

# Package ‘lpcde’

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**Type** Package

**Title** Boundary Adaptive Local Polynomial Conditional Density Estimator

**Version** 0.1.7

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**Description** Tools for estimation and inference of conditional densities, derivatives and functions. This is the companion software for Cattaneo, Chandak, Jansson and Ma (2024) <[doi:10.3150/23-BEJ1711](https://doi.org/10.3150/23-BEJ1711)>.

**Depends** R (>= 3.3.0)

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** false

**SystemRequirements** GNU make

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**RoxygenNote** 7.3.2

**Imports** Rcpp (>= 0.12.8),  
ggplot2,  
purrr,  
MASS,  
mvtnorm,  
combinat,  
Matrix,  
Rdpack,  
stats

**RdMacros** Rdpack

**LinkingTo** Rcpp,  
RcppArmadillo

**Suggests** testthat (>= 3.0.0),  
covr

**Config/testthat/edition** 3

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basis_vec	<i>Unit basis vector</i>
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**Description**

Function to generate unit basis vector according to polynomial order and derivative order. This function returns unit vector that is the same size as the vector returned by poly\_base(x, p).

**Usage**

```
basis_vec(x, p, mu)
```

**Arguments**

- x                      Sample input scalar or vector.
- p                      Polynomial order.
- mu                     Derivative order.

**Value**

Vector of appropriate length with ones corresponding to entries of order mu.

**Examples**

```
basis_vec(x = 2, p = 5, mu = 1)
```

coef.lpbwcde

*Coef method for local polynomial density bandwidth selection***Description**

The coef method for local polynomial density bandwidth selection objects.

**Usage**

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

**Arguments**

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

**Value**

Matrix            A matrix containing y\_grid points and selected bandwidths.

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**See Also**

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

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

```
n=100 x_data = as.matrix(rnorm(n, mean=0, sd=1)) y_data = as.matrix(rnorm(n, mean=0, sd=1))
y_grid = stats::quantile(y_data, seq(from=0.1, to=0.9, by=0.1))bandwidth selection
```

```
y_grid = stats::quantile(y_data, seq(from=0.1, to=0.9, by=0.1)) model2 = lpcde::lpbwcde(y_data=y_data,
x_data=x_data, x=0, y_grid = y_grid, bw_type = "mse-rot") coef(model2)
```

coef.lpcde

*Coef method for local polynomial density conditional estimation***Description**

The coef method for local polynomial conditional density objects.

**Usage**

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

**Arguments**

object	Class "lpcde" object, obtained by calling <a href="#">lpcde</a> .
...	Additional options.

**Value**

outputs	A matrix containing the estimates
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**See Also**

[lpcde](#) for local polynomial conditional density estimation.

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

**Examples**

```
n <- 100
x_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
# density estimation
model1 <- lpcde::lpcde(x_data = x_data, y_data = y_data, y_grid = y_grid, x = 0, bw = 0.5)
coef(model1)
```

confint.lpcde

*Confint method for local polynomial density conditional estimation***Description**

The confint method for local polynomial conditional density objects.

**Usage**

```
## S3 method for class 'lpcde'
confint(
  object,
  parm = NULL,
  level = 0.95,
  CIuniform = FALSE,
  CIsimul = 2000,
  ...
)
```

**Arguments**

object	Class "lpcde" object, obtained by calling <code>lpcde</code> .
parm	Integer, indicating which parameters are to be given confidence intervals.
level	Numeric scalar between 0 and 1, the confidence level for computing confidence intervals/bands. Equivalent to (1-significance level).
CIuniform	TRUE or FALSE (default), plotting either pointwise confidence intervals (FALSE) or uniform confidence bands (TRUE).
CIsimul	Positive integer, specifies the number of simulations used to construct critical values (default is 2000). This option is ignored if CIuniform=FALSE.
...	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) CIuniform specifies whether displaying pointwise confidence intervals (FALSE, default) or the uniform confidence band (TRUE); (iv) CIsimul specifies the number of simulations used to construct critical values (default is 2000).

**Value**

Estimate	A matrix containing grid points, estimates and confidence interval end points using p- and q-th order local polynomials as well as bias-corrected estimates and corresponding confidence intervals.
crit_val	The critical value used in computing the confidence interval end points.

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## See Also

[lpcde](#) for local polynomial conditional density estimation.

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

## Examples

```
n <- 100
x_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
# density estimation
model1 <- lpcde::lpcde(x_data = x_data, y_data = y_data, y_grid = y_grid, x = 0, bw = 0.5)
confint(model1)
```

---

lpbwcde

*Data-driven bandwidth selection for local polynomial conditional density estimators*


---

## Description

[lpbwcde](#) implements the bandwidth selection methods for local polynomial based conditional density (and derivatives) estimation proposed and studied in (Cattaneo et al. 2024).

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

Related Stata and R packages useful for nonparametric estimation and inference are available at <https://nppackages.github.io/>.

## Usage

```
lpbwcde(
  y_data,
  x_data,
  x,
  y_grid = NULL,
  p = NULL,
  q = NULL,
  grid_spacing = "",
  ng = NULL,
  mu = NULL,
  nu = NULL,
  kernel_type = c("epanechnikov", "triangular", "uniform"),
  bw_type = c("imse-rot", "mse-rot"),
  regularize = NULL
)
```

**Arguments**

y_data	Numeric matrix/data frame, the raw data of independent.
x_data	Numeric matrix/data frame, the raw data of covariates.
x	Numeric, specifies the evaluation point in the x-direction. Default is median of the dataset.
y_grid	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.
p	Nonnegative integer, specifies the order of the local polynomial for Y used to construct point estimates. (Default is 2.)
q	Nonnegative integer, specifies the order of the local polynomial for X used to construct point estimates. (Default is 1.)
grid_spacing	String, If equal to "quantile" will generate quantile-spaced grid evaluation points, otherwise will generate equally spaced points.
ng	Int, number of grid points to be used in generating bandwidth estimates.
mu	Nonnegative integer, specifies the derivative with respect to Y of the distribution function to be estimated. 0 for the distribution function, 1 (default) for the density function, etc.
nu	Nonnegative integer, specifies the derivative with respect to X of the distribution function to be estimated.
kernel_type	String, specifies the kernel function, should be one of "triangular", "uniform" or "epanechnikov".
bw_type	String, specifies the method for data-driven bandwidth selection. This option will be ignored if bw is provided. Implementable with "mse-rot" (default, mean squared error-optimal bandwidth selected for each grid point)
regularize	Boolean (default TRUE). Option to regularize bandwidth selection to have at least $20 + \max(p, q) + 1$ datapoints when evaluating the estimator.

**Value**

BW	A matrix containing (1) y_grid (grid point), (2) bw (bandwidth)
opt	A list containing options passed to the function.

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**References**

Cattaneo MD, Chandak R, Jansson M, Ma X (2024). "Local Polynomial Conditional Density Estimators." *Bernoulli*.

**See Also**

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

## Examples

```
# Generate a random sample
set.seed(42)
x_data <- rnorm(2000)
y_data <- rnorm(2000, mean = x_data)
x <- 0

# Construct bandwidth
bw1 <- lpbwcd(y_data = y_data, x_data = x_data, x = x, bw_type = "mse-rot")
summary(bw1)

# Display bandwidths for a subset of y_grid points
summary(bw1, y_grid = bw1$BW[2:5, "y_grid"])
```

---

lpcde

*Local polynomial conditional density estimation*


---

## Description

[lpcde](#) implements the local polynomial regression based conditional density (and derivatives). The estimator proposed in (Cattaneo et al. 2024). Robust bias-corrected inference methods, both point-wise (confidence intervals) and uniform (confidence bands), are also implemented.

## Usage

```
lpcde(
  x_data,
  y_data,
  y_grid = NULL,
  x = NULL,
  bw = NULL,
  p = NULL,
  q = NULL,
  p_RBC = NULL,
  q_RBC = NULL,
  mu = NULL,
  nu = NULL,
  rbc = TRUE,
  ng = NULL,
  cov_flag = c("full", "diag", "off"),
  normalize = FALSE,
  nonneg = FALSE,
  grid_spacing = "",
  kernel_type = c("epanechnikov", "triangular", "uniform"),
  bw_type = NULL
)
```

## Arguments

`x_data`                      Numeric matrix/data frame, the raw data of covariates.



y_data	Numeric matrix/data frame, the raw data of independent.
y_grid	Numeric, specifies the grid of evaluation points in the y-direction. 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 in y-direction.
x	Numeric, specifies the grid of evaluation points in the x-direction. When set to default, the evaluation point will be chosen as the median of the x data. To generate estimates for multiple conditioning values, please loop over the x values and evaluate the lpcde function at each point.
bw	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/matrix specifying bandwidths for each grid point (should be the same dimension as grid).
p	Nonnegative integer, specifies the order of the local polynomial for Y used to construct point estimates. (Default is 2.)
q	Nonnegative integer, specifies the order of the local polynomial for X used to construct point estimates. (Default is 1.)
p_RBC	Nonnegative integer, specifies the order of the local polynomial for Y used to construct bias-corrected point estimates. (Default is p+1.)
q_RBC	Nonnegative integer, specifies the order of the local polynomial for X used to construct bias-corrected point estimates. (Default is q+1.)
mu	Nonnegative integer, specifies the derivative with respect to Y of the distribution function to be estimated. 0 for the distribution function, 1 (default) for the density function, etc.
nu	Nonnegative integer, specifies the derivative with respect to X of the distribution function to be estimated. Default value is 0.
rbc	Boolean. TRUE (default) for rbc calculations, required for valid uniform inference.
ng	Int, number of grid points to be used. generates evenly space points over the support of the data.
cov_flag	String, specifies covariance computation. Must be one of "full" (default), "diag" or "off".
normalize	Boolean, False (default) returns original estimator, True normalizes estimates to integrate to 1.
nonneg	Boolean, False (default) returns original estimator, True returns maximum of estimate and 0.
grid_spacing	String, If equal to "quantile" will generate quantile-spaced grid evaluation points, otherwise will generate equally spaced points.
kernel_type	String, specifies the kernel function, should be one of "triangular", "uniform", and "epanechnikov"(default).
bw_type	String, specifies the method for data-driven bandwidth selection. This option will be ignored if bw is provided. Implementable with "mse-dpi" (default, mean squared error-optimal bandwidth selected for each grid point)

## Details

Bias correction is only used for the construction of confidence intervals/bands, but not for point estimation. The point estimates, denoted by `est`, are constructed using local polynomial estimates of

order  $p$  and  $q$ , while the centering of the confidence intervals/bands, denoted by `est_RBC`, are constructed using local polynomial estimates of order  $p\_RBC$  and  $q\_RBC$ . The confidence intervals/bands take the form:  $[\text{est\_RBC} - \text{cv} * \text{SE}(\text{est\_RBC}), \text{est\_RBC} + \text{cv} * \text{SE}(\text{est\_RBC})]$ , where  $\text{cv}$  denotes the appropriate critical value and  $\text{SE}(\text{est\_RBC})$  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  $p\_RBC$  equal to  $p$  and  $q\_RBC$  to  $q$ , 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) est (point estimates with $p$ -th and $q$ -th order local polynomial), (4) est_RBC (point estimates with $p\_RBC$ -th and $q\_RBC$ -th order local polynomial), (5) se (standard error corresponding to est. Set to NA if <code>cov_flag="off"</code> ). (6) se_RBC (standard error corresponding to est_RBC). Set to NA if <code>cov_flag="off"</code>
CovMat	The variance-covariance matrix corresponding to est. Will be 0 if <code>cov_flag="off"</code> or a diagonal matrix if <code>cov_flag="diag"</code> .
opt	A list containing options passed to the function.

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### References

- Cattaneo MD, Chandak R, Jansson M, Ma X (2024). “Local Polynomial Conditional Density Estimators.” *Bernoulli*.
- Calonico S, Cattaneo MD, Farrell MH (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, Cattaneo MD, Farrell MH (2022). “Coverage Error Optimal Confidence Intervals for Local Polynomial Regression.” *Bernoulli*, **28**(4), 2998–3022.
- Cattaneo MD, Jansson M, Ma X (2020). “Simple local polynomial density estimators.” *J. Amer. Statist. Assoc.*, **115**(531), 1449–1455.

### See Also

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

## Examples

```
# Density estimation example
n <- 500
x_data <- matrix(rnorm(n, mean = 0, sd = 1))
y_data <- matrix(rnorm(n, mean = x_data, sd = 1))
y_grid <- seq(from = -1, to = 1, length.out = 5)
model1 <- lpcde::lpcde(x_data = x_data, y_data = y_data, y_grid = y_grid, x = 0, bw = 0.5)
# summary of estimation
summary(model1)
```

---

mvec	<i>Polynomial order vector</i>
------	--------------------------------

---

## Description

Generates list of all combinations of length less than or equal to d of numbers that add up to n.

## Usage

```
mvec(n, d)
```

## Arguments

n	Total value of each combination
d	Maximum length of combinations

---

plot.lpcde	<i>Plot method for local polynomial density conditional estimation</i>
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---

## Description

The plot method for local polynomial density objects. A standard ggplot2 object is returned, hence can be used for further customization.

## Usage

```
## S3 method for class 'lpcde'
plot(
  ...,
  alpha = NULL,
  type = NULL,
  lty = NULL,
  lwd = NULL,
  lcol = NULL,
  pty = NULL,
  pwd = NULL,
  pcol = NULL,
  y_grid = NULL,
```

```

CItpe = NULL,
CIuniform = FALSE,
CIsimul = 2000,
Cishade = NULL,
Cicol = NULL,
title = NULL,
xlabel = NULL,
ylabel = NULL,
legendTitle = NULL,
legendGroups = NULL,
rbc = FALSE
)

```

### Arguments

...	Class "lpcde" object, obtained from calling <code>lpcde</code> .
alpha	Numeric scalar between 0 and 1, specifies the significance level for plotting confidence intervals/bands.
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 <code>ggplot2</code> . 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 <code>ggplot2</code> . 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 <code>ggplot2</code> . 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 <code>ggplot2</code> . 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 <code>ggplot2</code> . If more than one is provided, they will be applied to each data series accordingly.
y_grid	Numeric vector, specifies a subset of grid points to plot point estimates. This option is effective only if type is "points" or "both"; or if CItpe is "ebar" or "all".
CItpe	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).
CIsimul	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 . If more than one is provided, they will be applied to each data series accordingly.
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.
rbc	TRUE or FALSE (default), plotting confidence intervals and bands with standard estimates (FALSE) or RBC estimates (TRUE).

**Value**

Figure	A standard ggplot2 object is returned, hence can be used for further customization.
--------	---

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 Xinwei Ma, University of California San Diego. <x1ma@ucsd.edu>.

**See Also**

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

---

poly_base	<i>Polynomial basis vector expansion</i>
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---

**Description**

Generate polynomial basis vector up to order p. has multivariate functionality as described in the main paper normalized by factorials in denominator. NOTE: currently works only up to 4th degree polynomial expansion for multivariate x.

**Usage**

```
poly_base(x, p)
```

**Arguments**

x                    A number or vector.  
 p                    A number (integer).

**Value**

Polynomial basis of x up to degree p.

**Examples**

```
poly_base(x = 2, p = 5)
```

---

print.lpbwcde	<i>Print method for local polynomial conditional density bandwidth selection</i>
---------------	--

---

**Description**

The print method for local polynomial conditional density bandwidth selection objects.

**Usage**

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

**Arguments**

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

**Value**

Display output    A list of specified options provided to the function.

**Author(s)**

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 Xinwei Ma, University of California San Diego. <x1ma@ucsd.edu>.

**See Also**

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

**Examples**

```
n <- 100
x_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
# bandwidth selection
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
model2 <- lpcde::lpbwcode(y_data = y_data, x_data = x_data, x = 0,
                        y_grid = y_grid, bw_type = "mse-rot")
print(model2)
```

---

print.lpcde	<i>Print method for local polynomial conditional density estimation</i>
-------------	---

---

**Description**

The print method for local polynomial conditional density objects.

**Usage**

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

**Arguments**

x	Class "lpcde" object, obtained from calling <a href="#">lpcde</a> .
...	Additional options.

**Value**

Display output    summary of inputs to lpcde

**Author(s)**

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**See Also**

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

**Examples**

```
n <- 100
x_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
# density estimation
model1 <- lpcde::lpcde(x_data = x_data, y_data = y_data, y_grid = y_grid, x = 0, bw = 0.5)
print(model1)
```

---

summary.lpbwcde	<i>Summary method for local polynomial conditional density bandwidth selection</i>
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---

**Description**

The summary method for local polynomial conditional density bandwidth selection objects.

**Usage**

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

**Arguments**

object	Class "lpbwcde" object, obtained by calling <a href="#">lpbwcde</a> .
...	Additional options, including (i) y_grid specifies a subset of y_grid points to display the bandwidth; (ii) gridIndex specifies the indices of y_grid points to display the bandwidth.

**Value**

Display output	A list of specified options and a matrix of grid points, bandwidth, and effective sample size.
----------------	--

**Author(s)**

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**See Also**

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



## Examples

```
n <- 100
x_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
# bandwidth selection
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
model2 <- lpcde::lpcwde(y_data = y_data, x_data = x_data, x = 0,
                       y_grid = y_grid, bw_type = "mse-rot")
summary(model2)
```

summary.lpcde

*Summary method for local polynomial density conditional estimation*

## Description

The summary method for local polynomial conditional density objects.

## Usage

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

## Arguments

object	Class "lpcde" object, obtained from calling <a href="#">lpcde</a> .
...	Additional options, including (i) y_grid specifies a subset of grid points in y-directions 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).

## Value

Display output    A list of specified options and a matrix of grid points and estimates.

## Author(s)

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## See Also

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

## Examples

```
n <- 100
x_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
# density estimation
model1 <- lpcde::lpcde(x_data = x_data, y_data = y_data, y_grid = y_grid, x = 0, bw = 0.5)
summary(model1)
```

---

vcov.lpcde	<i>Variance-Covariance</i>
------------	----------------------------

---

## Description

The vcov method for local polynomial conditional density objects.

## Usage

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

## Arguments

object	Class "lpcde" object, obtained by calling <a href="#">lpcde</a> .
...	Additional options.

## Details

Vcov method for local polynomial density conditional estimation

## Value

stdErr	A matrix containing grid points and standard errors using p- and q-th order local polynomials.
CovMat	The variance-covariance matrix corresponding to est.
CovMat_RBC	The variance-covariance matrix corresponding to est_RBC.

## Author(s)

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## See Also

[lpcde](#) for local polynomial conditional density estimation.  
 Supported methods: [plot.lpcde](#), [print.lpcde](#), [summary.lpcde](#),

**Examples**

```
n <- 100
x_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_data <- as.matrix(rnorm(n, mean = 0, sd = 1))
y_grid <- stats::quantile(y_data, seq(from = 0.1, to = 0.9, by = 0.1))
# density estimation
model1 <- lpcde::lpcde(x_data = x_data, y_data = y_data, y_grid = y_grid, x = 0, bw = 0.5)
vcov(model1)
```

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