



AGENCY FOR HEALTHCARE RESEARCH AND QUALITY



# MEPS – HC

## Estimation and Analysis

Sadeq Chowdhury, PhD

# Outline

- **Estimation from MEPS-HC**
  - ▶ **Computing/Producing Estimates**
  - ▶ **Computing Precision of Estimates**
- **Analysis of Subpopulations**
- **Family-Level Analysis**
- **Other Types of Analyses**

# **Estimation From MEPS**

**(Producing Estimates &  
Computing Precision or Reliability)**

# Producing Estimates - Weights Must be Used

- **Sample weight**
  - Indicates how many persons in the population a sample person represents
- **Unequal sample weights due to**
  - Oversampling of Blacks, Hispanics, Asians
  - Differential response rates
- **Weights must be used to produce unbiased estimates**
  - Unweighted estimates are biased

# Distribution of Final Positive Person Weights

Distribution of Weight	Year		
	2019	2020	2021
<b>Minimum</b>	<b>497</b>	<b>331</b>	<b>229</b>
<b>Average</b>	<b>10,574</b>	<b>12,238</b>	<b>12,119</b>
<b>Maximum</b>	<b>104,866</b>	<b>101,795</b>	<b>106,959</b>
<b>Variable Name</b>	<b>PERWT19F</b>	<b>PERWT20F</b>	<b>PERWT21F</b>

# Final Person Weights - Positive versus Zero

- **Weight > 0 (i.e., positive)**
  - ▶ **Persons Key and in-scope for survey**
  - ▶ **More than 95% cases**
- **Weight = 0**
  - ▶ **About 5% of cases every year**
  - ▶ **Persons not key or in-scope for survey but living in households with in-scope person(s)**
  - ▶ **Included for family analysis**

# Measures of Precision/ Reliability of Estimates

- **Sampling error, Variance or Standard error**
- **Standard Error (SE) =  $\sqrt{\text{Variance}}$**
- **Relative Standard Error (RSE)**
  - ▶ **SE of estimate ÷ estimate**
  - ▶ **also called Coefficient of Variation (CV)**
- **Confidence Interval (CI)**
  - ▶ **95% CI: Estimate  $\pm$  1.96xSE**

# Example: Precision of Average Total Expenses, 2021

- Sample Size **27,332**
- Estimate = **\$6,934** (Average Expense per Capita)
- Standard Error = **189**
- 95% Confidence Interval  
= (**\$6,934 ± 1.96x189, i.e., \$6,563 to \$7,306**)
- Relative Standard Error (RSE)  
= (**189 ÷ 6,934**) x 100 = **2.7%**



# Computing Variances of Estimates from Complex Sample Design

- **Appropriate method must be used to compute standard errors to account for complex sample design**
- **Assuming simple random sampling usually underestimates standard errors**

# Computing Standard Error (Precision of an Estimate)

- **Basic software procedures assume simple random sampling (SRS)**
  - ▶ **Estimates correct if weighted**
  - ▶ **Standard errors usually smaller than actual**
- **Software to account for complex design**
  - ▶ **SUDAAN (stand-alone or callable within SAS)**
  - ▶ **STATA (svy commands)**
  - ▶ **SAS 9.2 (survey procedures)**
  - ▶ **R (survey package)**
  - ▶ **Other (SPSS)**

# Example:

## Average Total Expenditures, 2021

- **Weighted mean = \$ 6,934 per capita**  
**Unweighted mean = \$ 7,870 (biased)**
- **SE complex survey procedure = 189**
  - ▶ **SAS: PROC SURVEYMEANS**
  - ▶ **SUDAAN: PROC DESCRIPT**
  - ▶ **Stata: svy: mean**
  - ▶ **R: svymean**
- **SE assuming SRS = 167 (too low)**
  - ▶ **SAS: PROC UNIVARIATE or MEANS**

# Example Codes to Produce Estimates and SEs

- **SAS V9.2**

```
proc surveymeans data=HC224 mean;  
stratum varstr; cluster varpsu;  
weight perwt20f; var totexp20;
```

- **Stata**

```
svyset varpsu [pweight=perwt20f], strata(varstr)  
svy: mean 2
```

- **SUDAAN (SAS-callable)**

First sort the file by varstr & varpsu

```
proc descript data= HC224 filetype=SAS design=wr;  
nest varstr varpsu; weight perwt20f;  
var totexp20;
```

- **R**

```
mepsdsgn = svydesign(id = ~varpsu, strata = ~varstr, weights = ~perwt20f,  
  data = HC224, nest = TRUE)  
svymean(~ totexp20, design = mepsdsgn)
```

# Computing Standard Errors for MEPS Estimates



## Document on MEPS website

[http://www.meps.ahrq.gov/mepsweb/survey\\_comp/standard\\_errors.jsp](http://www.meps.ahrq.gov/mepsweb/survey_comp/standard_errors.jsp)

# **Domain Estimation or Analysis of Subpopulations**

# Analysis of Subpopulations – Special Procedure Needed

- **Analysis within specific subpopulation say within a Race-ethnicity, Poverty or Insurance status categories**  
**Example: Asian 65+ years only or Uninsured Hispanics**
- **Special procedure or domain analysis must be used**

# Analysis of Subpopulations – Avoid Subsetting the File

- Analyzing a subset file may produce incorrect standard errors
- A subset file of the sample may not contain all variance estimation information
- Software may give error messages in some situations
- Particularly important for analyzing small subpopulations that are not available in all PSUs
- Subsetting is ok for large subpopulations which are likely to be available in all PSUs such as males, females, children, elderly, etc.



# Keywords for Specifying Subpopulations

- Each software has special facility for subpopulation analysis using the entire file
  - SAS: **domain**
  - SUDAAN: **subpopn**
  - Stata: **subpop**
  - R: **subset**

## *Example*

```
proc surveymeans data=HC233 mean;  
stratum varstr; cluster varpsu;  
weight perwt20f; var totexp20;  
domain racethnx;
```

# References on Analysis of Subpopulations

- **Variance Estimation from MEPS Event Files**
  - ▶ [http://meps.ahrq.gov/mepsweb/data\\_files/publications/mr26/mr26.pdf](http://meps.ahrq.gov/mepsweb/data_files/publications/mr26/mr26.pdf)

# Family-Level Estimation

# Family-Level Estimation

- **Create a family-level file containing one record per family from person-level file (see instructions in the PUF documentation)**
- **Two family type options:**
  - **MEPS: Includes unmarried couples** (starting with 2017, foster children excluded from MEPS)
  - **Current Population Survey (CPS): Unmarried couples not family unit**
- **Two timeframe options:**
  - **December 31 (MEPS, CPS)**
  - **Any time during year (MEPS only)**

# Family Sample Sizes, 2021 (MEPS Annual Household File)

<b>Number of Families</b>	<b>MEPS Anytime During Year</b>	<b>MEPS December 31</b>	<b>CPS December 31</b>
<b>Unweighted</b>	<b>13,014</b>	<b>12,910</b>	<b>13,461</b>
<b>Weighted (millions)</b>	<b>142.5</b>	<b>141.3</b>	<b>147.8</b>
<b>Family Weight Variable Name</b>	<b>FAMWT21F</b>	<b>FAMWT21F (subset to FMRS1231 = 1)</b>	<b>FAMWT21C</b>

# Family-Level Estimation

## Example: Average total healthcare expenses per MEPS family by family size, 2021

- The estimation is based on MEPS families in scope at any time during the year
- Average number of persons per family is about 2.3

Family size	Estimate	SE
All	\$15,526	\$436
1	\$10,719	\$443
2	\$18,772	\$748
3	\$18,121	\$1,757
4	\$17,741	\$1,293
5+	\$17,873	\$1,371

# Estimation and Analysis

- Event Level
- State Level
- MEPS-HC Supplements

# Medical Event as Unit of Analysis

- **Event files can be used to estimate average expense per event**
- **Some estimates are available in HC summary tables on MEPS website**

## **Example:**

**In 2021,**

- **the mean expense per office visit to a physician was \$365 (SE = \$16)**
- **the mean expense per emergency room (ER) visit was \$1,164 (SE = \$48)**
- **the mean expense per inpatient stay was \$16,809 (SE = \$639)**



# State-Level Estimation



- **Considerable interest in state-level estimates**
- **MEPS sample is not designed to produce estimates for all states**
  - Small sample sizes
  - Insufficient number of PSUs for variance estimation
- **To protect confidentiality, public use files (PUFs) do not include state identifiers**
  - Tables and reports available on website for larger states (selected estimates)
  - Access to identifiers in AHRQ Data Center
    - *Need to use state-level sample design and state identifiers*

# MEPS-HC Supplements

- Special supplement variables on person-level files
- Consult documentation for appropriate weight
  - Self-Administered Questionnaire (SAQ) → SAQWTyyF
  - Diabetes Care Survey (DCS) → DIABWyyF
  - Cancer SAQ (CSAQ) → CSAQWyyF
    - MEPS full-year files 2011, 2016 and 2017 only
  - Veteran SAQ (VSAQ) → VSAQWyyF
    - MEPS full-year files 2018 and 2019 only

# Thank you!



[Sadeq.Chowdhury@ahrq.hhs.gov](mailto:Sadeq.Chowdhury@ahrq.hhs.gov)