The leaps Package

March 24, 2009

Title regression subset selection

Version 2.8

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Description Regression subset selection including exhaustive search

Depends

Suggests biglm

License GPL version 2 or later

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Repository CRAN

Date/Publication

R topics documented:

`leaps`   *all-subsets regressiom*

Description

leaps() performs an exhaustive search for the best subsets of the variables in x for predicting y in linear regression, using an efficient branch-and-bound algorithm. It is a compatibility wrapper for [regsubsets](#) does the same thing better.

Usage

```
leaps(x=, y=, wt=rep(1, NROW(x)), int=TRUE, method=c("Cp", "adjr2", "r2"), nbest=1(
```

Arguments

x	A matrix of predictors
y	A response vector
wt	Optional weight vector
int	Add an intercept to the model
method	Calculate Cp, adjusted R-squared or R-squared
nbest	Number of subsets of each size to report
names	vector of names for columns of x
df	Total degrees of freedom to use instead of `nrow(x)` in calculating Cp and adjusted R-squared
strictly.compatible	
	Implement misfeatures of leaps() in S

Value

A list with components

which	logical matrix. Each row can be used to select the columns of x in the respective model
size	Number of variables, including intercept if any, in the model
cp	or `adjr2` or `r2` is the value of the chosen model selection statistic for each model
label	vector of names for the columns of x

Note

With `strictly.compatible=T` the function will stop with an error if x is not of full rank or if it has more than 31 columns. It will ignore the column names of x even if `names==NULL` and will replace them with "0" to "9", "A" to "Z".

References

Alan Miller "Subset Selection in Regression" Chapman & Hall

See Also

regsubsets, regsubsets.formula, regsubsets.default

Examples

```
x<-matrix(rnorm(100),ncol=4)
y<-rnorm(25)
leaps(x,y)
```

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leaps.setup	*Internal functions for leaps(), subsets()*

Description

These functions are used internally by regsubsets and leaps. They are wrappers for Fortran routines that construct and manipulate a QR decomposition.

Usage

```
leaps.setup(x,y,wt=rep(1,length(y)),force.in=NULL,force.out=NULL,intercept=TRUE,nvm
leaps.seqrep(leaps.obj)
leaps.exhaustive(leaps.obj,really.big=FALSE)
leaps.backward(leaps.obj)
leaps.forward(leaps.obj)
```

Arguments

x	A matrix of predictors
y	A response vector
wt	Optional weight vector
intercept	Add an intercept to the model
force.in	vector indicating variable that must be in the model
force.out	vector indicating variable that must not be in the model
nbest	Number of subsets of each size to report
nvmax	largest subset size to examine
warn.dep	warn if x is not of full rank
leaps.obj	An object of class leaps as produced by leaps.setup
really.big	required before R gets sent off on a long uninterruptible computation

See Also

regsubsets, leaps

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plot.regsubsets   *Graphical table of best subsets*

Description

Plots a table of models showing which variables are in each model. The models are ordered by the specified model selection statistic. This plot is particularly useful when there are more than ten or so models and the simple table produced by summary.regsubsets is too big to read.

Usage

```
## S3 method for class 'regsubsets':
plot(x, labels=obj$xnames, main=NULL, scale=c("bic", "Cp", "adjr2", "r2"), col=gray
```

Arguments

x	regsubsets object
labels	variable names
main	title for plot
scale	which summary statistic to use for ordering plots
col	Colors: the last color should be close to but distinct from white
...	other arguments

Value

None

Author(s)

Thomas Lumley, based on a concept by Merlise Clyde

See Also

regsubsets,summary.regsubsets

Examples

```
data(swiss)
a<-regsubsets(Fertility~.,nbest=3,data=swiss)
par(mfrow=c(1,2))
plot(a)
plot(a,scale="r2")
```

| regsubsets | *functions for model selection* |

Description

Model selection by exhaustive search, forward or backward stepwise, or sequential replacement

Usage

```
regsubsets(x=, ...)

## S3 method for class 'formula':
regsubsets(x=, data=, weights=NULL, nbest=1, nvmax=8, force.in=NULL, force.out=NULL

## Default S3 method:
regsubsets(x=, y=, weights=rep(1, length(y)), nbest=1, nvmax=8,
force.in=NULL, force.out=NULL, intercept=TRUE, method=c("exhaustive",
"backward", "forward", "seqrep"), really.big=FALSE,...)

## S3 method for class 'biglm':
regsubsets(x,nbest=1,nvmax=8,force.in=NULL,
method=c("exhaustive","backward", "forward", "seqrep"), really.big=FALSE,...)

## S3 method for class 'regsubsets':
summary(object,all.best=TRUE,matrix=TRUE,matrix.logical=FALSE,df=NULL,...)
```

Arguments

x	design matrix or model formula for full model, or `biglm` object
data	Optional data frame
y	response vector
weights	weight vector
nbest	number of subsets of each size to record
nvmax	maximum size of subsets to examine
force.in	index to columns of design matrix that should be in all models
force.out	index to columns of design matrix that should be in no models
intercept	Add an intercept?
method	Use exhaustive search, forward selection, backward selection or sequential replacement to search.
really.big	Must be TRUE to perform exhaustive search on more than 50 variables.
object	regsubsets object
all.best	Show all the best subsets or just one of each size

`matrix`	Show a matrix of the variables in each model or just summary statistics
`matrix.logical`	
	With `matrix=TRUE`, the matrix is logical TRUE/FALSE or string `"*"/" "`
`df`	Specify a number of degrees of freedom for the summary statistics. The default is `n-1`
`...`	Other arguments for future methods

Details

Since this function returns separate best models of all sizes up to `nvmax` and since different model selection criteria such as AIC, BIC, ... differ only in how models of different sizes are compared, the results do not depend on the choice of cost-complexity tradeoff.

When `x` is a `biglm` object it is assumed to be the full model, so `force.out` is not relevant. If there is an intercept it is forced in by default; specify a `force.in` as a logical vector with `FALSE` as the first element to allow the intercept to be dropped.

Value

An object of class "regsubsets" containing no user-serviceable parts. It is designed to be processed by `summary.regsubsets`.

See Also

`leaps`

Examples

```
data(swiss)
a<-regsubsets(as.matrix(swiss[,-1]),swiss[,1])
summary(a)
b<-regsubsets(Fertility~.,data=swiss)
summary(a)
```

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