

Sampling II

LPO 9951 | Fall 2020

PURPOSE

In the last lecture, we discussed a number of ways to properly estimate the means and variances of complex survey designs. In this lecture, we'll discuss how to use Stata's internal `svy` commands and various variance estimation methods to more easily and correctly estimate what we want.

```
. capture log close                                // closes any logs, should they be open

. set linesize 90

. log using "sampling_part2.log", replace           // open new log
-----
name: <unnamed>
log: /Users/doylewr/lpo_prac/lessons/s1-06-sampling/sampling_part2.log
log type: text
opened on: 6 Oct 2021, 10:59:45
```

Complex survey designs: Cluster sampling and stratification

In the NCES surveys you'll be using this semester, the designers combined a design that includes multistage cluster sampling with stratification. In ECLS, for example, the designers designated counties as *PSUs*. They next stratified the sample by creating strata that combined census region with msa status, percent minority, and per capita income. They then randomly selected schools within each *PSU* (schools were the *SSUs*) and then randomly selected kindergarteners within each school (students were the *TSUs*). They then created two strata for each school with Asian and Pacific Islander students in one stratum and all other students in the other. Students were randomly sampled within this second stratum. The target number of children per school was 24.

Weights in complex survey designs such as the one employed with ECLS are calculated via the same that we discussed in the last lecture. Nothing changes except for the layers of complexity. The good news, however, is that we as researchers don't have to compute the weights ourselves. Instead, we can use information provided by the survey makers.

The *PSUs* that are provided by NCES are what is known as "analysis *PSUs*". They aren't the identifier for the actual school or student. Instead, they are allocated within strata (many times 2 *PSU* per strata). Strata themselves may be analysis strata, that is, not the same strata that were used to run the survey. Oftentimes, this is done in service of further protecting the anonymity of participants. As far as your analyses go, the end result is the same, but sometimes this can be a source of confusion.

Variance estimation in complex survey designs

There are four common options for estimating variance in complex survey designs:

1. Taylor series linearized estimates
2. Balanced repeated replication (BRR) estimates
3. Jackknife estimates
4. Bootstrap estimates

Remember that these are all estimates: you cannot directly compute the variance of quantities of interest from complex surveys. Instead, you must use one of these techniques, with trade-offs for each. We'll be using a couple of datasets for this lesson:

- *nhanes*, which is a health survey conducted using a complex survey design that comes with a variety of weights
- *nmihs_bs*, which is a survey of births that comes with bootstrap replicate weights

Let's start with the *nhanes* dataset from which we'd like to get average height weight and age for the US population. First, let's get the naive estimate:

```
. clear all                                // clear memory

. set more off                             // turn off annoying "__more__" feature

. webuse nhanes2f, clear

. preserve

. keep sampl stratid psuid

. save nhanes2f_s, replace
file nhanes2f_s.dta saved

. restore

. mean age height weight
```

Mean estimation Number of obs = 10,337

		Mean	Std. err.	[95% conf. interval]
-----+				
age		47.5637	.1693381	47.23177 47.89564
height		167.6512	.0950124	167.465 167.8375
weight		71.90088	.1510277	71.60484 72.19692

We can also take a look at the sampling design, particularly the designation of strata and *PSUs*:

```
. tab stratid psuid
```

Stratum identifier , 1-32	Primary sampling unit		Total
	PSU 1	PSU 2	
1	215	165	380
2	118	67	185
3	199	149	348
4	231	229	460
5	147	105	252
6	167	131	298
7	270	206	476
8	179	158	337
9	143	100	243
10	143	119	262
11	120	155	275
12	170	144	314
13	154	188	342
14	205	200	405
15	189	191	380
16	177	159	336
17	180	213	393
18	144	215	359
20	158	125	283
21	102	111	213
22	173	128	301
23	182	158	340
24	202	232	434
25	139	115	254
26	132	129	261
27	144	139	283
28	135	163	298
29	287	215	502
30	166	199	365
31	143	165	308
32	239	211	450
Total	5,353	4,984	10,337

It's important to remember that these are *analysis* PSUs and strata, not the exact ones that were used in the survey design itself. Essentially the original

strata are reassigned names that allow for deidentification, and then psus are assigned within the strata.

We can use the weights supplied with *nhanes* to get accurate estimates of the means, but the variance estimates will be off:

```
. mean age height weight [pw = finalwgt]
```

Mean estimation Number of obs = 10,337

		Mean	Std. err.	[95% conf. interval]	
	age	42.23732	.1617236	41.92031	42.55433
	height	168.4625	.1139787	168.2391	168.686
	weight	71.90869	.1802768	71.55532	72.26207

svyset and svy: <command>

To aid in the analysis of complex survey data, Stata has incorporated the **svyset** command and the **svy:** prefix, with its suite of commands. With **svyset**, you can set the *PSU* (and *SSU* and *TSU* if applicable), the weights, and the type of variance estimation along with the variance weights (if applicable). Once set, most Stata estimation commands such as **mean** can be combined with **svy:** in order to produce correct estimates.

Variance estimators

Taylor series linearized estimates

Taylor series linearized estimates are based on the general strategy of Taylor series estimation, which is used to linearize a non-linear function in order to describe the function in question. In this case, a Taylor series is used to approximate the function, and the variance of the result is the estimate of the variance.

The basic intuition behind a linearized estimate is that the variance in a complex survey will be a nonlinear function of the set of variances calculated within each stratum. We can calculate these, then use the first derivative of the function that would calculate the actual variance as a first order approximation of the actual variance. This works well enough in practice. To do this, you absolutely must have multiple *PSUs* in each stratum so you can calculate variance within each stratum.

This is the most common method and is used as the default by Stata. You must, however, have within-stratum variance among *PSUs* for this to work, which means that you must have at least two *PSUs* per stratum. This lonely PSU

problem is common and difficult to deal with. We'll return the lonely PSU later.

To set up a dataset to use linearized estimates in Stata, we use the `svyset` command:

```
. svyset psuid [pweight = finalwgt], strata(stratid)

Sampling weights: finalwgt
                  VCE: linearized
                  Single unit: missing
                  Strata 1: stratid
                  Sampling unit 1: psuid
                  FPC 1: <zero>
```

Now that we've set the data, every time we want estimates that reflect the sampling design, we use the `svy: <command>` format:

```
. svy: mean age height weight
(running mean on estimation sample)
```

Survey: Mean estimation

```
Number of strata = 31          Number of obs   =      10,337
Number of PSUs   = 62          Population size = 117,023,659
                                Design df        =          31
```

		Linearized		
		Mean	std. err.	[95% conf. interval]
age		42.23732	.3034412	41.61844 42.85619
height		168.4625	.1471709	168.1624 168.7627
weight		71.90869	.1672315	71.56762 72.24976

As you can see, the parameter estimates (means) are exactly the same as using the weighted sample, but the standard errors are quite different: nearly twice as large for age, but actually smaller for weight.

Balanced repeated replication (BRR) estimates

In a balanced repeated replication (BRR) design, the quantity of interests is estimated repeatedly by using half the sample at a time. In a survey which is designed with BRR in mind, each sampling stratum contains two *PSUs*. BRR proceeds by estimating the quantity of interest from one of the *PSUs* within each stratum. For H strata, $2H$ replications are done, and the variance of the quantity of interest across these strata forms the basis for the estimate.

BRR weights are usually supplied with a survey. These weights result in appropriate half samples being formed across strata. BRR weights should generally be used when the sample was designed with them in mind, and not elsewhere. This can be a serious complication when survey data are subset.

To get variance estimates using BRR in Stata, you either need to have a set of replicate weights set up or you need to create a set of balanced replicates yourself. If the data has BRR weights estimates can be obtained as follows:

```
. webuse nhanes2brr, clear

. svyset [pw=finalwgt], brrweight(brr*) vce(brr)

Sampling weights: finalwgt
                  VCE: brr
                  MSE: off
                  BRR weights: brr_1 .. brr_32
                  Single unit: missing
                  Strata 1: <one>
Sampling unit 1: <observations>
                  FPC 1: <zero>

. svy: mean age height weight
(running mean on estimation sample)

BRR replications (32)
-----+----- 1 -----+----- 2 -----+----- 3 -----+----- 4 -----+----- 5
.....

Survey: Mean estimation          Number of obs   =       10,351
                                Population size = 117,157,513
                                Replications   =         32
                                Design df      =         31

-----+-----
              |              BRR
              |              Mean  std. err.    [95% conf. interval]
-----+-----+-----
              age |    42.25264    .3013406    41.63805    42.86723
              height |   168.4599    .14663    168.1608    168.7589
              weight |   71.90064    .1656452    71.5628    72.23847
-----+-----
```

The `brrweight` option specified which variables constitute the brr weights, while the `vce` option says that variance should be calculated using the balanced repeated replication approach.

It's helpful to take a look at how BRR weights are related to PSUs and strata

```
. merge 1:1 sampl using nhanes2f_s
```

Result	Number of obs	
Not matched	14	
from master	14	(_merge==1)
from using	0	(_merge==2)
Matched	10,337	(_merge==3)

```
. order sampl finalwgt psu stratid brr*
```

```
. gsort stratid psuid
```

Jackknife estimates

The Jackknife is a general strategy for variance estimation, so named by Tukey because of its general usefulness. The strategy for creating a jackknifed estimate is to delete every observation save one, then estimate the quantity of interest. This is repeated for every single observation in the dataset. The variance of every estimate computed provides an estimate of the variance for the quantity of interest.

In a complex sample, this is done by *PSUs*, deleting each *PSU* one at a time and re-weighting the observations within the stratum, then calculating the parameter of interest. The variance of these parameters estimates is the within-stratum variance estimate. The within stratum variances calculated this way are then averaged across strata to give the final variance estimate.

The jackknife is best used when Taylor series estimation cannot be done, for instance in the case of lonely *PSUs*.

```
. webuse nhanes2jknife, clear
```

In Stata, the command is:

```
. svyset [pweight = finalwgt], jkrweight(jkw_*) vce(jackknife)
```

```

Sampling weights: finalwgt
                  VCE: jackknife
                  MSE: off
Jackknife weights: jkw_1 .. jkw_62
Single unit: missing
Strata 1: <one>
Sampling unit 1: <observations>
FPC 1: <zero>

```

Now we can compare the naive estimates with the `svyset` estimates:

```
. mean age weight height
```

```
Mean estimation                                Number of obs = 10,351
```

		Mean	Std. err.	[95% conf. interval]	
	age	47.57965	.1692044	47.24798	47.91133
	weight	71.89752	.1509381	71.60165	72.19339
	height	167.6509	.0949079	167.4648	167.8369

```
. svy: mean age weight height
(running mean on estimation sample)
```

```
Jackknife replications (62)
```

```
-----+----- 1 -----+----- 2 -----+----- 3 -----+----- 4 -----+----- 5
..... 50
.....
```

```
Survey: Mean estimation
```

```
Number of strata = 31                Number of obs   =      10,351
                                     Population size = 117,157,513
                                     Replications   =         62
                                     Design df      =         31
```

		Mean	Jackknife std. err.	[95% conf. interval]	
	age	42.25264	.3026765	41.63533	42.86995
	weight	71.90064	.1654453	71.56321	72.23806
	height	168.4599	.1466141	168.1609	168.7589

```
. merge 1:1 sampl using nhanes2f_s
```

Result	Number of obs
Not matched	14
from master	14 (_merge==1)
from using	0 (_merge==2)
Matched	10,337 (_merge==3)


```

. order sampl finalwgt psu stratid jkw_*

. gsort stratid psuid

. browse sampl finalwgt psuid stratid jkw_*

```

Bootstrap estimates

The bootstrap is a more general method than the jackknife. Bootstrapping involves repeatedly resampling within the sample itself and generating estimates of the quantity of interest. The variance of these replications (usually many, many replications) provides an estimate of the total variance. In NCES surveys, within stratum bootstrapping can be used, with the sum of the variances obtained used as an estimate of the population variance. Bootstrapping is an accurate, but computationally intense method of variance estimation.

As with the jackknife, bootstrapping must be accomplished by deleting each *PSU* within the stratum one at a time, re-weighting, calculating the estimate, then calculating the bootstrap variance estimate from the compiled samples.

```

. webuse nmihs_bs, clear

. svyset idnum [pweight = finwgt], vce(bootstrap) bsrweight(bsrw*)

```

```

Sampling weights: finwgt
                  VCE: bootstrap
                  MSE: off
Bootstrap weights: bsrw1 .. bsrw1000
                  Single unit: missing
                  Strata 1: <one>
Sampling unit 1: idnum
                  FPC 1: <zero>

```

```

. gen birthwgtlbs = birthwgt * 0.0022046
(7 missing values generated)

```

```

. mean birthwgtlbs

```

Mean estimation Number of obs = 9,946

	Mean	Std. err.	[95% conf. interval]	
birthwgtlbs	6.272294	.0217405	6.229678	6.31491

```
. svy: mean birthwgtlbs
(running mean on estimation sample)
```

```
Bootstrap replications (1,000)
```

```
-----+----- 1 -----+----- 2 -----+----- 3 -----+----- 4 -----+----- 5
..... 50
..... 100
..... 150
..... 200
..... 250
..... 300
..... 350
..... 400
..... 450
..... 500
..... 550
..... 600
..... 650
..... 700
..... 750
..... 800
..... 850
..... 900
..... 950
..... 1,000
```

```
Survey: Mean estimation          Number of obs   =      9,946
                                Population size = 3,895,562
                                Replications   =      1,000
```

	Observed	Bootstrap	Normal-based	
	mean	std. err.	[95% conf. interval]	
birthwgtlbs	7.39743	.0143754	7.369255	7.425606

```
. gsort finwgt
```

```
. browse idnum finwgt bsrw*
```

Lonely *PSUs*

The most common problem that students have with complex surveys is what is known as “lonely *PSUs*.” When you subset the data, you may very well end up with a sample that does not have multiple *PSUs* per stratum. There are several

options for what do i n this case:

- Eliminate the offending data by dropping strata with singleton *PSUs*. This is a terrible idea.
- Reassign the *PSU* to a neighboring stratum. This is okay, but you must have a reason why you’re doing this.
- Assign a variance to the stratum with a singleton *PSU*. This could be the average of the variance across the other strata. This process is also known as “scaling” and generally is okat, but you should take a look at how different this stratum is fr om the others before proceeding.

The `svyset` command includes three possible options for dealing with loney *PSUs*. Based on the above, I recommend you use the `singleunit(scaled)` command, but with caution and full knowledge of the implications for your estimates.

Design Effects

Design effects are pretty old-school and shouldn’t be used. That said, you will see these used in some older articles. These were used because most statistical programming languages weren’t able to compute variance estimates from complex surveys up until about 2010. As a patchwork solution, the survey provider would calculate standard errors for some commonly used estimates from some common variables and look at how much bigger they were than naive estimates. The ratio between these would be averaged and called a design effect. For instance, if standard errors from a Taylor series linearized estimate were on average 1.3 times as big as naive standard errors then the design effect was 1.3. Do not use this approach, for hopefully obvious reasons.

Using variance estimation from different surveys

```
. use ../../data/plans.dta, clear

. svyset psu [pw=f1pnlwt],strata(strat_id)

Sampling weights: f1pnlwt
                  VCE: linearized
                  Single unit: missing
                  Strata 1: strat_id
Sampling unit 1: psu
                  FPC 1: <zero>

. mean byncls2m

Mean estimation                                Number of obs = 16,160

-----+-----
|              Mean   Std. err.   [95% conf. interval]
```

```
-----+-----
      bynels2m |    44.44327    .1187556    44.21049    44.67604
-----+-----
```

```
. svy: mean bynels2m
(running mean on estimation sample)
```

Survey: Mean estimation

```
Number of strata = 361          Number of obs   =    16,160
Number of PSUs   = 751          Population size = 3,388,462
                                   Design df         =      390
```

```
-----+-----
              |          Linearized
              |          Mean    std. err.    [95% conf. interval]
-----+-----
      bynels2m |    44.74391    .2618191    44.22915    45.25866
-----+-----
```

```
. use ../../data/hsls_belong.dta, clear
```

```
. renvars *, lower
```

```
. svyset [pw=w1parent], brr(w1parent???) vce(brr)
```

```
Sampling weights: w1parent
                  VCE: brr
                  MSE: off
BRR weights: w1parent001 .. w1parent200
Single unit: missing
Strata 1: <one>
Sampling unit 1: <observations>
FPC 1: <zero>
```

```
. prop x3hscompstat
```

Proportion estimation Number of obs = 808

```
_prop_1: x3hscompstat = High school diploma
_prop_2: x3hscompstat = GED, certificate of attendance,
_prop_3: x3hscompstat = Dropped out
_prop_4: x3hscompstat = Still enrolled
_prop_5: x3hscompstat = Status unknown
```

```
-----+-----
```

	Proportion	Std. err.	Logit [95% conf. interval]	
x3hscompstat				
_prop_1	.7376238	.0154765	.7061348	.7668526
_prop_2	.0433168	.0071615	.031245	.0597649
_prop_3	.0482673	.0075401	.0354439	.0654156
_prop_4	.0680693	.0088606	.0526049	.0876591
_prop_5	.1027228	.0106805	.0835711	.1256616

```
. svy: prop x3hscompstat
(running proportion on estimation sample)
```

```
BRR replications (200)
```

```
-----+----- 1 ----+----- 2 ----+----- 3 ----+----- 4 ----+----- 5
..... 50
..... 100
..... 150
..... 200
```

```
Survey: Proportion estimation      Number of obs   =      808
                                   Population size = 218,060.05
                                   Replications    =      200
                                   Design df       =      199
```

```
_prop_1: x3hscompstat = High school diploma
_prop_2: x3hscompstat = GED, certificate of attendance,
_prop_3: x3hscompstat = Dropped out
_prop_4: x3hscompstat = Still enrolled
_prop_5: x3hscompstat = Status unknown
```

	Proportion	BRR std. err.	Logit [95% conf. interval]	
x3hscompstat				
_prop_1	.7019053	.0249524	.6504961	.7486743
_prop_2	.0385162	.0101322	.0228223	.0642919
_prop_3	.0535511	.0134007	.0325002	.0870105
_prop_4	.0705174	.0125091	.0494954	.0995331
_prop_5	.1355101	.0183957	.1031456	.1760363

```
. use ../../data/nhes_example.dta, clear
```

```
. replace dpcolor=. if inlist(dpcolor, -8 ,-7, -6 ,-5 ,-4 ,-3 ,-2, -1)
(1,847 real changes made, 1,847 to missing)
```

```
. svyset epsu [pw=fewt] ,strat(estratum) singleunit(scaled)
```

```
Sampling weights: fewt
                  VCE: linearized
                  Single unit: scaled
                  Strata 1: estratum
                  Sampling unit 1: epsu
                  FPC 1: <zero>
```

```
. prop dpcolor
```

```
Proportion estimation                      Number of obs = 3,997
```

```
_prop_1: dpcolor = 1 No
_prop_2: dpcolor = 2 Yes, some of them
_prop_3: dpcolor = 3 Yes, all of them
```

				Logit
	Proportion	Std. err.	[95% conf. interval]	
dpcolor				
_prop_1	.0542907	.0035841	.0476777	.0617615
_prop_2	.1788842	.0060621	.1673063	.1910794
_prop_3	.7668251	.0066884	.7534565	.7796808

```
. svy: prop dpcolor
(running proportion on estimation sample)
```

```
Survey: Proportion estimation
```

```
Number of strata =      3          Number of obs   =      3,997
Number of PSUs   = 3,997          Population size = 13,693,230
                                   Design df        =      3,994
```

```
_prop_1: dpcolor = 1 No
_prop_2: dpcolor = 2 Yes, some of them
_prop_3: dpcolor = 3 Yes, all of them
```

		Linearized	Logit
	Proportion	std. err.	[95% conf. interval]

-----+-----				
dpcolor				
_prop_1		.0639356	.0056819	.0536596
_prop_2		.2238697	.0100379	.2048063
_prop_3		.7121947	.0106956	.6907792
				.732701

```
. rename fewt finalwgt
```

```
. svyset epsu [pw=finalwgt] , vce(brr) brrweight(fewt*)
```

```
Sampling weights: finalwgt
                  VCE: brr
                  MSE: off
                  BRR weights: fewt1 .. fewt80
                  Single unit: missing
                  Strata 1: <one>
                  Sampling unit 1: epsu
                  FPC 1: <zero>
```

```
. prop dpcolor
```

```
Proportion estimation                                Number of obs = 3,997
```

```
  _prop_1: dpcolor = 1 No
  _prop_2: dpcolor = 2 Yes, some of them
  _prop_3: dpcolor = 3 Yes, all of them
```

-----+-----				
				Logit
		Proportion	Std. err.	[95% conf. interval]
-----+-----				
dpcolor				
_prop_1		.0542907	.0035841	.0476777
_prop_2		.1788842	.0060621	.1673063
_prop_3		.7668251	.0066884	.7534565
				.7796808

```
. svy: prop dpcolor
(running proportion on estimation sample)
```

```
BRR replications (80)
```

```
-----+ 1 -----+ 2 -----+ 3 -----+ 4 -----+ 5
.....
.....
```

50

```

Survey: Proportion estimation      Number of obs   =      3,997
                                   Population size = 13,693,230
                                   Replications    =        80
                                   Design df       =        79

```

```

_prop_1: dpcolor = 1 No
_prop_2: dpcolor = 2 Yes, some of them
_prop_3: dpcolor = 3 Yes, all of them

```

		BRR	Logit	
	Proportion	std. err.	[95% conf. interval]	
dpcolor				
_prop_1	.0639356	.0005379	.0628732	.0650147
_prop_2	.2238697	.0009822	.2219208	.2258307
_prop_3	.7121947	.0010494	.7101015	.7142789

```

. log close
  name: <unnamed>
  log: /Users/doylewr/lpo_prac/lessons/s1-06-sampling/sampling_part2.log
  log type: text
  closed on: 6 Oct 2021, 11:00:28

```

```

. exit

```