

Package ‘REEMtree’

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Type Package

Title Regression Trees with Random Effects for Longitudinal (Panel)
Data

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Description A data mining approach for longitudinal and clustered data,
which combines the structure of mixed effects model with tree-based
estimation methods. See Sela, R.J. and Simonoff, J.S. (2012) RE-EM
trees: a data mining approach for longitudinal and clustered data
<doi:10.1007/s10994-011-5258-3>.

License GPL

URL <http://pages.stern.nyu.edu/~jsimonof/REEMtree/>

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REEMtree-package	<i>Regression Trees with Random Effects for Longitudinal (Panel) Data</i>
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Description

This package estimates regression trees with random effects as a way to use data mining techniques to describe longitudinal or panel data.

Details

Package: REEMtree
Type: Package
Version: 1.0
Date: 2009-05-07
License: GPL

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
print(REEMresult)
```

AutoCorrelationLRtest *Test for autocorrelation in the residuals of a RE-EM tree*

Description

This function tests for autocorrelation in the residuals of a RE-EM tree using a likelihood ratio test. The test keeps the tree structure of the RE-EM tree object fixed and uses a standard likelihood ratio test on the linear random effects model.

Usage

```
AutoCorrelationLRtest(object, newdata=NULL, correlation=corAR1())
```

Arguments

object	A RE-EM tree
newdata	Dataset on which the test is to be performed; if none is given, the original dataset is used
correlation	Type of correlation to be tested for in the residuals. The correlation can be any of type corClasses .

Details

In general, newdata is likely to be the data used to estimate object. The RE-EM tree can be estimated with or without allowing for autocorrelation. Because the estimated tree may differ depending on whether autocorrelation is allowed in the RE-EM tree estimation process, but we recommend testing based on the tree estimated with autocorrelation allowed and the tree estimated without autocorrelation allowed.

Value

correlation	Type of correlation used in testing
loglik0	Likelihood of the random effects model if there is no autocorrelation
loglikAR	Likelihood of the random effects model if autocorrelation (of type AR(1)) is estimated
pvalue	P-value of the likelihood ratio test

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also[corClasses](#)**Examples**

```

data(simpleREEMdata)

# Estimation without autocorrelation
simpleEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
# Estimation with autocorrelation
simpleEMresult2<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID, correlation=corAR1())

# Autocorrelation test based on the first tree
AutoCorrelationLRtest(simpleEMresult, simpleREEMdata)
# Autocorrelation test based on the second tree
AutoCorrelationLRtest(simpleEMresult2, simpleREEMdata)
# Autocorrelation test with an alternative correlation structure
AutoCorrelationLRtest(simpleEMresult, simpleREEMdata, correlation=corCAR1())

```

fitted.REEMtree

*Extract the fitted values from a RE-EM tree***Description**

This function extracts the fitted values from the LME object underlying the RE-EM tree. The fitted values are the fixed effects (from the tree) plus the estimated contributions of the random effects to the fitted values at grouping levels less or equal to the level given.

Usage

```

## S3 method for class 'REEMtree'
fitted(object, level, asList, ...)

```

Arguments

<code>object</code>	an object of class REEMtree
<code>level</code>	the level of random effects used in creating fitted values. Level 0 is fixed effects; levels increase with the grouping of random effects. Default is the highest level.
<code>asList</code>	an optional logical value. If TRUE and a single value is given in level, the returned object is a list with the fitted values split by groups; otherwise the returned value is either a vector or a data frame, according to the length of level. Defaults to FALSE.
<code>...</code>	some methods for this generic require additional arguments; none are used here.

Value

If the level is a single value, the result is a vector or list (depending on `asList`) with the fitted values. Otherwise, the result is a data frame with columns given by the fitted values at different levels.

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also

[fitted](#), [REEMtree.object](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
fitted(REEMresult)
```

is.REEMtree	<i>Is a RE-EM tree object</i>
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Description

This function tests whether an object is of the REEMtree class.

Usage

```
is.REEMtree(object)
```

Arguments

object any R object

Value

TRUE if the object is of the REEMtree type

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
is.REEMtree(REEMresult)
```

logLik	<i>Log-likelihood of a RE-EM tree</i>
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Description

This returns the log-likelihood of the effects model of a RE-EM tree. This is the log-likelihood of the random effects model estimated in the RE-EM tree. (The regression tree is not associated with a log-likelihood.)

Usage

```
## S3 method for class 'REEMtree'
logLik(object,...)
```

Arguments

object	an object of class REEMtree
...	further arguments passed to or from other methods

Value

the log-likelihood of the fitted effects model associated with x

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also

[REEMtree.object](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
logLik(REEMresult)
```

plot

*Plot the RE-EM tree***Description**

Plots the regression tree associated with a RE-EM tree.

Usage

```
## S3 method for class 'REEMtree'
plot(x, text = TRUE, ...)
```

Arguments

x	a fitted object of class REEMtree
text	if TRUE, the text of the tree will be plotted on the tree automatically.
...	further arguments passed to or from other methods

Value

the coordinates of the nodes are returned as a list, with components x and y.

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also

[REEMtree](#), [plot.rpart](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
plot(REEMresult)
```

predict

Predictions from a regression tree with individual-specific effects

Description

Returns a vector of predictions from a fitted RE-EM Tree. Predictions are based on the node of the tree in which the new observation would fall and (optionally) an estimated random effect for the observation.

Usage

```
## S3 method for class 'REEMtree'
predict(object, newdata, id = NULL,
        EstimateRandomEffects = TRUE, ...)
```

Arguments

object	a fitted REEMtree
newdata	an data frame to be used for obtaining the predictions. All variables used in the fixed and random effects models, including the group identifier, must be present in the data frame. New values of the group identifier are allowed. Unlike in <code>predict.lme</code> and <code>predict.rpart</code> , the data frame is required
id	a string containing the name of the variable that is used to identify the groups. This is required if <code>EstimateRandomEffects=TRUE</code> and <code>newdata</code> does not match the data used to estimate the random effects model that created object.
EstimateRandomEffects	if TRUE, the fitted effects will be included in the estimates and effects for new groups will be estimated wherever the target variable is not missing. If FALSE or if the random effect cannot be estimated, random effects are set to 0, so that only the fixed effects based on the regression tree are used.
...	additional arguments that will be passed through to <code>rpart</code>

Details

If `EstimateRandomEffects=TRUE` and a group was not used in the original estimation, its random effect must be estimated. If there are no non-missing values of the target variable for this group, then the new effect is set to 0.

If there are non-missing values of the target variable, then the random effect is estimated based on the estimated variance of the errors and variance of the random effects in the fitted model. See Equation 3.2 of Laird and Ware (1982) for the precise relationship.

Important note: In this implementation, estimation of group effects for new groups can be used only with group-specific intercepts are estimated with only one grouping variable.

Value

a vector containing the predicted values

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal Data”, *Machine Learning*, 2011; Laird, N. M., and J. H. Ware (1982), “Random-effects models for longitudinal data”, *Biometrics* 38: 963-974

See Also

[predict.nlme](#), [predict.rpart](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
predict(REEMresult, simpleREEMdata, EstimateRandomEffects=FALSE)
predict(REEMresult, simpleREEMdata, id=simpleREEMdata$ID, EstimateRandomEffects=TRUE)

# Estimation based on a subset that excludes the last two time series,
# with predictions for all observations
sub <- rep(c(rep(TRUE, 10), rep(FALSE, 2)), 50)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID,
subset=sub)
pred1 <- predict(REEMresult, simpleREEMdata, EstimateRandomEffects=FALSE)
pred2 <- predict(REEMresult, simpleREEMdata, id=simpleREEMdata$ID, EstimateRandomEffects=TRUE)

# Estimation based on a subset that excludes the last five individuals,
# with predictions for all observations
sub <- c(rep(TRUE, 540), rep(FALSE, 60))
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID,
subset=sub)
pred3 <- predict(REEMresult, simpleREEMdata, EstimateRandomEffects=FALSE)
pred4 <- predict(REEMresult, simpleREEMdata, id=simpleREEMdata$ID, EstimateRandomEffects=TRUE)
```

print

Print a RE-EM Tree object

Description

This function prints a description of a fitted RE-EM tree object.

Usage

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

Arguments

`x` fitted model of class REEMtree
`...` further arguments passed to or from other methods

Details

This function is a method for the generic function `print` for class REEMtree. It can be invoked by calling `print` for an object of class REEMtree, or by calling `print.REEMtree` directly for an object of the corresponding type.

Side Effects

Prints representations of the regression tree and the random effects model that comprise a RE-EM tree.

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also

[print.rpart](#), [REEMtree.object](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
print(REEMresult)
```

ranef.REEMtree

Extract the estimated random effects from a RE-EM tree

Description

This function extracts the estimated random effects from a fitted RE-EM tree.

Usage

```
## S3 method for class 'REEMtree'
ranef(object,...)
```

Arguments

object an object of class REEMtree
 ... further arguments passed to or from other methods

Value

a vector containing the estimated random effects

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also

[random.effects](#), [REEMtree.object](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
ranef(REEMresult)
```

REEMtree

Create a RE-EM tree

Description

Fit a RE-EM tree to data. This estimates a regression tree combined with a linear random effects model.

Usage

```
REEMtree(formula, data, random, subset=NULL,
          initialRandomEffects=rep(0,TotalObs),
          ErrorTolerance=0.001, MaxIterations=1000,
          verbose=FALSE, tree.control=rpart.control(cp=0.001),
          cv=TRUE, no.SE =1,
          lme.control=lmeControl(returnObject=TRUE),
          method="REML", correlation=NULL)
```

Arguments

<code>formula</code>	a formula, as in the <code>lm</code> or <code>rpart</code> function
<code>data</code>	a data frame in which to interpret the variables named in the formula (unlike in <code>lm</code> or <code>rpart</code> , this is not optional)
<code>random</code>	a description of the random effects, as a formula of the form $\sim 1 g$, where g is the grouping variable
<code>subset</code>	an optional logical vector indicating the subset of the rows of data that should be used in the fit. All observations are included by default.
<code>initialRandomEffects</code>	an optional vector giving initial values for the random effects to use in estimation
<code>ErrorTolerance</code>	when the difference in the likelihoods of the linear models of two consecutive iterations is less than this value, the RE-EM tree has converged
<code>MaxIterations</code>	maximum number of iterations allowed in estimation
<code>verbose</code>	if TRUE, the current estimate of the RE-EM tree will be printed after each iteration
<code>tree.control</code>	a list of control values for the estimation algorithm to replace the default values used to control the <code>rpart</code> algorithm. Defaults to be <code>rpart.control(cp=0.001)</code> .
<code>cv</code>	if TRUE then cross-validation will be used for estimating the tree at each iteration. Default is TRUE.
<code>no.SE</code>	number of standard errors used in pruning (0 if unused)
<code>lme.control</code>	a list of control values for the estimation algorithm to replace the default values returned by the function <code>lmeControl</code> . Defaults to an empty list.
<code>method</code>	whether the linear model should be estimated with ML or REML
<code>correlation</code>	an optional <code>corStruct</code> object describing the within-group correlation structure; the available classes are given in corClasses

Value

an object of class `REEMtree`

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also

[rpart](#), [nlme](#), [REEMtree.object](#), [corClasses](#)

Examples

```

data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)

# Estimation allowing for autocorrelation
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID,
correlation=corAR1())

# Random parameters model for the random effects
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1+X|ID)

# Estimation with a subset
sub <- rep(c(rep(TRUE, 10), rep(FALSE, 2)), 50)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID,
subset=sub)

# Dataset from the R library "AER"
data("Grunfeld", package = "AER")
REEMtree(invest ~ value + capital, data=Grunfeld, random=~1|firm)
REEMtree(invest ~ value + capital, data=Grunfeld, random=~1|firm, correlation=corAR1())
REEMtree(invest ~ value + capital, data=Grunfeld, random=~1+year|firm)
REEMtree(invest ~ value + capital, data=Grunfeld, random=~1|firm/year)

```

REEMtree.object

*Random Effects/Expectation Maximization (RE-EM) Tree Object***Description**

Object representing a fitted REEMtree.

Value

Tree	Fitted rpart tree associated with the fitted RE-EM tree
EffectModel	fitted lme object associated with the fitted RE-EM tree
RandomEffects	vector of estimated random effects
BetweenMatrix	estimated variance of the random effects
ErrorVariance	estimated variance of the errors
data	the data frame used to estimate the RE-EM tree
logLik	log likelihood of the linear model for the random effects
IterationsUsed	number of iterations required to fit the REEMtree
Formula	formula used in fitting the REEMtree
Random	description of the random effects used in fitting the REEMtree
Groups	the vector of group identifiers used in estimation
Subset	the logical vector indicating the subset of the rows of data used in the fit

ErrorTolerance the error tolerance used in estimation
 correlation the correlation structure used in fitting the linear model
 residuals estimated residuals
 method method (ML or REML) used in estimating the linear random effects model
 lme.control parameters used to control fitting the linear random effects model
 tree.control parameters used to control fitting the regression tree

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., “RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data”, *Machine Learning* (2011).

See Also

[rpart](#), [nlme](#), [REEMtree](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
```

residuals.REEMtree	<i>Extract the residuals from a RE-EM tree</i>
--------------------	--

Description

This function extracts the residuals from the LME object underlying the RE-EM tree. The residuals depend on the fixed effects (from the tree) plus the estimated contributions of the random effects to the fitted values at grouping levels less or equal to the level given.

Usage

```
## S3 method for class 'REEMtree'
residuals(object, level, type, asList, ...)
```

Arguments

object an object of class REEMtree
 level the level of random effects used in creating residuals. Level 0 is fixed effects only; levels increase with the grouping of random effects. Default is the highest level.

type	optional character string specifying the type of residuals to be used. If "response", the "raw" residuals (observed - fitted) are used. If "pearson", the standardized residuals (raw residuals divided by the corresponding standard errors) are used. If "normalized", the normalized residuals (standardized residuals pre-multiplied by the inverse square-root factor of the estimated error correlation matrix) are used. Only the first character needs to be provided. Defaults to "pearson".
asList	an optional logical value. If TRUE and a single value is given in level, the returned object is a list with the residuals split by groups; otherwise the returned value is either a vector or a data frame, according to the length of level. Defaults to FALSE.
...	some methods for this generic require additional arguments; none are used here.

Value

If the level is a single value, the result is a vector or list (depending on `asList`) with the residuals. Otherwise, the result is a data frame with columns given by the residuals at different levels.

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., "RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data", *Machine Learning* (2011).

See Also

[residuals](#), [REEMtree.object](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
residuals(REEMresult)
```

simpleREEMdata

Sample Data for RE-EM trees

Description

This data set is consists of a panel of 50 individuals with 12 observations per individual. The data is based on a regression tree with an initial split based on a dummy variable (D) and a second split based on time in the branch where D=1. The observations include both randomly generated individual-specific effects and observation-specific errors.

Format

The data has 600 rows and 5 columns. The columns are:

- Y - the target variable
- t - a numeric predictor ("time")
- D - a categorical predictor with two levels, 0 and 1
- ID - the identifier for each individual
- X - another covariate (which is intentionally unrelated to the target variable)

References

Sela, Rebecca J., and Simonoff, Jeffrey S., "RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data", *Machine Learning* (2011).

tree

Extract the regression tree associated with a RE-EM tree

Description

Returns the fitted rpart object associated with a REEMtree object.

Usage

```
tree(object, ...)
```

Arguments

object	an object of class REEMtree
...	further arguments passed to or from other methods

Value

the fitted regression tree associated with the REEMtree object

Author(s)

Rebecca Sela <rsela@stern.nyu.edu>

References

Sela, Rebecca J., and Simonoff, Jeffrey S., "RE-EM Trees: A Data Mining Approach for Longitudinal and Clustered Data", *Machine Learning* (2011).

See Also

[rpart.object](#), [REEMtree.object](#)

Examples

```
data(simpleREEMdata)
REEMresult<-REEMtree(Y~D+t+X, data=simpleREEMdata, random=~1|ID)
tree.REEMtree(REEMresult)
tree(REEMresult)
```

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