

Package ‘llrRcpp’

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Type Package

Title Local linear regression

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Description This package provides an implementation of multivariate local linear regression using kd-tree implementation. Both an exact and approximate algorithm for using the kd-tree are present. The package also provides a binned method for the implementation of up to 2d.

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Encoding UTF-8

Imports Rcpp, methods, metaheuristicOpt

LinkingTo Rcpp, RcppEigen

RoxygenNote 7.1.1

Suggests rmarkdown,
knitr,
testthat

VignetteBuilder knitr

R topics documented:

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autoloocv.llr

*Auto selection of bandwidth using metaheuristic algorithms***Description**

Auto selection of bandwidth using metaheuristic algorithms

Usage

```
autoloocv.llr(
  x,
  y,
  weight,
  kernel = "epanechnikov",
  bw_range = c(0.01, 0.5),
  approx = FALSE,
  epsilon = 0.05,
  control = list(numPopulation = 15, maxIter = 75, Vmax = 2, ci = 1.49445, cg =
    1.49445, w = 0.729),
  algorithm = "PSO",
  seed = 1
)
```

Arguments

x	a numeric vector or matrix of x data
y	a numeric vector of y data corresponding to x data
weight	a numeric vector of length(x) for weight of each data point
kernel	kernel type used; supported are 'epanechnikov', 'rectangular', 'triangular', 'quartic', 'triweight', 'tricube', 'cosine', 'gauss', 'logistic'. 'sigmoid' and 'silverman'.
bw_range	range of bandwidth for the metaheuristic algorithm to search
approx	boolean flag: if true kdtree approximation is used.
epsilon	margin of error allowed for llr approximation using kdtree. Only used when approx = TRUE.
control	control parameter from R package metaheuristicOpt
algorithm	algorithm parameter from R package metaheuristicOpt
seed	seed to use for algorithm

Examples

```
## Not run:
n <- 1000
x <- runif(n, 0, 1)
y <- sin(x) + rnorm(n, 0, 0.2)
w <- rep(1/n, n)
## Optimization using Grasshopper Optimisation Algorithm
algorithm <- 'GOA'
control <- list(numPopulation = 10, maxIter = 100)
```

```
h <- autoloocv.llr(x, y, w, control = control , algorithm = algorithm)

## End(Not run)
```

bin

Splitting the data into bins

Description

Splitting the data into bins

Usage

```
bin(x, y, bins = 400, weight)
```

Arguments

x	numeric vector or matrix of 2 dimenons of x data
y	numeric vector of y data with length corresponding to x
bins	number of bins to split the data
weight	numeric vector corresponding to weight of each observation

Value

returns an S3 object of the class 'bin' containing

x a numeric vector of x data of size bins

y a numeric vector of y data corresponding to 'bin' class x

weight weight corresponding to 'bin' class x

Examples

```
n <- 1000
x <- seq(0,10,length.out = n)
x1 <- rnorm(n,0,0.2)
y <- sin(x) + x1
w <- rep(1/n, n)
binned <- bin(x,y,bins=400, w)
```

cv.llr

Bandwidth selection using k-fold cross validation

Description

Bandwidth selection using k-fold cross validation

Usage

```
cv.llr(
  x,
  y,
  weight,
  kernel = "epanechnikov",
  bandwidth,
  kdtree = FALSE,
  approx = FALSE,
  epsilon = 0.05,
  N_min = 1,
  k = 5
)
```

Arguments

x	a numeric vector or matrix of x data
y	a numeric vector of y data corresponding to x data
weight	a numeric vector of length(x) for weight of each data point.
kernel	kernel type used; supported are 'epanechnikov', "rectangular", "triangular", "quartic", "triweight", "tricube", "cosine", "gauss", "logistic". "sigmoid" and "silverman".
bandwidth	a numeric vector or matrix of bandwidth considered for selection.
kdtree	boolean flag: If TRUE, a kdtree is used for computation of local linear regression.
approx	boolean flag: if TRUE kdtree approximation is used.
epsilon	margin of error allowed for llr approximation using kdtree. Only used when kdtree = TRUE and approx = TRUE.
N_min	minimum number of points stored in the kd-tree. Only used when kdtree = TRUE and approx = TRUE.
k	number of folds used for cross validation

Value

returns a single numeric value or vector of bandwidth that gives the smallest mean square error from cross validation.

Examples

```

n <- 1000
x <- seq(0, 10, length.out = n)
x1 <- rnorm(n, 0, 0.2)
y <- sin(x) + x1
w <- rep(1 / n, n)
binned <- bin(x, y, bins = 400, w)
bandwidth <- seq(0.02, 0.3, by = 0.02)
## Bandwidth selection of binned data
h_bin <- cv.llr(binned$x, binned$y, binned$weight, bandwidth = bandwidth)
## Bandwidth selection of exact local linear regression
h_exact <- cv.llr(x, y, w, bandwidth = bandwidth)
## Bandwidth selection of exact local linear regression with kdtree
h_kdexact <- cv.llr(x, y, w, kdtree = TRUE, approx = FALSE, bandwidth = bandwidth)
## Bandwidth selection of approx local linear regression with kdtree
h_kdapprox <- cv.llr(x, y, w, kdtree = TRUE, approx = TRUE, bandwidth = bandwidth)

```

get_num_procs	<i>Get number of processors</i>
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Description

Get number of processors

Usage

```
get_num_procs()
```

Value

number of processors

llr	<i>Local linear regression (Regular and binned version)</i>
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Description

Local linear regression (Regular and binned version)

Usage

```

llr(x, ...)

## Default S3 method:
llr(
  x,
  y,
  xpred,
  kernel = "epanechnikov",
  bandwidth,

```

```

    weight,
    kdtree = FALSE,
    approx = FALSE,
    epsilon = 0.05,
    N_min = 1,
    ...
)

## S3 method for class 'bin'
llr(x, xpred, kernel = "epanechnikov", bandwidth, ...)

```

Arguments

<code>x</code>	bin object
<code>...</code>	further arguments to be passed
<code>y</code>	a numeric vector of y data corresponding to x.
<code>xpred</code>	a numeric vector or matrix of same dimension as x. x values to be predicted.
<code>kernel</code>	kernel type used; supported are 'epanechnikov', 'rectangular', 'triangular', 'quartic', 'triweight', 'tricube', 'cosine', 'gauss', 'logistic'.
<code>bandwidth</code>	a numeric vector or single number of same dimension as x.
<code>weight</code>	a numeric vector of length(x) for weight of each data point.
<code>kdtree</code>	boolean flag: If TRUE, kdtree is used for computation of local linear regression.
<code>approx</code>	boolean flag: If TRUE, kdtree approximation is used . Only used when kdtree = TRUE.
<code>epsilon</code>	margin of error allowed for llr approximation using kdtree. Only used when both kdtree = TRUE and approx = TRUE.
<code>N_min</code>	minimum number of points stored in the kd-tree. Only used when both kdtree = TRUE and approx = TRUE. Currently not in use.

Value

returns a S3 object of class "llr" containing

- x** sorted numeric vector or matrix of xpred
- y** estimated values corresponding to 'llr' class x

Examples

```

## Not run:
n <- 1000
x <- seq(0, 10, length.out = n)
x1 <- rnorm(n, 0, 0.2)
y <- sin(x) + x1
w <- rep(1 / n, n)
binned <- bin(x, y, bins = 400, w)
## local linear regression for exact without kdtree
llr_exact <- llr(x, y, x, bandwidth = 0.2, weight = w)
## local linear regression for kdtree exact
llr_kdexact <- llr(x, y, x, bandwidth = 0.2, weight = w, kdtree = TRUE)
## local linear regression for kdtree approximation
llr_kdapprox <- llr(x, y, x, bandwidth = 0.2, weight = w, kdtree = TRUE, approx = TRUE)

```

```
## local linear regression for data after binning.
llr_bin <- llr(binned, x, bandwidth = 0.2)

## End(Not run)
```

loocv.llr

Bandwidth selection using leave one out cross validation

Description

Bandwidth selection using leave one out cross validation

Usage

```
loocv.llr(
  x,
  y,
  weight,
  kernel = "epanechnikov",
  approx = FALSE,
  epsilon = 0.05,
  N_min = 1,
  bandwidth
)
```

Arguments

x	a numeric vector or matrix of x data
y	a numeric vector of y data corresponding to x data
weight	a numeric vector of length(x) for weight of each data point.
kernel	kernel type used; supported are 'epanechnikov', 'rectangular', 'triangular', 'quartic', 'triweight', 'tricube', 'cosine', 'gauss', 'logistic'. 'sigmoid' and 'silverman'.
approx	boolean flag: if true kdtree approximation is used.
epsilon	margin of error allowed for llr approximation using kdtree. Only used when approx = TRUE.
N_min	minimum number of points stored in the kd-tree. Only used when approx = TRUE.
bandwidth	a numeric vector or matrix of bandwidth considered for selection.

Value

returns a single or numeric vector of bandwidth that gives the smallest mean square error.

Examples

```
## Not run:
n <- 1000
x <- seq(0, 10, length.out = n)
x1 <- rnorm(n, 0, 0.2)
y <- sin(x) + x1
w <- rep(1 / n, n)
bandwidth <- seq(0.02, 0.4, by = 0.01)
binned <- bin(x, y, bins = 400, w)
## Bandwidth selection of binned data
h_bin <- gcv.llr(binned$x, binned$y, binned$weight, bandwidth = bandwidth)
## Bandwidth selection of exact local linear regression
h_exact <- gcv.llr(x, y, w, bandwidth = bandwidth)
## Bandwidth selection of approx local linear regression with kdtree
h_kdapprox <- gcv.llr(x, y, w, approx = TRUE, bandwidth = bandwidth)

## End(Not run)
```

plot.llr

Plot class 'llr' object

Description

Plot class 'llr' object

Plot 'bin' object

Usage

```
## S3 method for class 'llr'
plot(x, xorig, yorig, ...)

## S3 method for class 'bin'
plot(x, ...)
```

Arguments

x	a object of class 'bin'
xorig	initial numeric vector x
yorig	initial numeric vector y
...	for consistency

set_num_threads	<i>Set number of threads to use</i>
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Description

Set number of threads to use

Usage

```
set_num_threads(threads)
```

Arguments

threads	number of threads to use
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Value

number of threads set

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