

**bstat** — Report bootstrap results

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## Syntax

*Bootstrap statistics from variables*

```
bstat [varlist] [if] [in] [, options]
```

*Bootstrap statistics from file*

```
bstat [namelist] [using filename] [if] [in] [, options]
```

<i>options</i>	Description
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### Main

<u>s</u> tat( <i>vector</i> )	observed values for each statistic
accel( <i>vector</i> )	acceleration values for each statistic
<u>t</u> ies	adjust BC/BCa confidence intervals for ties
mse	use MSE formula for variance estimation

### Reporting

<u>l</u> evel(#)	set confidence level; default is level(95)
n(#)	# of observations from which bootstrap samples were taken
notable	suppress table of results
<u>n</u> oheader	suppress table header
<u>n</u> olegend	suppress table legend
<u>v</u> erbose	display the full table legend
<u>t</u> itle( <i>text</i> )	use <i>text</i> as title for bootstrap results
<i>display_options</i>	control column formats and line width

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

## Menu

Statistics > Resampling > Report bootstrap results

## Description

**bstat** is a programmer's command that computes and displays estimation results from bootstrap statistics.

For each variable in *varlist* (the default is all variables), then **bstat** computes a covariance matrix, estimates bias, and constructs several different confidence intervals (CIs). The following CIs are constructed by **bstat**:

1. Normal CIs (using the normal approximation)
2. Percentile CIs
3. Bias-corrected (BC) CIs
4. Bias-corrected and accelerated ( $BC_a$ ) CIs (optional)

`estat bootstrap` displays a table of one or more of the above confidence intervals; see [\[R\] bootstrap postestimation](#).

If there are bootstrap estimation results in `e()`, `bstat` replays them. If given the `using` modifier, `bstat` uses the data in *filename* to compute the bootstrap statistics while preserving the data currently in memory. Otherwise, `bstat` uses the data in memory to compute the bootstrap statistics.

The following options may be used to replay estimation results from `bstat`:

`level(#)` `notable` `noheader` `nolegend` `verbose` `title(text)`

For all other options and the qualifiers `using`, `if`, and `in`, `bstat` requires a bootstrap dataset.

## Options

### Main

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`stat(vector)` specifies the observed value of each statistic (that is, the value of the statistic using the original dataset).

`accel(vector)` specifies the acceleration of each statistic, which is used to construct  $BC_a$  CIs.

`ties` specifies that `bstat` adjust for ties in the replicate values when computing the median bias used to construct BC and  $BC_a$  CIs.

`mse` specifies that `bstat` compute the variance by using deviations of the replicates from the observed value of the statistics. By default, `bstat` computes the variance by using deviations from the average of the replicates.

### Reporting

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

`n(#)` specifies the number of observations from which bootstrap samples were taken. This value is used in no calculations but improves the table header when this information is not saved in the bootstrap dataset.

`notable` suppresses the display of the output table.

`noheader` suppresses the display of the table header. This option implies `nolegend`.

`nolegend` suppresses the display of the table legend.

`verbose` specifies that the full table legend be displayed. By default, coefficients and standard errors are not displayed.

`title(text)` specifies a title to be displayed above the table of bootstrap results; the default title is `Bootstrap results`.

`display_options`: `cformat(%fmt)`, `pformat(%fmt)`, `sformat(%fmt)`, and `nolstretch`; see [\[R\] estimation options](#).

## Remarks and examples

Remarks are presented under the following headings:

*Bootstrap datasets*  
*Creating a bootstrap dataset*

### Bootstrap datasets

Although `bstat` allows you to specify the observed value and acceleration of each bootstrap statistic via the `stat()` and `accel()` options, programmers may be interested in what `bstat` uses when these options are not supplied.

When working from a bootstrap dataset, `bstat` first checks the data characteristics (see [P] `char`) that it understands:

`_dta[bs_version]` identifies the version of the bootstrap dataset. This characteristic may be empty (not defined), 2, or 3; otherwise, `bstat` will quit and display an error message. This version tells `bstat` which other characteristics to look for in the bootstrap dataset.

`bstat` uses the following characteristics from version 3 bootstrap datasets:

```
_dta[N]
_dta[N_strata]
_dta[N_cluster]
_dta[command]
varname[observed]
varname[acceleration]
varname[expression]
```

`bstat` uses the following characteristics from version 2 bootstrap datasets:

```
_dta[N]
_dta[N_strata]
_dta[N_cluster]
varname[observed]
varname[acceleration]
```

An empty bootstrap dataset version implies that the dataset was created by the `bstrap` command in a version of Stata earlier than Stata 8. Here `bstat` expects `varname[bstrap]` to contain the observed value of the statistic identified by `varname` (`varname[observed]` in version 2). All other characteristics are ignored.

`_dta[N]` is the number of observations in the observed dataset. This characteristic may be overruled by specifying the `n()` option.

`_dta[N_strata]` is the number of strata in the observed dataset.

`_dta[N_cluster]` is the number of clusters in the observed dataset.

`_dta[command]` is the command used to compute the observed values of the statistics.

`varname[observed]` is the observed value of the statistic identified by `varname`. To specify a different value, use the `stat()` option.

`varname[acceleration]` is the estimate of acceleration for the statistic identified by `varname`. To specify a different value, use the `accel()` option.

`varname[expression]` is the expression or label that describes the statistic identified by `varname`.

## Creating a bootstrap dataset

Suppose that we are interested in obtaining bootstrap statistics by resampling the residuals from a regression (which is not possible with the `bootstrap` command). After loading some data, we run a regression, save some results relevant to the `bstat` command, and save the residuals in a new variable, `res`.

```
. use http://www.stata-press.com/data/r13/auto
(1978 Automobile Data)
```

```
. regress mpg weight length
```

Source	SS	df	MS			
Model	1616.08062	2	808.040312	Number of obs = 74		
Residual	827.378835	71	11.653223	F( 2, 71) = 69.34		
Total	2443.45946	73	33.4720474	Prob > F = 0.0000		
				R-squared = 0.6614		
				Adj R-squared = 0.6519		
				Root MSE = 3.4137		

  

mpg	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
weight	-.0038515	.001586	-2.43	0.018	-.0070138	-.0006891
length	-.0795935	.0553577	-1.44	0.155	-.1899736	.0307867
_cons	47.88487	6.08787	7.87	0.000	35.746	60.02374

```
. matrix b = e(b)
```

```
. local n = e(N)
```

```
. predict res, residuals
```

We can resample the residual values in `res` by generating a random observation ID (`rid`), generate a new response variable (`y`), and run the original regression with the new response variables.

```
. set seed 54321
```

```
. generate rid = int(_N*runiform()+1)
```

```
. matrix score double y = b
```

```
. replace y = y + res[rid]
```

```
(74 real changes made)
```

```
. regress y weight length
```

Source	SS	df	MS			
Model	1773.23548	2	886.617741	Number of obs = 74		
Residual	608.747732	71	8.57391172	F( 2, 71) = 103.41		
Total	2381.98321	73	32.629907	Prob > F = 0.0000		
				R-squared = 0.7444		
				Adj R-squared = 0.7372		
				Root MSE = 2.9281		

  

y	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
weight	-.0059938	.0013604	-4.41	0.000	-.0087064	-.0032813
length	-.0127875	.0474837	-0.27	0.788	-.1074673	.0818924
_cons	42.23195	5.22194	8.09	0.000	31.8197	52.6442

Instead of programming this resampling inside a loop, it is much more convenient to write a short program and use the `simulate` command; see [R] [simulate](#). In the following, `mysim_r` requires the user to specify a coefficient vector and a residual variable. `mysim_r` then retrieves the list of predictor variables (removing `_cons` from the list), generates a new temporary response variable with the resampled residuals, and regresses the new response variable on the predictors.

```

program mysim_r
  version 13
  syntax name(name=bvector), res(varname)
  tempvar y rid
  local xvars : colnames 'bvector'
  local cons _cons
  local xvars : list xvars - cons
  matrix score double 'y' = 'bvector'
  gen long 'rid' = int(_N*runiform()) + 1
  replace 'y' = 'y' + 'res'['rid']
  regress 'y' 'xvars'
end

```

We can now give `mysim_r` a test run, but we first set the random-number seed (to reproduce results).

```

. set seed 54321
. mysim_r b, res(res)
(74 real changes made)

```

Source	SS	df	MS			
Model	1773.23548	2	886.617741	Number of obs =	74	
Residual	608.747732	71	8.57391172	F( 2, 71) =	103.41	
Total	2381.98321	73	32.629907	Prob > F =	0.0000	
				R-squared =	0.7444	
				Adj R-squared =	0.7372	
				Root MSE =	2.9281	

  

	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
weight	-.0059938	.0013604	-4.41	0.000	-.0087064	-.0032813
length	-.0127875	.0474837	-0.27	0.788	-.1074673	.0818924
_cons	42.23195	5.22194	8.09	0.000	31.8197	52.6442

Now that we have a program that will compute the results we want, we can use `simulate` to generate a bootstrap dataset and `bstat` to display the results.

```

. set seed 54321
. simulate, reps(200) nodots: mysim_r b, res(res)
  command: mysim_r b, res(res)
. bstat, stat(b) n('n')

```

	Observed Coef.	Bootstrap Std. Err.	z	P> z	Normal-based [95% Conf. Interval]	
_b_weight	-.0038515	.0015715	-2.45	0.014	-.0069316	-.0007713
_b_length	-.0795935	.0552415	-1.44	0.150	-.1878649	.0286779
_b_cons	47.88487	6.150069	7.79	0.000	35.83096	59.93879

Finally, we see that `simulate` created some of the data characteristics recognized by `bstat`. All we need to do is correctly specify the version of the bootstrap dataset, and `bstat` will automatically use the relevant data characteristics.

```

. char list
  _dta[seed]:                X681014b5c43f462544a474abacbdd93d00042842
  _dta[command]:            mysim_r b, res(res)
  _b_weight[is_eexp]:      1
  _b_weight[colname]:      weight
  _b_weight[coleq]:        -
  _b_weight[expression]:   _b[weight]
  _b_length[is_eexp]:      1
  _b_length[colname]:      length
  _b_length[coleq]:        -
  _b_length[expression]:   _b[length]
  _b_cons[is_eexp]:        1
  _b_cons[colname]:        _cons
  _b_cons[coleq]:          -
  _b_cons[expression]:     _b[_cons]

```

```
. char _dta[bs_version] 3
```

```
. bstat, stat(b) n('n')
```

```

Bootstrap results                Number of obs    =        74
                                Replications      =       200

```

```
command: mysim_r b, res(res)
```

	Observed Coef.	Bootstrap Std. Err.	z	P> z	Normal-based [95% Conf. Interval]	
weight	-.0038515	.0015715	-2.45	0.014	-.0069316	-.0007713
length	-.0795935	.0552415	-1.44	0.150	-.1878649	.0286779
_cons	47.88487	6.150069	7.79	0.000	35.83096	59.93879

See [Poi \(2004\)](#) for another example of residual resampling.

## Stored results

bstat stores the following in `e()`:

### Scalars

<code>e(N)</code>	sample size
<code>e(N_reps)</code>	number of complete replications
<code>e(N_misreps)</code>	number of incomplete replications
<code>e(N_strata)</code>	number of strata
<code>e(N_clust)</code>	number of clusters
<code>e(k_aux)</code>	number of auxiliary parameters
<code>e(k_eq)</code>	number of equations in <code>e(b)</code>
<code>e(k_exp)</code>	number of standard expressions
<code>e(k_eexp)</code>	number of extended expressions (i.e., <code>_b</code> )
<code>e(k_extra)</code>	number of extra equations beyond the original ones from <code>e(b)</code>
<code>e(level)</code>	confidence level for bootstrap CIs
<code>e(bs_version)</code>	version for bootstrap results
<code>e(rank)</code>	rank of <code>e(V)</code>

### Macros

<code>e(cmd)</code>	<code>bstat</code>
<code>e(command)</code>	from <code>_dta[command]</code>
<code>e(cmdline)</code>	command as typed
<code>e(title)</code>	title in estimation output
<code>e(exp#)</code>	expression for the #th statistic
<code>e(prefix)</code>	<code>bootstrap</code>
<code>e(ties)</code>	ties, if specified
<code>e(mse)</code>	mse, if specified
<code>e(vce)</code>	<code>bootstrap</code>
<code>e(vcetype)</code>	title used to label Std. Err.
<code>e(properties)</code>	<code>b V</code>

### Matrices

<code>e(b)</code>	observed statistics
<code>e(b_bs)</code>	bootstrap estimates
<code>e(reps)</code>	number of nonmissing results
<code>e(bias)</code>	estimated biases
<code>e(se)</code>	estimated standard errors
<code>e(z0)</code>	median biases
<code>e(accel)</code>	estimated accelerations
<code>e(ci_normal)</code>	normal-approximation CIs
<code>e(ci_percentile)</code>	percentile CIs
<code>e(ci_bc)</code>	bias-corrected CIs
<code>e(ci_bca)</code>	bias-corrected and accelerated CIs
<code>e(V)</code>	bootstrap variance-covariance matrix

## References

- Ng, E. S.-W., R. Grieve, and J. R. Carpenter. 2013. Two-stage nonparametric bootstrap sampling with shrinkage correction for clustered data. *Stata Journal* 13: 141–164.
- Poi, B. P. 2004. From the help desk: Some bootstrapping techniques. *Stata Journal* 4: 312–328.

## Also see

- [R] [bootstrap postestimation](#) — Postestimation tools for bootstrap
- [R] [bootstrap](#) — Bootstrap sampling and estimation
- [R] [bsample](#) — Sampling with replacement