

help binstest

<u>Title</u>

Syntax

where <u>depvar</u> is the dependent variable, <u>indvar</u> is the independent variable for binning, and <u>othercovs</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.
- At least one test has to be specified via testmodelparfit(), testmodelpoly(), testshapel(), testshaper() and/or testshape2().

fweights, aweights and pweights are allowed; see weight.

Description

- binstest implements binscatter-based hypothesis testing procedures for parametric
 functional forms of and nonparametric shape restrictions on the regression
 function estimators, following the results in Cattaneo, Crump, Farrell and
 Feng (2021a). If the binning scheme is not set by the user, the companion
 command binsregselect 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 binsreg, binsqreg, binslogit and binsprobit.
- A detailed introduction to this command is given in <u>Cattaneo, Crump, Farrell and Feng (2021b)</u>. Companion R and Python packages with the same capabilities are available (see website below).
- Companion commands: binsreg for binscatter regression with robust inference procedures and plots, binsqreg for binscatter quantile regression with robust inference procedures and plots, binslogit for binscatter logit estimation with robust inference procedures and plots, binsprobit for binscatter probit estimation with robust inference procedures and plots, and binsregselect for data-driven (optimal) binning selection.

Related Stata, R and Python 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 othercovs at which the estimated function is evaluated for plotting. The default is **at(mean)**, which corresponds to the mean of *othercovs*. Other options are: **at(median)** for the median of *othercovs*, at(0) for zeros, and at(filename) for particular values of othercovs saved in another file.
- Note: when at (mean) or at (median) is specified, all factor variables in othercovs (if specified) are excluded from the evaluation (set as zero).
- 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().

」 _{Reghdfe} ∟

- 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/.

 $^{
m flue}$ Parametric Model Specification Testing $^{
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- testmodel(p s) sets a piecewise polynomial of degree p with s smoothness constraints for parametric model specification testing. The default is testmodel(3 3), which corresponds to a cubic B-spline estimate of the regression function of interest for testing against the fitting from a parametric model specification.
- $\textbf{testmodelparfit} \textit{(filename)} \ \text{specifies a dataset which contains the evaluation grid}$ and fitted values of the model(s) to be tested against. The file must have a variable with the same name as indvar, which contains a series of evaluation points at which the binscatter model and the parametric model of interest are compared with each other. Each parametric model is represented by a variable named as binsreg_fit*, which must contain the fitted values at the corresponding evaluation points.
- testmodelpoly(p) specifies the degree of a global polynomial model to be tested against.

 igsqc Nonparametric Shape Restriction Testing igsqcup

- $\textbf{testshape} \ (p \ s) \ \text{sets a piecewise polynomial of degree} \ p \ \text{with} \ s \ \text{smoothness} \\ \text{constraints for nonparametric shape restriction testing.} \ \ \text{The default is}$ testshape(3 3), which corresponds to a cubic B-spline estimate of the regression function of interest for one-sided or two-sided testing.
- one-sided hypothesis test to the left of the form H0: $\sup_{x \in a} \sup_{x \in a}$
- testshaper(numlist) specifies a numlist of null boundary values for hypothesis testing. Each number a in the numlist corresponds to one boundary of a one-sided hypothesis test to the right of the form H0: $inf_x mu(x) >= a$.
- testshape2 (numlist) specifies a numlist of null boundary values for hypothesis testing. Each number a in the numlist corresponds to one boundary of a two-sided hypothesis test of the form H0: $\sup_{x} |mu(x) - a| = 0$.

 $^{f J}$ Metric for Hypothesis Testing

lp(metric) specifies an Lp metric used for (two-sided) parametric model specification testing and/or shape restriction testing. The default is lp(inf), which corresponds to the sup-norm. Other options are lp(q) for a positive integer q.

Partitioning/Binning Selection

- The default is bins (2 2), which corresponds to a quadratic spline estimate.
- nbins (#) sets the number of bins for partitioning/binning of indvar. If not specified, the number of bins is selected via the companion command <u>binsregselect</u> 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 <u>binsregselect</u>. 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.
- 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.

」 Simulation └

- nsims(#) specifies the number of random draws for 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].
- simsqrid(#) specifies the number of evaluation points of an evenly-spaced grid within each bin used for evaluation of the supremum (infimum or lp metric) operation needed for hypothesis testing procedures. The default is simsgrid(20), which corresponds to 20 evenly-spaced evaluation points within each bin for approximating the supremum (infimum or Lp metric) 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.

masspoints(off) sets masspoints(noadjust) and masspoints(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.

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☐ Other Options
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vce(vcetype) specifies the vcetype for variance estimation used by the commands regress, logit, probit, qreg or reghdfe. 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 othercovs is omitted. Default is asyvar(off), that is, the uncertainty related to othercovs is taken into account.

useqtools(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 https://gtools.readthedocs.io/en/latest/index.html.

Examples

```
Setup
```

. sysuse auto

Test for linearity

. binstest mpg weight foreign, testmodelpoly(1)

Test for monotonicity

. binstest mpg weight foreign, deriv(1) bins(1 1) testshapel(0)

Stored results

```
Scalars
 e (N)
                    number of observations
  e(Ndist)
                    number of distinct values
 e(Nclust)
                    number of clusters
                    number of bins
 e(nbins)
                    degree of polynomial for bin selection
  e (p)
  e (s)
                    smoothness of polynomial for bin selection
  e(testshape_p)
                    degree of polynomial for testing shape restrictions
                    smoothness of polynomial for testing shape restrictions degree of polynomial for testing model specifications
  e(testshape s)
  e(testmodel_p)
  e(testmodel_s)
                    smoothness of polynomial for testing model specifications
  e(testpolyp)
                    degree of polynomial regression model
                    statistic for testing global polynomial model
  e(stat_poly)
  e(pval_poly)
                    p value for testing global polynomial model
Macros
                    varlist found in testmodel()
  e(testvarlist)
  e(testvalue2)
                    values in testshape2()
  e(testvalueR)
                    values in testshaper()
  e(testvalueL)
                    values in testshapel()
Matrices
  e(pval_model)
                    p values for testmodel()
  e(stat_model)
                    statistics for testmodel()
  e(pval_shape2)
                    p values for testshape2()
  e(stat_shape2)
                    statistics for testshape2()
  e (pval_shapeR)
                    p values for testshaper()
```

```
e(stat_shapeR) statistics for testshaper()
e(pval_shapeL) p values for testshapel()
e(stat_shapeL) statistics for testshapel()
```

References

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

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

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