Package: nlmixr2targets (via r-universe)

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Title Targets for 'nlmixr2' Pipelines
Version 0.0.0.9000
Description 'nlmixr2' often has long runtimes. A pipeline toolkit tailored to 'nlmixr2' workflows leverages 'targets' and 'nlmixr2' to ease reproducible workflows. 'nlmixr2targets' ensures minimal rework in model development with 'nlmixr2' and 'targets' by simplifying and standardizing models and datasets.
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<pre>URL https://nlmixr2.github.io/nlmixr2targets/</pre>
VignetteBuilder knitr
Repository https://nlmixr2.r-universe.dev
RemoteUrl https://github.com/nlmixr2/nlmixr2targets
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assign_origData

Replace the fit data with the original data, then return the modified fit

Description

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This function is intended for use within nlmixr2targets target creation, and it's not typically invoked by users.

Usage

```
assign_origData(fit, data)
```

Arguments

fit an estimated nlmixr2 object data the data from the original fit

Value

The fit with the data added back in as fit\$env\$origData

nlmixr_data_simplify Standardize and simplify data for nlmixr2 estimation

Description

This function is typically not needed by end users.

Usage

```
nlmixr_data_simplify(data, object, table = list())
```

Arguments

data nlmixr data

object an nlmixr_ui object (e.g. the output of running nlmixr(object = model)

table The output table control object (like 'tableControl()')

Details

The standardization keeps columns that rxode2 and nlmixr2 use along with the covariates. Column order is standardized (rxode2 then nlmixr2 then alphabetically sorted covariates), and rxode2 and nlmixr2 column names are converted to lower case.

Value

The data with the nlmixr2 column lower case and on the left and the covariate columns on the right and alphabetically sorted.

See Also

```
Other Simplifiers: nlmixr_object_simplify()
```

```
nlmixr_object_simplify
```

Simplify an nlmixr object

Description

This function is typically not needed by end users.

Usage

```
nlmixr_object_simplify(object)
```

Arguments

object

Fitted object or function specifying the model.

Details

The object simplification removes comments (so please use label() instead of comments to label parameters) and then converts the object to a "nlmixrui" object.

Since setting initial conditions with cmt(0) does not work with targets, the function definition of the object must set it with cmt(initial). cmt(initial) will be converted to cmt(0) before passing to nlmixr2.

Value

object converted to a nlmixrui object. The model name is always "object".

See Also

```
Other Simplifiers: nlmixr_data_simplify()
```

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tar_nlmixr

Generate a set of targets for nlmixr estimation

Description

The targets generated will include the name as the final estimation step, paste(name, "object_simple", sep = "_tar_") (e.g. "pheno_tar_object_simple") as the simplified model object, and paste(name, "data_simple", sep = "_tar_") (e.g. "pheno_tar_data_simple") as the simplified data object.

Usage

```
tar_nlmixr(
  name,
 object,
  data,
  est = NULL,
  control = list(),
  table = nlmixr2est::tableControl(),
  env = parent.frame()
)
tar_nlmixr_raw(
  name,
  object,
  data,
  est,
  control,
  table,
  object_simple_name,
  data_simple_name,
  fit_simple_name,
  env
)
```

Arguments

name

Symbol, name of the target. A target name must be a valid name for a symbol in R, and it must not start with a dot. Subsequent targets can refer to this name symbolically to induce a dependency relationship: e.g. tar_target(downstream_target, f(upstream_target)) is a target named downstream_target which depends on a target upstream_target and a function f(). In addition, a target's name determines its random number generator seed. In this way, each target runs with a reproducible seed so someone else running the same pipeline should get the same results, and no two targets in the same pipeline share the same seed. (Even dynamic branches have different names and thus different seeds.) You can recover the seed of a completed target with tar_meta(your_target, seed) and

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	run $tar_seed_set()$ on the result to locally recreate the target's initial RNG state.			
object	Fitted object or function specifying the model.			
data	nlmixr data			
est	estimation method (all methods are shown by 'nlmixr2AllEst()'). Methods can be added for other tools			
control	The estimation control object. These are expected to be different for each type of estimation method			
table	The output table control object (like 'tableControl()')			
env	The environment where the model is setup (not needed for typical use)			
object_simple_name, data_simple_name, fit_simple_name				
	target names to use for the simplified object, simplified data, fit of the simplified object with the simplified data, and fit with the original data re-inserted.			

Details

For the way that the objects are simplified, see nlmixr_object_simplify() and nlmixr_data_simplify(). To see how to write initial conditions to work with targets, see nlmixr_object_simplify().

Value

A list of targets for the model simplification, data simplification, and model estimation.

Functions

• tar_nlmixr_raw(): An internal function to generate the targets

Examples

```
## Not run:
library(targets)
targets::tar_script({
pheno <- function() {</pre>
  ini({
    lcl <- log(0.008); label("Typical value of clearance")</pre>
    lvc <- log(0.6); label("Typical value of volume of distribution")</pre>
    etalcl + etalvc \sim c(1,
                           0.01, 1)
    cpaddSd <- 0.1; label("residual variability")</pre>
  })
  model({
    cl <- exp(lcl + etalcl)</pre>
    vc <- exp(lvc + etalvc)</pre>
    kel <- cl/vc
    d/dt(central) <- -kel*central</pre>
    cp <- central/vc</pre>
    cp ~ add(cpaddSd)
 })
}
```

```
list(
  tar_nlmixr(
    name = pheno_model,
    object = pheno,
    data = nlmixr2data::pheno_sd,
    est = "saem"
  )
)
})
targets::tar_make()
## End(Not run)
```

tar_nlmixr_multimodel Generate a list of models based on a single dataset and estimation method

Description

Generate a list of models based on a single dataset and estimation method

Usage

```
tar_nlmixr_multimodel(
  name,
    ...,
  data,
  est,
  control = list(),
  table = nlmixr2est::tableControl(),
  env = parent.frame()
)
```

Arguments

name

Symbol, name of the target. A target name must be a valid name for a symbol in R, and it must not start with a dot. Subsequent targets can refer to this name symbolically to induce a dependency relationship: e.g. tar_target(downstream_target, f(upstream_target)) is a target named downstream_target which depends on a target upstream_target and a function f(). In addition, a target's name determines its random number generator seed. In this way, each target runs with a reproducible seed so someone else running the same pipeline should get the same results, and no two targets in the same pipeline share the same seed. (Even dynamic branches have different names and thus different seeds.) You can recover the seed of a completed target with tar_meta(your_target, seed) and run tar_seed_set() on the result to locally recreate the target's initial RNG state.

.. Named arguments with the format "Model description" = modelFunction

data	nlmixr data
------	-------------

est estimation method (all methods are shown by 'nlmixr2AllEst()'). Methods can

be added for other tools

control The estimation control object. These are expected to be different for each type

of estimation method

table The output table control object (like 'tableControl()')

env The environment where the model is setup (not needed for typical use)

Value

A list of targets for the model simplification, data simplification, and model estimation.

```
tar_nlmixr_multimodel_has_self_reference
```

Does the model list refer to another model in the model list?

Description

Does the model list refer to another model in the model list?

Usage

```
tar_nlmixr_multimodel_has_self_reference(model_list, name)
tar_nlmixr_multimodel_has_self_reference_single(model, name)
```

Arguments

model_list A named list of calls for model targets to be created

name Symbol, name of the target. A target name must be a valid name for a symbol in

R, and it must not start with a dot. Subsequent targets can refer to this name symbolically to induce a dependency relationship: e.g. tar_target(downstream_target,

f(upstream_target)) is a target named downstream_target which depends on a target upstream_target and a function f(). In addition, a target's name determines its random number generator seed. In this way, each target runs with a reproducible seed so someone else running the same pipeline should get the same results, and no two targets in the same pipeline share the same seed. (Even dynamic branches have different names and thus different seeds.) You can recover the seed of a completed target with tar_meta(your_target, seed) and run tar_seed_set() on the result to locally recreate the target's initial RNG

state

model A single model call for the model target to be created

Value

A logical vector the same length as model_list indicating if the model is self-referential to another model in the list

Functions

• tar_nlmixr_multimodel_has_self_reference_single(): A helper function to look at each call for each model separately

tar_nlmixr_multimodel_parse

Generate nlmixr multimodel target set for all models in one call to tar_nlmixr_multimodel()

Description

Generate nlmixr multimodel target set for all models in one call to tar_nlmixr_multimodel()

Usage

```
tar_nlmixr_multimodel_parse(name, data, est, control, table, model_list, env)
```

Arguments

name	Symbol, name of the target. A	\ target name must be a	valid name for a symbol in

R, and it must not start with a dot. Subsequent targets can refer to this name symbolically to induce a dependency relationship: e.g. tar_target(downstream_target, f(upstream_target)) is a target named downstream_target which depends on a target upstream_target and a function f(). In addition, a target's name determines its random number generator seed. In this way, each target runs with a reproducible seed so someone else running the same pipeline should get the same results, and no two targets in the same pipeline share the same seed. (Even dynamic branches have different names and thus different seeds.) You can recover the seed of a completed target with tar_meta(your_target, seed) and run tar_seed_set() on the result to locally recreate the target's initial RNG

state.

data nlmixr data

est estimation method (all methods are shown by 'nlmixr2AllEst()'). Methods can

be added for other tools

control The estimation control object. These are expected to be different for each type

of estimation method

table The output table control object (like 'tableControl()')

model_list A named list of calls for model targets to be created

env The environment where the model is setup (not needed for typical use)

tar_nlmixr_multimodel_single

Generate a single nlmixr multimodel target set for one model

Description

Generate a single nlmixr multimodel target set for one model

Usage

```
tar_nlmixr_multimodel_single(object, name, data, est, control, table, env)
```

Arguments

object Fitted object or function specifying the model.

name Symbol, name of the target. A target name must be a valid name for a symbol in

R, and it must not start with a dot. Subsequent targets can refer to this name symbolically to induce a dependency relationship: e.g. tar_target(downstream_target,

f(upstream_target)) is a target named downstream_target which depends on a target upstream_target and a function f(). In addition, a target's name determines its random number generator seed. In this way, each target runs with a reproducible seed so someone else running the same pipeline should get the same results, and no two targets in the same pipeline share the same seed. (Even dynamic branches have different names and thus different seeds.) You can recover the seed of a completed target with tar_meta(your_target, seed) and

run tar_seed_set() on the result to locally recreate the target's initial RNG

state.

data nlmixr data

est estimation method (all methods are shown by 'nlmixr2AllEst()'). Methods can

be added for other tools

control The estimation control object. These are expected to be different for each type

of estimation method

table The output table control object (like 'tableControl()')

env The environment where the model is setup (not needed for typical use)

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