

Package ‘ItemResponseTrees’

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Title IR-Tree Modeling in 'mirt', 'Mplus', or 'TAM'

Version 0.2.5

Description Item response tree (IR-tree) models are a class of item response theory (IRT) models that assume that the responses to polytomous items can best be explained by multiple psychological processes; see Böckenholt (2012) <doi:10.1037/a0028111> for details. The package 'ItemResponseTrees' allows to fit such IR-tree models in 'mirt', 'Mplus', or 'TAM'. The package automates some of the hassle of IR-tree modeling by means of a consistent syntax. This allows new users to quickly adopt this model class, and this allows experienced users to fit many complex models effortlessly.

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URL <https://github.com/hplieninger/ItemResponseTrees>

BugReports <https://github.com/hplieninger/ItemResponseTrees/issues>

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augment.irtree_fit *Augment data with information from an irtree_fit object*

Description

Augment accepts a model object and a dataset and adds information about each observation in the dataset, namely, predicted values in the .fitted column. New columns always begin with a . prefix to avoid overwriting columns in the original dataset.

Usage

```
## S3 method for class 'irtree_fit'
augment(x = NULL, data = NULL, se_fit = TRUE, method = "EAP", ...)
```

Arguments

x	object of class irtree_fit as returned from fit().
data	Optional data frame that is returned together with the predicted values. Argument is not needed since the data are contained in the fitted object.
se_fit	Logical indicating whether standard errors for the fitted values should be returned as well.
method	This is passed to mirt::fscores() or TAM::IRT.factor.scores() (as argument type) if applicable.

... Additional arguments passed to `mirt::fscores()` or `TAM::IRT.factor.scores()` if applicable.

Details

Note that argument `method` is used only for engines `mirt` and `TAM`.

Value

Returns a [tibble](#) with one row for each observation and one (two) additional columns for each latent variable if `se_fit = FALSE` (if `se_fit = TRUE`). The names of the new columns start with `.fit` (and `.se.fit`).

See Also

[generics::augment\(\)](#)

Examples

```
data("jackson")
df1 <- jackson[1:234, paste0("C", 1:5)]
irtree_create_template(df1)
m1 <- "
IRT:
t BY C1@1, C2@1, C3@1, C4@1, C5@1;
Class:
GRM"
fit1 <- fit(irtree_model(m1), data = df1)

tidy(fit1, par_type = "difficulty")

glance(fit1)

augment(fit1)
```

control_mirt

Control aspects of fitting a model in mirt

Description

This function should be used to generate the `control` argument of the `fit()` function.

Usage

```
control_mirt(
  SE = TRUE,
  method = "EM",
  quadpts = NULL,
  control = list(),
```

```

    technical = list(),
    ...
  )

```

Arguments

SE, method, quadpts, ...

These arguments are passed to and documented in `mirt::mirt()`. They can be used to tweak the estimation algorithm.

control List of arguments passed to argument control of `mirt::mirt()`.

technical List of arguments passed to argument technical of `mirt::mirt()`.

Value

A list with one element for every argument of `control_mirt()`.

Examples

```

control_mirt(SE = FALSE,
             method = "QMCEM",
             quadpts = 4455,
             technical = list(NCYCLES = 567),
             TOL = .001)
control_mirt(method = "MHRM",
             draws = 5544)

```

control_mplus

Control aspects of fitting a model in Mplus

Description

This function should be used to generate the control argument of the `fit()` function.

Usage

```

control_mplus(
  file = tempfile("irtree_"),
  overwrite = FALSE,
  cleanup = run,
  run = TRUE,
  estimator = "MLR",
  quadpts = 15,
  save_fscores = TRUE,
  analysis_list = list(COVERAGE = "0"),
  Mplus_command = "Mplus",
  warnings2messages = FALSE
)

```

Arguments

file	String naming the file (or path) of the Mplus files and the data file. Do not provide file endings, those will be automatically appended.
overwrite	Logical value indicating whether data and input (if present) files should be overwritten.
cleanup	Logical, whether the Mplus files should be removed on exit.
run	Logical, whether to indeed run Mplus.
estimator	String, passed to argument 'ESTIMATOR' in Mplus.
quadpts	This is passed to argument 'INTEGRATION' of Mplus. Thus, it may be an integer specifying the number of integration points for the Mplus default of rectangular numerical integration (e.g., quadpts = 15). Or it may be a string, which gives more fine grained control (e.g., quadpts = "MONTECARLO(2000)").
save_fscores	Logical, whether to save FSCORES or not.
analysis_list	Named list of strings passed to Mplus' argument ANALYSIS. See examples below.
Mplus_command	optional. N.B.: No need to pass this parameter for most users (has intelligent defaults). Allows the user to specify the name/path of the Mplus executable to be used for running models. This covers situations where Mplus is not in the system's path, or where one wants to test different versions of the Mplus program.
warnings2messages	Logical, whether Mplus errors and warnings should be signaled as warnings (the default) or messages.

Value

A list with one element for every argument of control_mplus().

Examples

```
control_mplus(file = tempfile("irtree_", tmpdir = "."),
             quadpts = "GAUSS(10)",
             analysis_list = list(COVERAGE = "0",
                                MITERATIONS = "500",
                                MCONVERGENCE = ".001"))
```

control_tam

Control aspects of fitting a model in TAM

Description

This function should be used to generate the control argument of the `fit()` function.

Usage

```
control_tam(
  set_min_to_0 = FALSE,
  control = list(snodes = 0, maxiter = 1000, increment.factor = 1, fac.oldxsi = 0),
  ...
)
```

Arguments

`set_min_to_0` Logical. `TAM::tam.mml()` expects the data to be scored 0, ..., K. If `set_min_to_0 = TRUE`, the minimum of the data is subtracted from each response, which will likely both satisfy TAM and do no harm to the data.

`control` List of arguments passed to argument `control` of `TAM::tam.mml()`. See examples below.

... Other arguments passed to `TAM::tam.mml()`.

Value

A list with one element for every argument of `control_tam()`.

Examples

```
control_tam(set_min_to_0 = TRUE,
  control = list(snodes = 0,
    maxiter = 1000,
    increment.factor = 1,
    fac.oldxsi = 0),
  constraint = "items")
```

fit.irtree_model *Fit an ItemResponseTrees model*

Description

This function takes a data frame and an object of class `irtree_model` and runs the model in either `mirt`, `Mplus`, or `TAM`.

Usage

```
## S3 method for class 'irtree_model'
fit(
  object = NULL,
  data = NULL,
  engine = c("mirt", "mplus", "tam"),
  ...,
  link = c("logit", "probit"),
  verbose = interactive(),
```

```

    control = NULL,
    improper_okay = FALSE
  )

```

Arguments

object	Object of class <code>irtree_model</code> . See irtree_model for more information.
data	Data frame containing containing one row per respondent and one column per variable. The variable names must correspond to those used in object.
engine	String specifying whether to use <code>mirt</code> , <code>Mplus</code> , or <code>TAM</code> for estimation.
...	Not currently used. Use <code>control</code> instead.
link	String specifying the link function. May be either <code>logit</code> , or (in case of <code>Mplus</code>), <code>probit</code> .
verbose	Logical indicating whether output should be printed to the console.
control	List. The allowed elements of this list depend on the engine. Use control_mirt() , control_mplus() , or control_tam() for convenience. Note that the <code>fit()</code> function does not use <code>...</code> , but that you can use the <code>control_*</code> () functions to pass additional arguments.
improper_okay	Logical indicating whether the model should also be fit if it is not a proper IR-tree model. Set this only to <code>TRUE</code> if you really know what you are doing.

Value

Returns a list of class `irtree_fit`. The first list element is the return value of either [mirt::mirt\(\)](#), [MplusAutomation::readModels\(\)](#), or [TAM::tam.mml\(\)](#). Further information is provided in the element spec.

Methods

The methods `coef()`, `summary()`, and `print()` are implemented for objects of class `irtree_fit`, and those wrap the respective functions of [mirt](#), [MplusAutomation](#), or [TAM](#). However, [glance\(\)](#), [tidy\(\)](#), and [augment\(\)](#) may be more helpful.

Examples

```

# Running these examples may take a while

data("jackson")
df1 <- jackson[1:456, paste0("C", 1:5)]
df2 <- jackson[1:456, c(paste0("C", 1:5), paste0("E", 1:5))]

irtree_create_template(df1)

# Graded Response Model -----

m1 <- "
IRT:

```

```

t BY C1@1, C2@1, C3@1, C4@1, C5@1;

Class:
GRM
"

model1 <- irtree_model(m1)

fit1 <- fit(model1, data = df1)

glance(fit1)
tidy(fit1, par_type = "difficulty")
augment(fit1)

# IR-Tree Models -----

##### IR-tree model for 1 target trait #####

m2 <- "
Equations:
1 = (1-m)*(1-t)*e
2 = (1-m)*(1-t)*(1-e)
3 = m
4 = (1-m)*t*(1-e)
5 = (1-m)*t*e

IRT:
t BY C1@1, C2@1, C3@1, C4@1, C5@1;
e BY C1@1, C2@1, C3@1, C4@1, C5@1;
m BY C1@1, C2@1, C3@1, C4@1, C5@1;

Class:
Tree
"

model2 <- irtree_model(m2)

# See ?mirt::mirt for details on method argument
fit2 <- fit(model2, data = df1, control = control_mirt(method = "MHRM"))

##### IR-tree model for 2 target traits #####

m3 <- "
Equations:
1 = (1-m)*(1-t)*e
2 = (1-m)*(1-t)*(1-e)
3 = m
4 = (1-m)*t*(1-e)
5 = (1-m)*t*e

IRT:
t1 BY C1@1, C2@1, C3@1, C4@1, C5@1;
t2 BY E1@1, E2@1, E3@1, E4@1, E5@1;

```



```

e BY C1@1, C2@1, C3@1, C4@1, C5@1, E1@1, E2@1, E3@1, E4@1, E5@1;
m BY C1@1, C2@1, C3@1, C4@1, C5@1, E1@1, E2@1, E3@1, E4@1, E5@1;

Class:
Tree

Constraints:
t = t1 | t2
"

model3 <- irtree_model(m3)

fit3 <- fit(model3, data = df2, control = control_mirt(method = "MHRM"))

##### IR-tree model constrained to Steps Model #####

m4 <- "
Equations:
1 = (1-a1)
2 = a1*(1-a2)
3 = a1*a2*(1-a3)
4 = a1*a2*a3*(1-a4)
5 = a1*a2*a3*a4

IRT:
a1 BY C1@1, C2@1, C3@1, C4@1, C5@1;
a2 BY C1@1, C2@1, C3@1, C4@1, C5@1;
a3 BY C1@1, C2@1, C3@1, C4@1, C5@1;
a4 BY C1@1, C2@1, C3@1, C4@1, C5@1;

Class:
Tree

Constraints:
a1 = a2
a1 = a3
a1 = a4
"

model4 <- irtree_model(m4)

fit4 <- fit(model4, data = df1)

# Partial Credit Model -----

##### Ordinary PCM #####

m5 <- "
IRT:
t BY C1@1, C2@1, C3@1, C4@1, C5@1;

Weights:
t = c(0, 1, 2, 3, 4)

```

```

Class:
PCM
"

model5 <- irtree_model(m5)

fit5 <- fit(model5, data = df1 - 1, engine = "tam")

##### Multidimensional PCM with constraints #####

m6 <- "
IRT:
t1 BY C1@1, C2@1, C3@1, C4@1, C5@1;
t2 BY E1@1, E2@1, E3@1, E4@1, E5@1;
e BY C1@1, C2@1, C3@1, C4@1, C5@1, E1@1, E2@1, E3@1, E4@1, E5@1;
m BY C1@1, C2@1, C3@1, C4@1, C5@1, E1@1, E2@1, E3@1, E4@1, E5@1;

Weights:
t = c(0, 1, 2, 3, 4)
e = c(1, 0, 0, 0, 1)
m = c(0, 0, 1, 0, 0)

Class:
PCM

Constraints:
t = t1 | t2
"

model6 <- irtree_model(m6)

fit6 <- fit(model6, data = df2 - 1, engine = "tam",
            control = control_tam(control = list(snodes = 1234)))

```

glance.irtree_fit *Glance at an irtree_fit object*

Description

Glance accepts an `irtree_fit` object and returns a [tibble](#) with exactly one row of model summaries.

Usage

```
## S3 method for class 'irtree_fit'
glance(x = NULL, ...)
```

Arguments

`x` object of class `irtree_fit` as returned from `fit()`.
`...` Additional arguments. Not used.

Value

A one-row [tibble](#) with columns such as AIC and BIC.

Converged:

The column `converged` indicates whether the model converged or not. For Mplus, this is TRUE if the output contained the phrase "The model estimation terminated normally". For mirt, this is equal to the output of `mirt::extract.mirt(x, "converged")`. For TAM, this is NA if no clear signs of non-convergence were observed. You are encouraged to check any warnings or errors in any case.

Iterations:

`iterations` is NA for Mplus models since respective information is not easily obtained from the output.

See Also

`generics::glance()`, `mirt::extract.mirt(x, "secondordertest")`

Examples

```
data("jackson")
df1 <- jackson[1:234, paste0("C", 1:5)]
irtree_create_template(df1)
m1 <- "
IRT:
t BY C1@1, C2@1, C3@1, C4@1, C5@1;
Class:
GRM"
fit1 <- fit(irtree_model(m1), data = df1)

tidy(fit1, par_type = "difficulty")

glance(fit1)

augment(fit1)
```

irtree_create_template

Create a template of a model string

Description

This function prints a template of a model string to the console based on the supplied data frame. This template can be copy-pasted and modified to define an [irtree_model](#).

Usage

```
irtree_create_template(data = NULL, mapping_matrix = NULL, rasch = TRUE)
```

Arguments

<code>data</code>	Data frame.
<code>mapping_matrix</code>	Matrix of so-called pseudo-items, optional. The observed response categories must appear in the first column. The other columns contain the pseudo-items and each entry may be either 1, 0, or NA.
<code>rasch</code>	Logical. The string @1 will be appended to each variable name if TRUE with no effect otherwise.

Examples

```
irtree_create_template(jackson[, c(1, 6, 11)])
#> m1 <- "
#> Equations:
#> 1 = ...
#> 2 = ...
#> 3 = ...
#> 4 = ...
#> 5 = ...
#>
#> IRT:
#> ... BY E1@1, E2@1, E3@1;
#> ... BY E1@1, E2@1, E3@1;
#>
#> Class:
#> Tree
#> "
```

```
irtree_create_template(jackson[, c(1, 6, 11)],
                       cbind(1:5,
                              m = c(0, 0, 1, 0, 0),
                              t = c(1, 1, NA, 0, 0),
                              e = c(1, 0, NA, 0, 1)))
#> m1 <- "
#> Equations:
#> 1 = (1-m)*t*e
#> 2 = (1-m)*t*(1-e)
#> 3 = m
#> 4 = (1-m)*(1-t)*(1-e)
#> 5 = (1-m)*(1-t)*e
#>
#> IRT:
#> m BY E1@1, E2@1, E3@1;
#> t BY E1@1, E2@1, E3@1;
#> e BY E1@1, E2@1, E3@1;
#>
#> Class:
#> Tree
```

```
#> "
```

```
irtree_gen_data      Generate data
```

Description

This function generates data from an ItemResponseTrees model.

Usage

```
irtree_gen_data(
  object = NULL,
  N = NULL,
  sigma = NULL,
  theta = NULL,
  itempar = NULL,
  link = c("logit", "probit"),
  na_okay = TRUE,
  skip = FALSE
)
```

Arguments

object	Object of class <code>irtree_model</code> . See irtree_model for more information.
N	Integer, the number of persons.
sigma	Either a matrix or a function that returns a matrix. This matrix is the variance-covariance matrix of the person parameters that is passed to <code>MASS::mvrnorm()</code> . Note that the order of the person parameters is taken from the section Processes in the model object (see irtree_model).
theta	Optional numeric matrix of person parameters with one row per person and one column per dimension (i.e., <code>object\$S</code>). If provided, this overrides N and sigma.
itempar	Either a list or a function that returns a list. The list has an element beta and an element alpha. Each of these is a matrix of item parameters. Note that the order of items (rows) is taken from the section Items and the order of processes (columns) is taken from the section Processes in the model (see irtree_model).
link	Character. Link function to use.
na_okay	Logical indicating whether variables with unobserved response categories are permitted. If FALSE, rejection sampling is used to ensure that all categories are observed.
skip	Logical. Some features of the irtree_model syntax, which are available for model fitting (e.g., Addendum), are not implemented for data generation. Those parts of the model are ignored if <code>skip = TRUE</code> .

Value

A list with element data containing the data and an element spec containing the true parameter values etc.

Examples

```
# IR-Tree Model -----

m1 <- "
Equations:
1 = (1-m)*(1-t)*e
2 = (1-m)*(1-t)*(1-e)
3 = m
4 = (1-m)*t*(1-e)
5 = (1-m)*t*e

IRT:
t BY x1, x2, x3;
e BY x1, x2, x3;
m BY x1, x2, x3;

Class:
Tree
"

model1 <- irtree_model(m1)

dat1 <- irtree_gen_data(model1, N = 5, sigma = diag(3),
                       itempar = list(beta = matrix(rnorm(9), 3, 3),
                                                alpha = matrix(1, 3, 3)))

dat1$data

# Partial Credit Model -----

m2 <- "
IRT:
t BY x1@1, x2@1, x3@1;
e BY x1@1, x2@1, x3@1;
m BY x1@1, x2@1, x3@1;

Weights:
t = c(0, 1, 2, 3, 4)
e = c(1, 0, 0, 0, 1)
m = c(0, 0, 1, 0, 0)

Class:
PCM
"

model2 <- irtree_model(m2)
dat2 <- irtree_gen_data(model2, N = 5, sigma = diag(3),
                       itempar = list(beta = matrix(sort(rnorm(12)), 3, 4)))

dat2$data
```

```

m3 <- "
IRT:
t BY x1@1, x2@1, x3@1;

Weights:
t = 0:4

Class:
PCM
"

model3 <- irtree_model(m3)

dat3 <- irtree_gen_data(model3, N = 5, sigma = diag(1),
                       itempar = list(beta = matrix(sort(rnorm(12)), 3, 4)))
dat3$data

```

irtree_model	<i>ItemResponseTrees model syntax</i>
--------------	---------------------------------------

Description

The *ItemResponseTrees* model syntax describes the statistical model. The function `irtree_model()` turns a user-defined model string into an object of class `irtree_model` that represents the full model as needed by the package.

Usage

```
irtree_model(model = NULL)
```

Arguments

`model` String with a specific structure as described below.

Value

List of class `irtree_model`. It contains the information extracted from `model`. Side note: The returned list contains an element `mapping_matrix` that contains the pseudoitems. This information is instructive, and it might be used as an input to the `dendriify()` function of the `irtrees` package.

Overview of the Model Syntax

1. The model string must contain at least the sections **IRT**, **Class**, and (if class is tree) **Equations**.
2. Section headings must appear on a separate line ending with a colon (:).
3. The model may contain empty lines and commented lines, which begin with # (do not use inline comments).

4. Line breaks are only allowed in section **IRT**.

Details for all the required and optional sections of the model string are given in the following.

Equations:

The model must contain a section with heading **Equations** if **Class** is **Tree**. Therein, the model equations are described. They have a structure similar to $Cat = p_1 * (1 - p_2)$, where *Cat* is any observed response category in the data set, and where p_1 is a freely chosen name of a parameter. These names must match the names of the latent variables in the section **IRT** (combined with **Constraints** if specified).

If you prefer to work with pseudo-items, `irtree_create_template()` can generate the equations for you.

The equations may contain only products and not sums. That is, it is not possible to estimate genuine mixture models as, for example, in the package `mpt2irt`.

Each equation must appear on a separate, non-broken line. For example:

Equations:

```
1 = (1-m)*(1-t)*e
2 = (1-m)*(1-t)*(1-e)
3 = m
4 = (1-m)*t*(1-e)
5 = (1-m)*t*e
```

IRT:

The model must contain a section with heading **IRT**. Therein, the IRT structure of the model is described in a way resembling the **MODEL** part of an **Mplus** input file. It has a structure of **LV BY item1*, item2@1**, where **LV** is the name of the latent variable/parameter/process, and **item** is the name of the observed variable in the data set, which is followed by the loading. The loading may either be fixed (e.g., to 1) using **@1** or it may be set free using ***** (which is equivalent to omitting a loading altogether).

Each measurement model (i.e., the LV and its items) must appear on a separate line ending with a semicolon. Items must be separated by commas. Line breaks are allowed. For example:

IRT:

```
t BY x1, x2, x3, x4, x5, x6;
e BY x1@1, x2@1, x3@1, x4@1, x5@1, x6@1;
m BY x1@1, x2@1, x3@1, x4@1, x5@1, x6@1;
```

Class:

The model must contain a section with heading **Class** to specify the type/class of IRT model to use. Currently, may be either **Tree**, **GRM**, or **PCM**. For example:

```
Class:
Tree
```

Constraints:

The model may contain a section with heading **Constraints** to specify equality constraints of latent variables. Constraints may be useful for multidimensional questionnaires to link **IRT** and **Equations** in a specific way. Or, latent variables in **IRT** may be constrained to equality.

Constraints in order to link sections IRT and Equations:

A process in the model equations (section **Equations**) may correspond to multiple latent variables (section **IRT**). For example, when analyzing a Big Five data set, one may wish to specify only one extremity process e for all items but multiple target traits t , namely, one for each of the five scales. In such a case, the section **Equations** would list only the parameter t , while the section **IRT** would list the parameters $t1, \dots, t5$.

In the framework of MPT, one would think of such a situation in terms of multiple albeit similar trees with specific parameter constraints across trees. For example, one would use one tree for each Big Five scale and fix the response style parameters to equality across trees but not the target trait parameters.

Each line in this section has a structure of Param = LV1 | LV2, where Param is the name of the process used only in section **Equations** and LV1 is the name of the process used only in section **IRT**. Use one line for each definition. For example:

Constraints:

t = t1 | t2 | t3 | t4 | t5

Constraints within section IRT:

For example, in a sequential model as proposed by Tutz as well as Verhelst, one would specify two processes for a 3-point item. The first process would correspond to a pseudoitem of $\theta-1-1$ and the second process to a pseudoitem of $NA-\theta-1$. However, the latent variables corresponding to these two processes would typically be assumed to be equal and need thus be constrained accordingly.

Each line in this section has a structure of LV1 = LV2, where LV1 and LV2 are the names of the latent variables used in section **IRT**. Use one line for each definition. For example:

Constraints:

LV1 = LV2

LV1 = LV3

Addendum:

The model may contain a section with heading **Addendum** if engine = "mplus" is used for estimation. Any code in this section is directly pasted in the MODEL section of the Mplus input file. Use a semicolon at the end of each line; lines must not exceed 90 characters. Note that the addendum is ignored in `irtree_gen_data()`. For example:

Addendum:

e WITH t@0;

m WITH t@0;

Weights:

The model may contain a section with heading **Weights** if model **Class** is PCM. This allows to specify (uni- and) multidimensional partial credit models. They have been proposed, for example, by Wetzel and Carstensen (2017), as an alternative to IR-tree models. Note that fitting these models is only implemented for engine = "tam".

Each line in this section has a structure of LV = weights, where LV is the name of the latent variable used in section **IRT**. `weights` must be valid R code, namely, a vector of weights (see, e.g., Table 1 in Wetzel & Carstensen, 2017, or Table 2 in Falk & Cai, 2015). Use one line for each definition. For example:

Weights:

t = c(0, 1, 2, 3, 4)

```
e = c(1, 0, 0, 0, 1)
m = c(0, 0, 1, 0, 0)
```

Examples

```
m1 <- "
# Random comment

Equations:
1 = (1-m)*(1-t)*e
2 = (1-m)*(1-t)*(1-e)
3 = m
4 = (1-m)*t*(1-e)
5 = (1-m)*t*e

IRT:
t1 BY x1@1, x2*, x3*;
t2 BY x4@1, x5*, x6*;
e BY x1@1, x2@1, x3@1, x4@1, x5@1, x6@1;
m BY x1@1, x2@1, x3@1, x4@1, x5@1, x6@1;

Constraints:
t = t1 | t2

Class:
Tree
"

model <- irtree_model(m1)
```

irtree_recode	<i>Recode data into pseudoitems</i>
---------------	-------------------------------------

Description

This function takes a data set with polytomous items and an `irtree_model` and returns the recoded items, the so-called pseudoitems.

Usage

```
irtree_recode(object = NULL, data = NULL, keep = FALSE)
```

Arguments

object	Object of class <code>irtree_model</code> . See irtree_model for more information.
data	Data frame containing containing one row per respondent and one column per variable. The variable names must correspond to those used in <code>object</code> .
keep	Logical indicating whether to append the original items to the data frame of the generated pseudoitems

Value

Data frame

Examples

```

m1 <- "
IRT:
t BY x1;
e BY x1;
m BY x1;
Equations:
1 = (1-m)*(1-t)*e
2 = (1-m)*(1-t)*(1-e)
3 = m
4 = (1-m)*t*(1-e)
5 = (1-m)*t*e
Class:
Tree
"

model1 <- irtree_model(m1)
dat <- data.frame(x1 = 1:5)
irtree_recode(model1, dat, keep = TRUE)

```

irtree_sim

Run a simulation by generating from and fitting an ItemResponseTrees model

Description

The function `irtree_sim()` generates data from an [irtree_model](#) and fits one or more models to these data. This process is repeated `R` times, and the argument `plan` allows to run the simulation in parallel.

Usage

```

irtree_sim(
  R = 1,
  gen_model = NULL,
  fit_model = gen_model,
  N = NULL,
  sigma = NULL,
  itempar = NULL,
  link = c("logit", "probit"),
  na_okay = TRUE,
  engine = c("mirt", "mplus", "tam"),
  verbose = FALSE,
  control = NULL,
  improper_okay = FALSE,

```

```

par_type = "difficulty",
plan = NULL,
plan_args = list(),
file = NULL,
dir = tempdir(),
save_rdata = TRUE,
in_memory = c("reduced", "everything", "nothing")
)

```

Arguments

R	Number of replications. Can be either a single number indicating the number of replications (e.g., R = 100), or can be a range (e.g., R = 1 : 100).
gen_model	Object of class <code>irtree_model</code> describing the data-generating model. See irtree_model for more information.
fit_model	Object of class <code>irtree_model</code> describing the model that should be fit to the data. May be a list of multiple objects of class <code>irtree_model</code> if different models should be fit to the same data set. See irtree_model for more information.
N	Integer, the number of persons.
sigma	Either a matrix or a function that returns a matrix. This matrix is the variance-covariance matrix of the person parameters that is passed to <code>MASS::mvrnorm()</code> . Note that the order of the person parameters is taken from the section Processes in the model object (see irtree_model).
itempar	Either a list or a function that returns a list. The list has an element beta and an element alpha. Each of these is a matrix of item parameters. Note that the order of items (rows) is taken from the section Items and the order of processes (columns) is taken from the section Processes in the model (see irtree_model).
link	Character. Link function to use.
na_okay	Logical indicating whether variables with unobserved response categories are permitted. If FALSE, rejection sampling is used to ensure that all categories are observed.
engine	String specifying whether to use mirt, Mplus, or TAM for estimation.
verbose	Logical indicating whether output should be printed to the console.
control	List. The allowed elements of this list depend on the engine. Use control_mirt() , control_mplus() , or control_tam() for convenience. Note that the <code>fit()</code> function does not use <code>...</code> , but that you can use the <code>control_*</code> () functions to pass additional arguments.
improper_okay	Logical indicating whether the model should also be fit if it is not a proper IR-tree model. Set this only to TRUE if you really know what you are doing.
par_type	Only used if the fit engine was mirt. Item parameters (or thresholds) can be either of type <code>easiness</code> (the mirt default) or <code>difficulty</code> (as in Mplus and TAM).
plan	Parameter passed as argument <code>strategy</code> to future::plan() . May be set to, for example, <code>multiprocess</code> in order to run the simulations in parallel.
plan_args	Named list. Parameters passed future::plan() .

file	String giving the file path used to save the output if <code>save_rdata = TRUE</code> . Note that the file ending is automatically set to <code>.rda</code> . This argument is also passed to <code>irtree_fit_mplus()</code> if applicable.
dir	Path name that is used to save the results of every run if <code>save_rdata = TRUE</code> .
save_rdata	Logical indicating whether to save the results to an RData file.
in_memory	Character string indicating what output should be kept in memory (note the argument <code>save_rdata</code> , which is not affected by <code>in_memory</code>). If "reduced", the output of <code>fit()</code> is discarded and only summary information is retained. The alternative is to keep "everything" in memory, or to keep "nothing" in memory (which makes only sense in combination with <code>save_rdata = TRUE</code>).

Value

Returns a list of length `R`. For each replication, a list is returned with two elements. The element `spec` contains various specifications (such as the data). The element `fits` is a list with one element for each `fit_model` that contains the output of `fit()` as well as the elements `glanced`, `tidied`, and `augmented` (see `glance()`, `tidy()`, and `augment()`). Thus, `res$sim3$fits$m2$glanced` gives model-fit information such as AIC for the second model in the third replication, and `res$sim3$spec$data` contains the corresponding data set.

If `in_memory = "nothing"`, returns `NULL`.

See Also

The data are generated via `irtree_gen_data()`, and the models are fit via `fit()`.

Examples

```
# Running these examples may take a while

m1 <- "
Equations:
1 = 1-a
2 = a*(1-b)
3 = a*b

IRT:
a BY x1@1, x2@1, x3@1, x4@1, X5@1, X6@1, X7@1;
b BY x1@1, x2@1, x3@1, x4@1, X5@1, X6@1, X7@1;

Class:
Tree
"

m2 <- "
IRT:
a BY x1@1, x2@1, x3@1, x4@1, X5@1, X6@1, X7@1;

Class:
GRM
```

```

"

model1 <- irtree_model(m1)
model2 <- irtree_model(m2)

res <- irtree_sim(
  ### Data generation ###
  gen_model = model1,
  link = "logit",
  N = 500,
  sigma = function(x) diag(2),
  itempar = function(x) list(
    beta = matrix(sort(runif(model1$J*model1$P, -2, 2)),
                  model1$J, model1$P),
    alpha = matrix(1, model1$J, model1$P)),
  na_okay = FALSE,

  ### Estimation ###
  fit_model = list(model1, model2),
  engine = "mirt",
  control = control_mirt(SE = FALSE),
  par_type = "difficulty",

  ### Replications ###
  R = 2,
  save_rdata = FALSE,

  ### Optional parallelization ###
  plan = "multiprocess",
  plan_args = list(workers = future::availableCores() - 1)
)

tab1 <- matrix(NA, 0, 4, dimnames = list(NULL, c("Rep", "Model", "AIC", "BIC")))

for (ii in seq_along(res)) {
  for (jj in seq_along(res[[ii]]$fits)) {
    IC <- res[[ii]]$fits[[jj]]$glanced
    tab1 <- rbind(tab1, c(ii, jj, round(IC$AIC, -1), round(IC$BIC, -1)))
  }
}

tab1
#>      Rep Model  AIC  BIC
#> [1,]   1     1 6900 6970
#> [2,]   1     2 7000 7060
#> [3,]   2     1 6810 6880
#> [4,]   2     2 6880 6940

```

Description

"This is data from an online big five personality test: <http://personality-testing.info/tests/BIG5.php>. The following items were rated on a likert scale from 1=disagree to 5=agree in relation to how much they applied to the test taker, they were presented to the taker 5 per page" (Jackson, 2012).

The following items are reverse keyed and were already recoded: A1, E2, C2, N2, O2, A3, E4, C4, N4, O4, A5, E6, C6, O6, A7, E8, C8, and E10.

Usage

```
data("jackson")
```

Format

A data frame with 9051 rows and 58 variables:

- E1** Am the life of the party.
- A1** Feel little concern for others.
- C1** Am always prepared.
- N1** Get stressed out easily.
- O1** Have a rich vocabulary.
- E2** Don't talk a lot.
- A2** Am interested in people.
- C2** Leave my belongings around.
- N2** Am relaxed most of the time.
- O2** Have difficulty understanding abstract ideas.
- E3** Feel comfortable around people.
- A3** Insult people.
- C3** Pay attention to details.
- N3** Worry about things.
- O3** Have a vivid imagination.
- E4** Keep in the background.
- A4** Sympathize with others' feelings.
- C4** Make a mess of things.
- N4** Seldom feel blue.
- O4** Am not interested in abstract ideas.
- E5** Start conversations.
- A5** Am not interested in other people's problems.
- C5** Get chores done right away.
- N5** Am easily disturbed.
- O5** Have excellent ideas.
- E6** Have little to say.

- A6** Have a soft heart.
- C6** Often forget to put things back in their proper place.
- N6** Get upset easily.
- O6** Do not have a good imagination.
- E7** Talk to a lot of different people at parties.
- A7** Am not really interested in others.
- C7** Like order
- N7** Change my mood a lot.
- O7** Am quick to understand things.
- E8** Don't like to draw attention to myself.
- A8** Take time out for others.
- C8** Shirk my duties.
- N8** Have frequent mood swings.
- O8** Use difficult words.
- E9** Don't mind being the center of attention.
- A9** Feel others' emotions.
- C9** Follow a schedule.
- N9** Get irritated easily.
- O9** Spend time reflecting on things.
- E10** Am quiet around strangers.
- A10** Make people feel at ease.
- C10** Am exacting in my work.
- N10** Often feel blue.
- O10** Am full of ideas.
- gender** Gender, either female, male, or other
- age** Age
- SecondsElapsed** Seconds elapsed
- E** Scale mean for extraversion
- C** Scale mean for conscientiousness
- N** Scale mean for neuroticism
- O** Scale mean for openness
- A** Scale mean for agreeableness

Details

The data set included here is Version 3 from 15.10.2012. It was released by Andrew Jackson under the [CC BY 4.0](https://creativecommons.org/licenses/by/4.0/) license.

Source

Jackson, A. (2012). IPIP Big Five personality test answers. <https://doi.org/10.6084/m9.figshare.96542.v3>

Description

IR-tree models can be fit on the basis of so-called pseudo-items. To this end, the original, polytomous items are recoded into binary pseudo-items. Whether a pseudo-item is coded as 1, 0, or NA depends on the model equations (e.g., Böckenholt, 2012; Plieninger, 2020).

The ItemResponseTrees package internally works with pseudo-items as well. However, the user has to specify the model equations rather than the pseudo-items in the `irtree_model` syntax. Internally, the original responses are recoded on the basis of the model supplied by the user by the function `irtree_recode()`. This function may also be used directly if desired.

As an alternative to specifying the model equations themselves, users may also use the function `irtree_create_template()` with a mapping matrix (that specifies the structure of the pseudo-items) to generate the model equations automatically.

References

Böckenholt, U. (2012). Modeling multiple response processes in judgment and choice. *Psychological Methods*, 17(4), 665–678. <https://doi.org/10.1037/a0028111>

Plieninger, H. (2020). Developing and applying IR-tree models: Guidelines, caveats, and an extension to multiple groups. *Organizational Research Methods*. Advance online publication. <https://doi.org/10.1177/1094428120911096>

Examples

```
# Mapping matrix for data with three response categories:
(mm <- cbind(cat = 0:2,
             p1 = c(0, 1, 1),
             p2 = c(NA, 0, 1)))
#>      cat p1 p2
#> [1,]  0  0 NA
#> [2,]  1  1  0
#> [3,]  2  1  1

irtree_create_template(data.frame(x1 = 0:2, x2 = 0:2),
                      mapping_matrix = mm)
#>
#> m1 <- "
#> Equations:
#> 0 = (1-p1)
#> 1 = p1*(1-p2)
#> 2 = p1*p2
#>
#> IRT:
#> p1 BY x1@1, x2@1;
#> p2 BY x1@1, x2@1;
#>
```

```
#> Class:
#> Tree
#> "
#>
```

tidy.irtree_fit	<i>Tidy an irtree_fit object</i>
-----------------	----------------------------------

Description

Tidy summarizes information about the parameter estimates of an ItemResponseTrees model.

Usage

```
## S3 method for class 'irtree_fit'
tidy(x = NULL, par_type = NULL, ...)
```

Arguments

x	object of class <code>irtree_fit</code> as returned from <code>fit()</code> .
par_type	Only used if the fit engine was <code>mirt</code> . Item parameters (or thresholds) can be either of type <code>easiness</code> (the <code>mirt</code> default) or <code>difficulty</code> (as in <code>Mplus</code> and <code>TAM</code>).
...	Not currently used.

Value

A [tibble](#) with one row for each model parameter and the following columns:

`term` The name of the model parameter.

`estimate` The estimated value of the term.

`std.error` The standard error of the term.

`statistic` The value of the test statistic of the term (`Mplus` only).

`p.value` The p-value associated with the statistic (`Mplus` only).

See Also

[generics::tidy\(\)](#)

Examples

```
data("jackson")
df1 <- jackson[1:234, paste0("C", 1:5)]
irtree_create_template(df1)
m1 <- "
IRT:
t BY C1@1, C2@1, C3@1, C4@1, C5@1;
Class:
GRM"
fit1 <- fit(irtree_model(m1), data = df1)

tidy(fit1, par_type = "difficulty")

glance(fit1)

augment(fit1)
```

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