Package ‘rdd’
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Title Regression Discontinuity Estimation
Description Provides the tools to undertake estimation in
Regression Discontinuity Designs. Both sharp and fuzzy designs are
supported. Estimation is accomplished using local linear regression.
A provided function will utilize Imbens-Kalyanaraman optimal
bandwidth calculation. A function is also included to test the
assumption of no-sorting effects.
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**rdd-package**

*Regression Discontinuity Estimation Package*

**Description**

Regression discontinuity estimation package

**Details**

*rdd* supports both sharp and fuzzy RDD utilizing the *AER* package for 2SLS regression under the fuzzy design. Local linear regressions are performed to either side of the cutpoint using the Imbens-Kalyanaraman optimal bandwidth calculation, *ikbandwidth*.

**Author(s)**

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**See Also**

*RDestimate, DCdensity, IKbandwidth, summary.RDplot.RD, kernelwts*

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**DCdensity**

*McCrary Sorting Test*

**Description**

*DCdensity* implements the McCrary (2008) sorting test.

**Usage**

```
DCdensity(runvar, cutpoint, bin = NULL, bw = NULL, verbose = FALSE, plot = TRUE, ext.out = FALSE, htest = FALSE)
```

**Arguments**

- **runvar**: numerical vector of the running variable
- **cutpoint**: the cutpoint (defaults to 0)
- **bin**: the binwidth (defaults to 2*sd(runvar)*length(runvar)^(-.5))
- **bw**: the bandwidth to use (by default uses bandwidth selection calculation from McCrary (2008))
- **verbose**: logical flag specifying whether to print diagnostic information to the terminal. (defaults to FALSE)
- **plot**: logical flag indicating whether to plot the histogram and density estimations (defaults to TRUE). The user may wrap this function in additional graphical options to modify the plot.
ext.out logical flag indicating whether to return extended output. When FALSE (the default) DCdensity will return only the p-value of the test. When TRUE, DCdensity will return the additional information documented below.

htest logical flag indicating whether to return an "htest" object compatible with base R's hypothesis test output.

Value

If ext.out is FALSE, only the p value will be returned. Additional output is enabled when ext.out is TRUE. In this case, a list will be returned with the following elements:

- theta the estimated log difference in heights at the cutpoint
- se the standard error of theta
- z the z statistic of the test
- p the p-value of the test. A p-value below the significance threshold indicates that the user can reject the null hypothesis of no sorting.
- binsize the calculated size of bins for the test
- bw the calculated bandwidth for the test
- cutpoint the cutpoint used
- data a dataframe for the binning of the histogram. Columns are cellmp (the midpoints of each cell) and cellval (the normalized height of each cell)

Author(s)

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References


Examples

#No discontinuity
x<-runif(1000,-1,1)
DCdensity(x,0)

#Discontinuity
x<-runif(1000,-1,1)
x<-x+2*(runif(1000,-1,1)>0&x<0)
DCdensity(x,0)
Description

IKbandwidth calculates the Imbens-Kalyanaraman optimal bandwidth for local linear regression in Regression discontinuity designs.

Usage

IKbandwidth(X, Y, cutpoint = NULL, verbose = FALSE, kernel = "triangular")

Arguments

X a numerical vector which is the running variable
Y a numerical vector which is the outcome variable
cutpoint the cutpoint
verbose logical flag indicating whether to print more information to the terminal. Default is FALSE.
kernel string indicating which kernel to use. Options are "triangular" (default and recommended), "rectangular", "epanechnikov", "quartic", "triweight", "tricube", "gaussian", and "cosine".

Value

The optimal bandwidth

Author(s)

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References

Description

This function will calculate the appropriate kernel weights for a vector. This is useful when, for instance, one wishes to perform local regression.

Usage

```r
kernelwts(X, center, bw, kernel = "triangular")
```

Arguments

- **X**: input x values. This variable represents the axis along which kernel weighting should be performed.
- **center**: the point from which distances should be calculated.
- **bw**: the bandwidth.
- **kernel**: a string indicating the kernel to use. Options are "triangular" (the default), "epanechnikov", "quartic", "triweight", "tricube", "gaussian", and "cosine".

Value

A vector of weights with length equal to that of the X input (one weight per element of X).

Author(s)

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Examples

```r
require(graphics)
X<-seq(-1,1,.01)
triang.wts<-kernelwts(X,0,1,kernel="triangular")
plot(X,triang.wts,type="l")

cos.wts<-kernelwts(X,0,1,kernel="cosine")
plot(X,cos.wts,type="l")
```
plot.RD  

Plot of the Regression Discontinuity

Description

Plot the relationship between the running variable and the outcome

Usage

```r
## S3 method for class 'RD'
plot(x, gran = 400, bins = 100, which = 1, range, ...)
```

Arguments

- `x`: rd object, typically the result of `RDestimate`
- `gran`: the granularity of the plot. This specifies the number of points to either side of the cutpoint for which the estimate is calculated.
- `bins`: if the dependent variable is binary, include the number of bins within which to average
- `which`: identifies which of the available plots to display. For a sharp design, the only possibility is 1, the plot of the running variable against the outcome variable. For a fuzzy design, an additional plot, 2, may also be displayed, showing the relationship between the running variable and the treatment variable. Both plots may be displayed with `which=c(1,2)``
- `range`: the range of values of the running variable for which to plot. This should be a vector of length two of the format `c(min,max)`. To plot from the minimum to the maximum value, simply enter `c("min","max")`. The default is a window 20 times wider than the first listed bandwidth from the `rd` object, truncated by the min/max values of the running variable from the data.
- ... unused

Details

It is important to note that this function will only plot the discontinuity using the bandwidth which is first in the vector of bandwidths passed to `RDestimate`

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### print.RD

**Print the Regression Discontinuity**

**Description**

Print a very basic summary of the regression discontinuity

**Usage**

```r
## S3 method for class 'RD'
print(x, digits = max(3, getOption("digits") - 3), ...)
```

**Arguments**

- `x` - rd object, typically the result of `RDestimate`
- `digits` - number of digits to print
- `...` - unused

**Author(s)**

Drew Dimmery `<drewd@nyu.edu>`

### RDestimate

**Regression Discontinuity Estimation**

**Description**

RDestimate supports both sharp and fuzzy RDD utilizing the AER package for 2SLS regression under the fuzzy design. Local linear regressions are performed to either side of the cutpoint using the Imbens-Kalyanaraman optimal bandwidth calculation, `ikbandwidth`.

**Usage**

```r
RDestimate(formula, data, subset = NULL, cutpoint = NULL, bw = NULL,
            kernel = "triangular", se.type = "HC1", cluster = NULL,
            verbose = FALSE, model = FALSE, frame = FALSE)
```

**Arguments**

- `formula` - the formula of the RDD. This is supplied in the format of `y ~ x` for a simple sharp RDD, or `y ~ x | c1 + c2` for a sharp RDD with two covariates. Fuzzy RDD may be specified as `y ~ x + z` where `x` is the running variable, and `z` is the endogenous treatment variable. Covariates are then included in the same manner as in a sharp RDD.
- `data` - an optional data frame
RDestimate

subset
an optional vector specifying a subset of observations to be used

cutpoint
the cutpoint. If omitted, it is assumed to be 0.

bw
a numeric vector specifying the bandwidths at which to estimate the RD. If omitted, the bandwidth is calculated using the Imbens-Kalyanaraman method, and then estimated with that bandwidth, half that bandwidth, and twice that bandwidth. If only a single value is passed into the function, the RD will similarly be estimated at that bandwidth, half that bandwidth, and twice that bandwidth.

kernel
a string specifying the kernel to be used in the local linear fitting. "triangular" kernel is the default and is the "correct" theoretical kernel to be used for edge estimation as in RDD (Lee and Lemieux 2010). Other options are "rectangular", "epanechnikov", "quartic", "triweight", "tricube", "gaussian" and "cosine".

se.type
this specifies the robust SE calculation method to use. Options are, as in vcovHC, "HC3", "const", "HC", "HC0", "HC1", "HC2", "HC4", "HC4m", "HC5". This option is overriden by cluster.

cluster
an optional vector specifying clusters within which the errors are assumed to be correlated. This will result in reporting cluster robust SEs. This option overrides anything specified in se.type. It is suggested that data with a discrete running variable be clustered by each unique value of the running variable (Lee and Card 2008).

verbose
will provide some additional information printed to the terminal.

model
logical. If TRUE, the model object will be returned.

frame
logical. If TRUE, the data frame used in model fitting will be returned.

Details
Covariates are problematic for inclusion in the regression discontinuity design. This package allows their inclusion, but cautions against them insomuch as is possible. When covariates are included in the specification, they are simply included as exogenous regressors. In the sharp design, this means they are simply added into the regression equation, uninteracted with treatment. Likewise for the fuzzy design, in which they are added as regressors in both stages of estimation.

Value
RDestimate returns an object of class "RD". The functions summary and plot are used to obtain and print a summary and plot of the estimated regression discontinuity. The object of class RD is a list containing the following components:

type
a string denoting either "sharp" or "fuzzy" RDD.
est
numeric vector of the estimate of the discontinuity in the outcome under a sharp design, or the Wald estimator in the fuzzy design for each corresponding bandwidth

se
numeric vector of the standard error for each corresponding bandwidth

z
numeric vector of the z statistic for each corresponding bandwidth

p
numeric vector of the p value for each corresponding bandwidth

ci
the matrix of the 95 for each corresponding bandwidth
bw numeric vector of each bandwidth used in estimation
obs vector of the number of observations within the corresponding bandwidth
call the matched call
na.action the observations removed from fitting due to missingness
model (if requested) For a sharp design, a list of the lm objects is returned. For a fuzzy design, a list of lists is returned, each with two elements: firststage, the first stage lm object, and iv, the ivreg object. A model is returned for each corresponding bandwidth.
frame (if requested) Returns the model frame used in fitting.

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References


See Also
summary.RD, plot.RD, DCDensity, kernelwts, vcovHC, ivreg.lm

Examples
x<-runif(1000,-1,1)
cov<-rnorm(1000)
y<-3+2*x+3*cov+10*(x>=0)+rnorm(1000)
RDestimate(y~x)
# Efficiency gains can be made by including covariates
RDestimate(y~x|cov)
Description

summary method for class "RD"

Usage

```r
## S3 method for class 'RD'
summary(object, digits = max(3, getOption("digits") - 3), ...)
```

Arguments

- `object` an object of class "RD", usually a result of a call to RDestimate
- `digits` number of digits to display
- `...` unused

Value

summary.RD returns an object of class "summary.RD" which has the following components:

- `coefficients` A matrix containing bandwidths, number of observations, estimates, SEs, z-values and p-values for each estimated bandwidth.
- `fstat` A global F-test of the corresponding model

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