Small Area Estimation of Health Insurance Coverage in 2010-2012

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February 3, 2014

1 Introduction

The Small Area Health Insurance Estimates (SAHIE) program at the U.S. Census Bureau produces estimates of numbers and proportions of those with and without health insurance coverage for demographic groups within states and counties. The demographic groups are defined by age, sex, and income, and in addition, for states by race and ethnicity. Income groups are defined in terms of income-to-poverty ratio (IPR), which is the family income divided by the appropriate federal poverty level.

For 2010 through 2012, SAHIE publishes estimates for states for the following domains:

(1) The full cross classification of:
   - Four age categories: 0-64, 18-64, 40-64, 50-64.
   - Four race/ethnicity categories: all races, Hispanic, White not Hispanic, Black not Hispanic.
   - Three sex categories: all sexes, male, female.
   - Five income groups: all incomes, and IPR categories 0-138%, 0-200%, 0-250%, and 0-400%. A sixth IPR category, 138-400%, was added in 2012.

(2) Age category 0-18 in all incomes and in IPR categories 0-138%, 0-200%, 0-250%, 0-400%, and 138-400%(added in 2012).

The domains for which SAHIE produces county estimates are nearly the same as those for states except SAHIE does not produce county estimates by race/ethnicity.
The choice of domains is motivated by the needs of one of SAHIE’s sponsors, the Centers for Disease Control and Prevention (CDC). The CDC has cancer screening programs for which the eligible population is low-income, uninsured women in specified age groups (SAHIE Team 2008). In addition, the age 0-18 low-income categories are relevant to the Children’s Health Insurance Program (CHIP). Because the SAHIE models produce estimates for disjoint groups covering virtually everyone under age 65, we release estimates for men and women as well as children, and for other aggregates of possible interest.

The choice of the income groups 0 to 200 percent IPR and 0 to 250 percent IPR is motivated by the needs of the CDC and CHIP. The income groups 0 to 138 percent IPR, 0 to 400 percent IPR, and 138 to 400 percent IPR are responsive to the needs of recent heath care legislation. The Patient Protection and Affordable Care Act helps families gain access to health care by allowing Medicaid to cover families within income group 0 to 138 percent of the federal poverty level. Also, families with incomes from 138 to 400 percent of the federal poverty level can receive tax credits that will help them pay for health coverage in the new health insurance exchanges.

In the sections to follow, we describe in detail the models used to produce the SAHIE estimates.

2 Overview of SAHIE modeling

2.1 The “base” level

We publish estimates for groups that sometimes overlap or are contained in one or another domain. However, actual modeling is done at a “base” level at which domains are disjoint, and are chosen so that the estimates needed for publication can be obtained as needed by aggregation.

For example for states, we do the actual modeling for the full cross-classification of:

- Five age categories: 0-17, 18, 19-39, 40-49, 50-64.
- Four race/ethnicity categories: White not Hispanic, Black not Hispanic, Hispanic, and not Hispanic.
- Two sex categories: male, female.
- Five income groups: 0-138% IPR, 138-200% IPR, 200-250% IPR, 250-400% IPR and 400% and above IPR.

2.2 Two proportions to estimate

Let the acronym ARSH (age, race, sex, Hispanic origin) denote, for states, age by race/ethnicity by sex, and for counties, age by sex.

For states and counties, we have demographic population estimates for ARSH groups that we treat as unbiased and measured without error. To obtain estimates
of the numbers with and without health insurance for states and counties within ARSH groups, we estimate two sets of proportions. Within each state or county by ARSH group, denoted by \( a \), we estimate the proportions in each of the five income groups, \( p_{ai}^{PR}, i = 1, \ldots, 5 \). In each ARSH by income group \((a, i)\), we estimate the proportion insured, \( p_{ai}^{IC} \). The number in ARSH by income group \((a, i)\) is the product of \( p_{ai}^{PR} \) and the population for ARSH group \( a \). The number insured is the product of \( p_{ai}^{IC} \) and the number in the ARSH/income group.

The SAHIE model consists of two largely distinct parts corresponding to these two proportions. We refer to the two parts of the model as the “income” and “insurance” parts.

### 2.3 Modeling survey data

The SAHIE model is an “area level” model (Rao 2003) in that it uses survey estimates for areas or domains of interest rather than individual responses, and it uses other data that are aggregates rather than for individuals. Each of the two parts of the SAHIE model is similar to a well-known small area-level model, the Fay-Herriot model. The Fay-Herriot model is a hierarchical model in which the variables of interest occupy a “middle” level, between high-level parameters such as regression coefficients, and observed data. Let \( \theta_i, i = 1 \ldots n \) be the variable of interest, and \( \hat{\theta}_i \) be a survey estimate of \( \theta_i \). A simple version of the normal Fay-Herriot model can be written

\[
\hat{\theta}_i = \theta_i + \varepsilon_i \tag{1}
\]

\[
\theta_i = x_i^T \beta + u_i \tag{2}
\]

where \( \varepsilon_i \sim \mathcal{N}(0, v_i^S) \) and \( u_i \sim \mathcal{N}(0, v_i^M) \).

The \( \varepsilon_i \) are sampling errors with sampling variances \( v_i^S \). The \( u_i \) can be viewed as model errors, or as area-specific random effects, with model variance \( v_i^M \). All the \( \varepsilon_i \)'s are independent of all the \( u_i \)'s. The equation in (1) is referred to as the “sampling model” and the equation in (2) is the “linking model.”

In a frequentist context, assuming the sampling and model variances are known, it can be shown that the best linear unbiased predictor (BLUP) of each \( \theta_i \) is a weighted average of the survey estimate and the regression prediction

\[
\hat{\theta}_i^{BLUP} = \gamma_i x_i^T \hat{\beta} + (1 - \gamma_i) \hat{\theta}_i \tag{3}
\]

with \( \gamma_i = \frac{v_i^S}{v_i^S + v_i^M} \). \( \hat{\beta} \) is the usual weighted least squares estimate from the regression of the \( \hat{\theta}_i \)'s on the \( x_i \)'s. Note that \( \gamma_i \) becomes closer to one as the model variance becomes smaller relative to the sampling variance, so that the BLUP estimate of \( \theta_i \) is primarily the regression prediction estimate \( x_i^T \hat{\beta} \). Complementarily, \( \gamma_i \) becomes closer to 0 when
the model variance becomes large relative to the sampling variance, so that the
BLUP estimate is primarily the survey estimate $\hat{\theta}_i$.

A similar result holds in the Bayesian context. Conditional on the $\beta$’s and
variances, the mean of each $\theta_i$ is a weighted average of $\hat{\theta}_i$ and a regression prediction $x_i^T \beta$, with the weight on $x_i^T \beta$ being $\frac{v_i^S}{v_i^S + v_i^M}$.

In addition to the fact that the SAHIE model contains two parts, each of which
is similar to a Fay-Herriot model, there are several differences between the SAHIE
model and a standard Fay-Herriot model:

- In both the income and insurance parts of the model, we model a survey
  estimate $\hat{p}$ of the proportion $p$, but we assume that the logit of $p$ satisfies
  a normal linear model. The sampling model and the linking model are not
  “matched.”

- The survey estimates of the proportions in the income groups within an ARSH
  group are not independent. There are five income groups within each ARSH
  group. The survey estimates for those five must add to one. We model four
  of them, and assume the correlations correspond to those of a multinomial
  distribution.

- In the insurance part of the model, we do not assume that the logits of the pro-
  portions are independent. We instead assume that they have a block diagonal
  variance matrix with identical blocks.

- In the insurance part of the model, we do not assume that the survey estimates
  $\hat{p}$ are normally distributed, but instead that they follow a mixture of discrete
  and continuous distributions.

- We model as random some auxiliary data rather than treat them as fixed
  predictors in a regression.

In later sections, we give details of the model, including details of the items above.
In the next section, we give a fuller discussion of the last item above.

2.4 Modeling auxiliary data

One large difference between the SAHIE model and the standard Fay-Herriot model
in (1) and (2) is in how some non-survey data are used. In small-area models such
as the Fay-Herriot model, “auxiliary” (i.e, non-survey) data are typically used as
covariates to help predict the variables of interest. In the standard Fay-Herriot
model, these covariates occur as fixed predictors as in the $x_i$ in (2).

Fisher and Gee (Fisher 2003 and Fisher and Gee 2004) proposed an alternative
in the context of estimating poverty. In their research for the Small Area Income
and Poverty Estimates program, they developed an alternative to the usual Fay-Herriot model, that they refer to as an “error-in-variables” model. In their model, they treat the covariates as measures of the quantity of interest, $\theta$ (log poverty in their example), that are possibly biased and have random error. Let $i$ index the observations and $j = 1, \ldots, p$ index the $p$ auxiliary data. Their model is

\[
\hat{\theta}_i = \theta_i + e_i
\]

\[
A_{ij} = b_j + c_j \theta_i + u_{ij} \quad j = 1, \ldots, p
\]

\[
\theta_i \sim N(\mu_i, v_i^\theta)
\]

where the $e_i$ and $u_{ij}$ are mean zero error terms that are normal with variances that possibly depend on parameters. In this approach, the auxiliary data $A_{ij}$ are treated in a way very similar to survey estimates. The primary difference is that they are not unbiased measures of $\theta_i$, and thus the model includes the unknown parameters $b_j$ and $c_j$.

A feature of the model in (5) - (7) is that the influence of $\hat{\theta}_i$ and the $A_{ij}$ on the estimate of $\theta_i$ can vary observation to observation, depending on the relative sizes of the variances. This is an extension of the property noted in (3) and (4) in which the influences of the survey estimate and a regression prediction vary depending on the relative magnitude of their variances.

The approach of Fisher and Gee was extended to small area estimates of insurance coverage in Fisher, O’Hara, and Riesz (2006). The SAHIE model includes both fixed predictors $x_i$ of the quantities of interest and auxiliary data to be modeled, $A_i = (A_{i1}, \ldots, A_{ip})^T$.

The $A_i$ are possibly nonlinear regressions of the $\theta_i$, and the $\theta_i$ are modeled by a generalized linear model. The two parts, income and insurance, of the SAHIE model each have the form

\[
\hat{\theta}_i = \theta_i + e_i
\]

\[
A_{ij} = h_j(\theta_i) + u_{ij} \quad j = 1, \ldots, p
\]

\[
g(\theta_i) = x_i^T \beta + v_i
\]

where the $e_i, u_{ij}$, and $v_i$ are error terms terms with mean zero and variances that depend on parameters, and that are independent except for exceptions noted later.

## 3 The primary data

We use the following primary data sources for states and counties.

**Single-year ACS direct estimates.** We have two sets of direct estimates from the American Community Survey:
• Estimates of the number in each of the five IPR categories by ARSH (age by race/ethnicity by sex categories for states, and by age and sex categories for counties).
• Estimates of the proportions insured in ARSH by income categories.

5-year ACS direct estimates. We use ACS 5-year direct estimates spanning the five years prior to the release year of the number in geography/ARSH/IPR categories.

Federal Tax Returns data. We use the number of Internal Revenue Service (IRS) exemptions in age by IPR categories in each state and county. The age categories are 0-18 and 19-64. We do not have actual ages for the IRS data. We use the number of child exemptions as a proxy for age 0-18.

Supplementary Nutrition Assistance Program data. For each state and county, we use counts of the number of people participating in the Supplemental Nutrition Assistance Program (SNAP, formerly Food Stamps) from the United States Department of Agriculture.

Medicaid/CHIP participation data. We use Medicaid participation records from the Centers for Medicare and Medicaid Services (CMS). States submit their data to the CMS quarterly. Individuals are in the file if Medicaid covered them for at least one day during the quarter. We have Children’s Health Insurance Program (CHIP) participation counts from states and counties gathered from a web page of the Centers for Medicare and Medicaid Services (CMS). We combine the Medicaid and CHIP participation data, and use the combined data for each state and county in cross-classifications of age by sex. Beginning in 2012, we use Medicaid/CHIP data from the year prior to the release year.

Demographic population estimates. We use demographic estimates of the resident population from the U.S. Census Bureau’s Population Estimates Program. These estimates are published for the nation, states, and counties by age, sex, race, and Hispanic origin.

County Business Patterns. For counties, we use the proportions of adults employed by firms with 19 employees or less and firms with 100 employees or more.

Census 2010. For counties, we use the Census 2010 proportions of housing units which are rural and housing units which are resident-owned.

See http://www.census.gov/did/www/sahie/methods/inputs/index.html for more information about these data sources.
4 Model details

In this section, we describe in detail the components of the SAHIE model for states. There are some differences in the modeling of counties that we describe later. We use the following notation:

- “ACS estimate” refers to the single year ACS estimate unless it is specified to be the ACS 5-year estimate.
- ARSH (age, race, sex, Hispanic origin) refers to either an age by race by sex (for states) or an age by sex (for counties) category.
- $a$ indexes state or county by ARSH category.
- $i$ indexes IPR category.
- $S_j$ denotes the sample size (number of persons for which data were collected) for the $j^{th}$ category.
- $POP$ denotes a demographic population estimate.
- $N$ denotes a number of people. $N^{IPR}_{ai}$ denotes the number of people in the $a^{th}$ state or county by ARSH and $i^{th}$ IPR category, and $N^{IC}_{ai}$ denotes the number of people with health insurance coverage in the $a^{th}$ state or county by ARSH and $i^{th}$ IPR category. $N^{UI}_{ai} = N^{IPR}_{ai} - N^{IC}_{ai}$ is the number uninsured.
- $p^{IPR}_{ai} \equiv N^{IPR}_{ai}/POP_a$ is the proportion among those in the $a^{th}$ state or county by ARSH group who are in the $i^{th}$ IPR category.
- $p^{IC}_{ai} \equiv N^{IC}_{ai}/N^{IPR}_{ai}$ is the proportion among those in the state or county by ARSH by IPR category $(a, i)$ who have health insurance coverage.
- $\beta$ denotes regression parameters that appear in a model for the proportion in an IPR category or a model for the proportion insured.
- $\alpha$ denotes a mean parameter, i.e., a parameter that appears in a model for the mean of the ACS 5-year or administrative record data.
- $\lambda$ denotes a variance parameter, i.e., a parameter that appears in the model for the sampling variance of the ACS estimates, or in a model for the variance of the ACS 5-year or administrative record data.
- Hatted variables such as $\hat{p}^{IPR}_{ai}$ denote direct survey estimates.
- Overlines such as $\overline{MED}$ denote means.

The parameters $\alpha$ and $\lambda$ typically depend on one or more of the age, race/ethnicity, sex, or IPR categories. We suppress indices that show these dependencies.

The income part of the model allows us to estimate $p^{IPR}_{ai}$, the proportion of people in IPR category $i$, within state or county by ARSH category $a$. The number of people within the state or county by ARSH by income group is given by $N^{IPR}_{ai} = p^{IPR}_{ai}POP_a$. 
The insurance part of the model allows us to estimate $p_{ai}^{IC}$, the proportion insured within state or county by ARSH by IPR category $(a, i)$. We combine these to estimate the primary quantities of interest, $N_{ai}^{IC}$ and $N_{ai}^{UI}$, the number insured and the number uninsured, where

\[
N_{ai}^{IC} = p_{ai}^{IC} N_{ai}^{IPR} \\
N_{ai}^{UI} = (1 - p_{ai}^{IC}) N_{ai}^{IPR}.
\]

### 4.1 The income part of the state model

#### 4.1.1 Modeling ACS estimates of proportions in income groups

In the first part of the income model, we model $\hat{p}_{ai}^{IPR}$, the ACS estimate of the proportion in IPR category $i$ within state by ARSH category $a$. We assume that these ACS estimates are unbiased and normally distributed. Note that for any $a$, $\sum_{i=1}^{5} \hat{p}_{ai}^{IPR} = 1$. For this reason, we model four of the five IPR categories, and do not treat those four as independent. We assume a parametric model for the variances, and assume the correlations correspond to those of a multinomial distribution. The model for the ACS estimates of proportions in IPR categories is as follows:

\[
(\hat{p}_{a1}^{IPR}, \ldots, \hat{p}_{a4}^{IPR}) \mid p^{IPR}, \lambda \sim \mathcal{N}(\left(p_{a1}^{IPR}, \ldots, p_{a4}^{IPR}\right), \Sigma^{IPR})
\]

\[
(\Sigma^{IPR})_{ii} = \frac{\lambda_0 p_{ai}^{IPR} (1 - p_{ai}^{IPR})}{S_{ai}^{\lambda_1}}, \quad i = 1, \ldots, 4
\]

\[
(\Sigma^{IPR})_{ij} = \rho_{aij} \sqrt{(\Sigma^{IPR})_{ii} (\Sigma^{IPR})_{jj}}
\]

where $\rho_{aij} = -\sqrt{\frac{p_{ai}^{IPR} p_{aj}^{IPR}}{(1 - p_{ai}^{IPR})(1 - p_{aj}^{IPR})}}$, $i \neq j$

where the parameters $\lambda_0$ and $\lambda_1$ vary by age. When the sample size is 1, $(\Sigma^{IPR})_{ii}$ is set to $p_{ai}^{IPR}(1 - p_{ai}^{IPR})$.

#### 4.1.2 The regression part of the income model

We assume that $p_{ai}^{IPR}$, the proportion of those in state by ARSH group $a$ who are in IPR group $i$, follows a five-category logistic model with normal errors. Let $X_{ai}^{IPR}$ be a matrix of fixed predictors with rows $(x_{ai}^{IPR})^T$. Then the following is the model
for $p_{ai}^{IPR}$:

$$p_{ai}^{IPR} = \frac{\exp(\mu_{ai}^{IPR})}{\sum_{i=1}^{5} \exp(\mu_{ai}^{IPR})}$$

$$\mu_{ai}^{IPR} \beta^{IPR}, v^{M,IPR} \overset{\text{indep}}{\sim} \mathcal{N}\left((x_{ai}^{IPR})^T \beta^{IPR}, v^{M,IPR}\right)$$

with $(x_{ai}^{IPR})^T = 0$ for all $a$.

The model variance $v^{M,IPR}$ is the same for all $a$ and $i$.

The predictors in $X^{IPR}$ for states are as follows:

- Main effects for IPR.
- Two-way interactions between age and IPR, between race/ethnicity and IPR, and between sex and IPR.
- Two-way interactions between IPR and the following continuous variables:
  - The logit of the proportion who are Hispanic in the state (from demographic population estimates).
  - The median log of income relative to poverty level, as measured by tax records.
  - The variance of log of income relative to poverty level, as measured by tax records.
- Three-way interactions among age, race/ethnicity, and IPR.
- Three-way interactions among age, sex, and IPR.
- Three-way interactions between age, IPR, and the previously mentioned continuous variables.

4.1.3 Modeling state ACS 5-year estimates, IRS exemptions, and SNAP counts

We model the means of the ACS 5-year estimates, the IRS exemptions, and the SNAP counts as functions of $N_{ai}^{IPR}$, the number of people in IPR category $i$, within state by ARSH, $a$.

4.1.4 Modeling ACS 5-year estimates for states

We model the ACS 5-year estimates, $ACS5_{ai}^{IPR}$, of the number of people in state by ARSH by IPR categories. We assume these estimates have means, $\overline{ACS5}_{ai}$, that are linear functions of the $N_{ai}^{IPR}$, and are conditionally independent. Then

$$ACS5_{ai}^{IPR} \overset{\text{indep}}{\sim} \mathcal{N}(\overline{ACS5}_{ai}, v_{ai}),$$

where

$$\overline{ACS5}_{ai} = (\alpha_0 + \alpha_1) N_{ai}^{IPR}$$

$$v_{ai} = \lambda_0 \overline{ACS5}_{ai} \lambda_1.$$
The $\alpha$’s and $\lambda$’s are parameters to be estimated. The parameter $\alpha_0$ varies by age and race/ethnicity, while $\alpha_1$ varies by age and IPR. The variance parameter $\lambda_0$ varies by age and IPR, and $\lambda_1$ does not vary by category.

4.1.5 Modeling IRS exemptions for states

From the IRS, we have the number of IRS exemptions by state by two approximate age categories (0-18, 19-64) by IPR categories. The age categories are approximate because the number that we use for the age 0-18 category is actually the number of child exemptions.

We assume that the numbers of exemptions are normally distributed with a mean that is a linear function of aggregate $N_{ai}^{IPR}$’s, and are conditionally independent. Let $\text{TAX}_{ti}$ be the number of IRS exemptions in state by age category $t$ and IPR category $i$. Then

$$\text{TAX}_{ti} \mid N_{ai}^{IPR}, \alpha, \lambda \quad \text{indep} \quad \mathcal{N} (\text{TAX}_{ti}, v_{ti})$$

$$\text{TAX}_{ti} = \alpha N_{ti}^{IPR}$$

$$v_{ti} = \lambda_0 \text{TAX}_{ti}^{\lambda_1}$$

where $N_{ti}^{IPR}$ is the number of people in state by age by IPR category $ti$. $N_{ti}^{IPR}$ is obtained by summing $N_{ai}^{IPR}$ over the appropriate age, race/ethnicity, and sex categories. The parameters $\alpha$ and $\lambda_0$ vary by the two ages by IPR, and $\lambda_1$ does not vary by category.

4.1.6 Modeling SNAP participation for states

$\text{SNAP}_s$ is the number of SNAP participants by state. We model the mean, $\overline{\text{SNAP}}_s$, as a linear function of the number of people in the state in the 0-138 percent IPR category. We assume that the $\text{SNAP}_s$’s are normally distributed and conditionally independent. Let $s$ index state. Then

$$\text{SNAP}_s \mid N_{i1}^{IPR}, \alpha, \lambda \quad \text{indep} \quad \mathcal{N} (\text{SNAP}_s, v_s)$$

$$\overline{\text{SNAP}}_s = \alpha N_{s1}^{IPR}$$

$$v_s = \lambda_0 \overline{\text{SNAP}}_s^{\lambda_1}.$$ 

Here $N_{s1}^{IPR}$ is the number of people in a state in the 0 - 138% IPR category. The parameters $\alpha$, $\lambda_0$, and $\lambda_1$ do not vary by category.

4.2 The insurance part of the state model

In the insurance part of the model, we model ACS estimates of $p_{ai}^{IC}$, the proportion insured in the state by ARSH by IPR category, and the combined Medicaid/CHIP data. From this part of the model, we obtain estimates of $p_{ai}^{IC}$, which enables us to
estimate our primary quantities of interest, $N_{ai}^{IC}$ and $N_{ai}^{UI}$, the number insured and the number uninsured in state by age by race/ethnicity by sex by IPR category $ai$, by $N_{ai}^{IC} = p_{ai}^{IC}N_{ai}^{IPR}$ and $N_{ai}^{UI} = (1 - p_{ai}^{IC})N_{ai}^{IPR}$.

### 4.2.1 Modeling the ACS estimates of the proportion insured

Proportions insured are often close to one. ACS estimates of proportions insured are often one, sometimes zero, and are bounded between zero and one. Rather than assume normality, we model the ACS estimates of proportions insured in a way to capture that they are bounded and have positive probability mass at zero and one. We use the term “three-part model” for the model we use, following Pfeffermann et al. (2008) who use the term “two-part model” to refer to a similar model.

We model the probability that $\hat{p}_{ai}^{IC}$ is one, the probability that $\hat{p}_{ai}^{IC}$ is zero, and conditional on $0 < \hat{p}_{ai}^{IC} < 1$, we assume that $\hat{p}_{ai}^{IC}$ follows a beta distribution. Let $p_{ai}^{(0)}$ and $p_{ai}^{(1)}$ be the probabilities that $\hat{p}_{ai}^{IC}$ is zero and one. The model is

$$\hat{p}_{ai}^{IC} | p_{ai}^{IC}, \lambda, \zeta \begin{cases} = 0 & \text{with probability } p_{ai}^{(0)} \\ = 1 & \text{with probability } p_{ai}^{(1)} \\ \sim \text{Beta}(a_{ai}, b_{ai}) & \text{with probability } 1 - p_{ai}^{(0)} - p_{ai}^{(1)} \end{cases}$$

with

$$\text{var}(\hat{p}_{ai}^{IC}) = \lambda_{0}p_{ai}^{IC}(1 - p_{ai}^{IC}) \frac{S_{ai}^{\lambda_{1}}}{r_{ai}}$$

(9)

$$p_{ai}^{(0)} = (1 - p_{ai}^{IC})^{1+\zeta_{0}(S_{ai}-1)}$$

(10)

$$p_{ai}^{(1)} = (p_{ai}^{IC})^{1+\zeta_{1}(S_{ai}-1)}$$

(11)

where Beta denotes the beta distribution. When sample size is 1, we set (9) to $\text{var}(\hat{p}_{ai}^{IC}) = p_{ai}^{IC}(1 - p_{ai}^{IC})$. The parameters $\lambda_{0}$ and $\lambda_{1}$ vary by age and IPR, as do the parameters $\zeta_{0}$ and $\zeta_{1}$. Note that the parameters, $a_{ai}$ and $b_{ai}$, of the beta distribution in (8) are functions of $p_{ai}^{IC}$, $p_{ai}^{(0)}$, $p_{ai}^{(1)}$ and var(\hat{p}_{ai}^{IC})$. We chose the functions for the variance and the probabilities of zero and one in (9) - (11) by starting with what the variances and probabilities of zero and one would be under simple random sampling, and introducing parameters to accommodate the effects of non-independence due to the sample design and correlated responses. Bauder and Szepelka (2011) considered various groups of observations and compared within groups the predicted and actual frequencies of survey estimates of zero and one. They found close agreement, confirming the choice of functions in (10) and (11).

### 4.2.2 The regression part of the insurance model

The model for the proportions insured is logistic-normal with a multivariate error structure. Let $\mu_{a}^{IC} = (\mu_{a1}^{IC}, \ldots, \mu_{a5}^{IC})^T$, and let $X^{(a)}$ be the matrix made up of the five
rows \((x_{a1}^{IC})^T, \ldots, (x_{a5}^{IC})^T\), and \(X^{IC}\) the data matrix obtained by stacking the \(X^{(a)}\)’s.

\[
p_{ai}^{IC} = \logit^{-1}(\mu_{ai}^{IC})
\]

\[
\mu_{ai}^{IC} \mid \beta, \Sigma^{IC} \overset{indep}{\sim} \mathcal{N}(X^{(a)}\beta, \Sigma^{IC})
\]

where \(\Sigma^{IC}\) is a 5 \(\times\) 5 matrix whose elements are estimated. Define age2 to take two values: one for age groups 0-17 and 18, the other for any of the other age groups: 19-38, 40-49, 59-64. The predictors in \(X^{IC}\) for states are as follows:

- Main effects for age, race/ethnicity, sex and IPR.
- All two-way interactions among age, race/ethnicity, sex and IPR.
- The two-way interaction between state and age2.
- Three-way interactions among:
  - Age, race, and IPR.
  - Age, sex, and IPR.
  - Race, sex, and IPR.
- The state median log of income relative to poverty level, as measured by tax data, interacted with IPR and with age2 by IPR.
- The variance within a state of log of income relative to poverty level, as measured by tax data, interacted with IPR and age2 by IPR.

### 4.2.3 Modeling Medicaid/CHIP enrollees

Let \(MED_m\) be the number of people enrolled in Medicaid or CHIP in a state by age by sex category, denoted \(m\). We assume that the mean, \(\overline{MED}_m\), is a function of the number insured in 0-250 percent IPR. We assume that the Medicaid counts \(MED_m\) are independent, conditional on all \(N_{IC}^{m1}\) and parameters. We have

\[
MED_m \mid \gamma, \alpha, \lambda \overset{indep}{\sim} \mathcal{N}(\overline{MED}_m, v_m)
\]

\[
\overline{MED}_m = \gamma_{sk} \alpha N_{m1}^{IC}
\]

\[
\gamma_{sk} \mid \delta \sim \text{Gamma (mean = 1, var = } \delta)
\]

\[
v_m = \lambda_0 \overline{MED}^{\lambda_1}
\]

where \(s\) is the state and \(k\) is the age2 category (as defined in section 4.2.2) of the \(m^{th}\) observation. \(N_{m1}^{IC}\) is obtained by summing \(N_1^{IC}\) over the race/ethnicity categories and the IPR categories 0-138 percent, 138-200 percent, and 200-250 percent. The parameter \(\alpha\) varies by age by sex, \(\lambda_0\) varies by age, and \(\lambda_1\) does not vary by category. The \(\gamma_{sk}\)’s are state by age2 random effects with variance \(\delta\), and are independent given \(\delta\). The \(\gamma_{sk}\)’s are multiplicative, rather than additive, effects to ensure that the coefficients of \(N_{m1}^{IC}\) are always positive, while still allowing the possibility that the \(\gamma_{sk}\)’s reduce the coefficient on \(N_{m1}^{IC}\).
5 The county model

For counties, the models for the ACS estimates of proportions in IPR categories and of proportions insured are like those in sections 4.1.1 and 4.2.1. The model for Medicaid/CHIP participation is like that in 4.2.3.

The regressions in the income and insurance parts of the model for counties have different predictors than for states.

5.1 Predictors for county IPR and IC regressions

The predictor matrix $X^{IPR}$ for counties (as in Section 4.1.2 for states) includes the following:

- Main effects for IPR.
- Two-way interactions between age and IPR, and between sex and IPR.
- The three-way interaction among age, sex, IPR.
- Log county population interacted with IPR, and with age by IPR (the coefficients can differ for small and large counties).
- Logit of the proportion Hispanic based on demographic estimates, interacted with IPR and with age by IPR.
- County median log IPR, interacted with IPR and with age by IPR.
- Variance of log IPR, interacted with IPR and with age by IPR.
- State, interacted with three IPR categories (0-200%, 200-400%, and 400%+).

As in Section 4.2.2, age2 takes two values: one for age groups 0-17 and 18, the other for any of the other age groups: 19-38, 40-49, 59-64. The predictor matrix $X^{IC}$ for counties (as in Section 4.2.2 for states) includes the following:

- IPR, age and sex categories and all their two- and three-way interactions.
- The three-way interaction among state, age2, and two IPR categories (0-200%, 200%+).
- Each of the following county level variables, and its interactions with age, IPR, and age2 by IPR:
  - Log population, from Demographic Population Estimates.
  - Median log of income relative to poverty level.
  - Variance of log of income relative to poverty level.
  - Logit of the ACS 5-year estimate of the proportion of adults with less than high school education.
- Each of the following county level variables, and its interaction with age2, IPR, and age2 by IPR:
  - Logit of the ACS 5-year estimate of the proportion who are non-citizens.
- Logit of the proportion who are American Indian/Alaskan Native, from Demographic Population Estimates.
- Logit of the proportion of adults who are employed by firms of size 19 or less, from County Business Patterns.
- Logit of the proportion of adults who are employed by firms of size 100 or more, from County Business Patterns.
- Logit of the proportion of housing units that are rural, from Census 2010.
- Logit of the proportion of housing units that are resident-owned, from Census 2010.

5.2 Modeling county ACS 5-year estimates, IRS exemptions, and SNAP counts

As with states, we model the means of the ACS 5-year estimates, the IRS exemptions, and the SNAP counts as functions of the $N_{ai}^{IPR}$’s, summed to the appropriate level. However, there are notable differences in how we model these data for counties.

5.2.1 Modeling the ACS 5-year estimates for counties

For counties, we model the ACS 5-year estimates, $ACS_{5ai}$, of numbers in county by ARSH group $a$ and IPR category $i$ as follows:

$$ACS_{5ai} | \alpha, \lambda \sim T(\nu, \text{mean} = \bar{ACS}_{5ai}, \text{var} = v_{ai})$$

$$\bar{ACS}_{5ai} = \alpha_0 N_{ai}^{IPR}$$

$$v_{ai} = \lambda_0 \bar{ACS}_{5ai}^{\lambda_1}$$

where $ACS_{5ai}$ is the ACS 5-year estimate and $T$ is the t-distribution, parameterized in terms of the degree of freedom parameter, $\nu$, and the mean and variance. We use a t-distribution because when we fit the model assuming normality, some residuals were too large to be consistent with the normality assumption. The parameter $\nu$ does not vary by category, $\alpha_0$ and $\lambda_0$ vary by age and IPR, and $\lambda_1$ varies by age.

5.2.2 Modeling IRS exemptions for counties

Let $t$ index county by the two tax age categories. For counties, we have

$$TAX_{ti} | \nu, \alpha, \lambda \sim T(\nu, \text{mean} = \bar{TAX}_{ti}, \text{var} = v_{ti})$$

$$\bar{TAX}_{ti} = \alpha_0 N_{ti}^{IPR}$$

$$v_{ti} = \lambda_0 \bar{TAX}_{ti}^{\lambda_1}$$

where $TAX_{ti}$ is the number of IRS exemptions in county by age category $t$ and IPR category $i$, and $T$ is the t-distribution, parameterized in terms of the degree of
freedom parameter, $\nu$, and the mean and variance. $N_{ti}^{IPR}$ is obtained by summing $N_{ai}^{IPR}$ over the appropriate age and sex categories. As with the ACS 5-year estimates, we use a $t$-distribution because when we fit the model assuming normality, some residuals were too large to be consistent with the normality assumption. We did not observe this with states. The parameters $\alpha_0$, $\lambda_0$ and $\nu$ vary by the two age by IPR categories. $\lambda_1$ does not vary by category.

5.2.3 Modeling SNAP participation for counties

Let $c$ index county. For SNAP data, we have

\[
SNAP_c \mid N^{IPR}, \alpha, \lambda \sim \mathcal{N}(SNAP_c, v_c)
\]

\[
SNAP_c = \alpha_0 (N_{cl}^{IPR})^{\alpha_1}
\]

\[
v_c = \lambda_0 SNAP_c^{\lambda_1}.
\]

Here, $N_{cl}^{IPR}$ is the number of people in a county in the 0-138% IPR category. The parameters $\alpha_0$, $\alpha_1$, $\lambda_0$, and $\lambda_1$ do not vary by category.

5.3 Prior distributions

For the Bayesian modeling, we generally use vague priors for the high level parameters. For the regression coefficients $\beta^{IPR}$ and $\beta^{IC}$, we use the (improper) uniform prior over the real numbers of appropriate dimension. For degrees of freedom parameters and multiplicative parameters in functions for means, we use truncated normal distributions with large variances. For multiplicative variance parameters ($\lambda_0$ in most cases above) we use the (improper) prior $\frac{1}{\sqrt{\lambda_0}}$. For parameters that are exponents in variance functions, we use a uniform prior on $(0, 3)$. For $\Sigma^{IC}$ in Section (4.2.2), we use a Wishart prior on $(\Sigma^{IC})^{-1}$. The prior mean is the previous year’s estimate of $(\Sigma^{IC})^{-1}$. The prior degrees of freedom is 6, which is one more than the dimension of $\Sigma^{IC}$.

6 Model selection

We made many modeling decisions to arrive at the current SAHIE models. In addition to the overall form of the model, these decisions include choices of predictors, mean and variance functions, and distributions. We describe some of the criteria we used in the next sections.
6.1 Model diagnostics

6.1.1 Standardized residuals

Some choices of mean, variance, and density functions resulted from perceived lack of fit based on diagnostics we use. Our primary model diagnostic is a certain type of standardized residual. For the survey estimates, ACS 5-year, and administrative data that we model, we predict means and variances so that for any data, \( y \), that we model, we can obtain a form of standardized residual and squared residual

\[
E_{\theta|data} \left[ \frac{y - E(y|\theta)}{\sqrt{\text{var}(y|\theta)}} \right] \quad \text{and} \quad E_{\theta|data} \left[ \frac{(y - E(y|\theta))^2}{\text{var}(y|\theta)} \right]
\]

from the Markov chain Monte Carlo (MCMC) output used to fit the model. See Chib and Greenberg (1995) for an explanation of MCMC. If the model is correct and \( y \) is normally distributed, this standardized residual is distributed as approximately normal(0,1). The standardized squared residuals should have a mean of approximately one. We check that averages of these residuals over large groups of observations are close to zero, and check for extremely small or large values. We look at plots against various quantities such as the predicted mean, population, predicted variance, and where appropriate, sample size. We also look at boxplots of standardized residuals for different values of categorical variables such as age and IPR, and against quantiles of population. We check that the averages over large groups of squared standardized residuals are reasonably close to one.

6.1.2 Posterior predictive p-values

Another model diagnostic that we use is the posterior predictive p-value (PPP-value) (Gelman, Meng, and Stern (1996)). A posterior predictive p-value is a measure of how surprising or improbable some function of the data (and possibly parameters) is, under the posterior predictive distribution of that data. Let \( y \) represent all of the data and \( \theta \) represent all of the parameters. A PPP-value is defined as

\[
P_{y^{rep},\theta|y}(T(y^{rep}, \theta) \geq T(y, \theta)) \quad \text{for some function} \quad T \quad \text{where the probability is with respect to} \quad p(y^{rep} | \theta)p(\theta|y) \quad \text{the joint distribution of a replication of the data,} \quad y^{rep}, \quad \text{and} \quad \theta, \quad \text{conditional on} \quad y.
\]

Let \( y_i \) represent a single data point. We use the functions \( T_1(y, \theta) = y_i \) and \( T_2(y) = (y_i - E(y_i|\theta))^2 \). Thus, the PPP-value corresponding to \( T_1 \) is \( P_{y^{rep},\theta|y}(y_i^{rep} \geq y_i) \). We refer to this PPP-value as the PPP-value for the mean because many values near 0 or near 1 suggest that means given by the model are generally too low, or too high, respectively. We refer to the PPP-value corresponding to \( T_2 \) as the PPP-value for the variance since it measures the surprise in the squared distance between the data and its mean. We compute PPP-values for each of the data sources in the model. We look at plots of PPP-values against various quantities, such as population, posterior means, posterior variances, and sample sizes. Our approach is to use the PPP-values informally to check for evidence of
model failure. Many values near zero or near one would suggest problems with the model.

6.2 Selecting predictors for the regression parts of the income and insurance models

In order to select predictors for the income and insurance parts of the model, we generally consider the posterior means and variances of the regression coefficients. We form an approximate 95 percent credible interval for the regression coefficient by taking its posterior mean plus or minus two times its posterior standard deviation. Generally speaking, we include a predictor in the model if the approximate 95 percent credible interval does not include zero.

7 Benchmarking

We benchmark SAHIE estimates of the numbers insured and uninsured in order to make them consistent with a set of national ACS estimates, and to make county estimates consistent with state estimates. We benchmark state estimates to a relatively small set of national direct estimates of numbers insured and uninsured. We benchmark all possible county estimates to the corresponding state estimates. The benchmarking procedure for counties is a simple proportional adjustment. The procedure for states is more complex.

7.1 State to national benchmarking.

We benchmark the state estimates to ACS national estimates of insured and uninsured for the following categories:

- IPR 0-200%.
- Age 0-18, IPR 0-200%.
- Age 19-64.
- Hispanic.
- Not Hispanic.
- White not Hispanic.
- Black not Hispanic.

7.1.1 Methodology for state to national benchmarking

The benchmarking procedure we use was developed by Luery (1986) in the context of controlling survey weights to control totals. The procedure is as follows. Let \( B \) be the number of benchmarks (here, 14), and let \( \hat{N} = (\hat{N}_1, \hat{N}_2, \ldots, \hat{N}_B)' \), be the benchmarks. Let \( A \) be the number of small area, or model, estimates, and let
\( \hat{Y} = (\hat{Y}_1, \hat{Y}_2, \ldots, \hat{Y}_A)' \) be those estimates. We want to adjust the model estimates so that their sums over states equal the benchmarks. Let \( b \) index the benchmarks, let \( i \) index the area (here, state by ARSH by IPR by insured/uninsured). Let \( X = (x_{ib}) \) be the \( A \times B \) matrix such that \( x_{ib} = 1 \) when area \( i \) contributes to benchmark \( b \), and 0 otherwise. Then the adjusted estimates \( \hat{Y}_i^* \) meet the constraints when \( \sum_{i=1}^{S} x_{ib} \hat{Y}_i^* = \hat{N}_b \) for all \( b \).

We want a set of benchmarked estimates that are, in some sense, optimal. Generally, benchmarked estimates are preferable when they are close to the original estimates. We choose to minimize the relative quadratic loss function

\[
\sum_{i=1}^{S} (\hat{Y}_i^* - \hat{Y}_i)^2 / \hat{Y}_i.
\]

That is, we minimize the squared change from the original to the benchmarked estimate, relative to the size of the original estimate. It can be shown that there exists a unique set of \( \hat{Y}_i^* \) that sum to the benchmarks and minimize (13). This optimal set of benchmarked estimates, \( \hat{Y}^* = (\hat{Y}_1^*, \hat{Y}_2^*, \ldots, \hat{Y}_A^*)' \) is given by

\[
\hat{Y}^* = \hat{Y} + D(\hat{Y})XP(\hat{N} - X^T\hat{Y})
\]

where \( D(\hat{Y}) \) is a diagonal matrix with the entries of \( \hat{Y} \) along the diagonal and \( P = [X^TD(\hat{Y})X]^{-1} \).

For the \( i^{th} \) area, this can be written as

\[
\hat{Y}_i^* = \hat{Y}_i \left( 1 + \sum_{b=1}^{B} f_b x_{ib} \right)
\]

where the \( f_b \) are the \( B \) factors given by \( F = (f_b) = P(\hat{N} - X^T\hat{Y}) \). Thus, the choice of the relative quadratic loss function ensures that if two areas \( i \) and \( i' \) have the same indicators, that is, if \( x_{ib} = x_{i'b} \) for all \( b \), then they receive the same proportional change to their estimates, as given in (15).

7.1.2 Variance of state benchmarked estimates

We estimate the models using MCMC methods in which a procedure for generating values from the posterior distribution of all unknown variables is repeated for many iterations. We can obtain an estimate of the variance of the benchmarked estimates by repeating the benchmarking procedure at each iteration of the MCMC, using each time a newly generated set of unbenchmarked estimates. However, the benchmarking totals themselves are estimates, and have some uncertainty. If we treat them as fixed in the benchmarking procedure, we will likely underestimate the uncertainty in the benchmarked estimates.

We address this issue as follows. We have variance estimates for the ACS national estimates we benchmark to. These estimates are large, so they should be close
to normally distributed. We approximate the distribution of the benchmark by assuming that it has a posterior distribution that is normal, with mean at the estimate, with the estimated variance. Then, in each iteration of the MCMC, we draw a value for each benchmark total from this approximate distribution. We then do the benchmarking procedure, controlling to these generated totals. In this way, the variability in the benchmarked estimates will come from both the variability of the unbenchmarked estimates and the variability of the benchmark totals, as it should.

7.2 Methodology for county to state benchmarking

We benchmark county estimates so that in each state, the county estimates for insured and uninsured in each age by sex by IPR group sum to the benchmarked state estimates. For each cross-classification of age, sex, and income, we apply an adjustment factor to the county estimates of the number insured and the number uninsured so that the sum of the county estimates equals the state estimate. Let $c$ index counties, $i$ index age by sex categories, $i$ index income categories, and $s$ index states. The adjusted estimate of the numbers insured and uninsured are given by

$$
\hat{N}^{IC,\text{adjusted}}_{cji} = \hat{N}^{IC}_{sji} \frac{\hat{N}^{IC}_{cji}}{\sum_c \hat{N}^{IC}_{cji}}
$$

$$
\hat{N}^{UI,\text{adjusted}}_{cji} = \hat{N}^{UI}_{sji} \frac{\hat{N}^{UI}_{cji}}{\sum_c \hat{N}^{UI}_{cji}}
$$

where $\hat{N}^{IC}_{sji}$ and $\hat{N}^{UI}_{sji}$ are state estimates of the insured and uninsured for age by sex by income categories, and the sums are over the counties, $c$, in state $s$.

For variance estimation, in order to take into account the fact that the state estimates have error, we perform the adjustment procedure in each iteration of the MCMC similar to that for state to national benchmarking. In each iteration of the MCMC, we simulate the state control from a normal distribution whose mean is the state estimate and whose variance is the variance of the state estimate.
References


Methods, 325-330.

