

Package ‘lpdensity’

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Title Local Polynomial Density Estimation and Inference

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Description Without imposing stringent distributional assumptions or shape restrictions, nonparametric estimation has been popular in economics and other social sciences for counterfactual analysis, program evaluation, and policy recommendations. This package implements a novel density (and derivatives) estimator based on local polynomial regressions: `lpdensity()` to construct local polynomial based density (and derivatives) estimator, and `lpbwdensity()` to perform data-driven bandwidth selection.

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lpdensity-package

lpdensity: Local Polynomial Density Estimation and Inference

Description

Without imposing stringent distributional assumptions or shape restrictions, nonparametric estimation has been popular in economics and other social sciences for counterfactual analysis, program evaluation, and policy recommendations. This package implements a novel density (and derivatives) estimator based on local polynomial regressions, documented in Cattaneo, Jansson and Ma (2020a) and Cattaneo, Jansson and Ma (2020b).

`lpdensity` implements the local polynomial regression based density (and derivatives) estimator. Robust bias-corrected inference methods, both pointwise (confidence intervals) and uniform (confidence bands), are also implemented. `lpbwdensity` implements the bandwidth selection methods. See Cattaneo, Jansson and Ma (2020c) for more implementation details and illustrations.

Related Stata and R packages useful for nonparametric estimation and inference are available at <https://sites.google.com/site/nppackages/>.

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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.

Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2020. **Coverage Error Optimal Confidence Intervals for Local Polynomial Regression**. Working paper.

Cattaneo, M. D., M. Jansson, and X. Ma. 2020a. **Simple Local Polynomial Density Estimators**. *Journal of the American Statistical Association*, forthcoming.

Cattaneo, M. D., M. Jansson, and X. Ma. 2020b. **Local Regression Distribution Estimators**. Working paper.

Cattaneo, M. D., M. Jansson, and X. Ma. 2020c. **lpdensity: Local Polynomial Density Estimation and Inference**. Working paper.

coef.lpbwdensity

Coef Method for Local Polynomial Density Bandwidth Selection

Description

The coef method for local polynomial density bandwidth selection objects.

Usage

```
## S3 method for class 'lpbwdensity'
coef(object, ...)
```

Arguments

object Class "lpbdensity" object, obtained by calling [lpbdensity](#).
 ... Other arguments.

Value

A matrix containing grid points and selected bandwidths.

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See Also

[lpbdensity](#) for data-driven bandwidth selection.
 Supported methods: [coef.lpbwdensity](#), [print.lpbwdensity](#), [summary.lpbwdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Construct bandwidth
coef(lpbwdensity(X))
```

coef.lpdensity

Coef Method for Local Polynomial Density Estimation and Inference

Description

The coef method for local polynomial density objects.

Usage

```
## S3 method for class 'lpdensity'
coef(object, ...)
```

Arguments

object Class "lpdensity" object, obtained by calling [lpdensity](#).
 ... Additional options.

Value

A matrix containing grid points and density estimates using p- and q-th order local polynomials.

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See Also

[lpdensity](#) for local polynomial density estimation.

Supported methods: [coef.lpdensity](#), [confint.lpdensity](#), [plot.lpdensity](#), [print.lpdensity](#), [summary.lpdensity](#), [vcov.lpdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Estimate density and report results
coef(lpdensity(data = X, bwselect = "imse-dpi"))
```

confint.lpdensity	<i>Confint Method for Local Polynomial Density Estimation and Inference</i>
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Description

The confint method for local polynomial density objects.

Usage

```
## S3 method for class 'lpdensity'
confint(object, parm = NULL, level = NULL, ...)
```

Arguments

object	Class "lpdensity" object, obtained by calling lpdensity .
parm	Integer, indicating which parameters are to be given confidence intervals.
level	Numeric scalar between 0 and 1, the significance level for computing confidence intervals
...	Additional options, including (i) grid specifies a subset of grid points to display the bandwidth; (ii) gridIndex specifies the indices of grid points to display the bandwidth (this is the same as parm); (iii) alpha specifies the significance level (this is 1-level); (iv) CIuniform specifies whether displaying pointwise confidence intervals (FALSE, default) or the uniform confidence band (TRUE); (v) CISimul specifies the number of simulations used to construct critical values (default is 2000).

Value

A matrix containing grid points and confidence interval end points using p- and q-th order local polynomials.

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See Also

[lpdensity](#) for local polynomial density estimation.

Supported methods: [coef.lpdensity](#), [confint.lpdensity](#), [plot.lpdensity](#), [print.lpdensity](#), [summary.lpdensity](#), [vcov.lpdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Estimate density and report 95% confidence intervals
est1 <- lpdensity(data = X, bwselect = "imse-dpi")
confint(est1)

# Report results for a subset of grid points
confint(est1, parm=est1$Estimate[4:10, "grid"])
confint(est1, grid=est1$Estimate[4:10, "grid"])
confint(est1, gridIndex=4:10)

# Report the 99% uniform confidence band
# Fix the seed for simulating critical values
set.seed(42); confint(est1, level=0.99, CIuniform=TRUE)
set.seed(42); confint(est1, alpha=0.01, CIuniform=TRUE)
```

lpbwdensity

Data-driven Bandwidth Selection for Local Polynomial Density Estimators

Description

[lpbwdensity](#) implements the bandwidth selection methods for local polynomial based density (and derivatives) estimation proposed and studied in Cattaneo, Jansson and Ma (2020a) and Cattaneo, Jansson and Ma (2020b). See Cattaneo, Jansson and Ma (2020c) for more implementation details and illustrations.

Companion command: [lpdensity](#) for estimation and robust bias-corrected inference.

Related Stata and R packages useful for nonparametric estimation and inference are available at <https://sites.google.com/site/nppackages/>.

Usage

```
lpbwdensity(data, grid = NULL, p = NULL, v = NULL,
  kernel = c("triangular", "uniform", "epanechnikov"),
  bwselect = c("mse-dpi", "imse-dpi", "mse-rot", "imse-rot"),
  massPoints = TRUE, stdVar = TRUE, regularize = TRUE,
  nLocalMin = NULL, nUniqueMin = NULL, Cweights = NULL,
  Pweights = NULL)
```

Arguments

<code>data</code>	Numeric vector or one dimensional matrix/data frame, the raw data.
<code>grid</code>	Numeric, specifies the grid of evaluation points. When set to default, grid points will be chosen as 0.05-0.95 percentiles of the data, with a step size of 0.05.
<code>p</code>	Nonnegative integer, specifies the order of the local polynomial used to construct point estimates. (Default is 2.)
<code>v</code>	Nonnegative integer, specifies the derivative of the distribution function to be estimated. 0 for the distribution function, 1 (default) for the density function, etc.
<code>kernel</code>	String, specifies the kernel function, should be one of "triangular", "uniform" or "epanechnikov".
<code>bwselect</code>	String, specifies the method for data-driven bandwidth selection. This option will be ignored if <code>bw</code> is provided. Can be (1) "mse-dpi" (default, mean squared error-optimal bandwidth selected for each grid point); or (2) "imse-dpi" (integrated MSE-optimal bandwidth, common for all grid points); (3) "mse-rot" (rule-of-thumb bandwidth with Gaussian reference model); and (4) "imse-rot" (integrated rule-of-thumb bandwidth with Gaussian reference model).
<code>massPoints</code>	TRUE (default) or FALSE, specifies whether point estimates and standard errors should be adjusted if there are mass points in the data.
<code>stdVar</code>	TRUE (default) or FALSE, specifies whether the data should be standardized for bandwidth selection.
<code>regularize</code>	TRUE (default) or FALSE, specifies whether the bandwidth should be regularized. When set to TRUE, the bandwidth is chosen such that the local region includes at least <code>nLocalMin</code> observations and at least <code>nUniqueMin</code> unique observations.
<code>nLocalMin</code>	Nonnegative integer, specifies the minimum number of observations in each local neighborhood. This option will be ignored if <code>regularize=FALSE</code> . Default is $20+p+1$.
<code>nUniqueMin</code>	Nonnegative integer, specifies the minimum number of unique observations in each local neighborhood. This option will be ignored if <code>regularize=FALSE</code> . Default is $20+p+1$.
<code>Cweights</code>	Numeric vector, specifies the weights used for counterfactual distribution construction. Should have the same length as the data. This option will be ignored if <code>bwselect</code> is "mse-rot" or "imse-rot".
<code>Pweights</code>	Numeric vector, specifies the weights used in sampling. Should have the same length as the data. This option will be ignored if <code>bwselect</code> is "mse-rot" or "imse-rot".

Value

<code>BW</code>	A matrix containing (1) <code>grid</code> (grid point), (2) <code>bw</code> (bandwidth), (3) <code>nh</code> (number of observations in each local neighborhood), and (4) <code>nhu</code> (number of unique observations in each local neighborhood).
<code>opt</code>	A list containing options passed to the function.

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References

- Cattaneo, M. D., M. Jansson, and X. Ma. 2020a. [Simple Local Polynomial Density Estimators](#). *Journal of the American Statistical Association*, forthcoming.
- Cattaneo, M. D., M. Jansson, and X. Ma. 2020b. [Local Regression Distribution Estimators](#). Working paper.
- Cattaneo, M. D., M. Jansson, and X. Ma. 2020c. [lpdfensity: Local Polynomial Density Estimation and Inference](#). Working paper.

See Also

Supported methods: [coef.lpbwdensity](#), [print.lpbwdensity](#), [summary.lpbwdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Construct bandwidth
bw1 <- lpbwdensity(X)
summary(bw1)

# Display bandwidths for a subset of grid points
summary(bw1, grid=bw1$BW[4:10, "grid"])
summary(bw1, gridIndex=4:10)
```

lpdfensity

Local Polynomial Density Estimation and Inference

Description

[lpdfensity](#) implements the local polynomial regression based density (and derivatives) estimator proposed in Cattaneo, Jansson and Ma (2020a). Robust bias-corrected inference methods, both pointwise (confidence intervals) and uniform (confidence bands), are also implemented following the results in Cattaneo, Jansson and Ma (2020a) and Cattaneo, Jansson and Ma (2020b). See Cattaneo, Jansson and Ma (2020c) for more implementation details and illustrations.

Companion command: [lpbwdensity](#) for bandwidth selection.

Related Stata and R packages useful for nonparametric estimation and inference are available at <https://sites.google.com/site/nppackages/>.

Usage

```
lpdfensity(data, grid = NULL, bw = NULL, p = NULL, q = NULL,
  v = NULL, kernel = c("triangular", "uniform", "epanechnikov"),
  scale = NULL, massPoints = TRUE, bwselect = c("mse-dpi",
  "imse-dpi", "mse-rot", "imse-rot"), stdVar = TRUE, regularize = TRUE,
  nLocalMin = NULL, nUniqueMin = NULL, Cweights = NULL,
  Pweights = NULL)
```

Arguments

<code>data</code>	Numeric vector or one dimensional matrix/data frame, the raw data.
<code>grid</code>	Numeric, specifies the grid of evaluation points. When set to default, grid points will be chosen as 0.05-0.95 percentiles of the data, with a step size of 0.05.
<code>bw</code>	Numeric, specifies the bandwidth used for estimation. Can be (1) a positive scalar (common bandwidth for all grid points); or (2) a positive numeric vector specifying bandwidths for each grid point (should be the same length as <code>grid</code>).
<code>p</code>	Nonnegative integer, specifies the order of the local polynomial used to construct point estimates. (Default is 2.)
<code>q</code>	Nonnegative integer, specifies the order of the local polynomial used to construct confidence intervals/bands (a.k.a. the bias correction order). Default is $p+1$. When set to be the same as <code>p</code> , no bias correction will be performed. Otherwise it should be strictly larger than <code>p</code> .
<code>v</code>	Nonnegative integer, specifies the derivative of the distribution function to be estimated. 0 for the distribution function, 1 (default) for the density function, etc.
<code>kernel</code>	String, specifies the kernel function, should be one of "triangular", "uniform", and "epanechnikov".
<code>scale</code>	Numeric, specifies how estimates are scaled. For example, setting this parameter to 0.5 will scale down both the point estimates and standard errors by half. Default is 1. This parameter is useful if only part of the sample is employed for estimation, and should not be confused with <code>Cweights</code> or <code>Pweights</code> .
<code>massPoints</code>	TRUE (default) or FALSE, specifies whether point estimates and standard errors should be adjusted if there are mass points in the data.
<code>bwselect</code>	String, specifies the method for data-driven bandwidth selection. This option will be ignored if <code>bw</code> is provided. Options are (1) "mse-dpi" (default, mean squared error-optimal bandwidth selected for each grid point); (2) "imse-dpi" (integrated MSE-optimal bandwidth, common for all grid points); (3) "mse-rot" (rule-of-thumb bandwidth with Gaussian reference model); and (4) "imse-rot" (integrated rule-of-thumb bandwidth with Gaussian reference model).
<code>stdVar</code>	TRUE (default) or FALSE, specifies whether the data should be standardized for bandwidth selection.
<code>regularize</code>	TRUE (default) or FALSE, specifies whether the bandwidth should be regularized. When set to TRUE, the bandwidth is chosen such that the local region includes at least <code>nLocalMin</code> observations and at least <code>nUniqueMin</code> unique observations.
<code>nLocalMin</code>	Nonnegative integer, specifies the minimum number of observations in each local neighborhood. This option will be ignored if <code>regularize=FALSE</code> . Default is $20+p+1$.
<code>nUniqueMin</code>	Nonnegative integer, specifies the minimum number of unique observations in each local neighborhood. This option will be ignored if <code>regularize=FALSE</code> . Default is $20+p+1$.
<code>Cweights,</code>	Numeric, specifies the weights used for counterfactual distribution construction. Should have the same length as the data.
<code>Pweights</code>	Numeric, specifies the weights used in sampling. Should have the same length as the data.

Details

Bias correction is only used for the construction of confidence intervals/bands, but not for point estimation. The point estimates, denoted by f_p , are constructed using local polynomial estimates of order p , while the centering of the confidence intervals/bands, denoted by f_q , are constructed using local polynomial estimates of order q . The confidence intervals/bands take the form: $[f_q - cv * SE(f_q), f_q + cv * SE(f_q)]$, where cv denotes the appropriate critical value and $SE(f_q)$ denotes an standard error estimate for the centering of the confidence interval/band. As a result, the confidence intervals/bands may not be centered at the point estimates because they have been bias-corrected. Setting q and p to be equal results on centered at the point estimate confidence intervals/bands, but requires undersmoothing for valid inference (i.e., (I)MSE-optimal bandwidth for the density point estimator cannot be used). Hence the bandwidth would need to be specified manually when $q=p$, and the point estimates will not be (I)MSE optimal. See Cattaneo, Jansson and Ma (2020a, 2020b) for details, and also Calonico, Cattaneo, and Farrell (2018, 2020) for robust bias correction methods.

Sometimes the density point estimates may lie outside of the confidence intervals/bands, which can happen if the underlying distribution exhibits high curvature at some evaluation point(s). One possible solution in this case is to increase the polynomial order p or to employ a smaller bandwidth.

Value

Estimate	A matrix containing (1) grid (grid points), (2) bw (bandwidths), (3) nh (number of observations in each local neighborhood), (4) nhu (number of unique observations in each local neighborhood), (5) f_p (point estimates with p -th order local polynomial), (6) f_q (point estimates with q -th order local polynomial, only if option q is nonzero), (7) se_p (standard error corresponding to f_p), and (8) se_q (standard error corresponding to f_q).
CovMat_p	The variance-covariance matrix corresponding to f_p .
CovMat_q	The variance-covariance matrix corresponding to f_q .
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.
- Calonico, S., M. D. Cattaneo, and M. H. Farrell. 2020. **Coverage Error Optimal Confidence Intervals for Local Polynomial Regression**. Working paper.
- Cattaneo, M. D., M. Jansson, and X. Ma. 2020a. **Simple Local Polynomial Density Estimators**. *Journal of the American Statistical Association*, forthcoming.
- Cattaneo, M. D., M. Jansson, and X. Ma. 2020b. **Local Regression Distribution Estimators**. Working paper.
- Cattaneo, M. D., M. Jansson, and X. Ma. 2020c. **lpdfensity: Local Polynomial Density Estimation and Inference**. Working paper.

See Also

Supported methods: [coef.lpdensity](#), [confint.lpdensity](#), [plot.lpdensity](#), [print.lpdensity](#), [summary.lpdensity](#), [vcov.lpdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Estimate density and report results
est1 <- lpdensity(data = X, bwselect = "imse-dpi")
summary(est1)

# Report results for a subset of grid points
summary(est1, grid=est1$Estimate[4:10, "grid"])
summary(est1, gridIndex=4:10)

# Report the 99% uniform confidence band
set.seed(42) # fix the seed for simulating critical values
summary(est1, alpha=0.01, CIuniform=TRUE)

# Plot the estimates and confidence intervals
plot(est1, legendTitle="My Plot", legendGroups=c("X"))

# Plot the estimates and the 99% uniform confidence band
set.seed(42) # fix the seed for simulating critical values
plot(est1, alpha=0.01, CIuniform=TRUE, legendTitle="My Plot", legendGroups=c("X"))

# Adding a histogram to the background
plot(est1, legendTitle="My Plot", legendGroups=c("X"),
     hist=TRUE, histData=X, histBreaks=seq(-1.5, 1.5, 0.25))
```

lpdensity.plot

Plot Method for Local Polynomial Density Estimation and Inference

Description

This has been replaced by [plot.lpdensity](#).

Usage

```
lpdensity.plot(..., alpha = NULL, type = NULL, lty = NULL,
  lwd = NULL, lcol = NULL, pty = NULL, pwd = NULL, pcol = NULL,
  grid = NULL, CIttype = NULL, CIuniform = FALSE, CISimul = 2000,
  CIs shade = NULL, Cicol = NULL, hist = FALSE, histData = NULL,
  histBreaks = NULL, histFillCol = 3, histFillShade = 0.2,
  histLineCol = "white", title = "", xlabel = "", ylabel = "",
  legendTitle = NULL, legendGroups = NULL)
```

Arguments

...	Class "lpdfensity" object, obtained from calling lpdfensity .
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.
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.
grid	Numeric vector, specifies a subset of grid points to plot point estimates.
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.
CIuniform	TRUE or FALSE (default), plotting either pointwise confidence intervals (FALSE) or uniform confidence bands (TRUE).
CIsimul	Positive integer, the number of simulations used to construct critical values (default is 2000). This option is ignored if CIuniform=FALSE.
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 each data series accordingly.
CIcol	Color of the 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.
hist	TRUE or FALSE (default), whether adding a histogram to the background.
histData	Numeric vector, the data used to construct the histogram plot.
histBreaks	Numeric vector, giving the breakpoints between histogram cells.

histFillCol	Color of the histogram cells.
histFillShade	Opaqueness of the histogram cells, should be between 0 (transparent) and 1. Default is 0.2.
histLineCol	Color of the histogram lines.
title, xlabel, ylabel	Strings, title of the plot and labels for x- and y-axis.
legendTitle	String, title of legend.
legendGroups	String Vector, group names used in legend.

Value

A standard ggplot object is returned, hence can be used for further customization.

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plot.lpdensity

Plot Method for Local Polynomial Density Estimation and Inference

Description

The plot method for local polynomial density objects.

Usage

```
## S3 method for class 'lpdensity'
plot(..., alpha = NULL, type = NULL, lty = NULL,
      lwd = NULL, lcol = NULL, pty = NULL, pwd = NULL, pcol = NULL,
      grid = NULL, CItpe = NULL, CIuniform = FALSE, CIsimul = 2000,
      CIs shade = NULL, Cicol = NULL, hist = FALSE, histData = NULL,
      histBreaks = NULL, histFillCol = 3, histFillShade = 0.2,
      histLineCol = "white", title = "", xlabel = "", ylabel = "",
      legendTitle = NULL, legendGroups = NULL)
```

Arguments

...	Class "lpdensity" object, obtained from calling lpdensity .
alpha	Numeric scalar between 0 and 1, specifies the significance level for plotting confidence intervals/bands. If more than one is provided, they will be applied to each data series accordingly.
type	String, one of "line" (default), "points" and "both", specifies how the point estimates are plotted. If more than one is provided, they will be applied to each data series accordingly.

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 each 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 each 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 each 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 each 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 each 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 each data series accordingly.
grid	Numeric vector, specifies a subset of grid points to plot point estimates.
CItype	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 each data series accordingly.
CIuniform	TRUE or FALSE (default), plotting either pointwise confidence intervals (FALSE) or uniform confidence bands (TRUE).
CIstimul	Positive integer, specifies the number of simulations used to construct critical values (default is 2000). This option is ignored if CIuniform=FALSE.
CIshade	Numeric, specifies the 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 each data series accordingly.
CIcol	Color of the 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 each data series accordingly.
hist	TRUE or FALSE (default), specifies whether a histogram should be added to the background.
histData	Numeric vector, specifies the data used to construct the histogram plot.
histBreaks	Numeric vector, specifies the breakpoints between histogram cells.
histFillCol	Color of the histogram cells.
histFillShade	Opaqueness of the histogram cells, should be between 0 (transparent) and 1. Default is 0.2.
histLineCol	Color of the histogram lines.
title, xlabel, ylabel	Strings, specifies the title of the plot and labels for the x- and y-axis.
legendTitle	String, specifies the legend title.
legendGroups	String vector, specifies the group names used in legend.

Value

A standard ggplot object is returned, hence can be used for further customization.

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See Also

[lpdensity](#) for local polynomial density estimation.

Supported methods: [coef.lpdensity](#), [confint.lpdensity](#), [plot.lpdensity](#), [print.lpdensity](#), [summary.lpdensity](#), [vcov.lpdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Generate a density discontinuity at 0
X <- X - 0.5; X[X>0] <- X[X>0] * 2

# Density estimation, left of 0 (scaled by the relative sample size)
est1 <- lpdensity(data = X[X<=0], grid = seq(-2.5, 0, 0.05), bwselect = "imse-dpi",
  scale = sum(X<=0)/length(X))
# Density estimation, right of 0 (scaled by the relative sample size)
est2 <- lpdensity(data = X[X>0], grid = seq(0, 2, 0.05), bwselect = "imse-dpi",
  scale = sum(X>0)/length(X))

# Plot
plot(est1, est2, legendTitle="My Plot", legendGroups=c("Left", "Right"))

# Plot uniform confidence bands
set.seed(42) # fix the seed for simulating critical values
plot(est1, est2, legendTitle="My Plot", legendGroups=c("Left", "Right"), CIuniform=TRUE)

# Adding a histogram to the background
plot(est1, est2, legendTitle="My Plot", legendGroups=c("Left", "Right"),
  hist=TRUE, histBreaks=seq(-2.4, 2, 0.2), histData=X)

# Plot point estimates for a subset of evaluation points
plot(est1, est2, legendTitle="My Plot", legendGroups=c("Left", "Right"),
  type="both", CItype="all", grid=seq(-2, 2, 0.5))
```

print.lpbwdensity

Print Method for Local Polynomial Density Bandwidth Selection

Description

The print method for local polynomial density bandwidth selection objects.

Usage

```
## S3 method for class 'lpbdensity'
print(x, ...)
```

Arguments

x Class "lpbdensity" object, obtained by calling [lpbdensity](#).
 ... Other arguments.

Author(s)

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See Also

[lpbdensity](#) for data-driven bandwidth selection.
 Supported methods: [coef.lpbwdensity](#), [print.lpbwdensity](#), [summary.lpbwdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Construct bandwidth
print(lpbwdensity(X))
```

print.lpdensity

Print Method for Local Polynomial Density Estimation and Inference

Description

The print method for local polynomial density objects.

Usage

```
## S3 method for class 'lpdensity'
print(x, ...)
```

Arguments

x Class "lpdensity" object, obtained from calling [lpdensity](#).
 ... Additional options.

Author(s)

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See Also

[lpdensity](#) for local polynomial density estimation.

Supported methods: [coef.lpdensity](#), [confint.lpdensity](#), [plot.lpdensity](#), [print.lpdensity](#), [summary.lpdensity](#), [vcov.lpdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Estimate density and report results
print(lpdensity(data = X, bwselect = "imse-dpi"))
```

summary.lpbwdensity	<i>Summary Method for Local Polynomial Density Bandwidth Selection</i>
---------------------	--

Description

The summary method for local polynomial density bandwidth selection objects.

Usage

```
## S3 method for class 'lpbwdensity'
summary(object, ...)
```

Arguments

object	Class "lpbwdensity" object, obtained by calling lpbwdensity .
...	Additional options, including (i) <code>grid</code> specifies a subset of grid points to display the bandwidth; (ii) <code>gridIndex</code> specifies the indices of grid points to display the bandwidth.

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See Also

[lpbwdensity](#) for data-driven bandwidth selection.

Supported methods: [coef.lpbwdensity](#), [print.lpbwdensity](#), [summary.lpbwdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Construct bandwidth
bw1 <- lpbwdensity(X)
summary(bw1)

# Display bandwidths for a subset of grid points
summary(bw1, grid=bw1$BW[4:10, "grid"])
summary(bw1, gridIndex=4:10)
```

summary.lpdensity	<i>Summary Method for Local Polynomial Density Estimation and Inference</i>
-------------------	---

Description

The summary method for local polynomial density objects.

Usage

```
## S3 method for class 'lpdensity'
summary(object, ...)
```

Arguments

object	Class "lpdensity" object, obtained from calling lpdensity .
...	Additional options, including (i) grid specifies a subset of grid points to display results; (ii) gridIndex specifies the indices of grid points to display results; (iii) alpha specifies the significance level; (iv) CIuniform specifies whether displaying pointwise confidence intervals (FALSE, default) or the uniform confidence band (TRUE); (v) CIsimul specifies the number of simulations used to construct critical values (default is 2000).

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See Also

[lpdensity](#) for local polynomial density estimation.

Supported methods: [coef.lpdensity](#), [confint.lpdensity](#), [plot.lpdensity](#), [print.lpdensity](#), [summary.lpdensity](#), [vcov.lpdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Estimate density and report results
est1 <- lpdensity(data = X, bwselect = "imse-dpi")
summary(est1)

# Report results for a subset of grid points
summary(est1, grid=est1$Estimate[4:10, "grid"])
summary(est1, gridIndex=4:10)

# Report the 99% uniform confidence band
set.seed(42) # fix the seed for simulating critical values
summary(est1, alpha=0.01, CIuniform=TRUE)
```

vcov.lpdensity

*Vcov Method for Local Polynomial Density Estimation and Inference***Description**

The vcov method for local polynomial density objects.

Usage

```
## S3 method for class 'lpdensity'
vcov(object, ...)
```

Arguments

object	Class "lpdensity" object, obtained by calling lpdensity .
...	Additional options.

Value

stdErr	A matrix containing grid points and standard errors using p- and q-th order local polynomials.
CovMat_p	The variance-covariance matrix corresponding to f_p.
CovMat_q	The variance-covariance matrix corresponding to f_q.

Author(s)

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See Also

[lpdensity](#) for local polynomial density estimation.
 Supported methods: [coef.lpdensity](#), [confint.lpdensity](#), [plot.lpdensity](#), [print.lpdensity](#), [summary.lpdensity](#), [vcov.lpdensity](#).

Examples

```
# Generate a random sample
set.seed(42); X <- rnorm(2000)

# Estimate density and report results
vcov(lpdensity(data = X, bwselect = "imse-dpi"))
```

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