

# Package ‘zoom’

May 8, 2026

**Type** Package

**Title** A Spatial Data Visualization Tool

**Depends** R (>= 2.10.0)

**Suggests** testthat

**Description** You can call `zm()`, when displaying any active plot to enter an interactive session to zoom/navigate any plot. The development version, as well as binary releases can be found at <https://github.com/cbarbu/R-package-zoom>.

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyLoad** yes

**URL** <https://github.com/cbarbu/R-package-zoom>

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zoom-package

*The zoom Package*

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**Description**

A spatial data visualization tool.

**Details**

Package: zoom  
Type: Package  
Version: 2.0.6  
Date: 2014-03-18  
Depends: R (>= 2.10.0)  
Encoding: UTF-8  
License: GPL (>= 3)  
LazyLoad: yes  
URL: <https://github.com/cbarbu/R-package-zoom>

zm(), enters an interactive session to zoom and navigate the active plot. The development version, as well as binary releases can be found at <https://github.com/cbarbu/R-package-zoom>

**Author(s)**

Corentin M Barbu <[corentin.barbu@gmail.com](mailto:corentin.barbu@gmail.com)>, with contributions from Sebastian Gibb <[mail@sebastiangibb.de](mailto:mail@sebastiangibb.de)>

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in.zoom

*Direct access to zoom functionalities.*

---

**Description**

Direct selection of a zoom method of the "session" type. Possibly of use in scripts?

**Usage**

```
in.zoom(...)
```

```
move.to.click.zoom(...)
```

```
inout.zoom(...)
```

```
out.zoom(...)
```

```
set.zoom(...)
```

```
sq.zoom(...)
```

**Arguments**

... Extra arguments to zoomplot.zoom.

**Note**

Each function starts a different interactive sequence

- `inout.zoom()`: left click within bounds zooms in, outside bounds zoom out
- `move.to.click.zoom()`: center plot around left click
- `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, Sebastian Gibb

**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 Hold left + right click if no working wheel.
- move: left click and move

**Usage**

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

**Arguments**

`type` the type of interaction with the plot. Possible types are:

- session for console menu
- navigation for mouse interaction

Or any short names for these. By default will try to launch a "navigation" session.

`rp` plot to navigate, saved using `rp<-recordPlot()`. 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
xyylim<-par("usr") # xmin,xmax,ymin,ymax of the final version of the plot
dev.off()
plot(rnorm(1000),rnorm(1000),xlim=xyylim[1:2],ylim=xyylim[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 allows 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,
  moveX = NULL,
  moveY = NULL,
  rp = NULL,
  x = NULL,
  y = NULL,
  xlimfn = NULL,
  ylimfn = 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)
moveX	Expected shift on X axis.
moveY	Expected shift on Y axis. corresponding warnings in ?recordPlot.
rp	A previously recorded plot with recordPlot(). With all the
x	x of a fix point when rescaling, by default the center.
y	y of a fix point when rescaling, by default the center.
xlimfn	a function using x, y and/or fact to generate new x lim if NULL and xlim/ylim not given will use multipancPoint
ylimfn	a function using x, y and/or fact to generate new y lim, if NULL will use xlimfn
...	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 guaranteed for now.

**Note**

This function is the heart of the zoom package and the one that can be affected by R version changes. It is inspired by the `zoomplot` function in TeachingDemos package

**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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