Bayesian Demographic Estimation and Population Reconstruction

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WPP Estimation Process



Bayesian Population Reconstruction was proposed as an improved method for reconstructing populations of the recent past.

Goals

- Quantify uncertainty probabilistically.
- Estimate all parameters consistently.
- Be easily replicable.
- Use all reliable data and expert opinion.

Current UN Practice



Bayesian Population Reconstruction



Indices

Age groups »*a*; $a \circ t^{\circ}$, $a = \bullet$; :::; \uparrow †, »‰•; 1° . Time periods »*t*; $t \circ t^{\circ}$, $t = t_{\bullet}$; :::; T †; t_{\bullet} is the "baseline" year. Sex `, *F*, *M*.

Parameters

Hierarchical Model

(= initial estimates and census counts which are fixed)

I. Likelihood $\log n_{a;t} j n_{a;t}; \frac{f}{n}$ Normal $\log n_{a;t}; \frac{f}{n}$

II. Projection Model

$$n_{a;t}$$
 j n_{t} +; f_{t} +; s_{t} +; g_{t} + = CCMPP¹ n_{t} +; f_{t} +; s_{t} +; g_{t} +

III. Priors on Inputs

 $\begin{array}{ccc} \log SRB_t \ j \ SRB_t; & {}^f_{SRB} & \text{Normal } \log SRB_t; & {}^f_{SRB} \\ & n_{a;t}, & \text{Unif } \bullet; K & ; K > \bullet \\ & \log f_{a;t} \ j & {}^f_t & \text{Normal } \log f_{a;t}; & {}^f_t \\ & \log t \ s_{a;t} \ j & {}^f_s & \text{Normal } \log t \ s_{a;t}; & {}^f_s \\ & g_{a;t} \ j & {}^f_g & \text{Normal } g_{a;t}; & {}^f_g \end{array}$

IV. Hyperparameters

$$\int_{V}^{f} InvGamma^{1} v; v^{\circ};$$

$$v 2 fn; f; s; g; SRBg$$

Bayesian Population Reconstruction

Hierarchical Model



The model has four levels. I shall start at level :

The true vital rates and baseline population are given priors, conditional on the initial estimates and variance parameters which account for measurement error.

The variance parameters are, themselves, given distributions at level

The priors on the vital rates and baseline counts induce a prior on subsequent counts through the deterministic projection model in level .

Level is a likelihood for the census counts which is parameterized by the projected counts and a variance parameter which accounts for measurement error.

- . Here are the things we have data on
- . Here are the things we have to specify
- . Here are the things we do inference on

Inference

Hierarchical Model: Key Relationships

(inputs are boxed)



 f_{V} InvGamma¹ $V; V^{\circ}; V = f; s; g; n;$

Bayesian Population Reconstruction

Measurement Error Variance

Measurement Error Variance

 $\begin{cases} InvGamma^{1} y; y^{0}; y = f; s; g; n; SRB \end{cases}$

The (Java crandom. They reflect non-systematic error in the initial estimates. We specify a and a using expert opinion provibly invested by expirat edimination of measurement error, or elicited as mean absolute instance error (BARS) through volumerets like "RMB probability" porcent, the initial estimates are accurate to within approximately parameter.

This is a flexible approach: some data sources have information about their accuracy, but others do not.

We choose the hyperparameters using the MAE of the marginal prior distributions of the observables. For example, we use the joint marginal distribution of the vector of fertility rates, ${}^{1}f_{cir}$; ::: $f_{A;T}$ ° which is

$$t_{AT} \log f$$
; $f \bullet f |_{AT}; f$; $, \bullet f; > \bullet$

Then

$$MAE^{1}\log f^{\circ} \quad E j \log f \quad \log f \quad j \quad \mathbf{i}$$

$$= E \quad E \quad j \log f \quad \log f \quad j \quad \mathbf{f}$$

$$= , AT \quad E \quad \mathbf{f}$$

$$= , AT \quad \mathbf{f}$$

$$= , AT \quad \mathbf{f}$$

$$= \frac{1}{1 + \frac{1}{2}} \cdot \mathbf{f}^{\circ}$$

$$= \frac{1}{1 + \frac{1}{2}} \cdot \mathbf{f}^{\circ}$$

$$= \frac{1}{2} \cdot \mathbf{f}^{\circ}$$

$$= \frac{1}{2} \cdot \mathbf{f}^{\circ}$$

where art is a length AT vector of s

The method is most useful for countries with unreliable, fragmentary data (high uncertainty).

However it also works for countries with very good data.

Reconstructing the female populations of Laos (–) and New Zealand (–)

The reconstruction periods are determined by the available data.

Population sizes are similar (counts in millions):

Laos — . . New Zealand . . .

Data quality and availability are very di erent.

Total Fertility Rate



% Posterior Interval Half-widths

	Mean Half-Width
Laos	•
NZ	

Bayesian Population Reconstruction

└─ Total Fertility Rate

- Total Factility Rate
- These plots show prior and posterior medians and the limits of % credible intervals.
 - The prior on TFR is in red, the posterior is in blue.
 - The initial estimates correspond to the prior median, shown by the red line.
- TFR in Laos was much higher than in New Zealand.
- Posterior uncertainty is quantified using the mean half-width of the credible intervals:
 - Mean half-width for New Zealand was about / th that of Laos.
- I show the WPP estimates for comparison

Total Net Migration

% Posterior Interval Half-widths



Bayesian Population Reconstruction

Inputs

Computation speed needs to be fast—could be a challenge with single-year age and time.

Needs to expand beyond span of most census collections, back to

Some countries have only one or two censuses, or rely on admin. data, household surveys.

Inputs and/or measurement error could be modelled (but see first point). Extensive testing required.

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