

Package ‘KOMA’

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Type Package

Title Kinetic Operating Microarray Analyzer (KOMA)

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Description Kinetic Operating Microarray Analyzer (KOMA) enables calibration and high-throughput analysis of quantitative microarray data collected by using kinetic detection protocol. This tool can be also helpful for analyzing data from any other analytical assays employing enzymatic signal amplification, in which a broader range of quantification is reached by the time-resolved recording of readouts.

License GPL-2

LazyLoad yes

Depends epicalc,fields,gplots,magic,rgl,tcltk2,tkrplot,R (>= 2.12)

OS_type windows

References James Wettenhall, Philippe Grosjean 2005 Felix Bonowski 2010 Pavlo Holenya 2011

R topics documented:

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analyte.choice	<i>Choosing an analyte</i>
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Description

This function opens a window showing all accessible analyte names of the specified array. Here the one to analyse or calibrate should be chosen.

Usage

```
analyte.choice(analyte.names)
```

Arguments

analyte.names	names of all the defined analytes
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Value

analyte	the name of the chosen analyte as character string
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Author(s)

Florian Heigwer

calculate.plot.mean.and.error	<i>Calculate and plot the mean and error surfaces</i>
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Description

This function gives the possibility to evaluate the goodness of calibration from the several independent replicates.

Usage

```
calculate.plot.mean.and.error(workpath, datasets.to.leeve.out)
```

Arguments

workpath	the specified working directory from the maindirectory
datasets.to.leeve.out	vector of dataset IDs to exclude, passed by from the function plotting.overlapping.cals

Note

please close the windows after analysis

Author(s)

Florian Heigwer

See Also

[plotting.overlapping.cals](#)

calibrationsettings

Calibration settings

Description

This function opens a panel to specify settings for the calibration analysis.

Usage

```
calibrationsettings()
```

Details

This function will then automatically run the subfunctions `timecourse` and `dilutionseries`.

Value

<code>times</code>	a vector of time points in the given interval
<code>concentrations</code>	a vector of concentrations

Author(s)

Florian Heigwer

See Also

[timecourse](#) [dilutionseries](#)

```
collect.and.save.cal
```

Save the analysed calibration run

Description

This functions generates a calibration file for the specified calibration run.

Usage

```
collect.and.save.cal(data, chosen.IDs, substance, run)
```

Arguments

<code>data</code>	a list of tables with confidence and signal values for each spotreplicate, timepoint and concentration
<code>chosen.IDs</code>	a vector of chosen IDs
<code>substance</code>	the analytes name
<code>run</code>	the name of the calibration run to name the files subsequently

Note

The file will be saved under the analytes name, calibration, and the run ID.

Author(s)

Florian Heigwer

See Also

[calculate.plot.mean.and.error](#)

```
dilutionseries
```

Calculate the dilutionseries

Description

This function calculates a series of concentrations depending on the highest concentration, the number of dilutionsteps and the diltionfactor.

Usage

```
dilutionseries(highestconc, dilutions, dilutionfactor)
```

Arguments

<code>highestconc</code>	the highest concentration as numeric object
<code>dilutions</code>	the number of diltionsteps performed in the experiment
<code>dilutionfactor</code>	the used factor of dilution

Value

concentrations

a series of ideal calculated concentrations used to calibrate

Author(s)

Florian Heigwer

Examples

```
dilutionseries(10,8,2) #give the highest concentration, the number of dilutions and the c
```

```
extractconfidence_strips
```

Extract confidence values

Description

This function parses all the folders of an experiment for the result.txt tables and sorts or reformats them.

Usage

```
extractconfidence_strips(workpath, substance, run, times, concentrations)
```

Arguments

workpath	the directory inheriting data to analyse
substance	the analytes name chosen by analyte.choice
run	the name of the certain run
times	timeseries as given from timecourse
concentrations	series of concentration as given from dilutionseries

Value

confidencedata

a list consisting of the sortet and formatted confidencedata, plus the timecourse and the dilutionseries

Author(s)

Florian Heigwer

See Also

[extractdata_strips](#)

`extractdata_strips` *Extract experimental data from folder with rawdata*

Description

This function parses all the folders of an experiment for the result.txt tables and sorts or reformats them.

Usage

```
extractdata_strips(workpath, substance, run, times, concentrations)
```

Arguments

<code>workpath</code>	the directory inheriting data to analyse
<code>substance</code>	the analytes name chosen by <code>analyte.choice</code>
<code>run</code>	the name of the certain run
<code>times</code>	timeseries as given from <code>timecourse</code>
<code>concentrations</code>	series of concentration as given from <code>dilutionseries</code>

Value

<code>signaldata</code>	a list consisting of the sortet and formatted signaldata, plus the timecourse and the dilutionseries
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Author(s)

Florian Heigwer

See Also

`extractconfidence_strips`

`extractdata_unknown_predict_conc`
Measure an concentration

Description

This function provides a workflow for calculation of an unknown concentration by using the analytes forprediction.RData file.

Usage

```
extractdata_unknown_predict_conc(workpath, substance)
```

Arguments

workpath	the directory containing the rawdata of the measurement
substance	the analytes name

Details

first it extracts the data second it plots the replicates and lets you choose excludable ones third it will compare the meaned data set with the topographic calibration and give back the estimated analyte concentration

Value

analyteconcentration
an estimate of the measured concentration

Author(s)

Florian Heigwer

See Also

[extractdata_unknown_predict_conc_wholestrip](#)

extractdata_unknown_predict_conc_wholestrip
Measure eigh folders at once

Description

This function provides a workflow for calculation of an unknown concentration by using the analytess forprediction file.

Usage

```
extractdata_unknown_predict_conc_wholestrip(foldernames, paths, substance, workpath)
```

Arguments

foldernames	the names of the single folders
paths	the directory containing the rawdata of the measurement
substance	the analytes name
workpath	the main working directory

Details

first it extracts the data, second it plots the replicates and lets you choose ecludable ones, third it will compare the meaned data set with the topographic calibration and give back the estimated analyte concentration

Value

`analyteconcentration`
an estimate of the measured concentration

Author(s)

Florian Heigwer

See Also

[extractdata_unknown_predict_conc](#)

GUI

Build the GUI

Description

This function generates a whole functioning graphical user interface.

Usage

`GUI ()`

Details

The user interface called KOMA consists of a mainmenu containing all necessary submenus for controlling the data analysis and a tabulary body for showing experimental results.

Author(s)

Florian Heigwer

References

http://www.sciviews.org/_rgui/tcltk/Tktable.html

See Also

[Separation](#)

Examples

`GUI ()`

`list.dir`*List Directories*

Description

This function creates a lists of directories and only directories in a certain path.

Usage

```
list.dir(path)
```

Arguments

<code>path</code>	the path to search the directories in
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Value

<code>Q</code>	the names of the found directories
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Author(s)

Florian Heigwer

Examples

```
list.dir(path=R.home())
```

`numeric.entry.window`*Open an interactive window asking for numeric input*

Description

This function opens an interactive window asking for numeric input.

Usage

```
numeric.entry.window(title, question, entryInit, entryWidth = 20, returnValOnCancel)
```

Arguments

<code>title</code>	the title of the window as character string
<code>question</code>	the question to ask in the window
<code>entryInit</code>	the default text entry when opening the window
<code>entryWidth</code>	the width of the textinput area
<code>returnValOnCancel</code>	what should be printed when Cancel has been pushed

Value

RetVal the typed character string

Author(s)

Florian Heigwer

References

James Wettenhall, Philippe Grosjean (2005)

See Also

[numeric.entry.window](#)

plotting.overlapping.cals

Plot replicates of several calibration runs.

Description

This function extracts datasets of different calibration replicates from files in the main working directory. These files are searched by regular expression matching with the analytes name and the word calibration.

Usage

```
plotting.overlapping.cals(workpath)
```

Arguments

workpath defines the directory to search in

Value

IDs.to.leeve.out
a list of dataset IDs which should be excluded from all further analysis

Note

At least two datasets has to be left over.If more should be left for analysis please stop the function by rightclick-> Stop.

Author(s)

Florian Heigwer

See Also

[collect.and.save.cal](#)

```
plotting.timecourses.and.choose.IDs.manually
```

Plot timecourses of signal and confidence values and choose IDs to exclude manually

Description

This function will open an tcltk interface panel which shows signal and confidence values time-resolved for all spot replicates. Giving as well a chance to choose failed spots for exclusion from further analysis.

Usage

```
plotting.timecourses.and.choose.IDs.manually(data, substance, confidencevalues,
```

Arguments

data	datasets as a list of matrices for each replicate, for each timepoint and eventually each concentration while calibration containing signal values
substance	the analytename as character string
confidencevalues	datasets as a list of matrices for each replicate, for each timepoint and eventually each concentration while calibration containing confidence values
workpath	the directory with the measured rawdata
foldernames	the single foldernames (each representing one well of an ArrayStrip)

Value

```
IDs.to.left.out
```

a vector of spot IDs to be exclude from further analysis

Author(s)

Florian Heigwer

See Also

[collect.and.save.cal](#)

```
Separation
```

Define everything at once

Description

This function is provided to define all the other needed functions by default and load the needed packages into the R workspace.

Usage

```
Separation()
```

Details

If there is a need to change the source code feel free to edit it in this file.

Author(s)

Florian Heigwer

See Also

[GUI](#)

```
textstring.entry.window
```

Open an interactive window asking for textinput

Description

This function opens an interactive window asking for textinput.

Usage

```
textstring.entry.window(title, question, entryInit, entryWidth = 20, returnValOn
```

Arguments

title	the title of the window as character string
question	the question to ask in the window
entryInit	the default text entry when opening the window
entryWidth	the width of the textinput area
returnValOnCancel	what should be printed when Cancel has been pushed

Value

ReturnVal	the typed character string
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Author(s)

Florian Heigwer

References

James Wettenhall, Philippe Grosjean (2005)

See Also

[numeric.entry.window](#)

timecourse	<i>Define timecourses</i>
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Description

This function calculates a series of timepoints.

Usage

```
timecourse(endpoint, measurementinterval)
```

Arguments

endpoint	endpoint of the measurement
measurementinterval	time interval between single measurements

Value

times	a vector of timespoints
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Warning

Time interval and the endpoint of the measurement got to be in the same unit.

Author(s)

Florian Heigwer

See Also

[dilutionseries](#)

Examples

```
time.gone=60 #minutes  
interval=1.5 #minutes  
  
timecourse(time.gone,interval)
```

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