

## Title

**bradout** — Outputs multiple estimation results in a clean table

## Syntax

**bradout** [string] [, options]

options	Description
Statistics	
<b>stats</b> ( <i>string</i> )	select which covariate statistics will be displayed
<b>stars</b> ( <i>numlist</i> )	creates up to 3 significance stars for p-values
<b>mstats</b> ( <i>string</i> )	select which model statistics will be displayed
<b>minfo</b> ( <i>string</i> )	select which model information will be displayed
Covariates	
<b>order</b> ( <i>string</i> )	choose which covariates are displayed first
<b>drop</b> ( <i>string</i> )	choose which covariates to drop, while keeping all others
<b>keep</b> ( <i>string</i> )	choose which covariates to keep, while dropping all others
<b>nobases</b>	do not display base levels of covariates CHECK THIS
<b>noomits</b>	do not display omitted covariates
<b>norefs</b>	do not display variable header for categorical variables CHECK THIS
Labelling	
<b>collabels</b> ( <i>string</i> )	specify column labels
<b>mlabels</b> ( <i>string</i> )	specify model labels
<b>novarlabels</b>	do not display variable labels
<b>refcats</b> ( <i>string</i> )	specify manual reference categories
<b>catfill</b> ( <i>string</i> )	specify fill string for category cells
<b>reffill</b> ( <i>string</i> )	specify fill string for reference cells
<b>omitfill</b> ( <i>string</i> )	specify fill string for omitted cells
Display	
<b>title</b> ( <i>string</i> )	optional custom title or "none" to display no title
<b>hseparator</b>	display horizontal separator lines
<b>vseparator</b>	display vertical separator lines
<b>wide</b>	display in a wide format
<b>bind</b>	bind models such that all their columns are displayed in one table
Output	
<b>excel</b> ( <i>string</i> )	Excel output options
System	
<b>clear</b>	clear stored estimates after command

## Description

**bradout** outputs multiple estimation results in a clean table that can be exported to Excel. It also has many customization options to structure the table how you'd like.

## Options

### Statistics

**stats** (*string*) allows users to choose from the following statistics:

<b>b</b>	b (Coefficient/OR/etc)
<b>se</b>	standard error
<b>tz</b>	t/z
<b>pvalue</b>	p-value
<b>ci</b>	confidence interval

**stars** (*numlist*) creates up to 3 significance stars for p-values less than *numlist* containing 0-3 values. Leaving *numlist* empty defaults to  $p < 0.05$  and  $p < 0.01$

**mstats** (*string*) allows users to choose from the following model statistics:

<b>n</b>	number of observations
<b>n_yes</b>	number of yes's
<b>n_pop</b>	size of population
<b>n_strata</b>	number of strata
<b>n_clust</b>	number of clusters
<b>n_psu</b>	number of primary sampling units
<b>n_cds</b>	number of completely determined successes
<b>n_cdf</b>	number of completely determined failures
<b>n_cd</b>	number of completely determined observations
<b>df_m</b>	model degrees of freedom
<b>mss</b>	model sum of squares
<b>df_r</b>	residual degrees of freedom
<b>rss</b>	residual sum of squares
<b>rmse</b>	root mean squared error
<b>r2</b>	R-squared
<b>r2_a</b>	adjusted R-squared
<b>r2_p</b>	pseudo R-squared
<b>ll</b>	log likelihood
<b>ll0</b>	log likelihood, constant-only model
<b>aic</b>	Akaike's information criterion
<b>bic</b>	Bayesian information criterion
<b>f</b>	F statistic
<b>chi2</b>	chi-squared
<b>p</b>	p-value for model test
<b>k</b>	number of parameters
<b>ic</b>	number of iterations
<b>rank</b>	rank of e(V)

**minfo**(*string*) allows users to choose from the following model information:

<b>mgroup</b>	model group
<b>depvar</b>	dependent variable
<b>cmd</b>	command
<b>weight</b>	weight

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### Covariates

**order**(*string*) specifies which covariates are displayed first.

**drop**(*string*) specifies which covariates to drop, while keeping all others.

**keep**(*string*) specifies which covariates to keep, while dropping all others.

**nobases** hides display of base levels of covariates.

**noomits** hides display of omitted covariates.

**norefs** hides display of variable headers for categorical variables.

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### Labelling

**collabels**(*string*) specifies the labels of columns. If there are fewer specified column labels than total columns, the labels will repeat.

**mlabels**(*string*) specifies the labels of models.

**novarlabels** hides display of variable labels.

**refcats**(*string*) specifies manual reference categories.

**catfill**(*string*) specifies the fill string for category cells.

**reffill**(*string*) specifies the fill string for reference cells.

**omitfill**(*string*) specifies the fill string for omitted cells.

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### Display

**title**(*string*) specifies an optional custom title or **"none"** to display no title.

**hseparator** displays horizontal separator lines.

**vseparator** displays vertical separator lines.

**wide** displays results in a wide format.

**bind** binds models such that all their columns are displayed in one table.

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## Output

**excel**(*string*) has the following options:

<b>file</b> ( <i>string</i> )	location of output file. Default is a file named <b>bradmean_output.xlsx</b> in the current working directory
<b>sheet</b> ( <i>string</i> )	name of sheet to be used. Default is the first file in the sheet or <b>Sheet1</b> in a new workbook
<b>replace</b>	replace the workbook
<b>sheetreplace</b>	replace the sheet
<b>modify</b>	append table to the end of the sheet
<b>font</b> ( <i>string</i> )	choose the font face from <b>Arial</b> , <b>Calibri</b> , <b>Garamond</b> , <b>Helvetica</b> , <b>TNR</b> (Times New Roman), or <b>Verdana</b> . Default is <b>Calibri</b>
<b>size</b> (#)	choose the font size between 9 and 12. Default is 11
<b>color</b> ( <i>string</i> )	choose the color styles from <b>bradmean</b> , <b>monochrome</b> , <b>rti</b> , <b>material_red</b> , <b>material_purple</b> , <b>material_indigo</b> , <b>material_blue</b> , <b>material_green</b> , and <b>material_orange</b>

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## System

**clear** clears stored results.