

# Package ‘KOMA’

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**Type** Package

**Title** Kinetic Operating Microarray Analyzer (KOMA)

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**Author** Florian Heigwer <florian\_heigwer@web.de>

**Maintainer** Stefan Woelfl <wolfl@uni-hd.de>

**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

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analyte.choice      *Choosing an analyte*

### 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

### Value

analyte      the name of the chosen analyte as character string

### Author(s)

Florian Heigwer

calculate.plot.mean.and.error  
*Calculate and plot the mean and error surfaces*

### 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**

`times` a vector of time points in the given interval

`concentrations`  
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**

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

**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

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

### Value

signaldata	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)
```



**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](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**

`path`            the path to search the directories in

**Value**

`Q`                the names of the found directories

**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**

`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

**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

<code>data</code>	datasets as a list of matrices for each replicate, for each timepoint and eventually each concentration while calibration containing signal values
<code>substance</code>	the analytename as character string
<code>confidencevalues</code>	datasets as a list of matrices for each replicate, for each timepoint and eventually each concentration while calibration containing confidence values
<code>workpath</code>	the directory with the measured rawdata
<code>foldernames</code>	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                      *Define timecourses*

---

**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
-------	-------------------------

**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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