



## help binstest

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### Title

**binstest** — Data-Driven Nonparametric Shape Restriction and Parametric Model Specification Testing using Binscatter.

### Syntax

```
binstest depvar indvar [covars] [if] [in] [weight] [ , estmethod(cmdname) deriv(v)
testmodel(p s) testmodelparfit(filename) testmodelpoly(p)
testshape(p s) testshapel(numlist) testshaper(numlist)
testshape2(numlist) lp(metric)
bins(p s) nbins(#) binspos(position) binsmethod(method) nbinsrot(#)
nsims(#) simsgrid(#) simsseed(seed)
dfcheck(n1 n2) masspoints(masspointsoption)
vce(vcetype) ]
```

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

*p*, *s* and *v* are integers satisfying  $0 \leq s, v \leq p$ , which can take different values in each case.

**fweights**, **aweight**s and **pweight**s are allowed; see [weight](#).

### Description

**binstest** implements binscatter-based hypothesis testing procedures for parametric functional forms and nonparametric shape restrictions on of 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 [Cattaneo, Crump, Farrell and Feng \(2021b\)](#). A companion R package with the same capabilities is 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](#) data-driven (optimal) binning selection.

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

<https://nppackages.github.io/>

### Options

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Estimand

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**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.

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### 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.

**testmodelparfit**(*filename*) 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.

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### Nonparametric Shape Restriction Testing

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**testshape**(*p s*) sets a piecewise polynomial of degree *p* with *s* smoothness constraints for nonparametric shape restriction testing. The default is **testmodel(3 3)**, which corresponds to a cubic B-spline estimate of the regression function of interest for one-sided or two-sided testing.

**testshapel**(*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 left of the form  $H_0: \sup_x \mu(x) \leq a$ .

**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  $H_0: \inf_x \mu(x) \geq a$ .

**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  $H_0: \sup_x |\mu(x) - a| = 0$ .

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### Metric for Hypothesis Testing

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**lp**(*metric*) specifies a *L<sub>p</sub>* 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*.

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### Partitioning/Binning Selection

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**bins**(*p s*) sets a piecewise polynomial of degree *p* with *s* smoothness constraints for data-driven (IMSE-optimal) selection of the partitioning/binning scheme. The default is **bins(0 0)**, which corresponds to piecewise constant (canonical binscatter).

**nbins**(*#*) sets the number of bins for partitioning/binning of *indvar*. 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.

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#### Simulation

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**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.

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#### Mass Points and Degrees of Freedom

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**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 (2019b) 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** or **qreg**. The default is **vce(robust)**.

### Examples

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Test linear model
. binstest y x w, testmodelpoly(1)
```

### Stored results

#### Scalars

<b>e(N)</b>	number of observations
<b>e(Ndist)</b>	number of distinct values
<b>e(Nclust)</b>	number of clusters
<b>e(nbins)</b>	number of bins
<b>e(p)</b>	degree of polynomial for bin selection
<b>e(s)</b>	smoothness of polynomial for bin selection
<b>e(testshape_p)</b>	degree of polynomial for testing shape
<b>e(testshape_s)</b>	smoothness of polynomial for testing shape
<b>e(testmodel_p)</b>	degree of polynomial for testing models
<b>e(testmodel_s)</b>	smoothness of polynomial for testing models
<b>e(testpolyp)</b>	degree of polynomial regression model
<b>e(stat_poly)</b>	statistic for testing global polynomial model
<b>e(pval_poly)</b>	p value for testing global polynomial model

#### Locals

<code>e(testvalueL)</code>	values in <code>testshapel()</code>
<code>e(testvalueR)</code>	values in <code>testshaper()</code>
<code>e(testvalue2)</code>	values in <code>testshape2()</code>
<code>e(testvarlist)</code>	varlist found in <code>testmodel()</code>

Matrices

<code>e(stat_shapeL)</code>	statistics for <code>testshapel()</code>
<code>e(pval_shapeL)</code>	p values for <code>testshapel()</code>
<code>e(stat_shapeR)</code>	statistics for <code>testshaper()</code>
<code>e(pval_shapeR)</code>	p values for <code>testshaper()</code>
<code>e(stat_shape2)</code>	statistics for <code>testshape2()</code>
<code>e(pval_shape2)</code>	p values for <code>testshape2()</code>
<code>e(stat_model)</code>	statistics for <code>testmodel()</code>
<code>e(pval_model)</code>	p values for <code>testmodel()</code>

## **References**

- Cattaneo, M. D., R. K. Crump, M. H. Farrell, and Y. Feng. 2021a. [On Binscatter](#). *arXiv:1902.09608*.
- Cattaneo, M. D., R. K. Crump, M. H. Farrell, and Y. Feng. 2021b. [Binscatter Regressions](#). *arXiv:1902.09615*.

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