Accounting for Uncertainty About Past Values In Probabilistic Projections of the Total Fertility Rate for All Countries *  

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Abstract

Since the 1940s, population projections have in most cases been produced using the deterministic cohort component method. However, in 2015, for the first time, in a major advance, the United Nations issued official probabilistic population projections for all countries based on Bayesian hierarchical models for total fertility and life expectancy. The estimates of these models and the resulting projections are conditional on the UN’s official estimates of past values. However, these past values are themselves uncertain, particularly for the majority of the world’s countries that do not have longstanding high-quality vital registration systems, when they rely on surveys and censuses with their own biases and measurement errors. This paper is a first attempt to remedy this for total fertility rates, by extending the UN model for the future to take account of uncertainty about past values. This is done by adding an additional level to the hierarchical model to represent the multiple data sources, in each case estimating their bias and measurement error variance. We assess the method by out-of-sample predictive validation. While the prediction intervals produced by the current method have somewhat less than nominal coverage, we find that our proposed method achieves close to nominal coverage. The prediction intervals become wider for countries for which the estimates of past total fertility rates rely heavily on surveys rather than on vital registration data.

Keywords: Bayesian hierarchical model, Markov chain Monte Carlo, Measurement error, Population projection, Total fertility rate, Vital registration
1 Introduction

Population projections or forecasts consist of forecasts of future population numbers and also the components of population change, namely births, deaths and migration, broken down by age and sex, and possibly also by other categories such as race. They are used by governments at all levels (local, regional, state, national and international) for planning and policy decision-making, since knowing the future numbers of people is key to government policy-making. They are also used by the private sector for strategic decisions, and by researchers in the health and social sciences.

The most widely used population projections for many individual countries are produced by their national statistical agency, such as the U.S. Census Bureau in the case of the United States [U.S. Census Bureau (2017)]. The United Nations publishes projections of population by age and sex, and mortality, fertility and migration rates for all countries by five-year age-groups in five-year periods to the year 2100, updated every two years in the UN’s World Population Prospects, whose most recent edition was published in 2017.
The UN’s population projections are widely viewed as the gold standard regularly updated projections for all countries (Lutz and Samir 2010). Since the 1940s, population projections have in most cases been produced by a deterministic method called the cohort-component method (Cannan 1895; Whelpton 1928 1936; Preston et al. 2000). This is based on the demographic balancing equation, namely

\[
\text{Population}_{t+1} = \text{Population}_t + \text{Births}_t - \text{Deaths}_t + \text{Immigrants}_t - \text{Emigrants}_t,
\]

where Population refers to the number at time \( t \), and Births, Deaths, Immigrants and Emigrants refer to the numbers in the time interval from time \( t \) to time \( t + 1 \). The cohort-component method uses an age-structured version of this, of which a simple form is

\[
\begin{align*}
\text{Population}_{a+1,t+1} &= \text{Population}_{a,t} \times \text{Survival Rate}_{a,t} + \text{Net Migration}_{a,t}, \\
\text{Population}_{0,t+1} &= \sum_a \text{Women}_{a,t} \times \text{Fertility Rate}_{a,t}.
\end{align*}
\]

This method is simple to implement, but it requires assumptions about future fertility, mortality and migration rates by age and sex. These are typically produced subjectively by experts, either in-house experts working at the agency producing the projections, or a panel of outside experts assembled by the agency. Uncertainty is communicated by scenarios; for example the UN traditionally published High, Medium and Low variants, in which the total fertility rates (TFR) for all countries and all future periods were increased or decreases by half a child per woman. This deterministic approach has been extensively criticized on the grounds that it has no probabilistic basis, and it can give implausible results over multiple projection periods (Keyfitz 1981; Stoto 1983; Lee and Tuljapurkar 1994); for a review and summary of this literature see the National Research Council report on the topic (Lee and Bulatao 2000).

Methods for probabilistic forecasting of future fertility rates have been proposed by Lee (1993), Alders et al. (2007), Alho et al. (2006), Alho et al. (2008) and Booth et al. (2009), in each case either for individual countries or groups of countries, typically in the developed world, but these were not easily applied to the U.N.’s task of producing forecasts for all countries.

In 2015, the U.N. adopted a different method for their official population projections for all countries (United Nations 2015b). This method was probabilistic and statistically-based, replacing the previous deterministic method, thus responding to the critiques. They
used Bayesian hierarchical models to produce probabilistic projections of the total fertility rate (Alkema et al., 2011, Raftery, Alkema and Gerland 2014, Fosdick and Raftery, 2014, Ševčíková et al., 2011), and life expectancy (Raftery et al., 2013, Raftery, Lalic and Gerland 2014). These projections were then simulated from, and each simulated trajectory was translated into age-specific fertility and mortality rates, which in turn were input into the cohort-component method to yield many possible future population trajectories of all countries (Ševčíková et al., 2016, Ševčíková and Raftery, 2016).

This method indicated that world population was likely to be higher than had previously been thought, reaching 11.2 billion (95% prediction interval 9.5 to 13.2 billion) in 2100, from 7.4 billion now (Gerland et al., 2014, United Nations 2017). The main reason for this is that fertility in high-fertility countries, many of them in Sub-Saharan Africa, has been declining more slowly than experts had expected, and the statistical approach took this into account more fully than the expert-based assumptions.

Although the new U.N. method takes account of uncertainty more systematically than previous methods, there are still sources of uncertainty that it does not account for. The Bayesian hierarchical model used by the U.N. is conditional on estimates of present and past population, and fertility and mortality rates. In countries with long-established high quality vital registration systems, and hence accurate counts of births and deaths, this is not a large source of uncertainty; this is the case for about 80 of the world’s 200 or so countries. However, the remaining 120 or so countries do not have longstanding high quality vital registration systems, and there fertility and mortality rates are typically estimated from surveys that can be subject to poor coverage in time and space, biases and measurement error. For example, the Demographic and Health Surveys (DHS) are one of the most important and reliable sources of data on fertility rates in countries without good vital registration [National Population Commission (2009), but they have suffered from large underestimates of TFR in some countries in Sub-Saharan Africa, according to Schoumaker (2010, 2011, 2014) and Pullum et al. (2013).

Thus the estimated present and past vital rates and population numbers for these countries are not exact, and the uncertainty about them is not accounted for in the projections. This may lead uncertainty in the projections to be underestimated (Abel et al., 2016). Demographers have developed methods for correcting estimates of TFR for specific forms of bias, such as recall errors, developing indirect estimation methods for this purpose (Brass, 1964, 2015). Bias and uncertainty of past and present estimates were modeled by Alkema et al. (2012), using multiple data quality indicators, such as source of the data, estimation
method (e.g. direct or indirect), recall time for retrospective birth history surveys, and so on. But these methods have not been used to account for the uncertainty in population projections that is due to uncertainty about past and present values.

In this paper we extend the UN probabilistic projection method to account for uncertainty about past and present total fertility rates, which may be the most important remaining unaccounted for source of uncertainty. This is made possible by the recent publication of a new dataset by the U.N. Population Division that contains not just estimates of past and present fertility rates for all countries, but also the data from all the data sources on which the estimates are based, including censuses, vital registration systems, partial and sample vital systems, international surveys such as the DHS and the Multiple Indicator Cluster Surveys, or MICS Fund (2016), and national, regional and local surveys (United Nations, 2015). We do this by developing a new Bayesian hierarchical model that extends the U.N. model to account for bias and measurement error in the different information sources.

The article is organized as follows. The data and proposed methodology are described in Section 2. In Section 3 we report the method’s performance using out-of-sample predictive validation. We then provide more detail in Section 4, which is a case study of how the method works for Nigeria, which is one of the most important countries for uncertainty about future world population, because it is the most populous country in Africa, has very high fertility, and does not have a long-established high-quality vital registration system. We conclude with a discussion in Section 5.

2 Method

2.1 Notation

We restrict our attention to estimation of the TFR of each country. The TFR is a period measure, defined as the number of children a woman would bear if she survived to the end of the reproductive interval and at each age she experienced the age-specific fertility rates prevalent in the period to which it refers. It is defined in units of children per woman.

We use the symbol $y$ to denote TFR estimates from different data sources and the symbol $f$ to denote the true (unobserved) TFR. Although the U.N. Population Division’s estimates of past TFR values do contain error, we assume that they are unbiased, in the sense that the errors do not tend to be systematically in one direction or the other; for discussion of this assumption see Alkema et al. (2012). These official U.N. estimates of past TFR
values will be denoted by $u$. All of these parameters will be indexed by country $c$ and time $t$. Data from different sources $y$ are also indexed by their source, denoted by $s$, and the bias and measurement error variance of these estimates are denoted by $\delta$ and $\rho^2$, respectively. The quantities of interest are the unknown past, present and future TFR $f$. We estimate past TFR for the time period $[t_0, t_1]$, while prediction will be for the period $[t_1, t_2]$. In practice in this article, $t_0 = 1950$, $t_1 = 2015$ and $t_2 = 2100$.

The three-phase Bayesian hierarchical model of Alkema et al. (2011) will be used to model the total fertility rates. For describing the Bayesian hierarchical model, country-specific parameters controlling the shape of total fertility rates of country $c$ are denoted by $\theta_c$, and the global parameters are denoted by $\psi$. In constructing the probabilistic projections of TFR for all countries, we are also interested in the country-specific parameters $\theta_c$.

2.2 Data

We use the World Fertility Data 2015 (United Nations, 2015a) from the U.N. Population Division for 201 countries in the world. This database is publicly available and includes estimates of TFR from surveys, censuses and sample or partial vital registration data for countries without high-quality vital registration systems. It includes data available as of November 2015 and covers the time period from 1950 to 2015. These data were used to produce the estimates of past TFR in the United Nations World Population Prospects (WPP) 2015 Revision. These estimates were in turn part of the basis for the U.N.’s 2015 population projections for all countries.

We use TFR estimates from national and international surveys, indirect estimates and vital registration for all 201 countries to estimate the bias and variance of different data sources. We take the estimates in the WPP 2015 revision as a baseline, assuming that they are unbiased (but not that they are without error). This assumption, also used by Alkema et al. (2012), is made because the analysts producing past estimates were often aware of sources of bias in datasets and corrected for them. While this assumption is not perfect, it seems reasonable to argue that WPP provides the least biased set of estimates available. In the 2015 revision of the WPP, the U.N. estimated the five-year average TFR, $u_{c,t}$, for country $c$ in time period $(t, t+5)$, for each five-year period from 1950 to 2015. The outcome in each five-year period $(t, t + 5)$ is an estimate of the average TFR between July 1 of year $t$ and July 1 of year $t + 5$, and so centered on January 1 of year $t + 3$. We construct trajectories and estimations in five-year intervals, and project TFR up to year 2100 probabilistically.
according to these estimated trajectories of the past.

2.3 Model

Three Phase Bayesian Hierarchical Model: Our methodology builds on that of Alkema et al. (2011) and Raftery, Alkema and Gerland (2014), as implemented by Ševčíková et al. (2011). This divides the evolution of TFR in a country into three phases: pre-demographic transition, transition and post-transition, as illustrated in Figure 1.

Figure 1: Illustration of the three phases in the typical evolution of fertility in a country: pre-transition high fertility (Phase I; grey), transition from high to low fertility — (Phase II; red), and post-transition fertility fluctuations and recovery (Phase III; green).

During the fertility transition or decline phase (Phase II), the total fertility rate is modeled as a random walk with negative drift, namely

\[ f_{c,t} = f_{c,t-5} - g(f_{c,t-5} | \theta_c) + \epsilon_{c,t}, \]  

where \( g(\cdot | \theta_c) \) is the expected five-year decrement in the TFR over the next period, modeled by a double logistic function governed by the country-specific parameter vector \( \theta_c = (\Delta_{c1}, \Delta_{c2}, \Delta_{c3}, \Delta_{c4}, d_c) \), and \( \epsilon_{c,t} \) is random noise around the expected decrement.
During the post-transition phase (Phase III), the total fertility rate is modeled by a Bayesian Hierarchical Autoregressive Model as:

\[ f_{c,t} = \mu_c + \rho_c(f_{c,t-5} - \mu_c) + \varepsilon_{c,t}, \]  

where \( \mu_c \) is the long-term mean of the TFR for country \( c \), and \( \varepsilon_{c,t} \) is the random noise similar to that in phase II.

Since all or almost all countries have already started the fertility transition, modeling the TFR during the pre-demographic transition Phase I was not necessary for projection purposes in previous work. However, for constructing probabilistic estimation of past TFR from 1950 to 2015, we do need to model the Phase I data. They are modeled by a random walk model from year 1950 to the start of fertility transition as:

\[ f_{c,t} = f_{c,t-5} + \varepsilon_{c,t}. \]  

The country-specific parameters in all three phases, \( (\theta_c, \mu_c) \), follow a world distribution, which is governed by world parameters \( \psi \), and these in turn have a prior distribution. The start and end of the fertility transition (phase II) are defined based on the UN estimates \( u_{c,t} \), by rules given in Alkema et al. (2011).

**Model of Imperfect Data:** The TFR estimates from different data sources \( y_{c,t,s} \) are modeled based on the unobserved true value \( f_{c,t} \). Building on Alkema et al. (2012), we distinguish between the bias and measurement error variance in our model. The estimated TFR are modeled by a conditional normal distribution as:

\[ y_{c,t,s} | f_{c,t} \sim \mathcal{N}(f_{c,t} + \delta_{c,s}, \rho_{c,s}^2), \]

\[ \mathbb{E}[\delta_{c,s}] = x_{c,s} \beta, \]  
\[ \mathbb{E}[\rho_{c,s}] = x_{c,s} \gamma. \]

The bias and measurement error variance, \( \delta_{c,s} \) and \( \rho_{c,s} \), are estimated using data quality indicators, denoted by \( x_{c,s} \). The estimation process is described in the following sections.

**Complete Model Layout:** We combine the three-phase Bayesian hierarchical model and imperfect data model into a four-level Bayesian hierarchical model with an additional level for the data sources. Estimation and prediction is then equivalent to getting the posterior
distribution of the unknown TFR values $f_{c,t}$ in the estimation period $[t_0, t_1]$ and the prediction period $[t_1, t_2]$, based on the observed TFR estimates from different data sources.

The observed estimates of TFR can be measured for any time between $t_0$ and $t_1$. However, we seek estimates of the average over five-year periods. We approximate the true TFR at any time by assuming that the TFR evolves linearly between the centers of any two successive five-year intervals. This is a reasonable assumption because most demographic quantities, including TFR, typically evolve relatively smoothly over time. Specifically, for any $t \in [t_\ell, t_\ell + 5]$, where $t_\ell$ and $t_\ell + 5$ are the centers of two successive five-year periods, we assume that

$$f_{c,t} = \frac{1}{5}[(t_{t+1} - t)f_{c,t_{t}} + (t - t_{t})f_{c,t_{t+1}}].$$

(7)

Then, we model the observed TFR estimates in Level 1, conditional on the true total fertility rates, which are modeled with the extant three-phase BHM in Level 2, conditional on the country-specific parameters. The country-specific parameters are then modeled conditionally on the global parameters in Level 3, which have a prior distribution specified by hyperparameters (Level 4). The overall model is specified as follows:

**Level 1:** $y_{c,t,s} | f_{c,t} \sim \mathcal{N}(f_{c,t} + \delta_{c,s}, \rho_{c,s}^2)$,

$$\mathbb{E}[\delta_{c,s}] = x_{c,s} \beta;$$

$$\mathbb{E}[\rho_{c,s}] = x_{c,s} \gamma;$$

$$f_{c,t} = \frac{1}{5}[(t_{t+1} - t)f_{c,t_{t}} + (t - t_{t})f_{c,t_{t+1}}] \text{ for } t \in [t_t, t_{t+5}];$$

**Level 2:**

**Phase I:** $f_{c,t} = f_{c,t-5} + \varepsilon_{c,t};$

**Phase II:** $f_{c,t} = f_{c,t-5} - g(f_{c,t-5} | \theta_c) + \varepsilon_{c,t};$

**Phase III:** $f_{c,t} = \mu_c + \rho_c(f_{c,t-5} - \mu_c) + \varepsilon_{c,t};$

$$\varepsilon_{c,t} \sim \mathcal{N}(0, \sigma_{c,t}^2);$$

**Level 3:**

$$\theta_c \sim h(\cdot | \psi),$$

$$\mu_c \sim \mathcal{N}(\bar{\mu}, \sigma_\mu^2),$$

$$\rho_c \sim \mathcal{N}(\bar{\rho}, \sigma_\rho^2);$$

**Level 4:**

$$\psi, \bar{\mu}, \sigma_\mu, \bar{\rho}, \sigma_\rho \sim \pi(\cdot).$$

Here, $g$ denotes the double logistic function, and $h$ and $\pi$ denote the conditional and
unconditional distributions of the parameters of interest, respectively. The parameter $\theta_c$ controls the shape of the double logistic curve. The functional form of the prior distribution $\pi(\cdot)$ is as specified by Alkema et al. (2011). A complete specification of the model, including some further details, is given in the Supplementary Information.

Inference is based on the joint posterior distribution of $(f_{c,t}, \theta_c)$. The model is summarized graphically in Figure 2.

![Figure 2: Model Specification](image)

Figure 2: Model Specification: $y_{c,t,s}$ are the observed TFR, $f_{c,t}$ are the unknown TFR values, $\theta_c$ are the country-specific parameters and $\psi$ are the global parameters.

### 2.4 Estimation

**Estimation of Bias and Measurement Error Variance:** The bias $\delta_{c,s}$ and measurement error variance, $\rho_{c,s}^2$, of the observed TFR estimates are estimated in a first stage, as input to the Bayesian hierarchical model, building on the method of Alkema et al. (2012).

We first estimate the bias of TFR. As we discussed in Section 2.3, the UN estimates will be treated as unbiased but not error-free, providing a baseline reference. Then, for each observation $y_{c,t,s}$, we have

$$\mathbb{E}[y_{c,t,s} - u_{c,t}] = f_{c,t} + \delta_{c,s} - f_{c,t} = \delta_{c,s}.$$ 

Thus we can use the difference between observed TFR and UN estimates, $(y_{c,t,s} - f_{c,t})$,
as samples for our estimation of the bias and measurement error variance of each source. The parameters $\beta$ are estimated by linear regression on data quality indicators $x_{c,s}$, as in equation [5]. The estimated biases $\hat{\delta}_{c,s}$ are then equal to the fitted values $x_{c,s}\hat{\beta}$.

We estimate the source-specific measurement error variance of the TFR estimates by regression on the data quality covariates $x_{c,s}$ of the plug-in estimate $\rho_{c,s} = \sqrt{\frac{\pi}{2}}E|z_{c,t,s} - u_{c,t}|$, where the unobserved true values $f_{c,t}$ are replaced by the UN estimates of TFR (taken to be unbiased), where the fitted values are used.

**Estimation of the Complete Model:** Given the estimated bias $\hat{\delta}_{c,s}$ and measurement error variance $\hat{\rho}_{c,s}^2$, we estimate the Bayesian hierarchical model for TFR using a purpose-built Markov Chain Monte Carlo (MCMC) algorithm, specially coded in R. The roughly 3,600 parameters and unknown TFR values are updated one at a time, using Gibbs steps, Metropolis-Hastings steps or slice sampling (Neal, 2003) for each parameter as appropriate. We monitored convergence by inspecting trace plots and using standard convergence diagnostics (Gelman and Rubin, 1992; Raftery and Lewis, 1996).

We thin enough for the thinned sample to be roughly independent. In practice, for the final results we ran 3 chains, each of length 12,000 iterations with a burn-in of 2,000, and we thinned the resulting chains by 10, to obtain a final, approximately independent sample of size 1,000 from the posterior distribution. More information about the convergence diagnostics used is provided in the Supplementary Information.

### 2.5 Prediction of Future TFR

Unlike the projection process developed by Alkema et al. (2011) and used by the U.N., we have probabilistic rather than point TFR estimates of past rates over the time period $[t_0, t_1]$. Thus, instead of just sampling from the posterior trajectories of country-specific parameters obtained from estimation process, we also generate posterior trajectories of past TFR values.

We proceed by repeating the following process many times. We first select a joint sample of model parameters and past and present TFR for all countries from the posterior distribution. Then, given the sampled model parameters and past and present TFR values, we simulate a trajectory of future TFR values, from 2015 to 2100 using the model specified by (1) and (2). This yields a sample from the joint posterior predictive distribution of future TFR in all countries and time periods considered, taking account of uncertainty about past
values.

Our method also differs slightly from the extant method in the way the end of the fertility transition, at which the model shifts from that for Phase II to that for Phase III, is determined. The current U.N. method uses deterministic rules based on the U.N. estimates (Alkema et al., 2011), and does not account for uncertainty about when the fertility transition ended. In our method, we retain the deterministic rules, but apply them separately to each sampled trajectory of past TFR values. Thus our method takes account of uncertainty about when the fertility transition ended in a particular country, and hence which phase the country is in at the end of the estimation period.

3 Results

We assess the predictive performance of our model using out-of-sample predictive validation, used for probabilistic forecasts, for example, by Raftery et al. (2005). We include all countries and regions in our validation exercise.

3.1 Study Design

The data we have cover the period from 1950 to 2015. We split this into the estimation period, \([t_0 = 1950, t_1 = 2005]\), and the prediction period, \([t_1 = 2005, t_2 = 2015]\). The inputs to our method consist of all TFR estimates from different sources referring to the estimation period.

For the U.N. estimates used as a reference, we take the values published in the WPP 2008 revision (United Nations, 2008). The U.N. estimates of the past have been refined since then as more data have become available, but we deliberately do not take advantage of this in our estimation. This makes our validation exercise more analogous to the real prediction task at hand, for which we are using U.N. estimates in the WPP 2015 revision of past TFR values up to 2015 to predict values past 2015. It can be expected that these estimates of TFR values up to 2015 will become more accurate in the future as data accumulate, but we are not able to take advantage of this for the present purpose.

We are making probabilistic projections, and so we evaluate not only the point predictions, but also the predictive intervals. Our aim is to account for an important source of uncertainty ignored by the present state of the art method, so the accuracy of the prediction intervals may be even more important than the point predictions. If our method is work-
ing well, we would expect the current state of the art intervals to have less than nominal coverage, and our method to give coverage closer to nominal. To evaluate our method, we compare our probabilistic projections with those produced by the U.N. in WPP 2015.

Our out-of-sample validation experiment proceeds as follows.

1. Choose the subset of the original data set \( D \) with TFR observed before year 2005 as the training data \( D_{\text{train}} \). We remove those observations before 2005 for those estimates in studies that provide series of estimates ending after 2005. For example, if a study lasts for 20 years and ends in 2008, yielding TFR observations for 1988 to 2008, we remove all observations from this study even though some of the estimates are for years before 2005.

2. Estimate bias and measurement error variance for all data points using UN estimates \( u_{c,t} \) from WPP 2008 revision as the reference.

3. Draw a sample from the joint posterior distribution of model parameters and past TFR values for 1950 to 2005, using MCMC.

4. For each sampled trajectory including the unobserved past TFR values and the model parameters, determine the TFR phase of country \( c \) for each time period for this trajectory, and make probabilistic projections for the projection period \([2005, 2015]\).

### 3.2 Out of Sample Validation Results

We produce results for all countries using our method. For comparison, we also produce results using the method of Alkema et al. (2011), which underlies the current U.N. methodology and does not take account of uncertainty about past TFR values.

We summarize the results in Table 1. This is based on the predictive intervals for each of the 201 countries and for both of the periods \([2005, 2010]\) and \([2010, 2015]\), so that each entry in Table 1 is an average over \(201 \times 2 = 402\) values. For each TFR value to be predicted, we take the predictive median as the point estimate, and we compute the quantile-based 80% and 95% prediction intervals. The table shows the mean absolute error (MAE) of the point estimates (the smaller the better), and the coverage of the prediction intervals (the closer to the nominal value the better).

The proposed method improves the point predictions slightly over the current method, as measured by the MAE. However, it improves the coverage of the prediction intervals.
Table 1: Mean Absolute Error and Coverage of Out of Sample TFR Point and Interval Predictions for Current Method (Alkema et al., 2011) and Proposed Method

<table>
<thead>
<tr>
<th></th>
<th>Current Method</th>
<th>Proposed Method</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mean Absolute Error</td>
<td>0.250</td>
<td>0.242</td>
</tr>
<tr>
<td>Coverage of 80% interval</td>
<td>74.0%</td>
<td>79.6%</td>
</tr>
<tr>
<td>Coverage of 95% interval</td>
<td>86.7%</td>
<td>94.5%</td>
</tr>
</tbody>
</table>

substantially. Under the current method, the coverage of the prediction intervals is somewhat below the nominal level, indicating that some of the uncertainty is being missed. Under the proposed method, the coverage of the prediction intervals is much closer to the nominal level, suggesting that the new method is capturing most of the missed uncertainty by taking account of uncertainty in past TFR values.

For illustration, results of the out-of-sample validation exercise are shown in Figure[3] for Argentina, Botswana, Nigeria and the United States. Of these, only the United States has had a high-quality vital registration system for the entire period, while Argentina has a vital registration that was of lower quality in the early years, and the other two countries have no comprehensive vital registration systems, relying instead on censuses and periodic surveys.

It can be seen that the posterior intervals of past TFR values are very narrow for the United States, reflecting the high quality vital registration data available for the entire period, while for Argentina they are somewhat wider. For both Botswana and Nigeria the intervals are far wider, reflecting the much lower quality of the available data. For the earlier years, from the 1950s to the 1970s, the intervals for Botswana and Nigeria were especially wide, reflecting the sparsity of the data for these decades. The predictive distributions cover the observations in all cases, although in some cases they lie towards the edge of the intervals, as expected if the intervals are well calibrated.

4 Case Study: TFR Estimation and Projection For Nigeria

In this section, we illustrate the method by producing probabilistic forecasts of the TFR of Nigeria from 2015 to 2100, using data available up to 2015. As we have discussed, the method first estimates the bias and measurement error variance of the different data sources. It then estimates the uncertainty about past TFR values, and takes this uncertainty
into account when making probabilistic projections.

4.1 Estimation of Bias and Measurement Error Variance of Different Data Sources

From 1950 to 2015, according to the U.N.’s WPP 2015 revision, the TFR in Nigeria reached its peak around 1980 at about 6.7 children per woman. It then declined slowly, reaching about 5.7 in 2015. However, the data on which these estimates are based are surprisingly noisy, as can be seen in Figure 4.

These data come from several sources, including national censuses, which are comprehensive but sparse in time and have issues of coverage. The other sources are mostly surveys, including the internationally organized Demographic and Health Surveys (DHS), the Multiple Indicator Cluster Surveys (MICS) run by UNICEF, and the Malaria Indicators Survey, or MIS, also run by DHS. There are also several occasional national cross-sectional and panel surveys. Some of the surveys, notably DHS and MICS, collect birth histories,
which allow one survey to generate estimates for several past years, in some cases using indirect methods.

We first estimate the bias and measurement error variance of the different data sources using the approach outlined in Section 2.3. From each observed TFR estimate $f_{c,t,s}$ we subtract the corresponding UN TFR estimate to obtain an estimate of the bias for that source, country and time, namely $z_{c,t,s} = f_{c,t,s} - u_{c,t}$. As data quality covariates, $x_{c,s}$, we use the source of the data and whether the estimate is direct or indirect. We then estimate the bias $\delta_{c,s}$ for country $c$ and data source $s$ as the fitted value from a regression of the $z_{c,t,s}$ on the data quality covariates $x_{c,s}$, as in Alkema et al. (2012).

The U.N. TFR estimates are for five-year periods, and we treat them as referring to the middle of the period. Thus, for example, we treat estimates for the 2010–2015 period as referring to the beginning of 2013. An observed TFR estimate can refer to any year between 1950 and 2015, and we use the convex combination of the two U.N. estimates closest to the time to which it refers as the corresponding U.N. estimate, $u_{c,t}$.

Similarly, after we get the fitted value of the bias estimates $\hat{\delta}_{c,s}$, we obtain the measurement error standard deviation estimates by regressing $|z_{c,t,s} - \delta_{c,s}|$ on the same data quality covariates. The fitted biases and measurement error standard deviations are summarized in Table 2.

We can see from Table 2 that direct estimates from the DHS are the highest quality estimates as measured by estimated mean squared error (equal to $\sqrt{\hat{\delta}^2 + \hat{\rho}^2}$). Direct estimates generally have smaller variances than indirect estimates. Figure 5 plots the fitted biases
Table 2: Estimates of Bias and Measurement Error Variance for All Combinations of Source and Estimate Types. Survey-NR are different Nigeria nationwide surveys and Survey represents other survey estimates. Under estimate type, D represents direct estimates, C cohort estimates and I indirect analysis. Here $\mu(\delta)$ and $\sigma(\delta)$ are the sample bias and measurement error standard deviations; when a hat is added they represent the estimates from the models. Estimated root mean squared errors are summarized in the column RMSE ($= \sqrt{\hat{\delta}^2 + \hat{\rho}^2}$). The number of observations for each combination is shown in the column $n$.

<table>
<thead>
<tr>
<th>Source</th>
<th>Estimate Type</th>
<th>$\mu(\delta)$</th>
<th>$\sigma(\delta)$</th>
<th>$\hat{\delta}$</th>
<th>$\hat{\sigma}(\delta)$ = $\hat{\rho}$</th>
<th>RMSE</th>
<th>$n$</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>DHS</td>
<td>D</td>
<td>0.04</td>
<td>0.48</td>
<td>0.11</td>
<td>0.38</td>
<td>0.40</td>
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<td>2</td>
<td>DHS</td>
<td>C</td>
<td>-0.26</td>
<td>0.51</td>
<td>-0.48</td>
<td>0.46</td>
<td>0.66</td>
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<tr>
<td>3</td>
<td>Census</td>
<td>D</td>
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<td>-0.43</td>
<td>0.50</td>
<td>0.66</td>
</tr>
<tr>
<td>4</td>
<td>Census</td>
<td>C</td>
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<td>0.43</td>
<td>-1.02</td>
<td>0.58</td>
<td>1.17</td>
</tr>
<tr>
<td>5</td>
<td>MICS</td>
<td>D</td>
<td>-1.10</td>
<td>1.03</td>
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<td>0.87</td>
</tr>
<tr>
<td>6</td>
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<td>0.89</td>
<td>1.28</td>
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<tr>
<td>7</td>
<td>MICS</td>
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<td>0.20</td>
<td>1.35</td>
<td>1.36</td>
</tr>
<tr>
<td>8</td>
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<td>0.22</td>
<td>0.56</td>
<td>0.60</td>
</tr>
<tr>
<td>9</td>
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<td>I</td>
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<td>1.37</td>
<td>0.75</td>
<td>1.09</td>
<td>1.32</td>
</tr>
<tr>
<td>10</td>
<td>Survey</td>
<td>D</td>
<td>-0.50</td>
<td>0.58</td>
<td>-0.47</td>
<td>0.42</td>
<td>0.63</td>
</tr>
<tr>
<td>11</td>
<td>Survey</td>
<td>C</td>
<td>-1.18</td>
<td>0.95</td>
<td>-1.06</td>
<td>0.49</td>
<td>1.17</td>
</tr>
<tr>
<td>12</td>
<td>Survey</td>
<td>I</td>
<td>0.14</td>
<td>0.98</td>
<td>0.06</td>
<td>0.95</td>
<td>0.95</td>
</tr>
<tr>
<td>13</td>
<td>Survey-NR</td>
<td>D</td>
<td>-0.40</td>
<td>0.18</td>
<td>-0.60</td>
<td>0.21</td>
<td>0.64</td>
</tr>
<tr>
<td>14</td>
<td>Survey-NR</td>
<td>C</td>
<td>-1.48</td>
<td>0.18</td>
<td>-1.18</td>
<td>0.29</td>
<td>1.22</td>
</tr>
</tbody>
</table>

and measurement error standard deviations against the observed ones; the model fit seems reasonably good.

### 4.2 Estimation of Past and Projection of Future TFR

The fertility transition, or Phase II, started in 1980 in Nigeria, according to the definition of Alkema et al. (2011). We initialize the MCMC algorithm with a warm start, simulating the starting values for the global parameters $\psi$ and the country-level parameters $\theta_c$ from their posterior distribution from the model that does not take account of uncertainty about past TFR values (Alkema et al., 2011; Raftery, Alkema and Gerland, 2014). The true past fertility rates are initialized as the U.N. estimates.

The results are shown in Figure 6. This is based on data up to 2015, and can be compared with Figure 3(c), which is based on data up to 2005. The posterior distribution for
Figure 5: Bias and Variance Estimates: Fitted against Observed. The size of the dots represents the number of observations. Most large dots are along the diagonal line.

the 2000-2005 period is tighter, because more data relevant to this period were available in 2015 than in 2005. The posterior distribution widens slightly for the past period, 2010–2015, again reflecting the relative paucity of data relevant to this period by 2015. One could expect that this posterior distribution will tighten as more data relevant to 2010–2015 become available in the future.

We make projections in two steps. In the first step, we will sample one trajectory from the MCMC results obtained in Section 4.2. Then given the sampled trajectory, the phase of the most recent year is determined by this trajectory, and then future TFR is sampled according to the country-specific parameters of this trajectory. The resulting projection is summarized in Figure 7.

The projections of future TFR from 2015 to 2100, taking account of uncertainty about the past, are shown in Figure 7. The black solid and dotted curves show the U.N.’s 2015 probabilistic projection (not taking account of uncertainty about the past), while the blue line and shaded region shows the projection from our method. Both project that Nigeria’s TFR will likely decline, with a great deal of uncertainty about how fast this will happen. Our proposed method yields a similar predictive median to the current U.N. method, but
somewhat wider prediction intervals. As we saw in the out-of-sample validation study, these wider intervals do incorporate an important additional source of uncertainty, and, on average, take the intervals from undercovering the truth to some extent, to close to nominal coverage.

Figure 7: TFR projections. The red shaded areas are the estimated TFR with 95% estimation intervals, and the blue shaded areas are the projected TFR with 95% prediction interval, where the present is taken to be 2015, marked by a dashed vertical line. The black line and the black dotted lines represent the U.N. WPP 2015 median and 95% predictions.
4.3 Model Validation: Simulation Study

We now run a simulation study with input data on past TFRs chosen to resemble the Nigerian data, to see how accurately the proposed method captures past TFR values. For each simulation, we sampled one TFR trajectory from the posterior distribution of our previous analysis as the true (unobserved) TFR. Then we randomly generated TFR estimates from the normal distribution in Level 1 of the model, by assuming the bias of data points are the previous estimated bias ($\hat{\delta}_{c,t,s}$), and measurement error variances are the previous estimated variances ($\hat{\rho}_{c,s}$). We then treated sampled data points as the input data for the estimation process. We still treated the U.N. estimates as unbiased, as before.

We repeated the simulation process 1000 times. The estimation results of one simulation are shown in Figure 8.

If we take the median of posterior TFR estimates as the point estimate, the mean absolute error (MAE) for all 13 time periods is 0.157. Breaking it down by the 13 time periods, the result are shown in Table 3.

The overall coverage rate of the 80% interval was 85.9%, and the overall coverage rate of the 95% interval is 95.1%. The overall coverage rate was close to the nominal rate, and the MAE was also low. Thus the model gave accurate point and intervals estimates of past values in the simulation study.
<table>
<thead>
<tr>
<th>80% Interval Coverage</th>
<th>95% Interval Coverage</th>
<th>Mean Absolute Error</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.865</td>
<td>0.917</td>
<td>0.259</td>
</tr>
<tr>
<td>0.888</td>
<td>0.976</td>
<td>0.263</td>
</tr>
<tr>
<td>0.895</td>
<td>0.982</td>
<td>0.222</td>
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<tr>
<td>0.794</td>
<td>0.914</td>
<td>0.177</td>
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<tr>
<td>0.847</td>
<td>0.967</td>
<td>0.139</td>
</tr>
<tr>
<td>0.718</td>
<td>0.898</td>
<td>0.152</td>
</tr>
<tr>
<td>0.797</td>
<td>0.933</td>
<td>0.118</td>
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<td>0.926</td>
<td>0.979</td>
<td>0.103</td>
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<tr>
<td>0.936</td>
<td>0.980</td>
<td>0.097</td>
</tr>
<tr>
<td>0.952</td>
<td>0.981</td>
<td>0.116</td>
</tr>
<tr>
<td>0.848</td>
<td>0.940</td>
<td>0.090</td>
</tr>
<tr>
<td>0.893</td>
<td>0.963</td>
<td>0.103</td>
</tr>
<tr>
<td>0.805</td>
<td>0.935</td>
<td>0.198</td>
</tr>
</tbody>
</table>

5 Discussion

Since 2015, the U.N. has been producing probabilistic projections of the total fertility rate as part of their official population projections for all countries of the world, using the Bayesian hierarchical model of Alkema et al. (2011) and Raftery, Alkema and Gerland (2014). However, one important source of uncertainty has not been accounted for so far in these projections, namely uncertainty about past fertility levels. This uncertainty is small for countries with long-standing vital registration systems; this is the case for less than half of the world’s roughly 200 countries. For the other countries, however, this uncertainty can be considerable.

We have developed a new method for projecting the total fertility rate probabilistic for all countries that extends the U.N. method to take account of uncertainty about past TFR values. In a validation experiment, we found that the existing U.N. method leads to prediction intervals whose coverage is somewhat lower than nominal, while for our new method the coverage is close to nominal. For the countries with the highest quality data on past rates, mostly in Europe and North America, our method gives results that are similar to the current method. However, for countries with lower quality data where TFR estimates have been based on surveys for at least part of the past 60 years, our method gives intervals that are noticeably wider than the current ones.

The long-term implications of these results could be far reaching. The countries with the most uncertainty about past TFR values are also largely those with the highest current
fertility levels and the greatest uncertainty about future levels, many of which are in Sub-Saharan Africa. Not surprisingly, therefore, our method indicates that these are also the countries for which the understatement of uncertainty was greatest. Thus our TFR results could lead to a considerable increase in uncertainty about long-term population in these countries, especially as the effects of differences in TFR compound over generations. The population of Sub-Saharan Africa is currently around 1 billion, and current projections are that it will increase to between 3.4 and 4.8 billion in 2100 with 80% probability (Gerland et al., 2014; United Nations, 2017). This interval will be wider still once uncertainty about past TFR has been factored in, with even more dramatic implications for future population levels in Africa, and hence for the world as a whole.

Our method is in two stages. In the first stage we estimate the bias and measurement error variance of the different data sources by country using a classical analysis of variance method. In the second stage we estimate a Bayesian hierarchical model taking the point estimates from the first stage as input. In principle it would be possible to unify these two stages by including the estimation of the bias and variance of the different sources in the Bayesian hierarchical model. However, this would complicate the model considerably, making it harder to specify, code, debug and interpret, and it seems unlikely that it would change the results appreciably. We feel that our modeling decision strikes a reasonable balance between complexity and performance. This is supported by the good assessment of predictive uncertainty provided by our method.

To use these projections of total fertility in population projections, one must convert them to age-specific fertility rates. The U.N. currently does this using the methodology described by Ševčíková et al. (2016). Each simulated future TFR value is converted to a corresponding age-specific fertility pattern, which is used with age-specific mortality and migration rates in the cohort-component projection method to project the corresponding future population by age and sex. A subtle point is that this takes account of uncertainty about future total fertility, but not about future age-specific fertility given total fertility, i.e. about the number but not the timing of future births. Because the age pattern of births is relatively concentrated regardless of their number, this is a much smaller source of uncertainty than uncertainty about the number of births. Nevertheless, it should be addressed in future research.

We have produced results for all the world’s countries with populations over 100,000 as of 2015, except for one: China, the most populous country. We did not include China in our analysis because the estimates for its TFR suffer from a unique form of bias, which would
require a different kind of analysis. This is due to the One Child Policy, introduced in 1979. As a result of this policy, many Chinese families did not report births to the authorities, with the hope of being able to circumvent the policy and have additional children. The underreporting was particularly severe in the late 1990s, and Goodkind (2004) has argued that this was because the 1991 Decree pushed the responsibility of implementing family planning rules, especially the one-child policy, to local governments, giving them a greater incentive to underreport the number of births.

There have been many efforts to correct for this underreporting. For example, Yi (1996); Retherford et al. (2005); Cai (2008) and Merli and Raftery (2000) attempted to correct estimates of TFR in 2000. The clearest evidence of this underreporting comes from primary school enrollments several years later, which were typically substantially larger than the reported number of births during this period. Zhai and Chen (2007) used these enrollment data to correct the TFR estimates for the late 1990s. The U.N. has also been using enrollment data to correct available estimates. Our method would not be sufficient to give good estimates of China’s TFR in the period of severe underreporting. Instead, for China it would be desirable to extend our method to include enrollment data, taking account of uncertainty in the enrollment data in the model. A simpler approach would be to include enrollment-corrected survey and census estimates as inputs to our method, but we felt that a more comprehensive approach was desirable given the great demographic importance of China and the unique data issues it presents, and so we omitted China from the present analysis.

**Acknowledgements**

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References


