

# Package: gms (via r-universe)

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**Type** Package

**Title** 'GAMS' Modularization Support Package

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**Description** A collection of tools to create, use and maintain modularized model code written in the modeling language 'GAMS' (<<https://www.gams.com/>>). Out-of-the-box 'GAMS' does not come with support for modularized model code. This package provides the tools necessary to convert a standard 'GAMS' model to a modularized one by introducing a modularized code structure together with a naming convention which emulates local environments. In addition, this package provides tools to monitor the compliance of the model code with modular coding guidelines.

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**Suggests** covr, curl, magclass, qgraph, testthat, callr

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

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**Author** Jan Philipp Dietrich [aut, cre] (Potsdam Institute for Climate Impact Research, <<https://orcid.org/0000-0002-4309-6431>>), David Klein [aut] (Potsdam Institute for Climate Impact Research), Anastasis Giannousakis [aut], Felicitas Beier [aut] (Potsdam Institute for Climate Impact Research), Johannes Koch [aut] (Potsdam Institute for Climate Impact Research), Lavinia Baumstark [aut] (Potsdam Institute for Climate Impact

Research), Mika Pflüger [aut], Oliver Richters [aut] (Potsdam  
Institute for Climate Impact Research)

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

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gms-package	<i>gms: 'GAMS' Modularization Support Package</i>
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## Description

A collection of tools to create, use and maintain modularized model code written in the modeling language 'GAMS' (<https://www.gams.com/>). Out-of-the-box 'GAMS' does not come with support for modularized model code. This package provides the tools necessary to convert a standard 'GAMS' model to a modularized one by introducing a modularized code structure together with a naming convention which emulates local environments. In addition, this package provides tools to monitor the compliance of the model code with modular coding guidelines.

## Author(s)

**Maintainer:** Jan Philipp Dietrich <[dietrich@pik-potsdam.de](mailto:dietrich@pik-potsdam.de)> ([ORCID](#)) (Potsdam Institute for Climate Impact Research)

Authors:

- David Klein (Potsdam Institute for Climate Impact Research)
- Anastasis Giannousakis
- Felicitas Beier (Potsdam Institute for Climate Impact Research)
- Johannes Koch (Potsdam Institute for Climate Impact Research)
- Lavinia Baumstark (Potsdam Institute for Climate Impact Research)
- Mika Pflüger
- Oliver Richters (Potsdam Institute for Climate Impact Research)

## See Also

Useful links:

- <https://github.com/pik-piam/gms>
- doi: [10.5281/zenodo.4390032](https://doi.org/10.5281/zenodo.4390032)
- Report bugs at <https://github.com/pik-piam/gms/issues>

---

checkAppearance	<i>checkAppearance</i>
-----------------	------------------------

---

### Description

Checks for all declared objects in which parts of the model they appear and calculates the type of each object (core object, interface object, module object of module xy,...)

### Usage

```
checkAppearance(x, capitalExclusionList = NULL)
```

### Arguments

x	A code list as returned by <a href="#">codeExtract</a>
capitalExclusionList	A vector of names that should be ignored when checking for unified capitalization of variables

### Value

A list with four elements: appearance, setappearance, type and warnings. Appearance is a matrix containing values which indicate whether an object appears in a part of the code or not (e.g. indicates whether "vm\_example" appears in realization "on" of module "test" or not.). 0 means that it does not appear, 1 means that it appears in the code and 2 means that it appears in the not\_used.txt. setappearance contains the same information but for sets instead of other objects. Type is a vector containing the type of each object (excluding sets). And warnings contains a list of warnings created during that process.

### Author(s)

Jan Philipp Dietrich

### See Also

[codeCheck](#), [readDeclarations](#)

---

checkDescription	<i>checkDescription</i>
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---

**Description**

Checks whether all Declarations of a GAMS code come with a Description, throws out a warning in case of a missing description.

**Usage**

```
checkDescription(x, w = NULL)
```

**Arguments**

x	GAMS declarations matrix as returned by
w	a vector of warnings the warnings should be added to <a href="#">readDeclarations</a>

**Value**

vector of warnings

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#)

---

checkNoTabs	<i>checkNoTabs</i>
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---

**Description**

Check all files (also in subdirectories) matching the given pattern for tabs. Will throw an error with a list of files where tabs were found if any.

**Usage**

```
checkNoTabs(pattern, exclude = NULL, excludeFolders = NULL)
```

**Arguments**

pattern	A regular expression. Only files matching this pattern will be checked for tabs.
exclude	A regular expression. Files matching this pattern will never be checked.
excludeFolders	Paths to folders that should not be checked.

**Value**

Invisibly, the list of files that were checked.

**Author(s)**

Pascal Sauer

**Examples**

```
## Not run:
gms::checkNoTabs(pattern = "\\.(R|Rprofile|gms|cfg|bib)$",
                  excludeFolders = c("output", "renv", ".git"))
gms::checkNoTabs(utils::glob2rx("*.R"))

## End(Not run)
```

---

checkSwitchAppearance *checkSwitchAppearance*

---

**Description**

Checks for all declared switches in which parts of the model they appear and calculates the type of each object (core object, interface object, module object of module xy,...)

**Usage**

```
checkSwitchAppearance(code)
```

**Arguments**

code                      Model code returned by [codeExtract](#)

**Value**

A list with three elements: switches, appearance and type. Switches is a vector containing all switches. The names of the vector contain the information where the switch is set. Appearance is a matrix containing values which indicate whether an object appears in a part of the code or not (e.g. indicates whether "vm\_example" appears in realization "on" of module "test" or not.). 0 means that it does not appear, 1 means that it appears in the code and 2 means that it appears in the not\_used.txt. Type is a vector containing the type of each object.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#), [readDeclarations](#), [codeExtract](#), [checkAppearance](#)

---

check_config	<i>Check config</i>
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### Description

Checks a model configuration file for consistency by comparing it to a reference config file and the given module structure of the model. The function will throw out an error if settings are missing in the config which exist in the reference config, or if settings are set in the config which do not exist in the reference config file or if a realization is chosen for a module which does not exist, not allowed setting combinations.

### Usage

```
check_config(  
    icfg,  
    reference_file = "config/default.cfg",  
    modulepath = "modules/",  
    settings_config = NULL,  
    extras = NULL,  
    saveCheck = FALSE  
)
```

### Arguments

icfg	Input config which should be checked for consistency (either as the config itself or as a file path linking to the config)
reference_file	Reference config which is having the right format (either as the config itself or as a file path linking to the config)
modulepath	The path where the modules are stored. If set to NULL the corresponding module check is deactivated.
settings_config	path where the table of possible setting combinations is stored, if NULL it is ignored
extras	vector of setting names that are allowed to appear in the input config even if they are missing in the reference config. That can be useful to allow for additional settings/information which does not necessarily have to exist
saveCheck	additional check which makes sure that saving and loading it to/from YAML format does not change its contents

### Value

The checked config as a config list ready for further usage.

### Author(s)

Jan Philipp Dietrich, Lavinia Baumstark

See Also

[getModules](#)

---

chooseFromList	<i>chooseFromList</i>
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---

Description

Allows the user to select single or multiple items from a list. Entries can be selected based on individual choice, groups, regex or all.

Usage

```
chooseFromList(  
  theList,  
  type = "items",  
  userinfo = NULL,  
  addAllPattern = TRUE,  
  returnBoolean = FALSE,  
  multiple = TRUE,  
  userInput = FALSE,  
  errormessage = NULL  
)
```

Arguments

theList	list or character vector to be selected from, names can specify groups
type	string in plural shown to user to understand what they have to choose
userinfo	string printed to the user before choosing
addAllPattern	boolean whether 'all' and 'Search by pattern' options are added
returnBoolean	TRUE: returns array with dimension of theList with FALSE and TRUE, which erases the order in which entries were selected FALSE: returns selected entries of theList, conserving the order in which entries were selected
multiple	TRUE: allows to select multiple entries. FALSE: no
userinput	string provided by the user. If not supplied, user is asked (mainly for testing)
errormessage	string used internally to tell the user before retrying that a selection does not make sense

Value

list or character vector, either a boolean with same length as theList or only the selected items.

Author(s)

Oliver Richters



Examples

```
## Not run:
chooseFromList(
  theList = c(Letter = "A", Letter = "B", Number = "1", Number = "2"),
  type = "characters",
  userinfo = "Please don't select B, it hurts.",
  returnBoolean = FALSE,
  multiple = TRUE)

## End(Not run)
```

---

codeCheck	<i>codeCheck</i>
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---

Description

Checks GAMS code for consistency. Throws out warnings if something is wrong in the code and returns a list containing the interfaces of each module of the code.

Usage

```
codeCheck(
  path = ".",
  modulepath = "modules",
  core_files = c("core/*.gms", "main.gms"),
  returnDebug = FALSE,
  interactive = FALSE,
  test_switches = TRUE,
  strict = FALSE,
  details = FALSE
)
```

Arguments

path	path of the main folder of the model
modulepath	path to the module folder relative to "path"
core_files	list of files that belong to the core (wildcard expansion is supported)
returnDebug	If TRUE additional information will be returned useful for debugging the codeCheck function
interactive	activates an interactive developer mode in which some of the warnings can be fixed interactively.
test_switches	(boolean) Should realization switches in model core be tested for completeness? Usually set to TRUE but should be set to FALSE for standalone models only using a subset of existing modules

strict	(boolean) test strictness. If set to TRUE warnings from codeCheck will stop calculations at the end of the analysis. Useful to enforce clean code.
details	(boolean) If activated the function will return more detailed output. Besides interface information it will provide a table containing all declarations in the code, an appearance table listing the appearance of all objects in the code and information about the existing modules. The format is list(interfaceInfo,declarations,appearance,modulesInfo). This setting will be ignored when returnDebug is set to TRUE.

### Details

Additional settings can be provided via a yaml file ".codeCheck" in the main folder of the model. Currently supported settings are: - capitalExclusionList: a list of names that should be ignored when checking for unified capitalization of variables

### Value

A list of all modules containing the interfaces for each module. Or more detailed output if either details or returnDebug is set to TRUE.

### Author(s)

Jan Philipp Dietrich

### See Also

[codeExtract](#), [readDeclarations](#)

### Examples

```
# check code consistency of dummy model
codeCheck(system.file("dummymodel", package = "gms"))
```

---

codeExtract	<i>codeExtract</i>
-------------	--------------------

---

### Description

Returns aggregated and cleaned GAMS code together with declaration matrix

### Usage

```
codeExtract(codeFiles, name)
```

### Arguments

codeFiles	A vector of file names of GAMS code files.
name	A name indicating what collection of code files this is (e.g. module name)

**Value**

A list with two elements: code and declarations. Code contains the cleaned up gams code and declarations contains the declarations matrix as returned by [readDeclarations](#)

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#), [readDeclarations](#)

---

`convert.modularGAMS`     *convert.modularGAMS*

---

**Description**

Converts modular GAMS code from an older modular definition to the newest one

**Usage**

```
convert.modularGAMS(path = ".", modulepath = "modules/")
```

**Arguments**

path	path to the main folder of the model
modulepath	Module path within the model (relative to the model main folder)

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#)

---

`copy_input`*copy\_input*

---

**Description**

Function to copy input files to their destination folders

**Usage**

```
copy_input(x, sourcepath, suffix = NULL, move = FALSE)
```

**Arguments**

<code>x</code>	Filepath or data frame containing the mapping of files to be deleted
<code>sourcepath</code>	Path to folder containing all input files
<code>suffix</code>	suffix that might be part of input names that should be deleted
<code>move</code>	If TRUE files will be moved instead of copied (default=FALSE)

**Author(s)**

Jan Philipp Dietrich, David Klein

---

`delete_olddata`*delete\_olddata*

---

**Description**

Delete data provided in mapping

**Usage**

```
delete_olddata(x)
```

**Arguments**

<code>x</code>	Filepath or data frame containing the mapping of files to be deleted
----------------	--

**Author(s)**

Jan Philipp Dietrich, David Klein

---

download\_distribute     *Download and unpack compressed data from repositories*

---

## Description

Downloads a list of tgz files from a list of repos and unpacks them

## Usage

```
download_distribute(
  files,
  repositories = list(`p/projects/rd3mod/inputdata/output` = NULL),
  modelfolder = ".",
  additionalDelete = NULL,
  debug = FALSE,
  stopOnMissing = FALSE
)
```

## Arguments

files	a vector of files containing input data, if a file is no .tgz it will be downloaded and not unpacked
repositories	a list of repositories (please pay attention to the list format!) in which the files should be searched for. Files will be searched in all repositories until found, always starting with the first repository in the list. The argument must have the format of a named list with the url of the repository as name and a corresponding list of options such as username or password to access the repository as value. If no options are required the value has to be NULL. (e.g. <code>list("ftp://my_pw_protected_server.de/data"=list(user="me",password=12345), "http://free_server.de/dat</code>
modelfolder	main model folder
additionalDelete	information which additional data should be deleted before new data are downloaded and distributed
debug	switch for debug mode with additional diagnostic information
stopOnMissing	Boolean passed along to <code>download_unpack()</code> to stop if any file in files could not be downloaded. Off (FALSE) by default.

## Value

Information about the download process in form of a data.frame with data sets as row names and repositories (where it was downloaded from) and corresponding md5sum as columns

## Author(s)

Jan Philipp Dietrich, Lavinia Baumstark

---

download_unpack	<i>Download and unpack compressed data from repositories</i>
-----------------	--

---

### Description

Downloads a list of tgz files from a list of repos and unpacks them. If a file is no .tgz-file it will be only downloaded.

### Usage

```
download_unpack(
  input,
  targetdir = "input",
  repositories = NULL,
  debug = FALSE,
  unpack = TRUE,
  stopOnMissing = FALSE
)
```

### Arguments

input	a vector of files to be downloaded or a cfg list with settings to be used (e.g. containing <code>cfg\$input</code> , <code>cfg\$repositories</code> ). Settings in the config list will be overwritten by other arguments of this function if they are not set to NULL
targetdir	directory the files should be downloaded and extracted to
repositories	a list of repositories (please pay attention to the list format!) in which the files should be searched for. Files will be searched in all repositories until found, always starting with the first repository in the list. The argument must have the format of a named list with the url of the repository as name and a corresponding list of options such as username or password to access the repository as value. If no options are required the value has to be NULL. (e.g. <code>list("ftp://my_pw_protected_server.de/data"=list(user="me",password=12345), "http://free_server.de/dat</code>
debug	switch for debug mode with additional diagnostic information
unpack	if switched off the source files are purley downloaded
stopOnMissing	Boolean indicating whether to stop if any file in <code>files</code> could not be downloaded. Off (FALSE) by default.

### Value

Information about the download process in form of a `data.frame` with data sets as row names and repositories (where it was downloaded from) and corresponding md5sum as columns

### Author(s)

Jan Philipp Dietrich

---

fulldataOutput	<i>fulldataOutput</i>
----------------	-----------------------

---

### Description

Creates GAMS code which stores automatically the levels, bounds and marginals of all equations and variables in time depending parameters.

### Usage

```
fulldataOutput(
  declarations_file = "declarations.gms",
  definitions_file = "postsolve.gms",
  warn = TRUE,
  types = c("level", "marginal"),
  ignore = "_dummy$",
  loopset = "t"
)
```

### Arguments

declarations_file	A GAMS file containing declarations. The function will read declarations from here and add own declarations in an R environment as used by <a href="#">replace_in_file</a> (used subject = OUTPUT DECLARATIONS)
definitions_file	A GAMS file which is executed after the solve statement but within the time step loop. Also here code will be added using <a href="#">replace_in_file</a> with subject OUTPUT DEFINITIONS
warn	Decides whether a warning should be thrown out, if the declarations file does not exist.
types	Types of outputs that should be written to.gdx file. Available types are level, marginal, upper and lower.
ignore	regular expression pattern for variables/equations which should be ignored by fulldataOutput
loopset	Set over which loop runs

### Author(s)

Jan Philipp Dietrich, Felicitas Beier

### See Also

[readDeclarations](#), [replace\\_in\\_file](#)

---

GAMScoreFilter

*GAMScoreFilter*


---

### Description

Cleans GAMS code supplied from empty lines and comments.

### Usage

```
GAMScoreFilter(x)
```

### Arguments

x                      A vector with lines of GAMS code (as you get by reading the code with readLines)

### Value

The cleaned GAMS code

### Author(s)

Jan Philipp Dietrich

### See Also

[readDeclarations](#)

### Examples

```
GAMScoreFilter(c("", "*comment", "a=12;", "", "b=13;"))
```

---

getfiledestinations

*getfiledestinations*


---

### Description

Create file2destination mapping based on information from the model, ignoring top-level directories listed in .gitignore.

### Usage

```
getfiledestinations(path = ".", ignoreFolders = "renv")
```



**Arguments**

path	main model folder
ignoreFolders	folders to be ignored by the function, additionally to directories listed in <code>.gitignore</code> (by default only "renv").

**Author(s)**

Jan Philipp Dietrich, David Klein

---

getLine	<i>getLine</i>
---------	----------------

---

**Description**

Get one line of user input regardless of whether running interactively or not (via Rscript). (base::readline does not wait for user input when running via Rscript.)

**Usage**

```
getLine()
```

**Value**

The user input as a string.

---

getModules	<i>getModules</i>
------------	-------------------

---

**Description**

Extract module information of a GAMS model.

**Usage**

```
getModules(modulepath)
```

**Arguments**

modulepath	The path where the modules are stored.
------------	--

**Value**

A matrix containing the different modules with name, corresponding module number and corresponding realizations

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#)

---

get_info	<i>get_info</i>
----------	-----------------

---

**Description**

Function to extract information from info.txt

**Usage**

```
get_info(file, grep_expression, sep, pattern = "", replacement = "")
```

**Arguments**

- |                 |  |
|-----------------|--|
| file            | path to info.txt (including info.txt)                  |
| grep_expression | String before the information that should be extracted |
| sep             | Separator between grep_expression and information      |
| pattern         | Optional pattern to be replaced (default pattern = "") |
| replacement     | Optional replacement (default replacement = "")        |

**Author(s)**

Jan Philipp Dietrich, David Klein

---

interfaceplot	<i>interfaceplot</i>
---------------	----------------------

---

**Description**

Creates an interface plot of a modular model using [qgraph](#) and returns the interface information.

**Usage**

```

interfaceplot(
  x = ".",
  modules_to_include = NULL,
  modules_to_exclude = NULL,
  links_to_include = NULL,
  links_to_exclude = NULL,
  items_to_include = NULL,
  items_to_exclude = NULL,
  items_to_display = NULL,
  default_groups = list(default1 = list(name = "core", nodes = "core", color = "black",
    shape = "rectangle"), default2 = list(name = "modules", nodes = NULL, color =
    "#6c9ebf", shape = "ellipse")),
  highlight_groups = NULL,
  max_length_node_names = NULL,
  add_nodeName_legend = FALSE,
  max_num_edge_labels = NULL,
  max_num_nodes_for_edge_labels = 30,
  ...
)

```

**Arguments**

**x** Either an interface list as returned by [codeCheck](#) or the path to the main folder of the model.

**modules\_to\_include** NULL (default value) or a vector of strings with names of modules to include, e.g. `c("core", "macro")`. If NULL all modules are included.

**modules\_to\_exclude** NULL (default value) or a vector of strings with names of modules to exclude, e.g. `c("core")`. If NULL no modules are excluded.

**links\_to\_include** NULL (default value) or list of lists with attributes "to" and "from", that each take a vector of module names, e.g. `list(list(to="macro", from="core"))`. If NULL all links are included.

**links\_to\_exclude** NULL (default value) or list of lists with attributes "to" and "from", that each take a vector of module names, e.g. `list(list(to="macro", from="core"))`. If NULL no links are excluded.

**items\_to\_include** NULL (default value) or a vector of strings with names of items to include, e.g. `c("vm_cesIO", "pm_pvp")`. Regex patterns can also be passed, e.g. `c("(vlp)m_.*")`. If NULL all items are included.

**items\_to\_exclude** NULL (default value) or a vector of strings with names of items to exclude, e.g. `c("vm_cesIO", "pm_pvp")`. Regex patterns can also be passed, e.g. `c("sm_.*")`. If NULL no items are excluded.

- `items_to_display` NULL (default value) or a vector of strings with names of items to display, e.g. `c("vm_cesIO", "pm_pvp")`. Regex patterns can also be passed, e.g. `c(.m\ S+)`. If NULL no items are displayed.
- `default_groups` List of lists with default group definitions. Defines the default formatting of the interface plot. By default, there are two groups, see usage, a "core" group made up of only the "core" module and a "modules" group made up of all the rest. If a "core" module doesn't exist, then that group is simply ignored.
- `highlight_groups` NULL (default value) or a list of lists with highlight-group definitions. By defining highlight groups, additional/specialized formatting can be applied to select modules. A group is defined by a list with the following attributes:
- `name`: a string with the group name. Will appear in legend.
  - `nodes`: a vector of strings with module names.
  - `shape`: a string with a valid qgraph shape.
  - `color`: a string with a valid qgraph color.
  - `edges_to_highlight`:
    - NULL = no edges are colored
    - "all" = edges starting from and ending at the highlight group's nodes are colored
    - "incoming" = edges ending at the highlight group's nodes are colored
    - "outgoing" = edges starting from the highlight group's nodes are colored
    - "within" = only edges that departed from the highlight group's nodes and end at them as well are colored
  - `edges_to_ignore`:
    - NULL = no edges are ignored
    - "outside" edges that neither start from or end at any of the group's nodes are ignored
    - "incoming" edges that do not depart from one of the group's nodes are ignored
    - "outgoing" edges that do not arrive at one of the group's nodes are ignored
    - "outgoing\_to\_no\_return" edges that departing from nodes outside of the group and not ending at nodes within the group are ignored
- An example: `list(list(name = "highlight", nodes = "welfare", color = "#ff8f00", shape = "ellipse", edges_to_highlight = "outgoing", edges_to_ignore = "outside"))`.
- `max_length_node_names` NULL (default value) or an integer `n` giving the maximum number of characters allowed in the node names. If not NULL, node names are truncated after `n` characters, e.g. `n=3: "example" -> "exa."`.
- `add_nodeName_legend` Logical (default FALSE) to add node names in legend, structured by group.

`max_num_edge_labels`  
 NULL (default value), an integer or the string "adjust". If NULL, all edge labels are displayed. If given an integer n, a maximum of n edge labels are shown. If set to "adjust", the number of edge labels displayed decreases with the number of nodes.

`max_num_nodes_for_edge_labels`  
 Integer, (default value = 30). The maximum number of nodes after which no edge labels are displayed.

... Optional arguments to [qgraph](#).

## Details

What modules (=nodes), links (=edges) and items (=what is passed along the edges) are taken into account when creating the plot can be fine-tuned with the "`_include`", "`_exclude`" arguments.

The "`default`"- and "`highlight_groups`" arguments control the formatting (and also the composition through "`highlight_group$edges_to_ignore`"). Groups in `qqgraph` are a way of clustering nodes together. The default formatting of the plot is defined with the "`default_groups`" argument. On top of that additional groups can be defined with the "`highlight_groups`" argument.

The rest of the arguments are pretty self-explanatory. Just remember that [qgraph](#) arguments can be passed on as well! Useful ones include: `fade=T/F`, `legend=T/F`, `legend.cex` (size of the legend font), `GLratio` (graph/legend size ratio, `edge.label.cex` (size of the edge label font)).

## Value

A tibble with the edge list and interface items.

## Author(s)

Johannes Koch

## See Also

[codeCheck](#), [qgraph](#)

---

is.modularGAMS

is.modularGAMS

---

## Description

Checks whether a folder seems to contain modular GAMS code or not.

## Usage

```
is.modularGAMS(path = ".", version = FALSE, modulepath = "modules/")
```

**Arguments**

path	path to the main folder of the model
version	if TRUE returns the version of the modular structure or FALSE, otherwise returns a boolean indicating whether it is modular or not.
modulepath	Module path within the model (relative to the model main folder)

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#)

**Examples**

```
is.modularGAMS(system.file("dummymodel", package="gms"))
```

---

loadConfig

*Load Config*

---

**Description**

Load config in YAML format as written via [saveConfig](#).

**Usage**

```
loadConfig(cfg)
```

**Arguments**

cfg	Either a character string naming a file which contains the config or a character string containing the config as YAML code.
-----	---

**Details**

To read in the default configuration (stored as R list in default.cfg or in a gams file), use [readDefaultConfig](#) instead.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[saveConfig](#)

### Examples

```
cfg <- list(input = c(data1 = "blub.tgz", data2 = "bla.tgz"), mode = "default")
yaml <- saveConfig(cfg)
loadConfig(yaml)
```

---

model_dependencies	<i>Function to detect R package dependencies</i>
--------------------	--

---

### Description

This function analyzes a model folder and all subfolders and searches for library and require statements.

### Usage

```
model_dependencies(mainfolder = ".")
```

### Arguments

mainfolder      main folder of the model to be analyzed

### Value

A list of dependencies sorted by appearances

### Author(s)

Jan Philipp Dietrich

---

model_lock	<i>Model lock/unlock</i>
------------	--------------------------

---

### Description

Lock the model to coordinate multiple processes accessing the model. This is necessary if you change files within the model and have to avoid that another process reads from your half-written changes in the mean time. `model_lock` creates an exclusive lock on the given file and returns a lock id. When the lock id is handed to `model_unlock` the lock is dropped again, and another process can take the lock. The lock is also dropped when the lock id goes out of scope and is garbage collected in R or when the R process terminates or crashes. However, it is good practice to explicitly call `model_unlock` when you don't need the lock anymore. If you want to check for informational purposes if the model is currently locked by another process, you can use the `is_model_locked()` function. Note however that you should never use `is_model_locked()` to coordinate access to the model folder. The return value of `is_model_locked()` is only a snapshot at the time when the function is run. Note additionally that `is_model_locked()` checks if the model is locked by another process. If you currently hold the lock of the model yourself, it will return `FALSE`.

**Usage**

```
model_lock(folder = ".", file = ".lock", timeout1 = 12, timeout2 = NULL,  
check_interval = NULL, oncluster = NULL)  
model_unlock(id, folder = NULL, file = NULL, oncluster = NULL)  
is_model_locked(folder = ".", file = ".lock")
```

**Arguments**

folder	path to the model folder
file	file name of the lock file. Don't use the lock file for anything else (i.e. don't read or write or open the lock file).
timeout1	Time in hours to wait for other processes to free the lock before giving up and exiting with an error.
timeout2	deprecated setting which will be ignored.
check_interval	deprecated setting which will be ignored.
oncluster	deprecated setting which will be ignored.
id	lock id as returned by model_lock.

**Value**

model\_lock returns the lock id which is needed to identify the lock in model\_unlock.

**Author(s)**

Jan Philipp Dietrich, David Klein, Mika Pflüger

**See Also**

[check\\_config](#)

**Examples**

```
#lock folder  
id <- model_lock(tempdir())  
  
#unlock folder  
model_unlock(id)
```



---

module.skeleton	<i>Create a Module skeleton</i>
-----------------	---------------------------------

---

## Description

This function creates you a module skeleton which you can use to easily create your own modules.

## Usage

```
module.skeleton(
    number,
    name,
    types,
    modelpath = ".",
    modulepath = "modules/",
    includefile = "modules/include.gms",
    version = is.modularGAMS(modelpath, version = TRUE)
)
```

## Arguments

number	Number of your module, typically something between 0-99. Sorts the execution of your modules. Please use a number which is not used, yet.
name	Name of your module (please choose a short name). If you want to extend an existing module (add a new realisation) use the name of the existing one.
types	Vector of names for the different module types (e.g. "on" or "off"). If you want to extend an existing module (add a new realisation), put here the additional type(s)
modelpath	Path of the MAgPIE version that should be updated (main folder).
modulepath	Module path within MAgPIE (relative to the MAgPIE main folder)
includefile	Name and location of the file which includes all modules (relative to main folder)
version	version of the modular GAMS code structure (1 or 2)

## Note

Module phases are automatically detected checking the main code of the model, but not checking code in modules. If you want to use additional phases which are only included within a module, you need to specify them manually by adding a comment into your gams code indicating that there is an additional phase. The syntax is `"* !add_phase!: <phase>"`, e.g. `"* !add_phase!: new_phase"`

## Author(s)

Jan Philipp Dietrich

**Examples**

```
# copy dummymodel to temporary directory and add new module "bla"
file.copy(system.file("dummymodel",package="gms"),tempdir(), recursive = TRUE)
model <- paste0(tempdir(),"/dummymodel")
module.skeleton(number="03", name="bla", types=c("on","off"), modelpath=model)
```

---

modules\_interfaceplot *modules\_interfaceplot*

---

**Description**

Function that applies [interfaceplot](#) for a whole model and all its modules.

**Usage**

```
modules_interfaceplot(
  x = ".",
  modulepath = "modules",
  filetype = "png",
  targetfolder = NULL,
  writetable = TRUE,
  includeCore = FALSE,
  ...
)
```

**Arguments**

x	Either the object returned by <a href="#">codeCheck</a> or the path to the main folder of the model.
modulepath	Path to the modules folder
filetype	Filetype that should be used (e.g. "png" or "pdf")
targetfolder	Folder outputs should be written to. If set to NULL outputs will be added to corresponding module folders
writetable	Logical deciding whether a csv containing the interfaces should be written as well.
includeCore	Logical to create plot for core or not, default FALSE.
...	Optional arguments to <a href="#">interfaceplot</a> .

**Value**

A list with interface tables for each module

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#), [interfaceplot](#)

---

path	<i>path</i>
------	-------------

---

**Description**

Small function to build a consistent path-string based on folder, filename and filetype. The function makes sure that slashes and the dot for the file ending are set correctly (you can supply your folder name either with or without a trailing slash in it. It does not matter.

**Usage**

```
path(..., ftype = NULL)
```

**Arguments**

...	the folders and the file name that should be pasted to a file/folder path
ftype	file type

**Value**

A string containing the path combined of folder, filename and filetype

**Author(s)**

Jan Philipp Dietrich

---

publish_data	<i>Publish data in a repository</i>
--------------	-------------------------------------

---

**Description**

Downloads a list of tgz files from a list of repos, merge them and publish it on another server

**Usage**

```
publish_data(
  input,
  name = NULL,
  target = Sys.getenv("PUBLISH_DATA_TARGET", unset = "."),
  ...
)
```

**Arguments**

input	a vector of files to be downloaded or a cfg list with settings to be used (e.g. containing <code>cfg\$input</code> , <code>cfg\$repositories</code> ). Settings in the config list will be overwritten by other arguments of this function if they are not set to NULL
name	name of the data to be published (will be used in as file name). If no name is given (default) source files will be published as is (separate tgz files with original name).
target	target the data should be published in (format <code>user@server:/folder/path</code> ) If a target vector, or targets separated by " " are provided the user will be asked interactively where the file should be written to. By default it will look for target information in the environment variable <code>PUBLISH_DATA_TARGET</code>
...	further options provided to <a href="#">download_unpack</a>

**Author(s)**

Jan Philipp Dietrich

**See Also**

[download\\_unpack](#), [tardir](#)

---

readDeclarations	<i>readDeclarations</i>
------------------	-------------------------

---

**Description**

Reads all declarations given in a GAMS code and returns them.

**Usage**

```
readDeclarations(
  file,
  unlist = TRUE,
  types = c("scalar", "(positive |)variable", "parameter", "table", "equation", "set")
)
```

**Arguments**

file	A gams file or a vector containing GAMS code.
unlist	A logical indicating whether the output should be returned as a list separated by object type or a matrix.
types	of declarations to be read.

**Value**

Either a list of declared objects or a matrix containing object name, the sets the object depends on and the description.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#)

---

<code>readDefaultConfig</code>	<i>readDefaultConfig</i>
--------------------------------	--------------------------

---

**Description**

Reads the default configuration of the model. Uses `default.cfg` or `main.cfg` as the source as appropriate. To read a configuration from YAML format, use [loadConfig](#) instead.

**Usage**

```
readDefaultConfig(path)
```

**Arguments**

<code>path</code>	path of the main folder of the model
-------------------	--------------------------------------

**Value**

A vector of parameter values and their names.

**Author(s)**

Mika Pflüger

**See Also**

[loadConfig](#)

---

readFileOrVector	<i>readFileOrVector</i>
------------------	-------------------------

---

**Description**

Input: a file name or a character vector. Output: a character vector of the contents of the file, or the character vector unchanged.

**Usage**

```
readFileOrVector(file)
```

**Arguments**

file	A file name or a character vector.
------	------------------------------------

**Value**

A character vector of the inputs of the file.

**Author(s)**

Mika Pflüger

---

readParameterValues	<i>readParameterValues</i>
---------------------	----------------------------

---

**Description**

Read values of given parameters from a given GAMS code section.

**Usage**

```
readParameterValues(code, parameters)
```

**Arguments**

code	A vector with GAMS code.
parameters	A vector with GAMS parameter names

**Details**

The GAMS code section should contain statements of the form parameter = value; for all the given parameters.

**Value**

A vector of values the parameters are set to with parameter names as names.

**Author(s)**

Mika Pflüger

**See Also**

[readDeclarations](#)

---

readSetglobals	<i>readSetglobals</i>
----------------	-----------------------

---

**Description**

Reads all setglobals given in a GAMS code and returns them.

**Usage**

```
readSetglobals(file)
```

**Arguments**

file	A gams file or a vector containing GAMS code.
------	---

**Value**

A vector of values the setglobal variables are set to with setglobal variables as names.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[readDeclarations](#)

---

readSettings	<i>readSettings</i>
--------------	---------------------

---

**Description**

Reads all settings (parameters, globals, and their respective values) in the given GAMS code.

**Usage**

```
readSettings(file)
```

**Arguments**

file	A gams file or a vector containing GAMS code.
------	---

**Value**

A vector of parameter values and their names.

**Author(s)**

Mika Pflüger

**See Also**

[readDeclarations](#)

---

read_yaml_header	<i>read_yaml_header</i>
------------------	-------------------------

---

**Description**

Reads header written in yaml format from a file

**Usage**

```
read_yaml_header(file, n = 20)
```

**Arguments**

file	path to the file which contains the YAML header
n	Number of lines to be read (header must be part of these lines in order to be read)

**Value**

A list containing the read in information



**Author(s)**

Jan Philipp Dietrich

---

replace_in_file	<i>Replace in File</i>
-----------------	------------------------

---

**Description**

Function to replace a marked paragraph in a text file. Paragraph has to be marked in the text file with an initial "##### R SECTION START (SUBJECT) #####" and "##### R SECTION END (SUBJECT) #####" as ending. The number of # symbols can be chosen by the user, but there has to be at least one at the beginning and one at the end. Furthermore it is allowed to add further symbols at the beginning or the end of the line. "SUBJECT" is chosen by the user and is used for identification, if a text file has more than one R section.

**Usage**

```
replace_in_file(
  file,
  content,
  subject = "CODE",
  add = FALSE,
  addfile = FALSE,
  comment = "*"
)
```

**Arguments**

file	a connection object or a character string describing the file, that should be manipulated.
content	the content that should be used as replacement stored as a vector of strings. Each vector component will be written as a line.
subject	A string used for identification of a paragraph.
add	Determines behavior when marking is missing in the code. add=FALSE will throw out an error, if the marking is missing, add="top" will add the markings automatically at the beginning of the file, add="bottom" or add=TRUE will do the same but at the end of the file.
addfile	Determines the behavior when the file does not exist. If addfile=TRUE, file will be created when missing.
comment	Symbol which is used to indicate a comment in the language the file is written that should be manipulated. Only relevant if add or addfile are used.

**Author(s)**

Jan Philipp Dietrich

---

saveConfig	<i>Save Config</i>
------------	--------------------

---

**Description**

Saves config in YAML format so that it can be read in again via [loadConfig](#).

**Usage**

```
saveConfig(cfg, file = NULL)
```

**Arguments**

cfg	Input config which should be saved
file	A character string naming a file. If set to NULL the YAML output will be returned by the function.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[loadConfig](#)

**Examples**

```
cfg <- list(input = c(data1 = "blub.tgz", data2 = "bla.tgz"), mode = "default")
saveConfig(cfg)
```

---

selectScript	<i>selectScript</i>
--------------	---------------------

---

**Description**

Functions which allows for interactive selection of scripts/files.

**Usage**

```
selectScript(folder = ".", ending = "R")
```

**Arguments**

folder	Folder in which the files/scripts are located which should be selected from.
ending	File ending of the files to be selected (without dot)

**Value**

A vector of paths to files selected by the user

**Author(s)**

Jan Philipp Dietrich

---

setScenario

*setScenario*

---

**Description**

setScenario is adapting a given config to a predefined scenario, meaning that all settings which are fixed for the given scenario are written to the config. Settings not defined by the scenario remain unchanged.

**Usage**

```
setScenario(cfg, scenario, scenario_config = "config/scenario_config.csv")
```

**Arguments**

cfg	Input config which should be adapted to the given scenario
scenario	name of scenario (e.g. "SSP2"). Can also be a vector of scenarios. In this case scenario settings are applied in the given order
scenario_config	The path where the scenario config table is stored.

**Value**

The updated config as a config list ready for further usage.

**Note**

The scenario config table is a table which contains as columns the different scenarios and as rows the different settings. Empty entries for a given scenario-setting combination indicate that this setting is not defined by the scenario and should not be changed by set Scenario!

**Author(s)**

Jan Philipp Dietrich, Anastasis Giannousakis

**See Also**

[check\\_config](#), [getModules](#)

---

settingsCheck	<i>settingsCheck</i>
---------------	----------------------

---

**Description**

Checks GAMS setglobals in code for consistency. Creates a warning if a setglobal command for an existing module is missing or a module is set to a realization which does not exist.

**Usage**

```
settingsCheck(path = ".", modulepath = "modules", fileName = "main.gms")
```

**Arguments**

path	path of the main folder of the model
modulepath	path to the module folder relative to "path"
fileName	name of the file containing setglobals, relative to "path"

**Value**

Nothing is returned.

**Author(s)**

Jan Philipp Dietrich

**See Also**

[codeCheck](#)

---

singleGAMSfile	<i>Merge GAMS code into single file</i>
----------------	---

---

**Description**

This function merges GAMS code which is distributed over several files into a single GAMS file. Optionally, it also embeds R scripts into the single GAMS file

**Usage**

```
singleGAMSfile(
  modelpath = ".",
  mainfile = "main.gms",
  output = "full.gms",
  embedRScripts = FALSE
)
```

**Arguments**

modelpath	The path where the model is stored
mainfile	The path to the main gams file (relative to the model path)
output	Name of the single output GAMS file.
embedRScripts	If TRUE, R scripts called by GAMS via Execute are also embedded. Default FALSE

**Author(s)**

Jan Philipp Dietrich, Anastasis Giannousakis

**Examples**

```
# copy dummymodel create single gms file out of it
file.copy(system.file("dummymodel", package = "gms"), tempdir(), recursive = TRUE)
model      <- paste0(tempdir(), "/dummymodel")
singlefile <- paste0(tempdir(), "/full.gms")
singleGAMSfile(modelpath = model, output = singlefile)
```

---

tardir	<i>Create tgz archive from directory</i>
--------	--

---

**Description**

Creates a tgz from all files in a directory

**Usage**

```
tardir(dir = ".", tarfile = "data.tgz")
```

**Arguments**

dir	directory from which the tar file should be generated
tarfile	name of the archive the data should be written to (tgz file)

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
# copy dummymodel to temporary directory and compress it
file.copy(system.file("dummymodel", package = "gms"), tempdir(), recursive = TRUE)
model <- paste0(tempdir(), "/dummymodel")
archive <- paste0(tempdir(), "/dummymodel.tgz")
tardir(model, archive)
```

---

```
updateInterfaceMapping
    updateInterfaceMapping
```

---

### Description

Function to update the mapping between interfaces and their origin modules.

### Usage

```
updateInterfaceMapping(path = ".", modulepath = "modules")
```

### Arguments

path	path of the main folder of the model
modulepath	path to the module folder relative to "path"

### Author(s)

Jan Philipp Dietrich

### See Also

[codeCheck](#)

---

```
update_fulldataOutput  update_fulldataOutput
```

---

### Description

Creates GAMS code which stores automatically the levels and marginals of all equations and variables in time depending parameters.

### Usage

```
update_fulldataOutput(
  modelpath = ".",
  modulepath = "modules",
  corepath = "core",
  loopset = "t"
)
```

**Arguments**

modelpath	Path of the Model version that should be updated (main folder).
modulepath	Module path within the model (relative to the model main folder)
corepath	Core path within the model (relative to the model main folder)
loopset	Set over which loop runs

**Author(s)**

Jan Philipp Dietrich, Felicitas Beier

**See Also**

[fulldataOutput,replace\\_in\\_file](#)

---

update\_modules\_embedding

*Update Modules Embedding in GAMS code*

---

**Description**

A function that updates in the GAMS code all include commands which are related to Modules. The function automatically checks which modules exist and which files in these modules exist and creates the corresponding include commands in GAMS

**Usage**

```
update_modules_embedding(
    modelpath = ".",
    modulepath = "modules/",
    includefile = "modules/include.gms",
    verbose = FALSE
)
```

**Arguments**

modelpath	Path to the model that should be updated (main folder).
modulepath	Module path within the model (relative to the model main folder)
includefile	Name and location of the file which includes all modules (relative to main folder)
verbose	Defines whether additional information should be printed or not.

**Note**

Module phases are automatically detected checking the main code of the model, but not checking code in modules. If you want to use additional phases which are only included within a module, you need to specify them manually by adding a comment into your gams code indicating that there is an additional phase. The syntax is `"* !add_phase!: <phase>"`, e.g. `"* !add_phase!: new_phase"`

**Author(s)**

Jan Philipp Dietrich

**Examples**

```
# copy dummymodel to temporary directory and update module embedding
file.copy(system.file("dummymodel", package="gms"), tempdir(), recursive = TRUE)
model <- paste0(tempdir(), "/dummymodel")
update_modules_embedding(model)
```

---

writeSets

*writeSets*

---

**Description**

Function to write a set declaration in GAMS syntax given either a character vector or a data.frame (latter to represent a mapping set)

**Usage**

```
writeSets(sets, file = NULL)
```

**Arguments**

sets	a list of sets where every set is another list consisting of three entries: name (set name as character), desc (set description as character) and items (set elements, either as character vector or data.frame)
file	Name of the file the set declarations should be written to. If NULL the result will just be returned by the function. The file must exist and contain a line of the form "##### R SECTION START (SETS) #####" followed by a line of the form ##### R SECTION END (SETS) ##### to indicate the part of the file that should be replaced with the set declaration!

**Value**

set declaration in GAMS syntax

**Author(s)**

Jan Philipp Dietrich, Florian Humpenöder



**Examples**

```
countries <- c("DEU", "FRA", "ENG", "ITA", "NLD", "POL")
map <- data.frame(region = rep(c("REG1", "REG2"), 3), countries = countries)
sets <- list(list(name = "countries",
                 desc = "list of countries",
                 items = countries),
            list(name = "country2region",
                 desc = "mapping between countries and regions",
                 items = map))
cat(writeSets(sets), sep="\n")
```

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