Package 'gDRtestData'

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Title gDRtestData - R data package with testing dose reponse data

Version 1.0.0

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Description R package with internal dose-response test data. Package provides functions to generate input testing data that can be used as the input for gDR pipeline. It also contains RDS files with MAE data processed by gDR.

Depends R (>= 4.2)

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R topics documented:

`gDRtestData-package` *gDRtestData: gDRtestData - R data package with testing dose reponse data*

Description

R package with internal dose-response test data. Package provides functions to generate input testing data that can be used as the input for gDR pipeline. It also contains RDS files with MAE data processed by gDR.

Value

package help page

Note

To learn more about functions start with `help(package = "gDRtestData")`

Author(s)

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- Dariusz Scigocki
- Allison Voung

`add_concentration` *Add concentrations*

Description

Add concentrations

Usage

```
add_concentration(df_layout, concentrations = 10^(seq(-3, 1, 0.5)))
```

Arguments

`df_layout`	data.table that should contains the cell line, drug, concentration, and replicate columns along with the annotations that needs to be propagated
`concentrations`	vector of numeric concentrations that will be added to df_layout

Value

data.table with concentrations

Examples

```
cell_lines <- create_synthetic_cell_lines()
add_concentration(cell_lines)
```

add_data_replicates *Add data replicates*

Description

Add data replicates

Usage

```
add_data_replicates(df_layout)
```

Arguments

df_layout	data.table that should contains the cell line, drug, concentration, and replicate columns along with the annotations that needs to be propagated

Value

data.table with replicates

Examples

```
cell_lines <- create_synthetic_cell_lines()
add_data_replicates(cell_lines)
```

add_day0_data *Add data with day 0*

Description

Add data with day 0

Usage

```
add_day0_data(df_merged, noise_level = 0.05)
```

Arguments

df_merged	data.table with merged data
noise_level	numeric scalar with the level of noise added to the data

Value

data.table with day0 data

Examples

```
cell_lines <- create_synthetic_cell_lines()
drugs <- create_synthetic_drugs()
data <- gDRtestData:::prepareData(cell_lines[seq_len(2), ], drugs[seq_len(4), ])
data$Duration <- 72
data$ReadoutValue <- 0
add_day0_data(data)
```

---

cell_lines *Cell lines*

Description

Cell lines

Value

data.table

Examples

```
path <- system.file("annotation_data", "cell_lines.csv", package = "gDRtestData")
data.table::fread(file = path)
```

---

create_synthetic_cell_lines *Create data.table with synthetic cell lines*

Description

Create data.table with synthetic cell lines

Usage

```
create_synthetic_cell_lines()
```

Value

data.table with synthetic cell lines

Examples

```
create_synthetic_cell_lines()
```

create_synthetic_drugs *Create data.table with synthetic drugs*

Description

Create data.table with synthetic drugs

Usage

```
create_synthetic_drugs()
```

Value

data.table with synthetic drugs

Examples

```
create_synthetic_drugs()
```

drugs *Drugs*

Description

Drugs

Value

data.table

Examples

```
path <- system.file("annotation_data", "drugs.csv", package = "gDRtestData")
data.table::fread(file = path)
```

generate_ec50 *Calculate EC50 metric*

Description

Calculate EC50 metric

Usage

```
generate_ec50(drugs, cell_lines)
```

Arguments

drugs	data.table with drugs
cell_lines	data.table with cell lines

Value

matrix with random EC50

Examples

```
generate_ec50(create_synthetic_drugs(), create_synthetic_cell_lines())
```

generate_e_inf *Calculate E inf metric*

Description

Calculate E inf metric

Usage

```
generate_e_inf(drugs, cell_lines)
```

Arguments

drugs	data.table with drugs
cell_lines	data.table with cell lines

Value

matrix with random E inf

Examples

```
generate_e_inf(create_synthetic_drugs(), create_synthetic_cell_lines())
```

`generate_hill_coef` *Generate hill coefficient*

Description

Generate hill coefficient

Usage

```
generate_hill_coef(drugs, cell_lines)
```

Arguments

`drugs`	data.table with drugs
`cell_lines`	data.table with cell lines

Value

matrix with random hill coefficient

Examples

```
generate_hill_coef(create_synthetic_drugs(), create_synthetic_cell_lines())
```

`generate_response_data`
Generate response data

Description

Generate response data

Usage

```
generate_response_data(df_layout, noise_level = 0.1)
```

Arguments

`df_layout`	data.table that should contains the cell line, drug, concentration, and replicate columns along with the annotations that needs to be propagated
`noise_level`	numeric scalar with the level of noise added to the data

Value

data.table with response data

Examples

```
cell_lines <- create_synthetic_cell_lines()
drugs <- create_synthetic_drugs()
gDRtestData:::prepareData(cell_lines[seq_len(2), ], drugs[seq_len(4), ])
```

`get_test_dataset_paths` *get_test_dataset_paths*

Description

Returns named vector of absolute paths to test datasets.

Usage

```
get_test_dataset_paths(datasets_dir = NULL, pattern = "finalMAE_")
```

Arguments

`datasets_dir`	path to directory with datasets (default `NULL`). If `NULL`, then `inst/testdata` directory from `gDRtestData` will be used.
`pattern`	used to: (1) filter to qs files from the dataset_dir path and (2) prettify the labels of the files

Value

named vector of absolute paths

Author(s)

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Examples

```
get_test_dataset_paths()
path <- system.file("testdata", package = "gDRtestData", mustWork = TRUE)
get_test_dataset_paths(path)
```

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