

Deterministic Bayesian inference methods for the Naomi model

HIV Inference Lab Group Meeting

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April 2023

Bayes

- 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 the posterior $p(\vartheta | y)$ which is **just**¹

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

- The central problem of Bayesian inference is that we can't compute $p(y)$

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

¹I've bolded this with sarcasm in mind

How might you do it?

- If you want to integrate something deterministically, you could use numerical integration methods
- Pick nodes $\vartheta \in \mathcal{Q} \subset \Theta$ and weights $\omega : \Theta \rightarrow \mathbb{R}$ then compute the sum

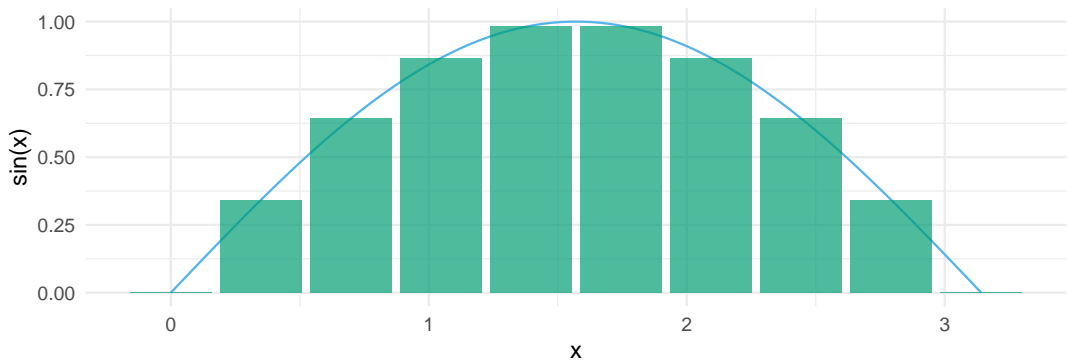
$$\tilde{p}(y) = \sum_{\vartheta \in \mathcal{Q}} p(y, \vartheta) \omega(\vartheta)$$

Naive quadrature example

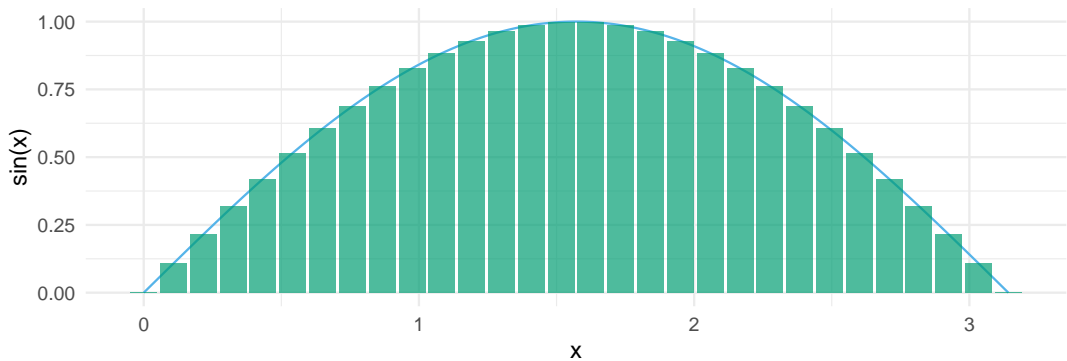
- Remember how integration is introduced? (Riemann sums)
- Try computing $\int_0^\pi \sin(x)dx = 2$ using trapezoid rule

```
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)  
}
```

Number of quadrature points: 10
Trapezoid rule estimate: 1.98
Truth: 2



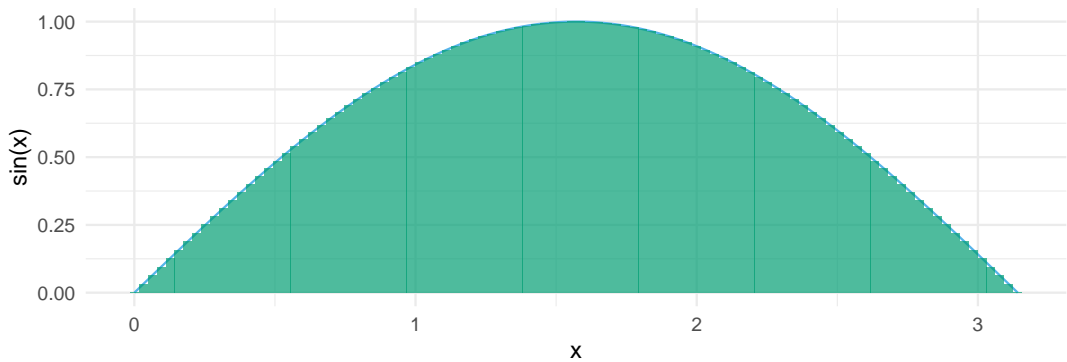
Number of quadrature points: 30
Trapezoid rule estimate: 1.998
Truth: 2



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

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

- 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

⇒ Bayesian quadrature, Bayesian survey design

For some half-baked thoughts, see athowes.github.io/fourth-gen/paper.pdf

Latent variables and hyperparameters

- Quadrature doesn't work very well when $\dim(\vartheta)$ gets even moderately sized
- Previously I had all of the parameters under the symbol ϑ
- What if we split them up as being $\vartheta = (x, \theta)$
- The key part about this is that $\dim(x)$ is big and $\dim(\theta)$ is small

Names for x	Names for θ
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
- 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 assumptions about how things vary over space-time³

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:

(Observations)	$y \sim p(y \mid x, \theta),$
(Latent field)	$x \sim \mathcal{N}(x \mid \mu(\theta), Q(\theta)^{-1}),$
(Hyperparameters)	$\theta \sim p(\theta).$

Laplace approximation

- Remember that we wanted to compute

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

- One trick for doing this is to pretend $p(\vartheta | y)$ 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 $\implies \tilde{p}_G(\vartheta | y) = \mathcal{N}(\vartheta | \hat{\vartheta}, H(\hat{\vartheta})^{-1})$

Example

- Consider the following model for $i = 1, \dots, n$ with fixed a and b

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

- It's conjugate so we directly know that

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

Laplace approximation

- Now

$$p(y) = \frac{p(\vartheta, y)}{p(\vartheta | y)} \approx \frac{p(\vartheta, y)}{p_G(\vartheta | y)}$$

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

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

Marginal Laplace approximation

- Hey wait a second, is it reasonable to just assume $p(\vartheta | y)$ is Gaussian?
 - No, not in general. But...
1. We just described a class of models (LGMs) where some subset of the parameters (the latent field 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 θ ! \implies most of the work in our integral can be done using a **marginal Laplace** approximation to get rid of x

Dichotomy in statistical inference methods:

1. Ones which aim to be completely general
2. Ones which aim to "exploit" properties of the problem at hand

We are taking approach 2.

Marginal Laplace approximation

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

$$\tilde{p}_G(x | \theta, y) = \mathcal{N}(x | \hat{x}, H(\hat{x}))^{-1}$$

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

- Now the marginal Laplace approximation is

$$p_{\text{LA}}(\theta, y) = \frac{p(x, \theta, y)}{\tilde{p}_G(x | \theta, y)} \Big|_{x=\hat{x}} = (2\pi)^{\dim(x)/2} \det(H(\hat{x}))^{-1/2} p(\hat{x}, \theta, y)$$

Integrated nested Laplace approximation

- Now we can compute $p_{\text{LA}}(\theta, y)$ but what we really want is still $p(y)$
- But hopefully⁴ the dimension of θ is small enough that we can now tackle this with quadrature
- So pick some nodes \mathcal{Q} and a weighting function ω and away we go

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

- This is the famous integrated nested Laplace approximation (INLA)

⁴Really: hopefully

Naomi evidence synthesis model

- Many people here have worked on Naomi 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 $p_{\text{LA}}(\theta, y)$
- Has to be pretty quick to allow for interactive review and development of estimates



Figure 1: A supermodel

1

2

3

4

5

6

7

Upload inputs

Review inputs

Model options

Fit model

Calibrate model

Review output

Save results

BACK / CONTINUE

Spectrum file (required)

Select new file

Browse

Area boundary file (required)

Select new file

Browse

Population (required)

Select new file

Browse

Household Survey (required)

Select new file

Browse

ART

Select new file

Browse

ANC Testing

Select new file

Browse

BACK / CONTINUE

Figure 2: Example of the user interface from <https://naomi.unaids.org/>

Why do we use TMB

- One answer: lack of other viable options
- Markov chain Monte Carlo (MCMC) 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 machinery 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 TMB), RSS discussion of Rue, Martino, and Chopin (2009)

Meanwhile in Canada...

- Alex Stringer in Toronto → 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!

⁵Amusingly similar to AGYW: adolescent girls and young women

What's AGHQ?

What's an ELGM?

Why is Naomi an ELGM?

ANC offset from household survey

$$\begin{aligned}\text{logit}(\rho_{x,a}^{\text{ANC}}) &= \text{logit}(\rho_{x,F,a}) + \beta^{\rho^{\text{ANC}}} + u_x^{\rho^{\text{ANC}}} + \eta_{R_x,a}^{\rho^{\text{ANC}}}, \\ \text{logit}(\alpha_{x,a}^{\text{ANC}}) &= \text{logit}(\alpha_{x,F,a}) + \beta^{\alpha^{\text{ANC}}} + u_x^{\alpha^{\text{ANC}}} + \eta_{R_x,a}^{\alpha^{\text{ANC}}}.\end{aligned}$$

Incidence depends on adult prevalence and coverage

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

ART attendance probability product

$$\dot{A}_{x',x,s,a} \sim \text{Bin}(N_{x',s,a}, \pi_{x',x,s,a})$$

where $\pi_{x',x,s,a} = \rho_{x',s,a} \cdot \alpha_{x',s,a} \cdot \gamma_{x',x,s,a}$.

ART attendance multinomial

Aggregation

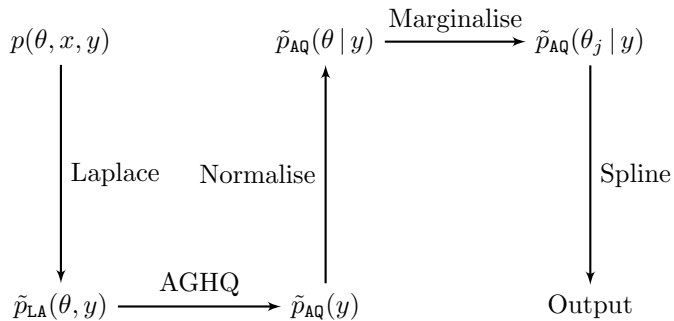


Figure 3: Inference for the hyperparameters. Shaped like a snake for no real reason.

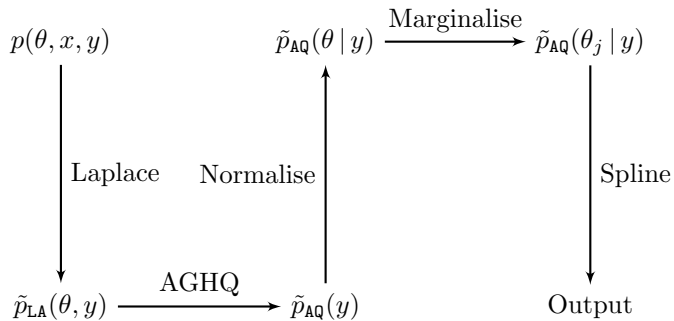


Figure 4: Inference for the latent field.

Latent field algorithm

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

- 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.