

Bayesian Modeling of Family Planning Indicators: The Family Planning Estimation Model (FPEM)

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- Introductions
- Family Planning Indicators
- Family Planning Estimation Model (FPEM)
- Demo

Links to all resources: http://herbsusmann.com/fp_map5

- PhD student, University of Massachusetts Amherst
- Research interests
 - Hierarchical spatio-temporal Bayesian modeling
 - Causal inference, targeted learning
- Website: <http://herbsusmann.com>, Twitter: [@herbps10](#)

- Leontine Alkema, University of Massachusetts Amherst
- Lab website: https://leontinealkema.github.io/alkema_lab/
- Bayesian modeling of demographic and health indicators
 - Family planning
 - Fertility
 - Sex ratio at birth
 - Maternal mortality
 - ...

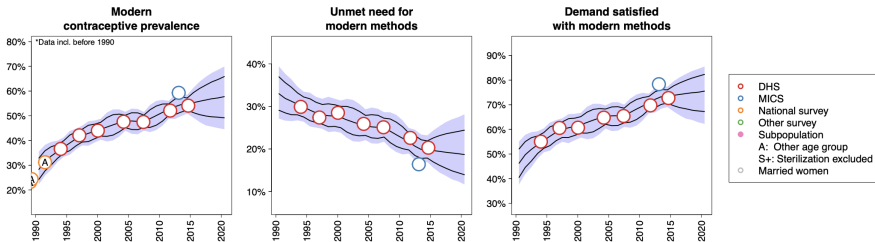
- Study population: women of childbearing age (15-49 years) currently married or in-union
- Modern contraceptive prevalence
 - Percentage of women who report themselves (or their partner) using a modern contraceptive method
- Traditional contraceptive prevalence
- Unmet need
 - Proportion of women who want to stop or delay childbearing but are not using any contraceptive method
- Demand satisfied with modern methods
 - Proportion of women who want to stop or delay childbearing and who use a modern contraceptive method

- International surveys (DHS, MICS, PMA)
- National Surveys
- Data are of varying quality and availability
- There may be multiple observations for the same country and year
- Data may exhibit systematic biases
 - Example: survey conducted for women of different age group than 15-49

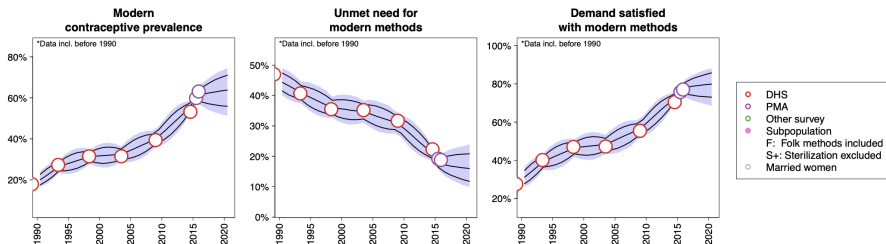


- Family Planning Estimation Model (FPEM): hierarchical Bayesian model of modern contraceptive use, traditional contraceptive use, and unmet need.
- Originally published as Alkema et al. (2013), updated by Cahill et al. (2018).
- Used by the United Nations Population Division and the Family Planning 2020 initiative.
- R packages: `FPEMglobal`, `FPEMlocal`.

Bangladesh



Kenya



- This presentation will focus on how FPEM models the modern contraceptive use rate indicator.
- True modern contraceptive use rate: $\eta_{c,t}$, for $c = 1, \dots, C$, $t = 1, \dots, T$.
- *Process model* describes evolution of $\eta_{c,t}$.
 - Systematic trends
 - Temporal smoothing
- Observed data y_i , with associated properties $c[i]$, $t[i]$, $s[i]$, ...
 - country $c[i]$, time point $t[i]$, data source $s[i]$
- *Data model* describes relationship between y_i and $\eta_{c[i],t[i]}$.



Statistics & Methodology

[Submitted on 19 Feb 2021]

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- The process model includes a systematic component and a smoothing component:

$$\text{logit}(\eta_{c,t}) = \underbrace{g(t, \eta_c, \alpha_c)}_{\text{systematic}} + \underbrace{\epsilon_{c,t}}_{\text{smoothing}}$$

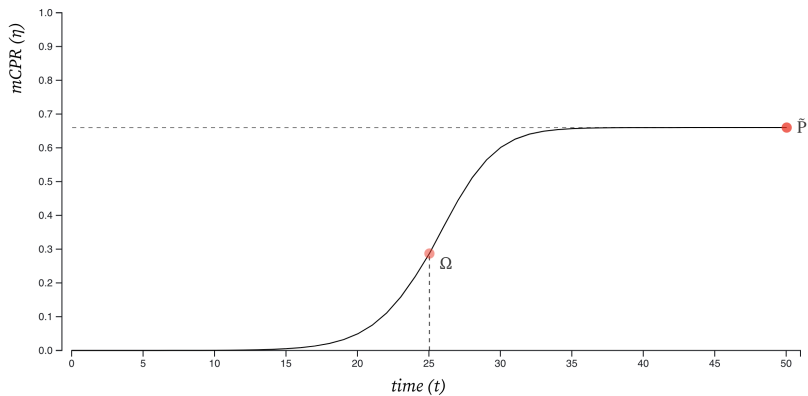
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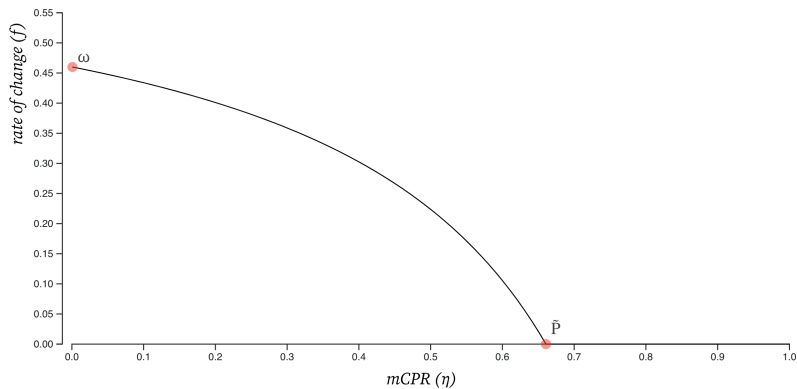
- The systematic component models logistic growth
- Three parameters: $\alpha_c = \{\Omega_c, \tilde{P}_c, \omega_c\}$
- We simulate forward and backward from a reference year t^* :

$$g(t, \eta_c, \alpha_c) = \begin{cases} \Omega_c, & t = t^*, \\ \text{logit}(\eta_{t-1}) + f(\eta_{t-1}) & t > t^* \\ \text{logit}(\eta_{t-1}) - f(\eta_{t-1}) & t < t^* \end{cases} \quad (1)$$

where

$$f(\eta_{c,t}, \tilde{P}_c, \omega_c) = \frac{(\tilde{P}_c - \eta_{c,t}) \cdot \omega_c}{\tilde{P}_c \cdot (\eta_{c,t} - 1)} \quad (2)$$





- Information is shared between countries through *hierarchical modeling*
- Example: pace parameter ω_c is given three-level hierarchical distribution (sub-regions, regions, and world).

$$\omega_c^* = \log \left(\frac{\omega_c - 0.01}{0.5 - \omega_c} \right) \quad (3)$$

$$\omega_c^* \sim N(\omega_{s[c]}^*, \kappa_{\omega}^{(c)}) \quad (4)$$

$$\omega_s^* \sim N(\omega_{r[s]}^*, \kappa_{\omega}^{(s)}) \quad (5)$$

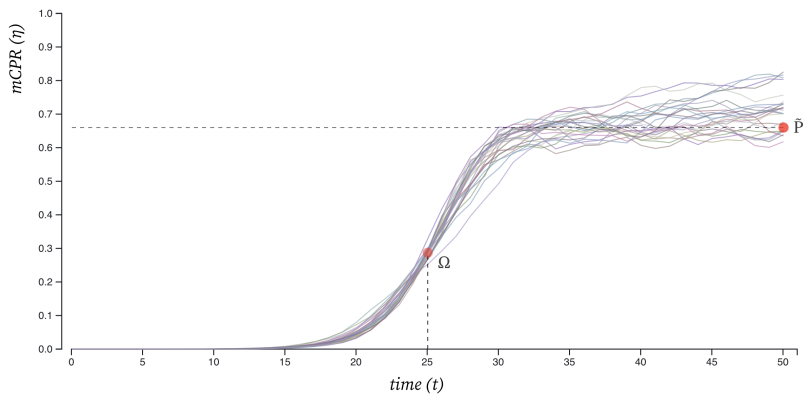
$$\omega_r^* \sim N(\omega_w^*, \kappa_{\omega}^{(r)}) \quad (6)$$

where $s[c]$ is the sub-region of country c and $r[s]$ is the region of sub-region s .

$$\text{logit}(\eta_{c,t}) = \underbrace{g(t, \eta_c, \alpha_c)}_{\text{systematic}} + \underbrace{\epsilon_{c,t}}_{\text{smoothing}}$$

- The smoothing component models trends not captured by logistic growth
- Affects the *rate of change* of the indicator
- FPEM assigns an AR(1) process to $\epsilon_{c,t}$, starting at $t = t^*$ and propagating forwards and backwards:

$$\epsilon_{c,t} \sim \begin{cases} \sim N(0, \frac{t^2}{1-\rho^2}), & t = t^* \\ \sim N(\rho \cdot \epsilon_{c,t-1}, \tau^2), & t > t^* \\ \sim N(\rho \cdot \epsilon_{c,t+1}, \tau^2), & t < t^* \end{cases} \quad (7)$$



- Demo: <https://observablehq.com/@herbps10/fpem>

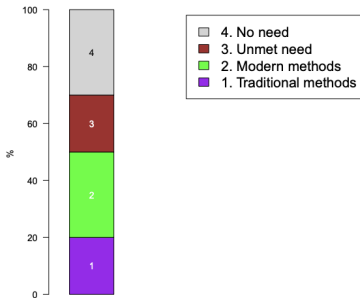
- Goal of the data model: describe the relationship between the truth and the observed data
 - The data model is "generative": given the true unobserved (latent) values of mCPR, it describes how the data is generated
- Toy data model for illustration:

$$y_i \sim N_{[0,1]}(\eta_{c[i],t[i]}, \sigma_{s[i]}^2) \quad (8)$$

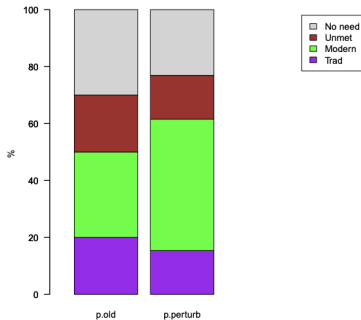
where $\sigma_{s[i]}^2$ is the estimated variance of data source $s[i]$.

- FPEM data model (in very brief): logistic normal for compositional data
- Incorporates sampling and non-sampling errors for each data source, as well as systematic biases
- See appendix of Alkema (2013) for detailed description

Composition of p



Perturbation of p



- Fitting FPEM to data from all countries is computationally intensive
- Individual countries may be interested in producing estimates for just their country using additional/different data
- General strategy
 - Fit FPEM to global data
 - For all non country-specific parameters, fix values to posterior estimates from global model
 - Perform full Bayesian inference using data from one country, using fixed global hyperparameters




- Fpemlocal R package provides routines for fitting FPEM to data from one country
- Demo code: <https://github.com/herbps10/fpemdmo>



Check for updates

SOFTWARE TOOL ARTICLE



Fpemlocal: Estimating family planning indicators in R for a single population of interest [version 1; peer review: 2 approved]

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<https://doi.org/10.12688/gatesopenres.13211.1>Latest published: 24 Feb 2021, 5:24
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Abstract

The global Family Planning Estimation model (FPEM) combines a Bayesian hierarchical model with country-specific time trends to yield estimates of contraceptive prevalence and unmet need for family planning for countries worldwide. In this paper, we introduce the R package *fpemlocal* that carries out the estimation of family planning indicators for a single population, for example, for a single country or smaller area. In this implementation of FPEM, all non-population-specific parameters are fixed at outcomes obtained in a prior global FPEM run. The development of this model was motivated by the demand for computational efficiency, without loss of model accuracy, when estimates and projections from FPEM were needed only for a single country. We present use cases to produce estimates for a single

Open Peer Review

Reviewer Status  

Invited Reviewers

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version 1		
24 Feb 2021	report	report
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1. Oliver Stevens 	Imperial College London, London, UK	
2. Qingfeng Li 	Johns Hopkins Bloomberg School of Public Health, Baltimore, USA	

Any reports and responses or comments on the