Deterministic Bayesian inference methods for the Naomi model

HIV Inference Lab Group Meeting

Adam Howes

Imperial College London

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Bayesian modelling and inference

- ullet As a statistical modeller, the bulk of our job is in constructing a generative model for data y using parameters ϑ
- This is the joint distribution $p(y, \vartheta) = p(y | \vartheta)p(\vartheta)$
- Then, we want to compute (ideally without having to think much about it) the posterior $p(\vartheta \mid y)$ which is just¹

$$p(\vartheta \mid y) = \frac{p(y,\vartheta)}{p(y)} = \frac{p(y \mid \vartheta)p(\vartheta)}{p(y)}$$

• The central problem of Bayesian inference is doing the following integral

$$p(y) = \int p(y, \vartheta) d\vartheta$$

 $^{^1\}mbox{l've}$ highlighted this with sarcasm in mind: it's a difficult problem

Numerical integration

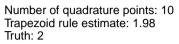
- If you want to integrate something deterministically, you could use numerical integration, otherwise called quadrature
- Select nodes $\theta \in \mathcal{Q} \subset \Theta$ and weights $\omega : \Theta \to \mathbb{R}$ then compute the sum

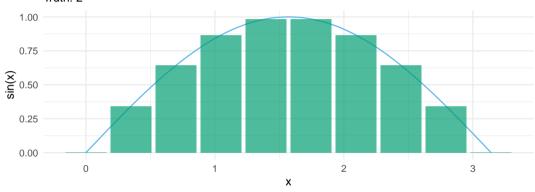
$$\tilde{p}(y) = \sum_{\vartheta \in \mathcal{Q}} p(y,\vartheta)\omega(\vartheta)$$

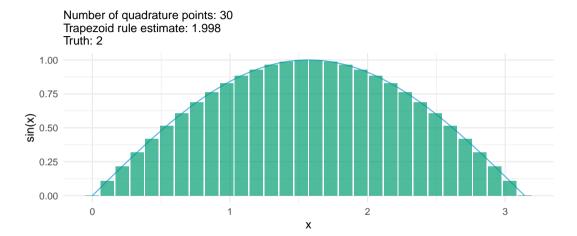
Naive quadrature example

- Remember how integration is taught? (Rienmann sums)
- Try computing $\int_0^{\pi} \sin(x) dx = 2$ using trapezoid rule

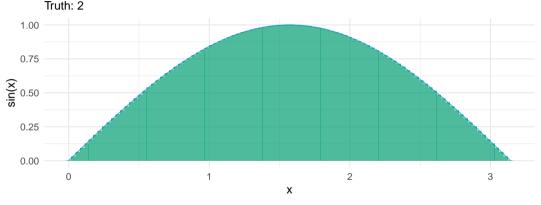
```
trapezoid_rule <- function(x, spacing) {
  w <- rep(spacing, length(x))
  w[1] <- w[1] / 2
  w[length(x)] <- w[length(x)] / 2
  sum(w * x)
}</pre>
```







Number of quadrature points: 100 Trapezoid rule estimate: 2 Truth: 2



Monte Carlo as an example of numerical integration

- Suppose we can sample $\vartheta_i \sim p(y, \vartheta)$ for $i = 1, \dots, N$
- If we set $\omega(\vartheta_i) = 1/N$ for all i then we get a **Monte Carlo** (MC) estimate

$$\tilde{p}(y) = \frac{1}{N} \sum_{i} p(y, \vartheta_i)$$

• For complicated models² it's not possible to sample directly from $p(y, \vartheta)$, but we can usually sample from a Markov chain which if you squint a bit is good enough (MCMC)

²Or not even that complicated

Monte Carlo is fundamentally unsound

- "Monte Carlo ignores information" according to O'Hagan (1987)
- Suppose N=3 and we sample $\vartheta_1,\vartheta_2,\vartheta_3$ with $\vartheta_2=\vartheta_3$ then our MC estimate is

$$\tilde{p}(y) = \frac{1}{3} \left(p(y, \vartheta_1) + p(y, \vartheta_2) + p(y, \vartheta_3) \right)$$

• This is despite the fact that nothing new about the function has been learned by adding $\{\vartheta_3, p(y, \vartheta_3)\}$

Application to HIV survey sampling

- This is a digression but...
- Say we're running a household survey, and sample the same individual twice
- We didn't learn anything new about HIV by surveying them again!
- This doesn't just bite for nodes or individuals which are exactly the same: an analogous argument can be made if they are close together and we expect their function evaluations to be similar

 \implies Bayesian quadrature, Bayesian survey design For some half-baked thoughts, see athowes.github.io/fourthgen/paper.pdf

Latent variables and hyperparameters

- Quadrature doesn't work very well when $\dim(\vartheta)$ gets even moderately sized If you're using k points per dimension it's $k^{\dim(\vartheta)}$
- Previously all of the parameters were under the symbol ϑ what if we split them up as being $\vartheta = (x, \theta)$
- The key part about this is that $\dim(x) = N$ is big and $\dim(\theta) = m$ is small

Names for x	Names for $ heta$
Latent variables, random effects, latent field	Hyperparameters, fixed effects

Spatio-temporal statistics

- There is nothing inherently special about spatio-temporal statistics, but we
 do often end up in similar situations because of the structures of the
 problems we tackle
- ullet We have observations indexed by space $s \in \mathcal{S}$ and time $t \in \mathcal{T}$
- Usually we associate parameters to spatio-temporal locations as well as observations
- This ends up with us having something like $\{x_{s,t}\}$

What's important about this?

- 1. There might be a lot of spatio-temporal locations, so $\dim(x)$ might be pretty big! If you have 100 districts and 10 years, that's already $100 \times 10 = 1000$ parameters
- 2. Perhaps we're willing to make quite strong assumptions about how things vary over space-time³
- Not IID anymore!

³Are there any slides about spatial statistics that don't describe Tobler's first law of geography?

Latent Gaussian models

 A latent Gaussian model (LGM) (Rue, Martino, and Chopin 2009) looks along these lines:

$$\begin{array}{ll} \text{(Observations)} & y \sim p(y \,|\, x, \theta), \\ \text{(Latent field)} & x \sim \mathcal{N}(x \,|\, \mu(\theta), \, Q(\theta)^{-1}), \\ \text{(Hyperparameters)} & \theta \sim p(\theta). \end{array}$$

- Many models are LGMs, especially in spatio-temporal statistics
- $\dim(x) = n$, $\dim(x) = N$, $\dim(\theta) = m$

• Remember that we wanted to compute

$$p(y) = \int p(y, \vartheta) d\vartheta$$

- One trick for doing this is to pretend $p(\vartheta \mid y)$ is Gaussian

 - Mode $\hat{\vartheta} = \arg\max_{\vartheta} \log p(y, \vartheta)$ Hessian $H(\hat{\vartheta}) = -\partial_{\vartheta}^2 \log p(y, \vartheta)|_{\vartheta = \hat{\vartheta}}$
 - Gaussian approximation $\implies \tilde{p}_{\mathbf{G}}(\vartheta \mid y) = \mathcal{N}(\vartheta \mid \hat{\vartheta}, H(\hat{\vartheta})^{-1})$

Example of computing the Laplace approximation

• Consider the following model for i = 1, ..., n with fixed a and b

$$y_i \sim \mathsf{Poisson}(\lambda), \quad \lambda \sim \mathsf{Gamma}(a, b).$$

• It's conjugate so we directly know that

$$\lambda \mid y \sim \mathsf{Gamma}(a + \sum_i y_i, b + n)$$

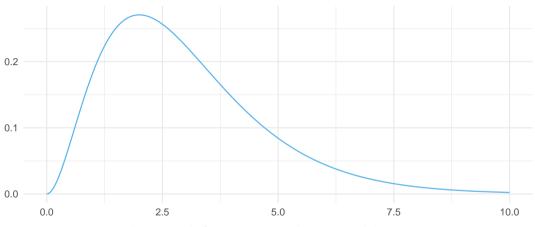


Figure 1: A Gamma prior with $\it a=3$ and $\it b=1$.

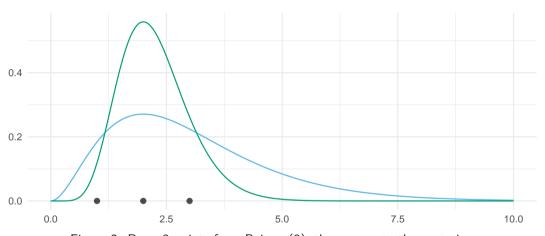


Figure 2: Draw 3 points from Poisson(3), then compute the posterior.

```
fn \leftarrow function(x) dgamma(x, a + sum(y), b + length(y), log = TRUE)
# Here we are using numerical derivatives rather than automatic
ff <- list(
 fn = fn
 gr = function(x) numDeriv::grad(fn, x),
 he = function(x) numDeriv::hessian(fn, x)
opt_bfgs <- aghq::optimize_theta(
 ff, 1, control = aghq::default_control(method = "BFGS")
```

```
laplace <- posterior +
 stat function(
  data = data.frame(x = c(0, 10)),
   aes(x).
   fun = dnorm.
   n = 500.
   args = list(mean = opt_bfgs$mode, sd = sqrt(1 / opt_bfgs$hessian)),
  col = cbpalette[3]
```

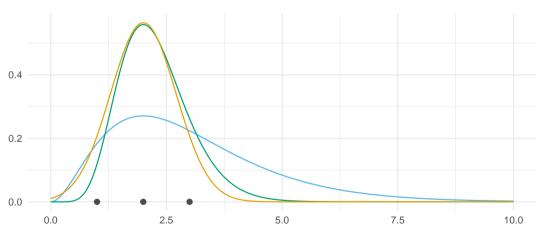


Figure 3: The Laplace approximation matches the true posterior near the mode but it's not great in the tails.

Now

$$p(y) = \frac{p(\vartheta, y)}{p(\vartheta \mid y)} \approx \frac{p(\vartheta, y)}{p_{G}(\vartheta \mid y)}$$

and we can evaluate RHS where we would like, so let's pick the point at which the Gaussian is most accurate, which is $\hat{\vartheta}$

$$p_{\mathrm{LA}}(y) = \frac{p(\vartheta, y)}{p_{\mathrm{G}}(\vartheta \mid y)}|_{\vartheta = \hat{\vartheta}} = (2\pi)^{\dim(\vartheta)/2} \det(H(\hat{\vartheta}))^{-1/2} p(\hat{\vartheta}, y)$$

Marginal Laplace approximation

- Hey wait a second, is it reasonable to just assume $p(\vartheta \mid y)$ is Gaussian?
- No, not in general. But...
- 1. We just described a class of models (LGMs) where some subset of the parameters (x) have a Gaussian prior \implies it's a lot more reasonable to think that they would have a marginal posterior which is close to Gaussian
- 2. We just talked about how big x is in comparison to $\theta! \implies$ most of the work in our integral can be done using a **marginal Laplace** approximation to get rid of x

Generalist versus specialist methods

Dichotomy in statistical inference methods:

- 1. Generalist: works in all situations
- 2. Specialist: "exploits" properties of the problem at hand

We are taking approach 2!

Marginal Laplace approximation

• What does this look like? Instead of assuming $p(\vartheta \mid y) = p(x, \theta \mid y)$ is Gaussian we assume $p(x \mid \theta, y)$ is

$$\tilde{p}_{\mathrm{G}}(x \mid \theta, y) = \mathcal{N}(x \mid \hat{x}, H(\hat{x}))^{-1}$$

where $\hat{x} = \hat{x}(\theta)$

Now the marginal Laplace approximation is

$$p_{\mathrm{LA}}(\theta, y) = \frac{p(x, \theta, y)}{\tilde{p}_{\mathrm{G}}(x \mid \theta, y)}|_{x = \hat{x}} = (2\pi)^{N/2} \det(H(\hat{x}))^{-1/2} p(\hat{x}, \theta, y)$$

Integrated nested Laplace approximation

- Now we can compute $p_{LA}(\theta, y)$ but what we really want is still p(y)
- ullet But hopefully 4 the dimension of heta is small enough that we can now tackle this with quadrature
- ullet So pick some nodes ${\mathcal Q}$ and a weighting function ω and away we go

$$p(y) \approx \sum_{\theta \in \mathcal{O}} p_{\text{LA}}(\theta, y) \omega(\theta)$$

This is the famous integrated nested Laplace approximation (INLA)

⁴Really: hopefully

Taking stock

- 1. Bayesian inference is integration
- 2. Spatial statistics has parameters (x, θ)
- 3. Integrate x cheaply using a Gaussian assumption
- 4. Try a bit harder with θ performing quadrature

Now let's apply this to a difficult problem in HIV inference!

The Naomi model

- Naomi is a spatial evidence synthesis model
- Used by countries to produce HIV estimates in a yearly process supported by UNAIDS
- Inference for Naomi is currently conducted using TMB by optimising $p_{LA}(\theta, y)$, and has to be pretty quick to allow for interactive review and development of estimates



Figure 4: A supermodel

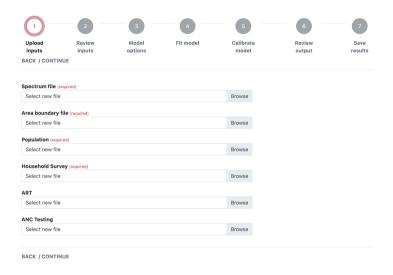


Figure 5: Example of the user interface from https://naomi.unaids.org/

Template Model Builder refresher

- TMB (Kristensen et al. 2015) is an R package which implements the Laplace approximation for latent variable models
- To use TMB, you write your objective function
 - $-\log p(\mathbf{y}\,|\,\mathbf{x},oldsymbol{ heta})p(\mathbf{x}\,|\,oldsymbol{ heta})p(oldsymbol{ heta})$ in TMB's C++ syntax
- For example, for the model $\mathbf{y} \sim \mathcal{N}(\mu, 1)$ with $p(\mu) \propto 1$ then the TMB user template looks as follows

```
#include <TMB.hpp>
template < class Type>
Type objective_function<Type>::operator()() {
 // Define data e.g.
 DATA_VECTOR(y);
 // Define parameters e.g.
 PARAMETER(mu);
 // Calculate negative log-likelihood e.g.
 nll = Type(0.0);
 nll -= dnorm(y, mu, 1, true).sum()
 return(nll);
```

Why do we use TMB

- It runs quickly, and is flexible enough to write the model
- Another answer: we don't have any better options
 - Markov chain Monte Carlo (MCMC) is accurate (eventually) but takes too long
 - Though Naomi is a spatial model with a large Gaussian latent field, it isn't technically a latent Gaussian model, and isn't compatible with the R-INLA implementation of INLA

Idea

- Implement an algorithm similar to INLA which
- 1. is compatible with Naomi
- 2. uses TMB to perform Laplace approximation
- Does this improves the quality of the inferences we get?

Not a new idea!

My main comment is that several aspects of the computational machineery that is presented by Rue and his colleagues could benefit from the use of a numerical technique known as automatic differentiation (AD) ... By the use of AD one could obtain a system that is automatic from a user's perspective... the benefit would be a fast, flexible and easy-to-use system for doing Bayesian analysis in models with Gaussian latent variables

– Hans J. Skaug (coauthor of ${\rm TMB}$), RSS discussion of Rue, Martino, and Chopin (2009)

Meanwhile in Canada...

- ullet Alex Stringer in Toronto ullet Waterloo had been thinking along similar lines, and made a lot of progress
- 1. Implementing an algorithm similar to INLA, using a specific quadrature rule $\mathcal Q$ called adaptive Gauss-Hermite quadrature (AGHQ) which he and others argue should be the default for this problem
- 2. Defining a class of models called extended latent Gaussian models (ELGMs) which Naomi fits into
- I will explain what both of these acronyms (AGHQ⁵ and ELGM) mean!

 $^{^5\}mbox{\sc Amusingly similar}$ to AGYW: adolescent girls and young women

Gauss-Hermite quadrature

Recall

$$\int f(z)\mathrm{d}z \approx \sum_{z\in\mathcal{Q}} f(z)\omega(z)$$

- Replace f(z) by $\phi(z)f(z)$ and say that f is a polynomial and ϕ is unknown
- Suppose that $\phi(z) = \exp(-z^2)$

Adaptation

- The nodes and weights we use should probably depend on the integrand
 Especially when the integrand is also a function of y as in p(y, θ)
- Let $z \in \mathcal{Q}(m,k)$ then $\theta(z) = \hat{\theta} + Lz$ where L is the lower Cholesky of $H(\hat{\theta})$

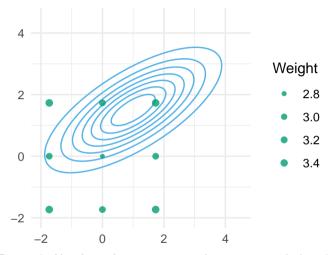
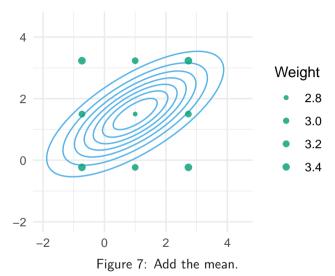


Figure 6: Unadapted points in two dimensions with k = 3.



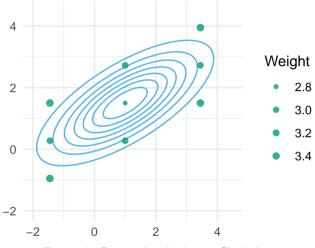


Figure 8: Rotate by the lower Cholesky.

Extended latent Gaussian models

- To explain what an extended latent Gaussian model (ELGM) is, we need a little more detail on LGMs
- In an LGM the conditional mean depends on exactly one structured additive predictor, specifically

$$y_i \sim p(y_i | \eta_i, \theta_1), \quad i \in [n]$$

 $\mu_i = \mathbb{E}(y_i | \eta_i) = g(\eta_i),$
 $\eta_i = \beta_0 + \sum_{l=1}^p \beta_j z_{ji} + \sum_{k=1}^r f_k(u_{ki}),$

Extended latent Gaussian models

• ELGM remove this requirement such that

$$\mu_{\it i} = {\it g}(\eta_{\mathcal{J}_{\it i}})$$

where $g_i: \mathbb{R}^{|\mathcal{J}_i|} \to \mathbb{R}$ and \mathcal{J}_i is some set of indices

• This allow for a higher degree of non-linearity in the model

Why is Naomi an ELGM?

	Things I thought made Naomi an ELGM	Does it actually?
1	ANC offset from household survey	?
2	Incidence depends on adult prevalence and coverage	?
3	ART attendance is a product	?
4	ART attendance uses a multinomial	?
5	Aggregation of finer processes	?

- I will explain each of these, and whether or not they make Naomi an ELGM, in more detail in slides to follow
 - The notation may not have been introduced properly, but hopefully the gist will still
 make sense

ANC offset from household survey

$$\begin{split} & \mathsf{logit}(\rho_{\mathsf{x},\mathsf{a}}^{\mathsf{ANC}}) = \mathsf{logit}(\rho_{\mathsf{x},\mathsf{F},\mathsf{a}}) + \beta^{\rho^{\mathsf{ANC}}} + u_{\mathsf{x}}^{\rho^{\mathsf{ANC}}} + \eta_{R_{\mathsf{x}},\mathsf{a}}^{\rho^{\mathsf{ANC}}}, \\ & \mathsf{logit}(\alpha_{\mathsf{x},\mathsf{a}}^{\mathsf{ANC}}) = \mathsf{logit}(\alpha_{\mathsf{x},\mathsf{F},\mathsf{a}}) + \beta^{\alpha^{\mathsf{ANC}}} + u_{\mathsf{x}}^{\alpha^{\mathsf{ANC}}} + \eta_{R_{\mathsf{x}},\mathsf{a}}^{\alpha^{\mathsf{ANC}}}. \end{split}$$

- Here $logit(\rho_{x,F,a})$ and $logit(\alpha_{x,F,a})$ are Gaussian
- They have their own structured additive predictors, but by use of the copy feature this should be possible in R-INLA
- Conclusion: does not make Naomi an ELGM

Incidence depends on adult prevalence and coverage

$$\log(\lambda_{x,s,a}) = \beta_0^{\lambda} + \beta_S^{\lambda,s=\mathsf{M}} + \log(\rho_x^{15\text{-}49}) + \log(1 - \omega \cdot \alpha_x^{15\text{-}49}) + u_x^{\lambda} + \eta_{R_x,s,a}^{\lambda}.$$

- Here $\log(\rho_x^{15-49})$ and $\log(1-\omega\cdot\alpha_x^{15-49})$ are not going to be Gaussian no matter your copy tricks
- Conclusion: does make Naomi an ELGM

ART attendance is a product

ART attendance uses the multinomial

Aggregation of finer processes

The algorithm

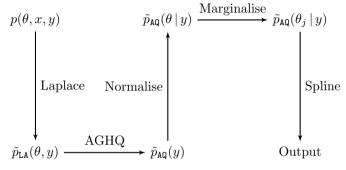


Figure 9: Inference for the hyperparameters. Shaped like a snake for no real reason.

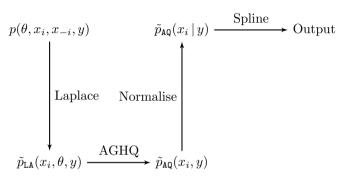


Figure 10: Inference for the latent field.

Thanks for listening!

- Working on a paper "Fast approximate Bayesian inference for small-area estimation of HIV indicators using the Naomi model" based on this work, joint with Alex Stringer (Waterloo) and my PhD supervisors Seth Flaxman (Oxford) and Jeff Eaton (Imperial)
- Let me know if you'd be up for being an early reader!
- Code for this project is at athowes.github.io/elgm-inf

References I

- Kristensen, Kasper, Anders Nielsen, Casper W Berg, Hans Skaug, and Brad Bell. 2015. "TMB: automatic differentiation and Laplace approximation." arXiv Preprint arXiv:1509.00660.
- O'Hagan, Anthony. 1987. "Monte Carlo Is Fundamentally Unsound." *The Statistician*, 247–49.
- Rue, Håvard, Sara Martino, and Nicolas Chopin. 2009. "Approximate Bayesian inference for latent Gaussian models by using integrated nested Laplace approximations." *Journal of the Royal Statistical Society: Series b* (Statistical Methodology) 71 (2): 319–92.