Title

bradout — Outputs multiple estimation results in a clean table

Syntax

```
bradout [string] [, options]
                        Description
options
Statistics
  stats(string)
                        select which covariate statistics will be displayed
                        creates up to 3 significance stars for p-values
  stars(numlist)
                        select which model statistics will be displayed
  mstats(string)
  minfo(string)
                        select which model information will be displayed
Covariates
                        choose which covariates are displayed first
  order(string)
                        choose which covariates to drop, while keeping all others
  drop (string)
                        choose which covariates to keep, while dropping all others
  keep (string)
                        do not display base levels of covariates
  <u>nobase</u>s
  <u>noomit</u>s
                        do not display omitted covariates
  norefs
                        do not display variable header for categorical variables
Labelling
  collabels(string)
                        specify column labels
  mlabels(string)
                        specify model labels
                        do not display variable labels specify manual reference categories
  <u>novarlab</u>els
  refcats (string)
  catfill(string)
                        specify fill string for category cells
                       specify fill string for reference cells specify fill string for omitted cells
  reffill (string)
  omitfill(string)
Display
                        optional custom title or "none" to display no title
  title(string)
                        display horizontal separator lines
  <u>hsep</u>arator
 <u>vsep</u>arator
                        display vertical separator lines
                        display in a wide format
  wide
                        bind models such that all their columns are displayed in one table
 bind
Output
  excel(string)
                        Excel output options
System
  clear
                        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 (Coefficient/OR/etc)
se standard error
tz t/z
pvalue p-value

ci 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

 $\underline{{\tt mstat}}{\tt s}\,(\underline{{\it string}})$ allows users to choose from the following model statistics:

number of observations n number of observations in subpopulation n sub number of yes's n yes size of population size of subpopulation n_pop n subpop number of primary sampling units n psu number of clusters n clust n_cds number of completely determined successes n_cdf number of completely determined failures number of completely determined observations n cd number of groups n q minimum group size g_min g avg average group size g_max maximum group size harmonic mean of group sizes number of dropped observations number of dropped groups model sum of squares tbar n drop g drop residual sum of squares rss tss total sum of squares df m model degrees of freedom df_r residual degrees of freedom df pear degrees of freedom for Pearson chi2 degrees of freedom for absorbed effect df a df b numerator degrees of freedom for F statistic root mean squared error
rank of e(V) rmse rank rank0 rank of e(V) for constant-only model rho
panel-level standard deviation
ancillary parameter rho sigma u sigma standard deviation of the error term (epsilon) sigma e deviance deviance deviance dispersion dispers corr correlation r2 R-squared adjusted R-squared r2 a r2_p pseudo R-squared R-squared for within model R-squared for overall model r2_w r2_0 R-squared for between model r2_b 11 log likelihood 110 log likelihood, constant-only model
log likelihood, comparison model 11c Akaike's information criterion Bayesian information criterion bic thta min minimum theta thta 5 5th percentile theta 50th percentile theta 95th percentile theta thta_50 thta 95 maximum theta thta max f F statistic f_f F statistic for test of u i=0 p-value for model test p p-value for test of u i=0 р chi2 chi-squared chi2 chi-squared for comparison test chi2 dev chi-squared test of deviance chi-squared test of deviance dispersion chi2 dis tol target tolerance dif achieved tolerance number of parameters k k aux number of auxiliary parameters number of equations in e(b) k eq k eq model number of equations in overall model test k_dv number of dependent variables number of quadrature points n quad phi scale parameter number of sampling stages stages number of iterations

mgroup model group

depvar dependent variable

cmd command
model model type

family distribution family

link link function

weight weight

___ 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.

Labelling L

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.

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.

Output

excel(string) has the following options:

file(string) location of output file. Default is a file named bradmean output.xlsx

in the current working directory

sheet(string) name of sheet to be used. Default is the first file in the sheet or

Sheet1 in a new workbook

modify append table to the end of the sheet

font(<u>string</u>) choose the font face from Arial, Calibri, Garamond, Helvetica, TNR

(Times New Roman), or Verdana. Default is Calibri

size(#) choose the font size between 9 and 12. Default is 11

color(string) choose the color styles from bradmean, monochrome, rti, material_red,

material_purple, material_indigo, material_blue, material_green, and

material orange

System

clear clears stored results.