

#### help binspwc

### <u>Title</u>

binspwc — Data-Driven Nonparametric Pairwise Group Comparison using Binscatter.

#### Syntax

where <u>depvar</u> is the dependent variable, <u>indvar</u> is the independent variable for binning, and <u>covars</u> are other covariates to be controlled for.

p, s and v are integers satisfying 0 <= s,v <= p, which can take different values in each case.

fweights, aweights and pweights are allowed; see weight.

### Description

binspwc implements binscatter-based hypothesis testing procedures for pairwise
 group comparison of binscatter estimators, following the results in <u>Cattaneo</u>,
 <u>Crump</u>, <u>Farrell and Feng (2021a)</u>. If the binning scheme is not set by the
 user, the companion command <u>binsregselect</u> is used to implement binscatter in a
 data-driven (optimal) way and inference procedures are based on robust bias
 correction. Binned scatter plots based on different models can be constructed
 using the companion commands <u>binsreg</u>, <u>binsqreg</u>, <u>binslogit</u> and <u>binsprobit</u>.

A detailed introduction to this command is given in  $\underline{\text{Cattaneo, Crump, Farrell and }}$   $\underline{\text{Feng (2021b)}}$ . A companion R package with the same capabilities is available (see website below).

Companion commands: <u>binsreg</u> for binscatter least squares regression with robust inference procedures and plots, <u>binsqreg</u> for binscatter quantile regression with robust inference procedures and plots, <u>binslogit</u> for binscatter logit estimation with robust inference procedures and plots, <u>binsprobit</u> for binscatter probit estimation with robust inference procedures and plots, and <u>binsregselect</u> data-driven (optimal) binning selection.

Related Stata and R packages are available in the following website:

https://nppackages.github.io/

### Options

```
Estimand
```

estmethod(cmdname) specifies the binscatter model. The default is estmethod(reg),
 which corresponds to the binscatter least squares regression. Other options
 are: estmethod(qreg #) for binscatter quantile regression where # is the
 quantile to be estimated, estmethod(logit) for binscatter logistic regression
 and estmethod(probit) for binscatter probit regression.

deriv(v) specifies the derivative order of the regression function for estimation,
 testing and plotting. The default is deriv(0), which corresponds to the
 function itself.

- at (position) specifies the values of covars at which the estimated function is evaluated for plotting. The default is at (mean), which corresponds to the mean of covars. Other options are: at (median) for the median of covars, at (0) for zeros, and at(filename) for particular values of covars saved in another file.
- Note: when at (mean) or at (median) is specified, all factor variables in covars (if specified) are excluded from the evaluation.
- nolink specifies that the function within the inverse link (logistic) function be reported instead of the conditional probability function. This option is used only if logit or probit model is specified in estmethod().
- by (varname) specifies the variable containing the group indicator to perform subgroup analysis; both numeric and string variables are supported. When by (varname) is specified, binspwc implements estimation by each subgroup separately and then conduct all pairwise comparison tests. By default, the binning structure is selected for each subgroup separately, but see the option samebinsby below for imposing a common binning structure across subgroups.

# ⊥ <sub>Reghdfe</sub> ∟

- absorb(absvars) specifies categorical variables (or interactions) representing the fixed effects to be absorbed. This is equivalent to including an indicator/dummy variable for each category of each absvar. When absorb() is specified, the community-contributed command reghdfe instead of the command regress is used.
- reghdfeopt (reghdfe\_option) options to be passed on to the command reghdfe. Important: absorb() and vce() should not be specified within this option.
- For more information about the community-contributed command reghdfe, please see http://scorreia.com/software/reghdfe/.

# ☐ Pairwise Group Comparison Testing

- $pwc(p \ s)$  sets a piecewise polynomial of degree p with s smoothness constraints for pairwise group comparison. The default is pwc(3 3), which corresponds to a cubic B-spline estimate of the function of interest for each group.
- ${\tt testtype}\ ({\it type})$  specifies the type of pairwise comparison test. The default is testtype(2), which corresponds to a two-sided test of the form HO:  $mu_1(x) = mu_2(x)$ . Other options are: **testtype(**1) for the one-sided test of the form H0:  $mu_1(x) <= mu_2(x)$  and **testtype(r)** for the one-sided test of the form H0:  $mu_1(x) > = mu_2(x)$ .
- lp(metric) specifies a Lp metric used for a (two-sided) test for the difference between two groups. The default is lp(inf), which corresponds to the sup-norm. Other options are Lp(q) for a positive integer q.

## $\square$ Partitioning/Binning Selection $^{oldsymbol{\mathsf{L}}}$

- The default is bins (0 0), which corresponds to piecewise constant (canonical binscatter).
- ordering of the group follows the result of the tabulate. If a single number of bins is specified, it applies to the estimation for all groups. If not specified, the number of bins is selected via the companion command binsregselect in a data-driven, optimal way whenever possible.

- binspos(position) specifies the position of binning knots. The default is
  binspos(qs), which corresponds to quantile-spaced binning (canonical
  binscatter). Other options are: es for evenly-spaced binning, or a numlist
  for manual specification of the positions of inner knots (which must be within
  the range of indvar).
- binsmethod(method) specifies the method for data-driven selection of the number of
   bins via the companion command binsregselect. The default is binsmethod(dpi),
   which corresponds to the IMSE-optimal direct plug-in rule. The other option
   is: rot for rule of thumb implementation.
- nbinsrot(#) specifies an initial number of bins value used to construct the DPI number of bins selector. If not specified, the data-driven ROT selector is used instead.
- samebinsby forces a common partitioning/binning structure across all subgroups
   specified by the option by(). The knots positions are selected according to
   the option binspos() and using the full sample. If nbins() is not specified,
   then the number of bins is selected via the companion command binsregselect
   and using the full sample.
- randcut(#) specifies the upper bound on a uniformly distributed variable used to
   draw a subsample for bins selection. Observations for which runiform()<=# are
   used. # must be between 0 and 1.</pre>

Simulation

- nsims(#) specifies the number of random draws for constructing confidence bands
  and hypothesis testing. The default is nsims(500), which corresponds to 500
  draws from a standard Gaussian random vector of size [(p+1)\*J (J-1)\*s].
- simsgrid(#) specifies the number of evaluation points of an evenly-spaced grid
  within each bin used for evaluation of the supremum (or infimum) operation
  needed to construct confidence bands and hypothesis testing procedures. The
  default is simsgrid(20), which corresponds to 20 evenly-spaced evaluation
  points within each bin for approximating the supremum (or infimum) operator.
- simsseed(#) sets the seed for simulations.

Mass Points and Degrees of Freedom

- dfcheck(n1 n2) sets cutoff values for minimum effective sample size checks, which
   take into account the number of unique values of indvar (i.e., adjusting for
   the number of mass points), number of clusters, and degrees of freedom of the
   different statistical models considered. The default is dfcheck(20 30). See
   Cattaneo, Crump, Farrell and Feng (2021b) for more details.
- masspoints(masspointsoption) specifies how mass points in indvar are handled. By
   default, all mass point and degrees of freedom checks are implemented.
   Available options:

masspoints(noadjust) omits mass point checks and the corresponding effective sample size adjustments.

masspoints(nolocalcheck) omits within-bin mass point and degrees of freedom
checks.

 ${\tt masspoints}\ ({\tt off})$  sets  ${\tt masspoints}\ ({\tt noadjust})$  and  ${\tt masspoints}\ ({\tt nolocalcheck})$  simultaneously.

masspoints(veryfew) forces the command to proceed as if indvar has only a few number of mass points (i.e., distinct values). In other words, forces the command to proceed as if the mass point and degrees of freedom checks were failed.

Other Options

vce(vcetype) specifies the vcetype for variance estimation used by the commands
 regress, logit or greg. The default is vce(robust).

- asyvar(on/off) specifies the method used to compute standard errors. If
  asyvar(on) is specified, the standard error of the nonparametric component is
  used and the uncertainty related to other control variables covars is omitted.
  Default is asyvar(off), that is, the uncertainty related to covars is taken
  into account.
- usegtools(on/off) forces the use of several commands in the community-distributed
   Stata package gtools to speed the computation up, if on is specified. Default
   is usegtools(off).
- For more information about the package **gtools**, please see <a href="https://gtools.readthedocs.io/en/latest/index.html">https://gtools.readthedocs.io/en/latest/index.html</a>.

### **Examples**

Setup

. sysuse auto

Generate two groups

. gen group=price>5000

Test for the difference between two groups . binspwc mpg weight foreign, by(group)

### Stored results

```
Scalars
 e (N)
                   number of observations
 e (p)
                  degree of polynomial for bin selection
                   smoothness of polynomial for bin selection
 e (s)
                  degree of polynomial for testing
 e (pwc_p)
                  smoothnes of polynomial for testing
 e (pwc_s)
Locals
 e(byvalue)
                  name of groups found in by ()
Matrices
 e(N_by)
                   number of observations for each group
 e(Ndist_by)
                   number of distinct values for each group
 e(Nclust_by)
                   number of clusters for each group
 e(nbins_by)
                  number of bins for each group
                  test statistics for all pairwise comparisons
 e(stat)
                  p values for all pairwise comparisons
 e(pval)
```

### References

```
Cattaneo, M. D., R. K. Crump, M. H. Farrell, and Y. Feng. 2021a. On Binscatter. arXiv:1902.09608.
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Cattaneo, M. D., R. K. Crump, M. H. Farrell, and Y. Feng. 2021b. <u>Binscatter Regressions</u>. arXiv:1902.09615.

### <u>Authors</u>

Matias D. Cattaneo, Princeton University, Princeton, NJ. <u>cattaneo@princeton.edu</u>.

Richard K. Crump, Federal Reserve Band of New York, New York, NY. richard.crump@ny.frb.org.

Max H. Farrell, University of Chicago, Chicago, IL. max.farrell@chicagobooth.edu.

Yingjie Feng, Tsinghua University, Beijing, China. fengyingjiepku@gmail.com.