Title

total — Estimate total

Description Options References Quick start Remarks and examples Also see Menu Stored results Syntax Methods and formulas

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]

options	Description	
if/in/over		
over(<i>varlist</i> _o)	group over subpopulations defined by varlist _o	
SE/Cluster		
vce(vcetype)	<i>vcetype</i> may be analytic, <u>cl</u> uster <i>clustvar</i> , <u>boot</u> strap, or <u>jack</u> knife	
Reporting		
<u>l</u> evel(#)	set confidence level; default is level(95)	
<u>noh</u> eader	suppress table header	
display_options	control column formats, line width, display of omitted variables and base and empty cells, and factor-variable labeling	
<u>coefl</u> egend	display legend instead of statistics	
varlist may contain factor varia	ables; see [U] 11.4.3 Factor variables.	

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

fif/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

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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/r18/total
(Fictional incidence of heart-attack data)
. total heartatk [pw=swgt], over(sex)
Total estimation
                                           Number of obs = 4,946
                      Totol
                              Std orr
                                            [95% conf interval]
```

	IUCAL	stu. err.	[95% com: mtervar]
c.heartatk@sex Male Female	944559 581590	104372.3	739943 1149175 419156.3 744023.7
romute	001000	02000.00	11010010 11102011

Stored results

total stores the following in e():

Scalars	
e(N)	number of observations
e(N_over)	number of subpopulations
e(N_clust)	number of clusters
e(k_eq)	number of equations in e(b)
e(df_r)	sample degrees of freedom
e(rank)	rank of e(V)
Macros	
e(cmd)	total
e(cmdline)	command as typed
e(varlist)	varlist
e(wtype)	weight type
e(wexp)	weight expression
e(title)	title in estimation output
e(clustvar)	name of cluster variable
e(over)	varlist from over()
e(vce)	vcetype specified in vce()
e(vcetype)	title used to label Std. err.
e(properties)	b V
e(estat_cmd)	program used to implement estat
e(marginsnotok)	predictions disallowed by margins

Matrices	
e(b)	vector of total estimates
e(V)	(co)variance estimates
e(_N)	vector of numbers of nonmissing observations
e(error)	error code corresponding to e(b)
Functions	
e(sample)	marks estimation sample

In addition to the above, the following is stored in r():

```
Matrices
```

r(table)

matrix containing the coefficients with their standard errors, test statistics, *p*-values, and confidence intervals

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, ..., 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 $Y_i, j = 1, ..., N$, the associated population total is

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

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

$$Y = N\overline{i}$$

The variance estimator for the total is

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

where $\widehat{V}(\overline{y})$ 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, \overline{x} , and \widehat{X} are similarly defined for another variable (observed jointly with y), the covariance estimator between \widehat{X} and \widehat{Y} is

$$\widehat{\operatorname{Cov}}(\widehat{X},\widehat{Y}) = \widehat{N}^2 \widehat{\operatorname{Cov}}(\overline{x},\overline{y})$$

where $\widehat{\text{Cov}}(\overline{x}, \overline{y})$ 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, ..., 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, ..., m and *m* is the number of observations in the sample.

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

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

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

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

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

$$z_j(Y) = y_j$$

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 $k = 1, \ldots, L_P$ 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

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

where

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

The score variable for the poststratified total is

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

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

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

and its score variable is

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

The estimator for the poststratified subpopulation total is

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

and its score variable is

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

References

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

Stuart, A., and J. K. Ord. 1994. Kendall's Advanced Theory of Statistics: Distribution Theory, Vol. 1. 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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