

svy jackknife — Jackknife estimation for survey data

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Description

`svy jackknife` performs jackknife estimation of specified statistics (or expressions) for a Stata command or a user-written program. The command is executed once for each replicate using sampling weights that are adjusted according to the jackknife methodology. Any Stata estimation command listed in [\[SVY\] svy estimation](#) may be used with `svy jackknife`. User-written programs that meet the requirements in [\[P\] program properties](#) may also be used.

Quick start

Estimate population mean of `v1` using jackknife standard-error estimates with sampling weight `wvar1` and sampling units identified by `su1`

```
svyset su1 [pweight = wvar1]
svy jackknife _b: mean v1
```

Same as above, but with jackknife replication weights in variables with prefix `rwvar`

```
svyset [pweight = wvar1], vce(jackknife) jkrweight(rwvar*)
svy: mean v1
```

Jackknife standard error of the difference between the means of `v2` and `v3` using either `svyset` command above

```
svy jackknife (_b[v2]-_b[v3]): mean v2 v3
```

Same as above, but name the result `diff` and save results from each replication to `mydata.dta`

```
svy jackknife diff=(_b[v2]-_b[v3]), saving(mydata): mean v2 v3
```

Note: Any estimation command meeting the requirements specified in the *Description* may be substituted for `mean` in the examples above.

Menu

Statistics > Survey data analysis > Resampling > Jackknife estimation

Syntax

`svy jackknife` *exp_list* [*, svy_options jackknife_options eform_option*] : *command*

<i>svy_options</i>	Description
<i>if/in</i>	
<code>subpop</code> ([<i>varname</i>] [<i>if</i>])	identify a subpopulation
Reporting	
<code>level</code> (#)	set confidence level; default is <code>level(95)</code>
<code>noheader</code>	suppress table header
<code>nolegend</code>	suppress table legend
<code>noadjust</code>	do not adjust model Wald statistic
<code>nocnsreport</code>	do not display constraints
<code>display_options</code>	control columns and column formats, row spacing, line width, display of omitted variables and base and empty cells, and factor-variable labeling
<code>coeflegend</code>	display legend instead of statistics

`coeflegend` is not shown in the dialog boxes for estimation commands.

<i>jackknife_options</i>	Description
Main	
<code>eclass</code>	number of observations is in <code>e(N)</code>
<code>rclass</code>	number of observations is in <code>r(N)</code>
<code>n</code> (<i>exp</i>)	specify <i>exp</i> that evaluates to number of observations used
Options	
<code>saving</code> (<i>filename</i> [<i>, ...</i>])	save results to <i>filename</i> ; save statistics in double precision; save results to <i>filename</i> every # replications
<code>keep</code>	keep pseudovalues
<code>mse</code>	use MSE formula for variance
Reporting	
<code>verbose</code>	display the full table legend
<code>nodots</code>	suppress replication dots
<code>dots</code> (#)	display dots every # replications
<code>noisily</code>	display any output from <i>command</i>
<code>trace</code>	trace <i>command</i>
<code>title</code> (<i>text</i>)	use <i>text</i> as title for jackknife results
Advanced	
<code>nodrop</code>	do not drop observations
<code>reject</code> (<i>exp</i>)	identify invalid results
<code>dof</code> (#)	design degrees of freedom

svy requires that the survey design variables be identified using `svyset`; see [SVY] [svyset](#).

`command` defines the statistical command to be executed. The `by` prefix cannot be part of `command`.

`collect` is allowed; see [U] [11.1.10 Prefix commands](#).

See [U] [20 Estimation and postestimation commands](#) for more capabilities of estimation commands.

Warning: Using `if` or `in` restrictions will often not produce correct variance estimates for subpopulations. To compute estimates for subpopulations, use the `subpop()` option.

`exp_list` specifies the statistics to be collected from the execution of `command`. `exp_list` is required unless `command` has the `svyj` program property, in which case `exp_list` defaults to `_b`; see [P] [program properties](#). The expressions in `exp_list` are assumed to conform to the following:

```
exp_list contains      (name: elist)
                      elist
                      eexp
elist contains        newvarname = (exp)
                      (exp)
eexp is               specname
                      [eqno]specname
specname is          _b
                    _b[]
                    _se
                    _se[]
eqno is              ##
                    name
```

`exp` is a standard Stata expression; see [U] [13 Functions and expressions](#).

Distinguish between `[]`, which are to be typed, and `[][]`, which indicate optional arguments.

Options

`svy_options`; see [SVY] [svy](#).

Main

`eclass`, `rclass`, and `n(exp)` specify where `command` stores the number of observations on which it based the calculated results. We strongly advise you to specify one of these options.

`eclass` specifies that `command` store the number of observations in `e(N)`.

`rclass` specifies that `command` store the number of observations in `r(N)`.

`n(exp)` allows you to specify an expression that evaluates to the number of observations used. Specifying `n(r(N))` is equivalent to specifying the `rclass` option. Specifying `n(e(N))` is equivalent to specifying the `eclass` option. If `command` stores the number of observations in `r(N1)`, specify `n(r(N1))`.

If you specify none of these options, `svy jackknife` will assume `eclass` or `rclass` depending upon which of `e(N)` and `r(N)` is not missing (in that order). If both `e(N)` and `r(N)` are missing, `svy jackknife` assumes that all observations in the dataset contribute to the calculated result. If that assumption is incorrect, then the reported standard errors will be incorrect. For instance, say that you specify

```
. svy jackknife coef=_b[x2]: myreg y x1 x2 x3
```

where `myreg` uses $e(n)$ instead of $e(N)$ to identify the number of observations used in calculations. Further assume that observation 42 in the dataset has `x3` equal to missing. The 42nd observation plays no role in obtaining the estimates, but `svy jackknife` has no way of knowing that and will use the wrong N . If, on the other hand, you specify

```
. svy jackknife coef=_b[x2], n(e(n)): myreg y x1 x2 x3
```

Then `svy jackknife` will notice that observation 42 plays no role. The `n(e(n))` option is specified because `myreg` is an estimation command, but it stores the number of observations used in $e(n)$ (instead of the standard $e(N)$). When `svy jackknife` runs the regression omitting the 42nd observation, `svy jackknife` will observe that $e(n)$ has the same value as when `svy jackknife` previously ran the regression by using all the observations. Thus `svy jackknife` will know that `myreg` did not use the observation.

Options

`saving(filename [, suboptions])` creates a Stata data file (`.dta` file) consisting of (for each statistic in `exp_list`) a variable containing the replicates.

`double` specifies that the results for each replication be saved as `doubles`, meaning 8-byte reals.

By default, they are saved as `floats`, meaning 4-byte reals. This option may be used without the `saving()` option to compute the variance estimates by using double precision.

`every(#)` specifies that results be written to disk every `#`th replication. `every()` should be specified in conjunction with `saving()` only when `command` takes a long time for each replication.

This will allow recovery of partial results should some other software crash your computer. See [P] [postfile](#).

`replace` specifies that `filename` be overwritten if it exists. This option does not appear in the dialog box.

`keep` specifies that new variables be added to the dataset containing the pseudovalues of the requested statistics. For instance, if you typed

```
. svy jackknife coef=_b[x2], eclass keep: regress y x1 x2 x3
```

Then the new variable `coef` would be added to the dataset containing the pseudovalues for `_b[x2]`. Let b be defined as the value of `_b[x2]` when all observations are used to fit the model, and let $b(j)$ be the value when the j th observation is omitted. The pseudovalues are defined as

$$\text{pseudovalue}_j = N \times \{b - b(j)\} + b(j)$$

where N is the number of observations used to produce b .

`keep` implies the `nodrop` option.

`mse` specifies that `svy jackknife` compute the variance by using deviations of the replicates from the observed value of the statistics based on the entire dataset. By default, `svy jackknife` computes the variance by using deviations of the pseudovalues from their mean.

Reporting

`verbose` requests that the full table legend be displayed.

`nodots` and `dots(#)` specify whether to display replication dots. By default, one dot character is printed for each successful replication. An “x” is displayed if `command` returns an error, “e” is displayed if at least one value in `exp_list` is missing, “n” is displayed if the sample size is not correct, and a yellow “s” is displayed if the dropped sampling unit is outside the subpopulation sample.

`nodots` suppresses display of the replication dots.

`dots(#)` displays dots every # replications. `dots(0)` is a synonym for `nodots`.

`noisily` requests that any output from *command* be displayed. This option implies the `nodots` option.

`trace` causes a trace of the execution of *command* to be displayed. This option implies the `noisily` option.

`title(text)` specifies a title to be displayed above the table of jackknife results; the default title is “Jackknife results”.

eform_option; see [R] [eform_option](#). This option is ignored if *exp_list* is not `_b`.

Advanced

`nodrop` prevents observations outside `e(sample)` and the `if` and `in` qualifiers from being dropped before the data are resampled.

`reject(exp)` identifies an expression that indicates when results should be rejected. When *exp* is true, the resulting values are reset to missing values.

`dof(#)` specifies the design degrees of freedom, overriding the default calculation, $df = N_{psu} - N_{strata}$.

Remarks and examples

stata.com

The jackknife is

- an alternative, first-order unbiased estimator for a statistic;
- a data-dependent way to calculate the standard error of the statistic and to obtain significance levels and confidence intervals; and
- a way of producing measures called pseudovalues for each observation, reflecting the observation’s influence on the overall statistic.

The idea behind the simplest form of the jackknife—the one implemented in [R] [jackknife](#)—is to repeatedly calculate the statistic in question, each time omitting just one of the dataset’s observations. Assume that our statistic of interest is the sample mean. Let y_j be the j th observation of our data on some measurement y , where $j = 1, \dots, N$ and N is the sample size. If \bar{y} is the sample mean of y using the entire dataset and $\bar{y}_{(j)}$ is the mean when the j th observation is omitted, then

$$\bar{y} = \frac{(N-1)\bar{y}_{(j)} + y_j}{N}$$

Solving for y_j , we obtain

$$y_j = N\bar{y} - (N-1)\bar{y}_{(j)}$$

These are the pseudovalues that `svy: jackknife` calculates. To move this discussion beyond the sample mean, let $\hat{\theta}$ be the value of our statistic (not necessarily the sample mean) using the entire dataset, and let $\hat{\theta}_{(j)}$ be the computed value of our statistic with the j th observation omitted. The pseudovalue for the j th observation is

$$\hat{\theta}_j^* = N\hat{\theta} - (N-1)\hat{\theta}_{(j)}$$

The mean of the pseudovalues is the alternative, first-order unbiased estimator mentioned above, and the standard error of the mean of the pseudovalues is an estimator for the standard error of $\hat{\theta}$ (Tukey 1958, Shao and Tu 1995).

When the jackknife is applied to survey data, primary sampling units (PSUs) are omitted instead of observations, N is the number of PSUs instead of the sample size, and the sampling weights are adjusted owing to omitting PSUs; see [SVY] **Variance estimation** for more details.

Because of privacy concerns, many public survey datasets contain jackknife replication-weight variables instead of variables containing information on the PSUs and strata. These replication-weight variables are the adjusted sampling weights, and there is one replication-weight variable for each omitted PSU.

► Example 1: Jackknife with information on PSUs and strata

Suppose that we were interested in a measure of association between the weight and height of individuals in the population represented by the NHANES II data (McDowell et al. 1981). To measure the association, we will use the slope estimate from a linear regression of `weight` on `height`. We also use `svy jackknife` to estimate the variance of the slope.

```
. use https://www.stata-press.com/data/r18/nhanes2
. svyset
Sampling weights: finalwgt
                  VCE: linearized
                  Single unit: missing
                  Strata 1: strata
Sampling unit 1: psu
                  FPC 1: <zero>

. svy jackknife slope = _b[height]: regress weight height
(running regress on estimation sample)
Jackknife replications (62): .....10.....20.....30.....40.....
> ..50.....60.. done

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

Command: regress weight height
slope: _b[height]
n(): e(N)
```

	Coefficient	Jackknife std. err.	t	P> t	[95% conf. interval]	
slope	.8014753	.0160281	50.00	0.000	.7687858	.8341648

◀

► Example 2: Jackknife replicate-weight variables

`nhanes2jknife.dta` is a privacy-conscious dataset equivalent to `nhanes2.dta`; all the variables and values remain, except that `strata` and `psu` are replaced with jackknife replicate-weight variables. The replicate-weight variables are already `svyset`, and the default method for variance estimation is `vce(jackknife)`.

```

. use https://www.stata-press.com/data/r18/nhanes2jknife
. svyset
  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>

```

Here we perform the same analysis as in the [previous example](#), using jackknife replication weights.

```

. svy jackknife slope = _b[height], nodots: regress weight height
Linear regression
Number of strata = 31
Number of obs   = 10,351
Population size = 117,157,513
Replications    = 62
Design df      = 31

Command: regress weight height
slope: _b[height]

```

	Coefficient	Jackknife std. err.	t	P> t	[95% conf. interval]	
slope	.8014753	.0160281	50.00	0.000	.7687858	.8341648

The `mse` option causes `svy jackknife` to use the MSE form of the jackknife variance estimator. This variance estimator will tend to be larger than the previous because of the addition of the familiar squared bias term in the MSE; see [\[SVY\] Variance estimation](#) for more details. The header for the column of standard errors in the table of results is `Jknife *` for the jackknife variance estimator, which uses the MSE formula.

```

. svy jackknife slope = _b[height], mse nodots: regress weight height
Linear regression
Number of strata = 31
Number of obs   = 10,351
Population size = 117,157,513
Replications    = 62
Design df      = 31

Command: regress weight height
slope: _b[height]

```

	Coefficient	Jknife * std. err.	t	P> t	[95% conf. interval]	
slope	.8014753	.0160284	50.00	0.000	.7687852	.8341654

□ Technical note

When the `svy jackknife` prefix is used with a user-defined program and when the expression list is `_b`, `svy jackknife` calls

```
set coeftabresults off
```

before entering the replication loop to prevent Stata from performing unnecessary calculations. This means that, provided option `noisily` is not specified, estimation commands will not build or post the coefficient table matrix `r(table)`.

If your program calls an estimation command and needs `r(table)` to exist to perform properly, then your program will need to call

```
set coeftabresults on
```

before calling other estimation commands.

□

Stored results

In addition to the results documented in [SVY] `svy`, `svy jackknife` stores the following in `e()`:

Scalars

<code>e(N_reps)</code>	number of replications
<code>e(N_misreps)</code>	number of replications with missing values
<code>e(k_exp)</code>	number of standard expressions
<code>e(k_eexp)</code>	number of <code>_b/_se</code> expressions
<code>e(k_extra)</code>	number of extra estimates added to <code>_b</code>

Macros

<code>e(cmdname)</code>	command name from <i>command</i>
<code>e(cmd)</code>	same as <code>e(cmdname)</code> or <code>jackknife</code>
<code>e(vce)</code>	<code>jackknife</code>
<code>e(exp#)</code>	<i>#</i> th expression
<code>e(jkrweight)</code>	<code>jkweight()</code> variable list

Matrices

<code>e(b_jk)</code>	jackknife means
<code>e(V)</code>	jackknife variance estimates

When *exp_list* is `_b`, `svy jackknife` will also carry forward most of the results already in `e()` from *command*.

Methods and formulas

See [SVY] **Variance estimation** for details regarding jackknife variance estimation.

References

- McDowell, A., A. Engel, J. T. Massey, and K. Maurer. 1981. Plan and operation of the Second National Health and Nutrition Examination Survey, 1976–1980. *Vital and Health Statistics* 1(15): 1–144.
- Shao, J., and D. Tu. 1995. *The Jackknife and Bootstrap*. New York: Springer.
- Tukey, J. W. 1958. Bias and confidence in not-quite large samples. Abstract in *Annals of Mathematical Statistics* 29: 614. <https://doi.org/10.1214/aoms/1177706647>.

Also see

- [SVY] [svy postestimation](#) — Postestimation tools for svy
- [SVY] [svy bootstrap](#) — Bootstrap for survey data
- [SVY] [svy brr](#) — Balanced repeated replication for survey data
- [SVY] [svy sdr](#) — Successive difference replication for survey data
- [SVY] [Calibration](#) — Calibration for survey data
- [SVY] [Poststratification](#) — Poststratification for survey data
- [SVY] [Subpopulation estimation](#) — Subpopulation estimation for survey data
- [SVY] [Variance estimation](#) — Variance estimation for survey data
- [R] [jackknife](#) — Jackknife estimation
- [U] [20 Estimation and postestimation commands](#)

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