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Description

`total` produces estimates of totals, along with standard errors.

Quick start

Total of continuous variable `v1`

```
total v1
```

Same as above, but restrict estimation to observations where `catvar = 1`

```
total v1 if catvar==1
```

Same as above, but using `svyset` data

```
svy, subpop(if catvar==1): total v1
```

Total of `v1` for each level of `catvar`

```
total v1, over(catvar)
```

With jackknife standard errors

```
total v1, vce(jackknife)
```

Menu

Statistics > Summaries, tables, and tests > Summary and descriptive statistics > Totals

Syntax

total *varlist* [*if*] [*in*] [*weight*] [, *options*]

<i>options</i>	Description
if/in/over	
over(<i>varlist</i> _o)	group over subpopulations defined by <i>varlist</i> _o
SE/Cluster	
vce(<i>vcetype</i>)	<i>vcetype</i> may be analytic, <u>cluster</u> <i>clustvar</i> , <u>bootstrap</u> , or <u>jackknife</u>
Reporting	
<u>level</u> (#)	set confidence level; default is level(95)
<u>noheader</u>	suppress table header
<i>display_options</i>	control column formats, line width, display of omitted variables and base and empty cells, and factor-variable labeling
<u>coeflegend</u>	display legend instead of statistics

varlist may contain factor variables; see [U] 11.4.3 Factor variables.

bayesboot, bootstrap, collect, jackknife, mi estimate, rolling, statsby, and svy are allowed; see [U] 11.1.10 Prefix commands.

vce(bootstrap) and vce(jackknife) are not allowed with the mi estimate prefix.

Weights are not allowed with the bootstrap prefix; see [R] bootstrap.

vce() and weights are not allowed with the svy prefix; see [SVY] svy.

fweights, iweights, and pweights are allowed; see [U] 11.1.6 weight.

coeflegend does not appear in the dialog box.

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

Options

if/in/over

over(*varlist*_o) specifies that estimates be computed for multiple subpopulations, which are identified by the different values of the variables in *varlist*_o. Only numeric, nonnegative, integer-valued variables are allowed in over(*varlist*_o).

SE/Cluster

vce(*vcetype*) specifies the type of standard error reported, which includes types that are derived from asymptotic theory (analytic), that allow for intragroup correlation (cluster *clustvar*), and that use bootstrap or jackknife methods (bootstrap, jackknife); see [R] vce_option.

vce(analytic), the default, uses the analytically derived variance estimator associated with the sample total.

Reporting

`level(#)`; see [R] [Estimation options](#).

`noheader` prevents the table header from being displayed.

display_options: `noomitted`, `vsquish`, `noemptycells`, `baselevels`, `allbaselevels`, `nofvlabel`, `fvwrap(#)`, `fvwrapon(style)`, `cformat(%fmt)`, and `nolstretch`; see [R] [Estimation options](#).

The following option is available with `total` but is not shown in the dialog box:

`coeflegend`; see [R] [Estimation options](#).

Remarks and examples

► Example 1

Suppose that we collected data on incidence of heart attacks. The variable `heartatk` indicates whether a person ever had a heart attack (1 means yes; 0 means no). We can then estimate the total number of persons who have had heart attacks for each sex in the population represented by the data we collected.

```
. use https://www.stata-press.com/data/r19/total
(Fictional incidence of heart-attack data)
```

```
. total heartatk [pw=swgt], over(sex)
```

Total estimation

Number of obs = 4,946

	Total	Std. err.	[95% conf. interval]	
c.heartatk@sex				
Male	944559	104372.3	739943	1149175
Female	581590	82855.59	419156.3	744023.7

Stored results

`total` stores the following in `e()`:

Scalars

<code>e(N)</code>	number of observations
<code>e(N_over)</code>	number of subpopulations
<code>e(N_clust)</code>	number of clusters
<code>e(k_eq)</code>	number of equations in <code>e(b)</code>
<code>e(df_r)</code>	sample degrees of freedom
<code>e(rank)</code>	rank of <code>e(V)</code>

Macros

<code>e(cmd)</code>	<code>total</code>
<code>e(cmdline)</code>	command as typed
<code>e(varlist)</code>	<i>varlist</i>
<code>e(wtype)</code>	weight type
<code>e(wexp)</code>	weight expression
<code>e(title)</code>	title in estimation output
<code>e(clustvar)</code>	name of cluster variable
<code>e(over)</code>	<i>varlist</i> from <code>over()</code>
<code>e(vce)</code>	<i>vcetype</i> specified in <code>vce()</code>
<code>e(vcetype)</code>	title used to label Std. err.
<code>e(properties)</code>	<code>b V</code>
<code>e(estat_cmd)</code>	program used to implement <code>estat</code>
<code>e(marginsnotok)</code>	predictions disallowed by margins

Matrices

<code>e(b)</code>	vector of total estimates
<code>e(V)</code>	(co)variance estimates
<code>e(_N)</code>	vector of numbers of nonmissing observations
<code>e(error)</code>	error code corresponding to <code>e(b)</code>

Functions

<code>e(sample)</code>	marks estimation sample
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In addition to the above, the following is stored in `r()`:

Matrices

<code>r(table)</code>	matrix containing the coefficients with their standard errors, test statistics, <i>p</i> -values, and confidence intervals
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Note that results stored in `r()` are updated when the command is replayed and will be replaced when any `r-class` command is run after the estimation command.

Methods and formulas

Methods and formulas are presented under the following headings:

The total estimator
Survey data
The survey total estimator
The poststratified total estimator
Subpopulation estimation

The total estimator

Let y denote the variable on which to calculate the total and $y_j, j = 1, \dots, n$, denote an individual observation on y . Let w_j be the frequency weight (or `iweight` or `pweight`), and if no weight is specified, define $w_j = 1$ for all j . The sum of the weights is an estimate of the population size:

$$\widehat{N} = \sum_{j=1}^n w_j$$

If the population values of y are denoted by $Y_j, j = 1, \dots, N$, the associated population total is

$$Y = \sum_{j=1}^N Y_j = N\bar{y}$$

where \bar{y} is the population mean. The total is estimated as

$$\hat{Y} = \widehat{N}\bar{y}$$

The variance estimator for the total is

$$\widehat{V}(\hat{Y}) = \widehat{N}^2 \widehat{V}(\bar{y})$$

where $\widehat{V}(\bar{y})$ is the variance estimator for the mean; see [\[R\] mean](#). The standard error of the total is the square root of the variance.

If x, x_j, \bar{x} , and \widehat{X} are similarly defined for another variable (observed jointly with y), the covariance estimator between \widehat{X} and \hat{Y} is

$$\widehat{\text{Cov}}(\widehat{X}, \hat{Y}) = \widehat{N}^2 \widehat{\text{Cov}}(\bar{x}, \bar{y})$$

where $\widehat{\text{Cov}}(\bar{x}, \bar{y})$ is the covariance estimator between two means; see [\[R\] mean](#).

Survey data

See [\[SVY\] Variance estimation](#) and [\[SVY\] Poststratification](#) for discussions that provide background information for the following formulas.

The survey total estimator

Let Y_j be a survey item for the j th individual in the population, where $j = 1, \dots, M$ and M is the size of the population. The associated population total for the item of interest is

$$Y = \sum_{j=1}^M Y_j$$

Let y_j be the survey item for the j th sampled individual from the population, where $j = 1, \dots, m$ and m is the number of observations in the sample.

The estimator \hat{Y} for the population total Y is

$$\hat{Y} = \sum_{j=1}^m w_j y_j$$

where w_j is a sampling weight. The estimator for the number of individuals in the population is

$$\hat{M} = \sum_{j=1}^m w_j$$

The score variable for the total estimator is the variable itself,

$$z_j(\hat{Y}) = y_j$$

The poststratified total estimator

Let P_k denote the set of sampled observations that belong to poststratum k , and define $I_{P_k}(j)$ to indicate if the j th observation is a member of poststratum k , where $k = 1, \dots, L_P$ and L_P is the number of poststrata. Also, let M_k denote the population size for poststratum k . P_k and M_k are identified by specifying the `poststrata()` and `postweight()` options on `svyset`; see [\[SVY\] svyset](#).

The estimator for the poststratified total is

$$\hat{Y}^P = \sum_{k=1}^{L_P} \frac{M_k}{\hat{M}_k} \hat{Y}_k = \sum_{k=1}^{L_P} \frac{M_k}{\hat{M}_k} \sum_{j=1}^m I_{P_k}(j) w_j y_j$$

where

$$\hat{M}_k = \sum_{j=1}^m I_{P_k}(j) w_j$$

The score variable for the poststratified total is

$$z_j(\hat{Y}^P) = \sum_{k=1}^{L_P} I_{P_k}(j) \frac{M_k}{\hat{M}_k} \left(y_j - \frac{\hat{Y}_k}{\hat{M}_k} \right)$$

Subpopulation estimation

Let S denote the set of sampled observations that belong to the subpopulation of interest, and define $I_S(j)$ to indicate if the j th observation falls within the subpopulation.

The estimator for the subpopulation total is

$$\hat{Y}^S = \sum_{j=1}^m I_S(j) w_j y_j$$

and its score variable is

$$z_j(\hat{Y}^S) = I_S(j) y_j$$

The estimator for the poststratified subpopulation total is

$$\hat{Y}^{PS} = \sum_{k=1}^{L_P} \frac{M_k}{\widehat{M}_k} \hat{Y}_k^S = \sum_{k=1}^{L_P} \frac{M_k}{\widehat{M}_k} \sum_{j=1}^{m_k} I_{P_k}(j) I_S(j) w_j y_j$$

and its score variable is

$$z_j(\hat{Y}^{PS}) = \sum_{k=1}^{L_P} I_{P_k}(j) \frac{M_k}{\widehat{M}_k} \left\{ I_S(j) y_j - \frac{\hat{Y}_k^S}{\widehat{M}_k} \right\}$$

References

Cochran, W. G. 1977. *Sampling Techniques*. 3rd ed. New York: Wiley.

Stuart, A., and J. K. Ord. 1994. *Distribution Theory*. Vol. 1 of *Kendall's Advanced Theory of Statistics*, 6th ed. London: Arnold.

Also see

[R] **total postestimation** — Postestimation tools for total

[R] **mean** — Estimate means

[R] **proportion** — Estimate proportions

[R] **ratio** — Estimate ratios

[MI] **Estimation** — Estimation commands for use with mi estimate

[SVY] **Direct standardization** — Direct standardization of means, proportions, and ratios

[SVY] **Poststratification** — Poststratification for survey data

[SVY] **Subpopulation estimation** — Subpopulation estimation for survey data

[SVY] **svy estimation** — Estimation commands for survey data

[SVY] **Variance estimation** — Variance estimation for survey data

[U] **20 Estimation and postestimation commands**

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