigenvalues are important in many applications, e.g.
 - stability analysis of dynamical systems,
 - web search engines,
 - principal component analysis, . . .
- **•** The basic idea of the **QR Algorithm** is to set $A_0 = A$ and iterate

$$A_{n-1} = Q_{n-1}R_{n-1}, \qquad A_n = R_{n-1}Q_{n-1}.$$

where Q_{n-1} is orthogonal and R_{n-1} is upper-triangular. The diagonal of A_k converges to the set of eigenvalues $\{\lambda_1, \ldots, \lambda_m\}$.

- To see why this works some calculation is needed.
- Also lots of details that are practically important.

Nonlinear information \implies difficult to envisage as a probabilistic numerical method.

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Example: Sparse matrices that arise as the discrete Laplace operator in the solution of the Poisson equation by a finite difference method with a five-point stencil

$$A = \begin{pmatrix} B & -I & & \\ -I & B & -I & \\ & \ddots & \ddots & -I \\ & & -I & B \end{pmatrix}, \qquad B = \begin{pmatrix} 4 & -1 & & \\ -1 & 4 & -1 & \\ & \ddots & \ddots & \\ & & \ddots & \ddots & -1 \\ & & & -1 & 4 \end{pmatrix},$$

where B is an $l \times l$ matrix and A is an $ml \times ml$ matrix, and we aim to recover the largest few eigenvalues of the matrices considered.

For GRE we took:

- \triangleright $x_n = 1/n$, where *n* is the number of iterations performed.
- entries of A_n are modelled as a priori independent; $f(x_n) = [A_n]_{i,i}$ (but this might be naïve).

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where B is an $l \times l$ matrix and A is an $ml \times ml$ matrix, and we aim to recover the largest few eigenvalues of the matrices considered.

For GRE we took:

- $x_n = 1/n$, where *n* is the number of iterations performed.
- entries of A_n are modelled as a priori independent; $f(x_n) = [A_n]_{i,i}$ (but this might be naïve).

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Figure: QR algorithm. All plots show red shaded $\pm 2\sigma$ credible intervals, numerical data as black circles, and true eigenvalues as blue stars. A total of n = 5 (left) and 15 (centre) iterations were used.

✓) No additional computational cost to GRE, since the sequence $(f(x_i))_{i=1}^n$ is generated during a single run of the iterative numerical method.

 (\checkmark) Overhead due to fitting the Gaussian process (GP) is negligible in this experiment.



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Importance of a Non-Stationary GP: Recall that *s* is *inferred* in these simulations - the estimated values were, respectively, 1.0186 and 1.0167.

Contrast with a *stationary* GP model (*i.e.* s = 0):



Figure: Comparison of non-stationary (left) and stationary (right) covariance kernels.

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3

Theory of Gauss-Richardson Extrapolation

Set-up for theoretical analysis of GRE

Recall $f(\mathbf{x}) = f(\mathbf{0}) + O(b(\mathbf{x}))$ for some $\mathbf{x} \in [0, 1]^d$. Design points $X_n = {\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_n} \subset [0, 1]^d$.



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Recall $f(\mathbf{x}) = f(\mathbf{0}) + O(b(\mathbf{x}))$ for some $\mathbf{x} \in [0, 1]^d$.

Error of the highest-fidelity experiment is $f(\mathbf{x}) - f(\mathbf{0}) = O(b(\mathbf{x}))$



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Recall $f(\mathbf{x}) = f(\mathbf{0}) + O(b(\mathbf{x}))$ for some $\mathbf{x} \in [0, 1]^d$.

Box fill distance ρ_{X_n} = size of the biggest cube not containing an element of X_n .



Recall $f(\mathbf{x}) = f(\mathbf{0}) + O(b(\mathbf{x}))$ for some $\mathbf{x} \in [0, 1]^d$.

Scaled design points $X_n^h = \{h\mathbf{x} : \mathbf{x} \in X_n\}$ where $h \in (0, 1]$.



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Theorem (Higher-order convergence with GRE) Assume that $b \in poly(r)$ and $k_e \in C^{2s}$.

Let $m_n^h[f]$ be the GRE estimator based on $f(X_n^h)$. Then there is an explicit constant C_s such that



whenever the box fill distance ρ_{X_n} is "small enough".

Remarks:

Proof via local polynomial reproduction, a la Wendland [2004].

Applies to s-smooth f, meaning that

$$|f|_{\mathcal{H}_{k}(\mathcal{X})} = \left\| \mathbf{x} \mapsto e(\mathbf{x}) := \frac{f(\mathbf{x}) - f(\mathbf{0})}{b(\mathbf{x})} \right\|_{\mathcal{H}_{k_{\mathbf{0}}}(\mathcal{X})} < \infty$$

- Extrapolation with convergence rate error $O(h^{s+r})$.
- Analogous result with exponential rates when f is smooth enough.

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$$\rho_{X_n} \leq \frac{1}{\gamma_d(r+2s)},$$

where $\gamma_d := 2d(1 + \gamma_{d-1})$ with base case $\gamma_1 := 2$.

Question: We lose the "one order per datum" of Richardson - is something lost in the proof?



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Illustration: Finite difference method

For $g : \mathbb{R} \to \mathbb{R}$ be s + 1 times continuously differentiable in an open neighbourhood of $t \in \mathbb{R}$, and consider the *central difference* method

$$f(x) = \frac{g(t+x) - g(t-x)}{2x}$$

for approximation of g'(t).

Central differences are second-order accurate, so we take $b(x) = x^2$.

$$X_n^h = \{0.2h, 0.4h, 0.6h, 0.8h, h\}.$$



Figure: Finite difference approximation; $g \in C^3$ (s = 2).

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Uncertainty quantification

For the maximum quasi likelihood estimator

$$\sigma_n^2[f] = \frac{1}{n} \left[f(X_n)^\top \mathbf{K}_b^{-1} f(X_n) - \frac{(\mathbf{1}^\top \mathbf{K}_b^{-1} f(X_n))^2}{\mathbf{1}^\top \mathbf{K}_b^{-1} \mathbf{1}} \right]$$

we can show that

$$\limsup_{h\to 0} \frac{|f(\mathbf{0}) - m_n^h[f]|}{\sqrt{v_n^h[f]}} < \infty$$

which is nice, but we seem to be a bit asymptotically over-confident when using the "right" kernel (s = 2).



Figure: Finite difference approximation; $g \in C^3$ (s = 2)

Optimal experimental design

The variance returned from GRE is

$$\mathsf{v}_n[\mathsf{f}] = \frac{\sigma_n^2[\mathsf{f}]}{\mathbf{1}^\top \mathsf{K}_b^{-1} \mathbf{1}}$$

An *a priori* optimal experimental design is

$$\underset{X}{\operatorname{arg\,max}} \quad \mathbf{1}^{\top}\mathbf{K}_{b}^{-1}\mathbf{1} \text{ s.t. } \sum_{\mathbf{x}\in X} c(\mathbf{x}) \leq C,$$

where $\mathbf{K}_b = [k_b(\mathbf{x}, \mathbf{x}')]_{\mathbf{x}, \mathbf{x}' \in X}$.

Let's vary the computational budget C and look at optimal designs:



Figure: Optimal experimental design; b(x) = x, $c(x) = x^{-1}$

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Case Study: Cardiac Modelling

Cardiac Model

Here $f(\mathbf{x})$ is a numerical simulation of a single heart beat with both a spatial (x_1) and a temporal (x_2) discretisation level [Strocchi et al., 2023].



Figure: Cardiac model: A subset of the mesh resolutions used in this case study. The finest resolution required 3×10^7 finite elements to be used.

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The computational cost $c(\mathbf{x})$ is measured in real computational time (seconds) and comprises

- setup time
- assembly time (the time taken to assemble linear systems of equations)
- solver time (the time taken to solve linear systems of equations)

with assembly time the main contributor to total computational cost.

To achieve a clinically-acceptable level of accuracy, it is typical for a simulation to be performed with $x_{default} \approx (0.4 \text{ mm}, 2 \text{ ms})$, at a cost $c(x_{default}) \approx 1.5 \times 10^4$ seconds (around ≈ 4 hours on 512 cores of ARCHER) for a single heart beat.

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Here is our workflow:

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Scalar quantities of interest

For assessment purposes we aim to predict $f(x_{hi-fi})$ as a ground truth, but in practice the goal is to predict f(0).

For each of 7 scalar quantities of interest associated with the cardiac model we display the relative error of GRE

$$\frac{|f(\mathbf{x}_{\mathsf{hi-fi}}) - m_n[f](\mathbf{x}_{\mathsf{hi-fi}})|}{|f(\mathbf{x}_{\mathsf{hi-fi}}) - f(\mathbf{x}_{\mathsf{default}})|}$$

with respect to the default approximation, as a function of the total computational budget C.



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Temporal quantities of interest

Extension of GRE to multi-output Gaussian process models allows temporal quantities of interest to be accurately extrapolated:



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- Black box probabilistic numerics leverages existing numerical methods in an extrapolation framework.
- Avoids the tricky issue of nonlinear information, by formulating a linear problem instead.
- Complementary to standard PN.

Full details in the report

Probabilistic Richardson Extrapolation, arXiv:2401.07562

where we discuss how Gauss-Richardson is capable also of

- estimation of uncertain convergence rates $b(\mathbf{x})$
- **i** simultaneous extrapolation and emulation for parametric models $f_{\theta}(\mathbf{x})$
- extrapolation of multivariate quantities $f(\mathbf{x}, \mathbf{t})$

Credit to: Onur Teymur, Chris Foley, Philip Breen, Toni Karvonen, Aretha Teckentrup, Marina Strocchi, Steve Niederer.

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