

# Fast approximate Bayesian inference for small-area estimation of HIV indicators using the Naomi model

Adam Howes<sup>1, 2</sup>, Alex Stringer<sup>3</sup>, Seth R. Flaxman<sup>4</sup>, Jeffrey W. Eaton<sup>2</sup>

Imperial College  
London

<sup>1</sup> Department of Mathematics, Imperial College London

<sup>2</sup> MRC Centre for Global Infectious Disease Analysis, School of Public Health, Imperial College London

<sup>3</sup> Department of Statistics and Actuarial Science, University of Waterloo

<sup>4</sup> Department of Computer Science, University of Oxford



## Summary

- Approximate Bayesian inference method combining Laplace approximations and adaptive Gauss-Hermite quadrature
- Motivated by a challenging, policy-relevant problem in small-area estimation of HIV indicators in sub-Saharan Africa
- Implemented as a part of the `aghq` R package, allowing use for any model with a `TMB` C++ user template

## The Naomi HIV model

- District-level model of HIV indicators which synthesises data from household surveys, antenatal care (ANC) clinics, and routine service provision of antiretroviral therapy (ART) (Eaton et al. 2021)
- Yearly estimation process: model run interactively by country teams using the web-app `naomi.unaids.org`
- Inference currently conducted with empirical Bayes and a Gaussian approximation to the latent field using Template Model Builder `TMB` (Kristensen et al. 2016)
- Days to get accurate answers with Markov chain Monte Carlo via `tmbstan` (Monnahan and Kristensen 2018)
- Motivates looking for a fast, approximate approach

## Extended latent Gaussian models

- In a latent Gaussian model (LGM) (Rue, Martino, and Chopin 2009) the conditional mean depends on exactly one structured additive predictor  $\mu_i = g(\eta_i)$  with  $g : \mathbb{R} \rightarrow \mathbb{R}$ 
  - The `R-INLA` implementation of integrated nested Laplace approximations applies only to LGMs
- Extended latent Gaussian models (ELGM) remove this requirement such that  $\mu_i = g(\eta_{\mathcal{J}_i})$  where  $g_i : \mathbb{R}^{|\mathcal{J}_i|} \rightarrow \mathbb{R}$
- Naomi is an ELGM, and not an LGM, because in our model:
  - ANC indicators are offset from household survey indicators
  - Incidence depends on prevalence and ART coverage
  - Observed data may be aggregated
  - Individuals may attend ART clinics outside their home district

## Inference procedure

### Background

- Laplace approximation** Integrate out variables using a Gaussian approximation to the denominator e.g.

$$p(\theta, y) \approx \tilde{p}_{\text{LA}}(\theta, y) = \frac{p(y, x, \theta)}{\tilde{p}_{\text{G}}(x \mid \theta, y)} \Big|_{x=\hat{x}(\theta)}$$

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

- Adaptive Gauss-Hermite Quadrature** Approximate integrals by  $\int_{\Theta} p(\theta) d\theta \approx |L| \sum_{z \in \mathcal{Q}(m, k)} p(\hat{\theta} + Lz) \omega(z)$  with Gauss-Hermite quadrature rule  $z \in \mathcal{Q}(m, k)$  adapted based upon the mode  $\hat{\theta} = \operatorname{argmax}_{\theta \in \Theta} p(\theta)$  and lower Cholesky  $LL^{\top} = -\partial_{\theta}^2 \log p(\theta) \Big|_{\theta=\hat{\theta}}$  of the target

### Our algorithm

- Given a C++ user template for  $-\log p(y, x, \theta)$  our algorithm is summarized by Figure 1

$$p(\theta, x, y) \xrightarrow{\text{Laplace}} \tilde{p}_{\text{LA}}(\theta, y)$$

$$\tilde{p}_{\text{LA}}(x_i, \theta, y) \qquad \tilde{p}_{\text{AQ}}(y)$$

Figure 1: Flowchart describing the algorithm

## Inference comparison

- Compare posterior inferences from `TMB`, `aghq`, `adam` and `tmbstan` using Kolmogorov-Smirnov tests on posterior marginals of hyperparameters, latent field, and outputs
- Figure 2 compares the distances between `TMB` and `adam` to `tmbstan`

## Future work

- More comprehensive inference comparison e.g. maximum mean discrepancy, Pareto-smoothed importance sampling
- Implement Laplace matrix algebra approximations e.g. Wood (2020)

**Interested to read more?** Working notebooks and R code available are available from `github.com/athowes/multi-agyw`.

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## References

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