

# Package ‘raster’

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parallel, rasterVis, MASS, sf, testthat, gstat, fields

**LinkingTo** Rcpp

**Imports** Rcpp, methods

**SystemRequirements** C++11

**Description** Reading, writing, manipulating, analyzing and modeling of spatial data. The package implements basic and high-level functions for raster data and for vector data operations such as intersections. See the manual and tutorials on <<https://rspatial.org/>> to get started.

**License** GPL (>= 3)

**URL** <https://rspatial.org/raster>

**BugReports** <https://github.com/rspatial/raster/issues/>

**NeedsCompilation** yes

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raster-package	<i>Overview of the functions in the raster package</i>
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## Description

The raster package provides classes and functions to manipulate geographic (spatial) data in 'raster' format. Raster data divides space into cells (rectangles; pixels) of equal size (in units of the coordinate reference system). Such continuous spatial data are also referred to as 'grid' data, and be contrasted with discrete (object based) spatial data (points, lines, polygons).

The package should be particularly useful when using very large datasets that can not be loaded into the computer's memory. Functions will work correctly, because they process large files in chunks, i.e., they read, compute, and write blocks of data, without loading all values into memory at once.

Below is a list of some of the most important functions grouped by theme. See the vignette for more information and some examples (you can open it by running this command: `vignette('Raster')`)

## Details

The package implements classes for Raster data (see [Raster-class](#)) and supports

- Creation of Raster\* objects from scratch or from file
- Handling extremely large raster files
- Raster algebra and overlay functions
- Distance, neighborhood (focal) and patch functions
- Polygon, line and point to raster conversion
- Model predictions

- Summarizing raster values
- Easy access to raster cell-values
- Plotting (making maps)
- Manipulation of raster extent, resolution and origin
- Computation of row, column and cell numbers to coordinates and vice versa
- Reading and writing various raster file types

## I. Creating Raster\* objects

RasterLayer, RasterStack, and RasterBrick objects are, as a group, referred to as Raster\* objects. Raster\* objects can be created, from scratch, files, or from objects of other classes, with the following functions:

raster	To create a RasterLayer
stack	To create a RasterStack (multiple layers)
brick	To create a RasterBrick (multiple layers)
subset	Select layers of a RasterStack/Brick
addLayer	Add a layer to a Raster* object
dropLayer	Remove a layer from a RasterStack or RasterBrick
unstack	Create a list of RasterLayer objects from a RasterStack

---

## II. Changing the spatial extent and/or resolution of Raster\* objects

merge	Combine Raster* objects with different extents (but same origin and resolution)
mosaic	Combine RasterLayers with different extents and a function for overlap areas
crop	Select a geographic subset of a Raster* object
extend	Enlarge a Raster* object
trim	Trim a Raster* object by removing exterior rows and/or columns that only have NAs
aggregate	Combine cells of a Raster* object to create larger cells
disaggregate	Subdivide cells
resample	Warp values to a Raster* object with a different origin or resolution
projectRaster	project values to a raster with a different coordinate reference system
shift	Move the location of Raster
flip	Flip values horizontally or vertically
rotate	Rotate values around the date-line (for lon/lat data)
t	Transpose a Raster* object

---

## III. Raster algebra

Arith-methods	Arith functions (+, -, *, ^, %, %/, /)
Math-methods	Math functions like abs, sqrt, trunc, log, log10, exp, sin, round
Logic-methods	Logic functions (!, &,  )
Summary-methods	Summary functions (mean, max, min, range, prod, sum, any, all)
Compare-methods	Compare functions (==, !=, >, <, <=, >=)

---

#### IV. Cell based computation

calc	Computations on a single Raster* object
overlay	Computations on multiple RasterLayer objects
cover	First layer covers second layer except where the first layer is NA
mask	Use values from first Raster except where cells of the mask Raster are NA
cut	Reclassify values using ranges
subs	Reclassify values using an 'is-becomes' matrix
reclassify	Reclassify using a 'from-to-becomes' matrix
init	Initialize cells with new values
stackApply	Computations on groups of layers in Raster* object
stackSelect	Select cell values from different layers using an index RasterLayer

---

#### V. Spatial contextual computation

distance	Shortest distance to a cell that is not NA
gridDistance	Distance when traversing grid cells that are not NA
distanceFromPoints	Shortest distance to any point in a set of points
direction	Direction (azimuth) to or from cells that are not NA
focal	Focal (neighborhood; moving window) functions
localFun	Local association (using neighborhoods) functions
boundaries	Detection of boundaries (edges)
clump	Find clumps (patches)
adjacent	Identify cells that are adjacent to a set of cells on a raster
area	Compute area of cells (for longitude/latitude data)
terrain	Compute slope, aspect and other characteristics from elevation data
Moran	Compute global or local Moran or Geary indices of spatial autocorrelation

---



**VI. Model predictions**

<code>predict</code>	Predict a non-spatial model to a RasterLayer
<code>interpolate</code>	Predict a spatial model to a RasterLayer

---

**VII. Data type conversion**

You can coerce Raster\* objects to Spatial\* objects using `as`, as in `as(object, 'SpatialGridDataFrame')`

<code>raster</code>	RasterLayer from SpatialGrid*, image, or matrix objects
<code>rasterize</code>	Rasterizing points, lines or polygons
<code>rasterToPoints</code>	Create points from a RasterLayer
<code>rasterToPolygons</code>	Create polygons from a RasterLayer
<code>rasterToContour</code>	Contour lines from a RasterLayer
<code>rasterFromXYZ</code>	RasterLayer from regularly spaced points
<code>rasterFromCells</code>	RasterLayer from a Raster object and cell numbers

---

**VIII. Summarizing**

<code>cellStats</code>	Summarize a Raster cell values with a function
<code>summary</code>	Summary of the values of a Raster* object (quartiles and mean)
<code>freq</code>	Frequency table of Raster cell values
<code>crosstab</code>	Cross-tabulate two Raster* objects
<code>unique</code>	Get the unique values in a Raster* object
<code>zonal</code>	Summarize a Raster* object by zones in a RasterLayer

---

**IX. Accessing values of Raster\* object cells**

Apart from the function listed below, you can also use indexing with `[]` for cell numbers, and `[[[]]` for row / column number combinations

<code>getValues</code>	Get all cell values (fails with very large rasters), or a row of values (safer)
<code>getValuesBlock</code>	Get values for a block (a rectangular area)
<code>getValuesFocal</code>	Get focal values for one or more rows
<code>as.matrix</code>	Get cell values as a matrix
<code>as.array</code>	Get cell values as an array
<code>extract</code>	Extract cell values from a Raster* object (e.g., by cell, coordinates, polygon)

<code>sampleRandom</code>	Random sample
<code>sampleRegular</code>	Regular sample
<code>minValue</code>	Get the minimum value of the cells of a Raster* object (not always known)
<code>maxValue</code>	Get the maximum value of the cells of a Raster* object (not always known)
<code>setMinMax</code>	Compute the minimum and maximum value of a Raster* object if these are not known

---

## X. Plotting

See the rasterVis package for additional plotting methods for Raster\* objects using methods from 'lattice' and other packages.

### Maps

<code>plot</code>	Plot a Raster* object. The main method to create a map
<code>plotRGB</code>	Combine three layers (red, green, blue channels) into a single 'real color' image
<code>splot</code>	Plot a Raster* with the splot function (sp package)
<code>image</code>	Plot a Raster* with the image function
<code>persp</code>	Perspective plot of a RasterLayer
<code>contour</code>	Contour plot of a RasterLayer
<code>filledContour</code>	Filled contour plot of a RasterLayer
<code>text</code>	Plot the values of a RasterLayer on top of a map

.

### Interacting with a map

<code>zoom</code>	Zoom in to a part of a map
<code>click</code>	Query values of Raster* or Spatial* objects by clicking on a map
<code>select</code>	Select a geometric subset of a Raster* or Spatial* object
<code>drawPoly</code>	Create a SpatialPolygons object by drawing it
<code>drawLine</code>	Create a SpatialLines object by drawing it
<code>drawExtent</code>	Create an Extent object by drawing it

.

### Other plots

<code>plot</code>	x-y scatter plot of the values of two RasterLayer objects
<code>hist</code>	Histogram of Raster* object values
<code>barplot</code>	barplot of a RasterLayer
<code>density</code>	Density plot of Raster* object values
<code>pairs</code>	Pairs plot for layers in a RasterStack or RasterBrick
<code>boxplot</code>	Box plot of the values of one or multiple layers

---

## XI. Getting and setting Raster\* dimensions

Basic parameters of existing Raster\* objects can be obtained, and in most cases changed. If there are values associated with a RasterLayer object (either in memory or via a link to a file) these are lost when you change the number of columns or rows or the resolution. This is not the case when the extent is changed (as the number of columns and rows will not be affected). Similarly, with **projection** you can set the projection, but this does not transform the data (see `projectRaster` for that).

<code>ncol</code>	The number of columns
<code>nrow</code>	The number of rows
<code>ncell</code>	The number of cells (can not be set directly, only via <code>ncol</code> or <code>nrow</code> )
<code>res</code>	The resolution (x and y)
<code>nlayers</code>	How many layers does the object have?
<code>names</code>	Get or set the layer names
<code>xres</code>	The x resolution (can be set with <code>res</code> )
<code>yres</code>	The y resolution (can be set with <code>res</code> )
<code>xmin</code>	The minimum x coordinate (or longitude)
<code>xmax</code>	The maximum x coordinate (or longitude)
<code>ymin</code>	The minimum y coordinate (or latitude)
<code>ymax</code>	The maximum y coordinate (or latitude)
<code>extent</code>	The extent (minimum and maximum x and y coordinates)
<code>origin</code>	The origin of a Raster* object
<code>crs</code>	The coordinate reference system (map projection)
<code>isLonLat</code>	Test if an object has a longitude/latitude coordinate reference system
<code>filename</code>	Filename to which a RasterLayer or RasterBrick is linked
<code>bandnr</code>	layer (=band) of a multi-band file that this RasterLayer is linked to
<code>nbands</code>	How many bands (layers) does the file associated with a RasterLayer object have?
<code>compareRaster</code>	Compare the geometry of Raster* objects
<code>NAvalue</code>	Get or set the NA value (for reading from a file)

---

## XII. Computing row, column, cell numbers and coordinates

Cell numbers start at 1 in the upper-left corner. They increase within rows, from left to right, and then row by row from top to bottom. Likewise, row numbers start at 1 at the top of the raster, and column numbers start at 1 at the left side of the raster.

<code>xFromCol</code>	x-coordinates from column numbers
<code>yFromRow</code>	y-coordinates from row numbers
<code>xFromCell</code>	x-coordinates from row numbers
<code>yFromCell</code>	y-coordinates from cell numbers
<code>xyFromCell</code>	x and y coordinates from cell numbers
<code>colFromX</code>	Column numbers from x-coordinates (or longitude)
<code>rowFromY</code>	Row numbers from y-coordinates (or latitude)
<code>rowColFromCell</code>	Row and column numbers from cell numbers
<code>cellFromXY</code>	Cell numbers from x and y coordinates
<code>cellFromRowCol</code>	Cell numbers from row and column numbers
<code>cellsFromExtent</code>	Cell numbers from extent object
<code>coordinates</code>	x and y coordinates for all cells
<code>validCell</code>	Is this a valid cell number?
<code>validCol</code>	Is this a valid column number?
<code>validRow</code>	Is this a valid row number?

---

### XIII. Writing files

#### Basic

<code>setValues</code>	Put new values in a Raster* object
<code>writeRaster</code>	Write all values of Raster* object to disk
<code>KML</code>	Save raster as KML file

#### Advanced

<code>blockSize</code>	Get suggested block size for reading and writing
<code>writeStart</code>	Open a file for writing
<code>writeValues</code>	Write some values
<code>writeStop</code>	Close the file after writing
<code>update</code>	Change the values of an existing file

---

### XIV. Manipulation of SpatialPolygons\* and other vector type Spatial\* objects

Some of these functions are in the `sp` package. The name in **bold** is the equivalent command in ArcGIS. These functions build on the geometry ("spatial features") manipulation functions in package `rgeos`. These functions are extended here by also providing automated attribute data handling.

<code>bind</code>	<b>append</b> combine Spatial* objects of the same (vector) type
<code>erase</code> or <code>"-"</code>	<b>erase</b> parts of a SpatialPolygons* object
<code>intersect</code> or <code>"*"</code>	<b>intersect</b> SpatialPolygons* objects
<code>union</code> or <code>"+"</code>	<b>union</b> SpatialPolygons* objects
<code>cover</code>	<b>update</b> and <b>identity</b> for a SpatialPolygons and another one
<code>symdif</code>	<b>symmetrical difference</b> of two SpatialPolygons* objects
<code>aggregate</code>	<b>dissolve</b> smaller polygons into larger ones
<code>disaggregate</code>	<b>explode</b> : turn polygon parts into separate polygons (in the <code>sp</code> package)
<code>crop</code>	<b>clip</b> a Spatial* object using a rectangle (Extent object)
<code>select</code>	<b>select</b> - interactively select spatial features
<code>click</code>	<b>identify</b> attributes by clicking on a map
<code>merge</code>	<b>Join table</b> (in the <code>sp</code> package)
<code>over</code>	spatial queries between Spatial* objects
<code>extract</code>	spatial queries between Spatial* and Raster* objects
<code>as.data.frame</code>	coerce coordinates of SpatialLines or SpatialPolygons into a data.frame

---

### XV. Extent objects

<code>extent</code>	Create an extent object
<code>intersect</code>	Intersect two extent objects
<code>union</code>	Combine two extent objects
<code>round</code>	round/floor/ceiling of the coordinates of an Extent object

<a href="#">alignExtent</a>	Align an extent with a Raster* object
<a href="#">drawExtent</a>	Create an Extent object by drawing it on top of a map (see plot)

---

## XVI. Miscellaneous

<a href="#">rasterOptions</a>	Show, set, save or get session options
<a href="#">getData</a>	Download and geographic data
<a href="#">pointDistance</a>	Distance between points
<a href="#">readIniFile</a>	Read a (windows) 'ini' file
<a href="#">hdr</a>	Write header file for a number of raster formats
<a href="#">trim</a>	Remove leading and trailing blanks from a character string
<a href="#">extension</a>	Get or set the extension of a filename
<a href="#">cv</a>	Coefficient of variation
<a href="#">modal</a>	Modal value
<a href="#">sampleInt</a>	Random sample of (possibly very large) range of integer values
<a href="#">showTmpFiles</a>	Show temporary files
<a href="#">removeTmpFiles</a>	Remove temporary files

---

## XVII. For programmers

<a href="#">canProcessInMemory</a>	Test whether a file can be created in memory
<a href="#">pbCreate</a>	Initialize a progress bar
<a href="#">pbStep</a>	Take a progress bar step
<a href="#">pbClose</a>	Close a progress bar
<a href="#">readStart</a>	Open file connections for efficient multi-chunk reading
<a href="#">readStop</a>	Close file connections
<a href="#">rasterTmpFile</a>	Get a name for a temporary file
<a href="#">inMemory</a>	Are the cell values in memory?
<a href="#">fromDisk</a>	Are the cell values read from a file?

---

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### Author(s)

Except where indicated otherwise, the functions in this package were written by Robert J. Hijmans

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addLayer	<i>Add or drop a layer</i>
----------	----------------------------

---

### Description

Add a layer to a Raster\* object or drop a layer from a RasterStack or RasterBrick. The object returned is always a RasterStack (unless nothing to add or drop was provided, in which case the original object is returned).

### Usage

```
addLayer(x, ...)
dropLayer(x, i, ...)
```

### Arguments

x	Raster* object
i	integer. Indices of the layers to be dropped
...	Additional arguments. The layers to add for addLayer. None implemented for dropLayer)

### Value

RasterStack

### See Also

[subset](#)

### Examples

```
file <- system.file("external/test.grd", package="raster")
s <- stack(file, file, file)
r <- raster(file)
s <- addLayer(s, r/2, r*2)
s
s <- dropLayer(s, c(3, 5))
nlayers(s)
```

adjacent

*Adjacent cells***Description**

Identify cells that are adjacent to a set of cells on a raster.

**Usage**

```
## S4 method for signature 'BasicRaster'
adjacent(x, cells, directions=4, pairs=TRUE, target=NULL, sorted=FALSE,
         include=FALSE, id=FALSE, ...)
```

**Arguments**

x	Raster* object
cells	vector of cell numbers for which adjacent cells should be found. Cell numbers start with 1 in the upper-left corner and increase from left to right and from top to bottom
directions	the number of directions in which cells should be connected: 4 (rook's case), 8 (queen's case), 16 (knight and one-cell queen moves), or 'bishop' to connect cells with one-cell diagonal moves. Or a neighborhood matrix (see Details)
pairs	logical. If TRUE, a matrix of pairs of adjacent cells is returned. If FALSE, a vector of cells adjacent to cells is returned
target	optional vector of target cell numbers that should be considered. All other adjacent cells are ignored
sorted	logical. Should the results be sorted?
include	logical. Should the focal cells be included in the result?
id	logical. Should the id of the cells be included in the result? (numbered from 1 to length(cells))
...	additional arguments. None implemented

**Details**

A neighborhood matrix identifies the cells around each cell that are considered adjacent. The matrix should have one, and only one, cell with value 0 (the focal cell); at least one cell with value 1 (the adjacent cell(s)); All other cells are not considered adjacent and ignored.

**Value**

matrix or vector with adjacent cells.

**Author(s)**

Robert J. Hijmans and Jacob van Etten

**Examples**

```

r <- raster(nrows=10, ncols=10)
adjacent(r, cells=c(1, 55), directions=8, pairs=TRUE)

a <- adjacent(r, cell = c(1,55,90), directions=4, sorted=TRUE)
a

r[c(1,55,90)] <- 1
r[a] <- 2
plot(r)

# same result as above
rook <- matrix(c(NA, 1, NA,
                1, 0, 1,
                NA, 1, NA), ncol=3, byrow=TRUE)

adjacent(r, cells = c(1,55,90), directions=rook, sorted=TRUE)

# Count the number of times that a cell with a certain value
# occurs next to a cell with a certain value
set.seed(0)
r <- raster(ncol=10, nrow=10)
values(r) <- round(runif(ncell(r)) * 5)
a <- adjacent(r, 1:ncell(r), 4, pairs=TRUE)
tb <- table(r[a[,1]], r[a[,2]])
tb
# make a matrix out of the 'table' object
tb <- unclass(tb)
plot(raster(tb, xmn=-0.5, xmx=5.5, ymn=-0.5, ymx=5.5))

```

---

aggregate

---

*Aggregate raster cells or SpatialPolygons/Lines*


---

**Description**

Raster\* objects:

Aggregate a Raster\* object to create a new RasterLayer or RasterBrick with a lower resolution (larger cells). Aggregation groups rectangular areas to create larger cells. The value for the resulting cells is computed with a user-specified function.

SpatialPolygon\*:

Aggregate a SpatialPolygon\* object, optionally by combining polygons that have the same attributes for one or more variables. If the polygons touch or overlap, internal boundaries are optionally "dissolved".



**Usage**

```
## S4 method for signature 'Raster'
aggregate(x, fact, fun=mean, expand=TRUE, na.rm=TRUE, filename='', ...)

## S4 method for signature 'SpatialPolygons'
aggregate(x, by, sums, dissolve=TRUE, vars=NULL, ...)
```

**Arguments**

x	Raster* object or SpatialPolygons* object
fact	postive integer. Aggregation factor expressed as number of cells in each direction (horizontally and vertically). Or two integers (horizontal and vertical aggregation factor) or three integers (when also aggregating over layers). See Details
fun	function used to aggregate values
expand	logical. If TRUE the output Raster* object will be larger then the input Raster* object if a division of the number of columns or rows with factor is not an integer
na.rm	logical. If TRUE, NA cells are removed from calculations
filename	character. Output filename (optional)
...	if x is a Raster* object, additional arguments as for <a href="#">writeRaster</a>
by	character or integer. The variables (column names or numbers) that should be used to aggregate (dissolve) the SpatialPolygons by only maintaining unique combinations of these variables. The default setting is to use no variables and aggregate all polygons. You can also supply a vector with a length of length(x)
sums	list with function(s) and variable(s) to summarize. This should be a list of lists in which each element of the main lists has two items. The first item is function (e.g. mean), the second element is a vector of column names (or indices) that need to summarize with that function. Be careful with character and factor variables (you can use, e.g. 'first' function(x)x[1] or 'last' function(x)x[length(x)] or modal for these variables
vars	deprecated. Same as by
dissolve	logical. If TRUE borders between touching or overlapping polygons are removed (requires package rgeos)

**Details**

Aggregation of a x will result in a Raster\* object with fewer cells. The number of cells is the number of cells of x divided by fact\*fact (when fact is a single number) or prod(fact) (when fact consists of 2 or 3 numbers). If necessary this number is adjusted according to the value of expand. For example, fact=2 will result in a new Raster\* object with 2\*2=4 times fewer cells. If two numbers are supplied, e.g., fact=c(2, 3), the first will be used for aggregating in the horizontal direction, and the second for aggregating in the vertical direction, and the returned object will have 2\*3=6 times fewer cells. Likewise, fact=c(2, 3, 4) aggregates cells in groups of 2 (rows) by 3 (columns) and 4 (layers).

Aggregation starts at the upper-left end of a raster (you can use `flip` if you want to start elsewhere). If a division of the number of columns or rows with `factor` does not return an integer, the extent of the resulting Raster object will either be somewhat smaller or somewhat larger than the original RasterLayer. For example, if an input RasterLayer has 100 columns, and `fact=12`, the output Raster object will have either 8 columns (`expand=FALSE`) (using  $8 \times 12 = 96$  of the original columns) or 9 columns (`expand=TRUE`). In both cases, the maximum x coordinate of the output RasterLayer would, of course, also be adjusted.

The function `fun` should take multiple numbers, and return a single number. For example `mean`, `modal`, `min` or `max`. It should also accept a `na.rm` argument (or ignore it as one of the 'dots' arguments).

### Value

RasterLayer or RasterBrick, or a SpatialPolygons\* object

### Author(s)

Robert J. Hijmans and Jacob van Etten

### See Also

[disaggregate](#), [resample](#). For SpatialPolygons\* [disaggregate](#)

### Examples

```
r <- raster()
# a new aggregated raster, no values
ra <- aggregate(r, fact=10)
r <- setValues(r, runif(ncell(r)))

# a new aggregated raster, max of the values
ra <- aggregate(r, fact=10, fun=max)

# multiple layers
s <- stack(r, r*2)
x <- aggregate(s,2)

#SpatialPolygons
if (require(rgdal) & require(rgeos)) {
p <- shapefile(system.file("external/lux.shp", package="raster"))
p
pa0 <- aggregate(p)
pa0
pa1 <- aggregate(p, by='NAME_1', sums=list(list(mean, 'ID_2')))
pa1
}
```

---

alignExtent	<i>Align an extent (object of class Extent)</i>
-------------	---

---

**Description**

Align an Extent object with the (boundaries of the) cells of a Raster\* object

**Usage**

```
alignExtent(extent, object, snap='near')
```

**Arguments**

extent	Extent object
object	Raster* object
snap	Character. One of 'near', 'in', or 'out', to determine in which direction the extent should be aligned. To the nearest border, inwards or outwards

**Details**

Aligning an Extent object to another object assures that it gets the same origin and resolution. This should only be used to adjust objects because of imprecision in the data. alignExtent should not be used to force data to match that really does not match (use e.g. [resample](#) or (dis)aggregate for this).

**Value**

Extent object

**See Also**

[extent](#), [drawExtent](#), [Extent-class](#)

**Examples**

```
r <- raster()
e <- extent(-10.1, 9.9, -20.1, 19.9)
ea <- alignExtent(e, r)
e
extent(r)
ea
```

---

`animate`*Animate layers of a Raster\* object*

---

**Description**

Animate (sequentially plot) the layers of a RasterStack or RasterBrick\* object to create a movie

**Usage**

```
## S4 method for signature 'RasterStackBrick'  
animate(x, pause=0.25, main, zlim, maxpixels=50000, n=10, ...)
```

**Arguments**

<code>x</code>	Raster* object
<code>pause</code>	numeric. How long should be the pause be between layers?
<code>main</code>	title for each layer. If not supplied the z-value is used if available. Otherwise the names are used.
<code>zlim</code>	numeric vector of length 2. Range of values to plot
<code>maxpixels</code>	integer > 0. Maximum number of cells to use for the plot. If <code>maxpixels &lt; ncell(x)</code> , <code>sampleRegular</code> is used before plotting
<code>n</code>	integer > 0. Number of loops
<code>...</code>	Additional arguments passed to <a href="#">plot</a>

**Value**

None

**See Also**

[plot](#), [spplot](#), [plotRGB](#)

**Examples**

```
b <- brick(system.file("external/rlogo.grd", package="raster"))  
animate(b, n=1)
```

---

approxNA	<i>Estimate values for cell values that are NA by interpolating between layers</i>
----------	--

---

### Description

approxNA uses the stats function [approx](#) to estimate values for cells that are NA by interpolation across layers. Layers are considered equidistant, unless an argument 'z' is used, or [getZ](#) returns values, in which case these values are used to determine distance between layers.

For estimation based on neighbouring cells see [focal](#)

### Usage

```
## S4 method for signature 'RasterStackBrick'
approxNA(x, filename="", method="linear", yleft, yright,
         rule=1, f=0, ties=mean, z=NULL, NArule=1, ...)
```

### Arguments

x	RasterStack or RasterBrick object
filename	character. Output filename (optional)
method	specifies the interpolation method to be used. Choices are "linear" or "constant" (step function; see the example in <a href="#">approx</a> )
yleft	the value to be returned before a non-NA value is encountered. The default is defined by the value of rule given below
yright	the value to be returned after the last non-NA value is encountered. The default is defined by the value of rule given below
rule	an integer (of length 1 or 2) describing how interpolation is to take place at for the first and last cells (before or after any non-NA values are encountered). If rule is 1 then NAs are returned for such points and if it is 2, the value at the closest data extreme is used. Use, e.g., rule = 2:1, if the left and right side extrapolation should differ
f	for method = "constant" a number between 0 and 1 inclusive, indicating a compromise between left- and right-continuous step functions. If y0 and y1 are the values to the left and right of the point then the value is $y_0*(1-f)+y_1*f$ so that $f = 0$ is right-continuous and $f = 1$ is left-continuous
ties	Handling of tied 'z' values. Either a function with a single vector argument returning a single number result or the string "ordered"
z	numeric vector to indicate the distance between layers (e.g., time, depth). The default is 1:nlayers(x)
NArule	single integer used to determine what to do when only a single layer with a non-NA value is encountered (and linear interpolation is not possible). The default value of 1 indicates that all layers will get this value for that cell; all other values do not change the cell values
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterBrick

**See Also**[focal](#)**Examples**

```

r <- raster(ncols=5, nrows=5)
r1 <- setValues(r, runif(ncell(r)))
r2 <- setValues(r, runif(ncell(r)))
r3 <- setValues(r, runif(ncell(r)))
r4 <- setValues(r, runif(ncell(r)))
r5 <- setValues(r, NA)
r6 <- setValues(r, runif(ncell(r)))
r1[6:10] <- NA
r2[5:15] <- NA
r3[8:25] <- NA
s <- stack(r1,r2,r3,r4,r5,r6)
s[1:5] <- NA
x1 <- approxNA(s)
x2 <- approxNA(s, rule=2)
x3 <- approxNA(s, rule=2, z=c(1,2,3,5,14,15))

```

---

**area***Size of cells*

---

**Description**

**Raster objects:** Compute the approximate surface area of cells in an unprojected (longitude/latitude) Raster object. It is an approximation because area is computed as the height (latitudinal span) of a cell (which is constant among all cells) times the width (longitudinal span) in the (latitudinal) middle of a cell. The width is smaller at the poleward side than at the equator-ward side of a cell. This variation is greatest near the poles and the values are thus not very precise for very high latitudes.

**SpatialPolygons:** Compute the area of the spatial features. Works for both planar and angular (lon/lat) coordinate reference systems

**Usage**

```
## S4 method for signature 'RasterLayer'
area(x, filename="", na.rm=FALSE, weights=FALSE, ...)
```

```
## S4 method for signature 'RasterStackBrick'
area(x, filename="", na.rm=FALSE, weights=FALSE, ...)
```

```
## S4 method for signature 'SpatialPolygons'
area(x, ...)
```

### Arguments

<code>x</code>	Raster* or SpatialPolygons object
<code>filename</code>	character. Filename for the output Raster object (optional)
<code>na.rm</code>	logical. If TRUE, cells that are NA are ignored
<code>weights</code>	logical. If TRUE, the area of each cells is divided by the total area of all cells that are not NA
<code>...</code>	additional arguments as for <a href="#">writeRaster</a>

### Details

If `x` is a RasterStack/Brick, a RasterBrick will be returned if `na.rm=TRUE`. However, if `na.rm=FALSE`, a RasterLayer is returned, because the values would be the same for all layers.

### Value

If `x` is a Raster\* object: RasterLayer or RasterBrick. Cell values represent the size of the cell in km<sup>2</sup>, or the relative size if `weights=TRUE`

If `x` is a SpatialPolygons\* object: area if each spatial object in squared meters if the CRS is longitude/latitude, or in squared map units (typically meter)

### Examples

```
r <- raster(nrow=18, ncol=36)
a <- area(r)

if (require(rgdal) & require(rgeos)) {
  p <- shapefile(system.file("external/lux.shp", package="raster"))
  p$area <- round(area(p) / 10000000,1)
  p$area
}
```

### Description

Standard arithmetic operators for computations with Raster\* objects and numeric values. The following operators are available: `+`, `-`, `*`, `/`, `^`, `%%`, `/%`

The input Raster\* objects should have the same extent, origin and resolution. If only the extent differs, the computation will continue for the intersection of the Raster objects. Operators are applied on a cell by cell basis. For a RasterLayer, numeric values are recycled by row. For a RasterStack or RasterBrick, recycling is done by layer. RasterLayer objects can be combined RasterStack/Brick

objects, in which case the RasterLayer is 'recycled'. When using multiple RasterStack or RasterBrick objects, the number of layers of these objects needs to be the same.

In addition to arithmetic with Raster\* objects, the following operations are supported for SpatialPolygons\* objects. Given SpatialPolygon objects x and y:

x+y is the same as [union\(x, y\)](#). For SpatialLines\* and SpatialPoints\* it is equivalent to [bind\(x, y\)](#)

x\*y is the same as [intersect\(x, y\)](#)

x-y is the same as [erase\(x, y\)](#)

### Details

If the values of the output Raster\* cannot be held in memory, they will be saved to a temporary file. You can use [options](#) to set the default file format, datatype and progress bar.

### Value

A Raster\* object, and in some cases the side effect of a new file on disk.

### See Also

[Math-methods](#), [overlay](#), [calc](#)

### Examples

```
r1 <- raster(ncols=10, nrows=10)
values(r1) <- runif(ncell(r1))
r2 <- setValues(r1, 1:ncell(r1) / ncell(r1) )
r3 <- r1 + r2
r2 <- r1 / 10
r3 <- r1 * (r2 - 1 + r1^2 / r2)

# recycling by row
r4 <- r1 * 0 + 1:ncol(r1)

# multi-layer object multiplication, no recycling
b1 <- brick(r1, r2, r3)
b2 <- b1 * 10

# recycling by layer
b3 <- b1 + c(1, 5, 10)

# addition of the cell-values of two RasterBrick objects
b3 <- b2 + b1

# summing two RasterBricks and one RasterLayer. The RasterLayer is 'recycled'
b3 <- b1 + b2 + r1
```



---

as.character	<i>Character representation of a Raster or Extent object</i>
--------------	--

---

### Description

as.character returns a text (R code) representation of a Raster\* or Extent object. The main purpose of this is to allow quick generation of objects to use in examples on, for example, stackoverflow.com.

### Usage

```
## S4 method for signature 'Raster'  
as.character(x, ...)  
## S4 method for signature 'Extent'  
as.character(x, ...)
```

### Arguments

x	Raster* or Extent object
...	additional arguments, none implemented

### Value

character

### Examples

```
r <- raster(ncol=3, nrow=3)  
values(r) <- 1:ncell(r)  
as.character(r)  
s <- stack(r, r)  
as.character(s)  
as.character(extent(s))  
  
x <- as.character(s)  
eval(parse(text=x))  
  
y <- as.character(extent(s))  
eval(parse(text=y))
```

---

as.data.frame	<i>Get a data.frame with raster cell values, or coerce SpatialPolygons, Lines, or Points to a data.frame</i>
---------------	--

---

## Description

as.matrix returns all values of a Raster\* object as a matrix. For RasterLayers, rows and columns in the matrix represent rows and columns in the RasterLayer object. For other Raster\* objects, the matrix returned by as.matrix has columns for each layer and rows for each cell.

as.array returns an array of matrices that are like those returned by as.matrix for a RasterLayer

If there is insufficient memory to load all values, you can use [getValues](#) or [getValuesBlock](#) to read chunks of the file. You could also first use [sampleRegular](#)

The methods for Spatial\* objects allow for easy creation of a data.frame with the coordinates and attributes; the default method only returns the attributes data.frame

## Usage

```
## S4 method for signature 'Raster'
as.data.frame(x, row.names=NULL, optional=FALSE, xy=FALSE,
             na.rm=FALSE, long=FALSE, ...)

## S4 method for signature 'SpatialPolygons'
as.data.frame(x, row.names=NULL, optional=FALSE,
             xy=FALSE, centroids=TRUE, sepNA=FALSE, ...)

## S4 method for signature 'SpatialLines'
as.data.frame(x, row.names=NULL, optional=FALSE,
             xy=FALSE, sepNA=FALSE, ...)
```

## Arguments

x	Raster* object
row.names	NULL or a character vector giving the row names for the data frame. Missing values are not allowed
optional	logical. If TRUE, setting row names and converting column names (to syntactic names: see <a href="#">make.names</a> ) is optional
xy	logical. If TRUE, also return the spatial coordinates
na.rm	logical. If TRUE, remove rows with NA values. This can be particularly useful for very large datasets with many NA values
long	logical. If TRUE, values are <a href="#">reshaped</a> from a wide to a long format
centroids	logical. If TRUE return the centroids instead of all spatial coordinates (only relevant if xy=TRUE)
sepNA	logical. If TRUE the parts of the spatial objects are separated by lines that are NA (only if xy=TRUE and, for polygons, if centroids=FALSE)
...	Additional arguments (none)

**Value**

data.frame

**Examples**

```
r <- raster(ncol=3, nrow=3)
values(r) <- sqrt(1:ncell(r))
r[3:5] <- NA
as.data.frame(r)
s <- stack(r, r*2)
as.data.frame(s)
as.data.frame(s, na.rm=TRUE)
```

---

as.list

*Create a list of RasterLayer objects*

---

**Description**

Create a list of RasterLayer objects from Raster\* objects

**Usage**

```
## S4 method for signature 'Raster'
as.list(x, ...)
```

**Arguments**

x	Raster* object
...	additional Raster* objects

**Value**

list

**Examples**

```
r <- raster(ncol=3, nrow=3)
values(r) <- 1:ncell(r)
as.list(r)

s <- stack(r,r*2,r*3)
as.list(s, r)
```

---

`as.logical`*Change cell values to logical or integer values*

---

### Description

Change values of a Raster\* object to logical or integer values. With `as.logical`, zero becomes FALSE, all other values become TRUE. With `as.integer` values are truncated.

### Usage

```
## S4 method for signature 'Raster'  
as.logical(x, filename='', ...)
```

```
## S4 method for signature 'Raster'  
as.integer(x, filename='', ...)
```

### Arguments

<code>x</code>	Raster* object
<code>filename</code>	character. Output filename (optional)
<code>...</code>	additional optional arguments as for <a href="#">writeRaster</a>

### See Also

[logical](#), [integer](#)

### Examples

```
r <- raster(nrow=10, ncol=10)  
set.seed(0)  
values(r) <- runif(ncell(r)) * 10  
r  
r <- as.integer(r)  
r  
as.logical(r)
```

---

`as.matrix`*Get a vector, matrix, or array with raster cell values*

---

**Description**

as.vector returns a vector of cell values. For a RasterLayer it is equivalent to `getValues(x)`.

as.matrix returns all values of a Raster\* object as a matrix. For RasterLayers, rows and columns in the matrix represent rows and columns in the RasterLayer object. For other Raster\* objects, the matrix returned by `as.matrix` has columns for each layer and rows for each cell.

as.array returns an array of matrices that are like those returned by `as.matrix` for a RasterLayer

If there is insufficient memory to load all values, you can use [getValues](#) or [getValuesBlock](#) to read chunks of the file.

as.matrix and as.vector can also be used to obtain the coordinates from an Extent object.

**Usage**

```
as.matrix(x, ...)
as.array(x, ...)

## S4 method for signature 'Extent'
as.vector(x, mode='any')

## S4 method for signature 'Raster'
as.vector(x, mode='any')
```

**Arguments**

x	Raster* or (for <code>as.matrix</code> and <code>as.vector</code> ) Extent object
mode	Character string giving an atomic mode (such as "numeric" or "character") or "list", or "any". Note: this argument is currently ignored!
...	additional arguments: maxpixels Integer. To regularly subsample very large objects transpose Logical. Transpose the data? (for <code>as.array</code> only)

**Value**

matrix, array, or vector

**Examples**

```
r <- raster(ncol=3, nrow=3)
values(r) <- 1:ncell(r)
as.matrix(r)
s <- stack(r,r)
as.array(s)
as.vector(extent(s))
```

---

as.raster *Coerce to a 'raster' object*

---

### Description

Implementation of the generic `as.raster` function to create a 'raster' (small r) object. NOT TO BE CONFUSED with the Raster\* (big R) objects defined by the raster package! Such objects can be used for plotting with the `rasterImage` function.

### Usage

```
as.raster(x, ...)
```

### Arguments

x RasterLayer object  
 ... Additional arguments.  
 maxpixels Integer. To regularly subsample very large objects  
 col Vector of colors. Default is `col=rev(terrain.colors(255))`

### Value

'raster' object

### Examples

```
r <- raster(ncol=3, nrow=3)
values(r) <- 1:ncell(r)
as.raster(r)
```

---

atan2 *Two argument arc-tangent*

---

### Description

For Raster\* objects x and y, `atan2(y, x)` returns the angle in radians for the tangent y/x, handling the case when x is zero. See [Trig](#)

See [Math-methods](#) for other trigonometric and mathematical functions that can be used with Raster\* objects.

### Usage

```
atan2(y, x)
```

**Arguments**

y	Raster* object
x	Raster* object

**See Also**

[Math-methods](#)

**Examples**

```
r1 <- r2 <- raster(nrow=10, ncol=10)
values(r1) <- (runif(ncell(r1))-0.5) * 10
values(r2) <- (runif(ncell(r1))-0.5) * 10
atan2(r1, r2)
```

---

autocorrelation	<i>Spatial autocorrelation</i>
-----------------	--------------------------------

---

**Description**

Compute Moran's I or Geary's C measures of global spatial autocorrelation in a RasterLayer, or compute the the local Moran or Geary index (Anselin, 1995).

**Usage**

```
Geary(x, w=matrix(c(1,1,1,1,0,1,1,1,1), 3,3))
Moran(x, w=matrix(c(1,1,1,1,0,1,1,1,1), 3,3))
MoranLocal(x, w=matrix(c(1,1,1,1,0,1,1,1,1), 3,3))
GearyLocal(x, w=matrix(c(1,1,1,1,0,1,1,1,1), 3,3))
```

**Arguments**

x	RasterLayer
w	Spatial weights defined by or a rectangular matrix with odd length (3, 5, ...) sides (as in <a href="#">focal</a> )

**Details**

The default setting uses a 3x3 neighborhood to compute "Queen's case" indices. You can use a filter (weights matrix) to do other things, such as "Rook's case", or different lags.

**Value**

A single value (Moran's I or Geary's C) or a RasterLayer (Local Moran or Geary values)

**Author(s)**

Robert J. Hijmans and Babak Naimi

## References

- Moran, P.A.P., 1950. Notes on continuous stochastic phenomena. *Biometrika* 37:17-23
- Geary, R.C., 1954. The contiguity ratio and statistical mapping. *The Incorporated Statistician* 5: 115-145
- Anselin, L., 1995. Local indicators of spatial association-LISA. *Geographical Analysis* 27:93-115  
[http://en.wikipedia.org/wiki/Indicators\\_of\\_spatial\\_association](http://en.wikipedia.org/wiki/Indicators_of_spatial_association)

## See Also

The `spdep` package for additional and more general approaches for computing indices of spatial autocorrelation

## Examples

```
r <- raster(nrows=10, ncols=10)
values(r) <- 1:ncell(r)

Moran(r)
# Rook's case
f <- matrix(c(0,1,0,1,0,1,0,1,0), nrow=3)
Moran(r, f)

Geary(r)

x1 <- MoranLocal(r)

# Rook's case
x2 <- MoranLocal(r, w=f)
```

---

bands

*Number of bands*

---

## Description

A 'band' refers to a single layer for a possibly multi-layer file. Most `RasterLayer` objects will refer to files with a single layer. The term 'band' is frequently used in remote sensing to refer to a variable (layer) in a multi-variable dataset as these variables typically represent reflection in different bandwidths in the electromagnetic spectrum. But in that context, bands could be stored in a single or in separate files. In the context of the raster package, the term band is equivalent to a layer in a raster file.

`nbands` returns the number of bands of the file that a `RasterLayer` points to (and 1 if it does not point at any file). This function also works for a `RasterStack` for which it is equivalent to `nlayers`.

`band` returns the specific band the `RasterLayer` refers to (1 if the `RasterLayer` points at single layer file or does not point at any file).



**Usage**

```
nbands(x)
bandnr(x, ...)
```

**Arguments**

```
x          RasterLayer
...        Additional arguments (none at this time)
```

**Value**

```
numeric >= 1
```

**See Also**

[nlayers](#)

**Examples**

```
f <- system.file("external/rlogo.grd", package="raster")
r <- raster(f, layer=2)
nbands(r)
bandnr(r)
```

---

barplot

*Bar plot of a RasterLayer*


---

**Description**

Create a barplot of the values of a RasterLayer. For large datasets a regular sample with a size of approximately `maxpixels` is used.

**Usage**

```
## S4 method for signature 'RasterLayer'
barplot(height, maxpixels=1000000, digits=0, breaks=NULL, col=rainbow, ...)
```

**Arguments**

```
height      RasterLayer
maxpixels   integer. To regularly subsample very large objects
digits      integer used to determine how to round the values before tabulating. Set to NULL
            or to a large number if you do not want any rounding
breaks      breaks used to group the data as in cut
col         a color generating function such as rainbow, or a vector of colors
...         additional arguments for plotting as in barplot
```

**Value**

A numeric vector (or matrix, when `beside = TRUE`) of the coordinates of the bar midpoints, useful for adding to the graph. See [barplot](#)

**See Also**

[hist](#), [boxplot](#)

**Examples**

```
f <- system.file("external/test.grd", package="raster")
r <- raster(f)
barplot(r, digits=-2, las=2, ylab='Frequency')

op <- par(no.readonly = TRUE)
par(mai = c(1, 2, .5, .5))
barplot(r, breaks=10, col=c('red', 'blue'), horiz=TRUE, digits=NULL, las=1)
par(op)
```

---

 bind

*Bind Spatial\* objects*


---

**Description**

Bind (append) `Spatial*` objects into a single object. All objects must be of the same vector type base class (`SpatialPoints`, `SpatialLines`, or `SpatialPolygons`)

**Usage**

```
## S4 method for signature 'SpatialPolygons,SpatialPolygons'
bind(x, y, ..., keepnames=FALSE)

## S4 method for signature 'SpatialLines,SpatialLines'
bind(x, y, ..., keepnames=FALSE)

## S4 method for signature 'SpatialPoints,SpatialPoints'
bind(x, y, ..., keepnames=FALSE)

## S4 method for signature 'data.frame,data.frame'
bind(x, y, ..., variables=NULL)

## S4 method for signature 'list,missing'
bind(x, y, ..., keepnames=FALSE)
```

**Arguments**

x	Spatial* object or data.frame, or a list of Spatial* objects
y	Spatial* object or data.frame, or missing
...	Additional Spatial* objects
keepnames	Logical. If TRUE the row.names are kept (if unique)
variables	character. Variable (column) names to keep, If NULL, all variables are kept

**Value**

Spatial\* object

**See Also**

[merge](#)

**Examples**

```
p <- readRDS(system.file("external/lux.rds", package="raster"))
mersch <- p[p$NAME_2=='Mersch', ]
diekirch <- p[p$NAME_2=='Diekirch', ]
remich <- p[p$NAME_2=='Remich', ]
remich$NAME_1 <- NULL
x <- bind(mersch, diekirch, remich)
plot(x)
data.frame(x)
```

---

blockSize

*Block size for writing files*

---

**Description**

This function can be used to suggest chunk sizes (always a number of entire rows), and corresponding row numbers, to be used when processing Raster\* objects in chunks. Normally used together with [writeValues](#).

**Usage**

```
blockSize(x, chunksize, n=nlayers(x), minblocks=4, minrows=1)
```

**Arguments**

x	Raster* object
chunksize	Integer, normally missing. Can be used to set the block size; unit is number of cells. Block size is then computed in units of number of rows (always >= 1)
n	Integer. number of layers to consider. The function divides chunksize by n to determine blocksize
minblocks	Integer. Minimum number of blocks
minrows	Integer. Minimum number of rows in each block

**Value**

A list with three elements:  
 rows, the suggested row numbers at which to start the blocks for reading and writing,  
 nrows, the number of rows in each block, and,  
 n, the total number of blocks

**See Also**

[writeValues](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
blockSize(r)
```

---

boundaries	<i>boundaries (edges) detection</i>
------------	-------------------------------------

---

**Description**

Detect boundaries (edges). boundaries are cells that have more than one class in the 4 or 8 cells surrounding it, or, if classes=FALSE, cells with values and cells with NA.

**Usage**

```
## S4 method for signature 'RasterLayer'
boundaries(x, type='inner', classes=FALSE, directions=8, asNA=FALSE, filename="", ...)
```

**Arguments**

x	RasterLayer object
type	character. 'inner' or 'outer'
classes	character. Logical. If TRUE all different values are (after rounding) distinguished, as well as NA. If FALSE (the default) only edges between NA and non-NA cells are considered
directions	integer. Which cells are considered adjacent? Should be 8 (Queen's case) or 4 (Rook's case)
asNA	logical. If TRUE, non-edges are returned as NA instead of zero
filename	character. Filename for the output RasterLayer (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer. Cell values are either 1 (a border) or 0 (not a border), or NA

**See Also**[focal](#), [clump](#)**Examples**

```
r <- raster(nrow=18, ncol=36, xmn=0)
r[150:250] <- 1
r[251:450] <- 2
plot( boundaries(r, type='inner') )
plot( boundaries(r, type='outer') )
plot( boundaries(r, classes=TRUE) )
```

boxplot

*Box plot of Raster objects***Description**

Box plot of layers in a Raster object

**Usage**

```
## S4 method for signature 'RasterStackBrick'
boxplot(x, maxpixels=100000, ...)
```

```
## S4 method for signature 'RasterLayer'
boxplot(x, y=NULL, maxpixels=100000, ...)
```

**Arguments**

x	Raster* object
y	If x is a RasterLayer object, y can be an additional RasterLayer to group the values of x by 'zone'
maxpixels	Integer. Number of pixels to sample from each layer of large Raster objects
...	Arguments passed to <code>graphics::boxplot</code>

**See Also**[pairs](#), [hist](#)**Examples**

```
r1 <- r2 <- r3 <- raster(ncol=10, nrow=10)
values(r1) <- rnorm(ncell(r1), 100, 40)
values(r2) <- rnorm(ncell(r1), 80, 10)
values(r3) <- rnorm(ncell(r1), 120, 30)
s <- stack(r1, r2, r3)
names(s) <- c('A', 'B', 'C')

boxplot(s, notch=TRUE, col=c('red', 'blue', 'orange'), main='Box plot', ylab='random' )
```

---

brick

---

*Create a RasterBrick object*


---

### Description

A RasterBrick is a multi-layer raster object. They are typically created from a multi-layer (band) file; but they can also exist entirely in memory. They are similar to a RasterStack (that can be created with [stack](#)), but processing time should be shorter when using a RasterBrick. Yet they are less flexible as they can only point to a single file.

A RasterBrick can be created from RasterLayer objects, from a RasterStack, or from a (multi-layer) file. They can also be created from SpatialPixels\*, SpatialGrid\*, and Extent objects, and from a three-dimensional array.

### Usage

```
## S4 method for signature 'character'
brick(x, ...)

## S4 method for signature 'RasterStack'
brick(x, values=TRUE, nl, filename='', ...)

## S4 method for signature 'RasterBrick'
brick(x, nl, ...)

## S4 method for signature 'RasterLayer'
brick(x, ..., values=TRUE, nl=1, filename='')

## S4 method for signature 'missing'
brick(nrows=180, ncols=360, xmn=-180, xmx=180, ymn=-90, ymx=90, nl=1, crs)

## S4 method for signature 'Extent'
brick(x, nrows=10, ncols=10, crs="", nl=1)

## S4 method for signature 'array'
brick(x, xmn=0, xmx=1, ymn=0, ymx=1, crs="", transpose=FALSE)

## S4 method for signature 'SpatialGrid'
brick(x)

## S4 method for signature 'SpatialPixels'
brick(x)
```

### Arguments

x character (filename, see Details); Raster\* object; missing; array; SpatialGrid\*; SpatialPixels\*; Extent; or list of Raster\* objects. Supported file types are the

	'native' raster package format and those that can be read via rgdal (see <a href="#">readGDAL</a> ), and NetCDF files (see details)
...	see Details
values	logical. If TRUE, the cell values of 'x' are copied to the RasterBrick object that is returned
n1	integer > 0. How many layers should the RasterBrick have?
filename	character. Filename if you want the RasterBrick to be saved on disk
nrows	integer > 0. Number of rows
ncols	integer > 0. Number of columns
xmn	minimum x coordinate (left border)
xmx	maximum x coordinate (right border)
ymn	minimum y coordinate (bottom border)
ymx	maximum y coordinate (top border)
crs	character or object of class CRS. PROJ4 type description of a Coordinate Reference System (map projection). If this argument is missing, and the x coordinates are within -360 .. 360 and the y coordinates are within -90 .. 90, "+proj=longlat +datum=WGS84" is used
transpose	if TRUE, the values in the array are transposed

## Details

If x is a RasterLayer, the additional arguments can be used to pass additional Raster\* objects.

If there is a filename argument, the additional arguments are as for [writeRaster](#).

If x represents a filename there is the following additional argument:

native: logical. If TRUE (not the default), reading and writing of IDRISI, BIL, BSQ, BIP, and Arc ASCII files is done with native (raster package) drivers, rather than via rgdal.

In addition, if x is a **NetCDF** filename there are the following additional arguments:

varname: character. The variable name (e.g. 'altitude' or 'precipitation'. If not supplied and the file has multiple variables a guess will be made (and reported))

lvar: integer > 0 (default=3). To select the 'level variable' (3rd dimension variable) to use, if the file has 4 dimensions (e.g. depth instead of time)

level: integer > 0 (default=1). To select the 'level' (4th dimension variable) to use, if the file has 4 dimensions, e.g. to create a RasterBrick of weather over time at a certain height.

dims: integer vector to indicated the order of the dimensions. Default is dims=c(1,2,3) (rows, cols, time).

To use NetCDF files the ncd4 package needs to be available. It is assumed that these files follow, or are compatible with the CF-1 convention.

## Value

RasterBrick

**See Also**[raster](#)**Examples**

```
b <- brick(system.file("external/rlogo.grd", package="raster"))
b
nlayers(b)
names(b)
extract(b, 870)
```

buffer

*buffer***Description**

Calculate a buffer around all cells that are not NA or around SpatialPoints, Lines, or Polygons.

Note that the distance unit of the buffer width parameter is meters if the RasterLayer is not projected (+proj=longlat), and in map units (typically also meters) when it is projected.

Except for SpatialLines and SpatialPolygons that are currently handled by rgeos, and can only deal with planar coordinate reference systems.

**Usage**

```
## S4 method for signature 'RasterLayer'
buffer(x, width=0, filename='', doEdge=FALSE, ...)
```

```
## S4 method for signature 'Spatial'
buffer(x, width=1, dissolve=TRUE, ...)
```

**Arguments**

x	RasterLayer or Spatial* object
width	numeric > 0. Unit is meter if x has a longitude/latitude CRS, or mapunits in other cases
filename	character. Filename for the output RasterLayer (optional)
doEdge	logical. If TRUE, the <a href="#">boundaries</a> function is called first. This may be efficient in cases where you compute a buffer around very large areas because boundaries determines the edge cells that matter for distance computation
dissolve	logical. If TRUE, buffer geometries of overlapping polygons are dissolved and all geometries are aggregated and attributes (the data.frame) are dropped
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer or SpatialPolygons\* object



**See Also**

[distance](#), [gridDistance](#), [pointDistance](#)

**Examples**

```
r <- raster(ncol=36,nrow=18)
values(r) <- NA
r[500] <- 1
b <- buffer(r, width=5000000)
#plot(b)
```

---

 calc

*Calculate*


---

**Description**

Calculate values for a new Raster\* object from another Raster\* object, using a formula.

If *x* is a RasterLayer, *fun* is typically a function that can take a single vector as input, and return a vector of values of the same length (e.g. `sqrt`). If *x* is a RasterStack or RasterBrick, *fun* should operate on a vector of values (one vector for each cell). `calc` returns a RasterLayer if *fun* returns a single value (e.g. `sum`) and it returns a RasterBrick if *fun* returns more than one number, e.g., `fun=quantile`.

In many cases, what can be achieved with `calc`, can also be accomplished with a more intuitive 'raster-algebra' notation (see [Arith-methods](#)). For example, `r <- r * 2` instead of

```
r <- calc(r, fun=function(x){x * 2}, or r <- sum(s) instead of
```

```
r <- calc(s, fun=sum). However, calc should be faster when using complex formulas on large datasets. With calc it is possible to set an output filename and file type preferences.
```

See ([overlay](#)) to use functions that refer to specific layers, like `(function(a,b,c){a + sqrt(b) / c})`

**Usage**

```
## S4 method for signature 'Raster,function'
calc(x, fun, filename='', na.rm, forcefun=FALSE, forceapply=FALSE, ...)
```

**Arguments**

<code>x</code>	Raster* object
<code>fun</code>	function
<code>filename</code>	character. Output filename (optional)
<code>na.rm</code>	Remove NA values, if supported by 'fun' (only relevant when summarizing a multilayer Raster object into a RasterLayer)
<code>forcefun</code>	logical. Force <code>calc</code> to not use <code>fun</code> with <code>apply</code> ; for use with ambiguous functions and for debugging (see <a href="#">Details</a> )
<code>forceapply</code>	logical. Force <code>calc</code> to use <code>fun</code> with <code>apply</code> ; for use with ambiguous functions and for debugging (see <a href="#">Details</a> )
<code>...</code>	Additional arguments as for <a href="#">writeRaster</a>

**Details**

The intent of some functions can be ambiguous. Consider:

```
library(raster)
r <- raster(volcano)
calc(r, function(x) x * 1:10)
```

In this case, the cell values are multiplied in a vectorized manner and a single layer is returned where the first cell has been multiplied with one, the second cell with two, the 11th cell with one again, and so on. But perhaps the intent was to create 10 new layers ( $x*1, x*2, \dots$ )? This can be achieved by using argument `forceapply=TRUE`

```
calc(r, function(x) x * 1:10), forceapply=TRUE
```

**Value**

a Raster\* object

**Note**

For large objects `calc` will compute values chunk by chunk. This means that for the result of `fun` to be correct it should not depend on having access to `_all_` values at once. For example, to scale the values of a Raster\* object by subtracting its mean value (for each layer), you would `_not_` do, for Raster object `x`:

```
calc(x, function(x) scale(x, scale=FALSE))
```

Because the mean value of each chunk will likely be different. Rather do something like

```
m <- cellStats(x, 'mean')
x - m
```

**Author(s)**

Robert J. Hijmans and Matteo Mattiuzzi

**See Also**

[overlay](#), [reclassify](#), [Arith-methods](#), [Math-methods](#)

**Examples**

```
r <- raster(ncols=36, nrows=18)
values(r) <- 1:ncell(r)

# multiply values with 10
fun <- function(x) { x * 10 }
rc1 <- calc(r, fun)

# set values below 100 to NA.
fun <- function(x) { x[x<100] <- NA; return(x) }
rc2 <- calc(r, fun)
```

```

# set NA values to -9999
fun <- function(x) { x[is.na(x)] <- -9999; return(x)}
rc3 <- calc(rc2, fun)

# using a RasterStack as input
s <- stack(r, r*2, sqrt(r))
# return a RasterLayer
rs1 <- calc(s, sum)

# return a RasterBrick
rs2 <- calc(s, fun=function(x){x * 10})
# recycling by layer
rs3 <- calc(s, fun=function(x){x * c(1, 5, 10)})

# use overlay when you want to refer to individual layer in the function
# but it can be done with calc:
rs4 <- calc(s, fun=function(x){x[1]+x[2]*x[3]})

##
# Some regression examples
##

# create data
r <- raster(nrow=10, ncol=10)
s1 <- lapply(1:12, function(i) setValues(r, rnorm(ncell(r), i, 3)))
s2 <- lapply(1:12, function(i) setValues(r, rnorm(ncell(r), i, 3)))
s1 <- stack(s1)
s2 <- stack(s2)

# regression of values in one brick (or stack) with another
s <- stack(s1, s2)
# s1 and s2 have 12 layers; coefficients[2] is the slope
fun <- function(x) { lm(x[1:12] ~ x[13:24])$coefficients[2] }
x1 <- calc(s, fun)

# regression of values in one brick (or stack) with 'time'
time <- 1:nlayers(s)
fun <- function(x) { lm(x ~ time)$coefficients[2] }
x2 <- calc(s, fun)

# get multiple layers, e.g. the slope _and_ intercept
fun <- function(x) { lm(x ~ time)$coefficients }
x3 <- calc(s, fun)

### A much (> 100 times) faster approach is to directly use
### linear algebra and pre-compute some constants

## add 1 for a model with an intercept
X <- cbind(1, time)

## pre-computing constant part of least squares
invXtX <- solve(t(X) %*% X) %*% t(X)

```

```
## much reduced regression model; [2] is to get the slope
quickfun <- function(y) (invXtX %*% y)[2]
x4 <- calc(s, quickfun)
```

---

cellFrom                      *Get cell, row, or column number*

---

### Description

Get cell number(s) of a Raster\* object from row and/or column numbers. Cell numbers start at 1 in the upper left corner, and increase from left to right, and then from top to bottom. The last cell number equals the number of cells of the Raster\* object.

### Usage

```
cellFromRowCol(object, row, col, ...)
cellFromRowColCombine(object, row, col, ...)
cellFromRow(object, rownr)
cellFromCol(object, colnr)
colFromX(object, x)
rowFromY(object, y)
cellFromXY(object, xy)
cellFromLine(object, lns)
cellFromPolygon(object, p, weights=FALSE)
fourCellsFromXY(object, xy, duplicates=TRUE)
```

### Arguments

object	Raster* object (or a SpatialPixels* or SpatialGrid* object)
colnr	column number; or vector of column numbers
rownr	row number; or vector of row numbers
col	column number; or vector of column numbers
row	row number; or vector of row numbers
x	x coordinate(s)
y	y coordinate(s)
xy	matrix of x and y coordinates, or a SpatialPoints or SpatialPointsDataFrame object
lns	SpatialLines object
p	SpatialPolygons object
weights	Logical. If TRUE, the fraction of each cell that is covered is also returned
duplicates	Logical. If TRUE, the same cell number can be returned twice (if the point in the middle of a division between two cells) or four times (if a point is in the center of a cell)
...	additional arguments (none implemented)

**Details**

cellFromRowCol returns the cell numbers obtained for each row / col number pair. In contrast, cellFromRowColCombine returns the cell numbers obtained by the combination of all row and column numbers supplied as arguments.

fourCellsFromXY returns the four cells that are nearest to a point (if the point falls on the raster). Also see [adjacent](#).

**Value**

vector of row, column or cell numbers. cellFromLine and cellFromPolygon return a list, fourCellsFromXY returns a matrix.

**See Also**

[xyFromCell](#), [cellsFromExtent](#), [rowColFromCell](#)

**Examples**

```
r <- raster(ncols=10, nrows=10)
cellFromRowCol(r, 5, 5)
cellFromRowCol(r, 1:2, 1:2)
cellFromRowColCombine(r, 1:3, 1:2)
cellFromCol(r, 1)
cellFromRow(r, 1)

colFromX(r, 0.5)
rowFromY(r, 0.5)
cellFromXY(r, cbind(c(0.5,5), c(15, 88)))
fourCellsFromXY(r, cbind(c(0.5,5), c(15, 88)))

cds1 <- rbind(c(-180,-20), c(-160,5), c(-60, 0), c(-160,-60), c(-180,-20))
cds2 <- rbind(c(80,0), c(100,60), c(120,0), c(120,-55), c(80,0))
pols <- SpatialPolygons(list(Polygons(list(Polygon(cds1)), 1), Polygons(list(Polygon(cds2)), 2)))
cellFromPolygon(r, pols)
```

---

cellsFromExtent

*Cells from extent, and vice versa*

---

**Description**

cellsFromExtent returns the cell numbers for a Raster\* object that are within a specified extent (rectangular area), supply an object of class Extent, or another Raster\* object.

extentFromCells returns an Extent object from a Raster\* object and cell numbers. All cells are within the returned Extent.

**Usage**

```
cellsFromExtent(object, extent, expand=FALSE)
extentFromCells(object, cells)
```

**Arguments**

object	A Raster* object
extent	An object of class Extent (which you can create with newExtent(), or another Raster* object )
expand	Logical. If TRUE, NA is returned for (virtual) cells implied by bndbox, that are outside the RasterLayer (object). If FALSE, only cell numbers for the area where object and bndbox overlap are returned (see <a href="#">intersect</a> )
cells	numeric. A vector of cell numbers

**Value**

a vector of cell numbers

**See Also**

[extent](#), [cellFromXY](#)

**Examples**

```
r <- raster()
bb <- extent(-5, 5, -5, 5)
cells <- cellsFromExtent(r, bb)
r <- crop(r, bb)
values(r) <- cells

e <- extentFromCells(r, 50:55)
```

---

cellStats

*Statistics across cells*

---

**Description**

Compute statistics for the cells of each layer of a Raster\* object. In the raster package, functions such as max, min, and mean, when used with Raster\* objects as argument, return a new Raster\* object (with a value computed for each cell). In contrast, cellStats returns a single value, computed from the all the values of a layer. Also see [layerStats](#)

**Usage**

```
## S4 method for signature 'RasterLayer'
cellStats(x, stat='mean', na.rm=TRUE, asSample=TRUE, ...)

## S4 method for signature 'RasterStackBrick'
cellStats(x, stat='mean', na.rm=TRUE, asSample=TRUE, ...)
```

**Arguments**

x	Raster* object
stat	The function to be applied. See Details
na.rm	Logical. Should NA values be removed?
asSample	Logical. Only relevant for stat=sd in which case, if TRUE, the standard deviation for a sample (denominator is n-1) is computed, rather than for the population (denominator is n)
...	Additional arguments

**Details**

cellStats will fail (gracefully) for very large Raster\* objects except for a number of known functions: sum, mean, min, max, sd, 'skew' and 'rms'. 'skew' (skewness) and 'rms' (Root Mean Square) must be supplied as a character value (with quotes), the other known functions may be supplied with or without quotes. For other functions you could perhaps use a sample of the RasterLayer that can be held in memory (see [sampleRegular](#) )

**Value**

Numeric

**See Also**

[freq](#), [quantile](#), [minValue](#), [maxValue](#), [setMinMax](#)

**Examples**

```
r <- raster(nrow=18, ncol=36)
values(r) <- runif(ncell(r)) * 10
# works for large files
cellStats(r, 'mean')
# same, but does not work for very large files
cellStats(r, mean)
# multi-layer object
cellStats(brick(r,r), mean)
```

---

clamp

*Clamp values*

---

**Description**

Clamp values to a minimum and maximum value. That is, all values below the lower clamp value and above the upper clamp value become NA (or the lower/upper value if useValue=TRUE)

**Usage**

```
## S4 method for signature 'Raster'
clamp(x, lower=-Inf, upper=Inf, useValues=TRUE, filename="", ...)
## S4 method for signature 'numeric'
clamp(x, lower=-Inf, upper=Inf, ...)
```

**Arguments**

x	RasterLayer, or numeric vector
lower	numeric. lowest value
upper	numeric. highest value
useValues	logical. If FALSE values outside the clamping range become NA, if TRUE, they get the extreme values
filename	character. Filename for the output RasterLayer (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

Raster object

**See Also**

[reclassify](#)

**Examples**

```
r <- raster(ncols=12, nrows=12)
values(r) <- 1:ncell(r)
rc <- clamp(r, 25, 75)
rc
```

---

clearValues

*Clear values*

---

**Description**

Clear cell values of a Raster\* object from memory

**Usage**

```
clearValues(x)
```

**Arguments**

x	Raster* object
---	----------------



**Value**

a Raster\* object

**See Also**

[values](#), [replacement](#)

**Examples**

```
r <- raster(ncol=10, nrow=10)
values(r) <- 1:ncell(r)
r <- clearValues(r)
```

---

click

*Query by clicking on a map*

---

**Description**

Click on a map (plot) to get values of a Raster\* or Spatial\* object at that location; and optionally the coordinates and cell number of the location. For SpatialLines and SpatialPoints you need to click twice (draw a box).

**Usage**

```
## S4 method for signature 'Raster'
click(x, n=Inf, id=FALSE, xy=FALSE, cell=FALSE, type="n", show=TRUE, ...)
```

```
## S4 method for signature 'SpatialGrid'
click(x, n=1, id=FALSE, xy=FALSE, cell=FALSE, type="n", ...)
```

```
## S4 method for signature 'SpatialPolygons'
click(x, n=1, id=FALSE, xy=FALSE, type="n", ...)
```

```
## S4 method for signature 'SpatialLines'
click(x, ...)
```

```
## S4 method for signature 'SpatialPoints'
click(x, ...)
```

**Arguments**

x	Raster*, or Spatial* object (or missing)
n	number of clicks on the map
id	Logical. If TRUE, a numeric ID is shown on the map that corresponds to the row number of the output

xy	Logical. If TRUE, xy coordinates are included in the output
cell	Logical. If TRUE, cell numbers are included in the output
type	One of "n", "p", "l" or "o". If "p" or "o" the points are plotted; if "l" or "o" they are joined by lines. See ?locator
show	logical. Print the values after each click?
...	additional graphics parameters used if type != "n" for plotting the locations. See ?locator

**Value**

The value(s) of x at the point(s) clicked on (or touched by the box drawn).

**Note**

The plot only provides the coordinates for a spatial query, the values are read from the Raster\* or Spatial\* object that is passed as an argument. Thus you can extract values from an object that has not been plotted, as long as it spatially overlaps with with the extent of the plot.

Unless the process is terminated prematurely values at at most n positions are determined. The identification process can be terminated by clicking the second mouse button and selecting 'Stop' from the menu, or from the 'Stop' menu on the graphics window.

**See Also**

[select, drawExtent](#)

**Examples**

```
## Not run:
r <- raster(system.file("external/test.grd", package="raster"))
plot(r)
click(r)
# now click on the plot (map)

## End(Not run)
```

---

clump

*Detect clumps*

---

**Description**

Detect clumps (patches) of connected cells. Each clump gets a unique ID. NA and zero are used as background values (i.e. these values are used to separate clumps). You can use queen's or rook's case, using the directions argument. For larger files that are processed in chunks, the highest clump number is not necessarily equal to the number of clumps (unless you use argument gaps=FALSE).

**Usage**

```
## S4 method for signature 'RasterLayer'  
clump(x, filename="", directions=8, gaps=TRUE, ...)
```

**Arguments**

x	RasterLayer
filename	Character. Filename for the output RasterLayer (optional)
directions	Integer. Which cells are considered adjacent? Should be 8 (Queen's case) or 4 (Rook's case)
gaps	Logical. If TRUE (the default), there may be 'gaps' in the chunk numbers (e.g. you may have clumps with IDs 1, 2, 3 and 5, but not 4). If it is FALSE, these numbers will be recoded from 1 to n (4 in this example)
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer

**Note**

This function requires that the `igraph` package is available.

**Author(s)**

Robert J. Hijmans and Jacob van Etten

**Examples**

```
r <- raster(ncols=12, nrows=12)  
set.seed(0)  
values(r) <- round(runif(ncell(r))*0.7 )  
rc <- clump(r)  
freq(rc)  
plot(rc)
```

---

cluster

*Use a multi-core cluster*

---

**Description**

`beginCluster` creates, and `endCluster` deletes a 'snow' cluster object. This object can be used for multi-core computing with those 'raster' functions that support it.

`beginCluster` determines the number of nodes (cores) that are available and uses all of them (unless the argument `n` is used).

NOTE: `beginCluster` may fail when the package `'nws'` is installed. You can fix that by removing the `'nws'` package, or by setting the cluster type manually, e.g. `beginCluster(type="SOCK")`

`endCluster` closes the cluster and removes the object.

The use of the cluster is automatic in these functions: `projectRaster`, `resample` and in `extract` when using polygons.

`clusterR` is a flexible interface for using cluster with other functions. This function only works with functions that have a `Raster*` object as first argument and that operate on a cell by cell basis (i.e., there is no effect of neighboring cells) and return an object with the same number of cells as the input raster object. The first argument of the function called must be a `Raster*` object. There can only be one `Raster*` object argument. For example, it works with `calc` and it also works with `overlay` as long as you provide a single `RasterStack` or `RasterBrick` as the first argument.

This function is particularly useful to speed up computations in functions like `predict`, `interpolate`, and perhaps `calc`.

Among other functions, it does not work with `merge`, `crop`, `mosaic`, `(dis)aggregate`, `resample`, `projectRaster`, `focal`, `distance`, `buffer`, `direction`. But note that `projectRaster` has a build-in capacity for clustering that is automatically used if `beginCluster()` has been called.

## Usage

```
beginCluster(n, type='SOCK', nice, exclude)
endCluster()
clusterR(x, fun, args=NULL, export=NULL, filename='', cl=NULL, m=2, ...)
```

## Arguments

<code>n</code>	Integer. The number of nodes to be used (optional)
<code>type</code>	Character. The cluster type to be used
<code>nice</code>	Integer. To set the priority for the workers, between -20 and 20 (UNIX like platforms only)
<code>exclude</code>	Character. Packages to exclude from loading on the nodes (because they may fail there) but are required/loaded on the master
<code>x</code>	<code>Raster*</code> object
<code>fun</code>	function that takes <code>x</code> as its first argument
<code>args</code>	list with the arguments for the function (excluding <code>x</code> , which should always be the first argument)
<code>export</code>	character. Vector of variable names to export to the cluster nodes such that the are visible to fun (e.g. a parameter that is not passed as an argument)
<code>filename</code>	character. Output filename (optional)
<code>cl</code>	cluster object (do not use it if <code>beginCluster()</code> has been called)
<code>m</code>	tuning parameter to determine how many blocks should be used. The number is rounded and multiplied with the number of nodes.
<code>...</code>	additional arguments as for <code>writeRaster</code>

**Value**

`beginCluster` and `endCluster`: None. The side effect is to create or delete a cluster object.  
`clusterR`: as for the function called with argument `fun`

**Note**

If you want to write your own cluster-enabled functions see [getCluster](#), [returnCluster](#), and the vignette about writing functions.

**Author(s)**

Matteo Mattiuzzi and Robert J. Hijmans

**Examples**

```
## Not run:
# set up the cluster object for parallel computing
beginCluster()

r <- raster()
values(r) <- 1:ncell(r)

x <- clusterR(r, sqrt, verbose=T)

f1 <- function(x) calc(x, sqrt)
y <- clusterR(r, f1)

s <- stack(r, r*2, r*3)
f2 <- function(d,e,f) (d + e) / (f * param)
param <- 122
ov <- clusterR(s, overlay, args=list(fun=f2), export='param')

pts <- matrix(c(0,0, 45,45), ncol=2, byrow=T)
d <- clusterR(r, distanceFromPoints, args=list(xy=pts))

values(r) <- runif(ncell(r))
m <- c(0, 0.25, 1, 0.25, 0.5, 2, 0.5, 1, 3)
m <- matrix(m, ncol=3, byrow=TRUE)
rc1 <- clusterR(r, reclassify, args=list(rcl=m, right=FALSE),
               filename=rasterTmpFile(), datatype='INT2S', overwrite=TRUE)

# equivalent to:
rc2 <- reclassify(r, rcl=m, right=FALSE, filename=rasterTmpFile(), datatype='INT2S', overwrite=TRUE)

# example with the calc function
a <- 10
f3 <- function(x) sum(x)+a

z1 <- clusterR(s, calc, args=list(fun=f3), export='a')
```

```

# for some raster functions that use another function as an argument
# you can write your own parallel function instead of using clusterR
# get cluster object created with beginCluster
cl <- getCluster()

library(parallel)
clusterExport(cl, "a")
z2 <- calc(s, fun=function(x){ parApply(cl, x, 1, f3)} )
# set flag that cluster is available again
returnCluster()
#

# done with cluster object
endCluster()

## End(Not run)

```

---

colortable

*colortable*


---

## Description

Get or set the colortable of a RasterLayer. A colortable is a vector of 256 colors in the RGB triple format as returned by the [rgb](#) function (e.g. "#C4CDDA").

When setting the colortable, it is assumed that the values are integers in the range [0,255]

## Usage

```

colortable(x)
colortable(x) <- value

```

## Arguments

x	RasterLayer object
value	vector of 256 character values

## See Also

[plotRGB](#)

## Examples

```

r <- raster(ncol=10, nrow=10)
values(r) <- sample(0:255, ncell(r), replace=TRUE)
ctab <- sample(rainbow(256))
colortable(r) <- ctab
plot(r)
head(colortable(r))

```

---

 Compare-methods

*Compare Raster\* objects*


---

**Description**

These methods compare the location and resolution of Raster\* objects. That is, they compare their spatial extent, projection, and number of rows and columns.

For BasicRaster objects you can use == and !=, the values returned is a single logical value TRUE or FALSE

For RasterLayer objects, these operators also compare the values associated with the objects, and the result is a RasterLayer object with logical (Boolean) values.

The following methods have been implemented for RasterLayer objects:

==, !=, >, <, <=, >=

**Value**

A logical value or a RasterLayer object, and in some cases the side effect of a new file on disk.

**Examples**

```
r1 <- raster()
r1 <- setValues(r1, round(10 * runif(ncell(r1))))
r2 <- setValues(r1, round(10 * runif(ncell(r1))))
as(r1, 'BasicRaster') == as(r2, 'BasicRaster')
r3 <- r1 == r2

b <- extent(0, 360, 0, 180)
r4 <- setExtent(r2, b)
as(r2, 'BasicRaster') != as(r4, 'BasicRaster')
# The following would give an error. You cannot compare RasterLayer
# that do not have the same BasicRaster properties.
#r3 <- r1 > r4
```

---

 compareCRS

*Partially compare two CRS objects*


---

**Description**

Compare CRS objects

**Usage**

```
compareCRS(x, y, unknown=FALSE, verbatim=FALSE, verbose=FALSE)
```

**Arguments**

x	CRS object, or object from which it can be extracted with <a href="#">projection</a> , or PROJ.4 format character string
y	same as x
unknown	logical. Return TRUE if x or y is TRUE
verbatim	logical. If TRUE compare x and y, verbatim (not partially)
verbose	logical. If TRUE, messages about the comparison may be printed

**Value**

logical

**See Also**

`sp::identicalCRS`, [crs](#)

**Examples**

```
compareCRS("+proj=lcc +lat_1=48 +lat_2=33 +lon_0=-100 +ellps=WGS84",
  "+proj=longlat +datum=WGS84")
compareCRS("+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0",
  "+proj=longlat +datum=WGS84")
compareCRS("+proj=longlat +datum=WGS84 +no_defs +ellps=WGS84 +towgs84=0,0,0",
  "+proj=longlat +datum=WGS84", verbatim=TRUE)
compareCRS("+proj=longlat +datum=WGS84", NA)
compareCRS("+proj=longlat +datum=WGS84", NA, unknown=TRUE)
```

---

compareRaster

*Compare Raster objects*

---

**Description**

Evaluate whether a two or more Raster\* objects have the same extent, number of rows and columns, projection, resolution, and origin (or a subset of these comparisons).

`all.equal` is a wrapper around `compareRaster` with options `values=TRUE`, `stopiffalse=FALSE` and `showwarning=TRUE`.

**Usage**

```
compareRaster(x, ..., extent=TRUE, rowcol=TRUE, crs=TRUE, res=FALSE, orig=FALSE,
  rotation=TRUE, values=FALSE, tolerance, stopiffalse=TRUE, showwarning=FALSE)
```



**Arguments**

x	Raster* object
...	Raster* objects
extent	logical. If TRUE, bounding boxes are compared
rowcol	logical. If TRUE, number of rows and columns of the objects are compared
crs	logical. If TRUE, coordinate reference systems are compared.
res	logical. If TRUE, resolutions are compared (redundant when checking extent and rowcol)
orig	logical. If TRUE, origins are compared
rotation	logical. If TRUE, rotations are compared
values	logical. If TRUE, cell values are compared
tolerance	numeric between 0 and 0.5. If not supplied, the default value is used (see <a href="#">rasterOptions</a> . It sets difference (relative to the cell resolution) that is permissible for objects to be considered 'equal', if they have a non-integer origin or resolution. See <a href="#">all.equal</a> .
stopiffalse	logical. If TRUE, an error will occur if the objects are not the same
showwarning	logical. If TRUE, a warning will be given if objects are not the same. Only relevant when stopiffalse is TRUE

**Examples**

```

r1 <- raster()
r2 <- r1
r3 <- r1
compareRaster(r1, r2, r3)
nrow(r3) <- 10

# compareRaster(r1, r3)
compareRaster(r1, r3, stopiffalse=FALSE)
compareRaster(r1, r3, rowcol=FALSE)

all.equal(r1, r2)
all.equal(r1, r3)

```

---

contour

*Contour plot*


---

**Description**

Contour plot of a RasterLayer.

**Usage**

```

## S4 method for signature 'RasterLayer'
contour(x, maxpixels=100000, ...)

```

**Arguments**

x	Raster* object
maxpixels	maximum number of pixels used to create the contours
...	any argument that can be passed to <a href="#">contour</a> (graphics package)

**See Also**

[persp](#), [filledContour](#), [rasterToContour](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
plot(r)
contour(r, add=TRUE)
```

---

corLocal

*Local correlation coefficient*

---

**Description**

Local correlation coefficient for two RasterLayer objects (using a focal neighborhood) or for two RasterStack or Brick objects (with the same number of layers (> 2))

**Usage**

```
## S4 method for signature 'RasterLayer,RasterLayer'
corLocal(x, y, ngb=5,
         method=c("pearson", "kendall", "spearman"), test=FALSE, filename='', ...)
```

```
## S4 method for signature 'RasterStackBrick,RasterStackBrick'
corLocal(x, y,
         method=c("pearson", "kendall", "spearman"), test=FALSE, filename='', ...)
```

**Arguments**

x	RasterLayer or RasterStack/RasterBrick
y	object of the same class as x, and with the same number of layers
ngb	neighborhood size. Either a single integer or a vector of two integers c(nrow, ncol)
method	character indicating which correlation coefficient is to be used. One of "pearson", "kendall", or "spearman"
test	logical. If TRUE, return a p-value
filename	character. Output filename (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer

**Note**

NA values are omitted

**See Also**

[cor](#), [cor.test](#)

**Examples**

```
b <- stack(system.file("external/rlogo.grd", package="raster"))
b <- aggregate(b, 2, mean)

set.seed(0)
b[[2]] <- flip(b[[2]], 'y') + runif(ncell(b))
b[[1]] <- b[[1]] + runif(ncell(b))

x <- corLocal(b[[1]], b[[2]], test=TRUE )
# plot(x)

# only cells where the p-value < 0.1
xm <- mask(x[[1]], x[[2]] < 0.1, maskvalue=FALSE)
plot(xm)

# for global correlation, use the cor function
x <- as.matrix(b)
cor(x, method="spearman")

# use sampleRegular for large datasets
x <- sampleRegular(b, 1000)
cor.test(x[,1], x[,2])

# RasterStack or Brick objects
y <- corLocal(b, flip(b, 'y'))
```

---

cover

*Replace NA values with values of other layers*

---

**Description**

For Raster\* objects: Replace NA values in the first Raster object (x) with the values of the second (y), and so forth for additional Rasters. If x has multiple layers, the subsequent Raster objects should have the same number of layers, or have a single layer only (which will be recycled).

For SpatialPolygons\* objects: Areas of x that overlap with y are replaced by (or intersected with) y.

## Usage

```
## S4 method for signature 'RasterLayer,RasterLayer'  
cover(x, y, ..., filename='')  
  
## S4 method for signature 'RasterStackBrick,Raster'  
cover(x, y, ..., filename='')  
  
## S4 method for signature 'SpatialPolygons,SpatialPolygons'  
cover(x, y, ..., identity=FALSE)
```

## Arguments

x	Raster* or SpatialPolygons* object
y	Same as x
filename	character. Output filename (optional)
...	Same as x. If x is a Raster* object, also additional arguments as for <a href="#">writeRaster</a>
identity	logical. If TRUE overlapping areas are intersected rather than replaced

## Value

RasterLayer or RasterBrick object, or SpatialPolygons object

## Examples

```
# raster objects  
r1 <- raster(ncols=36, nrows=18)  
values(r1) <- 1:ncell(r1)  
r2 <- setValues(r1, runif(ncell(r1)))  
r2[r2 < 0.5] <- NA  
r3 <- cover(r2, r1)  
  
#SpatialPolygons  
if (require(rgdal) & require(rgeos)) {  
  p <- shapefile(system.file("external/lux.shp", package="raster"))  
  b <- as(extent(6, 6.4, 49.75, 50), 'SpatialPolygons')  
  crs(b) <- crs(p)  
  b <- SpatialPolygonsDataFrame(b, data.frame(ID_1=9))  
  
  cv1 <- cover(p, b)  
  cv2 <- cover(p, b, identity=TRUE)  
}
```

---

 crop
*Crop*


---

**Description**

crop returns a geographic subset of an object as specified by an Extent object (or object from which an extent object can be extracted/created). If x is a Raster\* object, the Extent is aligned to x. Areas included in y but outside the extent of x are ignored (see [extend](#) if you want a larger area).

**Usage**

```
## S4 method for signature 'Raster'
crop(x, y, filename="", snap='near', datatype=NULL, ...)
```

```
## S4 method for signature 'Spatial'
crop(x, y, ...)
```

**Arguments**

x	Raster* object or SpatialPolygons*, SpatialLines*, or SpatialPoints* object
y	Extent object, or any object from which an Extent object can be extracted (see <a href="#">Details</a> )
filename	Character, output filename. Optional
snap	Character. One of 'near', 'in', or 'out', for use with <a href="#">alignExtent</a>
datatype	Character. Output <a href="#">dataType</a> (by default it is the same as the input datatype)
...	Additional arguments as for <a href="#">writeRaster</a>

**Details**

Objects from which an Extent can be extracted/created include RasterLayer, RasterStack, RasterBrick and objects of the Spatial\* classes from the sp package. You can check this with the [extent](#) function. New Extent objects can be also be created with function [extent](#) and [drawExtent](#) by clicking twice on a plot.

To crop by row and column numbers you can create an extent like this (for Raster x, row 5 to 10, column 7 to 12) `crop(x, extent(x, 5, 10, 7, 15))`

**Value**

RasterLayer or RasterBrick object; or SpatialLines or SpatialPolygons object.

**Note**

values within the extent of a Raster\* object can be set to NA with [mask](#)

**See Also**

[extend](#), [merge](#)

**Examples**

```

r <- raster(nrow=45, ncol=90)
values(r) <- 1:ncell(r)
e <- extent(-160, 10, 30, 60)
rc <- crop(r, e)

# use row and column numbers:
rc2 <- crop(r, extent(r, 5, 10, 7, 15))

# crop Raster* with Spatial* object
b <- as(extent(6, 6.4, 49.75, 50), 'SpatialPolygons')
crs(b) <- crs(r)
rb <- crop(r, b)

# crop a SpatialPolygon* object with another one
if (require(rgdal) & require(rgeos)) {
  p <- shapefile(system.file("external/lux.shp", package="raster"))
  pb <- crop(p, b)
}

```

crosstab

*Cross-tabulate***Description**

Cross-tabulate two RasterLayer objects, or multiple layers in a RasterStack or RasterBrick to create a contingency table.

**Usage**

```

## S4 method for signature 'Raster,Raster'
crosstab(x, y, digits=0, long=FALSE, useNA=FALSE, progress='', ...)

## S4 method for signature 'RasterStackBrick,missing'
crosstab(x, digits=0, long=FALSE, useNA=FALSE, progress='', ...)

```

**Arguments**

x	Raster* object
y	Raster* object if x is a RasterLayer; Can be missing if x is a RasterStack or RasterBrick
digits	integer. The number of digits for rounding the values before cross-tabulation
long	logical. If TRUE the results are returned in 'long' format data.frame instead of a table
useNA	logical, indicating if the table should include counts of NA values
progress	character. "text", "window", or "" (the default, no progress bar), only for large files that cannot be processed in one step
...	additional arguments. none implemented

**Value**

A table or data.frame

**See Also**

[freq](#), [zonal](#)

**Examples**

```
r <- raster(nc=5, nr=5)
values(r) <- runif(ncell(r)) * 2
s <- setValues(r, runif(ncell(r)) * 3)
crosstab(r,s)

rs <- r/s
r[1:5] <- NA
s[20:25] <- NA
x <- stack(r, s, rs)
crosstab(x, useNA=TRUE, long=TRUE)
```

---

cut

---

*Convert values to classes*


---

**Description**

Cut uses the base function [cut](#) to classify the values of a Raster\* object according to which interval they fall in. The intervals are defined by the argument breaks. The leftmost interval corresponds to level one, the next leftmost to level two and so on.

**Usage**

```
cut(x, ...)
```

**Arguments**

x                    A Raster\* object  
 ...                  additional arguments. See [cut](#)

**Value**

Raster\* object

**See Also**

[subs](#), [reclassify](#), [calc](#)

## Examples

```
r <- raster(ncols=36, nrows=18)
values(r) <- rnorm(ncell(r))
breaks <- -2:2 * 3
rc <- cut(r, breaks=breaks)
```

---

cv	<i>Coefficient of variation</i>
----	---------------------------------

---

## Description

Compute the coefficient of variation (expressed as a percentage). If there is only a single value, sd is NA and cv returns NA if aszero=FALSE (the default). However, if (aszero=TRUE), cv returns 0.

## Usage

```
## S4 method for signature 'ANY'
cv(x, ..., aszero=FALSE, na.rm = FALSE)

## S4 method for signature 'Raster'
cv(x, ..., aszero=FALSE, na.rm = FALSE)
```

## Arguments

x	A vector of numbers (typically integers for modal), or a Raster* object
...	additional (vectors of) numbers, or Raster objects
aszero	logical. If TRUE, a zero is returned (rather than an NA) if the cv of single value is computed
na.rm	Remove (ignore) NA values

## Value

vector or RasterLayer

## Examples

```
data <- c(0,1,2,3,3,3,3,4,4,4,5,5,6,7,7,8,9,NA)
cv(data, na.rm=TRUE)
```



---

datasource	<i>Are values in memory and/or on disk?</i>
------------	---

---

### Description

These are helper functions for programmers and for debugging that provide information about whether a Raster object has associated values, and if these are in memory or on disk.

`fromDisk` is TRUE if the data source is a file on disk; and FALSE if the object only exists in memory.

`inMemory` is TRUE if all values are currently in memory (RAM); and FALSE if not (in which case they either are on disk, or there are no values).

`hasValues` is TRUE if the object has cell values.

### Usage

```
fromDisk(x)
inMemory(x)
## S4 method for signature 'BasicRaster'
hasValues(x, ...)
```

### Arguments

x	Raster* object
...	additional arguments. None implemented

### Value

Logical

### Examples

```
rs <- raster(system.file("external/test.grd", package="raster"))
inMemory(rs)
fromDisk(rs)
rs <- readAll(rs)
inMemory(rs)
fromDisk(rs)
rs <- rs + 1
inMemory(rs)
fromDisk(rs)
rs <- raster(rs)
inMemory(rs)
fromDisk(rs)
rs <- setValues(rs, 1:ncell(rs))
inMemory(rs)
fromDisk(rs)
#rs <- writeRaster(rs, filename=rasterTmpFile(), overwrite=TRUE)
#inMemory(rs)
#fromDisk(rs)
```

---

 dataType

*Data type*


---

### Description

Get the datatype of a RasterLayer object. The datatype determines the interpretation of values written to disk. Changing the datatype of a Raster\* object does not directly affect the way they are stored in memory. For native file formats (.grd/.gri files) it does affect how values are read from file. This is not the case for file formats that are read via rgdal (such as .tif and .img files) or netcdf.

If you change the datatype of a RasterLayer and then read values from a native format file these may be completely wrong, so only do this for debugging or when the information in the header file was wrong. To set the datatype of a new file, you can give a 'datatype' argument to the functions that write values to disk (e.g. `writeRaster`).

### Usage

```
dataType(x)
dataType(x) <- value
```

### Arguments

x	A RasterLayer object
value	A data type (see below)

### Details

Setting the data type is useful if you want to write values to disk. In other cases use functions such as `round()`

Datatypes are described by 5 characters. The first three indicate whether the values are integers, decimal number or logical values. The fourth character indicates the number of bytes used to save the values on disk, and the last character indicates whether the numbers are signed (i.e. can be negative and positive values) or not (only zero and positive values allowed)

The following datatypes are available:

Datatype definition	minimum possible value	maximum possible value
LOG1S	FALSE (0)	TRUE (1)
INT1S	-127	127
INT1U	0	255
INT2S	-32,767	32,767
INT2U	0	65,534
INT4S	-2,147,483,647	2,147,483,647
INT4U	0	4,294,967,296
FLT4S	-3.4e+38	3.4e+38
FLT8S	-1.7e+308	1.7e+308

For all integer types, except the single byte types, the lowest (signed) or highest (unsigned) value is used to store NA. Single byte files do not have NA values. Logical values are stored as signed single byte integers, they do have an NA value (-127)

INT4U is available but they are best avoided as R does not support 32-bit unsigned integers.

### Value

Raster\* object

### Examples

```
r <- raster(system.file("external/test.grd", package="raster"))
dataType(r)
## Not run:
s <- writeRaster(r, 'new.grd', datatype='INT2U', overwrite=TRUE)
dataType(s)

## End(Not run)
```

---

density

*Density plot*

---

### Description

Create density plots of values in a Raster object

### Usage

```
## S4 method for signature 'Raster'
density(x, layer, maxpixels=100000, plot=TRUE, main, ...)
```

### Arguments

x	Raster object
layer	numeric. Can be used to subset the layers to plot in a multilayer object (Raster-Brick or RasterStack)
maxpixels	the maximum number of (randomly sampled) cells to be used for creating the plot
plot	if TRUE produce a plot, else return a density object
main	main title for each plot (can be missing)
...	Additional arguments passed to <a href="#">plot</a>

### Value

density plot (and a density object, returned invisibly if plot=TRUE)

## Examples

```
logo <- stack(system.file("external/rlogo.grd", package="raster"))
density(logo)
```

---

dim *Dimensions of a Raster\* object*

---

## Description

Get or set the number of rows, columns, and layers of a Raster\* object. You cannot use this function to set the dimensions of a RasterStack object.

When setting the dimensions, you can provide a row number, or a vector with the row and the column number (for a RasterLayer and a RasterBrick), or a row and column number and the number of layers (only for a RasterBrick)

## Usage

```
## S4 method for signature 'BasicRaster'
dim(x)
```

## Arguments

x Raster\* object

## Value

Integer or Raster\* object

## See Also

[ncell](#), [extent](#), [res](#)

## Examples

```
r <- raster()
dim(r)
dim(r) <- c(18)
dim(r)
dim(r) <- c(18, 36)
dim(r)
b <- brick(r)
dim(b)
dim(b) <- c(10, 10, 5)
dim(b)
```

---

direction	<i>Direction</i>
-----------	------------------

---

### Description

The direction (azimuth) to or from the nearest cell that is not NA. The direction unit is in radians, unless you use argument `degrees=TRUE`.

### Usage

```
## S4 method for signature 'RasterLayer'
direction(x, filename='', degrees=FALSE, from=FALSE, doEdge=FALSE, ...)
```

### Arguments

<code>x</code>	RasterLayer object
<code>filename</code>	Character. Output filename (optional)
<code>degrees</code>	Logical. If FALSE (the default) the unit of direction is radians.
<code>from</code>	Logical. Default is FALSE. If TRUE, the direction from (instead of to) the nearest cell that is not NA is returned
<code>doEdge</code>	Logical. If TRUE, the <a href="#">boundaries</a> function is called first. This may be efficient in cases where you compute the distance to large blobs. Calling <a href="#">boundaries</a> determines the edge cells that matter for distance computation
<code>...</code>	Additional arguments as for <a href="#">writeRaster</a>

### Value

RasterLayer

### See Also

[distance](#), [gridDistance](#)

For the direction between (longitude/latitude) points, see the `azimuth` function in the `geosphere` package

### Examples

```
r <- raster(ncol=36,nrow=18)
values(r) <- NA
r[306] <- 1
b <- direction(r)
#plot(b)
```

disaggregate

*Disaggregate***Description**

Disaggregate a RasterLayer to create a new RasterLayer with a higher resolution (smaller cells). The values in the new RasterLayer are the same as in the larger original cells unless you specify `method="bilinear"`, in which case values are locally interpolated (using the [resample](#) function).

**Usage**

```
## S4 method for signature 'Raster'
disaggregate(x, fact=NULL, method='', filename='', ...)
```

**Arguments**

<code>x</code>	a Raster object
<code>fact</code>	integer. amount of disaggregation expressed as number of cells (horizontally and vertically). This can be a single integer or two integers <code>c(x,y)</code> , in which case the first one is the horizontal disaggregation factor and <code>y</code> the vertical disaggregation factor. If a single integer value is supplied, cells are disaggregated with the same factor in <code>x</code> and <code>y</code> direction
<code>method</code>	Character. '' or 'bilinear'. If 'bilinear', values are locally interpolated (using the <a href="#">resample</a> function)
<code>filename</code>	Character. Output filename (optional)
<code>...</code>	Additional arguments as for <a href="#">writeRaster</a>

**Value**

Raster object

**Author(s)**

Robert J. Hijmans and Jim Regetz

**See Also**

[aggregate](#)

**Examples**

```
r <- raster(ncols=10, nrows=10)
rd <- disaggregate(r, fact=c(10, 2))
ncol(rd)
nrow(rd)
values(r) <- 1:ncell(r)
rd <- disaggregate(r, fact=c(4, 2), method='bilinear')
```

---

distance	<i>Distance</i>
----------	-----------------

---

### Description

For a single RasterLayer (y is missing) this method computes the distance, for all cells that are NA, to the nearest cell that is not NA. The distance unit is in meters if the RasterLayer is not projected (+proj=longlat) and in map units (typically also meters) when it is projected.

If two RasterLayer objects are provided, the cell-value distances are computed. If two Spatial vector type objects are provided, the distances between pairs of geographic object are computed.

### Usage

```
## S4 method for signature 'RasterLayer,missing'
distance(x, y, filename='', doEdge=TRUE, ...)
## S4 method for signature 'RasterLayer,RasterLayer'
distance(x, y, ...)
## S4 method for signature 'Spatial,Spatial'
distance(x, y, ...)
```

### Arguments

x	RasterLayer object
y	missing, RasterLayer or Spatial object
filename	Character. Filename for the output RasterLayer (optional)
doEdge	Logical. If TRUE, the <a href="#">boundaries</a> function is called first. This may be efficient in cases where you compute the distance to large blobs. Calling boundaries determines the edge cells that matter for distance computation
...	Additional arguments as for <a href="#">writeRaster</a>

### Value

RasterLayer

### See Also

[distanceFromPoints](#), [gridDistance](#), [pointDistance](#)

See the [gdistance](#) package for more advanced distances, and the [geosphere](#) package for great-circle distances (and more) between points in longitude/latitude coordinates.

### Examples

```
r <- raster(ncol=36,nrow=18)
values(r) <- NA
r[500] <- 1
dist <- distance(r)
#plot(dist / 1000)
```

---

distanceFromPoints      *Distance from points*

---

### Description

The function calculates the distance from a set of points to all cells of a Raster\* object.

The distance unit is in meters if the coordinate reference system (crs) of the Raster\* object is (+proj=longlat) or assumed to be if the crs is NA. In all other cases it is in the units defined by the crs (which typically is meters).

### Usage

```
distanceFromPoints(object, xy, filename='', ...)
```

### Arguments

object	Raster object
xy	matrix of x and y coordinates, or a SpatialPoints* object.
filename	character. Optional filename for the output RasterLayer
...	Additional arguments as for <a href="#">writeRaster</a>

### Details

Distances for longlat data are computed on the WGS84 spheroid using GeographicLib (Karney, 2013)

### Value

RasterLayer

### References

C.F.F. Karney, 2013. Algorithms for geodesics, J. Geodesy 87: 43-55. <https://dx.doi.org/10.1007/s00190-012-0578-z>.

### See Also

[crs](#), [distance](#), [gridDistance](#), [pointDistance](#)

### Examples

```
r <- raster(ncol=36,nrow=18)
xy <- c(0,0)
d1 <- distanceFromPoints(r, xy)
crs(r) = '+proj=utm +zone=12 +datum=WGS84'
d2 <- distanceFromPoints(r, xy)
par(mfrow=c(1,2))
plot(d1)
plot(d2)
```



---

draw	<i>Draw a line or polygon</i>
------	-------------------------------

---

**Description**

Draw a line or polygon on a plot (map) and save it for later use. After calling the function, start clicking on the map. To finish, right-click and select 'stop'.

**Usage**

```
drawPoly(sp=TRUE, col='red', lwd=2, ...)
drawLine(sp=TRUE, col='red', lwd=2, ...)
```

**Arguments**

sp	logical. If TRUE, the output will be a sp object (SpatialPolygons or SpatialLines). Otherwise a matrix of coordinates is returned
col	the color of the lines to be drawn
lwd	the width of the lines to be drawn
...	additional arguments padded to locator

**Value**

If sp==TRUE a SpatialPolygons or SpatialLines object; otherwise a matrix of coordinates

**See Also**

[locator](#)

---

drawExtent	<i>Create an Extent object by drawing on a map</i>
------------	--

---

**Description**

Click on two points of a plot (map) to obtain an object of class [Extent](#) ('bounding box')

**Usage**

```
drawExtent(show=TRUE, col="red")
```

**Arguments**

show	logical. If TRUE, the extent will be drawn on the map
col	sets the color of the lines of the extent

**Value**

Extent

**Examples**

```
## Not run:
r1 <- raster(nrow=10, ncol=10)
values(r1) <- runif(ncell(r1))
plot(r1)
# after running the following line, click on the map twice
e <- drawExtent()
# after running the following line, click on the map twice
mean(values(crop(r1, drawExtent()))))

## End(Not run)
```

---

erase

*Erase parts of a `SpatialPolygons*` or `SpatialLines*` object. The inverse of this can be done with [intersect](#)*

---

**Description**

Erase parts of a `SpatialPolygons*` or `SpatialLines*` object with a `SpatialPolygons*` object

**Usage**

```
## S4 method for signature 'SpatialPolygons,SpatialPolygons'
erase(x, y, ...)
## S4 method for signature 'SpatialLines,SpatialPolygons'
erase(x, y, ...)
```

**Arguments**

x	SpatialPolygons or SpatialLines object
y	SpatialPolygons object
...	Additional arguments (none)

**Value**

Spatial\*

**Author(s)**

Robert J. Hijmans

**See Also**

The equivalent for raster data is [mask](#)

**Examples**

```

if (require(rgdal) & require(rgeos)) {
# erase parts of polygons with other polygons
p <- shapefile(system.file("external/lux.shp", package="raster"))
b <- as(extent(6, 6.4, 49.75, 50), 'SpatialPolygons')
crs(b) <- crs(p)
e <- erase(p, b)
plot(e)

# erase parts of lines with polygons
r <- raster(extent(p) +c(-.1,.1,-.1,.1), crs=crs(p))
start <- xyFromCell(r, cellFromCol(r, 1))
end <- xyFromCell(r, cellFromCol(r, ncol(r)))
lines <- do.call(splines, lapply(1:10, function(i)rbind(start[i,], end[i,])))
crs(lines) <- crs(p)

e2 <- erase(lines, p)
plot(p)
lines(lines, col='blue', lwd=4, lty=3)
lines(e2, col='red', lwd=2)
}

```

---

extend

*Extend*


---

**Description**

Extend returns an Raster\* object with a larger spatial extent. The output Raster object has the outer minimum and maximum coordinates of the input Raster and Extent arguments. Thus, all of the cells of the original raster are included. See [crop](#) if you (also) want to remove rows or columns.

There is also an extend method for Extent objects to enlarge (or reduce) an Extent. You can also use algebraic notation to do that (see examples)

This function has replaced function "expand" (to avoid a name conflict with the Matrix package).

**Usage**

```
## S4 method for signature 'Raster'
extend(x, y, value=NA, filename='', ...)
```

```
## S4 method for signature 'Extent'
extend(x, y, ...)
```

**Arguments**

x                    Raster or Extent object

y	If x is a Raster object, y should be an Extent object, or any object that is or has an Extent object, or an object from which it can be extracted (such as sp objects). Alternatively, you can provide a numeric vector of length 2 indicating the number of rows and columns that need to be added (or a single number when the number of rows and columns is equal) If x is an Extent object, y should be a numeric vector of 1, 2, or 4 elements
value	value to assign to new cells
filename	Character (optional)
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer or RasterBrick, or Extent

**Author(s)**

Robert J. Hijmans and Etienne B. Racine (Extent method)

**See Also**

[crop](#), [merge](#)

**Examples**

```
r <- raster(xmn=-150, xmx=-120, ymx=60, ymn=30, ncol=36, nrow=18)
values(r) <- 1:ncell(r)
e <- extent(-180, 0, 0, 90)
re <- extend(r, e)

# extend with a number of rows and columns (at each side)
re2 <- extend(r, c(2,10))

# Extent object
e <- extent(r)
e
extend(e, 10)
extend(e, 10, -10, 0, 20)
e + 10
e * 2
```

---

extension

*Filename extensions*

---

**Description**

Get or change a filename extension

**Usage**

```
extension(filename, value=NULL, maxchar=10)
extension(filename) <- value
```

**Arguments**

filename	A filename, with or without the path
value	A file extension with or without a dot, e.g., ".txt" or "txt"
maxchar	Maximum number of characters after the last dot in the filename, for that string to be considered a filename extension

**Value**

A file extension, filename or path.

If `ext(filename)` is used without a value argument, it returns the file extension; otherwise it returns the filename (with new extensions set to value)

**Examples**

```
fn <- "c:/temp folder/filename.extension"
extension(fn)
extension(fn) <- ".txt"
extension(fn)
fn <- extension(fn, '.document')
extension(fn)
extension(fn, maxchar=4)
```

---

extent

*Extent*

---

**Description**

This function returns an Extent object of a Raster\* or Spatial\* object (or an Extent object), or creates an Extent object from a 2x2 matrix (first row: xmin, xmax; second row: ymin, ymax), vector (length=4; order= xmin, xmax, ymin, ymax) or list (with at least two elements, with names 'x' and 'y')

bbox returns a sp package like 'bbox' object (a matrix)

**Usage**

```
extent(x, ...)
```

**Arguments**

`x` Raster\* or Extent object, a matrix, or a vector of four numbers

`...` Additional arguments. When `x` is a single number representing 'xmin', you can pass three additional numbers (`xmax`, `ymin`, `ymax`)

When `x` is a Raster\* object, you can pass four additional arguments to crop the extent: `r1`, `r2`, `c1`, `c2`, representing the first and last row and column number

**Value**

Extent object

**Author(s)**

Robert J. Hijmans; Etienne Racine wrote the extent function for a list

**See Also**

[extent](#), [drawExtent](#)

**Examples**

```
r <- raster()
extent(r)
extent(c(0, 20, 0, 20))
#is equivalent to
extent(0, 20, 0, 20)
extent(matrix(c(0, 0, 20, 20), nrow=2))
x <- list(x=c(0,1,2), y=c(-3,5))
extent(x)

#crop the extent by row and column numbers
extent(r, 1, 20, 10, 30)
```

---

Extent math

*round Extent coordinates*

---

**Description**

use `round(x, digits=0)` to round the coordinates of an Extent object to the number of digits specified. This can be useful when dealing with a small imprecision in the data (e.g. 179.9999 instead of 180). `floor` and `ceiling` move the coordinates to the outer or inner whole integer numbers.

It is also possible to use Arithmetic functions with Extent objects (but these work perhaps unexpectedly!)

See [Math-methods](#) for these (and many more) methods with Raster\* objects.

**Usage**

```
## S4 method for signature 'Extent'  
floor(x)  
## S4 method for signature 'Extent'  
ceiling(x)
```

**Arguments**

x                    Extent object

**See Also**

[Math-methods](#)

**Examples**

```
e <- extent(c(0.999999, 10.000011, -60.4, 60))  
round(e)  
ceiling(e)  
floor(e)
```

---

Extent-class

*Class "Extent"*

---

**Description**

Objects of class Extent are used to define the spatial extent (extremes) of objects of the BasicRaster and Raster\* classes.

**Objects from the Class**

You can use the [extent](#) function to create Extent objects, or to extract them from Raster\* and Spatial\* objects.

**Slots**

xmin: minimum x coordinate  
xmax: maximum x coordinate  
ymin: minimum y coordinate  
ymax: maximum y coordinate

**Methods**

**show** display values of a Extent object

**See Also**

[extent](#), [setExtent](#)

**Examples**

```
ext <- extent(-180,180,-90,90)
ext
```

---

 extract

---

*Extract values from Raster objects*


---

**Description**

Extract values from a Raster\* object at the locations of other spatial data. You can use coordinates (points), lines, polygons or an Extent (rectangle) object. You can also use cell numbers to extract values.

If y represents points, extract returns the values of a Raster\* object for the cells in which a set of points fall. If y represents lines, the extract method returns the values of the cells of a Raster\* object that are touched by a line. If y represents polygons, the extract method returns the values of the cells of a Raster\* object that are covered by a polygon. A cell is covered if its center is inside the polygon (but see the weights option for considering partly covered cells; and argument small for getting values for small polygons).

It is also possible to extract values for point locations from SpatialPolygons.

**Usage**

```
## S4 method for signature 'Raster,matrix'
extract(x, y, method='simple', buffer=NULL, small=FALSE, cellnumbers=FALSE,
        fun=NULL, na.rm=TRUE, layer, nl, df=FALSE, factors=FALSE, ...)

## S4 method for signature 'Raster,SpatialLines'
extract(x, y, fun=NULL, na.rm=FALSE, cellnumbers=FALSE, df=FALSE, layer,
        nl, factors=FALSE, along=FALSE, sp=FALSE, ...)

## S4 method for signature 'Raster,SpatialPolygons'
extract(x, y, fun=NULL, na.rm=FALSE, weights=FALSE,
        normalizeWeights=TRUE, cellnumbers=FALSE, small=TRUE, df=FALSE, layer, nl,
        factors=FALSE, sp=FALSE, ...)

## S4 method for signature 'SpatialPolygons,SpatialPoints'
extract(x, y, ...)
```

**Arguments**

x	Raster* object
y	points represented by a two-column matrix or data.frame, or <a href="#">SpatialPoints*</a> ; <a href="#">SpatialPolygons*</a> ; <a href="#">SpatialLines</a> ; <a href="#">Extent</a> ; or a numeric vector representing cell numbers



method	character. 'simple' or 'bilinear'. If 'simple' values for the cell a point falls in are returned. If 'bilinear' the returned values are interpolated from the values of the four nearest raster cells.
buffer	numeric. The radius of a buffer around each point from which to extract cell values. If the distance between the sampling point and the center of a cell is less than or equal to the buffer, the cell is included. The buffer can be specified as a single value, or as a vector of the length of the number of points. If the data are not projected (latitude/longitude), the unit should be meters. Otherwise it should be in map-units (typically also meters).
small	logical. If TRUE and y represents points and a buffer argument is used, the function always return a number, also when the buffer does not include the center of a single cell. The value of the cell in which the point falls is returned if no cell center is within the buffer. If y represents polygons, a value is also returned for relatively small polygons (e.g. those smaller than a single cell of the Raster* object), or polygons with an odd shape, for which otherwise no values are returned because they do not cover any raster cell centers. In some cases, you could alternatively use the centroids of such polygons, for example using <code>extract(x, coordinates(y))</code> or <code>extract(x, coordinates(y), method='bilinear')</code> .
fun	function to summarize the values (e.g. mean). The function should take a single numeric vector as argument and return a single value (e.g. mean, min or max), and accept a <code>na.rm</code> argument. Thus, standard R functions not including an <code>na.rm</code> argument must be wrapped as in this example: <code>fun=function(x,...)length(x)</code> . If y represents points, fun is only used when a buffer is used (and hence multiple values per spatial feature would otherwise be returned).
na.rm	logical. Only useful when an argument fun is supplied. If <code>na.rm=TRUE</code> (the default value), NA values are removed before fun is applied. This argument may be ignored if the function used has a <code>...</code> argument and ignores an additional <code>na.rm</code> argument
cellnumbers	logical. If <code>cellnumbers=TRUE</code> , cell-numbers will also be returned (if no fun argument is supplied, and when extracting values with points, if buffer is NULL)
df	logical. If <code>df=TRUE</code> , results will be returned as a data.frame. The first column is a sequential ID, the other column(s) are the extracted values
weights	logical. If TRUE and <code>normalizeWeights=FALSE</code> , the function returns, for each polygon, a matrix with the cell values and the approximate fraction of each cell that is covered by the polygon (rounded to 1/100). If TRUE and <code>normalizeWeights=TRUE</code> the weights are normalized such that they add up to one. The weights can be used for averaging; see examples. This option can be useful (but slow) if the polygons are small relative to the cells size of the Raster* object
normalizeWeights	logical. If TRUE, weights are normalized such that they add up to one for each polygon
factors	logical. If TRUE, factor values are returned, else their integer representation is returned
layer	integer. First layer for which you want values (if x is a multilayer object)
n1	integer. Number of layers for which you want values (if x is a multilayer object)

along	boolean. Should returned values be ordered to go along the lines?
sp	boolean. Should the extracted values be added to the data.frame of the Spatial* object y? This only applies if y is a Spatial* object and, for SpatialLines and SpatialPolygons, if fun is not NULL. In this case the returned value is the expanded Spatial object
...	additional arguments (none implemented)

### Value

A vector for RasterLayer objects, and a matrix for RasterStack or RasterBrick objects. A list (or a data.frame if `df=TRUE`) if y is a SpatialPolygons\* or SpatialLines\* object or if a buffer argument is used (but not a fun argument). If `sp=TRUE` and y is a Spatial\* object and fun is not NULL a Spatial\* object is returned. The order of the returned values corresponds to the order of object y. If `df=TRUE`, this is also indicated in the first variable ('ID').

### See Also

[getValues](#), [getValuesFocal](#)

### Examples

```
r <- raster(ncol=36, nrow=18, vals=1:(18*36))

#####
# extract values by cell number
#####
extract(r, c(1:2, 10, 100))
s <- stack(r, sqrt(r), r/r)
extract(s, c(1, 10, 100), layer=2, n=2)

#####
# extract values with points
#####
xy <- cbind(-50, seq(-80, 80, by=20))
extract(r, xy)

sp <- SpatialPoints(xy)
extract(r, sp, method='bilinear')

# examples with a buffer
extract(r, xy[1:3,], buffer=1000000)
extract(r, xy[1:3,], buffer=1000000, fun=mean)

## illustrating the varying size of a buffer (expressed in meters)
## on a longitude/latitude raster
z <- extract(r, xy, buffer=1000000)
s <- raster(r)
for (i in 1:length(z)) { s[z[[i]]] <- i }

## compare with raster that is not longitude/latitude
crs(r) <- "+proj=utm +zone=17"
```

```

xy[,1] <- 50
z <- extract(r, xy, buffer=8)
for (i in 1:length(z)) { s[z[[i]]] <- i }
plot(s)
# library(maptools)
# data(wrld_simpl)
# plot(wrld_simpl, add=TRUE)

#####
# extract values with lines
#####
r <- raster(ncol=36, nrow=18, vals=1:(18*36))
cds1 <- rbind(c(-50,0), c(0,60), c(40,5), c(15,-45), c(-10,-25))
cds2 <- rbind(c(80,20), c(140,60), c(160,0), c(140,-55))
lines <- spLines(cds1, cds2)

extract(r, lines)

#####
# extract values with polygons
#####
cds1 <- rbind(c(-180,-20), c(-160,5), c(-60, 0), c(-160,-60), c(-180,-20))
cds2 <- rbind(c(80,0), c(100,60), c(120,0), c(120,-55), c(80,0))
polys <- spPolygons(cds1, cds2)

v <- extract(r, polys)
# mean for each polygon
unlist(lapply(v, function(x) if (!is.null(x)) mean(x, na.rm=TRUE) else NA ))

# v <- extract(r, polys, cellnumbers=TRUE)

# weighted mean
# v <- extract(r, polys, weights=TRUE, fun=mean)
# equivalent to:
# v <- extract(r, polys, weights=TRUE)
# sapply(v, function(x) if (!is.null(x)) {sum(apply(x, 1, prod)) / sum(x[,2])} else NA)

#####
# extract values with an extent
#####
e <- extent(150,170,-60,-40)
extract(r, e)
#plot(r)
#plot(e, add=T)

```

**Description**

These are shorthand methods that call other methods that should normally be used, such as [getValues](#), [extract](#), [crop](#).

`object[i]` can be used to access values of a Raster\* object, using cell numbers. You can also use row and column numbers as index, using `object[i, j]` or `object[i, ]` or `object[, j]`. In addition you can supply an Extent, SpatialPolygons, SpatialLines or SpatialPoints object.

If `drop=TRUE` (the default) cell values are returned (a vector for a RasterLayer, a matrix for a RasterStack or RasterBrick). If `drop=FALSE` a Raster\* object is returned that has the extent covering the requested cells, and with all other non-requested cells within this extent set to NA.

If you supply a RasterLayer, its values will be used as logical (TRUE/FALSE) indices if both Raster objects have the same extent and resolution; otherwise the cell values within the extent of the RasterLayer are returned.

Double brackets `'[[ ]]`' can be used to extract one or more layers from a multi-layer object.

**Methods**

`x[i]`

`x[i, j]`

Arguments

<code>x</code>	a Raster* object
<code>i</code>	cell number(s), row number(s), a (logical) RasterLayer, Spatial* object
<code>j</code>	column number(s) (only available if <code>i</code> is (are) a row number(s))
<code>drop</code>	If TRUE, cell values are returned. Otherwise, a Raster* object is returned

**See Also**

[getValues](#), [setValues](#), [extract](#), [crop](#), [rasterize](#)

**Examples**

```
r <- raster(ncol=10, nrow=5)
values(r) <- 1:ncell(r)
```

```
r[1]
r[1:10]
r[1,]
r[,1]
r[1:2, 1:2]
```

```
s <- stack(r, sqrt(r))
s[1:3]
s[[2]]
```

---

Extreme coordinates     *Coordinates of the Extent of a Raster object*

---

**Description**

These functions return or set the extreme coordinates of a Raster\* object; and return them for Spatial\* objects.

**Usage**

```
xmin(x)
xmax(x)
ymin(x)
ymax(x)

xmin(x, ...) <- value
xmax(x, ...) <- value
ymin(x, ...) <- value
ymax(x, ...) <- value
```

**Arguments**

x	Raster* or Extent object
value	numeric. x or y coordinate
...	additional arguments. None implemented

**Value**

numeric

**See Also**

[extent, dimensions](#)

**Examples**

```
r <- raster(xmn=-0.5, xmx = 9.5, ncols=10)
xmin(r)
xmax(r)
ymin(r)
ymax(r)
xmin(r) <- -180
xmax(r) <- 180
```

---

extremeValues	<i>Minimum and maximum values</i>
---------------	-----------------------------------

---

### Description

Returns the minimum or maximum value of a RasterLayer or layer in a RasterStack

### Usage

```
minValue(x, ...)  
maxValue(x, ...)
```

### Arguments

x	RasterLayer or RasterStack object
...	Additional argument: layer number (for RasterStack or RasterBrick objects)

### Details

If a Raster\* object is created from a file on disk, the min and max values are often not known (depending on the file format). You can use [setMinMax](#) to set them in the Raster\* object.

### Value

a number

### Examples

```
r <- raster()  
r <- setValues(r, 1:ncell(r))  
minValue(r)  
maxValue(r)  
r <- setValues(r, round(100 * runif(ncell(r)) + 0.5))  
minValue(r)  
maxValue(r)  
  
r <- raster(system.file("external/test.grd", package="raster"))  
minValue(r)  
maxValue(r)
```

factors

*Factors***Description**

These functions allow for defining a RasterLayer as a categorical variable. Such a RasterLayer is linked to other values via a "Raster Attribute Table" (RAT). Thus the cell values are an index, whereas the actual values of interest are in the RAT. The RAT is a data.frame. The first column in the RAT ("ID") has the unique cell values of the layer; this column should normally not be changed. The other columns can be of any basic type (factor, character, integer, numeric or logical). The functions documented here are mainly available such that files with a RAT can be read and processed; currently there is not too much further support. Whether a layer is defined as a factor or not is currently ignored by almost all functions. An exception is the 'extract' function (when used with option df=TRUE).

Function 'levels' returns the RAT for inspection. It can be modified and set using `levels <-value` (but use caution as it is easy to mess things up).

`as.factor` and `ratify` create a layer with a RAT table. Function 'deratify' creates a single layer for a (or each) variable in the RAT table.

**Usage**

```
is.factor(x)
as.factor(x)
levels(x)
```

```
## S4 method for signature 'Raster'
ratify(x, filename="", count=FALSE, ...)
```

```
factorValues(x, v, layer=1, att=NULL, append.names=FALSE)
deratify(x, att=NULL, layer=1, complete=FALSE, drop=TRUE, fun='mean', filename='', ...)
```

```
asFactor(x, ...)
```

**Arguments**

<code>x</code>	Raster* object
<code>v</code>	integer cell values
<code>layer</code>	integer > 0 indicating which layer to use (in a RasterStack or RasterBrick)
<code>att</code>	numeric or character. Which variable(s) in the RAT table should be used. If NULL, all variables are extracted. If using a numeric, skip the first two default columns
<code>append.names</code>	logical. Should names of data.frame returned by a combination of the name of the layer and the RAT variables? (can be useful for multilayer objects)
<code>filename</code>	character. Optional
<code>count</code>	logical. If TRUE, a columns with frequencies is added

...	additional arguments as for <code>writeRaster</code>
<code>complete</code>	logical. If TRUE, the layer returned is no longer a factor
<code>drop</code>	logical. If TRUE a factor is converted to a numerical value if possible
<code>fun</code>	character. Used to get a single value for each class for a weighted RAT table. 'mean', 'min', 'max', 'smallest', or 'largest'

### Value

Raster\* object; list (levels); boolean (is.factor); matrix (factorValues)

### Note

`asFactor` is deprecated and should not be used

### Examples

```
set.seed(0)
r <- raster(nrow=10, ncol=10)
values(r) <- runif(ncell(r)) * 10
is.factor(r)

r <- round(r)
f <- as.factor(r)
is.factor(f)

x <- levels(f)[[1]]
x
x$code <- letters[10:20]
levels(f) <- x
levels(f)
f

r <- raster(nrow=10, ncol=10)
values(r) = 1
r[51:100] = 2
r[3:6, 1:5] = 3
r <- ratify(r)

rat <- levels(r)[[1]]
rat$landcover <- c('Pine', 'Oak', 'Meadow')
rat$code <- c(12,25,30)
levels(r) <- rat
r

# extract values for some cells
i <- extract(r, c(1,2, 25,100))
i
# get the attribute values for these cells
factorValues(r, i)

# write to file:
```



```
# rr <- writeRaster(r, rasterTmpFile(), overwrite=TRUE)
# rr

# create a single-layer factor
x <- deratify(r, 'landcover')
x
is.factor(x)
levels(x)
```

---

filename

*Filename*


---

### Description

Get the filename of a Raster\* object. You cannot set the filename of an object (except for Raster-Stack objects); but you can provide a 'filename=' argument to a function that creates a new Raster-Layer or RasterBrick\* object.

### Usage

```
filename(x)
```

### Arguments

x                    A Raster\* object

### Value

a Raster\* object

### Examples

```
r <- raster( system.file("external/test.grd", package="raster") )
filename(r)
```

---

filledContour

*Filled contour plot*


---

### Description

Filled contour plot of a RasterLayer. This is a wrapper around [filled.contour](#) for RasterLayer objects.

### Usage

```
filledContour(x, y=1, maxpixels=100000, ...)
```

**Arguments**

x	A Raster* object
y	Integer. The layer number of x (if x has multiple layers)
maxpixels	The maximum number of pixels
...	Any argument that can be passed to <a href="#">filled.contour</a> (graphics package)

**See Also**

[filled.contour](#), [persp](#), [plot](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
filledContour(r)
```

---

flip

*Flip*

---

**Description**

Flip the values of a Raster\* object by inverting the order of the rows (direction=y) or the columns (direction='x').

**Usage**

```
flip(x, direction, ...)
```

**Arguments**

x	Raster* object
direction	Character. 'y' or 'x'; or 1 (=x) or 2 (=y)
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer or RasterBrick

**See Also**

transpose: [t](#), [rotate](#)

**Examples**

```
r <- raster(nrow=18, ncol=36)
m <- matrix(1:ncell(r), nrow=18)
values(r) <- as.vector(t(m))
rx <- flip(r, direction='x')
values(r) <- as.vector(m)
ry <- flip(r, direction='y')
```

---

flowPath	<i>Flow path</i>
----------	------------------

---

## Description

Compute the flow path (drainage path) starting at a given point. See package `gdistance` for more path computations.

## Usage

```
flowPath(x, p, ...)
```

## Arguments

<code>x</code>	RasterLayer of flow direction (as can be created with <a href="#">terrain</a> )
<code>p</code>	starting point. Either two numbers: <code>x</code> (longitude) and <code>y</code> (latitude) coordinates; or a single cell number
<code>...</code>	additional arguments (none implemented)

## Value

numeric (cell numbers)

## Author(s)

Ashton Shortridge

## Examples

```
data(volcano)
v <- raster(volcano, xmn=2667400, xmx=2668010, ymn=6478700, ymx=6479570, crs="+init=epsg:27200")
fd <- terrain(v, opt = "flowdir")

path <- flowPath(fd, 2407)
xy <- xyFromCell(fd, path)
plot(v)
lines(xy)
```

---

focal	<i>Focal values</i>
-------	---------------------

---

### Description

Calculate focal ("moving window") values for the neighborhood of focal cells using a matrix of weights, perhaps in combination with a function.

### Usage

```
## S4 method for signature 'RasterLayer'
focal(x, w, fun, filename='', na.rm=FALSE, pad=FALSE, padValue=NA, NAonly=FALSE, ...)
```

### Arguments

x	RasterLayer
w	matrix of weights (the moving window), e.g. a 3 by 3 matrix with values 1; see Details. The matrix does not need to be square, but the sides must be odd numbers. If you need even sides, you can add a column or row with weights of zero
fun	function (optional). The function fun should take multiple numbers, and return a single number. For example mean, modal, min or max. It should also accept a na.rm argument (or ignore it, e.g. as one of the 'dots' arguments. For example, length will fail, but function(x, ...){na.omit(length(x))} works.
filename	character. Filename for a new raster (optional)
na.rm	logical. If TRUE, NA will be removed from focal computations. The result will only be NA if all focal cells are NA. Except for some special cases (weights of 1, functions like min, max, mean), using na.rm=TRUE is generally not a good idea in this function because it will unbalance the effect of the weights
pad	logical. If TRUE, additional 'virtual' rows and columns are padded to x such that there are no edge effects. This can be useful when a function needs to have access to the central cell of the filter
padValue	numeric. The value of the cells of the padded rows and columns
NAonly	logical. If TRUE, only cell values that are NA are replaced with the computed focal values
...	Additional arguments as for <a href="#">writeRaster</a>

### Details

focal uses a matrix of weights for the neighborhood of the focal cells. The default function is sum. It is computationally much more efficient to adjust the weights-matrix than to use another function through the fun argument. Thus while the following two statements are equivalent (if there are no NA values), the first one is faster than the second one:

```
a <-focal(x, w=matrix(1/9, nc=3, nr=3))
```

```
b <- focal(x, w=matrix(1,3,3), fun=mean)
```

There is, however, a difference if NA values are considered. One can use the `na.rm=TRUE` option which may make sense when using a function like `mean`. However, the results would be wrong when using a weights matrix.

```
Laplacian filter: filter=matrix(c(0,1,0,1,-4,1,0,1,0),nrow=3)
```

```
Sobel filters: fx=matrix(c(-1,-2,-1,0,0,0,1,2,1) / 4,nrow=3) and fy=matrix(c(1,0,-1,2,0,-2,1,0,-1)/4,nrow=3)
```

see the [focalWeight](#) function to create distance based circular, rectangular, or Gaussian filters.

## Value

RasterLayer

## See Also

[focalWeight](#)

## Examples

```
r <- raster(ncols=36, nrows=18, xmn=0)
values(r) <- runif(ncell(r))

# 3x3 mean filter
r3 <- focal(r, w=matrix(1/9,nrow=3,ncol=3))

# 5x5 mean filter
r5 <- focal(r, w=matrix(1/25,nrow=5,ncol=5))

# Gaussian filter
gf <- focalWeight(r, 2, "Gauss")
rg <- focal(r, w=gf)

# The max value for the lower-right corner of a 3x3 matrix around a focal cell
f = matrix(c(0,0,0,0,1,1,0,1,1), nrow=3)
f
rm <- focal(r, w=f, fun=max)

# global lon/lat data: no 'edge effect' for the columns
xmin(r) <- -180
r3g <- focal(r, w=matrix(1/9,nrow=3,ncol=3))

## Not run:
## focal can be used to create a cellular automaton

# Conway's Game of Life
w <- matrix(c(1,1,1,1,0,1,1,1,1), nr=3,nc=3)
gameOfLife <- function(x) {
  f <- focal(x, w=w, pad=TRUE, padValue=0)
  # cells with less than two or more than three live neighbours die
  x[f<2 | f>3] <- 0
  # cells with three live neighbours become alive
```

```

x[f==3] <- 1
x
}

# simulation function
sim <- function(x, fun, n=100, pause=0.25) {
  for (i in 1:n) {
    x <- fun(x)
    plot(x, legend=FALSE, asp=NA, main=i)
    dev.flush()
    Sys.sleep(pause)
  }
  invisible(x)
}

# Gosper glider gun
m <- matrix(0, nc=48, nr=34)
m[c(40, 41, 74, 75, 380, 381, 382, 413, 417, 446, 452, 480,
     486, 517, 549, 553, 584, 585, 586, 619, 718, 719, 720, 752,
     753, 754, 785, 789, 852, 853, 857, 858, 1194, 1195, 1228, 1229)] <- 1
init <- raster(m)

# run the model
sim(init, gameOfLife, n=150, pause=0.05)

## Implementation of Sobel edge-detection filter
## for RasterLayer r
sobel <- function(r) {
  fy <- matrix(c(1,0,-1,2,0,-2,1,0,-1), nrow=3)
  fx <- matrix(c(-1,-2,-1,0,0,0,1,2,1) , nrow=3)
  rx <- focal(r, fx)
  ry <- focal(r, fy)
  sqrt(rx^2 + ry^2)
}

## End(Not run)

```

---

focalWeight

*Focal weights matrix*


---

### Description

Calculate focal ("moving window") weight matrix for use in the `focal` function. The sum of the values adds up to one.

### Usage

```
focalWeight(x, d, type=c('circle', 'Gauss', 'rectangle'))
```

**Arguments**

x	Raster* object
d	numeric. If type=circle, the radius of the circle (in units of the CRS). If type=rectangle the dimension of the rectangle (one or two numbers). If type=Gauss the size of sigma, and optionally another number to determine the size of the matrix returned (default is 3 times sigma)
type	character indicating the type of filter to be returned

**Value**

matrix that can be used in [focal](#)

**Examples**

```
r <- raster(ncols=36, nrows=18, xmn=0)
# Gaussian filter for square cells
gf <- focalWeight(r, 2, "Gauss")
```

---

freq	<i>Frequency table</i>
------	------------------------

---

**Description**

Frequency table of the values of a RasterLayer.

**Usage**

```
## S4 method for signature 'RasterLayer'
freq(x, digits=0, value=NULL, useNA='ifany', progress='', ...)

## S4 method for signature 'RasterStackBrick'
freq(x, digits=0, value=NULL, useNA='ifany', merge=FALSE, progress='', ...)
```

**Arguments**

x	RasterLayer
digits	non-negative integer for rounding the cell values. Argument is passed to round
value	numeric, logical or NA. An optional single value to only count the number of cells with that value
useNA	character. What to do with NA values? Options are "no", "ifany", "always". See to <a href="#">table</a>
progress	character to specify a progress bar. Choose from 'text', 'window', or "" (the default, no progress bar)
merge	logical. If TRUE the list will be merged into a single data.frame
...	additional arguments (none implemented)

**Value**

matrix (RasterLayer). List of matrices (one for each layer) or data.frame (if merge=TRUE) (RasterStack or RasterBrick)

**See Also**

[crosstab](#) and [zonal](#)

**Examples**

```
r <- raster(nrow=18, ncol=36)
values(r) <- runif(ncell(r))
r[1:5] <- NA
r <- r * r * r * 5
freq(r)

freq(r, value=2)

s <- stack(r, r*2, r*3)
freq(s, merge=TRUE)
```

---

Gain and offset

*Gain and offset of values on file*

---

**Description**

These functions can be used to get or set the gain and offset parameters used to transform values when reading them from a file. The gain and offset parameters are applied to the raw values using the formula below:

$$\text{value} <- \text{value} * \text{gain} + \text{offset}$$

The default value for gain is 1 and for offset is 0. 'gain' is sometimes referred to as 'scale'.

Note that setting gain and/or offset are intended to be used with values that are stored in a file. For a Raster\* object with values in memory, assigning gain or offset values will lead to the immediate computation of new values; in such cases it would be clearer to use [Arith-methods](#).

**Usage**

```
gain(x)
gain(x) <- value
offs(x)
offs(x) <- value
```

**Arguments**

x	Raster* object
value	Single numeric value



**Value**

Raster\* object or numeric value(s)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
gain(r)
offs(r)
r[1505:1510]
gain(r) <- 10
offs(r) <- 5
r[1505:1510]
```

---

geom

*Get the coordinates of a vector type Spatial\* object*

---

**Description**

Extract the coordinates of a Spatial object

**Usage**

```
## S4 method for signature 'SpatialPolygons'
geom(x, sepNA=FALSE, ...)
## S4 method for signature 'SpatialLines'
geom(x, sepNA=FALSE, ...)
## S4 method for signature 'SpatialPoints'
geom(x, ...)
## S4 method for signature 'data.frame'
geom(x, d, gt, crs, ...)
```

**Arguments**

x	SpatialPolygons*, SpatialLines*, or SpatialPoints* object; or a data.frame
sepNA	logical. If TRUE, geometries are separated by a row with NA values
...	additional arguments, none implemented
d	data.frame that matches the number of objects in data.frame x
gt	character. geometry type. Must be one of "polygons", "lines", "points"
crs	character. PROJ.4 crs string

**Value**

Matrix with 6, (5 SpatialLines), or 3 (SpatialPoints) columns. object (sequential object number) part (sequential part number within the object; not for SpatialPoints), cump (cumulative part number; not for SpatialPoints), hole (is this a hole or not; only for SpatialPolygons), x (x coordinate or longitude), y (y coordinate or latitude)

**See Also**

[coordinates](#), [geometry](#)

**Examples**

```
p <- readRDS(system.file("external/lux.rds", package="raster"))
x <- geom(p)
head(x)

# and back to a SpatialPolygonsDataFrame
x <- data.frame(x)
sp <- as(x, "SpatialPolygons")
crs(sp) <- crs(p)
spdf <- SpatialPolygonsDataFrame(sp, data.frame(p), match.ID=FALSE)
```

---

getData

*Get geographic data*

---

**Description**

Get geographic data for anywhere in the world. Data are read from files that are first downloaded if necessary. Function ccodes returns country names and the ISO codes

**Usage**

```
getData(name, download=TRUE, path=' ', ...)
ccodes()
```

**Arguments**

name	Data set name, currently supported are 'GADM', 'countries', 'SRTM', 'alt', and 'worldclim'. See Details for more info
download	Logical. If TRUE data will be downloaded if not locally available
path	Character. Path name indicating where to store the data. Default is the current working directory
...	Additional required (!) parameters. These are data set specific. See Details

**Details**

'alt' stands for altitude (elevation); the data were aggregated from SRTM 90 m resolution data between -60 and 60 latitude. 'GADM' is a database of global administrative boundaries. 'worldclim' is a database of global interpolated climate data. 'SRTM' refers to the hole-filled CGIAR-SRTM (90 m resolution). 'countries' has polygons for all countries at a higher resolution than the 'wrld\_simpl' data in the maptools package .

If name is 'alt' or 'GADM' you must provide a 'country=' argument. Countries are specified by their 3 letter ISO codes. Use `getData('ISO3')` to see these codes. In the case of GADM you must

also provide the level of administrative subdivision (0=country, 1=first level subdivision). In the case of alt you can set 'mask' to FALSE. If it is TRUE values for neighbouring countries are set to NA. For example:

```
getData('GADM',country='FRA',level=1)
getData('alt',country='FRA',mask=TRUE)
```

If name is 'SRTM' you must provide 'lon' and 'lat' arguments (longitude and latitude). These should be single numbers somewhere within the SRTM tile that you want.

```
getData('SRTM',lon=5,lat=45)
```

If name='worldclim' you must also provide arguments var, and a resolution res. Valid variables names are 'tmin', 'tmax', 'prec' and 'bio'. Valid resolutions are 0.5, 2.5, 5, and 10 (minutes of a degree). In the case of res=0.5, you must also provide a lon and lat argument for a tile; for the lower resolutions global data will be downloaded. In all cases there are 12 (monthly) files for each variable except for 'bio' which contains 19 files.

```
getData('worldclim',var='tmin',res=0.5,lon=5,lat=45)
getData('worldclim',var='bio',res=10)
```

To get (projected) future climate data (CMIP5), you must provide arguments var and res as above. Only resolutions 2.5, 5, and 10 are currently available. In addition, you need to provide model, rcp and year. For example,

```
getData('CMIP5',var='tmin',res=10,rcp=85,model='AC',year=70)
```

```
function (var, model, rcp, year, res, lon, lat, path, download = TRUE)
```

'model' should be one of "AC", "BC", "CC", "CE", "CN", "GF", "GD", "GS", "HD", "HG", "HE", "IN", "IP", "MI", "MR", "MC", "MP", "MG", or "NO".

'rcp' should be one of 26, 45, 60, or 85.

'year' should be 50 or 70

Not all combinations are available. See [www.worldclim.org](http://www.worldclim.org) for details.

## Value

A spatial object (Raster\* or Spatial\*)

## References

<http://www.worldclim.org>

<https://gadm.org>

<http://srtm.csi.cgiar.org/>

<http://diva-gis.org/gdata>

---

`getValues`*Get raster cell values*

---

**Description**

`getValues` returns all values or the values for a number of rows of a `Raster*` object. Values returned for a `RasterLayer` are a vector. The values returned for a `RasterStack` or `RasterBrick` are always a matrix, with the rows representing cells, and the columns representing layers

`values` is a shorthand version of `getValues` (for all rows).

**Usage**

```
getValues(x, row, nrows, ...)
```

```
values(x, ...)
```

**Arguments**

<code>x</code>	<code>Raster*</code> object
<code>row</code>	Numeric. Row number, should be between 1 and <code>nrow(x)</code> , or missing in which case all values are returned
<code>nrows</code>	Numeric. Number of rows. Should be an integer > 0, or missing
<code>...</code>	Additional arguments. When <code>x</code> is a <code>RasterLayer</code> : format to specify the output format. Either "matrix" or, the default "", in which case a vector is returned

**Value**

vector or matrix of raster values

**See Also**

[getValuesBlock](#), [getValuesFocal](#), [setValues](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
r
v <- getValues(r)
length(v)
head(v)
getValues(r, row=10)
```

---

getValuesBlock      *Get a block of raster cell values*

---

### Description

getValuesBlock returns values for a block (rectangular area) of values of a Raster\* object.

### Usage

```
## S4 method for signature 'RasterLayer'
getValuesBlock(x, row=1, nrows=1, col=1, ncols=(ncol(x)-col+1), format='', ...)

## S4 method for signature 'RasterBrick'
getValuesBlock(x, row=1, nrows=1, col=1, ncols=(ncol(x)-col+1), lyrs, ...)

## S4 method for signature 'RasterStack'
getValuesBlock(x, row=1, nrows=1, col=1, ncols=(ncol(x)-col+1), lyrs, ...)
```

### Arguments

x	Raster* object
row	positive integer. Row number to start from, should be between 1 and nrow(x)
nrows	positive integer. How many rows? Default is 1
col	positive integer. Column number to start from, should be between 1 and ncol(x)
ncols	positive integer. How many columns? Default is the number of columns left after the start column
format	character. When x is a RasterLayer, if format='matrix' or format='m', a matrix is returned instead of a vector. If format='matrix', it is a nrow x ncol matrix. If format='m' it is a 1 column matrix (the benefit is that the type of output is now the same for all Raster objects)
lyrs	integer (vector). Which layers? Default is all layers (1:nlayers(x))
...	additional arguments (none implemented)

### Value

matrix or vector (if (x=RasterLayer), unless format='matrix')

### See Also

[getValues](#)

**Examples**

```

r <- raster(system.file("external/test.grd", package="raster"))
b <- getValuesBlock(r, row=100, nrows=3, col=10, ncols=5)
b
b <- matrix(b, nrow=3, ncol=5, byrow=TRUE)
b

logo <- brick(system.file("external/rlogo.grd", package="raster"))
getValuesBlock(logo, row=35, nrows=3, col=50, ncols=3, lyrs=2:3)

```

---

getValuesFocal	<i>Get focal raster cell values</i>
----------------	-------------------------------------

---

**Description**

This function returns a matrix (or matrices) for all focal values of a number of rows of a Raster\* object

**Usage**

```

## S4 method for signature 'Raster'
getValuesFocal(x, row, nrows, ngb, names=FALSE, padValue=NA, array=FALSE, ...)

```

**Arguments**

x	Raster* object
row	Numeric. Row number, should be between 1 and nrow(x). Can be omitted to get all rows
nrows	Numeric. Number of rows, should be a positive integer smaller than row+nrow(x). Should be omitted if row is omitted
ngb	Neighbourhood size. Either a single integer or a vector of two integers c(nrow, ncol)
names	logical. If TRUE, the matrix returned has row and column names
padValue	numeric. The value of the cells of the "padded" rows and columns. That is 'virtual' values for cells within a neighbourhood, but outside the raster
array	logical. If TRUE and x has multiple layers, an array is returned in stead of a list of matrices
...	additional arguments (none implemented)

**Value**

If x has a single layer, a matrix with one row for each focal cell, and one column for each neighbourhood cell around it.

If x has multiple layers, an array (if array=TRUE) or a list of such matrices (one list element (matrix) for each layer)

**See Also**[getValues](#), [focal](#)**Examples**

```
r <- raster(nr=5, nc=5, crs='+proj=utm +zone=12')
values(r) <- 1:25
as.matrix(r)
getValuesFocal(r, row=1, nrows=2, ngb=3, names=TRUE)
getValuesFocal(stack(r,r), row=1, nrows=1, ngb=3, names=TRUE, array=TRUE)
```

---

gridDistance	<i>Distance on a grid</i>
--------------	---------------------------

---

**Description**

The function calculates the distance to cells of a RasterLayer when the path has to go through the centers of neighboring raster cells (currently only implemented as a 'queen' case in which cells have 8 neighbors).

The distance is in meters if the coordinate reference system (CRS) of the RasterLayer is longitude/latitude (+proj=longlat) and in the units of the CRS (typically meters) in other cases.

Distances are computed by summing local distances between cells, which are connected with their neighbours in 8 directions.

**Usage**

```
## S4 method for signature 'RasterLayer'
gridDistance(x, origin, omit=NULL, filename="", ...)
```

**Arguments**

x	RasterLayer
origin	value(s) of the cells from which the distance is calculated
omit	value(s) of the cells which cannot be traversed (optional)
filename	character. output filename (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Details**

If the RasterLayer to be processed is big, it will be processed in chunks. This may lead to errors in the case of complex objects spread over different chunks (meandering rivers, for instance). You can try to solve these issues by varying the chunk size, see function [setOptions\(\)](#).

**Value**

RasterLayer

**Author(s)**

Jacob van Etten and Robert J. Hijmans

**See Also**

See [distance](#) for 'as the crow flies' distance. Additional distance measures and options (directions, cost-distance) are available in the 'gdistance' package.

**Examples**

```
#world lon/lat raster
r <- raster(ncol=10,nrow=10, vals=1)
r[48] <- 2
r[66:68] <- 3
d <- gridDistance(r,origin=2,omit=3)
plot(d)

#UTM small area
crs(r) <- "+proj=utm +zone=15 +ellps=GRS80 +datum=NAD83 +units=m +no_defs"
d <- gridDistance(r,origin=2,omit=3)
plot(d)
```

---

 hdr

*Header files*


---

**Description**

Write header files to use together with raster binary files to read the data in other applications.

**Usage**

```
hdr(x, format, extension='.wld', filename='')
```

**Arguments**

x	RasterLayer or RasterBrick object associated with a binary values file on disk
format	Type of header file: 'VRT', 'BIL', 'ENVI', 'ErdasRaw', 'IDRISI', 'SAGA', 'RASTER', 'WORLDFILE', 'PRJ'
extension	File extension, only used with an ESRI worldfile (format='WORLDFILE')
filename	character. Need to be provided if x is not associated with a file



## Details

The RasterLayer object must be associated with a file on disk.

You can use `writeRaster` to save an existing file in another format. But if you have a file in a 'raster' format (or similar), you can also only export a header file, and use the data file (.gri) that already exists. The function can write a VRT (GDAL virtual raster) header (.vrt); an ENVI or BIL header (.hdr) file; an Erdas Raw (.raw) header file; an IDRISI (.rdc) or SAGA (.sgrd). This (hopefully) allows for reading the binary data (.gri), perhaps after changing the file extension, in other programs such as ENVI or ArcGIS.

## See Also

`writeRaster`, `writeGDAL`

## Examples

```
## Not run:
r <- raster(system.file("external/test.grd", package="raster"))
r <- writeRaster(r, filename='export.grd', overwrite=TRUE)
hdr(r, format="ENVI")

## End(Not run)
```

---

head

*Show the head or tail of a Raster\* object*

---

## Description

Show the head (first rows/columns) or tail (last rows/columns) of the cell values of a Raster\* object.

## Usage

```
head(x, ...)
tail(x, ...)
```

## Arguments

<code>x</code>	Raster* object
<code>...</code>	Additional arguments: <code>rows=10</code> and <code>cols=20</code> , to set the maximum number of rows and columns that are shown. For RasterStack and RasterBrick objects there is an additional argument <code>lyrs</code>

## Value

matrix

**See Also**[getValuesBlock](#)**Examples**

```
r <- raster(nrow=25, ncol=25)
values(r) = 1:ncell(r)
head(r)
tail(r, cols=10, rows=5)
```

hillShade

*Hill shading***Description**

Compute hill shade from slope and aspect layers (both in radians). Slope and aspect can be computed with function [terrain](#).

A hill shade layer is often used as a backdrop on top of which another, semi-transparent, layer is drawn.

**Usage**

```
hillShade(slope, aspect, angle=45, direction=0, filename='', normalize=FALSE, ...)
```

**Arguments**

slope	RasterLayer object with slope values (in radians)
aspect	RasterLayer object with aspect values (in radians)
angle	The the elevation angle of the light source (sun), in degrees
direction	The direction (azimuth) angle of the light source (sun), in degrees
filename	Character. Optional filename
normalize	Logical. If TRUE, values below zero are set to zero and the results are multiplied with 255
...	Standard additional arguments for writing RasterLayer files

**Author(s)**

Andrew Bevan, Robert J. Hijmans

**References**

Horn, B.K.P., 1981. Hill shading and the reflectance map. Proceedings of the IEEE 69(1):14-47

**See Also**[terrain](#)

**Examples**

```
## Not run:
alt <- getData('alt', country='CHE')
slope <- terrain(alt, opt='slope')
aspect <- terrain(alt, opt='aspect')
hill <- hillShade(slope, aspect, 40, 270)
plot(hill, col=grey(0:100/100), legend=FALSE, main='Switzerland')
plot(alt, col=rainbow(25, alpha=0.35), add=TRUE)

## End(Not run)
```

---

hist	<i>Histogram</i>
------	------------------

---

**Description**

Create a histogram of the values of a RasterLayer. For large datasets a sample is used.

**Usage**

```
## S4 method for signature 'Raster'
hist(x, layer, maxpixels=100000, plot=TRUE, main, ...)
```

**Arguments**

x	Raster* object
layer	integer (or character) to indicate layer number (or name). Can be used to subset the layers to plot in a multilayer Raster* object
maxpixels	integer. To regularly subsample very large objects
plot	logical. Plot the histogram or only return the histogram values
main	character. Main title(s) for the plot. Default is the value of <a href="#">names</a>
...	Additional arguments. See under Methods and at <a href="#">hist</a>

**Value**

This function is principally used for the side-effect of plotting a histogram, but it also returns an S3 object of class 'histogram' (invisibly if plot=TRUE).

**See Also**

[pairs](#), [boxplot](#)

**Examples**

```

r1 <- raster(nrows=50, ncols=50)
r1 <- setValues(r1, runif(ncell(r1)))
r2 <- setValues(r1, runif(ncell(r1)))
rs <- r1 + r2
rp <- r1 * r2
par(mfrow=c(2,2))
plot(rs, main='sum')
plot(rp, main='product')
hist(rs)
a = hist(rp)
a

```

---

image

*Image*


---

**Description**

Create an "image" type plot of a RasterLayer. This is an implementation of a generic function in the graphics package. In most cases the `plot` function would be preferable because it produces a legend (and has some additional options).

**Usage**

```

image(x, ...)
## S4 method for signature 'RasterLayer'
image(x, maxpixels=500000, useRaster=TRUE, ...)

## S4 method for signature 'RasterStackBrick'
image(x, y=1, maxpixels=100000, useRaster=TRUE, main, ...)

```

**Arguments**

x	Raster* object
maxpixels	integer > 0. Maximum number of cells to use for the plot. If maxpixels < ncell(x), sampleRegular is used before plotting
useRaster	If TRUE, the rasterImage function is used for plotting. Otherwise the image function is used. This can be useful if rasterImage does not work well on your system (see note)
main	character. Main plot title
...	Any argument that can be passed to <code>image</code> (graphics package)
y	If x is a RasterStack or RasterBrick: integer, character (layer name(s)), or missing to select which layer(s) to plot

**Note**

raster uses [rasterImage](#) from the graphics package. For unknown reasons this does not work on Windows Server and on a few versions of Windows XP. On that system you may need to use argument `useRaster=FALSE` to get a plot.

**See Also**

[plot](#), [image](#), [contour](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
image(r)
```

---

inifile

*Read a .ini file*

---

**Description**

This function reads '.ini' files. These are text file databases that are organized in [sections] containing pairs of "name = value".

**Usage**

```
readIniFile(filename, token='=', commenttoken=';', aslist=FALSE, case)
```

**Arguments**

filename	Character. Filename of the .ini file
token	Character. The character that separates the "name" (variable name) from the "value"
commenttoken	Character. This token and everything that follows on the same line is considered a 'comment' that is not for machine consumption and is ignored in processing
aslist	Logical. Should the values be returned as a list
case	Optional. Function that operates on the text, such as <a href="#">toupper</a> or <a href="#">tolower</a>

**Details**

This function allows for using inistrings that have "=" as part of a value (but the token cannot be part of the 'name' of a variable!). Sections can be missing.

**Value**

A n\*3 matrix of characters with columns: section, name, value; or a list if `aslist=TRUE`.

---

initialize

*Initialize a Raster object with values*


---

### Description

Create a new RasterLayer with values reflecting a cell property: 'x', 'y', 'col', 'row', or 'cell'. Alternatively, a function can be used. In that case, cell values are initialized without reference to pre-existing values. E.g., initialize with a random number (fun=`runif`). While there are more direct ways of achieving this for small objects (see examples) for which a vector with all values can be created in memory, the `init` function will also work for Raster\* objects with many cells.

### Usage

```
## S4 method for signature 'Raster'
init(x, fun, filename="", ...)
```

### Arguments

x	Raster* object
fun	function to be applied. This must be a function that can take the number of cells as a single argument to return a vector of values with a length equal to the number of cells, such as <code>fun=runif</code> . You can also supply one of the following character values: 'x', 'y', 'row', 'col', or 'cell' to get the x or coordinate, row, col or cell number; you can also use 'chess', to get a chessboard pattern
filename	character. Optional output filename
...	Additional arguments as for <code>writeRaster</code>

### Value

RasterLayer

### Note

For backwards compatibility, the character values valid for `fun` can also be passed as named argument `v`

### Examples

```
r <- raster(ncols=36, nrows=18)

x <- init(r, fun='cell')

y <- init(r, fun=runif)

# there are different ways to set all values to 1
# for large rasters:
# set1f <- function(x){rep(1, x)}
```

```
# z1 <- init(r, fun=set1f, filename=rasterTmpFile(), overwrite=TRUE)

# This is equivalent to (but not memory safe):
z2 <- setValues(r, rep(1, ncell(r)))
# or
values(r) <- rep(1, ncell(r))
# or
values(r) <- 1
```

---

interpolate

*Interpolate*


---

### Description

Make a RasterLayer with interpolated values using a fitted model object of classes such as 'gstat' (gstat package) or 'Krige' (fields package). That is, these are models that have location ('x' and 'y', or 'longitude' and 'latitude') as independent variables. If x and y are the only independent variables provide an empty (no associated data in memory or on file) RasterLayer for which you want predictions. If there are more spatial predictor variables provide these as a Raster\* object in the first argument of the function. If you do not have x and y locations as implicit predictors in your model you should use [predict](#) instead.

### Usage

```
## S4 method for signature 'Raster'
interpolate(object, model, filename="", fun=predict, xyOnly=TRUE,
  xyNames=c('x', 'y'), ext=NULL, const=NULL, index=1, na.rm=TRUE, debug.level=1, ...)
```

### Arguments

object	Raster* object
model	model object
filename	character. Output filename (optional)
fun	function. Default value is 'predict', but can be replaced with e.g. 'predict.se' (depending on the class of the model object)
xyOnly	logical. If TRUE, values of the Raster* object are not considered as co-variables; and only x and y (longitude and latitude) are used. This should match the model
xyNames	character. variable names that the model uses for the spatial coordinates. E.g., c('longitude', 'latitude')
ext	Extent object to limit the prediction to a sub-region of x
const	data.frame. Can be used to add a constant for which there is no Raster object for model predictions. This is particularly useful if the constant is a character-like factor value
index	integer. To select the column if 'predict.model' returns a matrix with multiple columns

`na.rm` logical. Remove cells with NA values in the predictors before solving the model (and return NA for those cells). In most cases this will not affect the output. This option prevents errors with models that cannot handle NA values

`debug.level` for gstat models only. See ?

... additional arguments passed to the `predict.model` function

**Value**

Raster\* object

**See Also**

[predict](#), [predict.gstat](#), [Tps](#)

**Examples**

```
## Thin plate spline interpolation with x and y only
# some example data
r <- raster(system.file("external/test.grd", package="raster"))
ra <- aggregate(r, 10)
xy <- data.frame(xyFromCell(ra, 1:ncell(ra)))
v <- getValues(ra)
# remove NAs
i <- !is.na(v)
xy <- xy[i,]
v <- v[i]

#### Thin plate spline model
library(fields)
tps <- Tps(xy, v)
p <- raster(r)

# use model to predict values at all locations
p <- interpolate(p, tps)
p <- mask(p, r)

plot(p)
## change the fun from predict to fields::predictSE to get the TPS standard error
se <- interpolate(p, tps, fun=predictSE)
se <- mask(se, r)
plot(se)

## another variable; let's call it elevation
elevation <- (init(r, 'x') * init(r, 'y')) / 100000000
names(elevation) <- 'elev'

z <- extract(elevation, xy)

# add as another independent variable
xyz <- cbind(xy, z)
tps2 <- Tps(xyz, v)
```



```

p2 <- interpolate(elevation, tps2, xyOnly=FALSE)

# as a linear covariate
tps3 <- Tps(xy, v, Z=z)

# Z is a separate argument in Krig.predict, so we need a new function
# Internally (in interpolate) a matrix is formed of x, y, and elev (Z)

pfun <- function(model, x, ...) {
  predict(model, x[,1:2], Z=x[,3], ...)
}
p3 <- interpolate(elevation, tps3, xyOnly=FALSE, fun=pfun)

#### gstat examples
library(gstat)
data(meuse)

## inverse distance weighted (IDW)
r <- raster(system.file("external/test.grd", package="raster"))
data(meuse)
mg <- gstat(id = "zinc", formula = zinc~1, locations = ~x+y, data=meuse,
            nmax=7, set=list(idp = .5))
z <- interpolate(r, mg)
z <- mask(z, r)

## kriging
coordinates(meuse) <- ~x+y
crs(meuse) <- crs(r)

## ordinary kriging
v <- variogram(log(zinc)~1, meuse)
m <- fit.variogram(v, vgm(1, "Sph", 300, 1))
gOK <- gstat(NULL, "log.zinc", log(zinc)~1, meuse, model=m)
OK <- interpolate(r, gOK)

# examples below provided by Maurizio Marchi
## universal kriging
vu <- variogram(log(zinc)~elev, meuse)
mu <- fit.variogram(vu, vgm(1, "Sph", 300, 1))
gUK <- gstat(NULL, "log.zinc", log(zinc)~elev, meuse, model=mu)
names(r) <- 'elev'
UK <- interpolate(r, gUK, xyOnly=FALSE)

## co-kriging
gCoK <- gstat(NULL, 'log.zinc', log(zinc)~1, meuse)
gCoK <- gstat(gCoK, 'elev', elev~1, meuse)
gCoK <- gstat(gCoK, 'cadmium', cadmium~1, meuse)
gCoK <- gstat(gCoK, 'copper', copper~1, meuse)
coV <- variogram(gCoK)
plot(coV, type='b', main='Co-variogram')
coV.fit <- fit.lmc(coV, gCoK, vgm(model='Sph', range=1000))
coV.fit
plot(coV, coV.fit, main='Fitted Co-variogram')

```

```
coK <- interpolate(r, coV.fit)
plot(coK)
```

---

 intersect

*Intersect*


---

### Description

It depends on the classes of the *x* and *y* what is returned.

If *x* is a *Raster\** object the extent of *y* is used, irrespective of the class of *y*, and a *Raster\** is returned. This is equivalent to [crop](#).

If *x* is a *Spatial\** object, a new *Spatial\** object is returned. If *x* or *y* has a *data.frame*, these are also returned (after merging if necessary) as part of a *Spatial\*DataFrame*, and this is how *intersect* is different from *rgeos::gIntersection* on which it depends.

Intersecting *SpatialPoints\** with *SpatialPoints\** uses the extent (bounding box) of *y* to get the intersection. Intersecting of *SpatialPoints\** and *SpatialLines\** is not supported because of numerical inaccuracies with that. You can use [buffer](#), to create *SpatialPoygons\** from *SpatialLines\** and use that in *intersect*. Or try [gIntersection](#).

### Usage

```
## S4 method for signature 'Extent,ANY'
intersect(x, y)

## S4 method for signature 'Raster,ANY'
intersect(x, y)

## S4 method for signature 'SpatialPoints,ANY'
intersect(x, y)

## S4 method for signature 'SpatialPolygons,SpatialPolygons'
intersect(x, y)

## S4 method for signature 'SpatialPolygons,SpatialLines'
intersect(x, y)

## S4 method for signature 'SpatialPolygons,SpatialPoints'
intersect(x, y)

## S4 method for signature 'SpatialLines,SpatialPolygons'
intersect(x, y)

## S4 method for signature 'SpatialLines,SpatialLines'
intersect(x, y)
```

**Arguments**

x                    Extent, Raster\*, SpatialPolygons\*, SpatialLines\* or SpatialPoints\* object  
 y                    same as for x

**Value**

if x is an Extent object: Extent  
 if x is a Raster\* object: Raster\*  
 if x is a SpatialPoints\* object: SpatialPoints\*  
 if x is a SpatialPolygons\* object: SpatialPolygons\*  
 if x is a SpatialLines\* object and if y is a SpatialLines\* object: SpatialPoints\*  
 if x is a SpatialLines\* object and if y is a SpatialPolygons\* object: SpatialLines\*

**See Also**

[union](#), [extent](#), [crop](#)

**Examples**

```
e1 <- extent(-10, 10, -20, 20)
e2 <- extent(0, 20, -40, 5)
intersect(e1, e2)

#SpatialPolygons
if (require(rgdal) & require(rgeos)) {
  p <- shapefile(system.file("external/lux.shp", package="raster"))
  b <- as(extent(6, 6.4, 49.75, 50), 'SpatialPolygons')
  projection(b) <- projection(p)
  i <- intersect(p, b)
  plot(p)
  plot(b, add=TRUE, col='red')
  plot(i, add=TRUE, col='blue', lwd=2)
}
```

---

 isLonLat

---

*Is this longitude/latitude data?*


---

**Description**

Test whether a Raster\* or other object has a longitude/latitude coordinate reference system (CRS) by inspecting the PROJ.4 coordinate reference system description. `couldBeLonLat` also returns TRUE if the CRS is NA but the x coordinates are within -365 and 365 and the y coordinates are within -90.1 and 90.1.

**Usage**

```
## S4 method for signature 'BasicRaster'
isLonLat(x, ...)
## S4 method for signature 'Spatial'
isLonLat(x, ...)
## S4 method for signature 'BasicRaster'
couldBeLonLat(x, warnings=TRUE, ...)
## S4 method for signature 'Spatial'
couldBeLonLat(x, warnings=TRUE, ...)
```

**Arguments**

x	Raster* or Spatial* object
warnings	logical. If TRUE, a warning is given if the CRS is NA or when the CRS is longitude/latitude but the coordinates do not match that
...	additional arguments. None implemented

**Value**

Logical

**Examples**

```
r <- raster()
isLonLat(r)
crs(r) <- "+proj=lcc +lat_1=48 +lat_2=33 +lon_0=-100 +ellps=WGS84"
isLonLat(r)
```

---

KML

*Write a KML or KMZ file*

---

**Description**

Export raster data to a KML file and an accompanying PNG image file. Multi-layer objects can be used to create an animation. The function attempts to combine these into a single (and hence more convenient) KMZ file (a zip file containing the KML and PNG files).

See package `plotKML` for more advanced functionality

**Usage**

```
## S4 method for signature 'RasterLayer'
KML(x, filename, col=rev(terrain.colors(255)),
    colNA=NA, maxpixels=100000, blur=1, zip='', overwrite=FALSE, ...)

## S4 method for signature 'RasterStackBrick'
KML(x, filename, time=NULL, col=rev(terrain.colors(255)),
```

```
colNA=NA, maxpixels=100000, blur=1, zip='', overwrite=FALSE, ...)

## S4 method for signature 'Spatial'
KML(x, filename, zip='', overwrite=FALSE, ...)
```

### Arguments

x	Raster* object
filename	output filename
time	character vector with time labels for multilayer objects. The length of this vector should be nlayers(x) to indicate "when" or nlayers(x)+1 to indicate "begin-end"
col	color scheme to be used (see <a href="#">image</a> )
colNA	The color to use for the background (default is transparent)
maxpixels	maximum number of pixels. If ncell(raster) > maxpixels, sampleRegular is used to reduce the number of pixels
blur	Integer (default=1). Higher values help avoid blurring of isolated pixels (at the expense of a png file that is blur^2 times larger)
zip	If there is no zip program on your path (on windows), you can supply the full path to a zip.exe here, in order to make a KMZ file
overwrite	logical. If TRUE, overwrite the file if it exists
...	If x is a Raster* object, additional arguments that can be passed to <a href="#">image</a>

### Value

None. Used for the side-effect files written to disk.

### Author(s)

This function was adapted for the raster package by Robert J. Hijmans, with ideas from Tony Fischbach, and based on functions in the mapproj package by Duncan Goulet, David Forrest and Roger Bivand.

### Examples

```
## Not run:
# Meuse data from the sp package
data(meuse.grid)
b <- rasterFromXYZ(meuse.grid)
projection(b) <- "+init=epsg:28992"
# transform to longitude/latitude
p <- projectRaster(b, crs="+proj=longlat +datum=WGS84", method='ngb')
KML(p, file='meuse.kml')

## End(Not run)
```

layerize

*Layerize***Description**

Create a RasterBrick with a Boolean layer for each class (value, or subset of the values) in a RasterLayer. For example, if the cell values of a RasterLayer indicate what vegetation type they are, this function will create a layer (presence/absence; dummy variable) for each of these classes. Classes and cell values are always truncated to integers.

You can supply a second spatially overlapping RasterLayer with larger cells (do not use smaller cells!). In this case the cell values are counts for each class. A similar result might be obtained more efficiently by using layerize with a single RasterLayer followed by [aggregate](#)(x, , sum).

**Usage**

```
## S4 method for signature 'RasterLayer,missing'
layerize(x, classes=NULL, falseNA=FALSE, filename='', ...)

## S4 method for signature 'RasterLayer,RasterLayer'
layerize(x, y, classes=NULL, filename='', ...)
```

**Arguments**

x	RasterLayer
y	RasterLayer or missing
classes	numeric. The values (classes) for which layers should be made. If NULL all classes are used
falseNA	logical. If TRUE, cells that are not of the class represented by a layer are NA rather than FALSE
filename	character. Output filename (optional)
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterBrick

**Examples**

```
r <- raster(nrow=20, ncol=20)
values(r) <- c(rep(NA, 50), rep(1:5, 70))
b <- layerize(r)

r2 <- raster(nrow=5, ncol=5)
b2 <- layerize(r, r2)
```

---

layerStats	<i>Correlation and (weighted) covariance</i>
------------	--

---

### Description

Compute correlation and (weighted) covariance for multi-layer Raster objects. Like [cellStats](#) this function returns a few values, not a Raster\* object (see [Summary-methods](#) for that).

### Usage

```
layerStats(x, stat, w, asSample=TRUE, na.rm=FALSE, ...)
```

### Arguments

x	RasterStack or RasterBrick for which to compute a statistic
stat	Character. The statistic to compute: either 'cov' (covariance), 'weighted.cov' (weighted covariance), or 'pearson' (correlation coefficient)
w	RasterLayer with the weights (should have the same extent, resolution and number of layers as x) to compute the weighted covariance
asSample	Logical. If TRUE, the statistic for a sample (denominator is n-1) is computed, rather than for the population (denominator is n)
na.rm	Logical. Should missing values be removed?
...	Additional arguments (none implemented)

### Value

List with two items: the correlation or (weighted) covariance matrix, and the (weighted) means.

### Author(s)

Jonathan A. Greenberg & Robert Hijmans. Weighted covariance based on code by Mort Canty

### References

For the weighted covariance:

- Canty, M.J. and A.A. Nielsen, 2008. Automatic radiometric normalization of multitemporal satellite imagery with the iteratively re-weighted MAD transformation. *Remote Sensing of Environment* 112:1025-1036.
- Nielsen, A.A., 2007. The regularized iteratively reweighted MAD method for change detection in multi- and hyperspectral data. *IEEE Transactions on Image Processing* 16(2):463-478.

### See Also

[cellStats](#), [cov.wt](#), [weighted.mean](#)

**Examples**

```

b <- brick(system.file("external/rlogo.grd", package="raster"))
layerStats(b, 'pearson')

layerStats(b, 'cov')

# weigh by column number
w <- init(b, v='col')
layerStats(b, 'weighted.cov', w=w)

```

---

localFun

*Local functions*


---

**Description**

Local functions for two RasterLayer objects (using a focal neighborhood)

**Usage**

```

## S4 method for signature 'RasterLayer,RasterLayer'
localFun(x, y, ngb=5, fun, filename='', ...)

```

**Arguments**

x	RasterLayer or RasterStack/RasterBrick
y	object of the same class as x, and with the same number of layers
ngb	integer. rectangular neighbourhood size. Either a single integer or a vector of two integers c(rows, cols), such as c(3,3) to have a 3 x 3 focal window
fun	function
filename	character. Output filename (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer

**Note**

The first two arguments that fun needs to accept are vectors representing the local cells of RasterLayer x and y (each of length ngb \* ngb). It also must have an ellipsis (...) argument

**See Also**

[corLocal](#), [localFun](#)



**Examples**

```
set.seed(0)
b <- stack(system.file("external/rlogo.grd", package="raster"))
x <- flip(b[[2]], 'y') + runif(ncell(b))
y <- b[[1]] + runif(ncell(b))

f <- localFun(x, y, fun=cor)

## Not run:
# local regression:
rfun <- function(x, y, ...) {
  m <- lm(y~x)
  # return R^2
  summary(m)$r.squared
}

ff <- localFun(x, y, fun=rfun)
plot(f, ff)

## End(Not run)
```

---

Logic-methods

*Logical operators and functions*

---

**Description**

The following logical (boolean) operators are available for computations with RasterLayer objects:

&, |, and !

The following functions are available with a Raster\* argument:

is.na, is.nan, is.finite, is.infinite

**Value**

A Raster object with logical (TRUE/FALSE values)

**Note**

These are convenient operators/functions that are most useful for relatively small RasterLayers for which all the values can be held in memory. If the values of the output RasterLayer cannot be held in memory, they will be saved to a temporary file. In that case it could be more efficient to use [calc](#) instead.

**See Also**

[Math-methods](#), [overlay](#), [calc](#)

**Examples**

```

r <- raster(ncols=10, nrows=10)
values(r) <- runif(ncell(r)) * 10
r1 <- r < 3 | r > 6
r2 <- !r1
r3 <- r >= 3 & r <= 6
r4 <- r2 == r3
r[r>3] <- NA
r5 <- is.na(r)
r[1:5]
r1[1:5]
r2[1:5]
r3[1:5]

```

---

mask

*Mask values in a Raster object*


---

**Description**

Create a new Raster\* object that has the same values as x, except for the cells that are NA (or other maskvalue) in a 'mask'. These cells become NA (or other updatevalue). The mask can be either another Raster\* object of the same extent and resolution, or a Spatial\* object (e.g. SpatialPolygons) in which case all cells that are not covered by the Spatial object are set to updatevalue. You can use inverse=TRUE to set the cells that are not NA (or other maskvalue) in the mask, or not covered by the Spatial\* object, to NA (or other updatevalue).

**Usage**

```

## S4 method for signature 'RasterLayer,RasterLayer'
mask(x, mask, filename="", inverse=FALSE,
      maskvalue=NA, updatevalue=NA, updateNA=FALSE, ...)

## S4 method for signature 'RasterStackBrick,RasterLayer'
mask(x, mask, filename="", inverse=FALSE,
      maskvalue=NA, updatevalue=NA, updateNA=FALSE, ...)

## S4 method for signature 'RasterLayer,RasterStackBrick'
mask(x, mask, filename="", inverse=FALSE,
      maskvalue=NA, updatevalue=NA, updateNA=FALSE, ...)

## S4 method for signature 'RasterStackBrick,RasterStackBrick'
mask(x, mask, filename="", inverse=FALSE,
      maskvalue=NA, updatevalue=NA, updateNA=FALSE, ...)

## S4 method for signature 'Raster,Spatial'
mask(x, mask, filename="", inverse=FALSE,
      updatevalue=NA, updateNA=FALSE, ...)

```

**Arguments**

x	Raster* object
mask	Raster* object or a Spatial* object
filename	character. Optional output filename
inverse	logical. If TRUE, areas on mask that are <i>_not_</i> the maskvalue are masked
maskvalue	numeric. The value in mask that indicates the cells of x that should become updatevalue (default = NA)
updatevalue	numeric. The value that cells of x should become if they are not covered by mask (and not NA)
updateNA	logical. If TRUE, NA values outside the masked area are also updated to the the updatevalue (only relevant if the updatevalue is not NA)
...	additional arguments as in <a href="#">writeRaster</a>

**Value**

Raster\* object

**See Also**

[rasterize](#), [crop](#)

**Examples**

```
r <- raster(ncol=10, nrow=10)
m <- raster(ncol=10, nrow=10)
values(r) <- runif(ncell(r)) * 10
values(m) <- runif(ncell(r))
m[m < 0.5] <- NA
mr <- mask(r, m)

m2 <- m > .7
mr2 <- mask(r, m2, maskvalue=TRUE)
```

---

match

*Value matching for Raster\* objects*

---

**Description**

match returns a Raster\* object with the position of the matched values. The cell values are the index of the table argument.

%in% returns a logical Raster\* object indicating if the cells values were matched or not.

**Usage**

```
match(x, table, nomatch = NA_integer_, incomparables = NULL)
```

```
x %in% table
```

**Arguments**

x	Raster* object
table	vector of the values to be matched against
nomatch	the value to be returned in the case when no match is found. Note that it is coerced to integer
incomparables	a vector of values that cannot be matched. Any value in x matching a value in this vector is assigned the nomatch value. For historical reasons, FALSE is equivalent to NULL

**Value**

Raster\* object

**See Also**

[calc](#), [match](#)

**Examples**

```
r <- raster(nrow=10, ncol=10)
values(r) <- 1:100
m <- match(r, c(5:10, 50:55))
n <- r %in% c(5:10, 50:55)
```

---

Math-methods

*Mathematical functions*

---

**Description**

Generic mathematical functions that can be used with a Raster\* object as argument:

"abs", "sign", "sqrt", "ceiling", "floor", "trunc", "cummax", "cummin",  
 "cumprod", "cumsum", "log", "log10", "log2", "log1p", "acos", "acosh", "asin",  
 "asinh", "atan", "atanh", "exp", "expm1", "cos", "cosh", "sin", "sinh", "tan", "tanh".

**Note**

You can use the, somewhat more flexible, function [calc](#) instead of the Math-methods.

**See Also**

[Arith-methods](#), [calc](#), [overlay](#), [atan2](#)

**Examples**

```
r1 <- raster(nrow=10, ncol=10)
r1 <- setValues(r1, runif(ncell(r1)) * 10)
r2 <- sqrt(r1)
s <- stack(r1, r2) - 5
b <- abs(s)
```

merge

*Merge Raster\* objects***Description**

Merge Raster\* objects to form a new Raster object with a larger spatial extent. If objects overlap, the values get priority in the same order as the arguments, but NA values are ignored (except when `overlap=FALSE`). See [subs](#) to merge a Raster\* object and a `data.frame`.

**Usage**

```
## S4 method for signature 'Raster,Raster'
merge(x, y, ..., tolerance=0.05, filename="", overlap=TRUE, ext=NULL)

## S4 method for signature 'RasterStackBrick,missing'
merge(x, ..., tolerance=0.05, filename="", ext=NULL)

## S4 method for signature 'Extent,ANY'
merge(x, y, ...)
```

**Arguments**

<code>x</code>	Raster* or Extent object
<code>y</code>	Raster* if <code>x</code> is a Raster* object (or missing). If <code>x</code> is an Extent, <code>y</code> can be an Extent or object from which an Extent can be extracted
<code>...</code>	additional Raster or Extent objects (and/or arguments for writing files as in <a href="#">writeRaster</a> )
<code>tolerance</code>	numeric. permissible difference in origin (relative to the cell resolution). See <a href="#">all.equal</a>
<code>filename</code>	character. Output filename (optional)
<code>overlap</code>	logical. If <code>FALSE</code> values of overlapping objects are based on the first layer, even if they are NA
<code>ext</code>	Extent object (optional) to limit the output to that extent

**Details**

The Raster objects must have the same origin and resolution. In areas where the Raster objects overlap, the values of the Raster object that is first in the sequence of arguments will be retained. If you would rather use the average of cell values, or do another computation, you can use [mosaic](#) instead of `merge`.

**Value**

RasterLayer or RasterBrick

**Examples**

```
r1 <- raster(xmx=-150, ymn=60, ncols=30, nrows=30)
values(r1) <- 1:ncell(r1)
r2 <- raster(xmn=-100, xmx=-50, ymx=50, ymn=30)
res(r2) <- c(xres(r1), yres(r1))
values(r2) <- 1:ncell(r2)
rm <- merge(r1, r2)

# if you have many RasterLayer objects in a list
# you can use do.call:
x <- list(r1, r2)
# add arguments such as filename
# x$filename <- 'test.tif'
m <- do.call(merge, x)
```

---

metadata

*Metadata*

---

**Description**

Get or set a metadata to a Raster object

**Usage**

```
metadata(x)
metadata(x) <- value
```

**Arguments**

x	Raster* object
value	list with named elements. Each element may be another list of named elements (but these nested lists are not allowed to be lists themselves)

**Value**

Raster\* object or list

**Note**

The metadata can contain single values or vectors of basic data types (character, integer, numeric) and Date. Some other types may also be supported. You cannot use a matrix or data.frame as a meta-data element.

**Examples**

```

r <- raster(nc=10, nr=10)
values(r) <- 1:ncell(r)

m <- list(wave=list(a=1, b=2, c=c('cool', 'important')), that=list(red='44', blue=1:5,
  days=as.Date(c('2014-1-15', '2014-2-15'))), this='888 miles from here', today=NA)

metadata(r) <- m

## Not run:

x <- writeRaster(r, rasterTmpFile(), overwrite=TRUE)
metax <- metadata(x)

identical(metax, m)

# nested too deep
badmeta1 <- list(wave=list(a=1, b=2, c='x'), that=list(red='4', blue=list(bad=5)))
metadata(r) <- badmeta1

# missing names
badmeta2 <- list(wave=list(1, 2, c='x'), that=list(red='44', blue=14, this='8m'))
metadata(r) <- badmeta2

# matrix not allowed
badmeta3 <- list(wave=list(a=1, b=matrix(1:4, ncol=2), c='x'), that=list(red='4'))
metadata(r) <- badmeta3

## End(Not run)

```

---

modal

*modal value*


---

**Description**

Compute the mode for a vector of numbers, or across raster layers. The mode, or modal value, is the most frequent value in a set of values.

**Usage**

```

## S4 method for signature 'ANY'
modal(x, ..., ties='random', na.rm=FALSE, freq=FALSE)

## S4 method for signature 'Raster'
modal(x, ..., ties='random', na.rm=FALSE, freq=FALSE)

```

**Arguments**

x	vector of numbers (typically integers), characters, logicals, or factors, or a Raster* object
...	additional argument of the same type as x
ties	character. Indicates how to treat ties. Either 'random', 'lowest', 'highest', 'first', or 'NA'
na.rm	logical. If TRUE, NA values are ignored. If FALSE, NA is returned if x has any NA values
freq	return the frequency of the modal value, instead of the modal value

**Value**

vector or RasterLayer. The vector has length 1 and is of the same type as x, except when x is a factor and additional arguments (values) are supplied, in which case the values are coerced to characters and a character value is returned.

**Examples**

```
data <- c(0,1,2,3,3,3,3,4,4,4,5,5,6,7,7,8,9,NA)
modal(data, na.rm=TRUE)
```

---

mosaic

---

*Merge Raster\* objects using a function for overlapping areas*


---

**Description**

Mosaic Raster\* objects to form a new object with a larger spatial extent. A function is used to compute cell values in areas where layers overlap (in contrast to the [merge](#) function which uses the values of the 'upper' layer). All objects must have the same origin, resolution, and coordinate reference system.

**Usage**

```
## S4 method for signature 'Raster,Raster'
mosaic(x, y, ..., fun, tolerance=0.05, filename="")
```

**Arguments**

x	Raster* object
y	Raster* object
...	Additional Raster or Extent objects (and/or arguments for writing files as in <a href="#">writeRaster</a> )
fun	Function. E.g. mean, min, or max. Must be a function that accepts a 'na.rm' argument
tolerance	Numeric. permissible difference in origin (relative to the cell resolution). See <a href="#">all.equal</a>
filename	Character. Output filename (optional)



**Details**

The Raster objects must have the same origin and resolution.

**Value**

RasterLayer or RasterBrick object.

**See Also**

[merge](#), [extend](#)

**Examples**

```
r <- raster(ncol=100, nrow=100)
r1 <- crop(r, extent(-10, 11, -10, 11))
r2 <- crop(r, extent(0, 20, 0, 20))
r3 <- crop(r, extent(9, 30, 9, 30))

values(r1) <- 1:ncell(r1)
values(r2) <- 1:ncell(r2)
values(r3) <- 1:ncell(r3)

m1 <- mosaic(r1, r2, r3, fun=mean)

s1 <- stack(r1, r1*2)
s2 <- stack(r2, r2/2)
s3 <- stack(r3, r3*4)
m2 <- mosaic(s1, s2, s3, fun=min)

# if you have a list of Raster objects, you can use do.call
x <- list(r1, r2, r3)
names(x)[1:2] <- c('x', 'y')
x$fun <- mean
x$na.rm <- TRUE

y <- do.call(mosaic, x)
```

---

movingFun

*Moving functions*

---

**Description**

Helper function to compute 'moving' functions, such as the 'moving average'

**Usage**

```
movingFun(x, n, fun=mean, type='around', circular=FALSE, na.rm=FALSE)
```

**Arguments**

x	A vector of numbers
n	Size of the 'window', i.e. the number of sequential elements to use in the function
fun	A function like mean, min, max, sum
type	Character. One of 'around', 'to', or 'from'. The choice indicates which values should be used in the computation. The focal element is always used. If type is 'around', the other elements are before and after the focal element. Alternatively, you can select the elements preceding the focal element ('to') or those coming after it 'from'. For example, to compute the movingFun with n=3 for element 5 of a vector; 'around' used elements 4,5,6; 'to' used elements 3,4,5, and 'from' uses elements 5,6,7
circular	Logical. If TRUE, the data are considered to have a circular nature (e.g. months of the year), and the last elements in vector x are used in the computation of the moving function of the first element(s) of the vector, and the first elements are used in the computation of the moving function for the last element(s)
na.rm	Logical. If TRUE, NA values should be ingored (by fun)

**Value**

Numeric

**Author(s)**

Robert J. Hijmans, inspired by Diethelm Wuertz' rollFun function in the fTrading package

**Examples**

```

movingFun(1:12, 3, mean)
movingFun(1:12, 3, mean, 'to')
movingFun(1:12, 3, mean, 'from')
movingFun(1:12, 3, mean, circular=TRUE)

v <- c(0,1,2,3,3,3,3,4,4,4,5,5,6,7,7,8,9,NA)
movingFun(v, n=5)
movingFun(v, n=5, na.rm=TRUE)

```

---

names

*Names of raster layers*

---

**Description**

Get or set the names of the layers of a Raster\* object

**Usage**

```
## S4 method for signature 'Raster'  
names(x)  
  
## S4 replacement method for signature 'Raster'  
names(x)<-value  
  
## S4 method for signature 'Raster'  
labels(object)
```

**Arguments**

x	Raster* object
object	Raster* object
value	character (vector)

**Value**

Character

**See Also**

[nlayers](#), [bands](#)

**Examples**

```
r <- raster(ncols=5, nrows=5)  
values(r) <- 1:ncell(r)  
s <- stack(r, r, r)  
nlayers(s)  
names(s)  
names(s) <- c('a', 'b', 'c')  
names(s)[2] <- 'hello world'  
names(s)  
s  
labels(s)
```

---

NAvalue

*Set the NA value of a RasterLayer*

---

**Description**

NAvalue returns the value that is used to write NA values to disk (in 'raster' type files). If you set the NA value of a Raster\* object, this value will be interpreted as NA when reading the values from a file. Values already in memory will not be affected.

If the NA value is smaller than zero, all values smaller or equal to that number will be set to NA.

**Usage**

```
NAvalue(x) <- value
NAvalue(x)
```

**Arguments**

x	A Raster object
value	the value to be interpreted as NA; set this before reading the values from the file. Integer values are matched exactly; for decimal values files any value <= the value will be interpreted as NA

**Value**

Returns or set the NA value used for storage on disk.

**Examples**

```
r1 <- raster(system.file("external/rlogo.grd", package="raster"))
r2 <- r1
NAvalue(r2)
NAvalue(r2) <- 255
#plot(r1)
#x11()
#plot(r2)
```

---

ncell

*Number or rows, columns, and cells of a Raster\* object*


---

**Description**

Get the number of rows, columns, or cells of a Raster\* object.

**Usage**

```
ncol(x)
nrow(x)
ncell(x)
ncol(x, ...) <- value
nrow(x, ...) <- value
```

**Arguments**

x	a Raster object
value	row or column number (integer > 0)
...	additional arguments. None implemented

**Value**

Integer

**See Also**[dim](#), [extent](#), [res](#)**Examples**

```
r <- raster()
ncell(r)
ncol(r)
nrow(r)
dim(r)

nrow(r) <- 18
ncol(r) <- 36
# equivalent to
dim(r) <- c(18, 36)
```

---

nlayers
---------

---

*Number of layers***Description**

Get the number of layers in a Raster\* object, typically used with a (multilayer) RasterStack or RasterBrick object

**Usage**

```
nlayers(x)
```

**Arguments**

x	Raster* object
---	----------------

**Value**

integer

**See Also**[names](#)

**Examples**

```
r <- raster(ncols=10, nrows=10)
values(r) <- 1:ncell(r)
s <- stack(r, r, r)
nlayers(s)
s <- stack(s,s)
nlayers(s)
s <- dropLayer(s, 2:3)
nlayers(s)
```

---

**Options***Global options for the raster package*

---

**Description**

Set, inspect, reset, save a number of global options used by the raster package.

Most of these options are used when writing files to disk. They can be ignored by specific functions if the corresponding argument is provided as an argument to these functions.

The default location is returned by `rasterTmpDir`. It is the same as that of the R temp directory but you can change it (for the current session) with `rasterOptions(tmpdir="path")`.

To permanently set any of these options, you can add them to `<your R installation>/etc/Rprofile.site`. For example, to change the default directory used to save temporary files, add a line like this: `options(rasterTmpDir='c:/temp/')` to that file. All temporary raster files in that folder that are older than 24 hrs are deleted when the raster package is loaded.

Function `tmpDir` returns the location of the temporary files

**Usage**

```
rasterOptions(format, overwrite, datatype, tmpdir, tmptime, progress,
              timer, chunksize, maxmemory, memfrac, todisk, setfileext, tolerance,
              standardnames, deprecatedwarnings, addheader, default=FALSE)
```

```
tmpDir(create=TRUE)
```

**Arguments**

<code>format</code>	character. The default file format to use. See <a href="#">writeFormats</a>
<code>overwrite</code>	logical. The default value for overwriting existing files. If TRUE, existing files will be overwritten
<code>datatype</code>	character. The default data type to use. See <a href="#">dataType</a>
<code>tmpdir</code>	character. The default location for writing temporary files; See <a href="#">rasterTmpFile</a>
<code>tmptime</code>	number > 1. The number of hours after which a temporary file will be deleted. As files are deleted when loading the raster package, this option is only useful if you save this option so that it is loaded when starting a new session

progress	character. Valid values are "text", "window" and "" (the default in most functions, no progress bar)
timer	Logical. If TRUE, the time it took to complete the function is printed
chunksize	integer. Maximum number of bytes to read/write in a single chunk while processing (chunk by chunk) disk based Raster* objects
maxmemory	numeric. Maximum number of bytes to read into memory. If a process is expected to require more than this value, <a href="#">canProcessInMemory</a> will return FALSE
memfrac	numeric. Fraction of available RAM that may be used by a process
todisk	logical. For debugging only. Default is FALSE and should normally not be changed. If TRUE, results are always written to disk, even if no filename is supplied (a temporary filename is used)
setfileext	logical. Default is TRUE. If TRUE, the file extension will be changed when writing (if known for the file type). E.g. GTiff files will be saved with the .tif extension
tolerance	numeric. The tolerance used when comparing the origin and resolution of Raster* objects. Expressed as the fraction of a single cell. This should be a number between 0 and 0.5
standardnames	logical. Default is TRUE. Should <a href="#">names</a> be standardized to be syntactically valid names (using <a href="#">make.names</a> )
depracatedwarnings	logical. If TRUE (the default) a warning is generated when a deprecated (obsolete) function is used
addheader	character. If not equal to '' (the default) an additional header file is written when a raster format file (grd/gri) is written. Supported formats are as in <a href="#">hdr</a>
default	logical. If TRUE, all options are set to their default values
create	logical. If TRUE, the temporary files directory is created if it does not exist

**Value**

list of the current options (invisibly). If no arguments are provided the options are printed.

**See Also**

[options](#), [rasterTmpFile](#)

**Examples**

```
## Not run:
rasterOptions()
rasterOptions(chunksize=2e+07)

## End(Not run)
```

---

origin	<i>Origin</i>
--------	---------------

---

### Description

Origin returns (or sets) the coordinates of the point of origin of a Raster\* object. This is the point closest to (0, 0) that you could get if you moved towards that point in steps of the x and y resolution.

### Usage

```
origin(x, ...)  
origin(x) <- value
```

### Arguments

x	Raster* object
value	numeric vector of length 1 or 2
...	additional arguments. None implemented

### Value

A vector of two numbers (x and y coordinates), or a changed origin for x.

### See Also

[extent](#)

### Examples

```
r <- raster(xmn=-0.5, xmx = 9.5, ncols=10)  
origin(r)  
r  
origin(r) <- 0  
r
```

---

overlay	<i>Overlay Raster objects</i>
---------	-------------------------------

---



## Description

Create a new Raster\* object, based on two or more Raster\* objects. (You can also use a single object, but perhaps `calc` is what you are looking for in that case).

You should supply a function `fun` to set the way that the RasterLayers are combined. The number of arguments in the function must match the number of Raster objects (or take any number). For example, if you combine two RasterLayers you could use `multiply: fun=function(x,y){return(x*y)}` `percentage: fun=function(x,y){return(100 * x / y)}`. If you combine three layers you could use `fun=function(x,y,z){return((x + y) * z)}`

Note that the function must work for vectors (not only for single numbers). That is, it must return the same number of elements as its input vectors. Alternatively, you can also supply a function such as `sum`, that takes `n` arguments (as `'...'`), and perhaps also has a `na.rm` argument, like in `sum(...,na.rm)`.

If a single multi-layer object is provided, its layers are treated as individual RasterLayer objects if the argument `unstack=TRUE` is used. If multiple objects are provided, they should have the same number of layers, or it should be possible to recycle them (e.g., 1, 3, and 9 layers, which would return a RasterBrick with 9 layers).

## Usage

```
## S4 method for signature 'Raster,Raster'
overlay(x, y, ..., fun, filename="", recycle=TRUE, forcefun=FALSE)

## S4 method for signature 'Raster,missing'
overlay(x, y, ..., fun, filename="", unstack=TRUE, forcefun=FALSE)
```

## Arguments

<code>x</code>	Raster* object
<code>y</code>	Raster* object, or missing (only useful if <code>x</code> has multiple layers)
<code>...</code>	Additional Raster objects (and/or arguments for writing files as in <code>writeRaster</code> )
<code>fun</code>	Function to be applied. When using RasterLayer objects, the number of arguments of the function should match the number of Raster objects, or it should take any number of arguments. When using multi-layer objects the function should match the number of layers of the RasterStack/Brick object (unless <code>unstack=FALSE</code> )
<code>filename</code>	Character. Output filename (optional)
<code>recycle</code>	Logical. Should layers from Raster objects with fewer layers be recycled?
<code>unstack</code>	Logical. Should layers be unstacked before computation (i.e. does the <code>fun</code> refer to individual layers in a multilayer object)?
<code>forcefun</code>	Boolean. If TRUE, <code>overlay</code> will not attempt to internally use <code>apply</code> (it is rarely necessary to use this argument)

## Details

Instead of the `overlay` function you can also use arithmetic functions such as `*`, `/`, `+`, `-` with Raster objects (see examples). In that case you cannot specify an output filename. Moreover, the `overlay` function should be more efficient when using large data files that cannot be loaded into memory, as the use of the complex arithmetic functions might lead to the creation of many temporary files.

While you can supply functions such as `sum` or `mean`, it would be more direct to use the Raster\* objects as arguments to those functions (e.g. `sum(r1,r2,r3)`)

See [rasterize](#) and [extract](#) for "overlays" involving Raster\* objects and polygons, lines, or points.

## Value

Raster\* object

## See Also

[calc](#), [Arith-methods](#)

## Examples

```
r <- raster(ncol=10, nrow=10)
r1 <- init(r, fun=runif)
r2 <- init(r, fun=runif)
r3 <- overlay(r1, r2, fun=function(x,y){return(x+y)})

# long version for multiplication
r4 <- overlay(r1, r2, fun=function(x,y){(x*y)} )

#use the individual layers of a RasterStack to get a RasterLayer
s <- stack(r1, r2)
r5 <- overlay(s, fun=function(x,y) x*y )
# equivalent to
r5c <- calc(s, fun=function(x) x[1]*x[2] )

#Combine RasterStack and RasterLayer objects (s2 has four layers.
# r1 (one layer) and s (two layers) are recycled)
s2 <- stack(r1, r2, r3, r4)
b <- overlay(r1, s, s2, fun=function(x,y,z){return(x*y*z)} )

# use a single RasterLayer (same as calc function)
r6 <- overlay(r1, fun=sqrt)

# multiplication with more than two layers
# (make sure the number of RasterLayers matches the arguments of 'fun')
r7 <- overlay(r1, r2, r3, r4, fun=function(a,b,c,d){return(a*b+c*d)} )
# equivalent function, efficient if values can be loaded in memory
r8 <- r1 * r2 + r3 * r4

# Also works with multi-layer objects.
s1 <- stack(r1, r2, r3)
x <- overlay(s1, s1, fun=function(x,y)x+y+5)
```

```
# in this case the first layer of the shorter object is recycled.
# i.e., s2 is treated as stack(r1, r3, r1)
s2 <- stack(r1, r3)
y <- overlay(s1, s2, fun=sum)
```

---

pairs

*Pairs plot (matrix of scatterplots)*

---

### Description

Pair plots of layers in a RasterStack or RasterBrick. This is a wrapper around graphics function [pairs](#).

### Usage

```
## S4 method for signature 'RasterStackBrick'
pairs(x, hist=TRUE, cor=TRUE, use="pairwise.complete.obs", maxpixels=100000, ...)
```

### Arguments

x	RasterBrick or RasterStack
hist	Logical. If TRUE a histogram of the values is shown on the diagonal
cor	Logical. If TRUE the correlation coefficient is shown in the upper panels
use	Argument passed to the <a href="#">cor</a> function
maxpixels	Integer. Number of pixels to sample from each layer of large Raster objects
...	Additional arguments (only cex and main)

### See Also

[boxplot](#), [hist](#), [density](#)

### Examples

```
r <- raster(system.file("external/test.grd", package="raster") )
s <- stack(r, 1/r, sqrt(r))
pairs(s)

## Not run:
# to make individual histograms:
hist(r)
# or scatter plots:
plot(r, 1/r)

## End(Not run)
```

---

persp                      *Perspective plot*

---

### Description

Perspective plot of a RasterLayer. This is an implementation of a generic function in the graphics package.

### Usage

```
## S4 method for signature 'RasterLayer'
persp(x, maxpixels=1e+05, ext=NULL, ...)

## S4 method for signature 'RasterStackBrick'
persp(x, y=1, maxpixels=10000, ext=NULL, ...)
```

### Arguments

x	Raster* object
y	integer > 0 & <= nlayers(x) to select the layer of x if x is a RasterLayer or RasterBrick
maxpixels	integer > 0. Maximum number of cells to use for the plot. If maxpixels < ncell(x), sampleRegular is used before plotting
ext	Extent. Can be used to zoom in to a region (see also <a href="#">zoom</a> and <a href="#">crop(x, drawExtent())</a> )
...	Any argument that can be passed to <a href="#">persp</a> (graphics package)

### See Also

[plot3D](#), [persp](#), [contour](#), [plot](#)

### Examples

```
r <- raster(system.file("external/test.grd", package="raster"))
persp(r)
```

---

plot *Plot a Raster\* object*

---

### Description

Plot (that is, make a map of) the values of a Raster\* object, or make a scatterplot of their values.

Points, lines, and polygons can be drawn on top of a map using `plot(..., add=TRUE)`, or with functions like `points`, `lines`, `polygons`

See the `rasterVis` package for more advanced (trellis/lattice) plotting of Raster\* objects.

### Usage

```
## S4 method for signature 'Raster,ANY'
plot(x, y, maxpixels=500000, col, alpha=NULL,
     colNA=NA, add=FALSE, ext=NULL, useRaster=TRUE, interpolate=FALSE,
     addfun=NULL, nc, nr, maxnl=16, main, npretty=0, ...)
```

```
## S4 method for signature 'Raster,Raster'
plot(x, y, maxpixels=100000, cex, xlab, ylab, nc, nr,
     maxnl=16, main, add=FALSE, gridded=FALSE, ncol=25, nrow=25, ...)
```

### Arguments

<code>x</code>	Raster* object
<code>y</code>	If <code>x</code> is a RasterStack or RasterBrick: integer, character (layer name(s)), or missing to select which layer(s) to plot. If missing, all RasterLayers in the RasterStack will be plotted (up to a maximum of 16). Or another Raster* object of the same extent and resolution, to produce a scatter plot of the cell values.
<code>maxpixels</code>	integer > 0. Maximum number of cells to use for the plot. If <code>maxpixels &lt; ncell(x)</code> , <code>sampleRegular</code> is used before plotting. If <code>gridded=TRUE</code> <code>maxpixels</code> may be ignored to get a larger sample
<code>col</code>	A color palette, i.e. a vector of <code>n</code> contiguous colors generated by functions like <a href="#">rainbow</a> , <a href="#">heat.colors</a> , <a href="#">topo.colors</a> , <a href="#">bpy.colors</a> or one or your own making, perhaps using <a href="#">colorRampPalette</a> . If none is provided, <code>rev(terrain.colors(255))</code> is used unless <code>x</code> has a 'color table'
<code>alpha</code>	Number between 0 and 1 to set transparency. 0 is entirely transparent, 1 is not transparent (NULL is equivalent to 1)
<code>colNA</code>	The color to use for the background (default is transparent)
<code>add</code>	Logical. Add to current plot?
<code>ext</code>	An extent object to zoom in a region (see also <a href="#">zoom</a> and <code>crop(x, drawExtent())</code> )
<code>useRaster</code>	If TRUE, the <code>rasterImage</code> function is used for plotting. Otherwise the <code>image</code> function is used. This can be useful if <code>rasterImage</code> does not work well on your system (see note)

interpolate	Logical. Should the image be interpolated (smoothed)? Only used when useRaster = TRUE
addfun	Function to add additional items such as points or polygons to the plot (map). Typically containing statements like "points(xy); plot(polygons, add=TRUE)". This is particularly useful to add something to each map when plotting a multi-layer Raster* object.
npretty	integer. Number of decimals for <a href="#">pretty</a> labels on the axes
...	Graphical parameters. Any argument that can be passed to <a href="#">image.plot</a> and to <a href="#">plot</a> , such as axes=FALSE, main='title', ylab='latitude'
xlab	Optional. x-axis label)
ylab	Optional. y-axis label)
nc	Optional. The number of columns to divide the plotting device in (when plotting multiple layers in a RasterLayer or RasterBrick object)
nr	Optional. The number of rows to divide the plotting device in (when plotting multiple layers in a RasterLayer or RasterBrick object)
maxnl	integer. Maximum number of layers to plot (for a multi-layer object)
main	character. Main plot title
cex	Symbol size for scatter plots
gridded	logical. If TRUE the scatterplot is gridded (counts by cells)
ncol	integer. Number of columns for gridding
nrow	integer. Number of rows for gridding

### Details

Most of the code for the plot function for a single Raster\* object was taken from `image.plot` (fields package).

Raster objects with a color-table (e.g. a graphics file) are plotted according to that color table.

### Note

`raster` uses `rasterImage` from the graphics package. For unknown reasons this does not work on Windows Server and on a few versions of Windows XP. On that system you may need to use argument `useRaster=FALSE` to get a plot.

### See Also

The `rasterVis` package has lattice based methods for plotting Raster\* objects (like [spplot](#)) red-green-blue plots (e.g. false color composites) can be made with [plotRGB](#) [barplot](#), [hist](#), [text](#), [persp](#), [contour](#), [pairs](#)

**Examples**

```

# RasterLayer
r <- raster(nrows=10, ncols=10)
r <- setValues(r, 1:ncell(r))
plot(r)

e <- extent(r)
plot(e, add=TRUE, col='red', lwd=4)
e <- e / 2
plot(e, add=TRUE, col='red')

# Scatterplot of 2 RasterLayers
r2 <- sqrt(r)
plot(r, r2)
plot(r, r2, gridded=TRUE)

# Multi-layer object (RasterStack / Brick)
s <- stack(r, r2, r/r)
plot(s, 2)
plot(s)

# two objects, different range, one scale:
values(r) <- runif(ncell(r))
r2 <- r/2
brks <- seq(0, 1, by=0.1)
nb <- length(brks)-1
cols <- rev(terrain.colors(nb))
par(mfrow=c(1,2))
plot(r, breaks=brks, col=cols, lab.breaks=brks, zlim=c(0,1), main='first')
plot(r2, breaks=brks, col=cols, lab.breaks=brks, zlim=c(0,1), main='second')

# breaks and labels
x <- raster(nc=10, nr=10)
values(x) <- runif(ncell(x))
brk <- c(0, 0.25, 0.75, 1)
arg <- list(at=c(0.12,0.5,0.87), labels=c("Low", "Med.", "High"))
plot(x, col=terrain.colors(3), breaks=brk)
plot(x, col=terrain.colors(3), breaks=brk, axis.args=arg)
par(mfrow=c(1,1))

# color ramp
plot(x, col=colorRampPalette(c("red", "white", "blue"))(255))

# adding random points to the map
xy <- cbind(-180 + runif(10) * 360, -90 + runif(10) * 180)
points(xy, pch=3, cex=5)

# for SpatialPolygons do
# plot(polys, add=TRUE)

```

```
# adding the same points to each map of each layer of a RasterStack
fun <- function() {
  points(xy, cex=2)
  points(xy, pch=3, col='red')
}
plot(s, addfun=fun)
```

---

plotRGB

*Red-Green-Blue plot of a multi-layered Raster object*


---

### Description

Make a Red-Green-Blue plot based on three layers (in a RasterBrick or RasterStack). Three layers (sometimes referred to as "bands" because they may represent different bandwidths in the electromagnetic spectrum) are combined such that they represent the red, green and blue channel. This function can be used to make 'true (or false) color images' from Landsat and other multi-band satellite images.

### Usage

```
## S4 method for signature 'RasterStackBrick'
plotRGB(x, r=1, g=2, b=3, scale, maxpixels=500000, stretch=NULL,
  ext=NULL, interpolate=FALSE, colNA='white', alpha, bgamma, addfun=NULL, zlim=NULL,
  zlimcol=NULL, axes=FALSE, xlab='', ylab='', asp=NULL, add=FALSE, margins=FALSE, ...)
```

### Arguments

x	RasterBrick or RasterStack
r	integer. Index of the Red channel, between 1 and nlayers(x)
g	integer. Index of the Green channel, between 1 and nlayers(x)
b	integer. Index of the Blue channel, between 1 and nlayers(x)
scale	integer. Maximum (possible) value in the three channels. Defaults to 255 or to the maximum value of x if that is known and larger than 255
maxpixels	integer > 0. Maximum number of pixels to use
stretch	character. Option to stretch the values to increase the contrast of the image: "lin" or "hist"
ext	An <a href="#">Extent</a> object to zoom in to a region of interest (see <a href="#">drawExtent</a> )
interpolate	logical. If TRUE, interpolate the image when drawing
colNA	color for the background (NA values)
alpha	transparency. Integer between 0 (transparent) and 255 (opaque)
bgalpha	Background transparency. Integer between 0 (transparent) and 255 (opaque)



addfun	Function to add additional items such as points or polygons to the plot (map). See <a href="#">plot</a>
zlim	numeric vector of length 2. Range of values to plot (optional)
zlimcol	If NULL the values outside the range of zlim get the color of the extremes of the range. If zlimcol has any other value, the values outside the zlim range get the color of NA values (see colNA)
axes	logical. If TRUE axes are drawn (and arguments such as main="title" will be honored)
xlab	character. Label of x-axis
ylab	character. Label of y-axis
asp	numeric. Aspect (ratio of x and y. If NULL, and appropriate value is computed to match data for the longitude/latitude coordinate reference system, and 1 for planar coordinate reference systems
add	logical. If TRUE add values to current plot
margins	logical. If TRUE standard whitespace margins are used. If FALSE, graphics::par(plt=c(0,1,0,1)) is used
...	graphical parameters as in <a href="#">plot</a> or <a href="#">rasterImage</a>

**Author(s)**

Robert J. Hijmans; stretch option based on functions by Josh Gray

**See Also**

[plot](#)

**Examples**

```
b <- brick(system.file("external/rlogo.grd", package="raster"))
plotRGB(b)
plotRGB(b, 3, 2, 1)
plotRGB(b, 3, 2, 1, stretch='hist')
```

---

pointDistance	<i>Distance between points</i>
---------------	--------------------------------

---

**Description**

Calculate the geographic distance between two (sets of) points on the WGS ellipsoid (lonlat=TRUE) or on a plane (lonlat=FALSE). If both sets do not have the same number of points, the distance between each pair of points is given. If both sets have the same number of points, the distance between each point and the corresponding point in the other set is given, except if allpairs=TRUE.

**Usage**

```
pointDistance(p1, p2, lonlat, allpairs=FALSE, ...)
```

**Arguments**

p1	x and y coordinate of first (set of) point(s), either as c(x, y), matrix(ncol=2), or SpatialPoints*.
p2	x and y coordinate of second (set of) second point(s) (like for p1). If this argument is missing, a distance matrix is computed for p1
lonlat	logical. If TRUE, coordinates should be in degrees; else they should represent planar ('Euclidean') space (e.g. units of meters)
allpairs	logical. Only relevant if the number of points in x and y is the same. If FALSE the distance between each point in x with the corresponding point in y is returned. If TRUE a full distance matrix is returned
...	Additional arguments. None implemented

**Value**

A single value, or a vector, or matrix of values giving the distance in meters (lonlat=TRUE) or map-units (for instance, meters in the case of UTM) If p2 is missing, a distance matrix is returned

**Author(s)**

Robert J. Hijmans and Jacob van Etten. The distance for longitude/latitude data uses GeographicLib by C.F.F. Karney

**See Also**

[distanceFromPoints](#), [distance](#), [gridDistance](#), [spDistsN1](#). The geosphere package has many additional distance functions and other functions that operate on spherical coordinates

**Examples**

```
a <- cbind(c(1,5,55,31),c(3,7,20,22))
b <- cbind(c(4,2,8,65),c(50,-90,20,32))

pointDistance(c(0, 0), c(1, 1), lonlat=FALSE)
pointDistance(c(0, 0), c(1, 1), lonlat=TRUE)
pointDistance(c(0, 0), a, lonlat=TRUE)
pointDistance(a, b, lonlat=TRUE)

#Make a distance matrix
dst <- pointDistance(a, lonlat=TRUE)
# coerce to dist object
dst <- as.dist(dst)
```

**Description**

Make a Raster object with predictions from a fitted model object (for example, obtained with `lm`, `glm`). The first argument is a Raster object with the independent (predictor) variables. The `names` in the Raster object should exactly match those expected by the model. This will be the case if the same Raster object was used (via `extract`) to obtain the values to fit the model (see the example). Any type of model (e.g. `glm`, `gam`, `randomForest`) for which a `predict` method has been implemented (or can be implemented) can be used.

This approach (`predict` a fitted model to raster data) is commonly used in remote sensing (for the classification of satellite images) and in ecology, for species distribution modeling.

**Usage**

```
## S4 method for signature 'Raster'
predict(object, model, filename="", fun=predict, ext=NULL,
        const=NULL, index=1, na.rm=TRUE, inf.rm=FALSE, factors=NULL,
        format, datatype, overwrite=FALSE, progress=' ', ...)
```

**Arguments**

<code>object</code>	Raster* object. Typically a multi-layer type ( <code>RasterStack</code> or <code>RasterBrick</code> )
<code>model</code>	fitted model of any class that has a 'predict' method (or for which you can supply a similar method as <code>fun</code> argument. E.g. <code>glm</code> , <code>gam</code> , or <code>randomForest</code> )
<code>filename</code>	character. Optional output filename
<code>fun</code>	function. Default value is 'predict', but can be replaced with e.g. <code>predict.se</code> (depending on the type of model), or your own custom function.
<code>ext</code>	Extent object to limit the prediction to a sub-region of <code>x</code>
<code>const</code>	<code>data.frame</code> . Can be used to add a constant for which there is no Raster object for model predictions. Particularly useful if the constant is a character-like factor value for which it is currently not possible to make a <code>RasterLayer</code>
<code>index</code>	integer. To select the column(s) to use if <code>predict.model</code> returns a matrix with multiple columns
<code>na.rm</code>	logical. Remove cells with NA values in the predictors before solving the model (and return a NA value for those cells). This option prevents errors with models that cannot handle NA values. In most other cases this will not affect the output. An exception is when predicting with a boosted regression trees model because these return predicted values even if some (or all!) variables are NA
<code>inf.rm</code>	logical. Remove cells with values that are not finite (some models will fail with -Inf/Inf values). This option is ignored when <code>na.rm=FALSE</code>

factors	list with levels for factor variables. The list elements should be named with names that correspond to names in object such that they can be matched. This argument may be omitted for standard models such as 'glm' as the predict function will extract the levels from the model object, but it is necessary in some other cases (e.g. cforest models from the party package)
format	character. Output file type. See <a href="#">writeRaster</a> (optional)
datatype	character. Output data type. See <a href="#">dataType</a> (optional)
overwrite	logical. If TRUE, "filename" will be overwritten if it exists
progress	character. "text", "window", or "" (the default, no progress bar)
...	additional arguments to pass to the predict.'model' function

**Value**

RasterLayer or RasterBrick

**Note**

For more on the use of the predict function see this resource on [species distribution modeling](#).

**See Also**

Use [interpolate](#) if your model has 'x' and 'y' as implicit independent variables (e.g., in kriging).

**Examples**

```
# A simple model to predict the location of the R in the R-logo using 20 presence points
# and 50 (random) pseudo-absence points. This type of model is often used to predict
# species distributions. See the dismo package for more of that.
```

```
# create a RasterStack or RasterBrick with with a set of predictor layers
logo <- brick(system.file("external/rlogo.grd", package="raster"))
names(logo)
```

```
## Not run:
# the predictor variables
par(mfrow=c(2,2))
plotRGB(logo, main='logo')
plot(logo, 1, col=rgb(cbind(0:255,0,0), maxColorValue=255))
plot(logo, 2, col=rgb(cbind(0,0:255,0), maxColorValue=255))
plot(logo, 3, col=rgb(cbind(0,0,0:255), maxColorValue=255))
par(mfrow=c(1,1))
```

```
## End(Not run)
```

```
# known presence and absence points
p <- matrix(c(48, 48, 48, 53, 50, 46, 54, 70, 84, 85, 74, 84, 95, 85,
  66, 42, 26, 4, 19, 17, 7, 14, 26, 29, 39, 45, 51, 56, 46, 38, 31,
  22, 34, 60, 70, 73, 63, 46, 43, 28), ncol=2)

a <- matrix(c(22, 33, 64, 85, 92, 94, 59, 27, 30, 64, 60, 33, 31, 9,
```

```

99, 67, 15, 5, 4, 30, 8, 37, 42, 27, 19, 69, 60, 73, 3, 5, 21,
37, 52, 70, 74, 9, 13, 4, 17, 47), ncol=2)

# extract values for points
xy <- rbind(cbind(1, p), cbind(0, a))
v <- data.frame(cbind(pa=xy[,1], extract(logo, xy[,2:3])))

#build a model, here an example with glm
model <- glm(formula=pa~., data=v)

#predict to a raster
r1 <- predict(logo, model, progress='text')

plot(r1)
points(p, bg='blue', pch=21)
points(a, bg='red', pch=21)

# use a modified function to get a RasterBrick with p and se
# from the glm model. The values returned by 'predict' are in a list,
# and this list needs to be transformed to a matrix

predfun <- function(model, data) {
  v <- predict(model, data, se.fit=TRUE)
  cbind(p=as.vector(v$fit), se=as.vector(v$se.fit))
}

# predfun returns two variables, so use index=1:2
r2 <- predict(logo, model, fun=predfun, index=1:2)

## Not run:
# You can use multiple cores to speed up the predict function
# by calling it via the clusterR function (you may need to install the snow package)
beginCluster()
r1c <- clusterR(logo, predict, args=list(model))
r2c <- clusterR(logo, predict, args=list(model=model, fun=predfun, index=1:2))

## End(Not run)

# principal components of a RasterBrick
# here using sampling to simulate an object too large
# to feed all its values to prcomp
sr <- sampleRandom(logo, 100)
pca <- prcomp(sr)

# note the use of the 'index' argument
x <- predict(logo, pca, index=1:3)
plot(x)

## Not run:
# partial least square regression
library(pls)
model <- plsrf(formula=pa~., data=v)

```

```

# this returns an array:
predict(model, v[1:5,])
# write a function to turn that into a matrix
pfun <- function(x, data) {
  y <- predict(x, data)
  d <- dim(y)
  dim(y) <- c(prod(d[1:2]), d[3])
  y
}

pp <- predict(logo, model, fun=pfun, index=1:3)

# Random Forest

library(randomForest)
rfmod <- randomForest(pa ~., data=v)

## note the additional argument "type='response'" that is
## passed to predict.randomForest
r3 <- predict(logo, rfmod, type='response', progress='window')

## get a RasterBrick with class membership probabilities
vv <- v
vv$pa <- as.factor(vv$pa)
rfmod2 <- randomForest(pa ~., data=vv)
r4 <- predict(logo, rfmod2, type='prob', index=1:2)
splot(r4)

# cforest (other Random Forest implementation) example with factors argument
v$red <- as.factor(round(v$red/100))
logo$red <- round(logo[[1]]/100)

library(party)
m <- cforest(pa~., control=cforest_unbiased(mtry=3), data=v)
f <- list(levels(v$red))
names(f) <- 'red'
# the second argument in party::predict.RandomForest
# is "OOB", and not "newdata" or similar. We need to write a wrapper
# predict function to deal with this
predfun <- function(m, d, ...) predict(m, newdata=d, ...)

pc <- predict(logo, m, OOB=TRUE, factors=f, fun=predfun)

# knn example, using calc instead of predict
library(class)
cl <- factor(c(rep(1, nrow(p)), rep(0, nrow(a))))
train <- extract(logo, rbind(p, a))
k <- calc(logo, function(x) as.integer(as.character(knn(train, x, cl))))

## End(Not run)

```

**Description**

These are low level functions that can be used by programmers to develop new functions. If in doubt, it is almost certain that you do not need these functions as these are already embedded in all other functions in the raster package.

`canProcessInMemory` is typically used within functions. In the raster package this function is used to determine if the amount of memory needed for the function is available. If there is not enough memory available, the function returns `FALSE`, and the function that called it will write the results to a temporary file.

`readStart` opens file connection(s) for reading, `readStop` removes it.

`pbCreate` creates a progress bar, `pbStep` sets the progress, and `pbClose` closes it.

**Usage**

```
canProcessInMemory(x, n=4, verbose=FALSE)
pbCreate(nsteps, progress, style=3, label='Progress', ...)
pbStep(pb, step=NULL, label='')
pbClose(pb, timer)
readStart(x, ...)
readStop(x)
getCluster()
returnCluster()
```

**Arguments**

<code>x</code>	RasterLayer or RasterBrick object (for connections) or RasterStack object ( <code>canProcessInMemory</code> )
<code>n</code>	integer. The number of copies of the Raster* object cell values that a function needs to be able to have in memory
<code>verbose</code>	logical. If <code>TRUE</code> the amount of memory needed and available is printed
<code>nsteps</code>	integer. Number of steps the progress bar will make from start to end (e.g. <code>nrow(raster)</code> )
<code>progress</code>	character. 'text', 'window', or ''
<code>style</code>	style for text progress bar. See <a href="#">txtProgressBar</a>
<code>label</code>	character. Label for the window type progress bar
<code>...</code>	additional arguments (None implemented, except for 'silent=TRUE' for <code>readStart</code> for files read with <code>gdal</code> , and other arguments passed to <code>gdal.open</code> )
<code>pb</code>	progress bar object created with <code>pbCreate</code>
<code>step</code>	which step is this ( $1 \leq \text{step} \leq \text{nsteps}$ ). If step is <code>NULL</code> , a single step is taken
<code>timer</code>	logical. If <code>TRUE</code> , time to completion will be printed. If missing, the value will be taken from the <code>rasterOptions</code>

**Value**

canProcessInMemory: logical  
 closeConnection: RasterLayer or RasterBrick object  
 getCluster: snow cluster object

**Examples**

```
r <- raster(nrow=100, ncol=100)
canProcessInMemory(r, 4)
r <- raster(nrow=50000, ncol=50000)
canProcessInMemory(r, 2, verbose=TRUE)
rasterOptions(maxmem=Inf, memfrac=.8)
rasterOptions(default=TRUE)
```

---

projection	<i>Get or set a coordinate reference system (projection)</i>
------------	--

---

**Description**

Get or set the coordinate reference system (CRS) of a Raster\* object.

**Usage**

```
## S4 method for signature 'ANY'
crs(x, asText=FALSE, ...)
## S4 method for signature 'Raster'
wkt(obj)
crs(x, ...) <- value

projection(x, asText=TRUE)
projection(x) <- value
```

**Arguments**

x	Raster* or Spatial object
obj	Raster*, Spatial, or CRS object
asText	logical. If TRUE, the projection is returned as text. Otherwise a <a href="#">CRS-class</a> object is returned
...	additional arguments. None implemented
value	CRS object or a character string describing a projection and datum in the PROJ.4 format

**Details**

projections are done by with the PROJ.4 library exposed by rgdal



**Value**

Raster\*, Spatial\*, or character object

**Note**

crs replaces earlier function projection. For compatibility with sp you can use proj4string instead of crs. wkt returns the "well-known-text" representation of the crs.

**See Also**

[projectRaster](#), [CRS-class](#), [spTransform](#), [projInfo](#)

**Examples**

```
r <- raster()
crs(r)
crs(r) <- "+proj=lcc +lat_1=48 +lat_2=33 +lon_0=-100 +datum=WGS84"
crs(r)
w <- wkt(r)
w
cat(w, "\n")
```

---

projectRaster

*Project a Raster object*

---

**Description**

Project the values of a Raster\* object to a new Raster\* object with another projection (coordinate reference system, (CRS)). You can do this by providing the new projection as a single argument in which case the function sets the extent and resolution of the new object. To have more control over the transformation, and, for example, to assure that the new object lines up with other datasets, you can provide a Raster\* object with the properties that the input data should be projected to.

projectExtent returns a RasterLayer with a projected extent, but without any values. This RasterLayer can then be adjusted (e.g. by setting its resolution) and used as a template 'to' in projectRaster.

**Usage**

```
projectRaster(from, to, res, crs, method="bilinear",
              alignOnly=FALSE, over=FALSE, filename="", ...)

projectExtent(object, crs)
```

**Arguments**

from	Raster* object
to	Raster* object with the parameters to which 'from' should be projected
res	single or (vector of) two numerics. To, optionally, set the output resolution if 'to' is missing
crs	character or object of class 'CRS'. PROJ.4 description of the coordinate reference system. In projectRaster this is used to set the output CRS if 'to' is missing, or if 'to' has no valid CRS
method	method used to compute values for the new RasterLayer. Either 'ngb' (nearest neighbor), which is useful for categorical variables, or 'bilinear' (bilinear interpolation; the default value), which is appropriate for continuous variables.
alignOnly	logical. Use to or other parameters only to align the output (i.e. same origin and resolution), but use the projected extent from from
over	logical. If TRUE wrapping around the date-line is turned off. This can be desirable for global data (to avoid mapping the same areas twice) but it is not desirable in other cases
filename	character. Output filename
...	additional arguments as for <a href="#">writeRaster</a>
object	Raster* object

**Details**

There are two approaches you can follow to project the values of a Raster object.

- 1) Provide a crs argument, and, optionally, a res argument, but do not provide a to argument.
- 2) Create a template Raster with the CRS you want to project to. You can use an existing object, or use projectExtent for this or an existing Raster\* object. Also set the number of rows and columns (or the resolution), and perhaps adjust the extent. The resolution of the output raster should normally be similar to that of the input raster. Then use that object as from argument to project the input Raster to. This is the preferred method because you have most control. For example you can assure that the resulting Raster object lines up with other Raster objects.

Projection is performed using the PROJ.4 library accessed through the rgdal package.

One of the best places to find PROJ.4 coordinate reference system descriptions is <http://www.spatialreference.org>.

You can also consult this page: [http://geotiff.maptools.org/proj\\_list](http://geotiff.maptools.org/proj_list) to find the parameter options and names for projections.

Also see projInfo('proj'), projInfo('ellps'), and projInfo('datum') for valid PROJ.4 values.

**Value**

RasterLayer or RasterBrick object.

**Note**

If the resolution of the output is much larger than that of the input, you should first aggregate the input such that the resolution of the input becomes more similar (perhaps a little smaller) to the output.

**Note**

Vector (points, lines, polygons) can be transformed with [spTransform](#).

`projectExtent` does not work very well when transforming projected circumpolar data to (e.g.) longitude/latitude. With such data you may need to adjust the returned object. E.g. `do ymax(object) <- 90`

**Author(s)**

Robert J. Hijmans and Joe Cheng

**See Also**

[resample](#), [CRS-class](#), [projInfo](#), [spTransform](#)

**Examples**

```
# create a new (not projected) RasterLayer with cellnumbers as values
r <- raster(xmn=-110, xmx=-90, ymn=40, ymx=60, ncols=40, nrows=40)
r <- setValues(r, 1:ncell(r))
crs(r)
# wkt(r)
# proj.4 projection description
newproj <- "+proj=lcc +lat_1=48 +lat_2=33 +lon_0=-100 +datum=WGS84"

# we need the rgdal package for this
if (require(rgdal)) {

#simplest approach
pr1 <- projectRaster(r, crs=newproj)

# alternatively also set the resolution
pr2 <- projectRaster(r, crs=newproj, res=20000)

# inverse projection, back to the properties of 'r'
inv <- projectRaster(pr2, r)

# to have more control, provide an existing Raster object, here we create one
# using projectExtent (no values are transferred)
pr3 <- projectExtent(r, newproj)
# Adjust the cell size
res(pr3) <- 20000
# now project
pr3 <- projectRaster(r, pr3)

## Not run:
```

```
# using a higher resolution
res(pr1) <- 10000
pr <- projectRaster(r, pr1, method='bilinear')
inv <- projectRaster(pr, r, method='bilinear')
dif <- r - inv
# small difference
plot(dif)

## End(Not run)

}
```

---

properties

*Raster file properties*

---

## Description

Properties of the values of the file that a RasterLayer object points to

dataSize returns the number of bytes used for each value (pixel, grid cell) dataSigned is TRUE for data types that include negative numbers.

## Usage

```
dataSize(object)
dataSigned(object)
```

## Arguments

object            Raster\* object

## Value

varies

## See Also

[filename](#)

## Examples

```
r <- raster(system.file("external/test.grd", package="raster"))
dataSize(r)
dataSigned(r)
dataType(r)
```

---

quantile	<i>Raster quantiles</i>
----------	-------------------------

---

**Description**

Compute quantiles for the cell values of a RasterLayer. If you want to compute quantiles for each cell across a number of layers, you can use `calc(x, fun=quantile)`.

**Usage**

```
quantile(x, ...)
```

**Arguments**

x	Raster object
...	Additional arguments: <code>na.rm=TRUE</code> , <code>ncells=NULL</code> , and additional arguments to the <code>stats::quantile</code> function, see <a href="#">quantile</a> <code>ncells</code> can be used to set the number of cells to be sampled, for very large raster datasets.

**Value**

A vector of quantiles

**See Also**

[density](#), [cellStats](#)

**Examples**

```
r <- raster(ncol=100, nrow=100)
values(r) <- rnorm(ncell(r), 0, 50)
quantile(r)
quantile(r, probs = c(0.25, 0.75), type=7, names = FALSE)
```

---

raster	<i>Create a RasterLayer object</i>
--------	------------------------------------

---

**Description**

Methods to create a RasterLayer object. RasterLayer objects can be created from scratch, a file, an Extent object, a matrix, an 'image' object, or from a Raster\*, Spatial\*, im (spatstat) asc, kasc (adehabitat\*), grf (geoR) or kde object.

In many cases, e.g. when a RasterLayer is created from a file, it does (initially) not contain any cell (pixel) values in (RAM) memory, it only has the parameters that describe the RasterLayer. You can access cell-values with [getValues](#), [extract](#) and related functions. You can assign new values with [setValues](#) and with [replacement](#).

For an overview of the functions in the raster package have a look here: [raster-package](#).

**Usage**

```
## S4 method for signature 'character'
raster(x, band=1, ...)

## S4 method for signature 'RasterLayer'
raster(x)

## S4 method for signature 'RasterStack'
raster(x, layer=0)

## S4 method for signature 'RasterBrick'
raster(x, layer=0)

## S4 method for signature 'missing'
raster(nrows=180, ncols=360, xmn=-180, xmx=180, ymn=-90, ymx=90,
crs, ext, resolution, vals=NULL)

## S4 method for signature 'Extent'
raster(x, nrows=10, ncols=10, crs="", ...)

## S4 method for signature 'matrix'
raster(x, xmn=0, xmx=1, ymn=0, ymx=1, crs="", template=NULL)

## S4 method for signature 'Spatial'
raster(x, origin, ...)

## S4 method for signature 'SpatialGrid'
raster(x, layer=1, values=TRUE)

## S4 method for signature 'SpatialPixels'
raster(x, layer=1, values=TRUE)

## S4 method for signature 'sf'
raster(x, origin, ...)
```

**Arguments**

x	filename (character), Extent, Raster*, sf, SpatialPixels*, SpatialGrid*, object, 'image', matrix, im, or missing. Supported file types are the 'native' raster package format and those that can be read via rgdal (see <a href="#">readGDAL</a> )
band	integer. The layer to use in a multi-layer file
...	Additional arguments, see Details
layer	integer. The layer (variable) to use in a multi-layer file, or the layer to extract from a RasterStack/Brick or SpatialPixelsDataFrame or SpatialGridDataFrame. An empty RasterLayer (no associated values) is returned if layer=0
values	logical. If TRUE, the cell values of 'x' are copied to the RasterLayer object that is returned

nrows	integer > 0. Number of rows
ncols	integer > 0. Number of columns
xmn	minimum x coordinate (left border)
xmx	maximum x coordinate (right border)
ymn	minimum y coordinate (bottom border)
ymx	maximum y coordinate (top border)
ext	object of class Extent. If present, the arguments xmn, xmx, ymn and ymx are ignored
crs	character or object of class CRS. PROJ.4 type description of a Coordinate Reference System (map projection). If this argument is missing, and the x coordinates are within -360 .. 360 and the y coordinates are within -90 .. 90, "+proj=longlat +datum=WGS84" is used. Also see under Details if x is a character (filename)
resolution	numeric vector of length 1 or 2 to set the resolution (see <a href="#">res</a> ). If this argument is used, arguments ncols and nrows are ignored
vals	optional. Values for the new RasterLayer. Accepted formats are as for <a href="#">setValues</a>
origin	minimum y coordinate (bottom border)
template	Raster* or Extent object used to set the extent (and CRS in case of a Raster* object). If not NULL, arguments xmn, xmx, ymn, ymx and crs (unless template is an Extent object) are ignored

### Details

If x is a filename, the following additional variables are recognized:

sub: positive integer. Subdataset number for a file with subdatasets

native: logical. Default is FALSE except when package rgdal is missing. If TRUE, reading and writing of IDRISI, BIL, BSQ, BIP, SAGA, and Arc ASCII files is done with native (raster package) drivers, rather than via rgdal. 'raster' and netcdf format files are always read with native drivers.

RAT: logical. The default is TRUE, in which case a raster attribute table is created for files that have one

offset: integer. To indicate the number of header rows on non-standard ascii files (rarely useful; use with caution)

crs: character. PROJ.4 string to set the CRS. Ignored when the file provides a CRS description that can be interpreted.

If x represents a **NetCDF** file, the following additional variable is recognized:

varname: character. The variable name, such as 'tasmax' or 'pr'. If not supplied and the file has multiple variables a guess will be made (and reported)

lvar: integer > 0 (default=3). To select the 'level variable' (3rd dimension variable) to use, if the file has 4 dimensions (e.g. depth instead of time)

level: integer > 0 (default=1). To select the 'level' (4th dimension variable) to use, if the file has 4 dimensions, e.g. to create a RasterBrick of weather over time at a certain height.

To use NetCDF files the `ncdf4` package needs to be available. It is assumed that these files follow, or are compatible with, the CF-1 convention (The GMT format may also work). If the `ncdf` file does not have a standard extension (which is used to recognize the file format), you can use argument `ncdf=TRUE` to indicate the format.

If `x` is a `Spatial` or an `Extent` object, additional arguments are for the method with signature `'missing'`

## Value

`RasterLayer`

## See Also

[stack](#), [brick](#)

## Examples

```
# Create a RasterLayer object from a file
# N.B.: For your own files, omit the 'system.file' and 'package="raster"' bits
# these are just to get the path to files installed with the package

f <- system.file("external/test.grd", package="raster")
f
r <- raster(f)

logo <- raster(system.file("external/rlogo.grd", package="raster"))

#from scratch
r1 <- raster(nrows=108, ncols=21, xmn=0, xmx=10)

#from an Extent object
e <- extent(r)
r2 <- raster(e)

#from another Raster* object
r3 <- raster(r)
s <- stack(r, r, r)
r4 <- raster(s)
r5 <- raster(s, 3)

## Not run:
# from NSIDC sea ice concentration file
baseurl <- "ftp://sidacs.colorado.edu/pub/DATASETS/"
# southern hemisphere
f1 <- paste(baseurl,
  "nsidc0051_gsfc_nasateam_seaice/final-gsfc/south/daily/2013/nt_20130114_f17_v1.1_s.bin",
  sep='')
# or northern hemisphere
f2 <- paste(baseurl,
  "nsidc0051_gsfc_nasateam_seaice/final-gsfc/north/daily/2013/nt_20130105_f17_v1.1_n.bin",
```



```

sep='')

if (!file.exists(basename(f1))) download.file(f1, basename(f1), mode = "wb")
ice1 <- raster(basename(f1))

if (!file.exists(basename(f2))) download.file(f2, basename(f2), mode = "wb")
ice2 <- raster(basename(f2))

## End(Not run)

```

---

Raster-class

*Raster\* classes*


---

## Description

A raster is a database organized as a rectangular grid that is sub-divided into rectangular cells of equal area (in terms of the units of the coordinate reference system). The 'raster' package defines a number of "S4 classes" to manipulate such data.

The main user level classes are RasterLayer, RasterStack and RasterBrick. They all inherit from BasicRaster and can contain values for the raster cells.

An object of the RasterLayer class refers to a single layer (variable) of raster data. The object can point to a file on disk that holds the values of the raster cells, or hold these values in memory. Or it can not have any associated values at all.

A RasterStack represents a collection of RasterLayer objects with the same extent and resolution. Organizing RasterLayer objects in a RasterStack can be practical when dealing with multiple layers; for example to summarize their values (see [calc](#)) or in spatial modeling (see [predict](#)).

An object of class RasterBrick can also contain multiple layers of raster data, but they are more tightly related. An object of class RasterBrick can refer to only a single (multi-layer) data file, whereas each layer in a RasterStack can refer to another file (or another band in a multi-band file). This has implications for processing speed and flexibility. A RasterBrick should process quicker than a RasterStack (irrespective if values are on disk or in memory). However, a RasterStack is more flexible as a single object can refer to layers that have values stored on disk as well as in memory. If a layer that does not refer to values on disk (they only exists in memory) is added to a RasterBrick, it needs to load all its values into memory (and this may not be possible because of memory size limitations).

Objects can be created from file or from each other with the following functions: [raster](#), [brick](#) and [stack](#).

Raster\* objects can also be created from SpatialPixels\* and SpatialGrid\* objects from the sp package using [as](#), or simply with the function [raster](#), [brick](#), or [stack](#). Vice versa, Raster\* objects can be coerced into a sp type object with [as](#)( , ), e.g. [as](#)(x, 'SpatialGridDataFrame') .

Common generic methods implemented for these classes include:

[summary](#), [show](#), [dim](#), and [plot](#), . . .

[ is implemented for RasterLayer.

The classes described above inherit from the BasicRaster class which inherits from BasicRaster. The BasicRaster class describes the main properties of a raster such as the number of columns and rows, and it contains an object of the link[raster]{Extent-class} to describe its spatial extent (coordinates). It also holds the 'coordinate reference system' in a slot of class CRS-class defined in the sp package. A BasicRaster cannot contain any raster cell values and is therefore seldomly used.

The Raster\* class inherits from BasicRaster. It is a virtual class; which means that you cannot create an object of this class. It is used only to define methods for all the classes that inherit from it (RasterLayer, RasterStack and RasterBrick). Another virtual class is the RasterStackBrick class. It is formed by a class union of RasterStack and RasterBrick. You cannot make objects of it, but methods defined for objects of this class as arguments will accept objects of the RasterLayer and RasterStack as that argument.

Classes RasterLayer and RasterBrick have a slot with an object of class RasterFile that describes the properties of the file they point to (if they do). RasterLayer has a slot with an object of class SingleLayerData, and the RasterBrick class has a slot with an object of class MultipleLayerData. These 'datalayer' classes can contain (some of) the values of the raster cells.

These classes are not further described here because users should not need to directly access these slots. The 'setter' functions such as setValues should be used instead. Using such 'setter' functions is much safer because a change in one slot should often affect the values in other slots.

### Objects from the Class

Objects can be created by calls of the form `new("RasterLayer", ...)`, or with the helper functions such as `raster`.

### Slots

Slots for RasterLayer and RasterBrick objects

`title`: Character

`file`: Object of class ".RasterFile"

`data`: Object of class ".SingleLayerData" or ".MultipleLayerData"

`history`: To record processing history, not yet in use

`legend`: Object of class .RasterLegend, Default legend. Should store preferences for plotting.  
Not yet implemented except that it stores the color table of images, if available

`extent`: Object of [Extent-class](#)

`ncols`: Integer

`nrows`: Integer

`crs`: Object of class "CRS", i.e. the coordinate reference system. In Spatial\* objects this slot is called 'proj4string'

### Examples

```
showClass("RasterLayer")
```

---

rasterFromCells      *Subset a raster by cell numbers*

---

### Description

This function returns a new raster based on an existing raster and cell numbers for that raster. The new raster is cropped to the cell numbers provided, and, if `values=TRUE` has values that are the cell numbers of the original raster.

### Usage

```
rasterFromCells(x, cells, values=TRUE)
```

### Arguments

x	Raster* object (or a SpatialPixels* or SpatialGrid* object)
cells	vector of cell numbers
values	Logical. If TRUE, the new RasterLayer has cell values that correspond to the cell numbers of x

### Details

Cell numbers start at 1 in the upper left corner, and increase from left to right, and then from top to bottom. The last cell number equals the number of cells of the Raster\* object.

### Value

RasterLayer

### See Also

[rowFromCell](#)

### Examples

```
r <- raster(ncols=100, nrows=100)
cells <- c(3:5, 210)
r <- rasterFromCells(r, cells)
cbind(1:ncell(r), getValues(r))
```

---

rasterFromXYZ                      *Create a Raster\* object from x, y, z values*

---

### Description

Create a Raster\* object from x, y and z values. x and y represent spatial coordinates and must be on a regular grid. If the resolution is not supplied, it is assumed to be the minimum distance between x and y coordinates, but a resolution of up to 10 times smaller is evaluated if a regular grid can otherwise not be created. z values can be single or multiple columns (variables) If the exact properties of the RasterLayer are known beforehand, it may be preferable to simply create a new RasterLayer with the raster function instead, compute cell numbers and assign the values with these (see example below).

### Usage

```
rasterFromXYZ(xyz, res=c(NA,NA), crs="", digits=5)
```

### Arguments

xyz	matrix or data.frame with at least three columns: x and y coordinates, and values (z). There may be several 'z' variables (columns)
res	numeric. The x and y cell resolution (optional)
crs	CRS object or a character string describing a projection and datum in PROJ.4 format
digits	numeric, indicating the requested precision for detecting whether points are on a regular grid (a low number of digits is a low precision)

### Value

RasterLayer or RasterBrick

### See Also

See [rasterize](#) for points that are not on a regular grid

### Examples

```
r <- raster(nrow=5, ncol=5, xmn=0, xmx=10, ymn=0, ymx=10, crs="")
set.seed(1)
values(r) <- sample(1:25)
r[r < 15] <- NA
xyz <- rasterToPoints(r)

rst <- rasterFromXYZ(xyz)

# equivalent to:
rr <- raster(nrow=5, ncol=5, xmn=0, xmx=10, ymn=0, ymx=10)
```

```

cells <- cellFromXY(rr, xyz[,1:2])
rr[cells] <- xyz[,3]

# multiple layers
xyzz <- cbind(xyz, a=1:nrow(xyz), b=nrow(xyz):1)
b <- rasterFromXYZ(xyzz)

```

---

rasterize

*Rasterize points, lines, or polygons*


---

### Description

Transfer values associated with 'object' type spatial data (points, lines, polygons) to raster cells.

For polygons, values are transferred if the polygon covers the center of a raster cell. For lines, values are transferred to all cells that are touched by a line. You can combine this behaviour by rasterizing polygons as lines first and then as polygons.

If x represents points, each point is assigned to a grid cell. Points that fall on a border between cells are placed in the cell to the right and/or in the cell below. The value of a grid cell is determined by the values associated with the points and function fun.

### Usage

```

## S4 method for signature 'matrix,Raster'
rasterize(x, y, field, fun='last', background=NA,
          mask=FALSE, update=FALSE, updateValue='all', filename="", na.rm=TRUE, ...)

## S4 method for signature 'SpatialPoints,Raster'
rasterize(x, y, field, fun='last', background=NA,
          mask=FALSE, update=FALSE, updateValue='all', filename="", na.rm=TRUE, ...)

## S4 method for signature 'SpatialLines,Raster'
rasterize(x, y, field, fun='last', background=NA,
          mask=FALSE, update=FALSE, updateValue='all', filename="", ...)

## S4 method for signature 'SpatialPolygons,Raster'
rasterize(x, y, field, fun='last', background=NA,
          mask=FALSE, update=FALSE, updateValue='all', filename="",
          getCover=FALSE, silent=TRUE, ...)

```

### Arguments

x	points (a <i>SpatialPoints*</i> object, or a two-column matrix (or data.frame)), <i>SpatialLines*</i> , <i>SpatialPolygons*</i> , or an Extent object
y	<i>Raster*</i> object

field	numeric or character. The value(s) to be transferred. This can be a single number, or a vector of numbers that has the same length as the number of spatial features (points, lines, polygons). If <code>x</code> is a <code>Spatial*DataFrame</code> , this can be the column name of the variable to be transferred. If missing, the attribute index is used (i.e. numbers from 1 to the number of features). You can also provide a vector with the same length as the number of spatial features, or a matrix where the number of rows matches the number of spatial features
fun	function or character. To determine what values to assign to cells that are covered by multiple spatial features. You can use functions such as <code>min</code> , <code>max</code> , or <code>mean</code> , or one of the following character values: <code>'first'</code> , <code>'last'</code> , <code>'count'</code> . The default value is <code>'last'</code> . In the case of <code>SpatialLines*</code> , <code>'length'</code> is also allowed (currently for planar coordinate systems only). If <code>x</code> represents points, <code>fun</code> must accept a <code>na.rm</code> argument, either explicitly or through <code>'dots'</code> . This means that <code>fun=length</code> fails, but <code>fun=function(x,...)length(x)</code> works, although it ignores the <code>na.rm</code> argument. To use the <code>na.rm</code> argument you can use a function like this: <code>fun=function(x,na.rm)if (na.rm) length(na.omit(x)) else length(x)</code> , or use a function that removes NA values in all cases, like this function to compute the number of unique values per grid cell "richness": <code>fun=function(x,...){length(unique(na.omit(x)))}</code> . If you want to count the number of points in each grid cell, you can use <code>fun='count'</code> or <code>fun=function(x,...){length(x)}</code> . You can also pass multiple functions using a statement like <code>fun=function(x,...)c(length(x),mean(x))</code> , in which case the returned object is a <code>RasterBrick</code> (multiple layers).
background	numeric. Value to put in the cells that are not covered by any of the features of <code>x</code> . Default is NA
mask	logical. If TRUE the values of the input Raster object are 'masked' by the spatial features of <code>x</code> . That is, cells that spatially overlap with the spatial features retain their values, the other cells become NA. Default is FALSE. This option cannot be used when <code>update=TRUE</code>
update	logical. If TRUE, the values of the Raster* object are updated for the cells that overlap the spatial features of <code>x</code> . Default is FALSE. Cannot be used when <code>mask=TRUE</code>
updateValue	numeric (normally an integer), or character. Only relevant when <code>update=TRUE</code> . Select, by their values, the cells to be updated with the values of the spatial features. Valid character values are <code>'all'</code> , <code>'NA'</code> , and <code>'!NA'</code> . Default is <code>'all'</code>
filename	character. Output filename (optional)
na.rm	If TRUE, NA values are removed if <code>fun</code> honors the <code>na.rm</code> argument
getCover	logical. If TRUE, the fraction of each grid cell that is covered by the polygons is returned (and the values of <code>field</code> , <code>fun</code> , <code>mask</code> , and <code>update</code> are ignored). The fraction covered is estimated by dividing each cell into 100 subcells and determining presence/absence of the polygon in the center of each subcell
silent	Logical. If TRUE, feedback on the polygon count is suppressed. Default is FALSE
...	Additional arguments for file writing as for <code>writeRaster</code>

**Value**

RasterLayer or RasterBrick

**See Also**[extract](#)**Examples**

```
#####
# rasterize points
#####
r <- raster(ncols=36, nrows=18)
n <- 1000
set.seed(123)
x <- runif(n) * 360 - 180
y <- runif(n) * 180 - 90
xy <- cbind(x, y)
# get the (last) indices
r0 <- rasterize(xy, r)
# presence/absence (NA) (is there a point or not?)
r1 <- rasterize(xy, r, field=1)
# how many points?
r2 <- rasterize(xy, r, fun=function(x,...)length(x))
vals <- runif(n)
# sum of the values associated with the points
r3 <- rasterize(xy, r, vals, fun=sum)

# with a SpatialPointsDataFrame
vals <- 1:n
p <- data.frame(xy, name=vals)
coordinates(p) <- ~x+y
r <- rasterize(p, r, 'name', fun=min)
#r2 <- rasterize(p, r, 'name', fun=max)
#plot(r, r2, cex=0.5)

#####
# rasterize lines
#####
cds1 <- rbind(c(-180,-20), c(-140,55), c(10, 0), c(-140,-60))
cds2 <- rbind(c(-10,0), c(140,60), c(160,0), c(140,-55))
cds3 <- rbind(c(-125,0), c(0,60), c(40,5), c(15,-45))

lines <- spLines(cds1, cds2, cds3)

r <- raster(ncols=90, nrows=45)
r <- rasterize(lines, r)

## Not run:
plot(r)
plot(lines, add=TRUE)

r <- rasterize(lines, r, fun='count')
plot(r)
```

```

values(r) <- 1:ncell(r)
r <- rasterize(lines, r, mask=TRUE)
plot(r)

values(r) <- 1
r[lines] <- 10
plot(r)

## End(Not run)

#####
# rasterize polygons
#####

p1 <- rbind(c(-180,-20), c(-140,55), c(10, 0), c(-140,-60), c(-180,-20))
hole <- rbind(c(-150,-20), c(-100,-10), c(-110,20), c(-150,-20))
p1 <- list(p1, hole)
p2 <- rbind(c(-10,0), c(140,60), c(160,0), c(140,-55), c(-10,0))
p3 <- rbind(c(-125,0), c(0,60), c(40,5), c(15,-45), c(-125,0))

pols <- spPolygons(p1, p2, p3)

r <- raster(ncol=90, nrow=45)
r <- rasterize(pols, r, fun=sum)

## Not run:

plot(r)
plot(pols, add=T)

# add a polygon
p5 <- rbind(c(-180,10), c(0,90), c(40,90), c(145,-10),
            c(-25, -15), c(-180,0), c(-180,10))
addpoly <- SpatialPolygons(list(Polygons(list(Polygon(p5)), 1)))
addpoly <- as(addpoly, "SpatialPolygonsDataFrame")
addpoly@data[1,1] <- 10
r2 <- rasterize(addpoly, r, field=1, update=TRUE, updateValue="NA")
plot(r2)
plot(pols, border="blue", lwd=2, add=TRUE)
plot(addpoly, add=TRUE, border="red", lwd=2)

# get the percentage cover of polygons in a cell
r3 <- raster(ncol=36, nrow=18)
r3 <- rasterize(pols, r3, getCover=TRUE)

## End(Not run)

```



## Description

Functions in the raster package create temporary files if the values of an output Raster\* object cannot be stored in memory (RAM). This can happen when no filename is provided to a function and in functions where you cannot provide a filename (e.g. when using 'raster algebra').

Temporary files are automatically removed at the start of each session. During a session you can use showTmpFiles to see what is there and removeTmpFiles to delete all the temporary files. rasterTmpFile returns a temporary filename. These can be useful when developing your own functions. These filenames consist of prefix\_date\_time\_pid\_rn where pid is the process id returned by `Sys.getpid` and rn is a 5 digit random number. This should make tempfiles unique if created at different times and also when created in parallel processes (different pid) that use `set.seed` and call rasterTmpFile at the same time. It is possible, however, to create overlapping names (see the examples), which is undesirable and can be avoided by setting the prefix argument.

## Usage

```
rasterTmpFile(prefix='r_tmp_')
showTmpFiles()
removeTmpFiles(h=24)
```

## Arguments

prefix	Character. Prefix to the filename (which will be followed by 10 random numbers)
h	Numeric. The minimum age of the files in number of hours (younger files are not deleted)

## Value

rasterTmpFile returns a valid file name

showTmpFiles returns the names (.grd only) of the files in the temp directory

removeTmpFiles returns nothing

## See Also

[rasterOptions](#), [tempfile](#)

## Examples

```
## Not run:
rasterTmpFile('mytemp_')
showTmpFiles()
removeTmpFiles(h=24)

## End(Not run)
```

---

rasterToContour	<i>Raster to contour lines conversion</i>
-----------------	---

---

**Description**

RasterLayer to contour lines. This is a wrapper around [contourLines](#)

**Usage**

```
rasterToContour(x, maxpixels=100000, ...)
```

**Arguments**

x	a RasterLayer object
maxpixels	Maximum number of raster cells to use; this function fails when too many cells are used
...	Any argument that can be passed to <a href="#">contourLines</a>

**Details**

Most of the code was taken from `maptools::ContourLines2SLDF`, by Roger Bivand & Edzer Pebesma

**Value**

SpatialLinesDataFrame

**Examples**

```
f <- system.file("external/test.grd", package="raster")
r <- raster(f)
x <- rasterToContour(r)
class(x)
plot(r)
plot(x, add=TRUE)
```

---

rasterToPoints	<i>Raster to points conversion</i>
----------------	------------------------------------

---

**Description**

Raster to point conversion. Cells with NA are not converted. A function can be used to select a subset of the raster cells (by their values).

**Usage**

```
rasterToPoints(x, fun=NULL, spatial=FALSE, ...)
```

**Arguments**

x	A Raster* object
fun	Function to select a subset of raster values
spatial	Logical. If TRUE, the function returns a SpatialPointsDataFrame object
...	Additional arguments. Currently only progress to specify a progress bar. "text", "window", or "" (the default, no progress bar)

**Details**

fun should be a simple function returning a logical value.

E.g.: fun=function(x){x==1} or fun=function(x){x>3}

**Value**

A matrix with three columns: x, y, and v (value), or a SpatialPointsDataFrame object

**Examples**

```
r <- raster(nrow=18, ncol=36)
values(r) <- runif(ncell(r)) * 10
r[r>8] <- NA
p <- rasterToPoints(r)
p <- rasterToPoints(r, fun=function(x){x>6})
#plot(r)
#points(p)
```

---

rasterToPolygons	<i>Raster to polygons conversion</i>
------------------	--------------------------------------

---

**Description**

Raster to polygons conversion. Cells with NA are not converted. A function can be used to select a subset of the raster cells (by their values).

**Usage**

```
rasterToPolygons(x, fun=NULL, n=4, na.rm=TRUE, digits=12, dissolve=FALSE)
```

**Arguments**

x	Raster* object
fun	function to select a subset of raster values (only allowed if x has a single layer)
n	integer. The number of nodes for each polygon. Only 4, 8, and 16 are allowed
na.rm	If TRUE, cells with NA values in all layers are ignored
digits	number of digits to round the coordinates to
dissolve	logical. If TRUE, polygons with the same attribute value will be dissolved into multi-polygon regions. This option requires the rgeos package

**Details**

fun should be a simple function returning a logical value.

E.g.: fun=function(x){x==1} or fun=function(x){x>3 & x<6}

**Value**

SpatialPolygonsDataFrame

**Examples**

```
r <- raster(nrow=18, ncol=36)
values(r) <- runif(ncell(r)) * 10
r[r>8] <- NA
pol <- rasterToPolygons(r, fun=function(x){x>6})

#plot(r > 6)
#plot(pol, add=TRUE, col='red')
```

---

Rcpp-class

*Rcpp classes*

---

**Description**

These classes are for internal use only

---

readAll

*Read values from disk*

---

**Description**

Read all values from a raster file associated with a Raster\* object into memory. This function should normally not be used. In most cases [getValues](#) or [getValuesBlock](#) is more appropriate as readAll will fail when there is no file associated with the RasterLayer (values may only exist in memory).

**Usage**

```
readAll(object)
```

**Arguments**

object            a Raster\* object

**See Also**

[getValues](#), [getValuesBlock](#), [extract](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
r <- readAll(r)
```

---

reclassify

*Reclassify*


---

**Description**

Reclassify values of a Raster\* object. The function (re)classifies groups of values to other values. For example, all values between 1 and 10 become 1, and all values between 11 and 15 become 2 (see functions [subs](#) and [cut](#) for alternative approaches).

Reclassification is done with matrix `rcl`, in the row order of the reclassify table. Thus, if there are overlapping ranges, the first time a number is within a range determines the reclassification value.

**Usage**

```
## S4 method for signature 'Raster'
reclassify(x, rcl, filename='', include.lowest=FALSE, right=TRUE, ...)
```

**Arguments**

<code>x</code>	Raster* object
<code>rcl</code>	matrix for reclassification. This matrix can have 3 or 2 columns. In a 3-column matrix the first two columns are "from" - "to" for the input values, and the third column "becomes" has the new value for that range. (You can also supply a vector that can be coerced into a n*3 matrix (with <code>byrow=TRUE</code> )). A 2-column matrix represents ("is", "becomes") which can be useful for integer values. In that case, the <code>right</code> argument is automatically set to NA
<code>filename</code>	character. Output filename (optional)
<code>include.lowest</code>	logical, indicating if a value equal to the lowest value in <code>rcl</code> (or highest value in the second column, for <code>right = FALSE</code> ) should be included. The default is FALSE
<code>right</code>	logical, indicating if the intervals should be closed on the right (and open on the left) or vice versa. The default is TRUE. A special case is to use <code>right=NA</code> . In this case both the left and right intervals are open
<code>...</code>	additional arguments as for <a href="#">writeRaster</a>

**Value**

Raster\* object

**See Also**

[subs](#), [clamp](#), [cut](#), [calc](#)

**Examples**

```
r <- raster(ncols=36, nrows=18)
values(r) <- runif(ncell(r))
# reclassify the values into three groups
# all values > 0 and <= 0.25 become 1, etc.
m <- c(0, 0.25, 1, 0.25, 0.5, 2, 0.5, 1, 3)
rclmat <- matrix(m, ncol=3, byrow=TRUE)
rc <- reclassify(r, rclmat)

# for values >= 0 (instead of > 0), do
rc <- reclassify(r, rclmat, include.lowest=TRUE)

# equivalent to
rc <- reclassify(r, c(-Inf,0.25,1, 0.25,0.5,2, 0.5,Inf,3))
```

---

rectify

*rectify a Raster object*

---

**Description**

rectify changes a rotated Raster\* object into a non-rotated (rectangular) object. This is wrapper function around [resample](#).

**Usage**

```
## S4 method for signature 'Raster'
rectify(x, ext, res, method='ngb', filename='', ...)
```

**Arguments**

x	Raster* object to be rectified
ext	Optional. Extent object or object from which an Extent object can be extracted
res	Optional. Single or two numbers to set the resolution
method	Method used to compute values for the new RasterLayer, should be "bilinear" for bilinear interpolation, or "ngb" for nearest neighbor
filename	Character. Output filename
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer or RasterBrick object

---

replacement	<i>Replace cell values or layers of a Raster* object</i>
-------------	--

---

### Description

You can set values of a Raster\* object, when i is a vector of cell numbers, a Raster\*, Extent, or Spatial\* object.

These are shorthand methods that work best for relatively small Raster\* objects. In other cases you can use functions such as [calc](#) and [rasterize](#).

### Methods

```
x[i] <-value
x[i, j] <-value
```

#### Arguments:

x	a Raster* object
i	cell number(s), row number(s), Extent, Spatial* object
j	columns number(s) (only available if i is (are) a row number(s))
value	new cell value(s)

### See Also

[calc](#), [rasterize](#)

### Examples

```
r <- raster(ncol=10, nrow=5)
values(r) <- 1:ncell(r) * 2
r[1,] <- 1
r[,1] <- 2
r[1,1] <- 3

s <- stack(r, sqrt(r))
s[s<5] <- NA
```

---

resample	<i>Resample a Raster object</i>
----------	---------------------------------

---

### Description

Resample transfers values between non matching Raster\* objects (in terms of origin and resolution). Use [projectRaster](#) if the target has a different coordinate reference system (projection).

Before using resample, you may want to consider using these other functions instead: [aggregate](#), [disaggregate](#), [crop](#), [extend](#), [merge](#).

**Usage**

```
## S4 method for signature 'Raster,Raster'
resample(x, y, method="bilinear", filename="", ...)
```

**Arguments**

x	Raster* object to be resampled
y	Raster* object with parameters that x should be resampled to
method	method used to compute values for the new RasterLayer, should be "bilinear" for bilinear interpolation, or "ngb" for using the nearest neighbor
filename	character. Output filename (optional)
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer or RasterBrick object

**Author(s)**

Robert J. Hijmans and Joe Cheng

**See Also**

[aggregate](#), [disaggregate](#), [crop](#), [extend](#), [merge](#), [projectRaster](#)

**Examples**

```
r <- raster(nrow=3, ncol=3)
values(r) <- 1:ncell(r)
s <- raster(nrow=10, ncol=10)
s <- resample(r, s, method='bilinear')
#par(mfrow=c(1,2))
#plot(r)
#plot(s)
```

---

resolution

*Resolution*

---

**Description**

Get (or set) the x and/or y resolution of a Raster\* object

**Usage**

```
xres(x)
yres(x)
res(x)
res(x) <- value
```



**Arguments**

x	Raster* object
value	Resolution (single number or vector of two numbers)

**Value**

A single numeric value or two numeric values.

**See Also**

[extent, ncell](#)

**Examples**

```
r <- raster(ncol=18, nrow=18)
xres(r)
yres(r)
res(r)

res(r) <- 1/120
# set yres differently
res(r) <- c(1/120, 1/60)
```

---

 RGB

---

*Create a Red-Green-Blue Raster object*


---

**Description**

Make a Red-Green-Blue object that can be used to create images.

**Usage**

```
## S4 method for signature 'RasterLayer'
RGB(x, filename='', col=rainbow(25), breaks=NULL, alpha=FALSE,
colNA='white', zlim=NULL, zlimcol=NULL, ext=NULL, ...)
```

**Arguments**

x	RasterLayer
filename	character. Output filename (optional)
col	A color palette, that is a vector of n contiguous colors generated by functions like <a href="#">rainbow</a> , <a href="#">heat.colors</a> , <a href="#">topo.colors</a> , <a href="#">bpy.colors</a> or one or your own making, perhaps using <a href="#">colorRampPalette</a> . If none is provided, <code>rev(terrain.colors(255))</code> is used unless x has a 'color table'

breaks	numeric. A set of finite numeric breakpoints for the colours: must have one more breakpoint than colour and be in increasing order
alpha	If TRUE a fourth layer to set the background transparency is added
colNA	color for the background (NA values)
zlim	vector of length 2. Range of values to plot
zlimcol	If NULL the values outside the range of zlim get the color of the extremes of the range. If zlimcol has any other value, the values outside the zlim range get the color of NA values (see colNA)
ext	An <a href="#">Extent</a> object to zoom in to a region of interest (see <a href="#">drawExtent</a> )
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterBrick

**See Also**

[plotRGB](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
x <- RGB(r)
plot(x, col=gray(0:9/10))
plotRGB(x)
```

---

rotate

*Rotate*

---

**Description**

Rotate a Raster\* object that has x coordinates (longitude) from 0 to 360, to standard coordinates between -180 and 180 degrees. Longitude between 0 and 360 is frequently used in global climate models.

**Usage**

```
## S4 method for signature 'Raster'
rotate(x, filename='', ...)
```

**Arguments**

x	Raster* object
filename	character. Output filename (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer or a RasterBrick object

**See Also**

[flip](#)

**Examples**

```
r <- raster(nrow=18, ncol=36)
m <- matrix(1:ncell(r), nrow=18)
values(r) <- as.vector(t(m))
extent(r) <- extent(0, 360, -90, 90)
rr <- rotate(r)
```

---

rotated

*Do the raster cells have a rotation?*

---

**Description**

Do the raster cells have a rotation?

**Usage**

```
rotated(x)
```

**Arguments**

x                    A Raster\* object

**Value**

Logical value

**See Also**

[rectify](#)

**Examples**

```
r <- raster()
rotated(r)
```

---

round	<i>Integer values</i>
-------	-----------------------

---

### Description

These functions take a single RasterLayer argument *x* and change its values to integers.

`ceiling` returns a RasterLayer with the smallest integers not less than the corresponding values of *x*.

`floor` returns a RasterLayer with the largest integers not greater than the corresponding values of *x*.

`trunc` returns a RasterLayer with the integers formed by truncating the values in *x* toward 0.

`round` returns a RasterLayer with values rounded to the specified number of digits (decimal places; default 0).

### Details

see `?base::round`

### Value

a RasterLayer object

### Methods

`ceiling(x)` `floor(x)` `trunc(x, ...)` `round(x, digits = 0)`

a RasterLayer object

**digits** integer indicating the precision to be used

**...** additional arguments

### Examples

```
r <- raster(ncol=10, nrow=10)
values(r) <- runif(ncell(r)) * 10
s <- round(r)
```

---

rowFromCell	<i>Row or column number from a cell number</i>
-------------	--

---

### Description

These functions get the row and/or column number from a cell number of a Raster\* object)

### Usage

```
colFromCell(object, cell)
rowFromCell(object, cell)
rowColFromCell(object, cell)
```

### Arguments

object	Raster* object (or a SpatialPixels* or SpatialGrid* object)
cell	cell number(s)

### Details

The colFromCell and similar functions accept a single value, or a vector or list of these values, Cell numbers start at 1 in the upper left corner, and increase from left to right, and then from top to bottom. The last cell number equals the number of cells of the Raster\* object.

### Value

row of column number(s)

### See Also

[cellFrom](#)

### Examples

```
r <- raster(ncols=10, nrows=10)
colFromCell(r, c(5,15))
rowFromCell(r, c(5,15))
rowColFromCell(r, c(5,15))
```

---

rowSums	<i>rowSums and colSums for Raster objects</i>
---------	---

---

### Description

Sum values of Raster objects by row or column.

### Usage

```
## S4 method for signature 'Raster'  
rowSums(x, na.rm=FALSE, dims=1L,...)  
## S4 method for signature 'Raster'  
colSums(x, na.rm=FALSE, dims=1L,...)
```

### Arguments

x	Raster* object
na.rm	logical. If TRUE, NA values are ignored
dims	this argument is ignored
...	additional arguments (none implemented)

### Value

vector (if x is a RasterLayer) or matrix

### See Also

See [cellStats](#) for summing all cells values

### Examples

```
r <- raster(ncols=2, nrows=5)  
values(r) <- 1:10  
as.matrix(r)  
rowSums(r)  
colSums(r)
```

---

SampleInt	<i>Sample integer values</i>
-----------	------------------------------

---

**Description**

Take a random sample from a range of integer values between 1 and n. Its purpose is similar to that of `sample`, but that function fails when n is very large.

**Usage**

```
sampleInt(n, size, replace=FALSE)
```

**Arguments**

n	Positive number (integer); the number of items to choose from
size	Non-negative integer; the number of items to choose
replace	Logical. Should sampling be with replacement?

**Value**

vector of integer numbers

**Examples**

```
sampleInt(1e+12, 10)

# this may fail:
# sample.int(1e+12, 10)
# sample.int(1e+9, 10)
```

---

sampleRandom	<i>Random sample</i>
--------------	----------------------

---

**Description**

Take a random sample from the cell values of a Raster\* object (without replacement).

**Usage**

```
## S4 method for signature 'Raster'
sampleRandom(x, size, na.rm=TRUE, ext=NULL,
             cells=FALSE, rowcol=FALSE, xy=FALSE, sp=FALSE, asRaster=FALSE, ...)
```

**Arguments**

x	Raster* object
size	positive integer giving the number of items to choose
na.rm	logical. If TRUE (the default), NA values are removed from random sample
ext	Extent object. To limit regular sampling to the area within the extent
cells	logical. If TRUE, sampled cell numbers are also returned
rowcol	logical. If TRUE, sampled row and column numbers are also returned
xy	logical. If TRUE, coordinates of sampled cells are also returned
sp	logical. If TRUE, a SpatialPointsDataFrame is returned
asRaster	logical. If TRUE, a Raster* object is returned with random cells with values, all other cells with NA
...	Additional arguments as in <a href="#">writeRaster</a> . Only relevant when asRaster=TRUE

**Details**

With argument `na.rm=TRUE`, the returned sample may be smaller than requested

**Value**

A vector, matrix (if `cells=TRUE` or `x` is a multi-layered object), or a `SpatialPointsDataFrame` (if `sp=TRUE`)

**See Also**

[sampleRegular](#), [sampleStratified](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
sampleRandom(r, size=10)
s <- stack(r, r)
sampleRandom(s, size=5, cells=TRUE, sp=TRUE)
```

---

sampleRegular

*Regular sample*

---

**Description**

Take a systematic sample from a Raster\* object.

**Usage**

```
## S4 method for signature 'Raster'
sampleRegular(x, size, ext=NULL, cells=FALSE, xy=FALSE, asRaster=FALSE,
              sp=FALSE, useGDAL=FALSE, ...)
```



**Arguments**

x	Raster object
size	positive integer giving the number of items to choose.
ext	Extent. To limit regular sampling to the area within that box
cells	logical. Also return sampled cell numbers (if asRaster=FALSE)
xy	logical. If TRUE, coordinates of sampled cells are also returned
asRaster	logical. If TRUE, a RasterLayer or RasterBrick is returned, rather than the sampled values
sp	logical. If TRUE, a SpatialPointsDataFrame is returned
useGDAL	logical. If TRUE, GDAL is used to sample in some cases. This is quicker, but can result in values for a different set of cells than when useGDAL=FALSE. Only for rasters that are accessed via rgdal, and are not rotated. When TRUE arguments cells, xy, and sp are ignored (i.e., FALSE)
...	additional arguments. None implemented

**Value**

A vector (single layer object), matrix (multi-layered object; or if cells=TRUE, or xy=TRUE), Raster\* object (if asRaster=TRUE), or SpatialPointsDataFrame (if sp=TRUE)

**See Also**

[sampleRandom](#), [sampleStratified](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
v <- sampleRegular(r, size=100)
x <- sampleRegular(r, size=100, asRaster=TRUE)
```

---

sampleStratified	<i>Stratified random sample</i>
------------------	---------------------------------

---

**Description**

Take a stratified random sample from the cell values of a Raster\* object (without replacement). An attempt is made to sample size cells from each stratum. The values in the RasterLayer x are rounded to integers; with each value representing a stratum.

**Usage**

```
## S4 method for signature 'RasterLayer'
sampleStratified(x, size, exp=10, na.rm=TRUE, xy=FALSE, ext=NULL, sp=FALSE, ...)
```

**Arguments**

x	Raster* object, with values (rounded to integers) representing strata
size	positive integer giving the number of items to choose
exp	numeric >= 1. 'Expansion factor' that is multiplied with size to get an initial sample. Can be increased when you get an insufficient number of samples for small strata
na.rm	logical. If TRUE (the default), NA values are removed from random sample
xy	logical. Return coordinates of cells rather than cell numbers
ext	Extent object. To limit regular sampling to the area within the extent
sp	logical. If TRUE, a SpatialPointsDataFrame is returned
...	Additional arguments. None implemented

**Details**

The function may not work well when the size (number of cells) of some strata is relatively small.

**Value**

matrix of cell numbers (and optionally coordinates) by stratum

**See Also**

[sampleRandom](#), [sampleRegular](#)

**Examples**

```
r <- raster(ncol=10, nrow=10)
names(r) <- 'stratum'
values(r) <- round((runif(ncell(r))+0.5)*3)
sampleStratified(r, size=3)
```

---

scale

*Scale values*

---

**Description**

Center and/or scale raster data

**Usage**

```
## S4 method for signature 'Raster'
scale(x, center=TRUE, scale=TRUE)
```

**Arguments**

x	Raster* object
center	logical or numeric. If TRUE, centering is done by subtracting the layer means (omitting NAs), and if FALSE, no centering is done. If center is a numeric vector with length equal to the nlayers(x), then each layer of x has the corresponding value from center subtracted from it.
scale	logical or numeric. If TRUE, scaling is done by dividing the (centered) layers of x by their standard deviations if center is TRUE, and the root mean square otherwise. If scale is FALSE, no scaling is done. If scale is a numeric vector with length equal to nlayers(x), each layer of x is divided by the corresponding value. Scaling is done after centering.

**Value**

Raster\* object

**See Also**

[scale](#)

**Examples**

```
b <- brick(system.file("external/rlogo.grd", package="raster"))
bs <- scale(b)
```

---

scalebar

*scalebar*

---

**Description**

Add a scalebar to a plot

**Usage**

```
scalebar(d, xy = NULL, type = "line", divs = 2, below = "",
         lonlat = NULL, label, adj=c(0.5, -0.5), lwd = 2, ...)
```

**Arguments**

d	distance covered by scalebar
xy	x and y coordinate to place the plot. Can be NULL. Use xy=click() to make this interactive
type	"line" or "bar"
divs	Number of divisions for a bar type. 2 or 4
below	Text to go below scalebar (e.g., "kilometers")

lonlat	Logical or NULL. If logical, TRUE indicates if the plot is using longitude/latitude coordinates. If NULL this is guessed from the plot's coordinates
adj	adjustment for text placement
label	Vector of three numbers to label the scale bar (beginning, midpoint, end)
lwd	line width for the "line" type scalebar
...	arguments to be passed to other methods

**Value**

None. Use for side effect of a scalebar added to a plot

**Author(s)**

Robert J. Hijmans; partly based on a function by Josh Gray

**See Also**

[plot](#)

**Examples**

```
f <- system.file("external/test.grd", package="raster")
r <- raster(f)
plot(r)
scalebar(1000)
scalebar(1000, xy=c(178000, 333500), type='bar', divs=4)
```

---

select

*Geometric subsetting*

---

**Description**

Geometrically subset Raster\* or Spatial\* objects by drawing on a plot (map).

**Usage**

```
## S4 method for signature 'Raster'
select(x, use='rec', ...)

## S4 method for signature 'Spatial'
select(x, use='rec', draw=TRUE, col='cyan', size=2, ...)
```

**Arguments**

x	Raster*, SpatialPoints*, SpatialLines*, or SpatialPolygons*
use	character: 'rec' or 'pol'. To use a rectangle or a polygon for selecting
draw	logical. Add the selected features to the plot?
col	color to use to draw the selected features (when draw=TRUE)
size	integer > 0. Size to draw the selected features with (when draw=TRUE))
...	additional arguments. None implemented

**Value**

Raster\* or Spatial\* object

**See Also**

[click](#), [crop](#)

**Examples**

```
## Not run:

# select a subset of a RasterLayer
r <- raster(nrow=10, ncol=10)
values(r) <- 1:ncell(r)
plot(r)
s <- select(r) # now click on the map twice

# plot the selection on a new canvas:
x11()
plot(s)

# select a subset of a SpatialPolygons object
p1 <- rbind(c(-180,-20), c(-140,55), c(10, 0), c(-140,-60), c(-180,-20))
hole <- rbind(c(-150,-20), c(-100,-10), c(-110,20), c(-150,-20))
p2 <- rbind(c(-10,0), c(140,60), c(160,0), c(140,-55), c(-10,0))
p3 <- rbind(c(-125,0), c(0,60), c(40,5), c(15,-45), c(-125,0))
pols <- SpatialPolygons( list( Polygons(list(Polygon(p1), Polygon(hole)), 1),
  Polygons(list(Polygon(p2)), 2), Polygons(list(Polygon(p3)), 3)))
pols@polygons[[1]]@Polygons[[2]]@hole <- TRUE

plot(pols, col=rainbow(3))
ps <- select(pols) # now click on the map twice
ps

## End(Not run)
```

---

setExtent	<i>Set the extent of a RasterLayer</i>
-----------	--

---

### Description

setExtent sets the extent of a Raster\* object. Either by providing a new Extent object or by setting the extreme coordinates one by one.

### Usage

```
setExtent(x, ext, keepres=FALSE, snap=FALSE)
extent(x) <- value
```

### Arguments

x	A Raster* object
ext	An object of class Extent (which you can create with <a href="#">extent</a> , or an object that has an extent (e.g. a Raster* or Spatial* object) )
keepres	logical. If TRUE, the resolution of the cells will stay the same after adjusting the bounding box (by adjusting the number of rows and columns). If FALSE, the number of rows and columns will stay the same, and the resolution will be adjusted.
snap	logical. If TRUE, the extent is adjusted so that the cells of the input and output RasterLayer are aligned
value	An object of class Extent (which you can create with <a href="#">extent</a> )

### Value

a Raster\* object

### See Also

[extent](#), [Extent-class](#)

### Examples

```
r <- raster()
bb <- extent(-10, 10, -20, 20)
extent(r) <- bb
r <- setExtent(r, bb, keepres=TRUE)
```

---

setMinMax	<i>Compute min and max values</i>
-----------	-----------------------------------

---

**Description**

The minimum and maximum value of a RasterLayer are computed (from a file on disk if necessary) and stored in the returned Raster\* object.

**Usage**

```
setMinMax(x, ...)
```

**Arguments**

x	Raster object
...	additional arguments, none implemented

**Value**

Raster object

**See Also**

[getValues](#)

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
r
r <- setMinMax(r)
r
```

---

setValues	<i>Set values of a Raster object</i>
-----------	--------------------------------------

---

**Description**

Assign (new) values to a Raster\* object.

**Usage**

```
## S4 method for signature 'RasterLayer'
setValues(x, values, ...)

## S4 method for signature 'RasterBrick'
setValues(x, values, layer=-1, ...)

## S4 method for signature 'RasterStack'
setValues(x, values, layer=-1, ...)

## S4 method for signature 'RasterLayerSparse'
setValues(x, values, index=NULL, ...)

values(x) <- value
```

**Arguments**

x	A Raster*
values	Cell values to associate with the Raster* object. There should be values for all cells
value	Cell values to associate with the Raster* object. There should be values for all cells
layer	Layer number (only relevant for RasterBrick and RasterStack objects). If missing, the values of all layers is set
index	Cell numbers corresponding to the values
...	Additional arguments (none implemented)

**Value**

a Raster\* object

**Note**

While you can access the 'values' slot of the objects directly, you would do that at your own peril because when setting values, multiple slots need to be changed; which is what setValues takes care of.

**See Also**

[replacement](#)

**Examples**

```
r <- raster(ncol=10, nrow=10)
vals <- 1:ncell(r)
r <- setValues(r, vals)
# equivalent to
values(r) <- vals
```



---

shapefile	<i>Read or write a shapefile</i>
-----------	----------------------------------

---

### Description

Reading and writing of "ESRI shapefile" format spatial data. Only the three vector types (points, lines, and polygons) can be stored in shapefiles. These are simple wrapper functions around readOGR and writeOGR (rgdal package). A shapefile should consist of at least four files: .shp (the geometry), .dbf (the attributes), .shx (the index that links the two, and .prj (the coordinate reference system). If the .prj file is missing, a warning is given. If any other file is missing an error occurs (although one could in principle recover the .shx from the .shp file). Additional files are ignored.

### Usage

```
## S4 method for signature 'character'
shapefile(x, stringsAsFactors=FALSE, verbose=FALSE, warnPRJ=TRUE, ...)

## S4 method for signature 'Spatial'
shapefile(x, filename='', overwrite=FALSE, ...)
```

### Arguments

x	character (a file name, when reading a shapefile) or Spatial* object (when writing a shapefile)
filename	character. Filename to write a shapefile
overwrite	logical. Overwrite existing shapefile?
verbose	logical. If TRUE, information about the file is printed
warnPRJ	logical. If TRUE, a warning is given if there is no .prj file
stringsAsFactors	logical. If TRUE, strings are converted to factors
...	Additional arguments passed to rgdal functions readOGR or writeOGR

### Value

Spatial\*DataFrame (reading). Nothing is returned when writing a shapefile.

### Examples

```
if (require(rgdal)) {
  filename <- system.file("external/lux.shp", package="raster")
  filename
  p <- shapefile(filename)

  ## Not run:
```

```
shapefile(p, 'copy.shp')

## End(Not run)
}
```

---

 shift

*Shift*


---

### Description

Shift the location of a Raster\* of vector type Spatial\* object in the x and/or y direction

### Usage

```
## S4 method for signature 'Raster'
shift(x, dx=0, dy=0, filename='', ...)

## S4 method for signature 'SpatialPolygons'
shift(x, dx=0, dy=0, ...)

## S4 method for signature 'SpatialLines'
shift(x, dx=0, dy=0, ...)

## S4 method for signature 'SpatialPoints'
shift(x, dx=0, dy=0, ...)
```

### Arguments

x	Raster* or Spatial* object
dx	numeric. The shift in horizontal direction
dy	numeric. The shift in vertical direction
filename	character file name (optional)
...	if x is a Raster* object: additional arguments as for <a href="#">writeRaster</a>

### Value

Same object type as x

### See Also

[flip](#), [rotate](#), and the [elide](#) function in the [maptools](#) package

### Examples

```
r <- raster()
r <- shift(r, dx=1, dy=-1)
```

---

Slope and aspect      *Slope and aspect*

---

### Description

DEPRACATED. Use [terrain](#) instead.

### Usage

```
slopeAspect(dem, filename='', out=c('slope', 'aspect'), unit='radians',
            neighbors=8, flatAspect, ...)
```

### Arguments

dem	DEPRACATED
filename	DEPRACATED
out	DEPRACATED
unit	DEPRACATED
neighbors	DEPRACATED
flatAspect	DEPRACATED
...	DEPRACATED

### See Also

[terrain](#)

---

sp      *Create SpatialLines\* or SpatialPolygons\**

---

### Description

Helper functions to simplify the creation of `SpatialLines*` or `SpatialPolygons*` objects from coordinates.

### Usage

```
spLines(x, ..., attr=NULL, crs="")
spPolygons(x, ..., attr=NULL, crs="")
```

**Arguments**

<code>x</code>	matrix of list with matrices. Each matrix must have two columns with x and y coordinates (or longitude and latitude, in that order). Multi-line or multi-polygon objects can be formed by combining matrices in a list
<code>...</code>	additional matrices and/or lists with matrices
<code>attr</code>	data.frame with the attributes to create a *DataFrame object. The number of rows must match the number of lines/polgyons
<code>crs</code>	the coordinate reference system (PROJ4 or WKT notation)

**Value**

SpatialLines\* or SpatialPolygons\*

**Examples**

```
x1 <- rbind(c(-180,-20), c(-140,55), c(10, 0), c(-140,-60))
x2 <- rbind(c(-10,0), c(140,60), c(160,0), c(140,-55))
x3 <- rbind(c(-125,0), c(0,60), c(40,5), c(15,-45))
x4 <- rbind(c(41,-41.5), c(51,-35), c(62,-41), c(51,-50))

a <- spLines(x1, x2, x3)
b <- spLines(x1, list(x2, x3), attr=data.frame(id=1:2), crs='+proj=longlat +datum=WGS84')
b

hole <- rbind(c(-150,-20), c(-100,-10), c(-110,20), c(-130,10))
d <- spPolygons(list(x1,hole), x2, list(x3, x4))

att <- data.frame(ID=1:3, name=c('a', 'b', 'c'))
e <- spPolygons(list(x1,hole), x2, list(x3, x4), attr=att, crs='+proj=longlat +datum=WGS84')
e
```

---

spplot

*Use spplot to plot a Raster\* or other object*

---

**Description**

A wrapper function around [spplot](#) (sp package). With spplot it is easy to map several layers with a single legend for all maps. `ssplot` is itself a wrapper around the [levelplot](#) function in the lattice package, and see the help for these functions for additional options.

One of the advantages of the wrapper function for Raster\* objects is the additional `maxpixels` argument to sample large objects for faster drawing.

There are also added spplot methods for Spatial objects that have no data.frame and for SpatVector (terra package)

**Usage**

```
## S4 method for signature 'Raster'
spplot(obj, ..., maxpixels=50000, as.table=TRUE, zlim)
```

**Arguments**

obj	Raster* object
...	Any argument that can be passed to <a href="#">splot</a> and <a href="#">levelplot</a>
maxpixels	integer. Number of pixels to sample from each layer of large Raster objects
as.table	If TRUE, the plots are ordered from top to bottom
zlim	Vector of two elements indicating the minimum and maximum values to be mapped (values outside that range are set to these limits)

**See Also**

[plot](#), [plotRGB](#)

The rasterVis package has more advanced plotting methods for Raster objects

**Examples**

```
r <- raster(system.file("external/test.grd", package="raster"))
s <- stack(r, r*2)
names(s) <- c('meuse', 'meuse x 2')

splot(s)

pts <- data.frame(sampleRandom(r, 10, xy=TRUE))
coordinates(pts) <- ~ x + y

splot(s, scales = list(draw = TRUE),
      xlab = "easting", ylab = "northing",
      col.regions = rainbow(99, start=.1),
      names.attr=c('original', 'times two'),
      sp.layout = list("sp.points", pts, pch=20, cex=2, col='black'),
      par.settings = list(fontsize = list(text = 12)), at = seq(0, 4000, 500))
```

---

stack

---

*Create a RasterStack object*


---

**Description**

A RasterStack is a collection of RasterLayer objects with the same spatial extent and resolution. A RasterStack can be created from RasterLayer objects, or from raster files, or both. It can also be created from a SpatialPixelsDataFrame or a SpatialGridDataFrame object.

**Usage**

```
## S4 method for signature 'character'
stack(x, ..., bands=NULL, varname="", native=FALSE, RAT=TRUE, quick=FALSE)

## S4 method for signature 'Raster'
stack(x, ..., layers=NULL)

## S4 method for signature 'missing'
stack(x)

## S4 method for signature 'list'
stack(x, bands=NULL, native=FALSE, RAT=TRUE, ...)
```

**Arguments**

x	filename (character), Raster* object, missing (to create an empty RasterStack), SpatialGrid*, SpatialPixels*, or list (of filenames and/or Raster* objects). If x is a list, additional arguments ... are ignored
bands	integer. which bands (layers) of the file should be used (default is all layers)
layers	integer (or character with layer names) indicating which layers of a RasterBrick should be used (default is all layers)
native	logical. If TRUE native drivers are used instead of gdal drivers (where available, such as for BIL and Arc-ASCII files)
RAT	logical. If TRUE a raster attribute table is created for files that have one
quick	logical. If TRUE the extent and resolution of the objects are not compared. This speeds up the creation of the RasterStack but should be use with great caution. Only use this option when you are absolutely sure that all the data in all the files are aligned, and you need to create RasterStack for many (>100) files
varname	character. To select the variable of interest in a NetCDF file (see <a href="#">raster</a> )
...	additional filenames or Raster* objects

**Value**

RasterStack

**See Also**

[addLayer](#), [dropLayer](#), [raster](#), [brick](#)

**Examples**

```
# file with one layer
fn <- system.file("external/test.grd", package="raster")
s <- stack(fn, fn)
r <- raster(fn)
s <- stack(r, fn)
nlayers(s)
```

```
# file with three layers
slogo <- stack(system.file("external/rlogo.grd", package="raster"))
nlayers(slogo)
slogo
```

---

stackApply

*Apply a function on subsets of a RasterStack or RasterBrick*


---

### Description

Apply a function on subsets of a RasterStack or RasterBrick. The layers to be combined are indicated with the vector `indices`. The function used should return a single value, and the number of layers in the output Raster\* equals the number of unique values in `indices`. For example, if you have a RasterStack with 6 layers, you can use `indices=c(1,1,1,2,2,2)` and `fun=sum`. This will return a RasterBrick with two layers. The first layer is the sum of the first three layers in the input RasterStack, and the second layer is the sum of the last three layers in the input RasterStack. Indices are recycled such that `indices=c(1,2)` would also return a RasterBrick with two layers (one based on the odd layers (1,3,5), the other based on the even layers (2,4,6)).

See [calc](#) if you want to use a more efficient function that returns multiple layers based on `_all_` layers in the Raster\* object.

### Usage

```
stackApply(x, indices, fun, filename='', na.rm=TRUE, ...)
```

### Arguments

<code>x</code>	Raster* object
<code>indices</code>	integer. Vector of length <code>nlayers(x)</code> (shorter vectors are recycled) containing all integer values between 1 and the number of layers of the output Raster*
<code>fun</code>	function that returns a single value, e.g. <code>mean</code> or <code>min</code> , and that takes a <code>na.rm</code> argument (or can pass through arguments via <code>...</code> )
<code>na.rm</code>	logical. If TRUE, NA cells are removed from calculations
<code>filename</code>	character. Optional output filename
<code>...</code>	additional arguments as for <a href="#">writeRaster</a>

### Value

A new Raster\* object, and in some cases the side effect of a new file on disk.

### See Also

[calc](#), [stackSelect](#)

## Examples

```
r <- raster(ncol=10, nrow=10)
values(r) <- 1:ncell(r)
s <- brick(r,r,r,r,r,r)
s <- s * 1:6
b1 <- stackApply(s, indices=c(1,1,1,2,2,2), fun=sum)
b1
b2 <- stackApply(s, indices=c(1,2,3,1,2,3), fun=sum)
b2
```

---

stackSave

*Save or open a RasterStack file*

---

## Description

A RasterStack is a collection of RasterLayers with the same spatial extent and resolution. They can be created from RasterLayer objects, or from file names. These two functions allow you to save the references to raster files and recreate a rasterStack object later. They only work if the RasterStack points to layers that have their values on disk. The values are not saved, only the references to the files.

## Usage

```
stackOpen(stackfile)
stackSave(x, filename)
```

## Arguments

stackfile	Filename for the RasterStack (to save it on disk)
x	RasterStack object
filename	File name

## Details

When a RasterStack is saved to a file, only pointers (filenames) to raster datasets are saved, not the data. If the name or location of a raster file changes, the RasterStack becomes invalid.

## Value

RasterStack object

## See Also

[writeRaster](#), [stack](#), [addLayer](#)



**Examples**

```

file <- system.file("external/test.grd", package="raster")
s <- stack(c(file, file))

## Not run:
s <- stackSave(s, "mystack")
# note that filename adds an extension .stk to a stackfile
s2 <- stackOpen("mystack.stk")
s2

## End(Not run)

```

stackSelect

*Select cell values from a multi-layer Raster\* object***Description**

Use a Raster\* object to select cell values from different layers in a multi-layer Raster\* object. The object to select values y should have values between 1 and nlayers(x). The values of y are rounded.

See [extract](#) for extraction of values by cell, point, or otherwise.

**Usage**

```

## S4 method for signature 'RasterStackBrick,Raster'
stackSelect(x, y, recycle=FALSE, type='index', filename='', ...)

```

**Arguments**

x	RasterStack or RasterBrick object
y	Raster* object
recycle	Logical. Recursively select values (default = FALSE. Only relevant if y has multiple layers. E.g. if x has 12 layers, and y has 4 layers, the indices of the y layers are used three times.
type	Character. Only relevant when recycle=TRUE. Can be 'index' or 'truefalse'. If it is 'index', the cell values of y should represent layer numbers. If it is 'truefalse' layer numbers are indicated by 0 (not used, NA returned) and 1 (used)
filename	Character. Output filename (optional)
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

Raster\* object

**See Also**

[stackApply](#), [extract](#)

**Examples**

```
r <- raster(ncol=10, nrow=10, vals=1)
s <- stack(r, r+2, r+5)
values(r) <- round((runif(ncell(r)))*3)
x <- stackSelect(s, r)
```

stretch

*Stretch***Description**

Linear stretch of values in a Raster object. Provide the desired output range (minv and maxv) and the lower and upper bounds in the original data, either as quintiles (if minq=0 and maxq=1 you use the minimum and maximum cell values), or as actual values (smin and smax; e.g. precomputed quantile values). If smin and smax are both not NA, minq and maxq are ignored.

**Usage**

```
## S4 method for signature 'Raster'
stretch(x, minv=0, maxv=255, minq=0, maxq=1, smin=NA, smax=NA,
        samplesize=1000000, filename='', ...)
```

**Arguments**

x	Raster object
minv	numeric $\geq 0$ and smaller than maxv. lower bound of stretched value
maxv	numeric $\leq 255$ and larger than minv. upper bound of stretched value
minq	numeric $\geq 0$ and smaller than maxq. lower quintile bound of original value. Ignored if smin is supplied
maxq	numeric $\leq 1$ and larger than minq. upper quintile bound of original value. Ignored if smax is supplied
smin	numeric $< smax$ . user supplied lower value for the layers, to be used instead of a quintile computed by the function itself
smax	numeric $> smin$ . user supplied upper value for the layers, to be used instead of a quintile computed by the function itself
samplesize	numeric $> 1$ . If samplesize $< ncell(x)$ , a regular sample of samplesize is taken from x to compute the quintiles (to speed things up)
filename	character. Filename for the output Raster object (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

Raster

**See Also**

stretch argument in [plotRGB](#)

**Examples**

```
r <- raster(nc=10, nr=10)
values(r) <- rep(1:2, 50)
stretch(r)
s <- stack(r, r*2)
stretch(s)
```

---

subset

*Subset layers in a Raster\* object*


---

**Description**

Extract a set of layers from a RasterStack or RasterBrick object.

**Usage**

```
## S4 method for signature 'Raster'
subset(x, subset, drop=TRUE, filename='', ...)

## S4 method for signature 'RasterStack'
subset(x, subset, drop=TRUE, filename='', ...)
```

**Arguments**

x	RasterBrick or RasterStack object
subset	integer or character. Should indicate the layers (represented as integer or by their name)
drop	If TRUE, a selection of a single layer will be returned as a RasterLayer
filename	character. Output filename (optional)
...	additional arguments as for <a href="#">writeRaster</a>

**Value**

Raster\* object

**See Also**

[dropLayer](#)

**Examples**

```
s <- stack(system.file("external/rlogo.grd", package="raster"))
sel <- subset(s, 2:3)

# Note that this is equivalent to
sel2 <- s[[2:3]]

# and in this particular case:
sel3 <- dropLayer(s, 1)

nlayers(s)
nlayers(sel)

# effect of 'drop=FALSE' when selecting a single layer
sel <- subset(s, 2)
class(sel)
sel <- subset(s, 2, drop=FALSE)
class(sel)
```

---

substitute

*Substitute values in a Raster\* object*


---

**Description**

Substitute (replace) values in a Raster\* object with values in a data.frame. The data.frame should have a column to identify the key (ID) to match with the cell values of the Raster\* object, and one or more columns with replacement values. By default these are the first and second column but you can specify other columns with arguments by and which. It is possible to match one table to multiple layers, or to use multiple layers as a single key, but not both.

**Usage**

```
## S4 method for signature 'Raster,data.frame'
subs(x, y, by=1, which=2, subsWithNA=TRUE, filename='', ...)
```

**Arguments**

x	Raster* object
y	data.frame
by	column number(s) or name(s) identifying the key (ID) to match rows in data.frame y to values of the Raster object
which	column number or name that has the new (replacement) values
subsWithNA	logical. If TRUE values that are not matched become NA. If FALSE, they retain their original value (which could also be NA). This latter option is handy when you want to replace only one or a few values. It cannot be used when x has multiple layers

filename            character. Optional output filename  
 ...                additional arguments as for [writeRaster](#)

### Details

You could obtain the same result with [reclassify](#), but `subs` is more efficient for simple replacement. Use `reclassify` if you want to replace ranges of values with new values.

You can also replace values using a fitted model. E.g. fit a model to `glm` or `loess` and then call [predict](#)

### Value

Raster object

### See Also

[reclassify](#), [clamp](#), [cut](#)

### Examples

```
r <- raster(ncol=10, nrow=10)
values(r) <- round(runif(ncell(r)) * 10)
df <- data.frame(id=2:8, v=c(10,10,11,11,12:14))
x <- subs(r, df)
x2 <- subs(r, df, subsWithNA=FALSE)

df$v2 <- df$v * 10
x3 <- subs(r, df, which=2:3)

s <- stack(r, r*3)
names(s) <- c('first', 'second')
x4 <- subs(s, df)
x5 <- subs(s, df, which=2:3)
```

---

Summary

*Summary*

---

### Description

Summarize a Raster\* object. A sample is used for very large files.

### Usage

```
## S4 method for signature 'RasterLayer'
summary(object, maxsamp=100000, ...)
```

**Arguments**

object	Raster* object
maxsamp	positive integer. Sample size used for large datasets
...	additional arguments. None implemented

**Value**

matrix with (an estimate of) the median, minimum and maximum values, the first and third quartiles, and the number of cells with NA values

**See Also**

[cellStats](#), [link\[raster\]{quantile}](#)

---

Summary-methods

*Summary methods*

---

**Description**

The following summary methods are available for Raster\* objects:

`mean`, `median`, `max`, `min`, `range`, `prod`, `sum`, `any`, `all`

All methods take `na.rm` as an additional logical argument. Default is `na.rm=FALSE`. If `TRUE`, NA values are removed from calculations. These methods compute a summary statistic based on cell values of RasterLayers and the result of these methods is always a single RasterLayer (except for `range`, which returns a RasterBrick with two layers). See [calc](#) for functions not included here (e.g. `median`) or any other custom functions.

You can mix RasterLayer, RasterStack and RasterBrick objects with single numeric or logical values. However, because generic functions are used, the method applied is chosen based on the first argument: `'x'`. This means that if `r` is a RasterLayer object, `mean(r, 5)` will work, but `mean(5, r)` will not work.

To summarize all cells within a single RasterLayer, see [cellStats](#) and [maxValue](#) and [minValue](#)

**Value**

a RasterLayer

**See Also**

[calc](#)

**Examples**

```
r1 <- raster(nrow=10, ncol=10)
r1 <- setValues(r1, runif(ncell(r1)))
r2 <- setValues(r1, runif(ncell(r1)))
r3 <- setValues(r1, runif(ncell(r1)))
r <- max(r1, r2, r3)
r <- range(r1, r2, r3, 1.2)

s <- stack(r1, r2, r3)
r <- mean(s, 2)
```

---

symdif

*Symmetrical difference*

---

**Description**

Symmetrical difference of SpatialPolygons\* objects

**Usage**

```
## S4 method for signature 'SpatialPolygons,SpatialPolygons'
symdif(x, y, ...)
```

**Arguments**

x	SpatialPolygons* object
y	SpatialPolygons* object
...	Additional SpatialPolygons* object(s)

**Value**

SpatialPolygons\*

**Author(s)**

Robert J. Hijmans

**See Also**

[erase](#)

## Examples

```
#SpatialPolygons
if (require(rgdal) & require(rgeos)) {
  p <- shapefile(system.file("external/lux.shp", package="raster"))
  b <- as(extent(6, 6.4, 49.75, 50), 'SpatialPolygons')
  crs(b) <- crs(p)
  sd <- symdif(p, b)
  plot(sd, col='red')
}
```

---

 terrain

*Terrain characteristics*


---

## Description

Compute slope, aspect and other terrain characteristics from a raster with elevation data. The elevation data should be in map units (typically meter) for projected (planar) raster data. They should be in meters when the coordinate reference system (CRS) is longitude/latitude.

## Usage

```
## S4 method for signature 'RasterLayer'
terrain(x, opt="slope", unit="radians", neighbors=8, filename="", ...)
```

## Arguments

x	RasterLayer object with elevation values. Values should have the same unit as the map units, or in meters when the crs is longitude/latitude
opt	Character vector containing one or more of these options: slope, aspect, TPI, TRI, roughness, flowdir (see Details)
unit	Character. 'degrees', 'radians' or 'tangent'. Only relevant for slope and aspect. If 'tangent' is selected that is used for slope, but for aspect 'degrees' is used (as 'tangent' has no meaning for aspect)
neighbors	Integer. Indicating how many neighboring cells to use to compute slope for any cell. Either 8 (queen case) or 4 (rook case). Only used for slope and aspect, see Details
filename	Character. Output filename (optional)
...	Standard additional arguments for writing Raster* objects to file

## Details

When neighbors=4, slope and aspect are computed according to Fleming and Hoffer (1979) and Ritter (1987). When neighbors=8, slope and aspect are computed according to Horn (1981). The Horn algorithm may be best for rough surfaces, and the Fleming and Hoffer algorithm may be better for smoother surfaces (Jones, 1997; Burrough and McDonnell, 1998). If slope = 0, aspect is set to  $0.5\pi$  radians (or 90 degrees if unit='degrees'). When computing slope or aspect, the CRS



([projection](#)) of the RasterLayer `x` must be known (may not be NA), to be able to safely differentiate between planar and longitude/latitude data.

`flowdir` returns the 'flow direction' (of water), i.e. the direction of the greatest drop in elevation (or the smallest rise if all neighbors are higher). They are encoded as powers of 2 (0 to 7). The cell to the right of the focal cell 'x' is 1, the one below that is 2, and so on:

```

32  64  128
16   x   1
 8   4   2

```

If two cells have the same drop in elevation, a random cell is picked. That is not ideal as it may prevent the creation of connected flow networks. ArcGIS implements the approach of Greenlee (1987) and I might adopt that in the future.

The terrain indices are according to Wilson et al. (2007), as in [gdaldem](#). TRI (Terrain Ruggedness Index) is the mean of the absolute differences between the value of a cell and the value of its 8 surrounding cells. TPI (Topographic Position Index) is the difference between the value of a cell and the mean value of its 8 surrounding cells. Roughness is the difference between the maximum and the minimum value of a cell and its 8 surrounding cells.

Such measures can also be computed with the [focal](#) function:

```

f <- matrix(1, nrow=3, ncol=3)
TRI <- focal(x, w=f, fun=function(x, ...) sum(abs(x[-5]-x[5]))/8, pad=TRUE, padValue=NA)
TPI <- focal(x, w=f, fun=function(x, ...) x[5] - mean(x[-5]), pad=TRUE, padValue=NA)
rough <- focal(x, w=f, fun=function(x, ...) max(x) - min(x), pad=TRUE, padValue=NA, na.rm=TRUE)

```

## References

- Burrough, P., and R.A. McDonnell, 1998. Principles of Geographical Information Systems. Oxford University Press.
- Fleming, M.D. and Hoffer, R.M., 1979. Machine processing of landsat MSS data and DMA topographic data for forest cover type mapping. LARS Technical Report 062879. Laboratory for Applications of Remote Sensing, Purdue University, West Lafayette, Indiana.
- Greenlee, D.D., 1987. Raster and vector processing for scanned linework. Photogrammetric Engineering and Remote Sensing 53:1383-1387
- Horn, B.K.P., 1981. Hill shading and the reflectance map. Proceedings of the IEEE 69:14-47
- Jones, K.H., 1998. A comparison of algorithms used to compute hill slope as a property of the DEM. Computers & Geosciences 24: 315-323
- Ritter, P., 1987. A vector-based slope and aspect generation algorithm. Photogrammetric Engineering and Remote Sensing 53: 1109-1111
- Wilson, M.F.J., O'Connell, B., Brown, C., Guinan, J.C., Grehan, A.J., 2007. Multiscale terrain analysis of multibeam bathymetry data for habitat mapping on the continental slope. Marine Geodesy 30: 3-35.

**See Also**[hillShade](#)**Examples**

```
## Not run:
elevation <- getData('alt', country='CHE')
x <- terrain(elevation, opt=c('slope', 'aspect'), unit='degrees')
plot(x)

# TPI for different neighborhood size:
tpiw <- function(x, w=5) {
  m <- matrix(1/(w^2-1), nc=w, nr=w)
  m[ceiling(0.5 * length(m))] <- 0
  f <- focal(x, m)
  x - f
}
tpi5 <- tpiw(elevation, w=5)

## End(Not run)
```

---

text

---

*Add labels to a map*


---

**Description**

Plots labels, that is a textual (rather than color) representation of values, on top an existing plot (map).

**Usage**

```
## S4 method for signature 'RasterLayer'
text(x, labels, digits=0, fun=NULL, halo=FALSE, ...)

## S4 method for signature 'RasterStackBrick'
text(x, labels, digits=0, fun=NULL, halo=FALSE, ...)

## S4 method for signature 'SpatialPolygons'
text(x, labels, halo=FALSE, ...)

## S4 method for signature 'SpatialPoints'
text(x, labels, halo=FALSE, ...)
```

**Arguments**

x Raster\*, SpatialPoints\* or SpatialPolygons\* object

labels	character. Optional. Vector of labels with length(x) or a variable name from names(x)
digits	integer. how many digits should be used?
fun	function to subset the values plotted (as in <a href="#">rasterToPoints</a> )
halo	logical. If TRUE a 'halo' is printed around the text. If TRUE, additional arguments hc='white' and hw=0.1 can be modified to set the colour and width of the halo
...	additional arguments to pass to graphics function <a href="#">text</a>

**See Also**

[text](#), [plot](#)

**Examples**

```
r <- raster(nrows=4, ncols=4)
r <- setValues(r, 1:ncell(r))
plot(r)
text(r)

plot(r)
text(r, halo=TRUE, hc='blue', col='white', hw=0.2)

plot(r, col=bpy.colors(5))
text(r, fun=function(x){x<5 | x>12}, col=c('red', 'white'), vfont=c("sans serif", "bold"), cex=2)
```

---

transpose

*Transpose*

---

**Description**

Transpose a Raster\* object

**Usage**

```
t(x)
```

**Arguments**

x                    a Raster\* object

**Value**

RasterLayer or RasterBrick

**See Also**

transpose: [flip](#), [rotate](#)

**Examples**

```
r <- raster(nrow=18, ncol=36)
values(r) <- 1:ncell(r)
rt <- t(r)
```

---

trim

*Trim*


---

**Description**

Trim (shrink) a Raster\* object by removing outer rows and columns that all have the same value (e.g. NA).

Or remove the whitespace before or after a string of characters (or a matrix, or the character values in a data.frame).

**Usage**

```
## S4 method for signature 'Raster'
trim(x, padding=0, values=NA, filename='', ...)
## S4 method for signature 'character'
trim(x, internal=FALSE, ...)
```

**Arguments**

x	Raster* object or a character string
values	numeric. Value(s) based on which a Raster* should be trimmed
padding	integer. Number of outer rows/columns to keep
filename	character. Optional output filename
internal	logical. If TRUE, sequential internal spaces are replaced by a single space
...	If x is a Raster* object: additional arguments as for <a href="#">writeRaster</a>

**Value**

A RasterLayer or RasterBrick object (if x is a Raster\* object) or a character string (if x is a character string).

**Author(s)**

Robert J. Hijmans and Jacob van Etten

**Examples**

```
r <- raster(ncol=18,nrow=18)
r[39:49] <- 1
r[113:155] <- 2
r[200] <- 6
s <- trim(r)
```

```
trim(" hi folks ! ")
```

---

union

*Union Extent or SpatialPolygons\* objects*


---

**Description**

Extent objects: Objects are combined into their union. See [crop](#) and [extend](#) to union a Raster object with an Extent object.

Two SpatialPolygons\* objects. Overlapping polygons (between layers, not within layers) are intersected, other spatial objects are appended. Tabular attributes are joined. See [bind](#) if you want to combine polygons without intersection.

Single SpatialPolygons\* object. Overlapping polygons are intersected. Original attributes are lost. New attributes allow for determining how many, and which, polygons overlapped.

Union for SpatialLines and SpatialPoints simply combines the two data sets; without any geometric intersections. This is equivalent to [bind](#).

**Usage**

```
## S4 method for signature 'Extent,Extent'
union(x, y)

## S4 method for signature 'SpatialPolygons,SpatialPolygons'
union(x, y)

## S4 method for signature 'SpatialPolygons,missing'
union(x, y)

## S4 method for signature 'SpatialLines,SpatialLines'
union(x, y)

## S4 method for signature 'SpatialPoints,SpatialPoints'
union(x, y)
```

**Arguments**

```
x          Extent or SpatialPolygons* object
y          Same as x or missing
```

**Value**

Extent or SpatialPolygons object

**See Also**

[intersect](#), [extent](#), [setExtent](#)

[merge](#) for merging a data.frame with attributes of Spatial objects and [+](#), [SpatialPolygons](#), [SpatialPolygons-method](#) for an algebraic notation

**Examples**

```
e1 <- extent(-10, 10, -20, 20)
e2 <- extent(0, 20, -40, 5)
union(e1, e2)

#SpatialPolygons
if (require(rgdal) & require(rgeos)) {
  p <- shapefile(system.file("external/lux.shp", package="raster"))
  p0 <- aggregate(p)
  b <- as(extent(6, 6.4, 49.75, 50), 'SpatialPolygons')
  crs(b) <- crs(p)
  u <- union(p0, b)
  plot(u, col=2:4)
}
```

---

unique

*Unique values*

---

**Description**

This function returns the unique values in a RasterLayer object or the unique combinations of the layers in a multilayer object.

**Usage**

```
## S4 method for signature 'RasterLayer,missing'
unique(x, incomparables=FALSE, na.last=NA, progress="", ...)

## S4 method for signature 'RasterStackBrick,missing'
unique(x, incomparables=FALSE, na.last=NA, progress="", ...)
```

**Arguments**

x	Raster object
incomparables	must be missing. The default value FALSE is used. See <a href="#">unique</a>
na.last	logical. for controlling the treatment of NAs. If TRUE, missing values in the data are put last; if FALSE, they are put first; if NA, they are removed.
progress	character. Use "text" or "window" for a progress indicator
...	additional arguments. as in <a href="#">unique</a>

**Value**

vector or matrix

**See Also**

[unique](#)

**Examples**

```
r <- raster(ncol=10, nrow=10)
values(r) <- round(runif(ncell(r))*10)
unique(r)
u <- unique(stack(r, round(r/2)))
do.call(rbind, u)
lapply(u, unique)
```

---

unstack

*Unstack*

---

**Description**

Create a list of RasterLayer objects from a RasterStack or RasterBrick

**Usage**

```
unstack(x, ...)
```

**Arguments**

x                    a RasterStack object  
...                    not used. further arguments passed to or from other methods

**Value**

A list of RasterLayer objects

**See Also**

[stack](#)

**Examples**

```
file <- system.file("external/test.grd", package="raster")
s <- stack(file, file)
list1 <- unstack(s)
b <- brick(s)
list2 <- unstack(b)
```

update

*Update raster cells of files (on disk)***Description**

Update cell values of a file (i.e., cell values on disk) associated with a RasterLayer or RasterBrick.

User beware: this function `_will_` make changes to your file (first make a copy if you are not sure what you are doing).

Writing starts at a cell number `cell`. You can write a vector of values (in cell order), or a matrix. You can also provide a vector of cell numbers (of the same length as vector `v`) to update individual cells.

See [writeFormats](#) for supported formats.

**Usage**

```
## S4 method for signature 'RasterLayer'
update(object, v, cell, ...)
## S4 method for signature 'RasterBrick'
update(object, v, cell, band, ...)
```

**Arguments**

<code>object</code>	RasterLayer or RasterBrick that is associated with a file
<code>v</code>	vector or matrix with new values
<code>cell</code>	cell from where to start writing. Or a vector of cell numbers if <code>v</code> is a vector of the same length.
<code>band</code>	band (layer) to update (for RasterBrick objects).
<code>...</code>	additional arguments. None implemented

**Value**

RasterLayer or RasterBrick

**Examples**

```
## Not run:
# setting up an example RasterLayer with file
r <- raster(nrow=5, ncol=10, vals=0)
r <- writeRaster(r, rasterTmpFile(), overwrite=TRUE, datatype='INT2S')
as.matrix(r)

# update with a vector starting a cell
r <- update(r, v=rep(1, 5), cell=6)
# 99.99 gets rounded because this is an integer file
r <- update(r, v=9.99, cell=50)
as.matrix(r)
```



```

# update with a vector of values and matching vector of cell numbers
r <- update(r, v=5:1, cell=c(5,15,25,35,45))
as.matrix(r)

# updating with a matrix, anchored at a cell number
m <- matrix(1:10, ncol=2)
r <- update(r, v=m, cell=2)
as.matrix(r)

## End(Not run)

```

---

validCell

*Validity of a cell, column or row number*


---

### Description

Simple helper functions to determine if a row, column or cell number is valid for a certain Raster\* object

### Usage

```

validCell(object, cell)
validCol(object, colnr)
validRow(object, rownr)

```

### Arguments

object	Raster* object (or a SpatialPixels* or SpatialGrid* object)
cell	cell number(s)
colnr	column number; or vector of column numbers
rownr	row number; or vector of row numbers

### Value

logical value

### Examples

```

#using a new default raster (1 degree global)
r <- raster()
validCell(r, c(-1, 0, 1))
validRow(r, c(-1, 1, 100, 10000))

```

---

validNames	<i>Create valid names</i>
------------	---------------------------

---

**Description**

Create a set of valid names (trimmed, no duplicates, not starting with a number).

**Usage**

```
validNames(x, prefix='layer')
```

**Arguments**

x	character
prefix	character string used if x is empty

**Value**

character

**See Also**

[make.names](#)

**Examples**

```
validNames(c('a', 'a', '', '1', NA, 'b', 'a'))
```

---

weighted.mean	<i>Weighted mean of rasters</i>
---------------	---------------------------------

---

**Description**

Computes the weighted mean for each cell of a number or raster layers. The weights can be spatially variable or not.

**Usage**

```
## S4 method for signature 'RasterStackBrick,vector'
weighted.mean(x, w, na.rm=FALSE, filename='', ...)

## S4 method for signature 'RasterStackBrick,RasterStackBrick'
weighted.mean(x, w, na.rm=FALSE, filename='', ...)
```

**Arguments**

x	RasterStack or RasterBrick
w	A vector of weights (one number for each layer), or for spatially variable weights, a RasterStack or RasterBrick with weights (should have the same extent, resolution and number of layers as x)
na.rm	Logical. Should missing values be removed?
filename	Character. Output filename (optional)
...	Additional arguments as for <a href="#">writeRaster</a>

**Value**

RasterLayer

**See Also**

[Summary-methods](#), [weighted.mean](#)

**Examples**

```
b <- brick(system.file("external/rlogo.grd", package="raster"))

# give least weight to first layer, most to last layer
wm1 <- weighted.mean(b, w=1:3)

# spatially varying weights
# weigh by column number
w1 <- init(b, v='col')

# weigh by row number
w2 <- init(b, v='row')
w <- stack(w1, w2, w2)

wm2 <- weighted.mean(b, w=w)
```

---

which	<i>Which cells are TRUE?</i>
-------	------------------------------

---

**Description**

Which returns a RasterLayer with TRUE or FALSE setting cells that are NA to FALSE (unless `na.rm=FALSE`). If the RasterLayer has numbers, all values that are 0 become FALSE and all other values become TRUE. The function can also return the cell numbers that are TRUE

**Usage**

```
## S4 method for signature 'RasterLayer'
Which(x, cells=FALSE, na.rm=TRUE, ...)
```

**Arguments**

x	RasterLayer
cells	logical. If TRUE, cell numbers are returned, otherwise a RasterLayer is returned
na.rm	logical. If TRUE, NA values are treated as FALSE, otherwise they remain NA (only when cells=FALSE)
...	Additional arguments (none implemented)

**Value**

RasterLayer

**See Also**

[which.max](#), [which.min](#)

**Examples**

```
r <- raster(ncol=10, nrow=10)
set.seed(0)
values(r) <- runif(ncell(r))
r[r < 0.2 ] <- 0
r[r > 0.8] <- 1
r[r > 0 & r < 1 ] <- 0.5

Which(r, cells=TRUE)
Which(r > 0.5, cells=TRUE)

s1 <- r > 0.5
s2 <- Which(r > 0.5)
s1[1:15]
s2[1:15]

# this expression
x1 <- Which(r, na.rm=FALSE)
# is the inverse of
x2 <- r==0
```

---

which.min

*Where is the min or max value?*

---

**Description**

Which cells have the minimum / maximum value (for a RasterLayer), or which layer has the minimum/maximum value (for a RasterStack or RasterBrick)?

which.min and which.max return the index of the first layer that has the min or max value for a cell. This can be problematic if there are ties.

In you want the index of all the layers that have the min or max value, use whiches.min or whiches.max (only for objects with less than 10 layers).

**Usage**

```
which.min(x)
which.max(x)
whiches.min(x, ...)
whiches.max(x, ...)
```

**Arguments**

x	Raster* object
...	additional arguments (none implemented)

**Value**

(which.\*): vector of cell numbers (if x is a RasterLayer). If x is a RasterStack or RasterBrick, a RasterLayer giving the number of the first layer with the minimum or maximum value for a cell.

(whiches.\*). An integer in which each digit represents a layer. For example, 35 means "layers 3 and 5"

**Note**

There is a limit to accurate integer number representation. Therefore, do not use `whiches.*` with more than 15 layers.

**See Also**

[Which](#)

**Examples**

```
b <- brick(system.file("external/rlogo.grd", package="raster"))

r <- which.min(b)

i <- which.min(b[[3]])
xy <- xyFromCell(b, i)
plot(b[[3]])
points(xy)

x <- welches.min(b)
freq(x)
```

---

writeFormats *File types for writing*

---

### Description

List supported file types for writing RasterLayer values to disk.

When a function writes a file to disk, the file format is determined by the 'format=' argument if supplied, or else by the file extension (if the extension is known). In other cases the default format is used. The 'factory-fresh' default format is 'raster', but this can be changed using [rasterOptions](#).

### Usage

```
writeFormats()
```

### Details

writeFormats returns a matrix of the file formats (the "drivers") that are supported.

Supported formats include:

File type	Long name	default extension	Multiband support
raster	'Native' raster package format	.grd	Yes
ascii	ESRI Ascii	.asc	No
SAGA	SAGA GIS	.sdatt	No
IDRISI	IDRISI	.rst	No
CDF	netCDF (requires ncdf4)	.nc	Yes
GTiff	GeoTiff (requires rgdal)	.tif	Yes
ENVI	ENVI .hdr Labelled	.envi	Yes
EHdr	ESRI .hdr Labelled	.bil	Yes
HFA	Erdas Imagine Images (.img)	.img	Yes

### See Also

[GDALDriver-class](#)

### Examples

```
writeFormats()
```

---

writeRaster *Write raster data to a file*

---

## Description

Write an entire Raster\* object to a file, using one of the many supported formats. See [writeValues](#) for writing in chunks (e.g. by row).

When writing a file to disk, the file format is determined by the 'format=' argument if supplied, or else by the file extension (if the extension is known). In other cases the default format is used. The default format is 'raster', but this setting can be changed (see [rasterOptions](#)).

## Usage

```
## S4 method for signature 'RasterLayer,character'
writeRaster(x, filename, format, ...)

## S4 method for signature 'RasterStackBrick,character'
writeRaster(x, filename, format, bylayer, suffix='numbers', ...)
```

## Arguments

x	Raster* object
filename	Output filename
format	Character. Output file type. See <a href="#">writeFormats</a> . If this argument is not provided, it is attempted to infer it from the filename extension. If that fails, the default format is used. The default format is 'raster', but this can be changed using <a href="#">rasterOptions</a>
...	Additional arguments: datatypeCharacter. Output data type (e.g. 'INT2S' or 'FLT4S'). See <a href="#">dataType</a> . If no datatype is specified, 'FLT4S' is used, unless this default value was changed with <a href="#">rasterOptions</a> overwrite: Logical. If TRUE, "filename" will be overwritten if it exists progress: Character. Set a value to show a progress bar. Valid values are "text" and "window". NAflag: Numeric. To overwrite the default value used to represent NA in a file bandorder: Character. 'BIL', 'BIP', or 'BSQ'. For 'native' file formats only. For some other formats you can use the 'options' argument (see below) options: Character. File format specific GDAL options. E.g., when writing a geotiff file you can use: options=c("COMPRESS=NONE", "TFW=YES") You can use options=c("PROFILE=BASELINE") to create a plain tif with no GeoTIFF tags. This can be useful when writing files to be read by applications intolerant of unrecognized tags. (see <a href="http://www.gdal.org/frmt_gtiff.html">http://www.gdal.org/frmt_gtiff.html</a> ) NetCDF files have the following additional, optional, arguments: varname, varunit, longname, xname, yname, zname, zunit prj: Logical. If TRUE, the crs is written to a .prj file. This can be useful when writing to an ascii file or another file type that does not store the crs
bylayer	if TRUE, write a separate file for each layer. You can provide a vector of filenames that matches the number of layers. Or you can provide a single filename that will get a unique suffix (see below)
suffix	'numbers' or 'names' to determine the suffix that each file gets when bylayer=TRUE; either a number between 1 and nlayers(x) or names(x)

## Details

See writeFormats for supported file types ("formats", "drivers").

The rgdal package is needed, except for these file formats: 'raster', 'BIL', 'BIP', 'BSQ', 'SAGA', 'ascii', 'IDRISI', and 'CDF'. Some of these formats can be used with or without rgdal (idrisi, SAGA, ascii). You need the 'ncdf4' library for the 'CDF' format.

In multi-layer files (i.e. files saved from RasterStack or RasterBrick objects), in the native 'raster' format, the band-order can be set to BIL ('Bands Interleaved by Line'), BIP ('Bands Interleaved by Pixels') or BSQ ('Bands SeQuential'). Note that bandorder is not the same as filetype here.

Supported file types include:

File type	Long name	default extension	Multiband support
raster	'Native' raster package format	.grd	Yes
ascii	ESRI Ascii	.asc	No
SAGA	SAGA GIS	.sdatt	No
IDRISI	IDRISI	.rst	No
CDF	netCDF (requires ncdf4)	.nc	Yes
GTiff	GeoTiff (requires rgdal)	.tif	Yes
ENVI	ENVI .hdr Labelled	.envi	Yes
EHdr	ESRI .hdr Labelled	.bil	Yes
HFA	Erdas Imagine Images (.img)	.img	Yes

## Value

This function is used for the side-effect of writing values to a file.

## See Also

[writeFormats](#), [writeValues](#)

## Examples

```
tmp <- tempdir()
r <- raster(system.file("external/test.grd", package="raster"))

# take a small part
r <- crop(r, extent(179880, 180800, 329880, 330840) )

# write to an integer binary file
rf <- writeRaster(r, filename=file.path(tmp, "allint.grd"), datatype='INT4S', overwrite=TRUE)

# make a brick and save multi-layer file
b <- brick(r, sqrt(r))
bf <- writeRaster(b, filename=file.path(tmp, "multi.grd"), bandorder='BIL', overwrite=TRUE)

# write to a new geotiff file (depends on rgdal)
if (require(rgdal)) {
  rf <- writeRaster(r, filename=file.path(tmp, "test.tif"), format="GTiff", overwrite=TRUE)
}
```



```

    bf <- writeRaster(b, filename=file.path(tmp, "multi.tif"),
options="INTERLEAVE=BAND", overwrite=TRUE)
}

# write to netcdf
if (require(ncdf4)) {
  rnc <- writeRaster(r, filename=file.path(tmp, "netCDF.nc"), format="CDF", overwrite=TRUE)
}

```

writeValues

*Write values to a file***Description**

Functions for writing blocks ( $\geq 1$  row(s)) of values to files. Writing has to start at the first cell of a row (identified with argument `start`) and the values written must represent 1 or more entire rows. Begin by opening a file with `writeStart`, then write values to it in chunks. When writing is done close the file with `writeStop`.

If you want to write all values of a `Raster*` object at once, you can also use `writeRaster` which is easier to use but more limited. The functions described here allow writing values to file using chunks of different sizes (e.g. 1 or 10 rows). Function `blockSize` can be used to suggest a chunk size to use.

**Usage**

```

## S4 method for signature 'RasterLayer,character'
writeStart(x, filename, options=NULL, format, prj=FALSE, ...)
## S4 method for signature 'RasterBrick,character'
writeStart(x, filename, options=NULL, format, prj=FALSE, ...)
## S4 method for signature 'RasterLayer,vector'
writeValues(x, v, start, ...)
## S4 method for signature 'RasterBrick,matrix'
writeValues(x, v, start, ...)
## S4 method for signature 'RasterLayer'
writeStop(x)
## S4 method for signature 'RasterBrick'
writeStop(x)

```

**Arguments**

<code>x</code>	<code>Raster*</code> object
<code>filename</code>	character. Output file name
<code>options</code>	character, see <code>writeRaster</code>
<code>format</code>	character, see <code>writeRaster</code>
<code>prj</code>	logical. If TRUE, a "prj" file is written
<code>...</code>	additional arguments as for <code>writeRaster</code>
<code>v</code>	vector ( <code>RasterLayer</code> ) or matrix ( <code>RasterBrick</code> ) of values
<code>start</code>	Integer. Row number (counting starts at 1) from where to start writing <code>v</code>

**Value**

RasterLayer or RasterBrick

**See Also**

[writeRaster](#), [blockSize](#), [update](#)

**Examples**

```
## Not run:
r <- raster(system.file("external/test.grd", package="raster"))
# write to a new binary file in chunks
s <- raster(r)
#
tr <- blockSize(r)
tr
s <- writeStart(s, filename='test.grd', overwrite=TRUE)
for (i in 1:tr$n) {
v <- getValuesBlock(r, row=tr$row[i], nrows=tr$nrows[i])
s <- writeValues(s, v, tr$row[i])
}
s <- writeStop(s)

if(require(rgdal)){
s2 <- writeStart(s, filename='test2.tif', format='GTiff', overwrite=TRUE)
# writing last row first
for (i in tr$n:1) {
v <- getValuesBlock(r, row=tr$row[i], nrows=tr$nrows[i])
s2 <- writeValues(s2, v, tr$row[i])
}
# row number 5 once more
v <- getValuesBlock(r, row=5, nrows=1)
writeValues(s2, v, 5)
s2 <- writeStop(s2)
}

## write values of a RasterStack to a RasterBrick
s <- stack(system.file("external/rlogo.grd", package="raster"))
# create empty brick
b <- brick(s, values=FALSE)
b <- writeStart(b, filename="test.grd", format="raster",overwrite=TRUE)
tr <- blockSize(b)
for (i in 1:tr$n) {
v <- getValuesBlock(s, row=tr$row[i], nrows=tr$nrows[i])
b <- writeValues(b, v, tr$row[i])
}
b <- writeStop(b)
# note that the above is equivalent to
# b <- writeRaster(s, filename="test.grd", format="raster",overwrite=TRUE)
```

```
## End(Not run)
```

---

xyFromCell	<i>Coordinates from a row, column or cell number</i>
------------	--

---

### Description

These functions get coordinates of the center of raster cells for a row, column, or cell number of a Raster\* object.

### Usage

```
## S4 method for signature 'Raster,numeric'
xFromCol(object, col)
## S4 method for signature 'Raster,numeric'
yFromRow(object, row)
## S4 method for signature 'Raster,numeric'
xFromCell(object, cell)
## S4 method for signature 'Raster,numeric'
yFromCell(object, cell)
## S4 method for signature 'BasicRaster,ANY'
xyFromCell(object, cell, spatial=FALSE, ...)
## S4 method for signature 'Raster'
coordinates(obj, ...)
## S4 method for signature 'Extent'
coordinates(obj, ...)
```

### Arguments

object	Raster* object (or a SpatialPixels* or SpatialGrid* object)
col	column number; or vector of column numbers. If missing, the x coordinates for all columns are returned
row	row number; or vector of row numbers. If missing, the y coordinates for all rows are returned
cell	cell number(s)
spatial	If spatial=TRUE, xyFromCell returns a SpatialPoints object instead of a matrix
...	additional arguments. None implemented
obj	Raster object

### Details

Cell numbers start at 1 in the upper left corner, and increase from left to right, and then from top to bottom. The last cell number equals the number of cells of the Raster\* object.

**Value**

xFromCol, yFromCol, xFromCell, yFromCell: vector of x or y coordinates  
 xyFromCell: matrix(x,y) with coordinate pairs  
 coordinates: xy coordinates for all cells

**See Also**

[cellFromXY](#)

**Examples**

```
#using a new default raster (1 degree global)
r <- raster()
xFromCol(r, c(1, 120, 180))
yFromRow(r, 90)
xyFromCell(r, 10000)
xyFromCell(r, c(0, 1, 32581, ncell(r), ncell(r)+1))

#using a file from disk
r <- raster(system.file("external/test.grd", package="raster"))
r
cellFromXY(r, c(180000, 330000))
#xy for corners of a raster:
xyFromCell(r, c(1, ncol(r), ncell(r)-ncol(r)+1, ncell(r)))
```

---

z-values

*Get or set z-values*


---

**Description**

Initial functions for a somewhat more formal approach to get or set z values (e.g. time) associated with layers of Raster\* objects. In development.

**Usage**

```
setZ(x, z, name='time')
getZ(x)
```

**Arguments**

x	Raster* object
z	vector of z values of any type (e.g. of class 'Date')
name	character label

**Value**

setZ: Raster\* object  
 getZ: vector

**Examples**

```

r <- raster(ncol=10, nrow=10)
s <- stack(lapply(1:3, function(x) setValues(r, runif(ncell(r)))))
s <- setZ(s, as.Date('2000-1-1') + 0:2)
s
getZ(s)

```

---

zApply

*z (time) apply*


---

**Description**

Experimental function to apply a function over a (time) series of layers of a Raster object

**Usage**

```
zApply(x, by, fun=mean, name='', ...)
```

**Arguments**

x	Raster* object
by	aggregation indices or function
fun	function to compute aggregated values
name	character label of the new time series
...	additional arguments

**Value**

Raster\* object

**Author(s)**

Oscar Perpinan Lamigueiro & Robert J. Hijmans

**Examples**

```

# 12 values of irradiation, 1 for each month
G0dm=c(2.766,3.491,4.494,5.912,6.989,7.742,7.919,7.027,5.369,3.562,2.814,2.179)*1000;
# RasterBrick with 12 layers based on G0dm + noise
r <- raster(nc=10, nr=10)
s <- brick(lapply(1:12, function(x) setValues(r, G0dm[x]+100*rnorm(ncell(r)) )))

# time
tm <- seq(as.Date('2010-01-15'), as.Date('2010-12-15'), 'month')
s <- setZ(s, tm, 'months')

# library(zoo)
# x <- zApply(s, by=as.yearqtr, fun=mean, name='quarters')

```

zonal

*Zonal statistics***Description**

Compute zonal statistics, that is summarized values of a Raster\* object for each "zone" defined by a RasterLayer.

If `stat` is a true function, `zonal` will fail (gracefully) for very large Raster objects, but it will in most cases work for functions that can be defined as by a character argument ('mean', 'sd', 'min', 'max', or 'sum'). In addition you can use 'count' to count the number of cells in each zone (only useful with `na.rm=TRUE`, otherwise `freq(z)` would be more direct.

If a function is used, it should accept a `na.rm` argument (or at least a ... argument)

**Usage**

```
## S4 method for signature 'RasterLayer,RasterLayer'
zonal(x, z, fun='mean', digits=0, na.rm=TRUE, ...)
```

```
## S4 method for signature 'RasterStackBrick,RasterLayer'
zonal(x, z, fun='mean', digits=0, na.rm=TRUE, ...)
```

**Arguments**

<code>x</code>	Raster* object
<code>z</code>	RasterLayer with codes representing zones
<code>fun</code>	function to be applied to summarize the values by zone. Either as character: 'mean', 'sd', 'min', 'max', 'sum'; or, for relatively small Raster* objects, a proper function
<code>digits</code>	integer. Number of digits to maintain in 'zones'. By default averaged to an integer (zero digits)
<code>na.rm</code>	logical. If TRUE, NA values in <code>x</code> are ignored
<code>...</code>	additional arguments. One implemented: <code>progress</code> , as in <a href="#">writeRaster</a>

**Value**

A matrix with a value for each zone (unique value in zones)

**See Also**

See [cellStats](#) for 'global' statistics (i.e., all of `x` is considered a single zone), and [extract](#) for summarizing values for polygons

**Examples**

```

r <- raster(ncols=10, nrows=10)
values(r) <- runif(ncell(r)) * 1:ncell(r)
z <- r
values(z) <- rep(1:5, each=20)
# for large files, use a character value rather than a function
zonal(r, z, 'sum')

# for smaller files you can also provide a function
## Not run:
zonal(r, z, mean)
zonal(r, z, min)

## End(Not run)

# multiple layers
zonal(stack(r, r*10), z, 'sum')
```

---

zoom

*Zoom in on a map*


---

**Description**

Zoom in on a map (plot) by providing a new extent, by default this is done by clicking twice on the map.

**Usage**

```

zoom(x, ...)
## S4 method for signature 'Raster'
zoom(x, ext=drawExtent(), maxpixels=100000, layer=1, new=TRUE, useRaster=TRUE, ...)

## S4 method for signature 'Spatial'
zoom(x, ext=drawExtent(), new=TRUE, ...)

## S4 method for signature 'missing'
zoom(x, ext=drawExtent(), new=TRUE, ...)
```

**Arguments**

x	Raster* or Spatial* (vector type) object
ext	Extent object, or other object from which an extent can be extracted
maxpixels	Maximum number of pixels used for the map
layer	Positive integer to select the layer to be used if x is a multilayer Raster object
new	Logical. If TRUE, the zoomed in map will appear on a new device (window)
useRaster	Logical. If TRUE, a bitmap raster is used to plot the image instead of polygons
...	additional parameters for <a href="#">plot</a>

**Value**

Extent object (invisibly)

**See Also**

[drawExtent](#), [plot](#)



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