

# Package ‘ade4TkGUI’

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**Title** 'ade4' Tcl/Tk Graphical User Interface

**Depends** ade4 (>= 1.4-3), adegraphics

**Imports** graphics, grDevices, lattice, stats, tcltk, utils

**Suggests** pixmap, tkrplot

**Description** A Tcl/Tk GUI for some basic functions in the 'ade4' package.

**License** GPL (>= 2)

**URL** <https://github.com/adeverse/ade4TkGUI>,  
<http://pbil.univ-lyon1.fr/ade4TkGUI/>

**BugReports** <https://github.com/adeverse/ade4TkGUI/issues>

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ade4TkGUI

*Tcl/Tk GUI for the ade4 package*


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## Description

A Tcl/Tk GUI for some basic functions in the ade4 package.

## Usage

```
ade4TkGUI(show = FALSE, history = FALSE)
```

## Arguments

show	a logical value. If TRUE, R commands executed in the GUI are echoed to the console.
history	a logical value. If TRUE, R commands executed in the GUI are stored in the R session history.

## Details

Right-click on ade4 function buttons to get help on these functions. You can also click on the questionhead icon in the dialog windows. This displays the R help for the corresponding function.

In all dialog windows, entries can either be filled with a Set button, or typed directly in the entry field. The Set button displays a listbox with a list of available objects in the global environment. This list is restricted to objects with an adequate type (dataframe, dudi, factor, etc). Entries can also be filled directly by typing the desired value. In this case, R expressions can also be entered, for example `c("red", "green", "blue")`, `doubs$mil[1:20, 1:5]`, `meau$plan$dat`, or `1:20`.

## Buttons

The first row of buttons (- Data sets -) is used to load data sets, either from a tab-delimited text file exported from Excel (Read data file button), or from the ade4 package built-in datasets (Load a data set button; right-click to get help on the selected dataset).

The second block of buttons (- One table analyses -) gives access to simple analyses (like `pca`), in which only one table is analysed.

The third block (- One table with groups -) is for analyses in which the rows of the table belong to several groups. This is the case for example for discriminant analysis (`discrimin`) and for the within and between analyses (see ade4 documentation).

The fourth block (- Two tables analyses -) gives access to three analyses in which the relationships between two data tables are analysed. `cca` is the well known (at least in ecology) canonical correspondence analysis, `coinertia` can be seen as a robust alternative to `cca` when the number of samples is low, and `pcaiv` is principal components analyses with respect to instrumental variables (see ade4 documentation).

The fifth block (- Graphic functions -) contains three buttons that launch the dialog windows for three basic graphic functions in ade4 : `s.label` (scatter diagram of a factor map), `s.class` (scatter diagram with groups), `s.value` (scatter diagram with squares proportional to one variable).

Other graphic functions are available in the Graphics menu : `s.arrow` (scatter diagram with arrows), `s.corcircle` (correlation circle in normed pca), `s.class` (scatter diagram with convex hulls, using the `chullSize` argument), and `s.match` (scatter diagram of two paired clouds of points).

The sixth block (- Advanced graphics -) can be used for several things:

- `dudi display` displays a dialog window grouping all the components of a `dudi`. Each of these components is represented by a button which action is to draw a graphical display of the corresponding component. The axes used to draw these graphics can be set by the user. The last row of buttons gives access to particular graphic functions that can be used according to the `dudi` type.
- `MCTests` displays a dialog window for computing Monte-Carlo tests after (e.g.) a between or coinertia analysis.
- `ordiClust` displays a dialog window for analysing cluster on ordination scores, providing a dynamic exploration of the clusters on the factor maps.

## Menu bar

The menu bar at the top of the window can be used to launch the same functions, plus several others. All these menus are tear-off.

The File menu can be used to read and load files and datasets, to edit a dataframe, and to quit R :

- Read text file allows to read a data file and store the result in a dataframe. It can be used to read standard Excel tab-delimited text files (with variable names on the first row and sample names in the first column, leaving the first cell empty). The name "clipboard" can be used to read a data table just copied from an Excel data sheet (so it is not necessary to save the data in a text file).
- Load data set allows to load a data set from the ade4 package.
- Edit data frame can be used to edit a dataframe.
- Quit R opens a dialog window to ask if the environment should be saved before quitting R.

The Windows menu allows to manage several graphical windows :

- New graphic window opens a new graphic window and makes it the active window. This is useful to compare easily several graphics.
- Change graphic window changes the active window (i.e., the one into which the next graphic will be drawn).
- Save graphic window saves the graphic drawn in the active window into a disk file in several formats (postscript, pdf, etc).

The `1table`, `1table+groups`, `2tables`, and Graphics menus give access to several other analysis methods and graphics.

## History and command line echoing

If the `show` argument is set to `TRUE`, then all the commands executed in the GUI are echoed to the console. This is handy for complex commands, as it allows to check the exact syntax of the command that was executed.

If the `history` argument is set to `TRUE`, then the commands executed in the GUI (the same as the commands which are echoed to the console) are stored in the R session history buffer. For each command, the history buffer is written to a temporary disk file with `savehistory()`, the command is appended to this file, and the file is reread in memory with `loadhistory()`.

The value (`TRUE` or `FALSE`) of both arguments is recalled in the title of the window (`ade4TkGUI`), so that different instances of the GUI launched simultaneously with different values for these arguments can be easily recognized.

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### Examples

```
## Not run:  
## Start the GUI  
ade4TkGUI()  
  
## End(Not run)
```

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ade4TkGUI-internal      *Internal functions for the ade4TkGUI package.*

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### Description

Internal functions for the `ade4TkGUI` package

### Details

These are not to be called by the user.

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ordiClust      *Ordination and Cluster analysis*

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### Description

This function does cluster analysis on ordination scores, providing a dynamic exploration of the clusters on the factor maps. As a first step, an ordination method computes the row scores of the input data table. In a second step, cluster analysis is used on these row scores to obtain groups. The `dudi` and the factor are available in the global environment under the names `ordiClust.dudi` and `ordiClust.factor`.

### Usage

```
ordiClust(datatab=NULL, hscalef=1.2, vscalef=1.2, maxgr=20)
```

**Arguments**

<code>datatab</code>	a dataframe containing the data table to analyse, or directly a dudi.
<code>hscalef</code>	horizontal scale factor used to resize the drawing in the tkrplot window.
<code>vscalef</code>	vertical scale factor used to resize the drawing in the tkrplot window.
<code>maxgr</code>	maximum number of groups displayed on the ordination factor map.

**Details**

The GUI is divided in a graph panel on the right and several dialog panels on the left.

Under the graph panel, a scale widget can be used to set the number of groups, and three buttons allow to change the graphical parameters (ellipses or convex hulls, and color or black & white). The Submit button under the graph (or the return key) draws the groups on the factor map. Up arrow and Down arrow keys (or the scale widget) can be used to increase or decrease the number of groups. Choosing another clustering algorithm or another distance automatically updates the graph, allowing easy comparisons. The Dismiss button at the bottom of the window closes the ordiClust window, and the Save button can be used to save the current graphic in a disk file.

From top to bottom the left panels are :

Input data frame : this is the data table that will be analysed. The user can set it with the Set button.

Input dudi : this is directly the dudi that will be analysed. The user can set it with the Set button. No ordination method should be used when a dudi is selected here: cluster analysis will be done on the row scores of this dudi.

Ordination method : the user can choose between principal component analysis (centered or normed), correspondence analysis, and multiple correspondence analysis. After computations are finished, the eigenvalues bar chart is displayed in the right panel. The user can then set the number of principal components (axes) on which cluster analysis will be done with the Set button (defaults to 2). If no number is chosen, then all axes are kept.

Ordination graph : draw the factor maps for rows or columns. Axis numbers can be chosen (default is 1 for X axis and 2 for Y axis).

Cluster analysis - Distance : choose the way distances are computed among rows. See `help("dist")`. The dudi option is a special case: computations are done with the `dist.dudi` function of the `ade4` package (i.e., for a normed PCA this is the Euclidean distance computed on normed variables).

Cluster analysis - Cluster method : choose the clustering algorithm. See `help("hclust")`. The Submit button starts the computations and the hierarchical tree is displayed in the right panel. Choosing another algorithm or another distance automatically updates the graph, allowing easy comparisons.

Number of groups : this is the number of groups used to cut the tree computed by cluster analysis. The Cut tree button cuts the tree and draws a red line on the right panel. Up arrow and Down

arrow keyboard keys (or the scale widget) can be used to increase or decrease this value respectively.

**Level height** : height of the cut level in the tree computed by cluster analysis. The Cut tree button cuts the tree and draws a red line on the right panel.

**Inertia ratio** : this is the between group to total inertia ratio (percentage of explained variance). The Draw curve button draws the curve of this ratio as a function of the number of groups. Steps on this curve can be used to choose the number of groups. The red cross on this graph gives the number of groups and the corresponding percentage of between-groups inertia. Up arrow and Down arrow keys (or the scale widget) can be used to increase or decrease the number of groups. Choosing another clustering algorithm or another distance automatically updates the graph, allowing easy comparisons.

**BGA MCTest p-value** : p-value of the Between Groups Analysis Monte-Carlo test (BGA, ade4 between function).

The dudi and the factor used to draw the graphs are available in the global environment, with the names `ordiClust.dudi` and `ordiClust.factor`.

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### Examples

```
## Not run:  
## Start the GUI form the console  
data(meau)  
ordiClust(dudi.pca(meau$mil,scan=F))  
  
## End(Not run)
```

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## \* **multivariate**

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