

# Package ‘zoom’

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**Type** Package

**Title** Allow to zoom/navigate in any plot.

**Version** 2.0.1

**Date** 2013-08-22

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**Description** zm(), called with any active plot allow to enter an interactive session to zoom/navigate any plot. If it sounds awesome it’s because it is. The development version, as well as binary releases can be found at

**https** //github.com/cbarbu/R-package-zoom

**License** LGPL

**Suggests** testthat

**Collate** ‘zoom-package.R’ ‘zoom.R’ ‘utils.R’

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zoom-package

*Zoom/navigate any plot directly in R.***Description**

The single `zm()` function allows to launch an interactive viewing session with any plot. You can zoom in and out and move the scope (just as in Google maps). Additionally, if a device contains several plots it will simultaneously navigate all the plots.

**Details**

Finally, it allows to print on the fly the current state and continue the navigation

It fully works under Windows standard GUI and Linux.

On Mac it fully works if launching R from the Xquartz terminal. It will default to the "session" mode if launched from Rgui. R launched from the system terminal may have a bug with X11 fonts that prevents even the simplest plots to be displayed, it will not get any better with this package.

Rstudio: should work fine on windows after replot of the graph in a native windows window. On Mac and linux it should fallback to the less interactive session, allowing to zoom in the Rstudio plotting region itself.

See `help(zm)` for more details on how to use the package.

```
Package:  zoom
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```

The only function end users should ever use is `zm()`. It orchestrate the access to other functions like:

- `navigation.zoom()`: launch a mouse interaction
- `session.zoom()`: launch a console menu interaction

Themselves orchestrating lower level functions:

- `in.zoom()`: zoom in
- `out.zoom()`: zoom out
- `set.zoom()`: zoom to a magnification factor
- `sq.zoom()`: zoom on a user defined square
- `zoomplot.zoom()`: the heart function replotting everything as directed by higher level functions.

**Note**

This package is maintained as part of the wider `spatcontrol` project on github: <https://github.com/cbarbu/spatcontrol>  
Bug reports/suggestions/patches can be directly submitted in this web interface.

**Known issues:**

- print to pdf: in navigation mode, print of a pdf generates a weird pdf. Exiting and printing using print.zoom() or simply dev.print() works fine.

**Author(s)**

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**Examples**

```
## Not run:
plot(rnorm(1000), rnorm(1000))
zm()

## End(Not run)
```

---

in.zoom

*Direct access to zoom functionalities.*

---

**Description**

Direct selection of a zoom method of the "session" type. Possibly of use in scripts?  
allow interactive in/out zoom in "session" mode

**Usage**

```
in.zoom(...)

inout.zoom(...)

out.zoom(...)

set.zoom(...)

sq.zoom(...)
```

**Arguments**

...                   Extra arguments to zoomplot.zoom.

**Value**

NULL

**Note**

Each function starts a different interactive sequence

- `inout.zoom()`: left click within bounds zooms in, outside bounds zoom out
- `in.zoom()`: each left click zooms in
- `out.zoom()`: each left click zooms out
- `set.zoom()`: ask for a magnification factor
- `sq.zoom()`: allow to click on the two corners of the desired region to zoom on

**Author(s)**

Corentin M. Barbu

**See Also**

`zm()`, `session.zoom()`.

---

`session.zoom`

*Opening of an interactive zoom/navigate session.*

---

**Description**

To launch an interactive session you should use `zm()` but if you are sure of your device you can launch directly one of these functions.

**Usage**

```
session.zoom(...)
```

```
navigation.zoom(...)
```

**Arguments**

... Everything that can be accepted by `sq.zoom`.

**Details**

`session.zoom` launch an interactive console menu to navigate a plot.

`navigation.zoom` allows to interactively navigate a plot with the mouse.

**Value**

Returns the final plot, as saved by `recordPlot()`.

**Author(s)**

Corentin M. Barbu

**See Also**

zm().

**Examples**

```
## Not run:
plot(rnorm(100),rnorm(100))
session.zoom()

## End(Not run)
```

---

zm

*Launch interaction on a plot*


---

**Description**

Allow to zoom/navigate in any open plot. The controls should be intuitive:

- zoom in: scroll up, or right click if no scrolling wheel.
- zoom out: scroll down, or double click if no scrolling wheel.
- move: left click and move

**Usage**

```
zm(type = "navigation", rp = NULL)
```

**Arguments**

type	the type of interaction with the plot. Possible types are: <ul style="list-style-type: none"> <li>• session for console menu</li> <li>• navigation for mouse interaction</li> </ul> Or any short names for these. By default will try to launch a "navigation" session.
rp	plot to navigate, saved using <code>rp&lt;-recordPlot()</code> . By default (NULL) will use the current device.

**Details**

By default, `zm()` try to open a mouse interactive session. If the current device is not interactive, will try to replot the current plot in a X11 (`type="Xlib"`) device. If it fails it will open a console menu based interactive session.

Zoom handle multiple plots on a device together. You need to navigate the last one plotted and all the other plots will be navigated according to the last one: that can be pretty amazing too if you want to explore multiple layers at the same time.

**Value**

The recording of the final plot. Can be replotted using `replayPlot()`. The most useful may be to get the `xlim` and `ylim` of the final plot. That can be simply got using: `par("usr")` after `zm()` ends.

**Note**

This function relies on pretty low level functions in R that change quite often with new versions. New version of R can break this package but I got used to it and fix it quickly.

In case you close the device before striking q, just hit Ctrl-C on the command line.

**Author(s)**

Corentin M. Barbu

**Examples**

```
## Not run:
# basic example
plot(rnorm(1000),rnorm(1000)) # could be any plot
zm() # navigate the plot

# use the same xlim/ylim as ended up in the zoom session
xylim<-par("usr") # xmin,xmax,ymin,ymax of the final version of the plot
dev.off()
plot(rnorm(1000),rnorm(1000),xlim=xylim[1:2],ylim=xylim[3:4])

# navigate two layers of data at the same time
par(mfrow=c(1,2))
plot(1,type="n",xlim=c(-3,3),ylim=c(-3,3),main="First Track")
polygon(c(-1,1,1,-1)*2,c(-1,-1,1,1)*2,col="blue")
lines(rnorm(100),rnorm(100))
plot(1,type="n",xlim=c(-3,3),ylim=c(-3,3),main="Second Track")
polygon(c(-1,1,1,-1)*2,c(-1,-1,1,1)*2,col="green")
lines(rnorm(100),rnorm(100))
zm() # it flickers quite a bit as it needs to replot everything every time...

# one might want to use the older interface
# if attached to cairo under linux or MacOS
# it is also sometimes helpful to just define a square you want to zoom on
zm(type="s")

## End(Not run)
```

---

zoomplot.zoom

*Central low level function of the zoom package.*

---

**Description**

This function allow to replot the current or a saved plot with specific boundaries, magnification factor and possibly around a user defined x/y.

**Usage**

```
zoomplot.zoom(xlim = NULL, ylim = NULL, fact = NULL,  
              rp = NULL, x = NULL, y = NULL, ...)
```

**Arguments**

xlim	A vector with min and max x
ylim	A vector with min and max y
fact	A scalar giving the magnification factor (>1 brings you closer)
rp	A previously recorded plot with recordPlot(). With all the corresponding warnings in ?recordPlot.
x	x of a fix point when rescaling, by default the center.
y	y of a fix point when rescaling, by default the center.
...	Additional parameters not implemented, just in case.

**Details**

This function is not necessarily easy to use by hand. It is designed to work well when called from higher level functions. End user should always use zm().

**Value**

Not guaranted for now.

**Note**

This function is the heart of the zoom package and the one that can be affected by R version changes.

**Author(s)**

Corentin M. Barbu

**See Also**

zm, in.zoom

**Examples**

```
plot(rnorm(1000),rnorm(1000))  
zoomplot.zoom(fact=2,x=0,y=0)
```

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