

Package ‘ORDER2PARENT’

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Type Package

Title Estimate parent distributions with data of several order statistics

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Description This package uses B-spline based nonparametric smooth estimators to estimate parent distributions given observations on multiple order statistics.

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support	a vector specifying the support of the parent distribution. If unknown, it can be omitted, and the interval of data will be used as the support.
weight.type	the type of weight matrix used in implementing the GMM estimator. The default is 1, i.e. the weight matrix based on sample size.

Details

The `dat` must be a list consisting of vectors of observations on order statistics. For example, there are three order statistics, and the observations on them are contained in three vectors, `dat.order1`, `dat.order2`, and `dat.order3`. Then a typical `dat` is `list(dat.order1, dat.order2, dat.order3)`. `\orderinfo` must be a matrix with two columns and `J` rows where `J` is the number of observed order statistics. For `j`-th row of `orderinfo`, the first column is the rank, and the second column is the size of the `j`-th order statistic. `\support` is vector whose first element is the lower bound of the support, and the second element is the upper bound. If you want to use the second type of weight matrix, which is based on mean square error of the first stage estimates, set `weight.type=2`.

Value

`bgmm` gives a list consisting of two element: `betahat` and `n.knots`. These two elements will be used in `parentest` for estimation of parent cdf.

See Also

[blr](#), [parentest](#)

Examples

```
n.order<-c(20, 20, 60) # number of observations for each order statistic below.
m<-5 # the size of random samples is 5.
# The three order statistics are 1:5 (the minimum), 3:5 (the sample median),
# and 5:5 (the maximum)
rank.x<-c(1, 3, 5)
data.example<-list()
for(i in 1:3){
  sorted.sample<-t(apply(matrix(rnorm(m*n.order[i]),nr=n.order[i],nc=m), 1, sort))
  data.example[[i]]<-sorted.sample[,rank.x[i]]
}
order.example<-rbind(c(1, 5), c(3, 5), c(5, 5), deparse.level=0)
gmm.example<-bgmm(data.example, order.example)
```

blr

Parent Distribution Estimation with B-Spline Linear Regression Estimator

Description

Given observations on several order statistics, this function use the B-Spline linear regression estimator (Chou and Tao, 2010) to estimate the corresponding parent distribution of these order statistics nonparametrically.

Usage

```
blr(dat, orderinfo, degree = 3, support = NULL, constraint = FALSE)
```

Arguments

<code>dat</code>	a list consisting of the vectors of observations on various order statistics.
<code>orderinfo</code>	a matrix about the ranks and the sizes of various order statistics.
<code>degree</code>	the degree of B-spline used for estimation. The default is 3, i.e. cubic B-spline.
<code>support</code>	a vector specifying the support of the parent distribution. If unknown, it can be omitted, and the interval of data will be used as the support.
<code>constraint</code>	whether add monotone constraints in estimation. The default is no (FALSE).

Details

The details about specifying the above arguments can be found in `bgmm`.

Value

`blr` gives a list consisting of two element: `betahat` and `n.knots`. These two elements will be used in `parentest` for estimation of parent cdf.

See Also

[bgmm](#), [parentest](#)

Examples

```
n.order<-c(20, 20, 60) # number of observations for each order statistic below.
m<-5 # the size of random samples is 5.
# The three order statistics are 1:5 (the minimum), 3:5 (the sample median),
# and 5:5 (the maximum)
rank.x<-c(1, 3, 5)
data.example<-list()
for(i in 1:3){
  sorted.sample<-t(apply(matrix(rnorm(m*n.order[i]),nr=n.order[i],nc=m), 1, sort))
  data.example[[i]]<-sorted.sample[,rank.x[i]]
}
order.example<-rbind(c(1, 5), c(3, 5), c(5, 5), deparse.level=0)
blr.example<-blr(data.example, order.example)
```

parentcdf

Transform CDF of Order Statistics to Its Parent CDF

Description

Using the well relationship between the cdf of order statistic and the cdf of parent distribution, this function can transform the the cdf of order statistic to the corresponding parent cdf.

Usage

```
parentcdf(F.order, k, m)
```

Arguments

F.order	CDF of order statistic, either a scalar or a vector.
k	the rank of the order statistic
m	the size of the order statistic

Details

Use the well known relationship (David and Nagaraja, 2003) between the cdf of order statistic and the parent distribution, parentcdf will return the corresponding parent distribution.

Value

The corresponding parent distribution.

References

David, H.A., and Nagaraja, H.N. (2003): Order Statistics, Third Edition. New York: Wiley.

parentest

Generate Smooth Estimator of Parent Distribution

Description

Using the output from b1r or bgmm, this function gives rise to estimates of parent cdf for any given value.

Usage

```
parentest(x0, beta.hat, n.knots, degree = 3, support = NULL)
```

Arguments

<code>x0</code>	the value whose parent cdf's are wanted. It can either be a scalar or a vector.
<code>beta.hat</code>	the estimate of control variables.
<code>n.knots</code>	the number of inner knots.
<code>degree</code>	the degree of B-spline. The default is 3, i.e. a cubic B-spline.
<code>support</code>	a vector specifying the support of the parent distribution. If unknown, it can be omitted, and the interval of data will be used as the support.

Details

Together with `blr` and/or `bgmm`, this function can be estimate the parent cdf of any given value.

Value

The estimates of parent cdf's of `x0`. NOTE that the degree used in `parentest` should be consistent with the degree used in estimation of control variables.

See Also

[blr](#), [bgmm](#)

Examples

```
n.order<-c(20, 20, 60) # number of observations for each order statistic below.
m<-5 # the size of random samples is 5.
# The three order statistics are 1:5 (the minimum), 3:5 (the sample median),
# and 5:5 (the maximum)
rank.x<-c(1, 3, 5)
data.example<-list()
for(i in 1:3){
  sorted.sample<-t(apply(matrix(rnorm(m*n.order[i]),nr=n.order[i],nc=m), 1, sort))
  data.example[[i]]<-sorted.sample[,rank.x[i]]
}
order.example<-rbind(c(1, 5), c(3, 5), c(5, 5), deparse.level=0)
blr.example<-blr(data.example, order.example)
# Based on 'blr.example', we can estimate the parent cdf of given values, like data.example[[3]]
parenthat<-parentest(data.example[[3]], blr.example$betahat, blr.example$n.knots)
```

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