

Package ‘nprobust’

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Type Package

Title Kernel Density and Local Polynomial Regression Methods

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Description Estimation, inference, bandwidth selection, and graphical procedures for kernel density and local polynomial regression methods, including robust bias-corrected confidence intervals as described in Calonico, Cattaneo and Farrell (2018, <[doi:10.1080/01621459.2017.1285776](https://doi.org/10.1080/01621459.2017.1285776)>). The package includes 'lprobust()' for local polynomial point estimation and robust bias-corrected inference, 'lpbwselect()' for local polynomial bandwidth selection, 'kdrobust()' for kernel density point estimation and robust bias-corrected inference, 'kdbwselect()' for kernel density bandwidth selection, and 'nprobust.plot()' for plotting results. The main methodological and numerical features are described in Calonico, Cattaneo and Farrell (2019, <[doi:10.18637/jss.v091.i08](https://doi.org/10.18637/jss.v091.i08)>).

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License GPL-3

URL <https://github.com/nppackages/nprobust>

BugReports <https://github.com/nppackages/nprobust/issues>

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nprobust-package	<i>Kernel Density and Local Polynomial Regression Methods</i>
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Description

This package implements estimation, inference, bandwidth selection, and graphical procedures for kernel density and local polynomial regression methods, including robust bias-corrected confidence intervals as described in Calonico, Cattaneo and Farrell (2018): [lprobust](#) for local polynomial point estimation and robust bias-corrected inference, [lpbwselect](#) for local polynomial bandwidth selection, [kdrobust](#) for kernel density point estimation and robust bias-corrected inference, [kdbwselect](#) for kernel density bandwidth selection, and [nprobust.plot](#) for plotting results. The main methodological and numerical features are described in Calonico, Cattaneo and Farrell (2019).

Details

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Function for LPR estimation and inference: [lprobust](#)
 Function for LPR bandwidth selection: [lpbwselect](#)
 Function for KDE estimation and inference: [kdrobust](#)
 Function for KDE bandwidth selection: [kdbwselect](#)
 Function for graphical analysis: [nprobust.plot](#)

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References

- Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2018. [On the Effect of Bias Estimation on Coverage Accuracy in Nonparametric Inference](#). *Journal of the American Statistical Association*, 113(522): 767-779. doi:10.1080/01621459.2017.1285776.
- Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2019. [nprobust: Nonparametric Kernel-Based Estimation and Robust Bias-Corrected Inference](#). *Journal of Statistical Software*, 91(8): 1-33. doi:10.18637/jss.v091.i08.
- Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2022. [Coverage Error Optimal Confidence Intervals for Local Polynomial Regression](#). *Bernoulli*, 28(4): 2998-3022.

kdbwselect	<i>Bandwidth Selection Procedures for Kernel Density Estimation and Inference</i>
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Description

`kdbwselect` implements bandwidth selectors for kernel density point estimators and inference procedures developed in Calonico, Cattaneo and Farrell (2018). See also Calonico, Cattaneo and Farrell (2022) for related optimality results. It also implements other bandwidth selectors available in the literature.

Companion commands are: `kdrobust` for kernel density point estimation and inference procedures.

A detailed introduction to this command is given in Calonico, Cattaneo and Farrell (2019). For more details and related software useful for empirical analysis, visit <https://nppackages.github.io/>.

Usage

```
kdbwselect(x, eval = NULL, neval = NULL, kernel = "epa",
           bwselect = "mse-dpi", bwcheck=21, imsegrid=30, subset = NULL, data = NULL)
```

Arguments

<code>x</code>	independent variable.
<code>eval</code>	vector of evaluation point(s). By default it uses 30 quantile-spaced points (deciles 0.1 to 0.9 in equal steps) over the support of <code>x</code> .
<code>neval</code>	number of quantile-spaced evaluation points on the support of <code>x</code> . Default is <code>neval=30</code> .
<code>kernel</code>	kernel function used to construct the kernel estimators. Options are <code>epa</code> for the epanechnikov kernel, and <code>uni</code> for the uniform kernel. Default is <code>kernel = epa</code> .
<code>bwselect</code>	bandwidth selection procedure to be used. Options are: <code>mse-dpi</code> second-generation DPI implementation of MSE-optimal bandwidth. Default option. <code>imse-dpi</code> second-generation DPI implementation of IMSE-optimal bandwidth (computed using grid of evaluation points selected).

imse-rot ROT implementation of IMSE-optimal bandwidth (computed using grid of evaluation points selected).

ce-dpi second generation DPI implementation of CE-optimal bandwidth.

ce-rot ROT implementation of CE-optimal bandwidth.

all reports all available bandwidth selection procedures.

Note: MSE = Mean Square Error; IMSE = Integrated Mean Squared Error; CE = Coverage Error; DPI = Direct Plug-in; ROT = Rule-of-Thumb. For details on implementation see Calonico, Cattaneo and Farrell (2019).

bwcheck	if a positive integer is provided, then the selected bandwidth is enlarged so that at least bwcheck effective observations are available at each evaluation point. Default is bwcheck = 21.
imsegrid	number of evaluations points used to compute the IMSE bandwidth selector. Default is imsegrid = 30.
subset	optional rule specifying a subset of observations to be used.
data	an optional data frame. When supplied, x and subset may be given as bare variable names referring to columns of data.

Value

bws	A matrix containing eval (grid points), h and b (selected bandwidths) for each evaluation point. When bwselect="all", the matrix is widened to include both MSE and CER bandwidths from each of the DPI and ROT estimators.
bws.imse	IMSE-optimal bandwidths returned when bwselect="all"; NULL otherwise.
opt	A list containing options passed to the function.

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References

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Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2019. [nprobust: Nonparametric Kernel-Based Estimation and Robust Bias-Corrected Inference](#). *Journal of Statistical Software*, 91(8): 1-33. doi:10.18637/jss.v091.i08.

Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2022. [Coverage Error Optimal Confidence Intervals for Local Polynomial Regression](#). *Bernoulli*, 28(4): 2998-3022.

See Also

[kdrobust](#)

Examples

```
x <- rnorm(500)
est <- kdbwselect(x)
summary(est)
```

kdrobust

Kernel Density Methods with Robust Bias-Corrected Inference

Description

`kdrobust` implements kernel density point estimators, with robust bias-corrected confidence intervals and inference procedures developed in Calonico, Cattaneo and Farrell (2018). See also Calonico, Cattaneo and Farrell (2022) for related optimality results. It also implements other estimation and inference procedures available in the literature.

Companion commands: `kdbwselect` for kernel density data-driven bandwidth selection, and `nprobust.plot` for plotting results.

A detailed introduction to this command is given in Calonico, Cattaneo and Farrell (2019). For more details and related software useful for empirical analysis, visit <https://nppackages.github.io/>.

Usage

```
kdrobust(x, eval = NULL, neval = NULL, h = NULL, b = NULL, rho = 1,
kernel = "epa", bwselect = NULL, bwcheck = 21, imsegrid=30, level = 95, subset = NULL,
data = NULL)
```

Arguments

<code>x</code>	independent variable.
<code>eval</code>	vector of evaluation point(s). By default it uses 30 quantile-spaced points (deciles 0.1 to 0.9 in equal steps) over the support of <code>x</code> .
<code>neval</code>	number of quantile-spaced evaluation points on the support of <code>x</code> . Default is <code>neval=30</code> .
<code>h</code>	main bandwidth used to construct the kernel density point estimator. Can be either scalar (same bandwidth for all evaluation points), or vector of same dimension as <code>eval</code> . If not specified, bandwidth <code>h</code> is computed by the companion command <code>kdbwselect</code> .
<code>b</code>	bias bandwidth used to construct the bias-correction estimator. Can be either scalar (same bandwidth for all evaluation points), or vector of same dimension as <code>eval</code> . By default it is set equal to <code>h</code> . If <code>rho</code> is set to zero, <code>b</code> is computed by the companion command <code>kdbwselect</code> .
<code>rho</code>	Sets $b=h/\rho$. Default is <code>rho = 1</code> .
<code>kernel</code>	kernel function used to construct kernel density estimators. Options are <code>epa</code> for the epanechnikov kernel and <code>uni</code> for the uniform kernel. Default is <code>kernel = epa</code> .

bwselect	<p>bandwidth selection procedure to be used via <code>kdbwselect</code>. By default it computes h and sets $b=h/\rho$ (with $\rho=1$ by default). It computes both h and b if ρ is set equal to zero. Options are:</p> <p><code>mse-dpi</code> second-generation DPI implementation of MSE-optimal bandwidth. Default option if only one evaluation point is chosen.</p> <p><code>imse-dpi</code> second-generation DPI implementation of IMSE-optimal bandwidth (computed using a grid of evaluation points). Default option if more than one evaluation point is chosen.</p> <p><code>imse-rot</code> ROT implementation of IMSE-optimal bandwidth (computed using a grid of evaluation points).</p> <p><code>ce-dpi</code> second generation DPI implementation of CE-optimal bandwidth.</p> <p><code>ce-rot</code> ROT implementation of CE-optimal bandwidth.</p> <p>Use <code>kdbwselect</code> with <code>bwselect="all"</code> to report all available bandwidth selection procedures.</p> <p>Note: MSE = Mean Square Error; IMSE = Integrated Mean Squared Error; CE = Coverage Error; DPI = Direct Plug-in; ROT = Rule-of-Thumb. For details on implementation see Calonico, Cattaneo and Farrell (2019).</p>
bwcheck	if a positive integer is provided, then the selected bandwidth is enlarged so that at least <code>bwcheck</code> effective observations are available at each evaluation point. Default is <code>bwcheck = 21</code> .
imsegrid	number of evaluations points used to compute the IMSE bandwidth selector. Default is <code>imsegrid = 30</code> .
level	confidence level used for confidence intervals; default is <code>level = 95</code> .
subset	optional rule specifying a subset of observations to be used.
data	an optional data frame. When supplied, <code>x</code> and <code>subset</code> may be given as bare variable names referring to columns of data.

Value

Estimate	A matrix containing <code>eval</code> (grid points), <code>h</code> , <code>b</code> (bandwidths), <code>N</code> (effective sample sizes), <code>tau.us</code> (conventional density estimate), <code>tau.bc</code> (bias-corrected density estimate), <code>se.us</code> (standard error corresponding to <code>tau.us</code>), and <code>se.rb</code> (robust standard error).
opt	A list containing options passed to the function.

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References

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Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2019. **nprobust: Nonparametric Kernel-Based Estimation and Robust Bias-Corrected Inference**. *Journal of Statistical Software*, 91(8): 1-33. doi:10.18637/jss.v091.i08.

Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2022. **Coverage Error Optimal Confidence Intervals for Local Polynomial Regression**. *Bernoulli*, 28(4): 2998-3022.

See Also

[kdbwselect](#)

Examples

```
x <- rnorm(500)
est <- kdrobust(x)
summary(est)
```

lpbwselect	<i>Bandwidth Selection Procedures for Local Polynomial Regression Estimation and Inference</i>
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Description

[lpbwselect](#) implements bandwidth selectors for local polynomial regression point estimators and inference procedures developed in Calonico, Cattaneo and Farrell (2018). See also Calonico, Cattaneo and Farrell (2022) for related optimality results. It also implements other bandwidth selectors available in the literature.

Companion commands: [lprobust](#) for local polynomial point estimation and inference procedures.

A detailed introduction to this command is given in Calonico, Cattaneo and Farrell (2019). For more details and related software useful for empirical analysis, visit <https://nppackages.github.io/>.

Usage

```
lpbwselect(y, x, eval = NULL, neval = NULL, p = NULL, deriv = NULL,
kernel = "epa", bwselect = "mse-dpi", bwcheck = 21, bwregul = 1,
imsegrid = 30, vce = "nn", cluster = NULL,
nnmatch = 3, interior = FALSE, subset = NULL,
weights = NULL, masspoints = "check", data = NULL)
```

Arguments

y	dependent variable.
x	independent variable.
eval	vector of evaluation point(s). By default it uses 30 equally spaced points over the support of x.
neval	number of equally spaced evaluation points on the support of x. Default is neval=30.

p	polynomial order used to construct point estimator; default is $p = 1$ (local linear regression).
deriv	derivative order of the regression function to be estimated. Default is <code>deriv=0</code> (regression function).
kernel	kernel function used to construct local polynomial estimators. Options are <code>epa</code> for the epanechnikov kernel, <code>tri</code> for the triangular kernel, <code>uni</code> for the uniform kernel and <code>gau</code> for the gaussian kernel. Default is <code>kernel = epa</code> .
bwselect	bandwidth selection procedure to be used. Options are: <code>mse-dpi</code> second-generation DPI implementation of MSE-optimal bandwidth. Default option. <code>mse-rot</code> ROT implementation of MSE-optimal bandwidth. <code>imse-dpi</code> second-generation DPI implementation of IMSE-optimal bandwidth (computed using grid of evaluation points selected). <code>imse-rot</code> ROT implementation of IMSE-optimal bandwidth (computed using grid of evaluation points selected). <code>ce-dpi</code> second generation DPI implementation of CE-optimal bandwidth. <code>ce-rot</code> ROT implementation of CE-optimal bandwidth. <code>all</code> reports all available bandwidth selection procedures. Note: MSE = Mean Square Error; IMSE = Integrated Mean Squared Error; CE = Coverage Error; DPI = Direct Plug-in; ROT = Rule-of-Thumb. For details on implementation see Calonico, Cattaneo and Farrell (2019).
bwcheck	if a positive integer is provided, then the selected bandwidth is enlarged so that at least <code>bwcheck</code> effective observations are available at each evaluation point. Default is <code>bwcheck = 21</code> .
bwregul	specifies scaling factor for the regularization term added to the denominator of bandwidth selectors. Setting <code>bwregul = 0</code> removes the regularization term from the bandwidth selectors. Default is <code>bwregul = 1</code> .
imsegrid	number of evaluations points used to compute the IMSE bandwidth selector. Default is <code>imsegrid = 30</code> .
vce	procedure used to compute the variance-covariance matrix estimator. Options are: <code>nn</code> heteroskedasticity-robust nearest neighbor variance estimator with <code>nnmatch</code> the (minimum) number of neighbors to be used. Default choice (when <code>cluster</code> is not supplied). <code>hc0</code> heteroskedasticity-robust plug-in residuals variance estimator without weights. <code>hc1</code> heteroskedasticity-robust plug-in residuals variance estimator with <code>hc1</code> weights. <code>hc2</code> heteroskedasticity-robust plug-in residuals variance estimator with <code>hc2</code> weights. <code>hc3</code> heteroskedasticity-robust plug-in residuals variance estimator with <code>hc3</code> weights. <code>cr1</code> cluster-robust variance estimator with $((n-1)/(n-k)) * (G/(G-1))$ multiplier (Stata-style CR1). Default choice when <code>cluster</code> is supplied. Requires <code>cluster</code> . <code>cr2</code> cluster-robust variance estimator with Bell-McCaffrey block-adjusted residuals. Requires <code>cluster</code> .

	cr3 cluster-robust variance estimator with block jackknife-style adjustment and $(G-1)/G$ multiplier. Requires <code>cluster</code> . When <code>cluster</code> is supplied, <code>vce=hc0</code> or <code>vce=hc1</code> is remapped to <code>cr1</code> , <code>vce=hc2</code> to <code>cr2</code> , and <code>vce=hc3</code> to <code>cr3</code> (with a warning). <code>vce=nn</code> with <code>cluster</code> silently defaults to <code>cr1</code> .
<code>cluster</code>	indicates the cluster ID variable used for cluster-robust variance estimation. When supplied, the default <code>vce</code> becomes <code>cr1</code> ; <code>cr2</code> and <code>cr3</code> are also available.
<code>nnmatch</code>	to be combined with <code>for</code> <code>vce=nn</code> for heteroskedasticity-robust nearest neighbor variance estimator with <code>nnmatch</code> indicating the minimum number of neighbors to be used. Default is <code>nnmatch=3</code> .
.	
<code>interior</code>	if TRUE, all evaluation points are assumed to be interior points. This option affects only data-driven bandwidth selection via <code>lpbwselect</code> . Default is <code>interior = FALSE</code> .
<code>subset</code>	optional rule specifying a subset of observations to be used.
<code>weights</code>	optional vector of non-negative observation weights (multiplicative with kernel weights in all bandwidth-selection steps).
<code>masspoints</code>	how to handle evaluation points with few unique <code>x</code> values within bandwidth. Options: "check" (default), "off".
<code>data</code>	an optional data frame. When supplied, <code>y</code> , <code>x</code> , <code>cluster</code> , <code>weights</code> , and <code>subset</code> may be given as bare variable names referring to columns of data.

Value

<code>bws</code>	A matrix containing <code>eval</code> (grid points), <code>h</code> and <code>b</code> (selected bandwidths) for each evaluation point. When <code>bwselect="all"</code> , the matrix is widened to include both MSE and CER bandwidths from each of the DPI and ROT estimators.
<code>bws.imse</code>	IMSE-optimal bandwidths returned when <code>bwselect="all"</code> ; NULL otherwise.
<code>opt</code>	A list containing options passed to the function.

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References

- Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2018. [On the Effect of Bias Estimation on Coverage Accuracy in Nonparametric Inference](#). *Journal of the American Statistical Association*, 113(522): 767-779. doi:10.1080/01621459.2017.1285776.
- Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2019. [nprobust: Nonparametric Kernel-Based Estimation and Robust Bias-Corrected Inference](#). *Journal of Statistical Software*, 91(8): 1-33. doi:10.18637/jss.v091.i08.
- Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2022. [Coverage Error Optimal Confidence Intervals for Local Polynomial Regression](#). *Bernoulli*, 28(4): 2998-3022.

See Also[lprobust](#)**Examples**

```
x <- runif(500)
y <- sin(4*x) + rnorm(500)
est <- lpbwselect(y,x)
summary(est)
```

lprobust

*Local Polynomial Methods with Robust Bias-Corrected Inference***Description**

[lprobust](#) implements local polynomial regression point estimators, with robust bias-corrected confidence intervals and inference procedures developed in Calonico, Cattaneo and Farrell (2018). See also Calonico, Cattaneo and Farrell (2022) for related optimality results. It also implements other estimation and inference procedures available in the literature.

Companion commands: [lpbwselect](#) for local polynomial data-driven bandwidth selection, and [nprobust.plot](#) for plotting results.

A detailed introduction to this command is given in Calonico, Cattaneo and Farrell (2019). For more details and related software useful for empirical analysis, visit <https://nppackages.github.io/>.

Usage

```
lprobust(y, x, eval = NULL, neval = NULL, p = NULL, deriv = NULL,
h = NULL, b = NULL, rho = 1, kernel = "epa", bwselect = NULL,
bwcheck = 21, bwregul = 1, imsegrid = 30, vce = "nn", covgrid = FALSE,
cluster = NULL, nmmatch = 3, level = 95, interior = FALSE, subset = NULL,
weights = NULL, masspoints = "check", data = NULL)
```

Arguments

y	dependent variable.
x	independent variable.
eval	vector of evaluation point(s). By default it uses 30 equally spaced points over the support of x.
neval	number of equally spaced evaluation points on the support of x. Default is neval=30.
p	polynomial order used to construct point estimator; default is p = 1 (local linear regression).
deriv	derivative order of the regression function to be estimated. Default is deriv=0 (regression function).

h	main bandwidth used to construct local polynomial point estimator. Can be either scalar (same bandwidth for all evaluation points), or vector of same dimension as eval. If not specified, bandwidth h is computed by the companion command <code>lpbwselect</code> .
b	bias bandwidth used to construct the bias-correction estimator. Can be either scalar (same bandwidth for all evaluation points), or vector of same dimension as eval. By default it is set equal to h. If rho is set to zero, b is computed by the companion command <code>lpbwselect</code> .
rho	Sets $b=h/\rho$. Default is $\rho = 1$.
kernel	kernel function used to construct local polynomial estimators. Options are <code>epa</code> for the epanechnikov kernel, <code>tri</code> for the triangular kernel, <code>uni</code> for the uniform kernel and <code>gau</code> for the gaussian kernel. Default is <code>kernel = epa</code> .
bwselect	bandwidth selection procedure to be used via <code>lpbwselect</code> . By default it computes h and sets $b=h/\rho$ (with $\rho=1$ by default). It computes both h and b if rho is set equal to zero. Options are: <code>mse-dpi</code> second-generation DPI implementation of MSE-optimal bandwidth. Default option if only one evaluation point is chosen. <code>mse-rot</code> ROT implementation of MSE-optimal bandwidth. <code>imse-dpi</code> second-generation DPI implementation of IMSE-optimal bandwidth (computed using a grid of evaluation points). Default option if more than one evaluation point is chosen. <code>imse-rot</code> ROT implementation of IMSE-optimal bandwidth (computed using a grid of evaluation points). <code>ce-dpi</code> second generation DPI implementation of CE-optimal bandwidth. <code>ce-rot</code> ROT implementation of CE-optimal bandwidth. Use <code>lpbwselect</code> with <code>bwselect="all"</code> to report all available bandwidth selection procedures. Note: MSE = Mean Square Error; IMSE = Integrated Mean Squared Error; CE = Coverage Error; DPI = Direct Plug-in; ROT = Rule-of-Thumb. For details on implementation see Calonico, Cattaneo and Farrell (2019).
bwcheck	if a positive integer is provided, then the selected bandwidth is enlarged so that at least bwcheck effective observations are available at each evaluation point. Default is <code>bwcheck = 21</code> .
bwregul	specifies scaling factor for the regularization term added to the denominator of bandwidth selectors. Setting <code>bwregul = 0</code> removes the regularization term from the bandwidth selectors. Default is <code>bwregul = 1</code> .
imsegrid	number of evaluations points used to compute the IMSE bandwidth selector. Default is <code>imsegrid = 30</code> .
vce	procedure used to compute the variance-covariance matrix estimator. Options are: <code>nn</code> heteroskedasticity-robust nearest neighbor variance estimator with <code>nnmatch</code> the (minimum) number of neighbors to be used. Default choice (when <code>cluster</code> is not supplied). <code>hc0</code> heteroskedasticity-robust plug-in residuals variance estimator without weights.

hc1 heteroskedasticity-robust plug-in residuals variance estimator with hc1 weights.
 hc2 heteroskedasticity-robust plug-in residuals variance estimator with hc2 weights.
 hc3 heteroskedasticity-robust plug-in residuals variance estimator with hc3 weights.
 cr1 cluster-robust variance estimator with $((n-1)/(n-k)) * (G/(G-1))$ multiplier (Stata-style CR1). Default choice when `cluster` is supplied. Requires `cluster`.
 cr2 cluster-robust variance estimator with Bell-McCaffrey block-adjusted residuals. Requires `cluster`.
 cr3 cluster-robust variance estimator with block jackknife-style adjustment and $(G-1)/G$ multiplier. Requires `cluster`.
 When `cluster` is supplied, `vce=hc0` or `vce=hc1` is remapped to `cr1`, `vce=hc2` to `cr2`, and `vce=hc3` to `cr3` (with a warning). `vce=nn` with `cluster` silently defaults to `cr1`.

<code>covgrid</code>	if TRUE, it computes two covariance matrices (<code>cov.us</code> and <code>cov.rb</code>) for classical and robust covariances across point estimators over the grid of evaluation points.
<code>cluster</code>	indicates the cluster ID variable used for cluster-robust variance estimation. When supplied, the default <code>vce</code> becomes <code>cr1</code> ; <code>cr2</code> and <code>cr3</code> are also available.
<code>nnmatch</code>	to be combined with <code>for</code> <code>vce=nn</code> for heteroskedasticity-robust nearest neighbor variance estimator with <code>nnmatch</code> indicating the minimum number of neighbors to be used. Default is <code>nnmatch=3</code>
.	
<code>level</code>	confidence level used for confidence intervals; default is <code>level = 95</code> .
<code>interior</code>	if TRUE, all evaluation points are assumed to be interior points. This option affects only data-driven bandwidth selection via <code>lpbwselect</code> . Default is <code>interior = FALSE</code> .
<code>subset</code>	optional rule specifying a subset of observations to be used.
<code>weights</code>	optional vector of non-negative observation weights of the same length as <code>x</code> . User weights multiply the kernel weights in estimation, bandwidth selection, and variance computation (weighted least squares interpretation).
<code>masspoints</code>	how to handle evaluation points whose bandwidth window contains few unique <code>x</code> values (the local polynomial may become unreliable in that case). Options: "check" (default) warns when there are fewer than <code>p+5</code> unique values inside the main bandwidth; "off" disables the check.
<code>data</code>	an optional data frame. When supplied, <code>y</code> , <code>x</code> , <code>cluster</code> , <code>weights</code> , and <code>subset</code> may be given as bare variable names referring to columns of data.

Value

<code>Estimate</code>	A matrix containing <code>eval</code> (grid points), <code>h</code> , <code>b</code> (bandwidths), <code>N</code> (effective sample sizes), <code>tau.us</code> (point estimates with <code>p</code> -th order local polynomial), <code>tau.bc</code> (bias-corrected point estimates with $(p+1)$ -th order local polynomial), <code>se.us</code> (standard error corresponding to <code>tau.us</code>), and <code>se.rb</code> (robust standard error).
<code>cov.us</code>	Conventional-estimator covariance matrix across the evaluation grid (<code>neval</code> by <code>neval</code>). Returned only when <code>covgrid=TRUE</code> ; NULL otherwise.

cov.rb	Robust-bias-corrected covariance matrix across the evaluation grid (neval by neval). Returned only when covgrid=TRUE; NULL otherwise.
opt	A list containing options passed to the function.

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References

Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2018. [On the Effect of Bias Estimation on Coverage Accuracy in Nonparametric Inference](#). *Journal of the American Statistical Association*, 113(522): 767-779. doi:10.1080/01621459.2017.1285776.

Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2019. [nprobust: Nonparametric Kernel-Based Estimation and Robust Bias-Corrected Inference](#). *Journal of Statistical Software*, 91(8): 1-33. doi:10.18637/jss.v091.i08.

Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2022. [Coverage Error Optimal Confidence Intervals for Local Polynomial Regression](#). *Bernoulli*, 28(4): 2998-3022.

See Also

[lpbwselect](#)

Examples

```
x <- runif(500)
y <- sin(4*x) + rnorm(500)
est <- lprobust(y,x)
summary(est)
```

nprobust.plot

Graphical Presentation of Results from nprobust Package.

Description

nprobust.plot plots estimated density and regression function using the nprobust package. A detailed introduction to this command is given in Calonico, Cattaneo and Farrell (2019).

Companion commands: [lprobust](#) for local polynomial point estimation and inference procedures, and [kdrobust](#) for kernel density point estimation and inference procedures.

For more details and related software useful for empirical analysis, visit <https://nppackages.github.io/>.

Usage

```
nprobust.plot(..., alpha = NULL, type = NULL, CItyp = NULL,
  title = "", xlabel = "", ylabel = "", lty = NULL, lwd = NULL,
  lcol = NULL, pty = NULL, pwd = NULL, pcol = NULL, CIshade = NULL,
  Cicol = NULL, legendTitle = NULL, legendGroups = NULL)
```

Arguments

...	Objects returned by kdrobust or lprobust .
alpha	Numeric scalar between 0 and 1, the significance level for plotting confidence regions. If more than one is provided, they will be applied to data series accordingly.
type	String, one of "line" (default), "points" or "both", how the point estimates are plotted. If more than one is provided, they will be applied to data series accordingly.
CItyp	String, one of "region" (shaded region, default), "line" (dashed lines), "ebar" (error bars), "all" (all of the previous) or "none" (no confidence region), how the confidence region should be plotted. If more than one is provided, they will be applied to data series accordingly.
title, xlabel, ylabel	Strings, title of the plot and labels for x- and y-axis.
lty	Line type for point estimates, only effective if type is "line" or "both". 1 for solid line, 2 for dashed line, 3 for dotted line. For other options, see the instructions for ggplot2 or par . If more than one is provided, they will be applied to data series accordingly.
lwd	Line width for point estimates, only effective if type is "line" or "both". Should be strictly positive. For other options, see the instructions for ggplot2 or par . If more than one is provided, they will be applied to data series accordingly.
lcol	Line color for point estimates, only effective if type is "line" or "both". 1 for black, 2 for red, 3 for green, 4 for blue. For other options, see the instructions for ggplot2 or par . If more than one is provided, they will be applied to data series accordingly.
pty	Scatter plot type for point estimates, only effective if type is "points" or "both". For options, see the instructions for ggplot2 or par . If more than one is provided, they will be applied to data series accordingly.
pwd	Scatter plot size for point estimates, only effective if type is "points" or "both". Should be strictly positive. If more than one is provided, they will be applied to data series accordingly.
pcol	Scatter plot color for point estimates, only effective if type is "points" or "both". 1 for black, 2 for red, 3 for green, 4 for blue. For other options, see the instructions for ggplot2 or par . If more than one is provided, they will be applied to data series accordingly.
CIshade	Numeric, opaqueness of the confidence region, should be between 0 (transparent) and 1. Default is 0.2. If more than one is provided, they will be applied to data series accordingly.

CIcol	color for confidence region. 1 for black, 2 for red, 3 for green, 4 for blue. For other options, see the instructions for ggplot2 or par . If more than one is provided, they will be applied to data series accordingly.
legendTitle	String, title of legend.
legendGroups	String Vector, group names used in legend.

Details

Companion commands: [lprobust](#) for local polynomial point estimation and inference, and [kdrobust](#) for kernel density point estimation and inference.

Value

A standard [ggplot2](#) object is returned, hence can be used for further customization.

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References

Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2019. [nprobust: Nonparametric Kernel-Based Estimation and Robust Bias-Corrected Inference](#). Journal of Statistical Software, 91(8): 1-33. [doi:10.18637/jss.v091.i08](https://doi.org/10.18637/jss.v091.i08).

See Also

[lprobust](#), [kdrobust](#), [ggplot2](#)

Examples

```
x <- runif(500)
y <- sin(4*x) + rnorm(500)
est <- lprobust(y,x)
nprobust.plot(est)
```

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