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Compensating for Provider Nonresponse Using Response Propensities to Form Adjustment Cells: The National Immunization Survey

Data From the National
Health Care Survey

U.S. DEPARTMENT OF HEALTH AND HUMAN SERVICES
Centers for Disease Control and Prevention
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Contents

Abstract	1
1. Introduction	1
2. Background and Outline of Report	2
3. Summary of NIS Sample Design	3
4. Original NIS Estimation Methodology	3
4.1 RDD Phase Sampling Weights	3
4.2 Nonresponse Adjustment for Second Phase of Sampling—Original Method Used Before 1998	4
5. Adjustment for Provider Vaccination History Nonresponse Using Adjustment Cells	5
5.1 Model Development	5
5.2 A Sequential Method for Choosing “Optimal” Number of Adjustment Cells	6
5.3 Raking to Control Bias	7
5.4 Estimators for Vaccination Coverage of a Subpopulation and Its Variance	7
6. Illustration and Evaluation of the Methods	8
6.1 Illustration of Sequential Method to Determine “Optimal” Number of Adjustment Cells and to Evaluate Provider-Verified Vaccination History Nonresponse Bias	8
6.2 Evaluation of Difference in Bias Reduction Obtained Using “Optimal” Choice for Number of Adjustment Cells, Compared With Using Five Adjustment Cells Within Each Stratum	8
6.3 Comparison of 4:3:1:3 Coverage Estimates Obtained Using the New NIS Methodology	8
7. Summary	8
References	9

Figure

Difference in State-specific coverage rates using new and old estimation methods: National Immunization Survey, 1998	5
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Detailed Tables

1. Percent of children with selected characteristics by provider data status: National Immunization Survey, 1998	11
2. Variables in the model selection for the response propensity model: National Immunization Survey, 1998	11
3. Results of the stepwise variable selection procedure for the response propensity model	12
4. Estimated log-odds and associated standard errors of the predictors selected in the stepwise procedure for the final model: National Immunization Survey, 1998	13
5. Variables used for raking the response propensity adjusted weights of children with provider data: National Immunization Survey, 1998	14
6. Illustration of method for determining optimal number of adjustment cells for Illinois-Rest of State: National Immunization Survey, 1998	14
7. Summary of nonresponse-adjusted estimates of 4:3:1:3 coverage and estimated bias reduction obtained by using optimal number of adjustment cells: National Immunization Survey, 1998	15
8. Statistical significance of the estimated difference in nonresponse-adjusted estimates of 4:3:1:3 coverage obtained by using optimal choice of k compared with using five adjustment cells in each IAP area: National Immunization Survey, 1998	17

Objectives

The National Immunization Survey (NIS) uses two phases of data collection to obtain vaccination information from a sample of young children: a random-digit-dialing (RDD) survey for identifying households with children 19–35 months of age, followed by a mail survey for obtaining provider-reported vaccination histories about these children. Vaccination histories are used to estimate vaccination coverage rates.

In 1998, provider-reported vaccination histories were not obtained for 32.9% of children with a completed RDD interview. This report describes the statistical methods adopted in 1998 to reduce the bias in vaccination coverage estimates that could result from “vaccination history nonresponse,” that is, differences between children for whom provider data was obtained and those for whom it was not obtained.

Methods

In the methods adopted in 1998, children with completed NIS RDD interviews are grouped into adjustment cells defined by their propensity to have adequate provider data. Sampling weights of children with adequate provider data are divided by the cell-specific weighted response rate to allow these children to represent all children in the cell.

Results

Using an “optimal” number of cells, the overall extent of bias reduction was 0.5%, suggesting that provider nonresponse bias was small. Authoritative literature suggests using five cells. No statistically significant differences were observed in vaccination coverage estimates when comparing estimates based on the “optimal” number of cells with five cells. Thus, five adjustment cells are used to reduce provider nonresponse bias in the NIS vaccination coverage estimates. No substantively important differences were observed between estimates based on the methodology used prior to 1998 and the methodology adopted in 1998.

Keywords: *Adjustment cell • National Immunization Survey • nonresponse bias • response propensity.*

Compensating for Provider Nonresponse Using Response Propensities to Form Adjustment Cells: The National Immunization Survey

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1. Introduction

One goal of the Childhood Immunization Initiative (1) is to achieve an immunization or vaccination coverage rate of 90% for four critical vaccines—diphtheria and tetanus toxoids and pertussis (DTP), poliovirus (polio), measles-mumps-rubella (MMR), and *Haemophilus Influenzae* type B (Hib).

The National Immunization Survey (NIS) was initiated in April 1994 to monitor progress toward achieving the vaccination coverage goals; to monitor national, State, and local vaccination coverage rates in the United States on an ongoing basis; and to assist the Centers for Disease Control (CDC) in directing resources. The NIS covers 78 Immunization Action Plan (IAP) areas that include the 50 States and 28 urban areas, including the District of Columbia. Each IAP area represents a stratum of the sampling design within which households are independently sampled. Within each IAP area, the design of the NIS includes two phases of data collection: a list-assisted random-digit-dialing (RDD) telephone survey to eligible households followed by a mail survey of vaccination providers of eligible sampled children. The mail survey is administered only to

those providers for whom verbal consent is obtained from the responding parent or guardian.

In the RDD sampling phase of the NIS, the most knowledgeable respondent in each sampled household with eligible children is asked to provide vaccination history information for all the eligible children in the household. Also, demographic and socioeconomic characteristics of the household are obtained.

When the NIS began in 1994, the intention was to estimate vaccination coverage rates using household reported vaccination histories. In late 1994, preliminary results from the 1994 National Immunization Provider Record Check Study (2) revealed a low correlation between household-reported vaccination histories and vaccination histories reported by children’s vaccination providers. The provider histories were thought to be more reliable, particularly for households in which no written vaccination histories were kept. As a result, in late 1994 the NIS began to ask RDD respondents for verbal consent to contact eligible children’s vaccination providers to obtain provider-reported vaccination histories. This practice still continues.

If verbal consent is obtained, the vaccination providers are mailed a questionnaire from which vaccination histories are obtained. Ezzati-Rice et

al. (3) and Zell et al. (4) give a more detailed description about the survey of vaccination providers. Because consent is not obtained from the parents/guardians of all age-eligible children and not all providers respond or provide adequate vaccination information, potential nonresponse bias in the survey estimates could be incurred if coverage estimates were prepared only from children with adequate provider data and without adjusting for differences between these children and children for whom inadequate or no vaccination provider histories were obtained.

This report describes the statistical methodology adopted in 1998 to adjust NIS sampling weights to compensate for failure to obtain adequate provider data from all children for whom completed random-digit-dialing (RDD) interviews are obtained. Estimates of the extent of the bias that would be incurred by not compensating for nonresponse are also included. Lastly, comparisons are made of the new methods that have been used since 1998 with the methods that were used before 1998 to obtain estimates of vaccination coverage.

2. Background and Outline of the Report

In 1998, 83.4% of respondents in RDD sampled households gave consent to NIS personnel to contact their children's vaccination providers. Among households giving consent and reporting provider name and address information, immunization history questionnaires were obtained for 95.0% of the vaccination providers receiving the NIS mail survey. Overall, adequate vaccination histories were obtained for 67.1% of age-eligible children with completed RDD interviews.

Data obtained from responding providers include information on whether sampled children received a sufficient number of doses to be considered up-to-date with vaccination recommendations for DTP, polio, MMR, and Hib. A child is said to be "4:3:1:3" up-to-date provided he/she has received four or more doses of DTP, three or

more doses of polio, one or more doses of MMR, and three or more doses of Hib.

Empirical results suggest that sampled children for whom vaccination histories are obtained have characteristics that are associated with a greater likelihood of being up-to-date for the 4:3:1:3 vaccination series, as compared with sampled children for whom vaccination histories are not obtained. Table 1 lists characteristics that distinguish these groups of children for the 1998 NIS. All differences in these tables are statistically significant at the $p=0.05$ level for each variable listed. Children whose providers give adequate information are more likely to live in households that have higher total incomes, more likely to have a white non-Hispanic mother, and more likely to live outside of a central city of a metropolitan statistical area. These factors are also known to be associated with higher vaccination rates. Also, children with adequate vaccination information are more likely to live in the same State as the one in which they were born and more likely to come from a household where the respondent could locate a written record (i.e., a shot card) of the child's vaccination history.

As a consequence, estimates of vaccination coverage that are not adjusted for these differences may be too high and may mislead immunization program managers about the success of their efforts to increase vaccination coverage rates. This report describes the statistical methods used in the NIS to adjust vaccination coverage rates for this potential "vaccination history nonresponse" bias.

The sampling design of the NIS is described in Section 3, and a synopsis of methods used to obtain estimates of vaccination coverage rates prior to 1998 is provided in Section 4. This has been described by Zell et al. (5). The method was based on adjusting for vaccination history nonresponse using a variant of the stratified two-phase estimation method described by Cochran (6). Section 4 includes a description of how first phase sampling weights are obtained and gives a synopsis of methods used to obtain estimates of vaccination coverage before 1998. It

also gives reasons why there was interest and opportunity to improve the original estimation methodology.

Section 5 describes a methodology that adjusts the first phase sampling weights to obtain coverage estimates with reduced "vaccination history nonresponse" bias. This method is based on grouping sampled children into adjustment cells according to the similarity of their response propensities to have a provider-reported vaccination history. The first phase sampling weight for children with adequate provider data is divided by the cell-specific weighted response rate. There have been previous discussions and applications of response propensities to adjust for nonresponse. In an econometric application, David et al. (7) describe the use of adjustment cells based on the use of response propensities. Little (8) describes how using response propensities to form adjustment cells can reduce nonresponse bias. Also, Yansaneh and Eltinge (9) and Eltinge and Yansaneh (10) described methods for determining the "optimal" number of adjustment cells that should be used in each stratum using response propensities. This section also describes the methods of Eltinge and Yansaneh in detail. It also describes the ratio estimators used within the context of the adjustment cell methodology to obtain estimates of vaccination coverage and their estimated standard errors.

In Section 6, these methods are applied to data from the 1998 NIS. Specifically, Section 6.1 includes demonstrations of the extent to which unadjusted estimates may be biased for the NIS coverage rate and how these rates' bias may be reduced using an "optimal" choice for the number of adjustment cells within each stratum. In Section 6.2, differences are evaluated in the bias reduction obtained using the "optimal" choice for the number of adjustment cells compared with using five adjustment cells within each stratum. Section 6.3 includes comparisons of the estimates obtained using five adjustment cells within each stratum with the original NIS methodology. The report concludes with a discussion of the findings and recommendations.

3. Summary of the NIS Sample Design

The NIS uses two phases of data collection to obtain vaccination information for a large national probability sample of young children. In the first phase of data collection, a random-digit-dialing (RDD) sample of telephone numbers is drawn. The NIS RDD design uses independent samples of telephone numbers in each of the 78 IAP areas for each calendar quarter. The quarterly samples make it possible to combine four consecutive quarters to form four-quarter estimates of vaccination coverage. The RDD sample in each IAP area is used to screen for households with one or more children age 19–35 months. In 1998, the target number of completed interviews for a quarter was 110 per IAP area. With an average sample size of 440 interviews for a four-quarter period, the total number of interviews for a four-quarter period is around 34,300.

The main goals of the sample design are: a) to ensure that a probability sample of telephone numbers is selected for each IAP area, b) to ensure that the desired target number of interviews is achieved in each IAP area, c) to minimize in a cost-effective manner the number of age-eligible children excluded from the sample frame, and d) to maintain an up-to-date sampling frame of telephone numbers.

To accomplish these goals, the NIS uses the list-assisted method of random-digit-dialing (11). This method is used to select a random sample of telephone numbers from banks of 100 consecutive telephone numbers (e.g., 617-495-0000 to 617-495-0099) in an IAP area that contain one or more directory-listed residential telephone numbers. When the IAP area is a city, county, or combination of counties, some prefix areas (e.g., 781-347) may cover part of the IAP area and part of an adjacent IAP area. In such situations, the NIS applies a “majority” rule that specifies that if at least 50% of the directory-listed households in a prefix area fall inside the IAP area, the prefix area is assigned to that IAP area. The

sampling frame of prefix areas is updated each quarter to account for newly created prefix areas.

The selection of a quarterly probability sample of telephone numbers to meet the target number of completed interviews with age-eligible children in an IAP area presents some challenges because the 78 IAP areas differ considerably along several dimensions. For example, they differ on a) the percent of telephone numbers that are working residential numbers, b) the likelihood of contacting a person among residential numbers, c) the percent of contacted households willing to complete the screener interview, d) the percent of households with an age-eligible child, and e) the willingness of parents to complete the telephone interview. These factors lead to considerable variation in the total sample size of telephone numbers needed to achieve a target sample size of completed interviews per IAP area.

Three design tools are used to address these issues. First, an automated procedure is used to eliminate a portion of the nonworking and business telephone numbers in the sample before it is dialed by the interviewers (12). Second, a statistical model to predict the number of sample telephone numbers needed in each IAP area for a given quarter of interviewing has been developed. Third, after drawing the required sample size of telephone numbers for an IAP area, that sample is divided into random subsamples called replicates. By administering the sample release on a replicate-by-replicate basis, the total number of interviews obtained is controlled and the interviews for each IAP area are spread evenly across the entire calendar quarter.

In the second phase of data collection, each child interviewed in the NIS is eligible for inclusion in the provider record-check survey. Permission to contact vaccination providers is requested from the respondent. The information collected from a child’s vaccination providers is used to determine whether a child is up-to-date on his or her vaccinations. Written requests for vaccination histories are mailed to the providers to obtain reports of vaccinations from medical

records. Providers have the option of responding via mail or facsimile. Postcard reminders and telephone followup are used to encourage nonresponding providers to participate in the study.

4. Original NIS Estimation Methodology

4.1 RDD Phase Sampling Weights

For children ages 19–35 months sampled in the 1998 random-digit-dialing (RDD) phase of the NIS (N = 32,511 completed interviews), RDD sampling weights were developed for each child for whom an interview was completed. These weights account for:

- the probability of selection of a telephone number within an IAP area,
- unit nonresponse arising from sample telephone numbers with an unresolved residential status,
- unit nonresponse arising from residential sample telephone numbers where the presence of age-eligible children is not determined,
- unit nonresponse attributable to incomplete and partial household interviews among eligible households, and
- multiple telephone numbers within eligible households.

To ensure that the final sampling weights for each child when aggregated represent all children 19–35 months (i.e., those in both telephone and nontelephone households), an adjustment for households without telephones was made. In particular, using data from the National Health Interview Survey (NHIS), the RDD phase sampling weights have been adjusted to account for expected differences in vaccination coverage levels of eligible children living in households with and without telephones.

Frankel et al. (13), Hoaglin and Battaglia (14), Battaglia et al. (15) give the details of this specific adjustment.

A poststratification ratio adjustment was also carried out within each IAP area using race/ethnicity of the mother, educational status of the mother, and age in months of the eligible child. The initial population controls were obtained from the NCHS natality file. However, special adjustments to the natality file population counts were made within each IAP area. The special adjustments account for a) infant mortality, b) foreign births, and c) geographic mobility among IAP areas. Further details on the original NIS RDD weighting methodology can be found in Smith et al. (16), Ezzati-Rice et al. (3), and Zell et al. (4).

In this report, these weights are referred to as the RDD phase sampling weights.

4.2 Nonresponse Adjustment for Second Phase of Sampling—Original Method Used Before 1998

In surveys like the NIS in which there are two phases of sampling and there is nonresponse at the second phase of sampling, it is necessary to adjust for any potential bias introduced by nonresponse at the second stage. Prior to 1998, a two-phase sampling estimation procedure described by Cochran (6) was used. This estimation methodology was adapted from that used in NIPRCS, the study that demonstrated that household reporting of vaccination status introduced an unacceptable level of response error. Initially there were no plans by CDC to release an NIS public use data file that would have required a single weight for each child with provider data. Over time, however, plans for release of NIS public use data files were made, and a different estimation strategy was needed because calculating single weights for children with provider data create a complication for the two-phase estimation procedure. A description of how the NIS estimation methodology was implemented to obtain vaccination coverage rates before 1998

is in this section.

Within an IAP area, each eligible child sampled was categorized into one of five strata depending upon whether the household respondent reported that the child was up-to-date on the 4:3:1:3 series and whether a shot card was used during the household interview. These were:

Stratum 1: household respondent had a shot card during the RDD interview and reported the eligible child as being 4:3:1:3 up-to-date,

Stratum 2: household respondent had a shot card during the RDD interview and reported the eligible child as not being 4:3:1:3 up-to-date,

Stratum 3: household respondent did not have a shot card during the RDD interview, but reported from memory/recall the eligible child as being 4:3:1:3 up-to-date,

Stratum 4: household respondent did not have a shot card during the RDD interview and reported from memory/recall the eligible child as not being 4:3:1:3 up-to-date, and

Stratum 5: missing information on 4:3:1:3 up-to-date status and/or shot card use.

Children belonging to stratum 1 were said to belong to the “top row.” These children were 4:3:1:3 up-to-date according to their shot card written record. Approximately 30% of the NIS children with completed household interviews belonged to the top row in 1998. Special treatment of top-row children is described below.

Let $\hat{p}_{1,ij}$ denote the estimated weighted proportion of children in IAP area i who are found to belong to stratum j , $j = 1, \dots, 5$, according to data obtained in the first phase of data collection. Also, let $\hat{p}_{2,ij}$ denote the estimated weighted proportion of children in IAP area i belonging to stratum j that are determined to be 4:3:1:3 up-to-date according to data obtained from the provider-verified vaccination history information in the second phase of data collection. Then, the stratified two-phase sampling estimator of children who are 4:3:1:3 up-to-date in IAP area i is

$$\hat{\pi}_i = \sum_{j=1}^5 \hat{p}_{1,ij} \hat{p}_{2,ij} \quad (1)$$

However, the NIS implementation of the two-phase sampling estimation methodology involved a special modification two-phase sampling estimator (1). Specifically, weighted data from the entire national sample of all children belonging to stratum 1 (i.e., the top row) whose vaccination providers responded were used to estimate $p_{2,ij}$ for every IAP area. This variant of two-phase sampling estimator was referred to as the “provider-adjusted top-row” estimator and was used in NIS before 1998. It was used to account for IAP areas in which a substantial proportion of top-row children were not 4:3:1:3 up-to-date according to provider data. This was thought to be caused primarily by incomplete provider records of vaccinations received by some children. A further description of this NIS estimation methodology used before 1998 can be found in Zell et al. (4).

One disadvantage of the provider-adjusted top-row estimator is that it tends to produce a slightly biased estimate of IAP area vaccination coverage rates. By using national data to estimate $p_{2,ij}$, coverage level estimates for IAP areas will tend to be biased toward the national figure.

An additional disadvantage of the two-phase sampling approach to account for provider-reported vaccination history nonresponse is that there are few methods tailored for statistical analyses of data from two-phase sampling designs when the analysis calls for more complicated models. For example, in epidemiologic analyses of complex survey data, logistic regression is routinely required. However, there are no standard statistical methods for logistic regression that have been developed and made accessible to analysts through commonly available commercial software that account for a two-phase sampling design. Therefore, although weighted estimates reflecting both phases of data collection can be produced, limitations exist for more complex epidemiologic analysis and for release of data on public use files.

Because of these disadvantages, a new enhanced statistical estimation methodology was implemented in 1998. The new estimation methodology adopted in 1998 is described in Section 5.

5. Adjustment for Provider Vaccination History Nonresponse Using Adjustment Cells

To adjust coverage estimates for “provider vaccination history nonresponse bias,” adjustment cells can be formed using response propensities obtained from a logistic regression model with a binary response variable indicating whether or not a child has adequate provider data.

Within each IAP area, sampled children can be grouped into adjustment cells according to the similarity of their response propensities to have a provider-reported vaccination history. A group of children who have similar response propensities will also be similar with respect to variables that are strongly associated with the probability of having a provider-reported vaccination history. In this important respect, children within each adjustment cell are comparable. Because of this, all sampled children in the cell may be represented by the sampled children within the cell who have adequate provider-reported immunization histories. In particular, by dividing the RDD phase sampling weights of children with adequate provider data by the cell’s weighted response rate, these children’s weights are adjusted to represent all of the children belonging to the cell. By doing this, the bias in estimated vaccination coverage rates attributable to differences between sampled children who have and do not have provider-reported immunization histories can be reduced. Within each adjustment cell, children without adequate provider-reported vaccination data are represented by children who are similar to them with respect to their response propensities and other associated variables.

Section 5.1 describes the variables that were found to be associated with their response propensities. Section 6 provides results that show how large the reduction in bias can be by using these

variables to estimate their response propensities.

5.1 Model Development

As a first step in forming the response propensity adjustment cells, a national response propensity model was developed using logistic regression. The response propensity is the probability that a sampled child has a provider-reported vaccination history.

Within each IAP area, the RDD phase sampling weights were first rescaled to add to the IAP area specific sample size. This was done by multiplying each sampling weight by the unweighted sample size within the IAP area and dividing by the sum of the weights within the IAP area. Inclusion of these rescaled weights as prior weights in the logistic regression ensures that as the sample size increases to the finite population size, the estimated logistic coefficients converge to the true finite population values. This property holds provided the logistic model and its linear predictors give a correct depiction for the target population of why provider vaccination histories are missing.

As candidates for predictors to the response propensity model, variables were used that have been found to be associated with immunization status in other research conducted by CDC (17) using the NIS. The variables used as candidates for the model are described in table 2. Forward stepwise logistic regression was used to select predictors among these candidates. S-PLUS 2000 (18) was used for all calculations.

At each step of the stepwise selection process, the logistic regression model examined all possible and allowable interactions between predictors. Also, at each step after adding regressors to the model, the model selection method reexamined each regressor included in the model to determine whether any predictor that entered at a previous step of the process could be dropped. Akaike’s (18) Information Criterion (AIC) statistic was used as the criterion by which the optimal set of candidate regressors was chosen or retained at each step of the process.

Results of the stepwise selection process applied to all 32,511 children for which completed RDD interviews were obtained in 1998 are given in table 3. Akaike’s statistic provides a measure of goodness of fit of a model, corrected for the number of parameters used in the model. Within this framework, a model with minimum AIC provides the best fit of the model to the data. Deviance provides a related measure of goodness of fit measured on the $-2 \times \log$ likelihood scale. The asymptotic distribution of the difference of deviances for two nested models is known to have a chi-square distribution with degrees of freedom equal to the difference in the number of parameters indexing the two models. Table 3 lists the chi-square values and degrees of freedom associated with adding a predictor at each step of the stepwise model selection process. McCullagh and Nelder (20) provide a further discussion on the use of the deviance as a way of evaluating the statistical significance of predictors in a model.

Using a reference cell parameterization in which the first level of each variable in table 2 is the variable’s reference level, table 4 lists the estimated odds ratios, estimated log-odds, and associated standard errors of the predictors selected in the stepwise procedure. Twenty-nine predictors were selected in the stepwise procedure. Examining the estimated odds ratios in table 4, children were found to be less likely to have adequate provider data if the respondent does not know or indicates that the child is not up-to-date on his or her vaccinations. Children for whom a shot card was not used during the interview are less likely to have adequate provider data. If the respondent was not the mother, the child is also less likely to have adequate provider data. With respect to socioeconomic and demographic factors, children below poverty are less likely to have adequate provider data, and children with a non-Hispanic white mother are more likely to have adequate provider data. If the mother was living in a different State at the time the child was born, the child is

less likely to have provider data, and children in households with more than one child are more likely to have adequate provider data.

Using the final national model obtained from the variable selection process, predicted response propensities were obtained for each sampled child within each IAP area, adjustment cells were formed, and the boundaries were defined by quantiles of the distribution of the predicted response propensities of children belonging to the IAP area. Within each IAP area, each sample child can be assigned to one of the IAP area-specific adjustment cells. The following subsection describes how an “optimal” choice for the number of adjustment cells can be determined.

5.2 A Sequential Method for Choosing the “Optimal” Number of Adjustment Cells

Yansaneh and Eltinge (9) and Eltinge and Yansaneh (10) have described methods for choosing the number of adjustment cells for the purposes of bias reduction for the case in which cells are constructed using response propensities. For the NIS data, this method proceeds in a sequence of stages within each IAP area.

At stage k , $k=2, \dots$, the estimated vaccination coverage rate \hat{Y}_k obtained by using k adjustment cells is compared with the estimated rate \hat{Y}_{k-1} obtained using $k-1$ adjustment cells. The estimated incremental bias reduction obtained by increasing the number of adjustment cells from $k-1$ to k is

$$\hat{\Delta}_k = \hat{Y}_k - \hat{Y}_{k-1}$$

and the estimated standard error of the estimated bias is

$$\hat{\sigma}(\hat{\Delta}_k) = [\hat{V}(\hat{Y}_k) + \hat{V}(\hat{Y}_{k-1}) - 2\hat{C}(\hat{Y}_k, \hat{Y}_{k-1})]^{1/2}$$

This estimated standard error is calculated using a Taylor series linearization to the estimated variance (21).

Let \hat{Y}_1 denote the estimated vaccination coverage rate that uses only

one adjustment cell. This estimate is obtained by dividing the first phase sample weights of children with adequate provider data by the overall weighted response rate. By using only one adjustment cell, no attempt is made to group children together that are comparable for achieving bias reduction. As a result, the estimate \hat{Y}_1 is referred to as the “unadjusted” estimate.

As discussed previously, estimates of vaccination coverage that are not adjusted for provider vaccination history may be too high. Because of this, there is particular interest in estimates \hat{Y}_k that yield negative values of $\hat{\Delta}_k$ that measures the extent to which the estimate corrects the positive bias by using one more adjustment cell. Because of this, one should compute 1-sided p -values

$$p_k = \Phi \left(\frac{\hat{\Delta}_k}{\hat{\sigma}(\hat{\Delta}_k)} \right)$$

to evaluate the significance of the incremental bias correction for each value of k . Here, $\Phi(\cdot)$ denotes the standard normal cumulative distribution function. The “best” value of k is defined as the smallest value k' such that $p_l < 0.05$ $l = 2, \dots, k'$ and $p_{k'+1} > 0.05$. One method of choosing the number of adjustment cells in each stratum is to find the “optimal” value of k for each stratum.

In addition to evaluating the incremental reduction in bias from stage to stage, one can use the incremental change in mean squared error (MSE) to evaluate the trade-off between bias and variance. Specifically, letting \hat{V}_n denote the estimated variance of the coverage estimate at stage n ,

$$\hat{Y}_n = \frac{\hat{V}_n}{\hat{V}_{n-1} + (\hat{Y}_n - \hat{Y}_{n-1})^2}$$

denotes the relative mean squared error using n adjustment cells compared to using $n-1$ adjustment cells.

5.3 Raking to Control Bias

As discussed previously, by dividing the RDD phase sampling weights of children who had

provider-reported immunization histories by their adjustment cell specific weighted response rate, these children should fairly represent all of the children in the cell as a whole. However, the revised weights may not match poststratification totals used to construct the RDD phase sampling weights. Also, the revised weights may not match the RDD phase sample weighted totals of variables that are known to be important predictors of being up-to-date.

To reduce bias attributable to these differences and to maintain the nonresponse bias adjustment, within each stratum, the revised weights are raked to match the poststratification totals and totals of a few other variables that are known to be associated with being up-to-date. Also, the adjustment cell specific RDD phase sampling weight totals are raked to maintain the effect of the nonresponse adjustment. Table 5 lists the poststratification and other variables used for raking.

5.4 Estimators for Vaccination Coverage of a Subpopulation and Its Variance

This section describes the estimators used for a ratio and its estimated variance for a subdomain of the target population when the sampling design corresponds to a stratified 1-stage cluster sampling design as is used in the NIS (i.e., households are sampled and all age-eligible children are selected).

Among households with eligible children with adequate provider data, let

N_h = the number of primary sampling units (PSUs, or households in the case of the NIS) in stratum h , $h=1, \dots, L$

n_h = the number of PSUs sampled in stratum h , $h=1, \dots, L$

M_{hi} = the number of children in PSU i of stratum h belonging to the target population

m_{hi} = the number of eligible children sampled in the i th PSU in stratum h

W_{hij} = the provider nonresponse adjusted sampling weight for the j th sampled child with adequate provider data sampled in the i th sampled PSU of stratum h

Y_{hij} = 0 when child j with adequate provider data in PSU i in stratum h is not up-to-date on a specific vaccination, and $Y_{hij}=1$ when the child is up-to-date

δ_{hij} = 1 when subject j with adequate provider data in PSU i in stratum h belongs to the domain of interest, and $\delta_{hij}=0$ otherwise

Y'_{hij} = 0 when the j th sampled child with adequate provider data in the i th sampled PSU in stratum h is not up-to-date on a specific vaccination, and $Y'_{hij}=1$ when the child is up-to-date

δ'_{hij} = 1 when the j th sampled child with adequate provider data in the i th sampled PSU in stratum h belongs to the domain of interest, and $\delta'_{hij}=0$, otherwise

Letting $Y_h = \sum_{i=1}^{N_h} \sum_{j=1}^{M_{hi}} \delta_{hij} Y_{hij}$,

and $T_h = \sum_{i=1}^{N_h} \sum_{j=1}^{M_{hi}} \delta_{hij}$,

the true but unknown vaccination coverage rate is

$$\theta = \frac{\sum_{h=1}^L Y_h}{\sum_{h=1}^L T_h}$$

Let $\hat{Y}_h = \sum_{i=1}^{n_h} \sum_{j=1}^{m_{hi}} \delta'_{hij} W_{hij} Y'_{hij}$

and $\hat{T}_h = \sum_{i=1}^{n_h} \sum_{j=1}^{m_{hi}} \delta'_{hij} W_{hij}$.

Then the combined ratio estimator of the vaccination coverage rate for the domain of interest is

$$\hat{\theta} = \frac{\sum_{h=1}^L \hat{Y}_h}{\sum_{h=1}^L \hat{T}_h} \tag{2}$$

To obtain an estimator of the variance of equation 2, a Taylor series approximation (21) is used. Letting

$$Z_{hij} = \frac{\delta'_{hij} W_{hij} (Y_{hij} - \hat{\theta})}{\hat{T}_h}$$

denote the linearized value of equation 2 and

letting $Z_{hi} = \sum_{j=1}^{m_{hi}} Z_{hij}$, and $\bar{Z}_h = \frac{\sum_{i=1}^{n_h} Z_{hi}}{n_h}$

the Taylor series approximation for the estimator of the variance of $\hat{\theta}$ is

$$\hat{V}_T(\hat{\theta}) = \sum_{h=1}^L \frac{n_h}{n_h-1} \sum_{i=1}^{n_h} (Z_{hi} - \bar{Z}_h)^2$$

6. Illustration and Evaluation of the Methods

6.1 Illustration of Sequential Method to Determine “Optimal” Number of Adjustment Cells and to Evaluate Provider-Verified Vaccination History Nonresponse Bias

Table 6 gives an example of the strategy of selecting the “optimal” value of k using the sequential method described in Section 5.3 for the 4:3:1:3 series for one IAP area—Illinois-Rest of State (that is, Illinois except for the city of Chicago). This table shows that a statistically significant reduction in bias can be accrued by using two adjustment cells and that no further significant reduction in bias is obtained by using

three or more adjustment cells. Overall, in going from no adjustment to two adjustment cells, bias is reduced by 0.5%. Also, table 6 shows the incremental change in relative efficiency, $\hat{\gamma}_k$. This table shows that the the relative efficiency decreases modestly by 3% by using two adjustment cells.

Table 7 summarizes the “optimal” choice of the k for each IAP area. This table shows that only two or three adjustment cells formed using response propensities are required when the “optimal” value of $k > 1$. Also, this table shows that the “optimal” value of k is equal to 1 for 57 of the 78 IAP areas. In 1998, 27% of all the children 19–35 months of age lived in the IAP areas in which more than one adjustment cell was designated as the “optimal” choice of k . Also, for the 1998 NIS, for IAP areas in which more than one adjustment cell is specified as the “optimal” value, the decrease in bias from the unadjusted value is on the order of 0.5%. This indicates that the extent to which bias can be reduced using adjustment cells formed using response propensities is small.

6.2 Evaluation of Difference in Bias Reduction Obtained Using “Optimal” Choice for Number of Adjustment Cells, Compared With Using Five Adjustment Cells Within Each Stratum

Cochran (22) published an influential paper describing the use of stratification to control bias in observational studies. Empirical results of his investigation indicated that approximately 90% of the bias is reduced by using five strata. As a result of this empirical finding, a “lore” has been developed among statistical practitioners that indicates that five is a good choice for the number of adjustment cells that should be used to reduce bias. Also, work

conducted by Rosenbaum and Rubin (23) has been effective in showing the advantages of constructing these strata using response propensities.

In table 8, results are given showing the difference in bias reduction $\hat{Y}_k - \hat{Y}_5$ and statistical significance p obtained by using the “optimal” value of k compared to using five strata formed using response propensities. Results in this table show that for the 1998 NIS data, there is little difference in the bias reduction obtained using the “optimal” value of k using the sequential method described in Section 5.3, compared with using five adjustment cells formed using response propensities. However, estimates formed using the “optimal” value of k tend to be a little more efficient as can be seen by the relative efficiencies $\hat{\gamma}_k$ listed in table 8.

6.3 Comparison of 4:3:1:3 Coverage Estimates Obtained Using the New NIS Methodology

The figure shows the differences between the provider-reported vaccination history nonresponse adjusted estimates (new NIS methodology) of coverage for 1998 minus the provider-adjusted top-row estimates (original NIS methodology) for the 50 States and the District of Columbia. This graph shows that across States, the difference tends to be positive, with a mean difference of 0.5 percent in 1998 (range = -4.1% +3.0%). This means that, on average, the provider-adjusted top row estimates tended to slightly underestimate 4:3:1:3 coverage. In general, differences in State level vaccination coverage rates estimated using the original NIS estimation methodology and the new NIS estimation methodology were small.

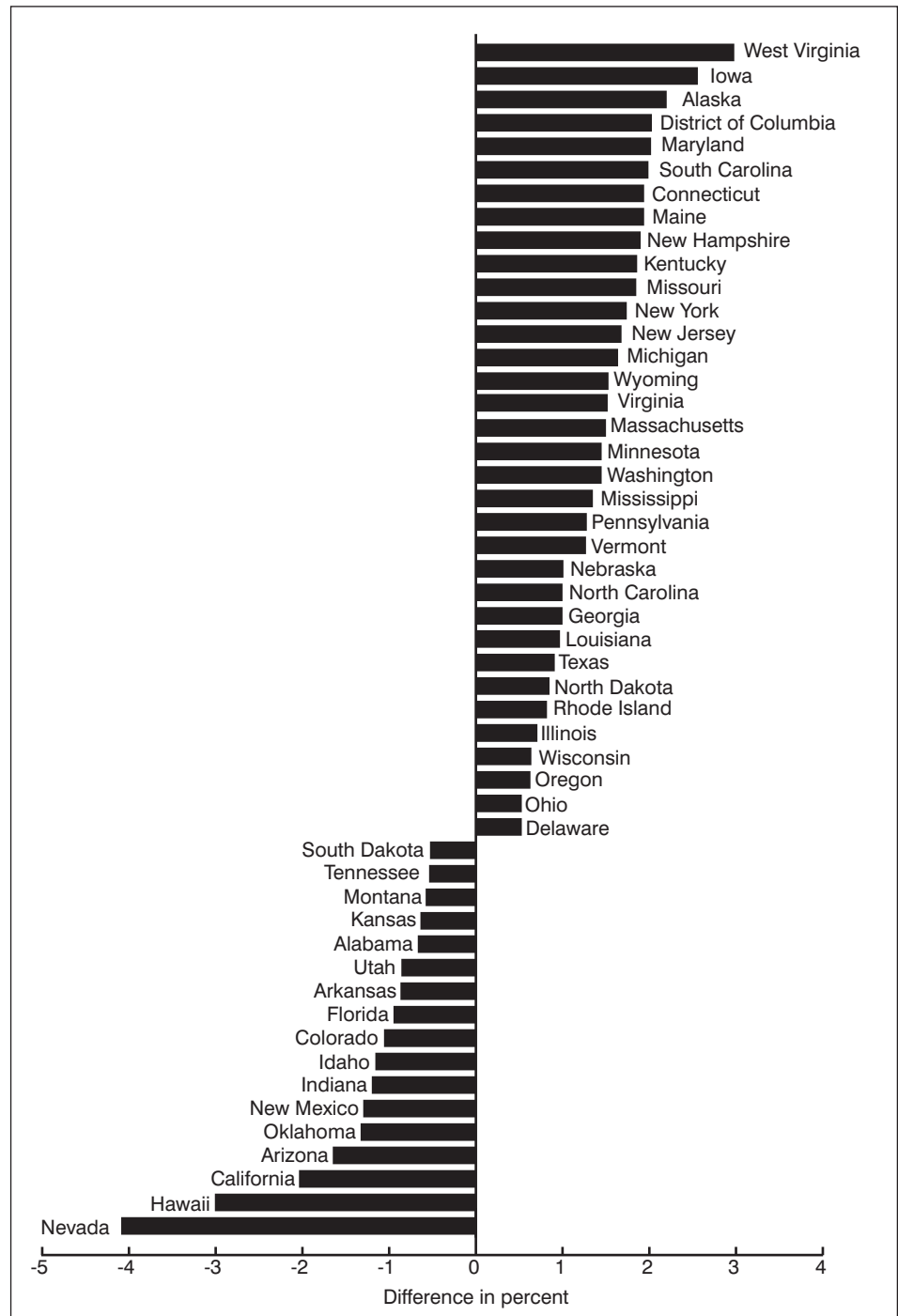


Figure. Difference in State-specific coverage rates using new and old estimation methods: National Immunization Survey, 1998

7. Summary

In 1998, 31,664 completed RDD interviews representing 32,511 age-eligible children were obtained. Among these, adequate provider data were obtained for 21,827 children allowing a determination of whether they were up-to-date according to the

recommended childhood immunization schedule. This report describes the statistical estimation methodology used to adjust for potential differences between sampled children in the NIS for whom provider-reported vaccination histories are obtained and sampled children for whom provider-reported vaccination histories are not obtained.

In the first step of the estimation methodology, a national logistic model of the propensity to have adequate provider data is obtained and estimated response propensities are obtained for all children with completed household RDD interviews. It is noted that the purpose of the logistic model is to rank children within each stratum with respect to their estimated response propensity. In this regard, a logistic model that includes stratum main effects would yield the same ranking within each stratum as a national model that does not include stratum main effects. Thus, a national model is sufficient to obtain this ranking provided there are no interactions of stratum effects with other variables that are predictive of the probability of having adequate provider data. Significant interactions with stratum effects were not detected in the analyses. Thus, a national model to estimate response propensities is suitable for obtaining a ranking of children within each stratum of the NIS sampling design.

In the second step of the estimation methodology, these ranked estimated response propensities were used to assign each sampled child to 1 of 5 adjustment cells formed using the quintiles of the distribution of estimated response propensities within each stratum. Next, within each adjustment cell in each stratum, sampling weights of children with adequate provider data were divided by the cell-specific weighted response rate to allow these children to represent all children in the adjustment cell. Finally, these adjusted sampling weights were raked to match the poststratification population control totals and totals of a few other variables that are known to be associated with being up-to-date.

Currently, official estimates of vaccination coverage rates are based on five adjustment cells within each stratum. To evaluate whether the use of five adjustment cells was sufficient to reduce bias attributable to differences between children with and without provider data, an investigation was conducted comparing the reduction in estimated bias obtained using five adjustment cells compared with an “optimal” number of cells in each

stratum.

Using an “optimal” choice for the number of adjustment cells within each stratum, results indicated that for the 1998 NIS, the overall bias reduction was 0.5 percent. These results indicate that in 1998, provider-vaccination history nonresponse bias was small. Also, no statistically significant differences were obtained in the vaccination coverage estimates by comparing estimates based on an “optimal” choice of the number of adjustment cells per stratum, compared with using five adjustment cells formed using the quintiles of the distribution of estimated response propensities within each stratum. Further, no important substantive differences were observed between estimates based on the methodology used prior to 1998 and the methodology adopted in 1998.

There are several methods to compensate for missing data and nonresponse in surveys. The level of nonresponse bias reduction may vary by the methodology used. The weighting adjustment procedure adopted in 1998 was selected, in part, to make the NIS estimation methodology as user-friendly as possible, thus permitting the use of commonly available and easy-to-use statistical software for the analysis of complex survey data. This was the primary motivation for adopting the new methodology.

In comparison with the two-phase estimation methodology used for the NIS before 1998, the approach described in this report offers several advantages. First, bias in NIS survey estimates attributable to using national information from top-row children in each State and IAP area estimates is reduced. However, it is noted that this bias was very small, in general. Secondly, compared with the two-phase sampling estimation approach, the method adopted in 1998 adjusts for provider nonresponse using a broad range of predictor variables. In the many strata in which only one adjustment cell is required, this feature makes no difference. However, for strata requiring more than one adjustment cell, this is important. These variables have been selected because they are highly predictive of the probability of provider nonresponse, the factor required to

adjust the sampling weights of children with adequate provider data so that they can represent the target population without incurring bias attributable to differences between children for whom adequate provider data were obtained and children for whom inadequate or no provider data were obtained.

The main conclusions of this report are that the vaccination coverage estimates based on the new estimation methodology are not that different from those based on the original methodology. In addition, little adjustment is required to account for differences between children for whom adequate provider data were obtained and children for whom inadequate or no provider data were obtained. Most importantly, however, the new methodology facilitates the preparation of analytical data files for use by analysts undertaking complex and important epidemiological analyses using available statistical software.

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Table 1. Percent of children with selected characteristics by provider data status: National Immunization Survey, 1998

Characteristic	Children with provider data	Children without provider data
	Percent	
Above poverty, household income less than \$75,000	56	47
Household used a shot card during RDD interview ¹	51	38
Household respondent was the mother	88	81
Household moved out of State since child's birth	8	14
Household is located in a central city	43	46
Race of the mother is white, non-Hispanic	67	60
	Number	
Unweighted sample size	21,827	10,684

¹RDD is random-digit-dialing.NOTE: All differences are statistically significant at the $p=0.05$ level.**Table 2. Variables in the model selection for the response propensity model: National Immunization Survey, 1998**

Name of variable	Predictor description and factor levels	Name of variable	Predictor description and factor levels
all4shot	Household report of up-to-date on 4:3:1:3 (4+ DTP, 3+ POL, 1+ MCV, and 3+ Hib)	marital	Marital status of the mother
1	Yes — R	1	Widowed/divorced/separated — R
2	No	2	Never married
3	Don't know, refused, missing	3	Married
		4	Deceased
c5	Relationship of the household respondent to the child (mother, father, or other)	mobil	Mobility status
1	Mother (step, foster, adoptive) or female guardian — R	1	Moved, different State — R
2	Father (step, foster, adoptive) or male guardian	2	Did not move from different State
3	Other	MSA ¹	Household within central city MSA, suburban, or nonmetropolitan area
4	Missing	1	MSA, central city — R
childnm	Number of children under 18 years of age living in the household	2	MSA, noncentral city
1	1 child — R	3	Non-MSA
2	2–3 children	racekid	Race of the child
3	4 or more children	1	Hispanic — R
educ1	Educational status of the mother	2	White, non-Hispanic
1	Less than 12 years — R	3	Black, non-Hispanic
2	12 years	4	American Indian, non-Hispanic
3	More than 12 years, noncollege graduate	5	Asian, non-Hispanic
4	College graduate	6	Other, non-Hispanic
full.cpo	Household report of up-to-date on varicella	racemom	Race of the mother
1	Yes — R	1	Hispanic — R
2	No	2	White, non-Hispanic
3	Missing	3	Black, non-Hispanic
full.hep	Household report of up-to-date on hepatitis	4	American Indian, non-Hispanic
1	Yes — R	5	Asian, non-Hispanic
2	No	6	Other, non-Hispanic
3	Missing	sex	Gender of the child
incpov1	Poverty status	1	Male — R
1	Above, \$75,000 or more — R	2	Female
2	Above, less than \$75,000	shotcard	Household reported immunization status was obtained using a "shot" card
3	Below	1	Shot card used during RDD interview — R
4	Unknown	2	Shot card not used during RDD interview
m.agegrp	Maternal age group		
1	Less than 19 years — R		
2	20–29 years		
3	30 years and over		

is metropolitan statistical area.

NOTE: R identifies reference level.

¹MSA

Table 3. Results of the stepwise variable selection procedure for the response propensity model

Step	Variable added	Degrees of freedom	Chi-square	Akaike's information criterion
1				36,909
2	+ incpov1	3	581.8	36,333
3	+ shotcard	1	431.9	35,903
4	+ c5	3	248.7	35,660
5	+ mobil	1	155.0	35,507
6	+ msa	2	109.2	35,402
7	+ full.hep	2	57.6	35,349
8	+ racemom	5	52.8	35,306
9	+ educ1	3	27.3	35,284
10	+ childnm	2	19.1	35,269
11	+ m.agegrp	2	15.7	35,258
12	+ full.cpo	2	12.0	35,250
13	+ all4shot	2	11.4	35,242
14	+ sex	1	2.6	35,242

NOTE: Final model for response variable y , indicating whether a sample child has adequate provider data: $y = \text{incpov1} + \text{shotcard} + \text{c5} + \text{mobil} + \text{msa} + \text{full.hep} + \text{racemom} + \text{educ1} + \text{childnm} + \text{m.agegrp} + \text{full.cpo} + \text{all4shot} + \text{sex}$.

Table 4. Estimated log-odds and associated standard errors of the predictors selected in the stepwise procedure for the final model: National Immunization Survey, 1998

Variable	Odds ratio	Log-odds ratio	Standard error	t-value	p-value ¹
Intercept	2.97	1.09	0.11	10.36	<0.01
Household report of up-to-date on 4:3:1:3					
Yes	1.00	0.00			
No	0.91	-0.09	0.03	-2.86	<0.01
Don't know, refused, missing	0.85	-0.16	0.04	-3.65	<0.01
Relationship of the household respondent to the child					
Mother	1.00	0.00			
Father	0.74	-0.30	0.04	-7.45	<0.01
Other (neither father nor mother)	0.62	-0.48	0.07	-6.70	<0.01
Missing	0.04	-3.18	0.43	-7.32	<0.01
Number of children under 18 years of age living in the household					
1 child	1.00	0.00			
2-3 children	1.16	0.15	0.03	4.97	<0.01
4 or more children	1.09	0.09	0.05	1.94	0.05
Educational status of the mother					
Less than 12 years	1.00	0.00			
12 years	0.84	-0.17	0.05	-3.80	<0.01
More than 12 years, noncollege graduate	0.85	-0.16	0.05	-3.31	<0.01
College graduate	0.88	-0.13	0.05	-2.64	0.01
Household report of up-to-date on varicella					
Yes	1.00	0.00			
No	0.91	-0.09	0.03	-3.35	<0.01
Missing	0.99	-0.01	0.05	-0.16	0.87
Household report of up-to-date on Hepatitis B					
Yes	1.00	0.00			
No	0.82	-0.20	0.03	-5.83	<0.01
Missing	0.91	-0.10	0.05	-1.94	0.05
Poverty status					
Above, \$75,000 or more	1.00	0.00			
Above, less than \$75,000	0.94	-0.06	0.04	-1.31	0.19
Below poverty	0.86	-0.15	0.05	-2.77	0.01
Unknown	0.44	-0.81	0.05	-15.60	<0.01
Maternal age group					
19 years or younger	1.00	0.00			
20-29 years	0.80	-0.22	0.07	-3.02	<0.01
30 years and over	0.77	-0.27	0.08	-3.52	<0.01
Mobility status					
Moved from different State	1.00	0.00			
Did not move from different State	1.67	0.51	0.04	12.18	<0.01
Household within central city MSA, ² suburban, or nonmetropolitan area					
MSA, ² central city	1.00	0.00			
MSA, ² noncentral city	0.94	-0.06	0.03	-1.92	0.05
Non-MSA ²	1.32	0.28	0.04	7.38	<0.01
Race of the mother					
Hispanic	1.00	0.00			
White, non-Hispanic	1.18	0.17	0.04	3.93	<0.01
Black, non-Hispanic	0.89	-0.12	0.05	-2.47	0.01
American Indian, non-Hispanic	1.06	0.06	0.11	0.53	0.59
Asian, non-Hispanic	0.98	-0.02	0.07	-0.24	0.81
Other race, non-Hispanic	1.33	0.28	0.23	1.22	0.22

See footnotes at end of table.

Table 4. Estimated log-odds and associated standard errors of the predictors selected in the stepwise procedure for the final model: National Immunization Survey, 1998—Con.

Variable	Odds ratio	Log-odds ratio	Standard error	t-value	p-value ¹
Gender					
Male	1.00	0.00			
Female	0.96	-0.04	0.03	-1.74	0.08
Household reported immunization status was obtained using a "shot" card					
Shot card used during RDD ³ interview	1.00	0.00			
Shot card not used during interview	0.67	-0.40	0.03	-12.46	<0.01

¹p-values correspond to two-sided tests.

²MSA is metropolitan statistical area.

³RDD is random-digit-dialing.

Table 5. Variables used for raking the response propensity adjusted weights of children with provider data: National Immunization Survey, 1998

Variable name	Variable description
educ1 ¹	Educational status of the mother
racekid ¹	Race of the child
agegrp ¹	Age group of the child
sex	Gender of the child
frstbrn	First born status of the child
adj.cell	Response propensity adjustment cell to which each child belongs within each IAP area

¹This variable was also used to provide a poststratified adjustment to the random-digit-dialing phase sampling weights.

Table 6. Illustration of method for determining optimal number of adjustment cells for Illinois-Rest of State: National Immunization Survey, 1998

k	$\hat{\Delta}_{Y_k}$	$\hat{\Delta}_{\Delta_k}$	p_k	$\hat{\gamma}_k$
2	83.13	-0.5	0.03	0.97
3	82.73	-0.4	0.13	0.92
4	82.87	0.14	0.76	1.03
5	82.84	-0.03	0.46	0.97

k is number of adjustment cells.

$\hat{\Delta}_{Y_k}$ is nonresponse adjusted estimate of 4:3:1:3 coverage based on k cells.

$\hat{\Delta}_{\Delta_k}$ is estimated bias reduction resulting from using k cells compared with k-1 cells, k=2, . . .

p_k is p-value of the null hypothesis $H: \Delta_k=0$.

$\hat{\gamma}_k$ is estimated relative efficiency of using k cells, compared to k-1 cells, k=2, . . .

Table 7. Summary of nonresponse adjusted estimates of 4:3:1:3 coverage and estimated bias reduction obtained by using optimal number of adjustment cells: National Immunization Survey, 1998

Area	k	$\hat{\Delta}_{Y_k}$	$\hat{\Delta}_k$
IL-Rest of State	2	83.13	-0.5
IL-City of Chicago	1	65.28	0
IN-Rest of State	1	77.88	0
IN-Marion County	2	78.2	-0.25
MI-Rest of State	1	79.17	0
MI-Detroit	2	69.66	-0.44
OH-Rest of State	1	78.5	0
OH-Cuyahoga County	1	74.63	0
OH-Franklin County	1	77.69	0
WI-Rest of State	1	79.19	0
WI-Milwaukee County	2	73.23	-0.22
AL-Rest of State	1	81.86	0
AL-Jefferson County	1	83.71	0
Kentucky	1	81.57	0
Mississippi	1	83.9	0
TN-Rest of State	2	84.73	-0.09
TN-Shelby County	1	71.23	0
TN-Davidson County	1	79.68	0
NJ-Rest of State	1	83.58	0
NJ-City of Newark	1	64.77	0
NY-Rest of State	1	87.85	0
NY-5 Counties	1	81.62	0
PA-Rest of State	2	83.97	-0.09
PA-Philadelphia	1	80.08	0
AZ-Rest of State	3	73.94	-0.5
AZ-Maricopa County	1	77.35	0
Colorado	1	75.8	0
Idaho	1	76.82	0
Montana	1	81.88	0
New Mexico	1	71.91	0
Nevada	1	76.39	0
Utah	1	75.83	0
Wyoming	1	80.19	0
Connecticut	1	89.48	0
MA-Rest of State	1	86.33	0
MA-City of Boston	1	89	0
Maine	1	86.36	0
New Hampshire	1	82.26	0
Rhode Island	1	86.16	0
Vermont	2	85.47	-0.51
Alaska	2	80.72	-0.41
CA-Rest of State	2	74.47	-0.71
CA-Los Angeles	1	76.35	0
CA-Santa Clara	2	84.23	-0.81
CA-San Diego County	1	77.19	0
Hawaii	2	79.59	-1.14
Oregon	2	76.19	-0.4
WA-Rest of State	1	79.33	0
WA-King County	1	86.04	0
District of Columbia	1	71.76	0
Delaware	1	78.82	0
FL-Rest of State	1	79.32	0
FL-Duval County	1	78.78	0
FL-Dade County	1	75.07	0
GA-Rest of State	1	82.54	0
GA-Fulton/Dekalb	1	70.76	0
MD-Rest of State	2	76.18	-0.86
MD-Baltimore City	1	81.31	0
North Carolina	1	82.77	0
South Carolina	1	88.1	0
Virginia	1	80.93	0
West Virginia	1	82.46	0
Iowa	2	81.64	-0.58
Kansas	1	82.09	0
Minnesota	1	82.27	0
Missouri	2	84.07	-1.21
North Dakota	2	79.58	-0.58

Table 7. Summary of nonresponse adjusted estimates of 4:3:1:3 coverage and estimated bias reduction obtained by using optimal number of adjustment cells: National Immunization Survey, 1998—Con.

Area	k	$\hat{\bar{Y}}_k$	$\hat{\Delta}_k$
Nebraska	2	76.55	-0.41
South Dakota	1	73.52	0
Arkansas	3	73.16	-0.8
LA-Rest of State	1	78.86	0
LA-Orleans Parish	2	79.15	-0.42
Oklahoma	1	77.15	0
TX-Rest of State	1	77.52	0
TX-Dallas County	1	72.85	0
TX-El Paso County	1	77.95	0
TX-City Houston	3	60.56	-1.04
TX-Bexar County	1	78.91	0

k is number of adjustment cells.

$\hat{\bar{Y}}_k$ is nonresponse-adjusted estimate of 4:3:1:3 coverage based on k cells.

$\hat{\Delta}_k$ is estimated bias reduction resulting from using k cells compared with $k-1$ cells, $k=2, \dots$

Table 8. Statistical significance of the estimated difference in nonresponse-adjusted estimates of 4:3:1:3 coverage obtained by using optimal choice of k compared with using five adjustment cells in each IAP area: National Immunization Survey, 1998

State	$\hat{\Delta}_{Y_{k'}}$	$\hat{\Delta}_{Y_{k'}} - \hat{\Delta}_{Y_5}$	p	$\hat{\gamma}_{k'}$
National	79.4	0.19	0.01	1.19
AK	80.72	-0.58	0.1	1
AL	82.14	0.02	0.95	1.03
AR	73.16	0.03	0.91	1.01
AZ	76.04	0.19	0.55	1.04
CA	75.77	-0.08	0.79	1.01
CO	75.79	-0.04	0.9	1.01
CT	89.47	-0.57	0.15	0.99
DC	71.76	0.33	0.61	1.05
DE	78.82	-0.12	0.69	1.01
FL	78.56	-0.09	0.75	1
GA	80.28	0.19	0.51	1.03
HI	79.59	0.31	0.27	1.03
IA	81.64	-0.02	0.84	1
ID	76.82	0.38	0.25	1.05
IL	77.97	0.46	0.13	1.11
IN	77.93	0.43	0.23	1.08
KS	82.09	0.33	0.12	1.05
KY	81.56	0.01	0.97	1.01
LA	78.89	0.53	0.17	1.11
MA	86.6	-0.1	0.64	1
MD	76.95	-0.07	0.8	1
ME	86.36	0.02	0.95	1.02
MI	77.96	0.23	0.61	1.06
MN	82.27	0.11	0.74	1.03
MO	84.07	-0.48	0.37	0.98
MS	83.9	0.21	0.37	1.03
MT	81.88	-0.04	0.81	1.01
NC	82.77	-0.03	0.85	1
ND	79.58	0.53	0.08	1.09
NE	76.55	0.14	0.61	1.02
NH	82.26	0.07	0.85	1.02
NJ	82.7	0.42	0.2	1.11
NM	71.91	0.81	0.26	1.11
NV	76.39	0.67	0.24	1.1
NY	84.93	0.39	0.22	1.07
OH	77.91	-0.12	0.82	1.04
OK	77.15	1.88	0.02	1.51
OR	76.19	0.66	0.17	1.1
PA	83.36	0.19	0.23	1.03
RI	86.16	-0.15	0.37	0.98
SC	88.09	0.21	0.63	1.04
SD	73.52	0.05	0.85	1.01
TN	81.46	0.01	0.96	1
TX	74.94	0.64	0.03	1.15
UT	75.83	0.19	0.47	1.01
VA	80.92	0.5	0.17	1.05
VT	85.47	-0.29	0.3	0.94
WA	81.18	0.34	0.36	1.06
WI	77.83	0.09	0.51	1.01
WV	82.46	0.08	0.77	1.03
WY	80.2	0.27	0.42	1.04

$\hat{\Delta}_{Y_{k'}}$ is nonresponse-adjusted estimate of 4:3:1:3 coverage based on k cells.

$\hat{\Delta}_{Y_{k'}} - \hat{\Delta}_{Y_5}$ is difference in estimated bias in nonresponse-adjusted estimates of 4:3:1:3 coverage obtained by using optimal choice of k adjustment cells compared with using five adjustment cells.

$p_{k'}$ is p -value of the null hypothesis $H: \hat{\Delta}_{Y_{k'}} - \hat{\Delta}_{Y_5} = 0$.

$\hat{\gamma}_{k'}$ is estimated relative efficiency of using k' cells, compared to five cells, $k=2, \dots$

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