



Survey Methods II: Statistical Analysis of Surveys in R

CSDE Workshop

Jessica Godwin

January 27, 2026

Resources and Materials
oooo

Why survey statistics?
oooo

Survey Designs
oooooooooooooooooooo

Estimation with Survey Data
oooooooooooooooooooo

Data Visualization
oooooooo

Etc.
oooo

Resources and Materials

Why survey statistics?

Survey Designs

Estimation with Survey Data

Data Visualization

Etc.

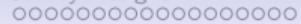
Resources and Materials



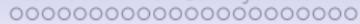
Why survey statistics?



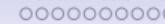
Survey Designs



Estimation with Survey Data



Data Visualization



Etc.



Resources and Materials

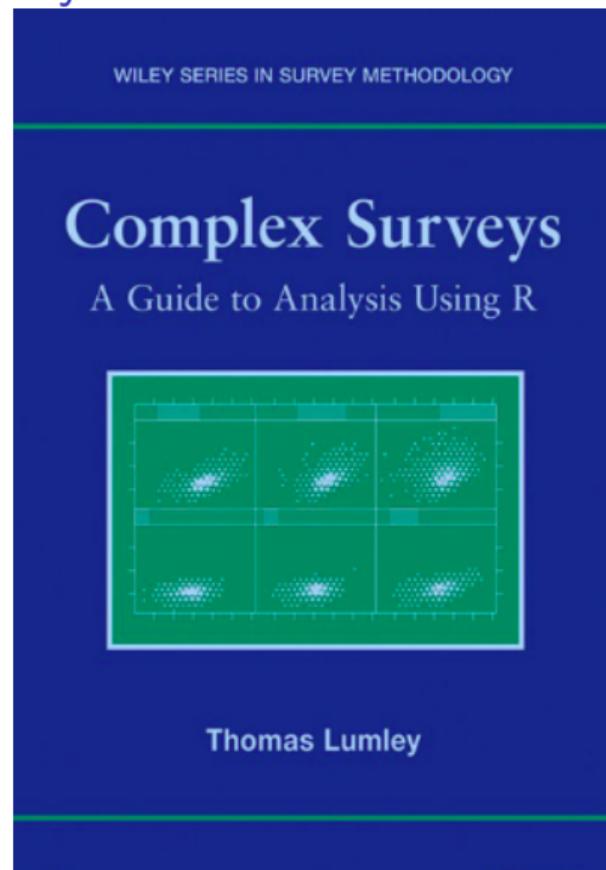
Install survey

- Survey Package Documentation

```
install.packages("survey", dependencies = TRUE)
```

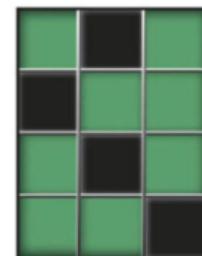
Complex Surveys in R

- Lumley, Thomas (2004). Analysis of Complex Survey Samples. *Journal of Statistical Software*, 9(8), 1–19.
<https://doi.org/10.18637/jss.v009.i08>
- Lumley, Thomas (2011). Complex surveys: a guide to analysis using R (Vol. 565). John Wiley & Sons.



Survey Statistics

- Lohr, Sharon L (1999). Sampling: Design and analysis. Pacific Grove, CA: Duxbury Press. <https://doi.org/10.1201/9780429298899>
- Särndal, C. E., Swensson, B., & Wretman, J. (2003). Model assisted survey sampling. Springer Science & Business Media. <https://link.springer.com/book/9780387406206>
- CSSS/STAT 529 (Spring, taught by Elena Erosheva or Jon Wakefield)



Sampling: Design and Analysis

SECOND EDITION

Sharon L. Lohr

Resources and Materials
oooo

Why survey statistics?
●ooo

Survey Designs
oooooooooooooooooooo

Estimation with Survey Data
oooooooooooooooooooo

Data Visualization
ooooooo

Etc.
oooo

Why survey statistics?

Why survey statistics?

- If our outcome is binary:

Why survey statistics?

- If our outcome is binary:
 - Did our data arise from flipping a coin? or
 - Did our data arise from drawing from an urn?

Why survey statistics?

- If our outcome is binary:
 - Did our data arise from flipping a coin? or
 - Did our data arise from drawing from an urn?
- What does flipping a coin or drawing from an urn have to do with surveys with human respondents?

Finite vs. superpopulations

For observations $i = 1, \dots, n$, let

$$y_i = \begin{cases} 1, & \text{success,} \\ 0, & \text{failure.} \end{cases}$$

- Superpopulation: If $y \sim \text{Bernoulli}(p)$,
 - $E[y] = p$ $\text{Var}(y) = p(1 - p)$.
- Finite population: If $y \sim \text{Hypergeometric}(N, K, n)$,
 - $E[y] = \frac{K}{N}$ $\text{Var}(y) = \frac{K}{N} \left(1 - \frac{K}{N}\right) \left(1 - \frac{n}{N}\right)$. How do we say what \hat{p} means in either case? Is it the same?

Finite vs. superpopulations, cont'd

If $y_i \stackrel{\text{iid}}{\sim} \text{Bernoulli}(p)$,

$$\hat{p} = \frac{\sum_{i=1}^n y_i}{n},$$

$$\widehat{Var}(\hat{p}) = \frac{\hat{p}(1 - \hat{p})}{n}.$$

If $y_i \stackrel{\text{iid}}{\sim} \text{Hypergeometric}(N, K, n)$,

$$\hat{p} = \frac{\sum_{i=1}^n y_i}{n} = \frac{k}{n}$$

$$\widehat{Var}(\hat{p}) = \frac{\hat{p}(1 - \hat{p})}{n} \times \left(1 - \frac{n}{N}\right).$$

How do we say what \hat{p} means in either case? Is it the same?

Resources and Materials
oooo

Why survey statistics?
oooo

Survey Designs
●oooooooooooooooooooo

Estimation with Survey Data
oooooooooooooooooooo

Data Visualization
oooooooo

Etc.
oooo

Survey Designs

Simple random sampling(SRS)

- Under an **SRS** of n observations
- Under an **SRSWOR** of n observation

$$\Pr(\text{subject } k \in \text{sample}, S) =$$

$$\pi_k = \frac{1}{N}$$

$$\Pr(\text{subjects } k, k' \in \text{sample}, S) =$$

$$\pi_{k,k'} = \frac{1}{N} \times \frac{1}{N}.$$

$$\Pr(\text{subject } k \in \text{sample}, S) =$$

$$\pi_k = \frac{n}{N}$$

$$\Pr(\text{subjects } k, k' \in \text{sample}, S) =$$

$$\pi_{k,k'} = \frac{n}{N} \times \frac{n-1}{N-1}.$$

Specifying the Design: `svydesign`

- `svydesign()` needs to know what columns in your data (if any) represent
 - sampling weights (`weights`)
 - strata (`strata`)
 - clusters (`ids`)
 - units
 - finite population correct (`fpc`)

```
library(survey)
data(api)
?api
?svydesign
```

Specifying the Design: `svydesign`

- `apisrs$pw`: sampling weight
- `apisrs$fpc`: finite population correction, i.e. N for an SRS
- `apisrs$stype`: school type (elementary, middle, high)

```
table(apisrs$pw,  
      apisrs$fpc)
```

```
##  
##          6194  
##    30.97 200
```

6194/200

```
## [1] 30.97
```

Specifying the Design: `svydesign`

- `apisrs$pw`: sampling weight
- `apisrs$fpc`: finite population correction, i.e. N for an SRS
- `apisrs$type`: school type (elementary, middle, high)

```
table(apisrs$pw,  
      apisrs$type)
```

```
##  
##          E    H    M  
##    30.97 142   25   33
```

Specifying the Design: `svydesign`

```
srs_des <- svydesign(ids = ~1, weights = ~pw,  
                      fpc = ~fpc, data = apisrs)  
srs_des
```

```
## Independent Sampling design  
## svydesign(ids = ~1, weights = ~pw, fpc = ~fpc, data = apisrs)
```

Systematic sampling

- Select every r^{th} sampling unit from the sampling frame of length N : $r \times n \leq N < r \times (n + 1)$
 - What is π_k for individual $k = r$? $k = r + 1$?

Systematic sampling

- Select every r^{th} sampling unit from the sampling frame of length N : $r \times n \leq N < r \times (n + 1)$
 - What is π_k for individual $k = r$? $k = r + 1$?
 - Can a systematic sample be implemented so that it is the equivalent of an SRS?

Systematic sampling

- Select every r^{th} sampling unit from the sampling frame of length N : $r \times n \leq N < r \times (n + 1)$
 - What is π_k for individual $k = r$? $k = r + 1$?
 - Can a systematic sample be implemented so that it is the equivalent of an SRS?
 - What is $\pi_{r,r+1}$?

Systematic sampling

- Select every r^{th} sampling unit from the sampling frame of length N : $r \times n \leq N < r \times (n + 1)$
 - What is π_k for individual $k = r$? $k = r + 1$?
 - Can a systematic sample be implemented so that it is the equivalent of an SRS?
 - What is $\pi_{r,r+1}$?
- Random single start → what changes?

Systematic sampling

- Select every r^{th} sampling unit from the sampling frame of length N : $r \times n \leq N < r \times (n + 1)$
 - What is π_k for individual $k = r$? $k = r + 1$?
 - Can a systematic sample be implemented so that it is the equivalent of an SRS?
 - What is $\pi_{r,r+1}$?
- Random single start \rightarrow what changes?
- Multiple starts
 - No individual sampling probabilities are 0 or 1
 - Joint sampling probabilities defined

Stratified simple random sampling (strSRS)

- Consider $h = 1, \dots, H$ strata from each of which you want to sample n_h individuals.

$$\Pr(\text{subject } k \in S_h) = \pi_k = \frac{n_h}{N_h}$$

$$\Pr(\text{subjects } k, k' \in S_h) = \pi_{k,k'} = \frac{n_h}{N_h} \times \frac{n_h - 1}{N_h - 1}$$

$$\Pr(\text{subjects } k \in S_h, k' \in S_{h'}) = \pi_{k,k'} = \frac{n_h}{N_h} \times \frac{n_{h'}}{N_{h'}}.$$

strSRS, cont'd

- Why stratify? Why not an SRS or SRSWOR?

strSRS, cont'd

- Why stratify? Why not an SRS or SRSWOR?
 - Availability of **sampling frame**
 - Cost, convenience, speed
 - N_1, \dots, N_h vary widely
 - Rare outcomes within certain strata
 - We know strata are related to outcome of interest → precision gains!
- What happens if we ignore the stratification?
 - Waste a lot of folks' money!!
 - Implicit assumption that outcome of interest doesn't differ by strata
 - → obscure differences in outcomes by strata
 - → OVERESTIMATE variance/standard errors
 - → worsens variability in outcomes between strata grows and within strata shrinks
 - → worsens as variability in $\pi_{k \in S_h}$ between strata grows

Specifying the Design: `svydesign`

- `apisrs$pw`: sampling weight
- `apisrs$fpc`: finite population correction, i.e. N_h for an strSRS
- `apisrs$stype`: strata; chool type (elementary, middle, high)

```
table(apistrat$pw, apistrat$fpc)
```

```
##  
## 755 1018 4421  
## 15.1000003814697 50 0 0  
## 20.3600006103516 0 50 0  
## 44.2099990844727 0 0 100
```

755/50

```
## [1] 15.1
```

Specifying the Design: `svydesign`

- `apisrs$pw`: sampling weight
- `apisrs$fpc`: finite population correction, i.e. N_h for an strSRS
- `apisrs$stype`: strata; chool type (elementary, middle, high)

```
table(apistrat$pw,  
      apistrat$stype)
```

```
##          E      H      M
## 15.1000003814697 0 50 0
## 20.3600006103516 0 0 50
## 44.20999990844727 100 0 0
```

Specifying the Design: svydesign

```
strsrs_des <- svydesign(ids = ~1,  
                         strata = ~stype,  
                         weights = ~pw,  
                         fpc = ~fpc,  
                         data = apistrat)  
strsrs_des
```

```
## Stratified Independent Sampling design  
## svydesign(ids = ~1, strata = ~stype, weights = ~pw, fpc = ~fpc,  
##             data = apistrat)
```

Cluster sampling

Consider sampling $c = 1, \dots, C$ clusters or **primary sampling units (PSU)** from your population of N_C clusters and N **units**.

Individuals k are the **observation units** contained within clusters on which we will make measurements.

One-stage cluster sampling

Two-stage cluster sampling

Sample m_c from M_c units in cluster c .

$$\Pr(\text{PSU } c \in S) = \frac{C}{N_c}$$

$$\pi_{k \in S_c} = \begin{cases} 1, & \text{PSU } c \in S, \\ 0, & \text{otherwise.} \end{cases}$$

$$\Pr(\text{PSU } c \in S) = \frac{C}{N_c}$$

$$\pi_{k \in S_c} = \begin{cases} \frac{m_c}{M_c}, & \text{PSU } c \in S, \\ 0, & \text{otherwise.} \end{cases}$$

Cluster sampling, cont'd

- Probability proportional to size (PPS) sampling
 - $\pi_c \propto M_c$
 - When does this make sense?
- Why implement a cluster sample?
 - The only sampling frame we have is a list of groups of observation units
 - Cost and convenience
- What happens if we ignore clustering in our sample?
 - The m_c observation units sampled in cluster c are **not** independent samples
 - → we have LESS information than m_c observations from an SRS
 - → we will UNDERESTIMATE variances and standard errors if we ignore this dependence
 - → this underestimation worsens as the correlation between outcomes from individuals in a cluster increases

Specifying the Design: svydesign

```
clus_des <- svydesign(ids = ~dnum,  
                      weights = ~pw,  
                      fpc = ~fpc,  
                      data = apiclus1)  
clus_des
```

```
## 1 - level Cluster Sampling design  
## With (15) clusters.  
## svydesign(ids = ~dnum, weights = ~pw, fpc = ~fpc, data = apiclus1)
```

Specifying the Design: `svydesign`

```
twoclus_des <- svydesign(ids = ~dnum + snum,  
                           weights = ~pw,  
                           fpc = ~fpc1 + fpc2,  
                           data = apiclus2)  
twoclus_des
```

```
## 2 - level Cluster Sampling design  
## With (40, 126) clusters.  
## svydesign(ids = ~dnum + snum, weights = ~pw, fpc = ~fpc1 + fpc2,  
##           data = apiclus2)
```

Complex surveys

Multi-stage sampling

- **Example:** DHS (among others) stratify clusters by administrative divisions × urban/rural → select women within households within clusters within strata
- **Stratified two-stage cluster sampling**
- PSUs → **secondary sampling units (SSUs)** → observation units
- One could stratify within clusters if a sampling frame necessitates (never encountered this yet)

Multi-phase sampling

- Fancy term for trying again to reach non-respondents!!
- Sub-sample (perhaps fully) your nonrespondents in attempts to get a response.

Designs of Common Surveys

- Demographic and Health Surveys (DHS)
 - <https://dhsprogram.com/>
 - Kenya DHS 2014 Final Report <https://dhsprogram.com/pubs/pdf/FR308/FR308.pdf>
- Youth Risk Behavior Survey (YRBS)
 - <https://www.cdc.gov/healthyyouth/data/yrbs/index.htm>
 - 2019 National YRBS Data User's Guide https://www.cdc.gov/healthyyouth/data/yrbs/pdf/2019/2019_National_YRBS_Data_Users_Guide.pdf
- American Community Survey (ACS)
 - <https://www.census.gov/programs-surveys/acs>
 - Design & Methodology https://www2.census.gov/programs-surveys/acs/methodology/design_and_methodology/acs_design_methodology_ch04_2014.pdf

Resources and Materials
oooo

Why survey statistics?
oooo

Survey Designs
oooooooooooooooooooo

Estimation with Survey Data
●oooooooooooooooooooo

Data Visualization
ooooooo

Etc.
oooo

Estimation with Survey Data

Horvitz-Thompson estimators

- Each individual k has their responses weighted by their **sampling weight** $w_k = \frac{1}{\pi_k}$
 - i.e. an individual with low chance of being sampled $\rightarrow \pi_k$ small $\rightarrow w_k$ big
 - w_k can be interpreted as number of individuals in the finite population that individual k 's response represents
 - **Caveat:** nonresponse
- **Average or arithmetic mean**

$$\begin{aligned}\frac{\sum_{k=1}^n y_k}{n} &\stackrel{?}{=} \frac{\sum_{k=1}^n w_k y_k}{\sum_{k=1}^n w_k} &= \frac{\sum_{k=1}^n \frac{N}{n} y_k}{\sum_{k=1}^n \frac{N}{n}} \\ &= \frac{\frac{N}{n} \sum_{k=1}^n y_k}{\frac{N}{n} \sum_{k=1}^n 1} &= \frac{N}{n} \left(\frac{\sum_{k=1}^n y_k}{\frac{N}{n} \times n} \right) \\ &= \frac{N}{n} \left(\frac{\sum_{k=1}^n y_k}{N} \right) &= \frac{\sum_{k=1}^n y_k}{n}\end{aligned}$$

Horvitz-Thompson estimators

- Each individual k has their responses weighted by their **sampling weight** $w_k = \frac{1}{\pi_k}$
 - i.e. an individual with low chance of being sampled $\rightarrow \pi_k$ small $\rightarrow w_k$ big
 - w_k can be interpreted as number of individuals in the finite population that individual k 's response represents
 - **Caveat:** nonresponse
- **Weighted average**

$$\sum_{k=1}^n w_k y_k \text{ such that } w_k \in [0, 1] \text{ and } \sum_k w_k = 1$$

Totals

- Consider a population of size N , a sample of size n , where each individual has outcome Y_k
- Y_k is **not** random, but Z_k is

$$Z_k = \begin{cases} 1, & k \in S \\ 0, & \text{otherwise.} \end{cases}$$

- Once sample taken $y_k = Y_k \times Z_k$ denotes an individual's observed response (may contain measurement error)
 - $E[y_k] = E[Y_k \times Z_k] = Y_k E[Z_k] = Y_k \times \pi_k$

Totals

- The population total of outcomes Y is

$$T = \sum_{k=1}^N Y_k$$

$$\hat{T} = \sum_{k=1}^n w_k y_k = \sum_{k=1}^n \frac{y_k}{\pi_k}$$

$$\widehat{Var}(\hat{T}) = \sum_{k,k'} \frac{y_k y_{k'}}{\pi_k \pi_{k'}} - \frac{y_k y_{k'}}{\pi_{kk'}}$$

Totals: Stratified sampling

$$\hat{T} = \sum_{h=1}^H \hat{T}_h = \sum_{h=1}^H \sum_{k=1}^{n_h} w_{hk} y_{hk},$$

$$\widehat{Var}(\hat{T}) = \sum_{h=1}^H \widehat{Var}(\hat{T}_h) = \sum_{h=1}^H \sum_{k,k'} \frac{y_{hk} y_{hk'}}{\pi_{hk} \pi_{hk'}} - \frac{y_{hk} y_{hk'}}{\pi_{hkk'}},$$

- Calculate variance in terms of each individual's difference from their respective strata total.

Totals: Cluster sampling

$$\hat{T} = \sum_{c=1}^C T_c = \sum_{c=1}^C \sum_{k=1}^{N_c} w_{ck} y_{ck} = \sum_{c=1}^C w_c \sum_{k=1}^{N_c} y_{ck},$$

- Calculate the variance in terms of each cluster total's difference from the overall population total

Totals: Stratified two-stage cluster sampling

$$\hat{T} = \sum_{h=1}^H \hat{T}_h = \sum_{h=1}^H \sum_{c_1=1}^{C_{1h}} \hat{T}_{h[c_1]}$$

$$= \sum_{h=1}^H \sum_{c_1=1}^{C_{1h}} \sum_{c_2=1}^{C_{2h}} \hat{T}_{h[c_1:c_2]} = \sum_{h=1}^H \sum_{c_1=1}^{C_{1h}} \sum_{c_2=1}^{C_{2h}} \sum_{k=1}^{n_{c_2}} w_{h[c_1:c_2]k} y_{h[c_1:c_2]k}$$

$$\widehat{Var}(\hat{T}) = \sum_{h=1}^H \widehat{Var}(\hat{T}_h).$$

- Apply methods from previous two in appropriate summation order

Totals: svytotal

```
?svytotal
svytotal(~enroll, design = srs_des)

##           total      SE
## enroll 3621074 169520

svytotal(~enroll, design = strsrs_des)

##           total      SE
## enroll 3687178 114642

svytotal(~enroll, design = twoclus_des, na.rm = TRUE)

##           total      SE
## enroll 2639273 799638
```

Means

- The population mean of outcomes Y is

$$\bar{Y} = \frac{\sum_{k=1}^N Y_k}{N}$$

$$\hat{Y} = \frac{\sum_{k=1}^n w_k y_k}{N} = \frac{1}{N} \sum_{k=1}^n \frac{y_k}{\pi_k}$$

$$\widehat{Var}(\hat{Y}) = \frac{\widehat{Var}(\hat{T})}{N^2}$$

$$SRS \left(1 - \frac{n}{N}\right) \times \frac{1}{n} \sum_{k=1}^n (y_k - \bar{y})$$

Means: svymean

```
?svymean  
svymean(~enroll, design = srs_des)
```

```
##           mean      SE  
## enroll 584.61 27.368
```

```
svymean(~enroll, design = strsrs_des)
```

```
##           mean      SE  
## enroll 595.28 18.509
```

```
svymean(~enroll, design = twoclus_des, na.rm = TRUE)
```

```
##           mean      SE  
## enroll 526.26 80.341
```

Proportions

- The population mean of binary outcomes Y or **prevalence** is

$$P = \frac{\sum_{k=1}^N Y_k}{N}$$

$$\hat{P} = \frac{\sum_{k=1}^n w_k y_k}{N} = \frac{1}{N} \sum_{k=1}^n \frac{y_k}{\pi_k}$$

$$\widehat{Var}(\hat{P}) = \frac{(\hat{P}(1 - \hat{P}))}{N}$$

Proportions: svyciprop

```
confint(svymean(~sch.wide, design = srs_des))
```

```
##                   2.5 %    97.5 %
## sch.wideNo  0.1319288 0.2380712
## sch.wideYes 0.7619288 0.8680712
```

```
svyciprop(~sch.wide, design = srs_des)
```

```
##                   2.5% 97.5%
## sch.wide 0.815 0.756 0.863
```

Proportions: svyciprop

```
confint(svymean(~sch.wide, design = strsrs_des))
```

```
##                   2.5 %    97.5 %
## sch.wideNo  0.1243371 0.2197669
## sch.wideYes 0.7802331 0.8756629
```

```
svyciprop(~sch.wide, design = strsrs_des)
```

```
##                   2.5% 97.5%
## sch.wide 0.828 0.775 0.871
```

Ratio Estimation

- What if we don't know N or don't have the finite population corrections?

$$\widehat{\bar{Y}} = \frac{\widehat{T}}{\widehat{N}} = \frac{\sum_k w_k y_k}{\sum_k w_k}$$

- Now what is $\widehat{Var}(\widehat{\bar{Y}})$? survey uses Taylor linearization.
- What if we have some other variable X that we measured in our survey and know population totals for?

$$\widehat{T_Y} = \widehat{T_Y} \frac{T_X}{\widehat{T_X}} = \sum_k w_k y_k \times \frac{T_X}{\sum_k w_k x_k}$$

- If we're over(under)estimating T_X , then maybe we're over(under)estimating T_Y

Ratio Estimation: svyratio

```
srs_des_nofpc <- svydesign(ids = ~1, weights = ~pw,  
                           data = apisrs)
```

```
svymean(~enroll, design = srs_des)
```

```
##           mean      SE  
## enroll 584.61 27.368
```

```
svymean(~enroll, design = srs_des_nofpc)
```

```
##           mean      SE  
## enroll 584.61 27.821
```

```
strsrs_des_nofpc <- svydesign(ids = ~1,
                                strata = ~stype,
                                weights = ~pw,
                                data = apistrat)

svymean(~enroll, design = strsrs_des)
```

```
##           mean      SE
## enroll 595.28 18.509
```

```
svymean(~enroll, design = strsrs_des_nofpc)
```

```
##           mean      SE
## enroll 595.28 18.941
```

```
srs_des_nofpc <- update(srs_des_nofpc, counter = 1)
svyratio(numerator = ~enroll, denominator = ~counter, design = srs_des_nofpc)

## Ratio estimator: svyratio.survey.design2(numerator = ~enroll, denominator = ~counter, design = srs_des_nofpc)
## Ratios=
##       counter
## enroll 584.61
## SEs=
##       counter
## enroll 27.82121

svymean(~enroll, design = srs_des_nofpc)

##       mean      SE
## enroll 584.61 27.821
```

Small Area Estimation: svyby

```
svyby(~enroll, by = ~stype, design = srs_des, svytotal)
```

```
##    stype    enroll        se
## E      E 1849900.0 99738.62
## H      H  890666.2 187717.67
## M      M  880508.1 151805.23
```

```
svyby(~enroll, by = ~stype, design = strsrs_des, svytotal)
```

```
##    stype    enroll        se
## E      E 1842584.3 72581.33
## H      H  997128.5 69239.40
## M      M  847464.7 55502.96
```

General Linear Models: svyglm

```
srs_mod <- svyglm(sch.wide ~ ell + meals + mobility,  
                    design = srs_des, family = quasibinomial())  
strsrs_mod <- svyglm(sch.wide ~ ell + meals + mobility,  
                    design = strsrs_des, family = quasibinomial())
```

General Linear Models: svyglm

```
coefs_table <- data.frame(SRS_Coef = coef(srs_mod),  
                           SRS_SE = SE(srs_mod),  
                           StrSRS_Coef = coef(strsrs_mod),  
                           StrSRS_SE = SE(strsrs_mod))  
round(coefs_table, digits = 3)  
  
##           SRS_Coef  SRS_SE StrSRS_Coef  StrSRS_SE  
## (Intercept)  1.744   0.456   0.836     0.456  
## ell        -0.022   0.011   -0.002     0.013  
## meals       0.011   0.009   -0.003     0.009  
## mobility   -0.015   0.022    0.061     0.032
```

Post-stratification

What if we don't have a **probability survey**? OR What if our ideal stratification scheme was not possible to implement given our sampling frame?

```
pop.types <- apipop %>%
  group_by(stype) %>%
  summarize(Freq = n())
```

```
srs_post <- postStratify(srs_des, ~stype, pop.types)
```

Resources and Materials
oooo

Why survey statistics?
oooo

Survey Designs
oooooooooooooooooooo

Estimation with Survey Data
oooooooooooooooooooo

Data Visualization
●oooooooooooo

Etc.
oooo

Data Visualization

Functions in the survey package

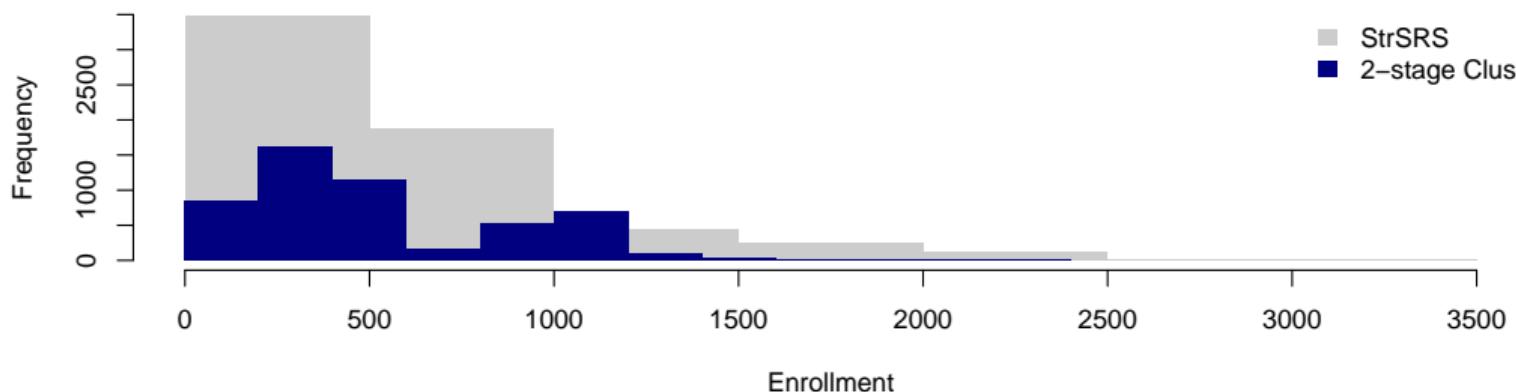
Instead of plotting the data in your sample, the following visualizations give you a sense of the data in your sample AND their relative contribution to the population.

- Histograms (`svyhist`) of weighted outcomes
- Boxplots (`svyboxplot`) of weighted outcomes (by group, if desired)
- Scatterplots (`svyplot`) of two variables showing relative weight of observations (e.g. with transparency or character size)
- Scatterplots by group (`svycoplot`) of two variables conditional on the value of other variables (e.g. binary measure of exposure) using the hexagonal binning method available in `svyplot`

Histograms: svyhist

```
svyhist(~enroll, design = strsrs_des,
        main = "", xlab = "Enrollment",
        probability = FALSE, col = 'grey80', border = FALSE)
svyhist(~enroll, design = twoclus_des,
        main = "", xlab = "Enrollment",
        probability = FALSE, col = 'navy', border = FALSE, add = TRUE)
legend('topright', bty = 'n',
       fill = c("grey80", "navy"), border = FALSE,
       legend = c("StrSRS", "2-stage Clus"))
```

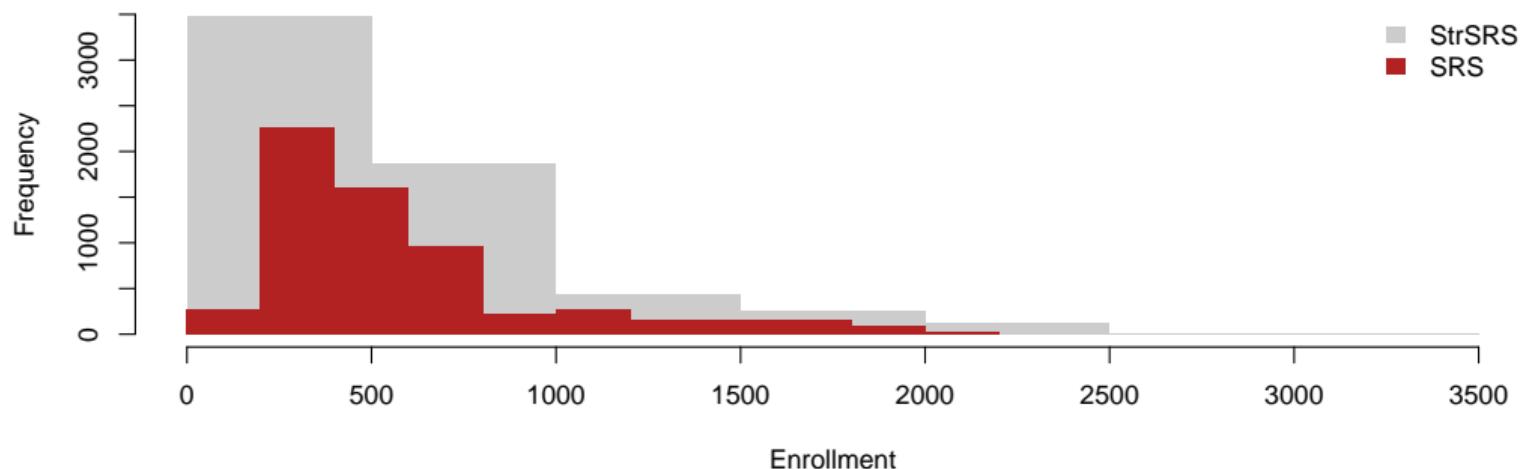
Histograms: `svyhist`



Histograms: svyhist

```
svyhist(~enroll, design = strsrs_des,
        main = "", xlab = "Enrollment",
        probability = FALSE, col = 'grey80', border = FALSE)
svyhist(~enroll, design = srs_des,
        main = "", xlab = "Enrollment",
        probability = FALSE, col = 'firebrick', border = FALSE,
        add = TRUE)
legend('topright', bty = 'n',
       fill = c("grey80", "firebrick"), border = FALSE,
       legend = c("StrSRS", "SRS"))
```

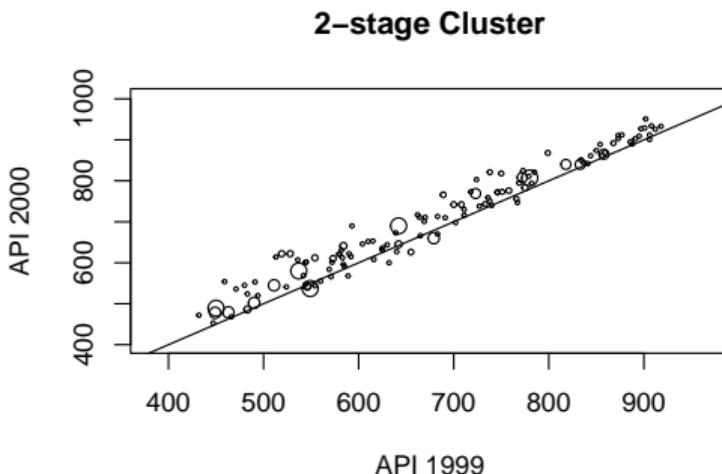
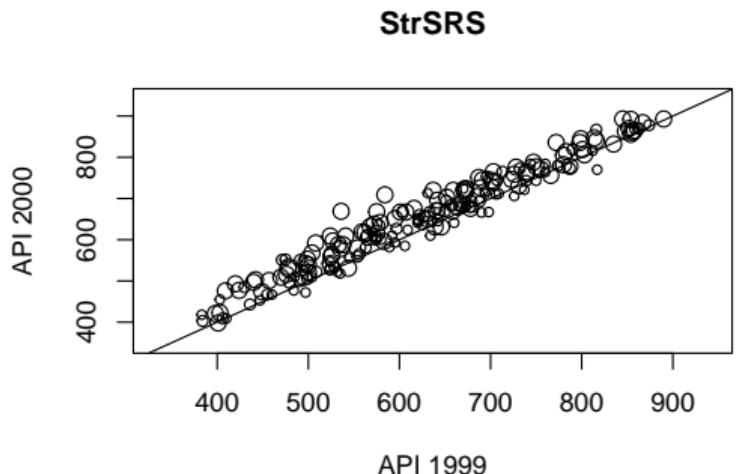
Histograms: svyhist



Scatterplots: svyplot

```
par(mfrow = c(1,2))
svyplot(api00~api99, design=strsrs_des, style="bubble",
        col = "firebrick",
        xlab = "API 1999", ylab = "API 2000", main = "StrSRS")
abline(0,1)
svyplot(api00~api99, design=twoclus_des, style="bubble",
        col = "firebrick",
        xlab = "API 1999", ylab = "API 2000", main = "2-stage Cluster")
abline(0,1)
```

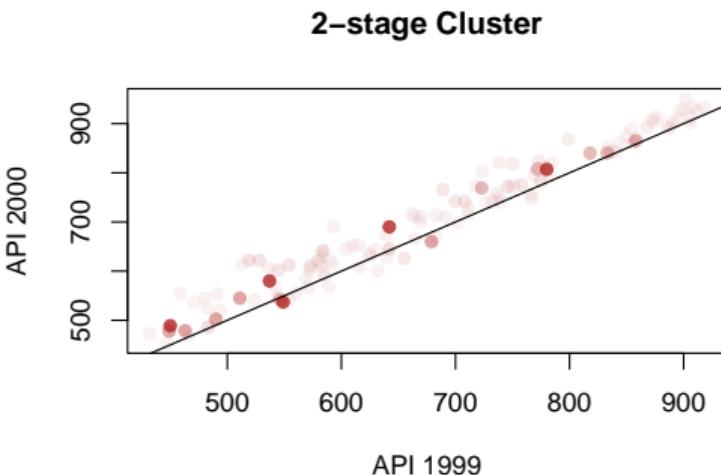
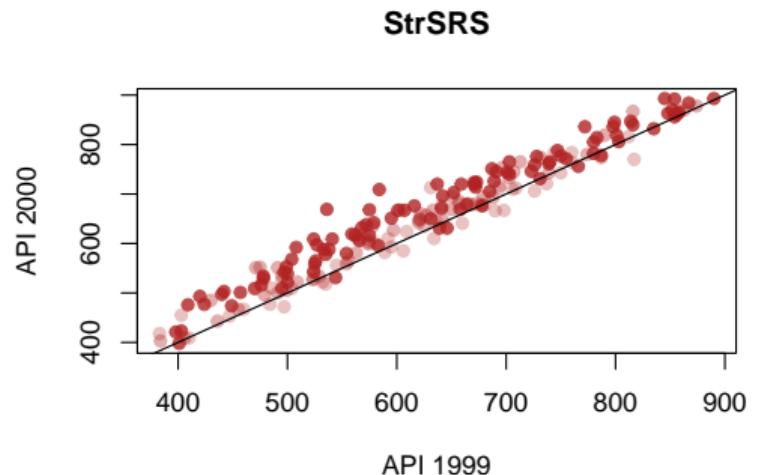
Scatterplots: `svyplot`



Scatterplots: svyplot

```
par(mfrow = c(1,2))
svyplot(api00~api99, design=strsrs_des, style="transparent",
        basecol = "firebrick", pch=19,
        xlab = "API 1999", ylab = "API 2000", main = "StrSRS")
abline(0,1)
svyplot(api00~api99, design=twoclus_des, style="transparent",
        basecol = "firebrick", pch=19,
        xlab = "API 1999", ylab = "API 2000", main = "2-stage Cluster")
abline(0,1)
```

Scatterplots: `svyplot`



Resources and Materials
oooo

Why survey statistics?
oooo

Survey Designs
oooooooooooooooooooo

Estimation with Survey Data
oooooooooooooooooooo

Data Visualization
ooooooo

Etc.
●ooo

Etc.

What didn't we cover?

- Post-stratification and raking (`survey::postStratify`)
- Replicate weights (`survey::svrepdesign`)
- Non-response
- Multi-phase sampling
- Model-based estimation

Specific Question: Contrasts

- Contrasts are just linear combinations of random variables where the weights add up to 0, e.g. averages and differences.

$$E[aX + bY] = aE[X] + bE[Y]$$

$$E[X - Y] = 1 \times E[X] + (-1) \times E[Y]$$

$$Var(aX + bY) = a^2 Var(X) + b^2 Var(Y) + 2abCov(X, Y)$$

$$Var(X - Y) = 1^2 \times Var(X) + (-1)^2 \times Var(Y) + 2(1)(-1) \times Cov(X, Y)$$

Specific Question: Contrasts

```
cont_total <- svytotal(~api00+api99, strata_des)
svycontrast(cont_total, list(diff=c(1,-1)))
```

```
##      contrast      SE
## diff    203736 12705
```

```
vcov(cont_total)
```

```
##            api00      api99
## api00 3396439386 3521991247
## api99 3521991247 3808949720
```

```
sqrt(vcov(cont_total)[1,1] + vcov(cont_total)[2,2] +
  2*1*(-1)*vcov(cont_total)[1,2])
```

```
## [1] 12704.59
```