

Package ‘toxpiR’

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Type Package

Title Create ToxPi Prioritization Models

Version 1.0.3

Description Enables users to build 'ToxPi' prioritization models and provides functionality within the grid framework for plotting ToxPi graphs. 'toxpiR' allows for more customization than the 'ToxPi GUI' (<<https://toxpi.org>>) and integration into existing workflows for greater ease-of-use, reproducibility, and transparency. toxpiR package behaves nearly identically to the GUI; the package documentation includes notes about all differences. The vignettes download example files from <<https://github.com/ToxPi/ToxPi-example-files>>.

Imports grDevices, methods, S4Vectors, grid, rlang, stats, BiocGenerics, pryr, tidyr, utils

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BugReports <https://github.com/ToxPi/toxpiR/issues>

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boxLegendGrob	<i>Create a filled-box legend</i>
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Description

Create a filled-box legend

Usage

```
boxLegendGrob(labels, fills, name = NULL, vp = NULL, gp = NULL)
```

Arguments

labels	Character, the legend labels
fills	Colors to fill the slices
name, vp, gp	Passed to grid::frameGrob

Details

Not yet exported. Need to break out the creation of viewports and grobs as done in the exported grobs. This will allow better grobEdit methods, which also needs to be created for the boxLegendGrob. Also need to do some input checks.

Also, if `grid::legendGrob` gets updated to use the 'has.fill' option this function should be removed and `grid::legendGrob` can be used instead.

pieGridGrob	<i>Make grid of pieGrobs</i>
-------------	------------------------------

Description

Make grid of pieGrobs

Usage

```
pieGridGrob(  
  radMat,  
  wts = NULL,  
  fills = NULL,  
  labels = NULL,  
  showRadSum = FALSE,  
  ncol = NULL,  
  nrow = NULL,  
  byrow = TRUE,  
  name = NULL,  
  gp = NULL,  
  vp = NULL  
)
```

```
grid.pieGridGrob(  
  radMat,  
  wts = NULL,  
  fills = NULL,  
  labels = NULL,  
  showRadSum = FALSE,  
  ncol = NULL,  
  nrow = NULL,  
  byrow = TRUE,  
  name = NULL,  
  gp = NULL,  
  vp = NULL  
)
```

Arguments

radMat	matrix(<numeric>), observations by slice radii
wts	vector(<numeric>), relative weights of each slice
fills	Vector of colors to fill slices
labels	vector(<character>), (optional) label for each observation
showRadSum	Logical scalar, when TRUE show the weighted sum of slices below the label
nrow, ncol	Integer scalar, number of rows and columns for the grid
byrow	Logical scalar, fill the grid by rows when TRUE
name, gp, vp	Passed to grid::gTree

Value

pieGrob [grid::grob](#) object

Examples

```
library(grid)

s <- seq(0.2, 1, by = 0.1)
smat <- do.call("rbind", replicate(20, s, simplify = FALSE))
grid.newpage()
grid.pieGridGrob(radMat = smat)

rownames(smat) <- sprintf("obs%02d", 1:20)
grid.newpage()
grid.pieGridGrob(radMat = smat, wts = s)
grid.newpage()
grid.pieGridGrob(radMat = smat, wts = s, showRadSum = TRUE, labels = FALSE)
grid.newpage()
grid.pieGridGrob(radMat = smat, labels = "hello")
grid.newpage()
grid.pieGridGrob(radMat = smat, labels = 1:20)

## Can edit like normal grid objects
grid.newpage()
grid.pieGridGrob(radMat = smat, wts = s, showRadSum = TRUE)
grid.ls() ## shows grid elements
grid.edit("pie-20", fills = 1:9)
grid.edit("pie-19-label", gp = gpar(font = 2, col = "red"))
grid.edit("pie-1", wts = rep(1, 9), rads = rep(1, 9))
for (s in sprintf("pie-%d-radSum", 2:4)) {
  grid.edit(s, gp = gpar(font = 2, col = "blue"))
}
```

pieGrob	<i>Create a pie grob</i>
---------	--------------------------

Description

Create a pie grob

Usage

```
pieGrob(rads, fills = NULL, wts = NULL, name = NULL, vp = NULL, gp = NULL)
```

```
grid.pieGrob(rads, fills = NULL, wts = NULL, name = NULL, vp = NULL, gp = NULL)
```

Arguments

rads	Numeric, radius values for each slice from 0 to 1
fills	Colors to fill the slices
wts	Numeric, the relative portion of each slice
name, vp, gp	Passed to grid::gTree

Details

The default coloring can be set with `options("txp.fills")`.

Value

pieGrob [grid::grob](#) object

Examples

```
library(grid)

s <- seq(0.2, 1, by = 0.1)
grid.newpage()
grid.pieGrob(rads = s)
grid.newpage()
grid.pieGrob(rads = s, wts = s)

curr_txp_fills <- options()$txp.fills
options(txp.fills = 1:8)
grid.newpage()
grid.pieGrob(rads = s)
options(txp.fills = curr_txp_fills)

## Can edit
grid.newpage()
grid.pieGrob(rads = s, name = "myPie")
grid.ls() ## show the grid elements
```

```
grid.edit("myPie", fills = 1:9, wts = 9:1)
```

toxiR-datasets	<i>toxiR data objects</i>
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Description

Objects included in the toxiR package, loaded with `utils::data`

Usage

```
data(txp_example_input, package = "toxiR")
```

```
data(txp_example_model, package = "toxiR")
```

txp_example_input

Small example input data to be used with `txpCalculateScores` in creating `TxpResult` objects. A `base::data.frame` with 10 rows and 9 variables

name Observation names

metric# Input data for ToxPi models

txp_example_model

Example `TxpModel` object intended for `txp_example_data`; model with 4 slices.

Source

<https://github.com/ToxPi/ToxPi-example-files>

Examples

```
data(txp_example_input, package = "toxiR")
data(txp_example_model, package = "toxiR")
txp_example_input
txp_example_model

## Code to create txp_example_model
tf1 <- TxpTransFuncList(linear = function(x) x)
s1 <- TxpSliceList(s1 = TxpSlice(sprintf("metric%d", 1:2)),
                  s2 = TxpSlice("metric3"),
                  s3 = TxpSlice(sprintf("metric%d", 4:7),
                                tf1[rep("linear", 4)]),
                  s4 = TxpSlice("metric8", tf1))
tf2 <- TxpTransFuncList(NULL, linear = function(x) x, NULL, NULL)
TxpModel(txpSlices = s1, txpWeights = c(2, 1, 3, 2), txpTransFuncs = tf2)
```

txpCalculateScores *Calculate ToxPi Scores for the given model and input data*

Description

Calculate ToxPi Scores for the given model and input data

Usage

```
txpCalculateScores(model, input, ...)  
  
## S4 method for signature 'TxpModel,data.frame'  
txpCalculateScores(  
  model,  
  input,  
  id.var = NULL,  
  rank.ties.method = c("average", "first", "last", "random", "max", "min")  
)  
  
## S4 method for signature 'TxpModelList,data.frame'  
txpCalculateScores(  
  model,  
  input,  
  id.var = NULL,  
  rank.ties.method = c("average", "first", "last", "random", "max", "min")  
)  
  
## S4 method for signature 'list,data.frame'  
txpCalculateScores(  
  model,  
  input,  
  id.var = NULL,  
  rank.ties.method = c("average", "first", "last", "random", "max", "min")  
)
```

Arguments

model	TxpModel object or TxpModelList object
input	data.frame object containing the model input data
...	Included for extendability; not currently used
id.var	Character scalar, column in 'input' to store in
rank.ties.method	Character scalar, passed to base::rank

Details

txpCalculateScores is implemented as an S4 generic function with methods for [TxpModel](#) and [TxpModelList](#).

Ranks are calculated such that the highest ToxPi score has a rank of 1.

Value

[TxpResult](#) or [TxpResultList](#) object

See Also

[TxpModel](#), [TxpResult](#)

Examples

```
## Load example dataset & model; see ?TxpModel for building model objects
data(txp_example_input, package = "toxiR")
data(txp_example_model, package = "toxiR")

## Calculate scores for single model; returns TxpResult object
res <- txpCalculateScores(model = txp_example_model,
                          input = txp_example_input,
                          id.var = "name")

## Calculate scores for list of models; returns TxpResultList object
txpCalculateScores(model = TxpModelList(m1 = txp_example_model,
                                       m2 = txp_example_model),
                  input = txp_example_input,
                  id.var = "name")
resLst <- txpCalculateScores(model = list(m1 = txp_example_model,
                                       m2 = txp_example_model),
                            input = txp_example_input,
                            id.var = "name")
```

txpGenerics

toxiR package generics

Description

toxiR package generics; see class man pages for associated methods

Usage

```
txpValueNames(x, ...)
```

```
txpValueNames(x, ...) <- value
```



```
txpTransFuncs(x, ...)  
txpTransFuncs(x, ...) <- value  
txpSlices(x, ...)  
txpSlices(x, ...) <- value  
txpWeights(x, ...)  
txpWeights(x, ...) <- value  
txpScores(x, ...)  
txpSliceScores(x, ...)  
txpModel(x, ...)  
txpIDs(x, ...)  
txpIDs(x, ...) <- value  
txpRanks(x, ...)
```

Arguments

x	toxiR S4 object
...	Included for extensibility; not currently used
value	Replacement value

Value

See specific methods for details.

txpImportGui	<i>Import data file generated by ToxPi GUI</i>
--------------	--

Description

Import data file generated by ToxPi GUI

Usage

```
txpImportGui(guiDataFile)
```

Arguments

guiDataFile	Character scalar, the path to a 'data' export from the ToxPi GUI
-------------	--

Details

This function takes the '_data.csv' files generated by the GUI. See <https://toxpi.org> for more information.

Because of the way toxpiR implements transformation functions, there is not a way currently to use the GUI 'hitcount' function.

Value

list with \$model containing [TxpModel](#) object; \$input containing data.frame with input data; \$fills containing a vector of fill colors.

TxpModel-class	<i>ToxPi Model</i>
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Description

S4 class to store ToxPi models

Usage

```
TxpModel(txpSlices, txpWeights = NULL, txpTransFuncs = NULL)
```

```
## S4 method for signature 'TxpModel'
txpSlices(x)
```

```
## S4 replacement method for signature 'TxpModel'
txpSlices(x) <- value
```

```
## S4 method for signature 'TxpModel'
txpWeights(x, adjusted = FALSE)
```

```
## S4 replacement method for signature 'TxpModel'
txpWeights(x) <- value
```

```
## S4 method for signature 'TxpModel'
txpTransFuncs(x)
```

```
## S4 replacement method for signature 'TxpModel'
txpTransFuncs(x) <- value
```

```
## S4 method for signature 'TxpModel'
names(x)
```

```
## S4 replacement method for signature 'TxpModel'
names(x) <- value
```

```
## S4 method for signature 'TxpModel'
length(x)

## S4 method for signature 'TxpModel,TxpModel'
merge(x, y)
```

Arguments

txpSlices	Passed to txpSlices slot
txpWeights	Passed to txpWeights slot
txpTransFuncs	Passed to txpTransFuncs slot
x, y	TxpModel object
value	Replacement value
adjusted	Scalar logical, should the returned weights be adjusted such that they sum to 1?

Functions

- txpSlices, TxpModel-method: Return txpSlices slot
- txpWeights, TxpModel-method: Return txpWeights slot
- txpTransFuncs, TxpModel-method: Return txpTransFuncs slot
- names, TxpModel-method: Return slice names; shortcut for names(txpSlices(x))
- length, TxpModel-method: Return number of slices in model; shortcut for length(txpSlices(x))
- merge, TxpModel, TxpModel-method: Merge two TxpModel objects into a single model

Slots

txpSlices [TxpSliceList](#) object

txpWeights numeric vector specifying the relative weight of each slice; when NULL, defaults to 1 (equal weighting) for each slice

txpTransFuncs [TxpTransFuncList](#) object (or list of functions coercible to TxpTransFuncList)

Examples

```
## Create TxpSliceList & TxpTransFuncList objects
s1 <- list(S1 = TxpSlice("inpt1"), S2 = TxpSlice("inpt2"))
tf <- list(NULL, sqrt = function(x) sqrt(x))

## Create TxpModel object
m1 <- TxpModel(txpSlices = s1, txpWeights = 2:1, txpTransFuncs = tf)
m1

## Access TxpModel slots
txpSlices(m1)
txpWeights(m1)
txpWeights(m1, adjusted = TRUE)
txpTransFuncs(m1)
```

```

## length
length(m1) ## equal to length(txpSlices(m1))
length(m1) == length(txpSlices(m1))

## names
names(m1) ## equal to names(txpSlices(m1))
all(names(m1) == names(txpSlices(m1)))

## Replacement
m2 <- m1
txpSlices(m2) <- list(S3 = TxpSlice("inpt3"), S4 = TxpSlice("inpt4"))
m2
names(m2)[2] <- "hello"
names(m2)
txpTransFuncs(m2) <- NULL
m2
txpTransFuncs(m2)[[1]] <- function(x) x^2
names(txpTransFuncs(m2))[1] <- "sq"
m2

## merge
m3 <- merge(m1, m2)
m3

```

TxpModelList-class *List of TxpModel objects*

Description

Extension of [S4Vectors::SimpleList](#) that holds only [TxpModel](#) objects.

Usage

```

TxpModelList(...)

## S4 method for signature 'TxpModelList'
duplicated(x)

as.TxpModelList(x)

```

Arguments

...	TxpModel object to create TxpModelList object
x	TxpModelList object

Functions

- `duplicated, TxpModelList-method`: Returns logical vector of `length(x)`, where TRUE indicates a duplicate model in the list; see [base::duplicated](#)
- `as.TxpModelList`: Coerce list or [TxpModel](#) objects to TxpModelList

Examples

```
## Create some TxpModel objects; see ?TxpModel for more details
s1 <- list(S1 = TxpSlice("inpt1"), S2 = TxpSlice("inpt2"))
tf <- list(NULL, sqrt = function(x) sqrt(x))
m1 <- TxpModel(txpSlices = s1, txpWeights = 2:1, txpTransFuncs = tf)
m2 <- m1
txpSlices(m2) <- list(S3 = TxpSlice("inpt3"), S4 = TxpSlice("inpt4"))
m3 <- merge(m1, m2)

## Build a TxpModelList object
TxpModelList(m1 = m1, m2 = m2, m3 = m3)

## Note: names are printed as '' when all are NULL
TxpModelList(m1, m2, m3)
names(TxpModelList(m1, m2, m3))

## Test for duplicates
duplicated(TxpModelList(m1 = m1, m2 = m2, m3 = m3))
duplicated(TxpModelList(m1 = m1, m2 = m1, m3 = m3))

## Coerce lists/TxpModel objects to TxpModelList
as(list(m1 = m1, m2 = m2, m3 = m3), "TxpModelList")
as.TxpModelList(list(m1 = m1, m2 = m2, m3 = m3))

as(m1, "TxpModelList")
as.TxpModelList(m1)
```

TxpResult-class

*ToxPi Result***Description**

S4 class to store ToxPi results

Usage

```
## S4 method for signature 'TxpResult'
txpScores(x)

## S4 method for signature 'TxpResult'
txpSliceScores(x, adjusted = TRUE)

## S4 method for signature 'TxpResult'
txpRanks(x)

## S4 method for signature 'TxpResult'
txpModel(x)

## S4 method for signature 'TxpResult'
```

```
txpIDs(x)

## S4 replacement method for signature 'TxpResult'
txpIDs(x) <- value

## S4 method for signature 'TxpResult'
txpWeights(x, adjusted = FALSE)

## S4 method for signature 'TxpResult'
txpSlices(x)

## S4 method for signature 'TxpResult'
txpTransFuncs(x, level, simplify = FALSE)

## S4 method for signature 'TxpResult'
txpValueNames(x, simplify = FALSE)

## S4 method for signature 'TxpResult,logical,missing'
x[i, j, ..., drop = FALSE]

## S4 method for signature 'TxpResult,integer,missing'
x[i, j, ..., drop = FALSE]

## S4 method for signature 'TxpResult,numeric,missing'
x[i, j, ..., drop = FALSE]

## S4 method for signature 'TxpResult,character,missing'
x[i, j, ..., drop = FALSE]

## S4 method for signature 'TxpResult'
length(x)

## S4 method for signature 'TxpResult'
sort(x, decreasing = TRUE, na.last = TRUE, ...)

## S4 method for signature 'TxpResult'
names(x)

## S4 replacement method for signature 'TxpResult'
names(x) <- value

## S4 method for signature 'TxpResult'
as.data.frame(
  x,
  row.names = NULL,
  optional = FALSE,
  ...,
  id.name = "id",
```

```

    score.name = "score",
    rank.name = "rank",
    adjusted = FALSE
  )

```

Arguments

x	TxpResult object
adjusted	Logical scalar, when TRUE the weights are adjusted to sum to 1 or the slice scores are scaled to their respective weight
value	Replacement value
level	c('model', 'slices'); indicates whether to retrieve txpTransFuncs slot from the model or underlying slices
simplify	Logical scalar, flatten txpValueNames or txpTransFunc slots when retrieving slice-level information
i	Subsetting index
j, drop, optional	Not currently implemented
...	Passed to base::data.frame in <code>as.data.frame</code> or base::sort in <code>sort</code>
decreasing, na.last	Passed to base::sort
row.names	Passed to base::data.frame
id.name, score.name, rank.name	Character scalar; when coercing to base::data.frame , the name for the txpIDs, txpScores, and txpRanks columns, respectively

Functions

- `txpScores`, `TxpResult`-method: Return `txpScores` slot
- `txpSliceScores`, `TxpResult`-method: Return `txpSliceScores` slot; default `adjusted = TRUE`, i.e. return slice scores adjusted for weight
- `txpRanks`, `TxpResult`-method: Return `txpRanks` slot
- `txpModel`, `TxpResult`-method: Return `txpModel` slot
- `txpIDs`, `TxpResult`-method: Return `txpIDs` slot
- `txpWeights`, `TxpResult`-method: Return `txpWeights` slot from model – shortcut for `txpWeights(txpModel(x))`; default `adjusted = FALSE`, i.e. return unadjusted weights
- `txpSlices`, `TxpResult`-method: Return `txpSlices` slot from model – shortcut for `txpSlices(txpModel(x))`
- `txpTransFuncs`, `TxpResult`-method: Return `txpTransFuncs` slot from model – shortcut for `txpTransFuncs(txpModel(x))`
- `txpValueNames`, `TxpResult`-method: Return `txpValueNames` slot from slices – shortcut for `txpValueNames(txpSlices(txpModel(x)))`
- `length`, `TxpResult`-method: Return the number of observations; shortcut for `length(txpScores(x))`
- `sort`, `TxpResult`-method: Sort the “TxpResult” object by their ranks
- `names`, `TxpResult`-method: Returns IDs; equal to `txpIDs(x)`
- `as.data.frame`, `TxpResult`-method: Coerce `TxpResult` to [base::data.frame](#) object with IDs, scores, ranks, and slice scores

Slots

txpScores vector(<numeric>) of model scores
 txpSliceScores matrix(<numeric>), sample by slice matrix with individual slice scores
 txpRanks vector(<numeric>) with rank of scores
 txpModel [TxpModel](#) object
 txpIDs vector(<character>) of observation IDs

See Also

[txpCalculateScores](#), [plot](#), [TxpResultList](#)

Examples

```
## Load example dataset & model; see ?TxpModel for building model objects
data(txp_example_input, package = "toxiR")
data(txp_example_model, package = "toxiR")

## Calculate scores for single model; returns TxpResult object
res <- txpCalculateScores(model = txp_example_model,
  input = txp_example_input,
  id.var = "name")

## Accessors
txpScores(res)

txpSliceScores(res) ## adjusted for weight, by default
apply(txpSliceScores(res), 2, max, na.rm = TRUE)

txpSliceScores(res, adjusted = FALSE) ## each score should have maximum of 1
apply(txpSliceScores(res, adjusted = FALSE), 2, max, na.rm = TRUE)

txpRanks(res)

txpModel(res)
identical(txpModel(res), txp_example_model)

txpIDs(res)
names(res) ## identical to txpIDs(res)
identical(txpIDs(res), names(res))

# Can access TxpModel slots directly
txpWeights(res)
txpWeights(res, adjusted = TRUE)
txpSlices(res)
# When retrieving transform functions, must specify level because both
# models and slices have transform functions
txpTransFuncs(res, level = "model")

# Can access TxpSliceList slots directly
txpValueNames(res)
txpValueNames(res, simplify = TRUE)
```



```
txpTransFuncs(res, level = "slices")
txpTransFuncs(res, level = "slices", simplify = TRUE)

## Subsetting
res[1]
res[c("chem01", "chem09")]
res[grepl("4|6", txpIDs(res))]
## Not run:
res[c(TRUE, FALSE)] ## gets recycled with warning

## End(Not run)

## length -- returns number of observations
length(res)
length(res[1:5])

## sort
names(res)
names(sort(res))

txpScores(res)
txpScores(sort(res))
txpScores(sort(res, decreasing = FALSE))

## as.data.frame
as.data.frame(res)
as.data.frame(res, id.name = "nm", score.name = "scr", rank.name = "rnk")
```

TxpResult-plot

Plot TxpResult objects

Description

Plot [TxpResult](#) objects

Usage

```
## S4 method for signature 'TxpResult,missing'
plot(
  x,
  fills = NULL,
  showScore = TRUE,
  gp = NULL,
  vp = NULL,
  name = NULL,
  newpage = TRUE,
  ...
)
```

```
## S4 method for signature 'TxpResult,numeric'
plot(
  x,
  y,
  labels = NULL,
  newpage = TRUE,
  margins = c(4, 0, 1, 1),
  name = NULL,
  gp = NULL,
  vp = NULL,
  ...
)
```

Arguments

x	TxpResult object
fills	Vector of colors to fill slices
showScore	Logical scalar, overall score printed below the name when TRUE
gp, vp, name	Passed to grid::frameGrob when creating the plotting area
newpage	Logical scalar, grid::grid.newpage called prior to plotting when TRUE
...	Passed to pieGridGrob when plotting ToxPi and to pointsGrob when plotting ranks
y	Rank vector, i.e. <code>txpRanks(x)</code>
labels	Integer vector, indices of x to label in the rank plot
margins	Passed to grid::plotViewport ; only affects the scatterplot region margins

Details

It is strongly recommended to use a specific device (e.g., [grDevices::png](#), [grDevices::pdf](#)) when creating rank plots. Using a GUI device will likely lead to inaccurate labeling, and any changes to the device size WILL lead to inaccurate labeling.

The plotting is built on the [grid::grid-package](#), and can be adjusted or edited as such.

If the labels are running of the device, the top or bottom margins can be increased with the `margins` parameter.

Value

No return value; called for side effect (i.e. drawing in current graphics device.)

Functions

- `plot, TxpResult, missing-method`: Plot ToxPi diagrams
- `plot, TxpResult, numeric-method`: Plot ToxPi ranks

Examples

```
## Load example dataset & model; see ?TxpModel for building model objects
data(txp_example_input, package = "toxiR")
data(txp_example_model, package = "toxiR")

## Calculate scores for single model; returns TxpResult object
res <- txpCalculateScores(model = txp_example_model,
                          input = txp_example_input,
                          id.var = "name")

library(grid)
plot(res)

plot(res, txpRanks(res))
plot(res, txpRanks(res), pch = 16, size = unit(0.75, "char"))

## Will likely make inaccurate labels within a GUI, e.g. RStudio
## use png, pdf, etc. to get accurate labels
## Not run:
tmpPdf <- tempfile()
pdf(tmpPdf)
plot(res, txpRanks(res), labels = c(10, 4, 2), pch = 16)
dev.off()

## End(Not run)
```

TxpResultList-class *List of TxpResult objects*

Description

Extension of [S4Vectors::SimpleList](#) that holds only [TxpResult](#) objects.

Usage

```
TxpResultList(...)

## S4 method for signature 'TxpResultList'
duplicated(x)

as.TxpResultList(x)
```

Arguments

```
...                    TxpResult object to create TxpResultList object
x                      TxpResultList object
```

See Also

[TxpResult](#), [txpCalculateScores](#)

Examples

```
## Load example dataset & model; see ?TxpModel for building model objects
data(txp_example_input, package = "toxiR")
data(txp_example_model, package = "toxiR")

## Calculate scores for list of models; returns TxpResultList object
txpCalculateScores(model = TxpModelList(m1 = txp_example_model,
                                         m2 = txp_example_model),
                  input = txp_example_input,
                  id.var = "name")
resLst <- txpCalculateScores(model = list(m1 = txp_example_model,
                                         m2 = txp_example_model),
                           input = txp_example_input,
                           id.var = "name")

## duplicated
duplicated(resLst)

## Coercion
as(list(resLst[[1]], resLst[[2]]), "TxpResultList")
as.TxpResultList(list(res1 = resLst[[1]], res2 = resLst[[2]]))

as(resLst[[1]], "TxpResultList")
as.TxpResultList(resLst[[1]])
```

TxpSlice-class

ToxPi Slice

Description

S4 class to store ToxPi slices

Usage

```
TxpSlice(txpValueNames, txpTransFuncs = NULL)

## S4 method for signature 'TxpSlice'
txpValueNames(x)

## S4 replacement method for signature 'TxpSlice'
txpValueNames(x) <- value

## S4 method for signature 'TxpSlice'
txpTransFuncs(x)
```

```
## S4 replacement method for signature 'TxpSlice'
txpTransFuncs(x) <- value

## S4 method for signature 'TxpSlice'
length(x)

## S4 method for signature 'TxpSlice,TxpSlice'
merge(x, y)
```

Arguments

txpValueNames	Passed to txpValueNames slot
txpTransFuncs	Passed to txpTransFuncs slot
x, y	TxpSlice object
value	Replacement value

Details

If the user supplies txpTransFuncs a single function/[TxpTransFunc](#) object, the given function will be recycled for each input with a warning.

Functions

- txpValueNames, TxpSlice-method: Return txpValueNames slot
- txpTransFuncs, TxpSlice-method: Return txpTransFuncs slot
- length, TxpSlice-method: Return number of inputs in slice; shortcut for length(txpValueNames(x))
- merge, TxpSlice, TxpSlice-method: Merge two TxpSlice objects into a single slice

Slots

txpValueNames vector(<character>) specifying the input columns to include in the slice
txpTransFuncs [TxpTransFuncList](#) with one function per entry in txpValueNames or an object that can be coerced to TxpTransFuncList; when NULL, no transformation function applied

Examples

```
## Create TxpSlice object
# Without transform functions
TxpSlice(txpValueNames = c("sqrData", "expData"))
# With transform functions
TxpSlice(txpValueNames = c("sqrData", "expData"),
         txpTransFuncs = c(sq = function(x) x^2, log = function(x) log(x)))

# Transformation function recycled with warning when single function given
TxpSlice(txpValueNames = c("sqrData", "expData"),
         txpTransFuncs = function(x) x^2)

## Access TxpSlice slots
```

```

s1 <- TxpSlice(txpValueNames = c("sqrData", "expData"),
              txpTransFuncs = c(sq = function(x) x^2,
                                log = function(x) log(x)))

txpValueNames(s1)
txpTransFuncs(s1)

## Replacement
txpValueNames(s1)[1] <- "hello"
s1

txpTransFuncs(s1)[[2]](exp(1))
txpTransFuncs(s1)[[2]] <- function(x) sqrt(x)
txpTransFuncs(s1)[[2]](exp(1))

# Note that replacing a single list element does NOT update the name
s1
names(txpTransFuncs(s1))[2] <- "sqrt"
s1

# Replacing the whole list DOES update the names
txpTransFuncs(s1) <- list(sqrt = function(x) sqrt(x),
                          log = function(x) log(x))
s1

## length -- returns number of inputs
length(TxpSlice(letters))

## merge
s1 <- TxpSlice("hello")
s2 <- TxpSlice("data")
merge(s1, s2)

# Note, input names still must be unique
## Not run: merge(s1, s1) ## produces error

```

TxpSliceList-class *List of TxpSlice objects*

Description

Extension of [S4Vectors::SimpleList](#) that requires uniquely-named elements and holds only [TxpSlice](#) objects.

Usage

```

TxpSliceList(...)

## S4 method for signature 'TxpSliceList'
txpValueNames(x, simplify = FALSE)

```

```
## S4 method for signature 'TxpSliceList'
txpTransFuncs(x, simplify = FALSE)

## S4 method for signature 'TxpSliceList'
duplicated(x)

as.TxpSliceList(x)
```

Arguments

...	TxpSlice object to create TxpSlicelist object; MUST give unique names to each slice
x	TxpSliceList object
simplify	Scalar logical, when TRUE the returned list is simplified to a vector/ TxpTransFuncList object

Details

Note, there is no coercion for [TxpSlice](#) to TxpSliceList because unique names are required.

Functions

- txpValueNames, TxpSlicelist-method: Return list of txpValueNames slots for the contained [TxpSlice](#) objects, or vector when simplify = TRUE
- txpTransFuncs, TxpSlicelist-method: Return list of txpTransFuncs slots for the contained [TxpSlice](#) objects, or [TxpTransFuncList](#) when simplify = TRUE
- duplicated, TxpSliceList-method: Returns logical vector of length(x), where TRUE indicates a duplicate slice in the list; see [base::duplicated](#)

Examples

```
## Create TxpSlice objects
s1 <- TxpSlice("input1", list(linear = function(x) x))
s2 <- TxpSlice(c("input2", "input3"),
              list(log = function(x) log(x), sqrt = function(x) sqrt(x)))

## Create TxpSliceList
s1 <- TxpSliceList(s1 = s1, s2 = s2)

## Accessors
txpValueNames(s1)
txpValueNames(s1, simplify = TRUE)

txpTransFuncs(s1)
txpTransFuncs(s1, simplify = TRUE)

## Coercion
as(list(s1 = TxpSlice("hello"), s2 = TxpSlice("user")), "TxpSliceList")
as.TxpSliceList(c(s1 = TxpSlice("hello"), s2 = TxpSlice("user")))
```

```
## Concatenation
c(s1, TxpSliceList(s3 = TxpSlice("input4")))

## Reduce TxpSliceList to single slice
Reduce(merge, s1)
```

TxpTransFunc-class *Numeric transformation function*

Description

S4 class to store numeric transformation functions

Usage

```
TxpTransFunc(x)
```

Arguments

x function, see details

Details

TxpTransFunc inherits from a standard R function, but specifies a single input and a numeric output of the same length.

Functions can be passed directly to TxpTransFuncList list and the functions will be coerced to TxpTransFunc.

We have an imperfect system for dealing with primitive functions (e.g., `base::sqrt`). To coerce primitives to TxpTransFunc's, we wrap them in another function `cal`; wrapping the primitives obscures the original function and requires the user to explore the function environment to understand the primitive called. We recommend wrapping primitives in separate functions to make the intent clear, e.g., `mysqrt <- function(x) sqrt(x)`.

Examples

```
f1 <- function(x) "hello"
f2 <- function(x) 3
f3 <- function(x) x + 5
## Not run:
t1 <- TxpTransFunc(x = f1) ## Produces error
t2 <- TxpTransFunc(x = f2) ## Produces error

## End(Not run)
t3 <- TxpTransFunc(x = f3)

## TxpTransFunc objects act as any other function
body(t3)
formals(t3)
```



```

t3(1:10)

## Coercion from functions
## Not run:
TxpTransFuncList(f1, f2, f3) ## Produces error because f1, f3 not valid

## End(Not run)

```

TxpTransFuncList-class

List of TxpTransFunc objects

Description

Extension of [S4Vectors::SimpleList](#) that holds only NULL or [TxpTransFunc](#) objects.

Usage

```

TxpTransFuncList(...)

as.TxpTransFuncList(x)

```

Arguments

...	TxpTransFunc object or function to create TxpTransFuncList object
x	list, function, or TxpTransFunc object to coerce to TxpTransFuncList

Details

When ... includes function objects, [TxpTransFuncList](#) will attempt to coerce them to [TxpTransFunc](#) and return an error if any of the elements cannot be coerced to [TxpTransFunc](#).

Examples

```

## Create TxpTransFunc objects
tf1 <- TxpTransFunc(function(x) x)
tf2 <- TxpTransFunc(function(x) sqrt(x))

## Create TxpTransFuncList
tf1 <- TxpTransFuncList(linear = tf1, sqrt = tf2, cube = function(x) x^3)
tf1[[3]](3) == 27
tf1[["sqrt"]](4) == 2

## Concatenate
c(tf1, tf1)

## names
names(c(tf1, tf1))

```

```
# note: names are printed as '' when missing; NULL is printed when list item
# is NULL
names(TxpTransFuncList(function(x) x, NULL))
TxpTransFuncList(function(x) x, NULL)

## coercion
as(function(x) x, "TxpTransFuncList")
as.TxpTransFuncList(function(x) x)

as(TxpTransFunc(function(x) x), "TxpTransFuncList")
as.TxpTransFuncList(TxpTransFunc(function(x) x))

as(list(function(x) x, sqrt = function(x) sqrt(x)), "TxpTransFuncList")
as.TxpTransFuncList(list(function(x) x, sqrt = function(x) sqrt(x)))
```

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