

Fast approximate Bayesian inference of HIV indicators using PCA adaptive Gauss-Hermite quadrature

Adam Howes^{1, 2}, Alex Stringer³, Seth R. Flaxman⁴, Jeffrey W. Eaton^{5, 2}



¹ Department of Mathematics, Imperial College London

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

³ Department of Statistics and Actuarial Science, University of Waterloo

⁴ Department of Computer Science, University of Oxford

⁵ Center for Communicable Disease Dynamics, Department of Epidemiology, Harvard T.H. Chan School of Public Health

Summary

- We developed an approximate Bayesian inference method using Laplace approximation, adaptive Gauss-Hermite quadrature and principal component analysis
- Motivation: an evidence synthesis model for small-area estimation of HIV indicators in >35 countries in sub-Saharan Africa
- Implemented using the `aghq` package (Stringer 2021), and compatible with any model with a Template Model Builder `TMB` (Kristensen et al. 2016) C++ user template

1. The Naomi HIV model

- District-level model of HIV indicators which synthesises data from 1) household surveys, 2) antenatal care (ANC) clinics, and 3) routine service provision of antiretroviral therapy (ART) (Eaton et al. 2021)
 - Combining evidence from multiple data sources helps overcome the limitations of any one
 - Small-area estimation methods (structured random effects) to overcome limited district-level sample sizes
- Yearly estimation process: model run interactively by country teams using `naomi.unaids.org` web-app
 - Figure 1 illustrates the seven stages of using the app
- Inference conducted in minutes using empirical Bayes and a Gaussian approximation
- Days to get accurate answers with MCMC via `tmbstan` (Monnahan and Kristensen 2018): not practical!
- Naomi has a large latent field x : controlled by a smaller number of hyperparameters θ
- Extended latent Gaussian model (Stringer, Brown, and Stafford 2022): more complex dependency structures than a latent Gaussian model
- Looking for a fast, approximate approach, that properly takes uncertainty in hyperparameters into account

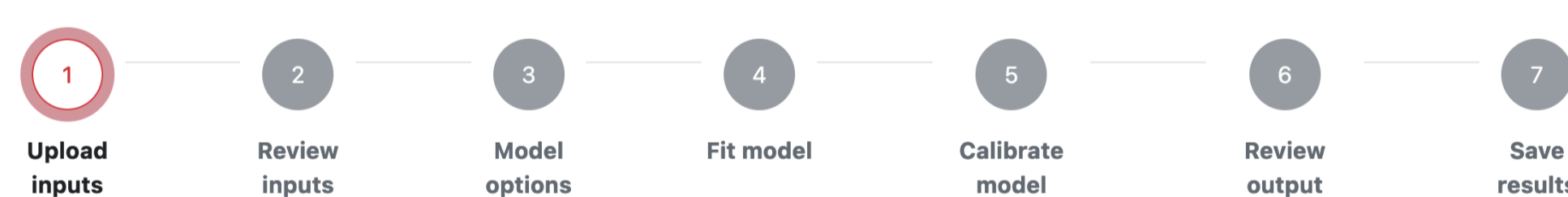


Figure 1: Model fitting occurs interactively in stages

2. Inference procedure

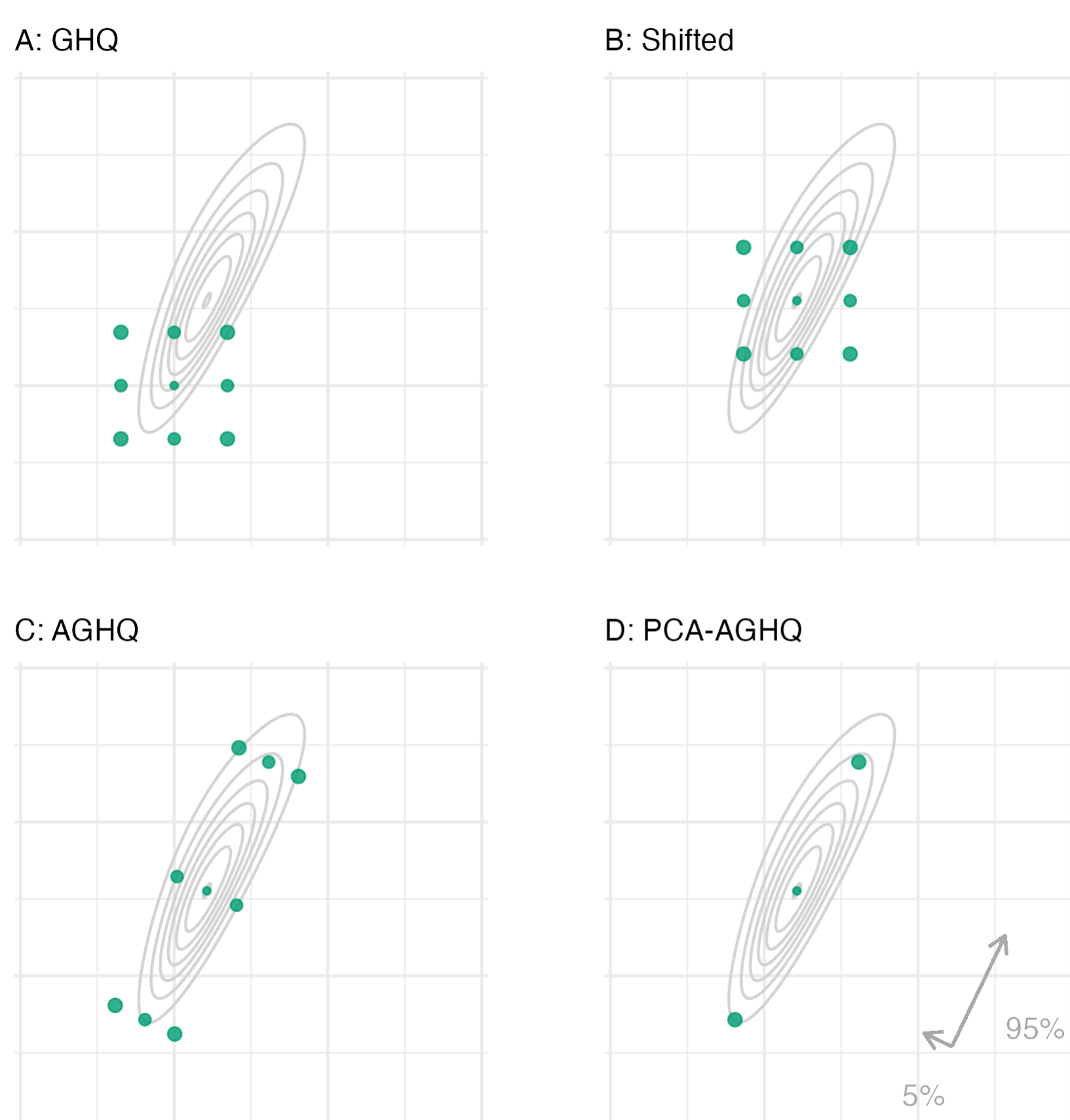


Figure 2: Demonstration of Gauss-Hermite quadrature, adaption, and our principal components approach.

- Laplace approximation** Integrate out latent field using a Gaussian approximation to the denominator

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

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

- Use automatic differentiation via `CppAD` in `TMB`

- Adaptive Gauss-Hermite Quadrature (AGHQ)** perform quadrature over the hyperparameters

$$\int_{\theta} p_{LA}(\theta, y) d\theta \approx |L| \sum_{z \in \mathcal{Q}(m, k)} p_{LA}(\hat{\theta} + Lz, y) \omega(z),$$

where $m = \dim(\theta)$ and the Gauss-Hermite quadrature rule $z \in \mathcal{Q}(m, k)$ with weights $\omega : \mathcal{Q} \mapsto \mathbb{R}$ and k points per dimension is adapted (Figure 2) based upon

- The mode $\hat{\theta} = \operatorname{argmax}_{\theta \in \Theta} p_{LA}(\theta, y)$
- A matrix decomposition $LL^T = -\partial_{\theta}^2 \log p_{LA}(\theta, y) |_{\theta=\hat{\theta}}$

- PCA-AGHQ** To integrate over large spaces, use the spectral decomposition $L = EA^{1/2}$ and keep only the first $s < m$ principal components

3. Application to Malawi

- Malawi is a relatively small country but still has latent field $\dim(x) = 467$ and hyperparameters $\dim(\theta) = 24$

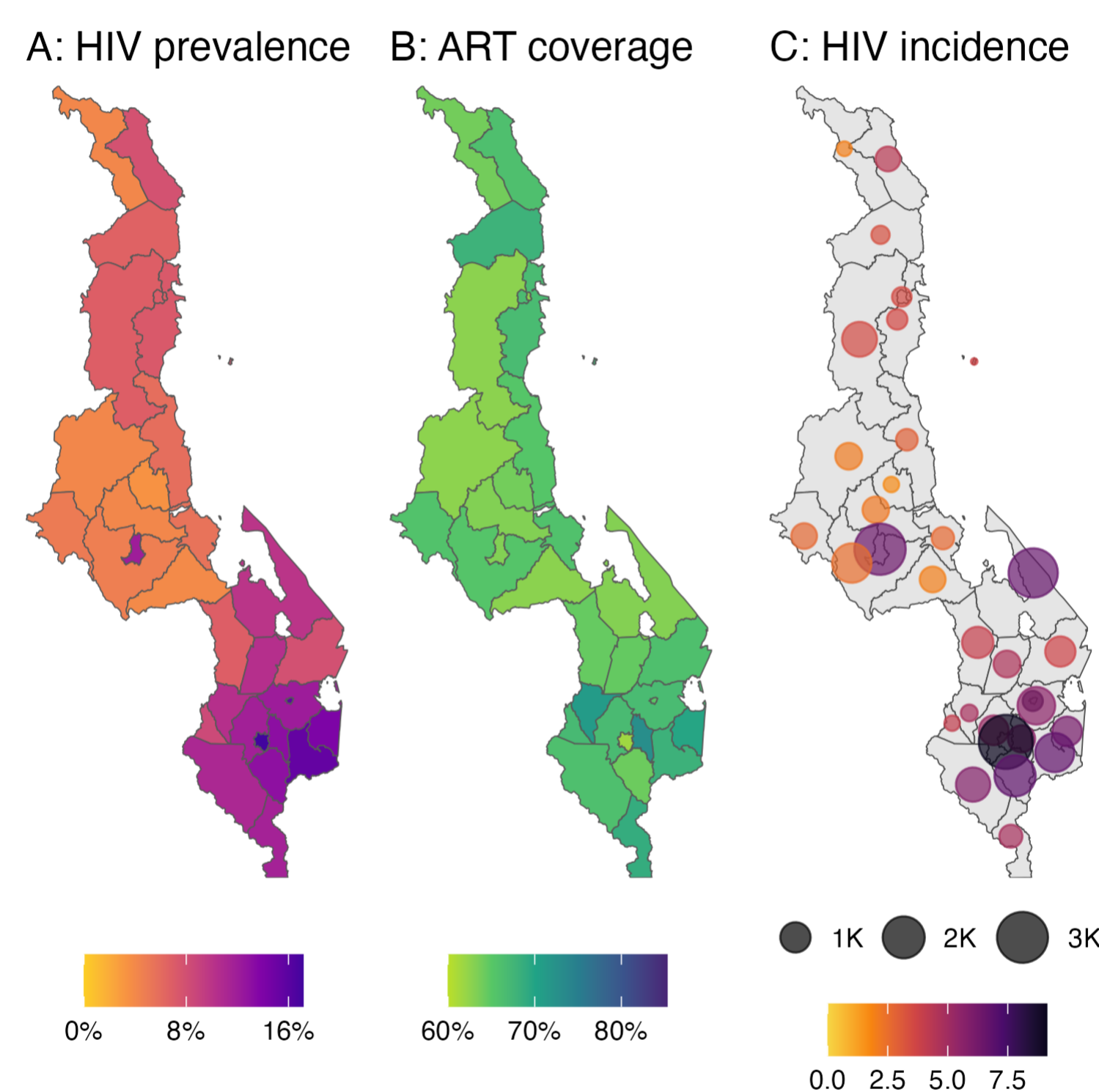


Figure 3: District-level model outputs for adults 15-49.

- Run three methods: 1. `TMB` (baseline, 54 secs), 2. `PCA-AGHQ` (new, 1 hour), 3. `NUTS` (gold-standard, 3.3 days)
 - For `PCA-AGHQ` $k = 3$ and $s = 8$ chosen using Scree plot to explain ~90% of variance
 - For `NUTS` 4 chains of 100,000 thinned by 40 were required for good diagnostics e.g. all $\hat{R} < 1.025$

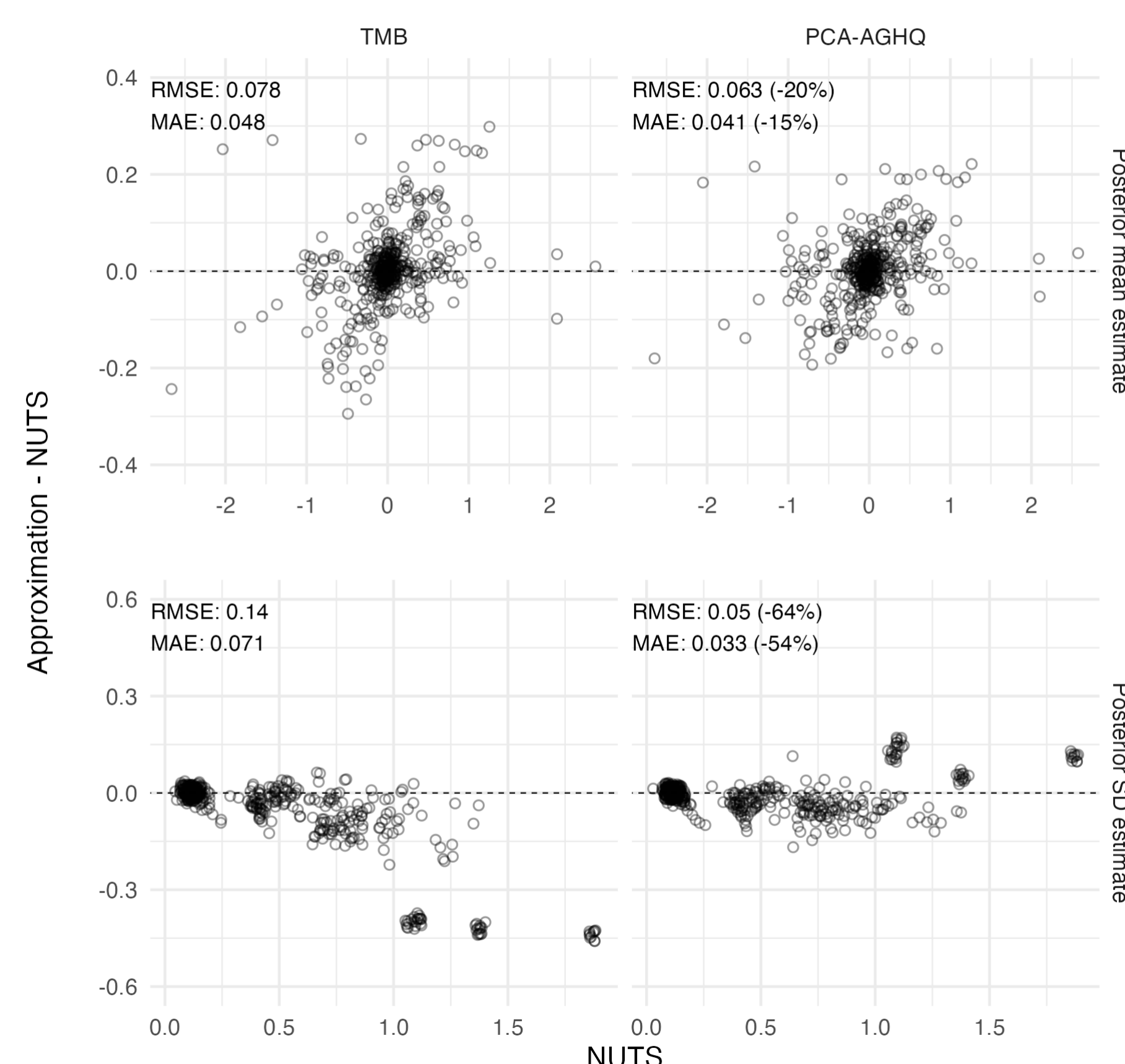


Figure 4: PCA-AGHQ moderately improves the posterior mean and substantially improves the posterior standard deviation

- `PCA-AGHQ` improves latent field point estimates (Figure 4) and distributional quantities like the Kolmogorov-Smirnov (KS, Figure 5) test statistic (-9% on average)

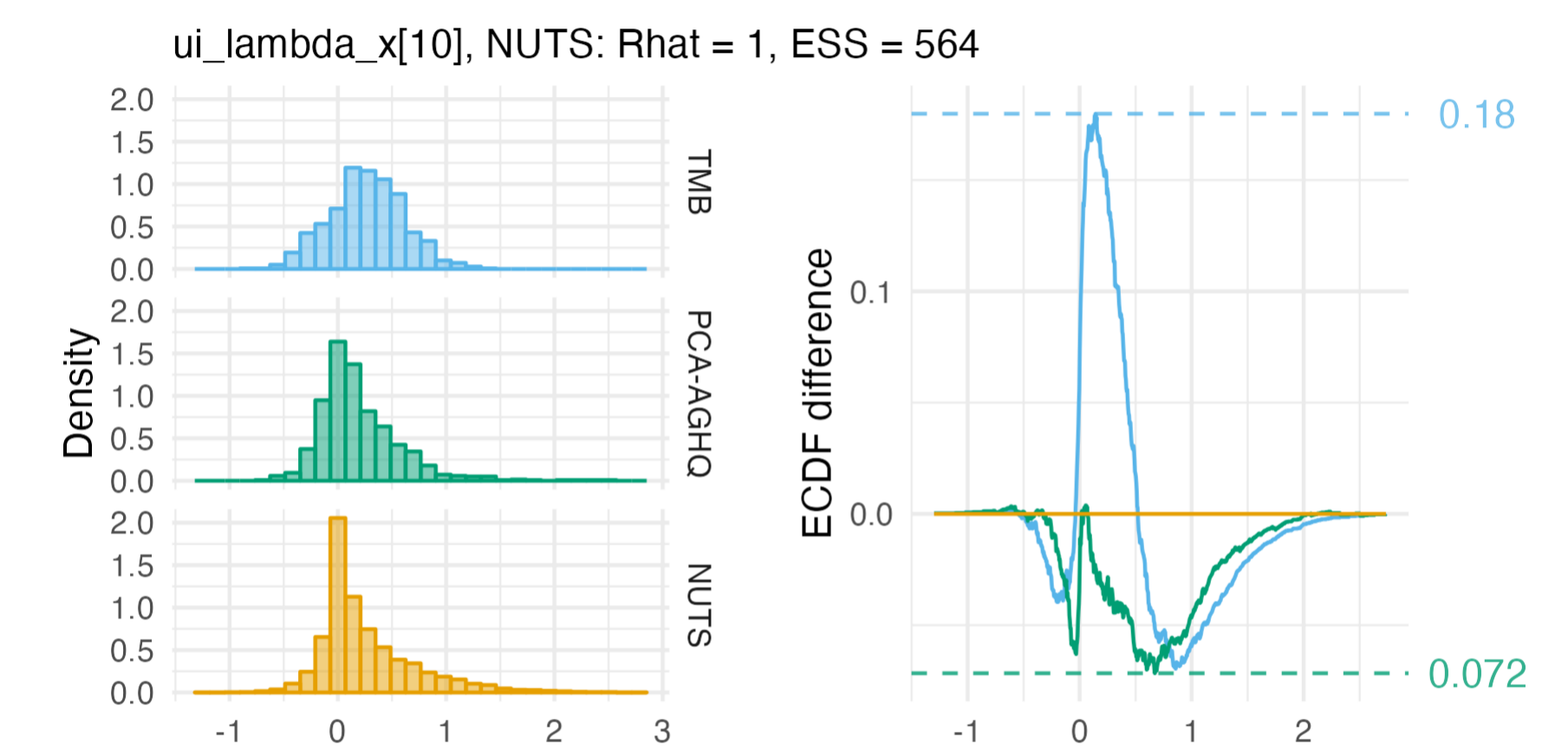


Figure 5: KS test demonstration.

- Naomi can be used to assess probabilities targets have been met e.g. 90% of those who know their HIV status are on ART ("second 90")
 - Though `PCA-AGHQ` is better (-16% RMSE), both `TMB` and `PCA-AGHQ` are biased (Figure 6)

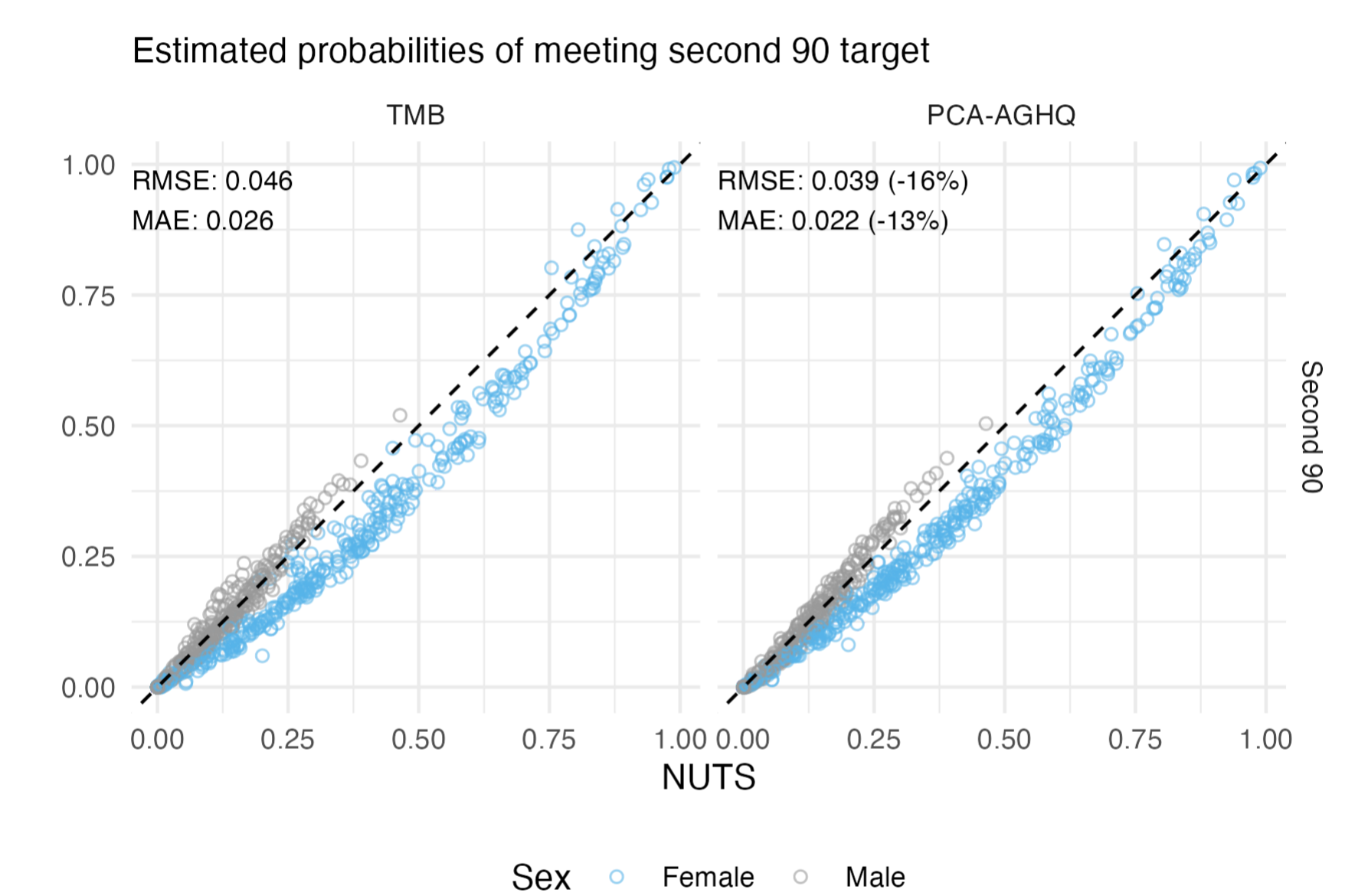


Figure 6: PCA-AGHQ moderately improves the second 90 bias.

4. Future directions

- Can we do better than moderate improvements?
 - Especially for the quantities of policy interest
- Laplace marginals with matrix algebra approximations (Wood 2020) to speed up calculations
- Further methods for allocation of effort to "important" dimensions of hyperparameter grid

Contact

- `</>` github.com/athowes/naomi-aghq
- `↗` athowes.github.io
- `✉` ath19@ic.ac.uk

Funding AH was supported by the EPSRC and Bill & Melinda Gates Foundation. This research was supported by the MRC Centre for Global Infectious Disease Analysis.

References

- Eaton, Jeffrey W., Laura Dwyer-Lindgren, Steve Gutreuter, Megan O'Driscoll, Oliver Stevens, Sumali Bajaj, Rob Ashton, et al. 2021. "Naomi: A New Modelling Tool for Estimating HIV Epidemic Indicators at the District Level in Sub-Saharan Africa." *Journal of the International AIDS Society* 24 (S5): e25788.
- Kristensen, Kasper, Anders Nielsen, Casper W Berg, Hans Skaug, Bradley M Bell, et al. 2016. "TMB: Automatic Differentiation and Laplace Approximation." *Journal of Statistical Software* 70 (i05).
- Monnahan, Cole C, and Kasper Kristensen. 2018. "No-U-turn sampling for fast Bayesian inference in ADMB and TMB: Introducing the `adnuts` and `tmbstan` R packages." *PLoS One* 13 (5): e0197954.
- Stringer, Alex. 2021. "Implementing Approximate Bayesian Inference using Adaptive Quadrature: the `aghq` Package." *arXiv Preprint arXiv:2101.04468*.
- Stringer, Alex, Patrick Brown, and Jamie Stafford. 2022. "Fast, Scalable Approximations to Posterior Distributions in Extended Latent Gaussian Models." *Journal of Computational and Graphical Statistics*, 1-15.
- Wood, Simon N. 2020. "Simplified integrated nested Laplace approximation." *Biometrika* 107 (1): 223-30.

