

# Package ‘magclass’

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**Type** Package

**Title** Data Class and Tools for Handling Spatial-Temporal Data

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**Description** Data class for increased interoperability working with spatial-temporal data together with corresponding functions and methods (conversions, basic calculations and basic data manipulation). The class distinguishes between spatial, temporal and other dimensions to facilitate the development and interoperability of tools build for it. Additional features are name-based addressing of data and internal consistency checks (e.g. checking for the right data order in calculations).

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**BugReports** <https://github.com/pik-piam/magclass/issues>

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magclass-package	<i>Data Class and Tools for Handling Spatial-Temporal Data</i>
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## Description

Data class for increased interoperability working with spatial-temporal data together with corresponding functions and methods (conversions, basic calculations and basic data manipulation). The class distinguishes between spatial, temporal and other dimensions to facilitate the development and interoperability of tools build for it. Additional features are name-based addressing of data and internal consistency checks (e.g. checking for the right data order in calculations).

## Author(s)

Maintainer: Jan Philipp Dietrich <dietrich@pik-potsdam.de>

## See Also

Useful links:

- <https://github.com/pik-piam/magclass>
- doi: [10.5281/zenodo.1158580](https://doi.org/10.5281/zenodo.1158580)
- Report bugs at <https://github.com/pik-piam/magclass/issues>

---

add_columns	<i>add_columns</i>
-------------	--------------------

---

## Description

Function adds new columns to the existing magpie object.

## Usage

```
add_columns(x, addnm = "new", dim = 3.1, fill = NA)
```

**Arguments**

x	MAGPIE object which should be extended.
addnm	The new elements that should be added to the (sub)dimension
dim	The (sub)dimension to be filled either identified via name or dimension code (see <a href="#">dimCode</a> for more information)
fill	fill value of length 1 for the newly added columns (NA by default)

**Value**

The extended MAGPIE object

**Author(s)**

Jan Philipp Dietrich, Benjamin Bodirsky

**See Also**

[add\\_dimension](#), [dimCode](#)

**Examples**

```
a <- maxample("animal")
a2 <- add_columns(a, addnm = c("horse", "magpie"), dim = "species", fill = 42)
getItems(a2, dim = 3)
getItems(a2, dim = 3, split = TRUE)
head(a2[, , "magpie"])
```

---

add_dimension	<i>add_dimension</i>
---------------	----------------------

---

**Description**

Function adds a name dimension as dimension number "dim" with the name "add" with an empty data column with the name "nm".

**Usage**

```
add_dimension(x, dim = 3.1, add = NULL, nm = "dummy")
```

**Arguments**

x	MAGPIE object which should be extended.
dim	The dimension number of the new dimension (e.g. 3.1)
add	The name of the new dimension
nm	The name of the first entry in dimension "add".

**Value**

The extended MAgPIE object

**Author(s)**

Jan Philipp Dietrich, Benjamin Bodirsky

**See Also**

[add\\_columns](#), [mbind](#)

**Examples**

```
a <- maxample("animal")
str(add_dimension(a, dim = 3.2))
str(add_dimension(a, dim = 2.3, nm = paste0("d", 1:3)))
```

---

as.array-methods

~~ *Methods for Function as.array* ~~

---

**Description**

~~ Methods for function as.array ~~

**Usage**

```
## S4 method for signature 'magpie'
as.array(x)
```

**Arguments**

x                    object which should be converted to an array

**Methods**

**list("signature(x = \"ANY\")")** standard as.array-method

**list("signature(x = \"magpie\")")** Conversion takes place just by removing MAgPIE-object specific elements

---

as.data.frame-methods *~~ Methods for Function as.data.frame ~~*

---

## Description

~~ Methods for function as.data.frame ~~

## Usage

```
## S4 method for signature 'magpie'
as.data.frame(x, rev = 1, raw = FALSE)
```

## Arguments

x	A MAgPIE-object
rev	The revision of the algorithm that should be used for conversion. rev=1 creates columns with the predefined names Cell, Region, Year, Data1, Data2,... and Value, rev=2 uses the set names of the MAgPIE object for naming and adds an attribute "dimtype" to the data.frame which contains information about the types of the different columns (spatial, temporal, data or value), rev=3 is identical to rev=2 except that characters are not being converted to factors (stringsAsFactors = FALSE).
raw	Logical to control whether years beginning with "y" should be converted to integers (without "y") and coordinates should be converted to numerics. If set to raw columns are returned as they are in the initial object.

## Methods

**list("signature(x = \"magpie\")")** Conversion creates columns for Cell, Region, Year, Data1, Data2,... and Value

## Examples

```
pop <- maxample("pop")
head(as.data.frame(pop))
head(as.data.frame(pop, rev = 2))

a <- maxample("animal")
head(as.data.frame(a, rev = 3))
head(as.data.frame(a, rev = 3, raw = TRUE))
attr(as.data.frame(a, rev = 3), "dimtype")
```

---

as.RasterBrick	<i>as.RasterBrick</i>
----------------	-----------------------

---

## Description

Convert magclass object to a RasterBrick object

## Usage

```
as.RasterBrick(x, res = NULL)
```

## Arguments

x	MAGPIE object
res	spatial data resolution. If not provided it will be guessed.

## Value

A RasterBrick object

## Author(s)

Jan Philipp Dietrich

## See Also

[getCoords](#)

## Examples

```
## Not run:
if (requireNamespace("raster", quietly = TRUE)) {
  r <- raster::brick(ncols = 36, nrows = 18, nl = 4)
  r[14:18, 25:28] <- (1:20 %% t(1:4))
  names(r) <- c("y2000..bla", "y2001..bla", "y2000..blub", "y2001..blub")
  m <- as.magpie(r)
  r2 <- as.RasterBrick(m)
}

## End(Not run)
```



---

as.SpatRaster	<i>as.SpatRaster</i>
---------------	----------------------

---

## Description

Convert magclass object to a SpatRaster object. Requires the terra package.

## Usage

```
as.SpatRaster(x, res = NULL)
```

## Arguments

x	MAGPIE object
res	spatial data resolution. If not provided it will be guessed.

## Value

A SpatRaster object

## Author(s)

Jan Philipp Dietrich

## See Also

[getCoords](#)

## Examples

```
if (requireNamespace("terra", quietly = TRUE)) {  
  r <- terra::rast(ncols = 360, nrows = 180, nl = 4)  
  r[85:89, 176:179] <- (1:20 %*% t(1:4))  
  r[15:19, 76:79] <- (10 + 1:20 %*% t(1:4))  
  names(r) <- c("y2000..bla", "y2001..bla", "y2000..blub", "y2001..blub")  
  m <- as.magpie(r)  
  r2 <- as.SpatRaster(m)  
}
```

---

as.SpatRasterDataset    *as.SpatRasterDataset*

---

**Description**

Convert magclass object to a SpatRasterDataset object. Requires the terra package.

**Usage**

```
as.SpatRasterDataset(...)
```

**Arguments**

...                    arguments passed to as.SpatRaster

**Details**

Calls [as.SpatRaster](#) and then [spatRasterToDataset](#).

**Value**

A SpatRasterDataset object

**Author(s)**

Pascal Sauer

---

as.SpatVector            *as.SpatVector*

---

**Description**

Convert magclass object to a SpatVector object. Requires the terra package and requires the mag-class object to provide the geometry of the spatial entities as "geometry" attribute in "WKT" format. (see object "m" in example).

**Usage**

```
as.SpatVector(x)
```

**Arguments**

x                      MAgPIE object

**Value**

A SpatVector object

**Author(s)**

Jan Philipp Dietrich

**See Also**[as.SpatRaster](#)**Examples**

```

if (requireNamespace("terra", quietly = TRUE)) {
  r <- terra::rast(ncols = 360, nrows = 180, nl = 4)
  r[85:89, 176:179] <- (1:20 %*% t(1:4))
  r[15:19, 76:79] <- (10 + 1:20 %*% t(1:4))
  names(r) <- c("y2000..bla", "y2001..bla", "y2000..blub", "y2001..blub")
  v <- terra::as.polygons(r)
  m <- as.magpie(v)
  attr(m, "geometry")
  attr(m, "crs")
  v2 <- as.SpatVector(m)
}

```

as\_tibble.magpie

*magpie method for tibble::as\_tibble***Description**

magpie method for tibble::as\_tibble

**Usage**

```

as_tibble.magpie(
  x,
  ...,
  .rows = NULL,
  .name_repair = c("check_unique", "unique", "universal", "minimal"),
  rownames = pkgconfig::get_config("tibble::rownames", NULL)
)

```

**Arguments**

x	A magpie object
...	Unused, for extensibility.
.rows	The number of rows, useful to create a 0-column tibble or just as an additional check.
.name_repair	Treatment of problematic column names: <ul style="list-style-type: none"> <li>• "minimal": No name repair or checks, beyond basic existence,</li> </ul>

- "unique": Make sure names are unique and not empty,
- "check\_unique": (default value), no name repair, but check they are unique,
- "universal": Make the names unique and syntactic
- a function: apply custom name repair (e.g., `.name_repair = make.names` for names in the style of base R).
- A purrr-style anonymous function, see `rlang::as_function()`

This argument is passed on as `repair` to `vctrs::vec_as_names()`. See there for more details on these terms and the strategies used to enforce them.

rownames

How to treat existing row names of a data frame or matrix:

- NULL: remove row names. This is the default.
- NA: keep row names.
- A string: the name of a new column. Existing rownames are transferred into this column and the `row.names` attribute is deleted. Read more in [rownames](#).

## Value

A tibble object

---

clean\_magpie

*MAGPIE-Clean*

---

## Description

Function cleans MAGPIE objects so that they follow some extended magpie object rules (currently it makes sure that the dimnames have names and removes cell numbers if it is purely regional data)

## Usage

```
clean_magpie(x, what = "all", maindim = 1:3)
```

## Arguments

x	MAGPIE object which should be cleaned.
what	term defining what type of cleaning should be performed. Current modes are "cells" (removes cell numbers if the data seems to be regional - this should be used carefully as it might remove cell numbers in some cases in which they should not be removed), "sets" (making sure that all dimensions have names), "items" (replace empty elements with single spaces " ") and "all" (performing all available cleaning methods)
maindim	main dimension(s) the cleaning should get applied to.

## Value

The eventually corrected MAGPIE object

**Author(s)**

Jan Philipp Dietrich

**See Also**["magpie"](#)**Examples**

```
pop <- maxample("pop")
a <- clean_magpie(pop)
```

collapseDim

*Collapse dataset dimensions***Description**

This function will remove names in the data dimension which are the same for each element (meaning that this data dimension contains exactly one element) or, if forced, remove any other subdimension. It is a generalized version of the function [collapseNames](#)

**Usage**

```
collapseDim(x, dim = NULL, keepdim = NULL)
```

**Arguments**

x	MAGPIE object
dim	Either NULL, dimension code or name of dimension or a vector of these. If set to NULL all single entry subdimensions will be removed as they are irrelevant to uniquely identify a data element. If specified, only the specified subdimensions will be removed (See <a href="#">dimCode</a> for more details how to specify a subdimension). CAUTION: The function also allows to specify subdimensions which are otherwise needed to clearly identify an entry. By removing these subdimensions duplicates in the data will be created potentially causing problems in the further use of the data set. Be careful in removing subdimensions.
keepdim	(only considered if dim is not specified) Can be used to converse single element subdimension which otherwise would get deleted. If dim is specified this setting will not have any effect.

**Value**

The provided MAGPIE object with collapsed dimensions

**Note**

This function has some similarities to [dimReduce](#), but serves a different purpose. While [collapseDim](#) only removes dimensions which contain only a single element or which it is specifically told to remove, [dimReduce](#) looks whether the entries of a multi-entry dimension are all the same and removes dimensions for which this is the case. In some cases both will lead to the same result but in many other cases the results will differ.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getItems](#) "magpie"

**Examples**

```
x <- new.magpie(c("GLO.1", "GLO.2"), 2000, c("bla.a", "bla.b"))
collapseDim(x)
collapseDim(x, keepdim = 1:2)
collapseDim(x, dim = 1.1)
collapseDim(x, dim = 3.2)
```

---

collapseNames

*Collapse dataset names*


---

**Description**

This function has been superseded by [collapseDim](#) which is a more generalized version of this function. Please use this one instead!

**Usage**

```
collapseNames(x, collapsedim = NULL, preservedim = NULL)
```

**Arguments**

x	MAGPIE object
collapsedim	If you want to remove the names of particular dimensions provide the dimensions here. Since the function only works in the third dimension, you have to count from there on (e.g. dim = 3.2 refers to collapsedim = 2). Alternatively, you can also specify the name of the dimension. Default: NULL. CAUTION with parameter collapsedim! You could also force him to remove dimnames, which are NOT the same for each element and so create duplicates in dimnames.
preservedim	If you want to remove the name of particular dimensions except some, you can specify the dimension(s) to preserve here. See collapsedim for naming convention. Note that preservedim will be ignored in the case, of a specified collapsedim

**Details**

This function will remove names in the data dimension which are the same for each element (meaning that this data dimension contains exactly one element)

**Value**

The provided MAgPIE object with collapsed names

**Author(s)**

Jan Philipp Dietrich, David Klein, Xiaoxi Wang

**See Also**

[collapseDim](#), [getItems](#), "magpie"

---

colSums-methods      *~~ Methods for Function colSums and colMeans ~~*

---

**Description**

*~~ Methods for function colSums and colMeans ~~*

**Usage**

```
## S4 method for signature 'magpie'
colSums(x, na.rm = FALSE, dims = 1, ...)
```

**Arguments**

x	object on which calculation should be performed
na.rm	logical. Should missing values (including NaN) be omitted from the calculations?
dims	integer: Which dimensions are regarded as "rows" or "columns" to sum over. For row*, the sum or mean is over dimensions dims+1, ...; for col* it is over dimensions 1:dims.
...	further arguments passed to other colSums/colMeans methods

**Methods**

**list("signature(x = \"ANY\")")** normal colSums and colMeans method

**list("signature(x = \"magpie\")")** classical method prepared to handle MAgPIE objects

complete\_magpie      *complete\_magpie*

---

### Description

MAGPIE objects can be incomplete to reduce memory. This function blows up a magpie object to its real dimensions, so you can apply unwrap.

### Usage

```
complete_magpie(x, fill = NA, dim = 3)
```

### Arguments

x	MAGPIE object which should be completed.
fill	Value that shall be written into the missing entries
dim	dimensions in which the completion should take place (1, 2 and/or 3). For full completion use 1:3

### Value

The completed MAGPIE object

### Author(s)

Jan Philipp Dietrich, Benjamin Bodirsky

### See Also

[add\\_dimension](#), [clean\\_magpie](#)

### Examples

```
pop <- maxample("pop")
complete_magpie(pop)

ani <- maxample("animal")
complete_magpie(ani)
```



---

convergence	<i>convergence</i>
-------------	--------------------

---

### Description

Cross-Fades the values of one MAGPIE object into the values of another over a certain time

### Usage

```
convergence(
  origin,
  aim,
  start_year = NULL,
  end_year = NULL,
  direction = NULL,
  type = "smooth",
  par = 1.5
)
```

### Arguments

origin	an object with one name-column
aim	Can be twofold: An magpie object or a numeric value.
start_year	year in which the convergence from origin to aim starts. If set to NULL the the first year of aim is used as start_year
end_year	year in which the convergence from origin to aim shall be (nearly) reached. If set to NULL the the last year of aim is used as end_year.
direction	NULL, "up" or "down". NULL means normal convergence in both directions, "up" is only a convergence if origin<aim, "down" means only a convergence if origin>aim
type	"smooth", "s", "linear" or "decay". Describes the type of convergence: linear means a linear conversion , s is an s-curve which starts from origin in start_year and reaches aim precisely in end_year. After 50 percent of the convergence time, it reaches about the middle of the two values. Its based on the function $\min(1, \text{pos}^4 / (0.07 + \text{pos}^4) * 1.07)$ smooth is a conversion based on the function $x^3 / (0.1 + x^3)$ . In the latter case only 90% of convergence will be reached in the end year, because full convergence is reached in infinity. decay is a conversion based on the function $x / (1.5 + x) * 2.5$ .
par	parameter value for convergence function; currently only used for type="decay"

### Value

returns a time-series with the same timesteps as origin, which lineary fades into the values of the aim object

**Author(s)**

Benjamin Bodirsky, Jan Philipp Dietrich

**Examples**

```
pop <- maxample("pop")
population <- add_columns(pop, "MIX")
population[, , "MIX"] <- convergence(population[, , "A2"], population[, , "B1"])
```

---

copy.attributes

*Copy Attributes*

---

**Description**

This function copies attributes from one object and assigns them to another.

**Usage**

```
copy.attributes(
  from,
  to,
  delete = c("names", "row.names", "class", "dim", "dimnames"),
  delete2 = NULL
)
```

```
copy.attributes(
  to,
  delete = c("names", "row.names", "class", "dim", "dimnames"),
  delete2 = NULL
) <- value
```

**Arguments**

from	object from which the attributes should be taken
to	object to which the attributes should be written
delete	attributes which should not be copied. By default this are class specific attributes which might cause problems if copied to another object. But you can add or remove attributes from the vector.
delete2	Identical to delete and just added for convenience for the case that you want to delete additional attributes but do not want to repeat the vector given in delete. In the function both vectors, delete and delete2, are just merged to one deletion vector.
value	Same as "from" (object from which the attributes should be taken)

**Functions**

- `copy.attributes( to, delete = c("names", "row.names", "class", "dim", "dimnames"), delete2 = NULL )` <- value: assign attributes from object "value"

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
from <- array(12)
attr(from,"blablub") <- "I am an attribute!"
attr(from,"blablub2") <- "I am another attribute!"

print(attributes(from))

to <- as.magpie(0)
print(attributes(to))

copy.attributes(to) <- from
print(attributes(to))
```

---

copy.magpie

*Copy MAgPIE-files*

---

**Description**

This function copies MAgPIE-files from one location to another. During the copying it is also possible to change the file type (e.g. from 'mz' to 'csv')

**Usage**

```
copy.magpie(input_file, output_file, round = NULL)
```

**Arguments**

<code>input_file</code>	file, that should be copied
<code>output_file</code>	copy destination
<code>round</code>	number of digits the values should be rounded. NULL means no rounding

**Author(s)**

Jan Philipp Dietrich

**See Also**

[read.magpie](#), [write.magpie](#)

**Examples**

```
# copy.magpie("bla.csv", "blub.mz")
```

---

dimCode	<i>dimCode</i>
---------	----------------

---

**Description**

Function converts a dimension name or number to a dimension Code used for MAgPIE objects

**Usage**

```
dimCode(dim, x, missing = 0, strict = FALSE, sep = ".")
```

**Arguments**

dim	A vector of dimension numbers or dimension names which should be translated
x	MAgPIE object in which the dimensions should be searched for.
missing	Either a value to which a dimension should be set in case that it is not found (default is 0), or "stop" indicating that the function should throw an error in these cases.
strict	if set to TRUE also properly set dimension names which refer to non-existing subdimensions will be treated as missing, otherwise these dimension codes will be returned, even if the subdimension does not exist
sep	A character separating joined dimension names

**Value**

A dimension code identifying the dimension. Either a integer which represents the main dimensions (1=spatial, 2=temporal, 3=data) or a numeric, representing the subdimensions of a dimension (e.g. 3.2 for the second data dimension).

**Author(s)**

Jan Philipp Dietrich, Kristine Karstens

**See Also**

[mselect](#), [getDim](#)

**Examples**

```
pop <- maxample("pop")
dimCode(c("t", "scenario", "blablub"), pop)
```

---

dimExists	<i>dimExists</i>
-----------	------------------

---

**Description**

Function checks whether a dimension exist in a MAgPIE objects

**Usage**

```
dimExists(dim, x, sep = ".")
```

**Arguments**

dim	A vector of dimension numbers or dimension names which should be checked for
x	MAgPIE object in which the dimensions should be searched for.
sep	A character separating joined dimension names

**Value**

Boolean indicating whether dimension exists or not

**Author(s)**

Jan Philipp Dietrich

**See Also**

[dimCode](#)

**Examples**

```
pop <- maxample("pop")
dimExists(c("t", "scenario", "blablub"), pop)
```

dimOrder

*dimOrder*

---

**Description**

Changes the order of the sub-dimension in a magpie object similar to unwrapping and applying the `aperm` command, but more efficient.

**Usage**

```
dimOrder(x, perm, dim = 3)
```

**Arguments**

<code>x</code>	magpie object
<code>perm</code>	vector with the new order of the sub-dimension. Missing sub-dimensions will be added automatically at the end
<code>dim</code>	main dimension in which the order of sub-dimensions should be changed (1, 2 or 3)

**Value**

magpie object

**Author(s)**

Jan Philipp Dietrich, Benjamin Leon Bodirsky

**Examples**

```
a <- maxample("animal")
head(a)
head(dimOrder(a, perm = 3:1, dim = 1))
head(dimOrder(a, perm = c(2,1,3), dim = 3))
```

---

dimReduce*dimReduce*

---

**Description**

Remove dimensions which contain identical data for all elements in it

**Usage**

```
dimReduce(x, dim_exclude = NULL)
```

**Arguments**

`x`                    MAgPIE object which should be reduced  
`dim_exclude`        Vector with names of dimensions which must not be reduced

**Value**

The reduced MAgPIE object

**Note**

This function has some similarities to [collapseDim](#), but serves a different purpose. While [collapseDim](#) only removes dimensions which contain only a single element or which it is specifically told to remove, [dimReduce](#) looks whether the entries of a multi-entry dimension are all the same and removes dimensions for which this is the case. In some cases both will lead to the same result but in many other cases the results will differ.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[add\\_dimension](#)

**Examples**

```
# create data with 5 identical scenarios
p <- add_dimension(maxample("pop")[1:3, 1:3, ], nm = paste0("scen", 1:2))
str(p)
str(dimReduce(p))

# set years to same value
p[, , ] <- setYears(p[, 1, ], NULL)
str(p)
str(dimReduce(p))

# set regions to same value
p[, , ] <- setCells(p[1, , ], "GLO")
str(p)
str(dimReduce(p))
```

---

dimSums	<i>Summation over dimensions</i>
---------	----------------------------------

---

**Description**

This function sums over any (sub-)dimension of a magpie object

**Usage**

```
dimSums(x, dim = 3, na.rm = FALSE)
```

**Arguments**

x	A MAGPIE-object
dim	The dimensions(s) to sum over. A vector of dimension codes or dimension names. See <a href="#">dimCode</a> for more information
na.rm	logical. Should missing values (including NaN) be omitted from the calculations?

**Value**

A MAGPIE object with values summed over the specified dimensions

**Author(s)**

Jan Philipp Dietrich

**See Also**

[rowSums](#), [getItems](#), [dimCode](#)

**Examples**

```
a <- maxample("animal")
dimSums(a, dim = c(1, 2, 3.2))
dimSums(a, dim = c("x", "y", "cell", "month"))
```



---

escapeRegex	<i>escapeRegex</i>
-------------	--------------------

---

**Description**

Escapes all symbols in a string which have a special meaning in regular expressions.

**Usage**

```
escapeRegex(x)
```

**Arguments**

x                      String or vector of strings that should be escaped.

**Value**

The escaped strings.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[grep](#)

---

fulldim	<i>Reconstructs full dimensionality of MAgPIE objects</i>
---------	---

---

**Description**

If a MAgPIE object is created from a source with more than one data dimension, these data dimensions are combined to a single dimension. `fulldim` reconstructs the original dimensionality and reports it.

**Usage**

```
fulldim(x, sep = ".")
```

**Arguments**

x                      A MAgPIE-object  
sep                    A character separating joined dimension names

**Value**

A list containing in the first element the dim output and in the second element the dimnames output of the reconstructed array.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[as.magpie](#), [unwrap](#), [wrap](#)

---

getCells

*Get Cells*

---

**Description**

Extracts cell names of a MAgPIE-object

**Usage**

```
getCells(x)
```

```
getCells(x) <- value
```

```
setCells(object, nm = "GLO")
```

**Arguments**

x, object	MAgPIE object
value, nm	cell names the data should be set to.

**Details**

setCells is a shortcut to use a MAgPIE object with manipulated cell names. setCells uses the variable names "object" and "nm" in order to be consistent to the already existing function setName.

**Value**

getCells returns cell names of the MAgPIE-object, whereas setCells returns the MAgPIE object with the manipulated cell names.

**Functions**

- `getCells(x) <- value`: set cell names
- `setCells()`: set cell names

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getRegions](#), [getNames](#), [setNames](#), [getCPR](#), [read.magpie](#), [write.magpie](#), ["magpie"](#)

**Examples**

```
a <- as.magpie(1)
getCells(a)
setCells(a, "AFR")
```

---

*getComment*

*getComment*

---

**Description**

Extracts the comment from a MAgPIE-object

**Usage**

```
getComment(x)

getComment(x) <- value

setComment(object, nm = NULL)
```

**Arguments**

<code>x, object</code>	MAgPIE object
<code>value, nm</code>	A vector containing the comment.

**Value**

`getComment` returns the comment attached to a MAgPIE-object, `NULL` if no comment is present.  
`setComment` returns the magpie object with the modified comment.

**Functions**

- `getComment(x) <- value`: set comment
- `setComment()`: set comment

**Author(s)**

Markus Bonsch

**See Also**

[getRegions](#), [getNames](#), [getYears](#), [getCPR](#), [read.magpie](#), [write.magpie](#), "magpie"

**Examples**

```
a <- as.magpie(1)
#returns NULL
getComment(a)
#set the comment
getComment(a)<-c("bla","blubb")
getComment(a)
```

---

getCoords

*Get Coordinates*

---

**Description**

Extracts spatial coordinates of a MAgPIE-object

**Usage**

```
getCoords(x, xlab = "x", ylab = "y")
```

```
getCoords(x, xlab = "x", ylab = "y") <- value
```

**Arguments**

x	MAgPIE object
xlab	label of x-dimension
ylab	label of y-dimension
value	coordinates as two column data.frame the data should be set to (first column = x, second column = y).

**Value**

coordinates of the MAgPIE-object

**Functions**

- `getCoords(x, xlab = "x", ylab = "y") <- value`: set coordinates

**Author(s)**

Jan Philipp Dietrich

**See Also**

[as.RasterBrick](#), [getItems](#), ["magpie"](#)

**Examples**

```
a <- maxample("animal")
getCoords(a)
```

---

getCPR

*Get cells per region*

---

**Description**

Counts how often each element of the provided subdimension exists in the given data set. Originally created to count the number of cells in a region (this is also where its name originates from) it can now be used to count elements of any subdimension via the `dim` argument.

**Usage**

```
getCPR(x, dim = 1.1)
```

**Arguments**

<code>x</code>	MAGPIE object or a resolution written as numeric (currently only data for 0.5 degree resolution is available).
<code>dim</code>	Dimension for which the items should be returned. Either number or name of dimension or a vector of these (in case of a vector all subdimensions must belong to the same main dimension!). See <a href="#">dimCode</a> for more details.

**Value**

cells per region

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getRegions](#), [read.magpie](#), [write.magpie](#)

**Examples**

```

getCPR(0.5)
a <- maxample("animal")
getCPR(a, dim = "color")
getCPR(a, dim = 3.2)
getCPR(a, dim = "country")
getCPR(a, dim = c("color", "species"))

```

---

getDim

*getDim*


---

**Description**

Function which tries to detect the dimension to which the given elems belong

**Usage**

```
getDim(elems, x, fullmatch = FALSE, dimCode = TRUE)
```

**Arguments**

elems	A vector of characters containing the elements that should be found in the MAg-PIE object
x	MAGPIE object in which elems should be searched for.
fullmatch	If enabled, only dimensions which match exactly the elements provided will be returned. Otherwise, it is sufficient if elems contains a subset of the dimension.
dimCode	If enabled, the dimCode will be returned, otherwise the name of the dimension.

**Value**

The name or dimCode of the dimensions in which elems were found.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[mcalc,dimCode](#)

**Examples**

```

pop <- maxample("pop")
getDim(c("AFR", "CPA"), pop)
getDim(c("AFR", "CPA"), pop, fullmatch=TRUE)
getDim(c("AFR", "CPA"), pop, dimCode=FALSE)

```

---

<code>getItems</code>	<i>Get Items</i>
-----------------------	------------------

---

**Description**

Extract items of a given (sub-)dimension of a MAgPIE-object

**Usage**

```
getItems(x, dim = NULL, split = FALSE, full = FALSE)
```

```
getItems(x, dim, full = NULL, maindim = NULL, raw = FALSE) <- value
```

**Arguments**

<code>x</code>	MAgPIE object
<code>dim</code>	Dimension for which the items should be returned. Either number or name of dimension or a vector of these. See <a href="#">dimCode</a> for more details.
<code>split</code>	Boolean which determines whether a main dimension should be split in subdimensions. Only applicable to main dimensions (1,2,3) and ignored for all other.
<code>full</code>	if TRUE dimension names are returned as they are (including repetitions), if FALSE only the dimension elements (unique list of entries) are returned.
<code>maindim</code>	main dimension the data should be added to (does not need to be set if <code>dim</code> exists in the data. Should be set if <code>dim</code> might not exist, or if <code>dim</code> might potentially exist in a different main dimension than the one anticipated).
<code>raw</code>	if set to FALSE inputs will be corrected (e.g. dots replaced by the letter "p") if necessary. If TRUE data will be written as is (risking the creation of inconsistent objects).
<code>value</code>	a vector with the length of the main dimension the dimnames should be replaced in / added to. If set to NULL the corresponding dimension will be removed.

**Value**

items of the requested dimension in the MAgPIE-object. If `split=TRUE` and applied to a main dimension (1,2,3) a list of items for each sub-dimension.

**Functions**

- `getItems(x, dim, full = NULL, maindim = NULL, raw = FALSE) <- value`: set dimension names

**Author(s)**

Jan Philipp Dietrich

**See Also**

[dimCode](#)

**Examples**

```
x <- maxample("pop")
getItems(x, "scenario")
getItems(x, 3.1)
getItems(x, "i") <- paste0("REG", seq_len(ncells(x)))
getItems(x, "i")
y <- x[, 1, ]
getItems(y, "t") <- NULL
```

---

getNames

*Get dataset names*


---

**Description**

Extracts dataset names of a MAgPIE-object

**Usage**

```
getNames(x, fulldim = FALSE, dim = NULL)
```

```
getNames(x, dim = NULL) <- value
```

**Arguments**

x	MAgPIE object
fulldim	specifies, how the object is treated. In case of FALSE, it is assumed that x is 3 dimensional and dimnames(x)[[3]] is returned. In case of TRUE, the dimnames of the real third dimension names are returned
dim	Argument to choose a specific data dimension either by name of the dimension or by number of the data dimension.
value	a vector of names current names should be replaced with. If only one data element exists you can also set the name to NULL.

**Details**

setNames is a shortcut to use a MAgPIE object with manipulated data names. The setNames method uses the variable names "object" and "nm" in order to be consistent to the already existing function setNames.

**Value**

getNames returns data names of the MAgPIE-object, whereas setNames returns the MAgPIE object with the manipulated data names.

**Functions**

- getNames(x, dim = NULL) <- value: set names



**Author(s)**

Jan Philipp Dietrich

**See Also**

[setNames-methods](#), [getRegions](#), [getYears](#), [getCPR](#), [read.magpie](#), [write.magpie](#), [ndata](#), ["magpie"](#)

**Examples**

```
a <- as.magpie(1)
getNames(a)
setNames(a, "bla")

x <- new.magpie("GLO", 2000, c("a.o1", "b.o1", "a.o2"))
getNames(x, dim = 2)

getSets(x, fulldim = FALSE)[3] <- "bla.blub"
getNames(x, dim = "bla")

getSets(x)[4] <- "ble"
getNames(x, dim = "ble") <- c("Hi", "Bye")
x
```

---

getRegionList

*Get a list of cellulare region-belongings*

---

**Description**

Extracts a vector containing the region of each cell of a MAgPIE-object

**Usage**

```
getRegionList(x)
```

```
getRegionList(x) <- value
```

**Arguments**

x	MAgPIE object
value	A vector with ncell elements containing the regions of each cell.

**Value**

A vector with ncell elements containing the region of each cell.

**Functions**

- `getRegionList(x) <- value`: set region names

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getRegions](#), [getYears](#), [getNames](#), [getCPR](#), [read.magpie](#), [write.magpie](#), "magpie"

**Examples**

```
# a <- read.magpie("example.mz")
# getRegionList(a)
```

---

getRegions	<i>Get regions</i>
------------	--------------------

---

**Description**

Extracts regions of a MAgPIE-object

**Usage**

```
getRegions(x)

getRegions(x) <- value
```

**Arguments**

x	MAgPIE object
value	Vector containing the new region names of the MAgPIE objects. If you also want to change the mapping of regions to cell please use <a href="#">getRegionList</a> instead.

**Value**

Regions of the MAgPIE-object

**Functions**

- `getRegions(x) <- value`: overwrite region names

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getYears](#), [getNames](#), [getCPR](#), [read.magpie](#), [write.magpie](#), "magpie"

**Examples**

```
# a <- read.magpie("example.mz")  
# getRegions(a)
```

---

getSets

*Get sets*

---

**Description**

Extracts sets of a MAgPIE-object if available

**Usage**

```
getSets(x, fulldim = TRUE, sep = ".")  
  
getSets(x, fulldim = TRUE, sep = ".") <- value
```

**Arguments**

x	MAgPIE object
fulldim	bool: Consider dimension 3 as a possible aggregate of more dimensions (TRUE) or stick to it as one dimension (FALSE)
sep	A character separating joined dimension names
value	A vector with set names you want to replace the current set names of the object with.

**Value**

Sets of the MAgPIE-object. If no information about contained sets is available NULL

**Functions**

- `getSets(x, fulldim = TRUE, sep = ".") <- value`: replace set names

**Author(s)**

Markus Bonsch, Jan Philipp Dietrich

**See Also**

[getRegions](#), [getNames](#), [getYears](#), [getCPR](#), [read.magpie](#), [write.magpie](#), "magpie"

**Examples**

```
a <- new.maggpie("GL0.1", 2000, c("a.o1", "b.o1", "a.o2"))
getSets(a) <- c("reg", "cell", "t", "bla", "blub")
getSets(a)

getSets(a)["d3.1"] <- "BLA"
getSets(a, fulldim = FALSE)
getSets(a)
```

---

getYears

*Get years*


---

**Description**

Extracts years of a MAgPIE-object

**Usage**

```
getYears(x, as.integer = FALSE)

getYears(x) <- value

setYears(object, nm = NULL)
```

**Arguments**

x, object	MAgPIE object
as.integer	Switch to decide, if output should be the used year-name (e.g. "y1995") or the year as integer value (e.g. 1995)
value, nm	Years the data should be set to. Either supplied as a vector of integers or a vector of characters in the predefined year format ("y0000"). If only 1 year exist you can also set the name of the year to NULL.

**Details**

setYears is a shortcut to use a MAgPIE object with manipulated year names. setYears uses the variable names "object" and "nm" in order to be consistent to the already existing function setName.

**Value**

getYears returns years of the MAgPIE-object, whereas setYears returns the MAgPIE object with the manipulated years.

**Functions**

- getYears(x) <- value: rename years
- setYears(): set years

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getRegions](#), [getNames](#), [setNames](#), [getCPR](#), [read.magpie](#), [write.magpie](#), ["magpie"](#)

**Examples**

```
a <- as.magpie(1)
getYears(a)
setYears(a, 1995)
```

---

hasCoords

*Has Coordinates*

---

**Description**

Checks, whether object contains coordinates.

**Usage**

```
hasCoords(x, xlab = "x", ylab = "y")
```

**Arguments**

x	MAGPIE object
xlab	label of x-dimension
ylab	label of y-dimension

**Value**

Boolean indicating whether coordinates were found or not

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getCoords](#)

**Examples**

```
hasCoords(maxample("pop"))  
hasCoords(maxample("animal"))
```

---

hasSets

*Has Sets*

---

**Description**

Checks, whether set names have been set

**Usage**

```
hasSets(x)
```

**Arguments**

x                    MAgPIE object

**Value**

Boolean indicating whether coordinates were found or not

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getCoords](#)

**Examples**

```
hasSets(maxample("pop"))  
hasSets(maxample("animal"))
```

---

head.magpie	<i>head/tail</i>
-------------	------------------

---

## Description

head and tail methods for MAgPIE objects to extract the head or tail of an object

## Usage

```
## S3 method for class 'magpie'  
head(x, n1 = 3L, n2 = 6L, n3 = 2L, ...)
```

## Arguments

x	MAgPIE object
n1, n2, n3	number of lines in first, second and third dimension that should be returned. If the given number is higher than the length of the dimension all entries in this dimension will be returned.
...	arguments to be passed to or from other methods.

## Value

head returns the first n1 x n2 x n3 entries, tail returns the last n1 x n2 x n3 entries.

## Author(s)

Jan Philipp Dietrich

## See Also

[head](#), [tail](#)

## Examples

```
pop <- maxample("pop")  
head(pop)  
tail(pop, 2, 4, 1)
```

---

is.temporal	<i>is.temporal, is.spatial</i>
-------------	--------------------------------

---

**Description**

Functions to find out whether a vector consists of strings consistent with the definition for auto-detection of temporal or spatial data.

**Usage**

```
is.temporal(x)
```

**Arguments**

x	A vector
---	----------

**Value**

Returns TRUE or FALSE

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
is.temporal(1991:1993)
is.spatial(c("GLO", "AFR"))
```

---

isYear	<i>isYear</i>
--------	---------------

---

**Description**

Function to find out whether a vector consists of strings in the format "yXXXXX" or "XXXXX" with X being a number

**Usage**

```
isYear(x, with_y = TRUE)
```

**Arguments**

x	A vector
with_y	indicates which dataformat years have to have (4-digit without y (e.g.1984) or 5digit including y (y1984))



**Value**

Returns a vector of the length of x with TRUE and FALSE

**Author(s)**

Benjamin Bodirsky

**Examples**

```
x <- c("1955", "y1853", "12a4")
isYear(x, with_y = TRUE)
isYear(x, with_y = FALSE)
```

---

log-methods

*log-method for MAgPIE objects*

---

**Description**

log-method for MAgPIE-objects respectively. Works exactly as for arrays.

**Usage**

```
## S4 method for signature 'magpie'
log(x, base = exp(1))
```

**Arguments**

x	a magpie object
base	ia positive or complex number: the base with respect to which logarithms are computed. Defaults to e=exp(1).

**Methods**

**x = "magpie"** works as log(x) for arrays.

---

 lowpass

*Lowpass Filter*


---

### Description

Filters high frequencies out of a time series. The filter has the structure  $x'(n) = (x(n-1) + 2 * x(n) + x(n+1)) / 4$

### Usage

```
lowpass(x, i = 1, fix = NULL, altFilter = NULL, warn = TRUE)
```

### Arguments

x	Vector of data points, that should be filtered or MAgPIE object
i	number of iterations the filter should be applied to the data
fix	Fixes the starting and/or ending data point. Default value is NULL which doesn't fix any point. Available options are: "start" for fixing the starting point, "end" for fixing the ending point and "both" for fixing both ends of the data.
altFilter	set special filter rule to indexes defined in this parameter. The special filter has the structure $x'(n) = (2 * x(n) + x(n+1)) / 3$
warn	boolean deciding whether lowpass issues a warning for critical parameter choices or not

### Value

The filtered data vector or MAgPIE object

### Author(s)

Jan Philipp Dietrich, Misko Stevanovic

### Examples

```
lowpass(c(1, 2, 11, 3, 4))
# to fix the starting point
lowpass(c(0, 9, 1, 5, 14, 20, 6, 11, 0), i = 2, fix = "start")
```

---

magclassdata	<i>magclassdata</i>
--------------	---------------------

---

**Description**

General magclass-dataset

**Details**

Please do not directly access that data. It should be only used by library functions.

**Author(s)**

Jan Philipp Dietrich

---

magpie-class	<i>Class "magpie" ~~~</i>
--------------	---------------------------

---

**Description**

The MAgPIE class is a data format for cellular MAgPIE data with a close relationship to the array data format. `is.magpie` tests if `x` is an MAgPIE-object, `as.magpie` transforms `x` to an MAgPIE-object (if possible).

**Arguments**

<code>x</code>	An object that should be either tested or transformed as/to an MAgPIE-object.
<code>...</code>	additional arguments supplied for the conversion to a MAgPIE object. Allowed arguments for arrays and dataframes are <code>spatial</code> and <code>temporal</code> both expecting a vector of dimension or column numbers which contain the spatial or temporal information. By default both arguments are set to <code>NULL</code> which means that the <code>as.magpie</code> will try to detect automatically the temporal and spatial dimensions. The arguments will just overwrite the automatic detection. If you want to specify that the data does not contain a spatial or temporal dimension you can set the corresponding argument to <code>0</code> . In addition <code>as.magpie</code> for <code>data.frames</code> is also expecting an argument called <code>datacol</code> which expects a number stating which is the first column containing data. This argument should be used if the dimensions are not detected correctly, e.g. if the last dimension column contains years which are then detected as values and therefore interpreted as first data column. In addition an argument <code>tidy=TRUE</code> can be used to indicate that the <code>data.frame</code> structure is following the rules of tidy data (last column is the data column all other columns contain dimension information). This information will help the conversion. <code>sep</code> defines the dimension separator (default is <code>."</code> ) and <code>replacement</code> defines how the separator as a reserved character should be converted in order to not mess up with the object (default <code>"_"</code> ). Another available argument for conversions of

data.frames and quitte objects to magpie is `filter` if set to TRUE (default) "." (separator) will be replaced with the replacement character and empty entries will be replaced with a single space. If set to FALSE no filter will be applied to the data.

### Objects from the Class

Objects can be created by calls of the form `new("magpie", data, dim, dimnames, ...)`. MAgPIE objects have three dimensions (cells,years,datatype) and the dimensionnames of the first dimension have the structure "REGION.cellnumber". MAgPIE-objects behave the same like array-objects with 2 exceptions:

1. Dimensions of the object will not collapse (e.g. `x[1,1,1]` will remain 3D instead of becoming 1D)
2. It is possible to extract full regions just by typing `x["REGIONNAME",,]`.

Please mind following standards:

Header must not contain any purely numeric entries, but combinations of characters and numbers are allowed (e.g. "bla","12" is forbidden, whereas "bla","b12" is allowed)

Years always have the structure "y" + 4-digit number, e.g. "y1995"

Regions always have the structure 3 capital letters, e.g. "AFR" or "GLO"

This standards are necessary to allow the scripts to detect headers, years and regions properly and to have a distinction to other data.

### Author(s)

Jan Philipp Dietrich

### See Also

[read.magpie](#), [write.magpie](#), [getRegions](#), [getYears](#), [getNames](#), [getCPR](#), [ncells](#), [nyears](#), [ndata](#)

### Examples

```
showClass("magpie")

pop <- maxample("pop")

# returning PA0 and PAS for 2025
pop["PA", 2025, , pmatch = "left"]

# returning CPA for 2025
pop["PA", 2025, , pmatch = "right"]

# returning CPA PA0 and PAS for 2025
pop["PA", 2025, , pmatch = TRUE]

# returning PAS and 2025
```

```

pop["PAS", 2025, ]

# return all entries for year 2025
pop[2025, dim = 2]

# returning everything but values for PAS or values for 2025
pop["PAS", 2025, , invert = TRUE]

# accessing subdimension via set name

a <- maxample("animal")
a[list(country = "NLD", y = "53p25"), , list(species = c("rabbit", "dog"))]

# please note that the list elements act as filter. For instance, the
# following example will not contain any dogs as the data set does
# not contain any dogs which are black.
a[list(country = "NLD", y = "53p25"), , list(species = c("rabbit", "dog"), color = "black")]

# it is also possible to extract given combinations of subdimensions
# via a data-frame
df <- data.frame(getItems(a, 3, split = TRUE, full = TRUE))[c(1, 3, 4), ][3:2]
getItems(a[df], 3)

# Unknown dimensions to be added in output!
df$blub <- paste0("bl", 1:dim(df)[1])
getItems(a[df], 3)

```

---

magpiesort

*MAGPIE-Sort*


---

## Description

Brings the spatial and temporal structure of MAGPIE objects in the right order. This function is especially useful when you create new MAGPIE objects as the order typically should be correct for MAGPIE objects.

## Usage

```
magpiesort(x)
```

## Arguments

x                   MAGPIE object which might not be in the right order.

## Value

The eventually corrected MAGPIE object (right order in spatial in temporal dimension)

**Author(s)**

Jan Philipp Dietrich

**See Also**["magpie"](#)**Examples**

```
pop <- maxample("pop")
a <- magpiesort(pop)
```

---

magpie\_expand

*magpie\_expand*


---

**Description**

Expands a MAgPIE object based on a reference

**Usage**

```
magpie_expand(x, ref)
```

**Arguments**

x	MAgPIE object that should be expanded
ref	MAgPIE object that serves as a reference

**Details**

Expansion means here that the dimensions of `x` are expanded accordingly to `ref`. Please note that this is really only about expansion. In the case that one dimension of `ref` is smaller than of `x` nothing happens with this dimension. At the moment `magpie_expand` is only internally available in the `magclass` library

You can influence the verbosity of this function by setting the option `"magclass.verbosity"`. By default verbosity is set to 1 which means that only warnings are returned. Setting verbosity to 2 means that warnings as well as additional notes are returned. This is done by `options(verbosity.level=2)`

With version 5 of the package `magpie_expand` has been updated to a newer version (currently 2.1) and since version 6 this is the only currently supported version. To switch to the old setup you have to install `magclass` in a version `< 6` and set `options(magclass_expand_version=1)`.

By default expansion is based on the elements in a dimension ignoring the set name of the dimension. To expand based on set names instead of contents (recommended) you can switch `options(magclass_setMatching=TRUE)`. Please be careful with this setting as it alters the behavior of `magclass` objects quite significantly! For more information have a look at `vignette("magclass-expansion")`.

**Value**

An expanded version of x.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[as.magpie](#), [options](#)

**Examples**

```
a <- new.magpie(c("AFR", "CPA"), "y1995", c("m", "n"))
b <- new.magpie("GLO", "y1995", c("bla", "blub"))
magpie_expand(b, a)
options(magclass.verbosity = 2)
magpie_expand(b, a)
```

---

magpie\_expand\_dim      *magpie\_expand\_dim*

---

**Description**

Expands a single MAgPIE object dimension

**Usage**

```
magpie_expand_dim(x, ref, dim = 1)
```

**Arguments**

x	MAgPIE object that should be expanded
ref	MAgPIE object that serves as a reference
dim	dimension that should be expanded

**Details**

Expansion means here that the dimensions of x are expanded accordingly to ref. Please note that this is really only about expansion. In the case that one dimension of ref is smaller than of x nothing happens with this dimension. At the moment magpie\_expand is only internally available in the magclass library

In contrast to [magpie\\_expand](#) this function is expanding only a single dimension. It is meant as a support function for [magpie\\_expand](#) itself.

**Value**

An expanded version of x.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[as.magpie](#), [options](#)

**Examples**

```
d <- new.magpie(c("AFR.BLUB.1", "AFR.BLUB.2", "EUR.BLUB.1",
                "AFR.BLA.1", "AFR.BLA.2", "EUR.BLA.1"), fill = 1)
getSets(d)[1:3] <- c("reg", "b", "i")
e <- new.magpie(c("BLA.AFR.A", "BLA.EUR.A", "BLUB.AFR.A", "BLUB.EUR.A",
                "BLA.AFR.B", "BLA.EUR.B", "BLUB.AFR.B", "BLUB.EUR.B"), fill = 2)
getSets(e)[1:3] <- c("b", "reg", "a")
magclass::magpie_expand_dim(d, e, dim = 1)
```

---

magpply

*magpply*

---

**Description**

apply command for magpieobjects. Very efficient for replacing loops.

**Usage**

```
magpply(X, FUN, MARGIN = NULL, DIM = NULL, ..., INTEGRATE = FALSE)
```

**Arguments**

X	magpie object
FUN	function that shall be applied X
MARGIN	dimension over which FUN shall be applied (like a loop over that dimension). This dimension will be preserved in the output object (see also DIM).
DIM	dimension in which FUN shall be applied. This dimension will be missing in the output. DIM and MARGIN are opposite ways of expressing the dimensions to be addressed and you must only use one of them with MARGIN excluding dimensions from the calculation and DIM including them.
...	further parameters passed on to FUN
INTEGRATE	if TRUE, the output will be filled into an magpie object of the same dimensionality as X



**Value**

magpie object

**Author(s)**

Jan Philipp Dietrich, Benjamin Leon Bodirsky

**Examples**

```
pop <- maxample("pop")
magpply(pop, FUN = sum, MARGIN = 2)
fourdim <- pop * setNames(pop, c("jkk", "lk"))
magpply(fourdim, FUN = sum, MARGIN = c(1, 3.1))
```

---

maxample

*maxample*

---

**Description**

A collection of magclass example data sets

**Usage**

```
maxample(data)
```

**Arguments**

data	name of the example data set. Currently available are "pop" (regional population data, previously named "population_magpie"), "animal" (fictional, high-dimensional animal sighting data set) and "bilateral" (fictional, bilateral trade cost data set).
------	---

**Value**

the chosen example data set

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
p <- maxample("pop")
str(p)

a <- maxample("animal")
str(a)
getItems(a, split = TRUE)
```

---

`mbind`*mbind*

---

**Description**

Merges MAgPIE-objects with identical structure in two dimensions. If data differs in the temporal or spatial dimension each year or region/cell must appear only once!

**Usage**

```
mbind(...)
```

**Arguments**

... MAgPIE objects or a list of MAgPIE objects that should be merged.

**Value**

The merged MAgPIE object

**Author(s)**

Jan Philipp Dietrich, Misko Stevanovic

**See Also**

["magpie"](#)

**Examples**

```
m <- new.magpie(c("AFR", "CPA", "EUR"), c(1995, 2005), "Data1", fill = c(1, 2, 3, 4, 5, 6))
ms <- dimSums(m, dim = 3.1)
mbind(m, ms)
my <- new.magpie(getRegions(m), 2010, getNames(m), fill = c(6, 6, 4))
mbind(m, my)
md <- new.magpie(getRegions(m), getYears(m), "Data2", fill = c(7, 6, 5, 7, 8, 9))
mbind(m, md)

pop <- maxample("pop")
a <- mbind(pop, pop)
dim(pop)
dim(a)
```

---

mcalc	<i>mcalc</i>
-------	--------------

---

**Description**

Select values from a MAgPIE-object

**Usage**

```
mcalc(x, f, dim = NULL, append = FALSE)
```

**Arguments**

x	MAgPIE object
f	A formula describing the calculation that should be performed
dim	The dimension in which the manipulation should take place. If set to NULL function tries to detect the dimension automatically.
append	If set to TRUE the result will be appended to x, otherwise the result will be returned.

**Details**

This functions only work for MAgPIE objects with named dimensions as the dimension name (set\_name) has to be used to indicate in which dimension the entries should be searched for!

**Value**

The calculated MAgPIE object in the case that append is set to FALSE. Otherwise nothing is returned (as x is appended in place)

**Author(s)**

Jan Philipp Dietrich

**See Also**

[mselect](#)

**Examples**

```
pop <- maxample("pop")
pop
mcalc(pop, X12 ~ A2 * B1, append = TRUE)
pop
mcalc(pop, `Nearly B1` ~ 0.5 * A2 + 99.5 * B1)
```

---

mselect	<i>MSelect</i>
---------	----------------

---

**Description**

Select values from a MAgPIE-object

**Usage**

```
mselect(x, ..., collapseNames = FALSE)
```

```
mselect(x, ...) <- value
```

**Arguments**

x	MAgPIE object
...	entry selections of the form <code>set_name=c(set_elem1,set_elem2)</code> . Alternatively a single list element containing these selections can be provided.
collapseNames	Boolean which decides whether names should be collapsed or not.
value	values on which the selected magpie entries should be set.

**Details**

This functions only work for MAgPIE objects with named dimensions as the dimension name (`set_name`) has to be used to indicate in which dimension the entries should be searched for!

**Value**

The reduced MAgPIE object containing only the selected entries or the full MAgPIE object in which a selection of entries was manipulated.

**Functions**

- `mselect(x, ...) <- value`: replace values in magpie object

**Author(s)**

Jan Philipp Dietrich

**See Also**

[collapseNames](#), ["magpie"](#)

**Examples**

```
pop <- maxample("pop")
mselect(pop, i = c("AFR", "EUR"), scenario = "A2", t = "y2035")
```

---

ncells	<i>Count elements</i>
--------	-----------------------

---

**Description**

Functions to count the number of cells/years/datasets/regions of an MAgPIE-object

**Usage**

```
ncells(x)
```

```
ndata(x)
```

```
nregions(x)
```

```
nyears(x)
```

**Arguments**

x	A MAgPIE-object
---	-----------------

**Value**

value	The number of cells/years/datasets/regions of x
-------	---

**Functions**

- `ndata()`: count datasets
- `nregions()`: count regions
- `nyears()`: count years

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
a <- is.magpie(NULL)
ncells(a)
nyears(a)
ndata(a)
nregions(a)
```

---

ndim	<i>Count sub-dimensions</i>
------	-----------------------------

---

**Description**

Functions to count the subdimensions of an MAgPIE-object

**Usage**

```
ndim(x, dim = NULL)
```

**Arguments**

x	A MAgPIE-object
dim	main dimension in which the sub-dimensions should be counted. If NULL the sum of all subdimensions is returned

**Value**

Number of subdimensions

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
a <- maxample("animal")
ndim(a)
ndim(a,1)
ndim(a,2)
ndim(a,3)
```

---

new.magpie	<i>new.magpie</i>
------------	-------------------

---

**Description**

Creates a new MAgPIE object

**Usage**

```
new.magpie(
  cells_and_regions = "GLO",
  years = NULL,
  names = NULL,
  fill = NA,
  sort = FALSE,
  sets = NULL,
  unit = NULL
)
```

**Arguments**

cells_and_regions	Either the region names (e.g. "AFR"), or the cells (e.g. 1:10), or both in combination (e.g. "AFR.1"). NULL means no spatial element.
years	dimnames for years in the format "yXXXX" or as integers. NULL means one year which is not further specified
names	dimnames for names. NULL means one data element which is not further specified
fill	Default value for the MAgPIE object
sort	Boolean. Decides, whether output should be sorted or not.
sets	A vector of dimension names. See <a href="#">getSets</a> for more information.
unit	deprecated

**Value**

an empty magpie object filled with fill, with the given dimnames

**Author(s)**

Benjamin Bodirsky, Jan Philipp Dietrich

**See Also**

[as.magpie](#)

**Examples**

```
a <- new.magpie(1:10, 1995:2000)
b <- new.magpie(c("AFR", "CPA"), "y1995", c("bla", "blub"), sets = c("i", "t", "value"))
c <- new.magpie()
```

---

population\_magpie      *population\_magpie*

---

**Description**

Example dataset for a regional MAgPIE object

**Value**

A2 and B1 population scenario from SRES

**Author(s)**

Benjamin Bodirsky

---

print.magpie      *print*

---

**Description**

print method for MAgPIE objects for convenient display of magpie data.

**Usage**

```
## S3 method for class 'magpie'
print(x, drop = TRUE, reshape = FALSE, ...)
```

**Arguments**

x	MAgPIE object
drop	argument which controls whether empty dimensions should be skipped or not.
reshape	argument that controls tabular representation of nested data dimension cross tables, FALSE will reproduce standard print behavior any pair of two dimension numbers will create a table for these two dims, and loop over the other dimensions
...	arguments to be passed to or from other methods.

**Value**

Invisibly, the MAgPIE object x.

**Author(s)**

Jan Philipp Dietrich, Kristine Karstens, Felicitas Beier



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

```
pop <- maxample("pop")
print(pop)
print(pop[, 1, ], drop = FALSE)
print(pop[, 1, ])
```

read.magpie

*Read MAgPIE-object from file***Description**

Reads a MAgPIE-file and converts it to a 3D array of the structure (cells,years,datacolumn)

**Usage**

```
read.magpie(
  file_name,
  file_folder = "",
  file_type = NULL,
  as.array = FALSE,
  comment.char = "*",
  check.names = FALSE,
  ...
)
```

**Arguments**

file_name	file name including file ending (wildcards are supported). Optionally also the full path can be specified here (instead of splitting it to file_name and file_folder)
file_folder	folder the file is located in (alternatively you can also specify the full path in file_name - wildcards are supported)
file_type	format the data is stored in. If file_type=NULL the file ending of the file_name is used as format. If format is different to the formats mentioned standard MAgPIE format is assumed. See <a href="#">write.magpie</a> for a list of supported file formats.
as.array	Should the input be transformed to an array? This can be useful for regional or global inputs, but all advantages of the magpie-class are lost.
comment.char	character: a character vector of length one containing a single character or an empty string. Use "" to turn off the interpretation of comments altogether. If a comment is found it will be stored in attr("comment"). In text files the comment has to be at the beginning of the file in order to be recognized by read.magpie.

check.names      logical. If TRUE then the names of the variables in the data frame are checked to ensure that they are syntactically valid variable names. Same functionality as in read.table.

...                additional arguments passed to specific read functions (e.g. varname for specifying the variable to be read in from a multi-variable NCDF file.)

### Details

See [write.magpie](#) for a list of supported file formats.

### Value

x                    MAgPIE-object

### Note

See [write.magpie](#) for the detailed structure of binary MAgPIE formats .m and .mz.

### Author(s)

Jan Philipp Dietrich, Stephen Bi, Florian Humpenoeder, Pascal Sauer

### See Also

"[magpie](#)", [write.magpie](#)

---

read.report	<i>Read file in report format</i>
-------------	-----------------------------------

---

### Description

This function reads the content of a reporting file (a file in the model intercomparison file format \*.mif) into a list of MAgPIE objects or a single MAgPIE object.

### Usage

```
read.report(file, as.list = TRUE, showSeparatorWarning = TRUE)
```

### Arguments

file                file name the object should be read from.

as.list            if TRUE a list is returned (default), if FALSE it is tried to merge all information in one MAgPIE object (still under development and works currently only if the entries for the different models and scenarios have exactly the same regions and years).

showSeparatorWarning      Boolean (default value TRUE) that decides whether the warning about the replacement of dots in variable names is displayed (default value) or not.

## Details

The **Model Intercomparison File Format (MIF)** is the default file format for data produced by Integrated Assessment Models. It is based on the common format used for Model Intercomparison Projects such as EMF and SSP with some slight changes/clarifications in its definition. For interactions between models this format should be used. For everything else it is at least recommended to use this format, too.

Aim of this standardization is to achieve a more flexible and smooth communication between models and to facilitate the creation of aggregated outputs from integrated assessment scenario runs which then can easily be uploaded to external databases such as the EMF or SSP database. By using this standard most of the required decisions for a working input output interface between models have already been specified which significantly reduces the required work to get a new interaction running.

### Definition

The format is characterized by the following features:

- The file ending is ".mif"
- The file is written in ASCII format
- Entries are separated with ";", every line ends with a ";"
- The file always contains a header
- The format of the header is: Model;Scenario;Region;Variable;Unit;<ADDITIONAL\_COLUMNS>;<YEARS>;

The first 5 entries always have to exist, <ADDITIONAL\_COLUMNS> is additional information which can be added optionally (e.g. "Description") and <YEARS> are the years for which data is delivered. <YEARS> are always written as 4 digit numbers. In the (very unlikely) case that a year before 1000 is used the number has to start with a 0, e.g. 0950. <ADDITIONAL\_COLUMNS> can be anything, there are no further rules at the moment what it can contain. However, there are strict rules for naming these columns. Allowed are single names starting with a capital letter without special characters in it except "\_" which is allowed. Examples: "Description" allowed, "More Description" not allowed, "More\_Description" allowed, "123Description" not allowed, "Description123" allowed. Scripts using this format must be able to ignore additional columns. For years there are no specific limitations/requirements which years should be reported. Scripts dealing with this data must be able to work with different temporal resolutions. For variables basically everything can be reported here. Missing values have to be marked with "N/A".

### Author(s)

Jan Philipp Dietrich

### See Also

[write.report](#)

### Examples

```
## Not run:  
read.report("report.csv")
```

```
## End(Not run)
```

---

replace\_non\_finite      *Replace Non-Finite Data*

---

### Description

Replaces all instances of non-finite data (NA, NaN, Inf, and -Inf).

### Usage

```
replace_non_finite(x, replace = 0)
```

### Arguments

x                      A vector or [magpie](#) object.  
replace                A value to replace non-finite data with.

### Value

A vector or [magpie](#) object, same as x.

### Author(s)

Michaja Pehl

### Examples

```
part <- new.magpie(letters[1:3], years = 'y1995', names = 'foo')
total <- new.magpie(letters[1:3], years = 'y1995', names = 'foo')

part[,,] <- c(0, 1, 2)
total[,,] <- c(0, 10, 10)

part / total

replace_non_finite(part / total)
```

---

round-methods	<i>Round-method for MAgPIE objects</i>
---------------	--

---

**Description**

Round-method for MAgPIE-objects respectively. Works exactly as for arrays.

**Usage**

```
## S4 method for signature 'magpie'
round(x, digits = 0)
```

**Arguments**

x	a magpie object
digits	integer indicating the number of decimal places (round) or significant digits (signif) to be used. Negative values are allowed.

**Methods**

`x = "magpie"` works as `round(x)` for arrays.

---

rowSums-methods	<i>~~ Methods for Function rowSums and rowMeans ~~</i>
-----------------	--

---

**Description**

~~ Methods for function rowSums and rowMeans~~

**Usage**

```
## S4 method for signature 'magpie'
rowSums(x, na.rm = FALSE, dims = 1, ...)
```

**Arguments**

x	object on which calculation should be performed
na.rm	logical. Should missing values (including NaN) be omitted from the calculations?
dims	integer: Which dimensions are regarded as "rows" or "columns" to sum over. For row*, the sum or mean is over dimensions dims+1, ...; for col* it is over dimensions 1:dims.
...	further arguments passed to other colSums/colMeans methods

**Methods**

`list("signature(x = \"ANY\")")` normal rowSums and rowMeans method

`list("signature(x = \"magpie\")")` classical method prepared to handle MAgPIE objects

---

setItems	<i>Set Items</i>
----------	------------------

---

**Description**

Set items of a given (sub-)dimension of a MAgPIE-object

**Usage**

```
setItems(x, dim, value, maindim = NULL, raw = FALSE)
```

**Arguments**

<code>x</code>	MAgPIE object
<code>dim</code>	Dimension for which the items should be returned. Either number or name of dimension or a vector of these. See <a href="#">dimCode</a> for more details.
<code>value</code>	a vector with the length of the main dimension the dimnames should be replaced in / added to. If set to NULL the corresponding dimension will be removed.
<code>maindim</code>	main dimension the data should be added to (does not need to be set if <code>dim</code> exists in the data. Should be set if <code>dim</code> might not exist, or if <code>dim</code> might potentially exist in a different main dimension than the one anticipated).
<code>raw</code>	if set to FALSE inputs will be corrected (e.g. dots replaced by the letter "p") if necessary. If TRUE data will be written as is (risking the creation of inconsistent objects).

**Value**

the manipulated MAgPIE object

**Author(s)**

Jan Philipp Dietrich

**See Also**

[getItems](#)

**Examples**

```
x <- maxample("pop")
setItems(x, "i", paste0("REG", 1:ncells(x)))
```

---

setNames-methods	<i>Get dataset names</i>
------------------	--------------------------

---

**Description**

Extracts dataset names of a MAgPIE-object

**Usage**

```
## S4 method for signature 'magpie'
setNames(object = nm, nm)
```

**Arguments**

object	MAgPIE object
nm	a vector of names current names should be replaced with. If only one data element exists you can also set the name to NULL.

**Details**

setNames is a shortcut to use a MAgPIE object with manipulated data names. The setNames method uses the variable names "object" and "nm" in order to be consistent to the already existing function setNames.

**Methods**

```
list("signature(object = \"ANY\")") normal setNames method
list("signature(object = \"magpie\")") setNames for MAgPIE objects
```

**See Also**

[getNames](#),

---

show-methods	<i>~~ Method for function show ~~</i>
--------------	---------------------------------------

---

**Description**

Show a magpie object by calling the default show method, print, or str, depending on options("magclass\_show\_func").

**Usage**

```
## S4 method for signature 'magpie'
show(object)
```

**Arguments**

object            A MAgPIE-object

**Author(s)**

Pascal Sauer

---

sizeCheck

*sizeCheck*

---

**Description**

Calculates expected magclass object length and checks that it stays below the limit defined with `magclass_sizeLimit` (default =  $10^9$ ). This is useful to prevent out of memory errors in case of unwanted object expansions Ignored if `getOption("magclass_sizeLimit")` is negative.

**Usage**

```
sizeCheck(dim)
```

**Arguments**

dim                dimensions of the current object as returned by function `dim`

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
pop <- maxample("pop")
magclass:::sizeCheck(dim(pop))

## Not run:
magclass:::sizeCheck(c(6765L, 10946L, 17711L))

## End(Not run)
```



spatRasterToDataset     *spatRasterToDataset*

**Description**

Convert a SpatRaster to a SpatRasterDataset.

**Usage**

spatRasterToDataset(x)

**Arguments**

x                      SpatRaster with names of the form "y[0-9]+..[varname]"

**Value**

SpatRasterDataset

**Author(s)**

Pascal Sauer

str.magpie                      *str*

**Description**

str method for MAGPIE objects for convenient display of the structure of a magpie object.

**Usage**

```
## S3 method for class 'magpie'
str(object, ...)
```

**Arguments**

object                      MAGPIE object  
 ...                          arguments to be passed to or from other methods.

**Details**

In contrast to the default str this will not show the attributes of object@.Data as these contain only a duplicate of dimnames. Also, dim is not shown, because the information it provides is implicitly included in dimnames.

**Author(s)**

Pascal Sauer

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

```
str(maxample("pop"))
```

---

`suppressSpecificWarnings`*suppressSpecificWarnings*

---

**Description**

Like [suppressWarnings](#), but instead of suppressing all warnings this only suppresses warnings if they match the given pattern.

**Usage**

```
suppressSpecificWarnings(expr, regularExpr, fixed = FALSE)
```

**Arguments**

<code>expr</code>	The expression/code to evaluate, can be a block of code inside curly braces.
<code>regularExpr</code>	Only warnings matching this regular expression are suppressed.
<code>fixed</code>	Match the literal string given by <code>regularExpr</code> instead of interpreting it as a regular expression. Passed to <a href="#">grepl</a> .

**Value**

The result of evaluating `expr`.

**Author(s)**

Pascal Sauer

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

---

time_interpolate	<i>time_interpolate</i>
------------------	-------------------------

---

### Description

Function to extrapolate missing years in MAgPIE objects.

### Usage

```
time_interpolate(  
  dataset,  
  interpolated_year,  
  integrate_interpolated_years = FALSE,  
  extrapolation_type = "linear"  
)
```

### Arguments

dataset	An MAgPIE object
interpolated_year	Vector of years, of which values are required. Can be in the formats 1999 or y1999.
integrate_interpolated_years	FALSE returns only the dataset of the interpolated year, TRUE returns the whole dataset, including all years of data and the interpolated year
extrapolation_type	Determines what happens if extrapolation is required, i.e. if a requested year lies outside the range of years in dataset. Specify "linear" for a linear extrapolation. "constant" uses the value from dataset closest in time to the requested year.

### Value

Uses linear extrapolation to estimate the values of the interpolated year, using the values of the two surrounding years. If the value is before or after the years in data, the two closest neighbours are used for extrapolation.

### Author(s)

Benjamin Bodirsky, Jan Philipp Dietrich

### See Also

[convergence](#)

**Examples**

```
p <- maxample("pop")
time_interpolate(p, "y2000", integrate = TRUE)
time_interpolate(p, c("y1980", "y2000"), integrate = TRUE, extrapolation_type = "constant")
```

---

unitsplit	<i>splits a data.frame or vector of strings with form 'variable (unit)' into a data.frame with variable and unit separated</i>
-----------	--

---

**Description**

splits a data.frame or vector of strings with form 'variable (unit)' into a data.frame with variable and unit separated

**Usage**

```
unitsplit(x, col = "variable")
```

**Arguments**

x	data.frame or vector of strings
col	column name. Default: variable

**Author(s)**

Jan Philipp Dietrich, Mika Pflüger, Oliver Richters

---

unwrap	<i>Unwrap</i>
--------	---------------

---

**Description**

Creates a higher dimensional array by separating all subdimensions in the third dimension of a MAgPIE object and returning them as separate dimension.

**Usage**

```
unwrap(x, sep = NULL)
```

**Arguments**

x	A MAgPIE object
sep	deprecated, please do not use anymore

**Value**

An array with the full dimensionality of the original data

**Author(s)**

Jan Philipp Dietrich

**See Also**

[wrap](#), [fulldim](#)

**Examples**

```
a <- as.magpie(array(1:6, c(3, 2), list(c("bla", "blub", "ble"), c("up", "down"))))
unwrap(a)
```

---

where

*where*

---

**Description**

Analysis function for magpie objects

**Usage**

```
where(x, plot = NULL)
```

**Arguments**

**x** A logical statement with a magpie object  
**plot** deprecated. Use the function whereplot in package luplot.

**Value**

A list of analysis parameters

**Author(s)**

Benjamin Leon Bodirsky, Jan Philipp Dietrich

**See Also**

whereplot in package luplot

**Examples**

```
p <- maxample("pop")
where(p > 500)
```

wrap

*Wrap*

---

**Description**

Reshape an array or a matrix by permuting and/or joining dimensions.

**Usage**

```
wrap(x, map = list(NA), sep = ".")
```

**Arguments**

x	An array
map	A list of length equal to the number of dimensions in the reshaped array. Each element should be an integer vectors specifying the dimensions to be joined in corresponding new dimension. One element may equal NA to indicate that that dimension should be a join of all non-specified (remaining) dimensions. Default is to wrap everything into a vector.
sep	A character separating joined dimension names

**Note**

This function is extracted from the R.utils library which is licensed under LGPL>=2.1 and written by Henrik Bengtsson.

**Author(s)**

Henrik Bengtsson, Jan Philipp Dietrich

**See Also**

[unwrap](#), [fulldim](#)

---

write.magpie*Write MAgPIE-object to file*

---

**Description**

Writes a magpie object to a file. The file type is determined by the filename extension. The written file can be read again using [read.magpie](#).

**Usage**

```

write.magpie(
  x,
  file_name,
  file_folder = "",
  file_type = NULL,
  append = FALSE,
  comment = NULL,
  comment.char = "*",
  mode = NULL,
  zname = "time",
  ...
)

```

**Arguments**

x	a magclass object. An exception is that formats written via the raster package (currently "nc", "asc", "grd" and "tif") also accept RasterBrick objects which have been previously created from a magclass object via as.RasterBrick)
file_name	file name including file ending (wildcards are supported). Optionally also the full path can be specified here (instead of splitting it to file_name and file_folder)
file_folder	folder the file should be written to (alternatively you can also specify the full path in file_name - wildcards are supported)
file_type	Format the data should be stored as. If file_type=NULL the file ending of the file_name is used as format. See detailed description for a list of available file types. Please be aware that the file_name is independent of the file_type you choose here, so no additional file ending will be added!
append	Decides whether an existing file should be overwritten (FALSE) or the data should be added to it (TRUE). Append = TRUE only works if the existing data can be combined with the new data using the mbind function
comment	Vector of strings: Optional comment giving additional information about the data. If different to NULL this will overwrite the content of attr(x,"comment")
comment.char	character: a character vector of length one containing a single character or an empty string. Use "" to turn off the interpretation of comments altogether.
mode	File permissions the file should be written with as 3-digit number (e.g. "777" means full access for user, group and all, "750" means full access for user, read access for group and no access for anybody else). Set to NULL system defaults will be used. Access codes are identical to the codes used in unix function chmod.
zname	name of the time variable for raster files like nc, asc, grd and tif
...	additional arguments passed to specific write functions

**Details**

This function supports writing the following file types:

- "cs2" is the new standard format for cellular data with or without header and the first columns (year,regiospatial) or only (regiospatial)
- "cs2b" is identical to "cs2" except that it will suppress the data name if it has only 1 element in the data dimension.
- "csv" is the standard format for regional data with or without header and the first columns (year,region,cellnumber) or only (region,cellnumber)
- "cs3" is another csv format which is specifically designed for multidimensional data for usage in GAMS.
- "cs4" alternative multidimensional format compatible to GAMS, in contrast to cs3 it can also handle sparse data
- "csvr", "cs2r", "cs3r", "cs4r" which are the same formats as the ones previously explained with the only difference that they have a REMIND compatible format
- "cs5" a more generalized version of cs4
- "rds" is an R-default format for storing R objects
- "m" (magpie) and "mz" (magpie zipped) are new formats developed to allow a less storage intensive management of MAgPIE-data. The only difference between both formats is that .mz is gzipped whereas .m is not compressed. So .mz needs less memory, whereas .m might have a higher compatibility to other languages
- "asc" is the ASCII grid format. It can only be applied for gridded data and writes one file per year per data column. In the case that more than one year and data column is supplied several files are written with the structure filename\_year\_datacolumn.asc
- "tif" is the GEOTiff format for gridded data.
- "grd" is the native raster format for gridded data.
- "nc" is the netCDF format for gridded data.

### Note

The binary MAgPIE formats .m and .mz have the following content/structure (you only have to care for that if you want to implement read.magpie/write.magpie functions in other languages):

```
[ FileFormatVersion | Current file format version number (currently 6) | integer | 2 Byte ]
[ nchar_comment | Number of character bytes of the file comment | integer | 4 Byte ]
[ nbyte_metadata | Number of bytes of the serialized metadata (currently = 0) | integer | 4 Byte ]
[ nchar_sets | Number of characters bytes of all regionnames + 2 delimiter | integer | 2 Byte]
[ nyears | Number of years | integer | 2 Byte ]
[ yearList | All years of the dataset (0, if year is not present) | integer | 2*nyears Byte ]
[ ncells | Number of cells | integer | 4 Byte ]
[ nchar_cell | Number of characters bytes of all regionnames + (nreg-1) for delimiters | integer | 4
Byte ]
[ cells | Cell names saved as cell1\cell2 (\n is the delimiter) | character | 1*nchar_cell Byte ]
[ nelelem | Total number of data elements | integer | 4 Byte ]
[ nchar_data | Number of char. bytes of all datanames + (ndata - 1) for delimiters | integer | 4 Byte ]
[ datanames | Names saved in the format data1\ndata2 (\n as del.) | character | 1*nchar_data Byte ]
[ data | Data of the MAgPIE array in vectorized form | numeric | 4*nelem Byte ]
[ comment | Comment with additional information about the data | character | 1*nchar_comment
```



Byte ]  
 [ sets | Set names with \n as delimiter | character | 1\*nchar\_sets Byte]  
 [ metadata | serialized metadata information (currently not in use) | bytes | 1\*nbyte\_metadata Byte]

### Author(s)

Jan Philipp Dietrich, Stephen Bi, Florian Humpenoeder, Pascal Sauer

### See Also

"magpie", [read.magpie](#), [mbind](#)

### Examples

```
pop <- maxample("pop")
path <- tempfile(fileext = ".mz")
write.magpie(pop, path)
pop2 <- read.magpie(path)
```

---

write.report

*Write file in report format*

---

### Description

This function writes the content of a MAgPIE object into a file or returns it directly using the reporting format as it is used for many model inter-comparisons.

### Usage

```
write.report(
  x,
  file = NULL,
  model = NULL,
  scenario = NULL,
  unit = NULL,
  ndigit = 4,
  append = FALSE,
  skipempty = TRUE,
  extracols = NULL
)
```

### Arguments

x MAgPIE object or a list of lists with MAgPIE objects as created by read.report. In the latter case settings for model and scenario are overwritten by the information given in the list.

file	file name the object should be written to. If NULL the formatted content is returned
model	Name of the model which calculated the results
scenario	The scenario which was used to get that results.
unit	Unit of the data. Only relevant if unit is not already supplied in Dimnames (format "name (unit)"). Can be either a single string or a vector of strings with a length equal to the number of different data elements in the MAgPIE object
ndigit	Number of digits the output should have
append	Logical which decides whether data should be added to an existing file or an existing file should be overwritten
skipempty	Determines whether empty entries (all data NA) should be written to file or not.
extracols	names of dimensions which should appear in the output as additional columns

**Author(s)**

Jan Philipp Dietrich

**See Also**

[read.report](#)

**Examples**

```
write.report(maxample("pop"))
```

---

write.report2	<i>Write file in report format</i>
---------------	------------------------------------

---

**Description**

This function is deprecated, please use [write.report](#) instead.

**Usage**

```
write.report2(...)
```

**Arguments**

... arguments are forwarded to [write.report](#)

**Author(s)**

Jan Philipp Dietrich

**See Also**

[write.report](#)

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