

# Package ‘icesTAF’

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**Description** Functions to support the ICES Transparent Assessment Framework  
<<https://taf.ices.dk>> to organize data, methods, and results used in ICES  
assessments. ICES is an organization facilitating international collaboration  
in marine science.

**License** GPL-3

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### Description

Functions to support the ICES Transparent Assessment Framework, to organize data, methods, and results used in ICES assessments.

### Details

#### *Initial TAF steps:*

<code>draft.data</code>	draft DATA.bib file
<code>draft.software</code>	draft SOFTWARE.bib file
<code>period</code>	paste period string for DATA.bib
<code>taf.bootstrap</code>	set up data files and software
<code>taf.skeleton</code>	create empty TAF template

#### *Running scripts:*

<code>clean</code>	clean TAF directories
<code>make</code>	run R script if needed
<code>makeAll</code>	run all TAF scripts as needed
<code>makeTAF</code>	run TAF script if needed
<code>msg</code>	show message
<code>sourceAll</code>	run all TAF scripts
<code>sourceTAF</code>	run TAF script

#### *File management:*

<code>convert.spaces</code>	convert spaces
<code>cp</code>	copy files

<code>mkdir</code>	create directory
<code>os.linux</code>	operating system
<code>os.macos</code>	operating system
<code>os.windows</code>	operating system
<code>read.taf</code>	read TAF table from file
<code>sourceDir</code>	read all *.R files
<code>taf.library</code>	load package from TAF library
<code>taf.unzip</code>	unzip file
<code>write.taf</code>	write TAF table to file

*Tables:*

<code>div</code>	divide column values
<code>flr2taf</code>	convert FLR to TAF
<code>long2taf</code>	convert long format to TAF
<code>long2xtab</code>	convert long format to crosstab
<code>plus</code>	rename plus group column
<code>rnd</code>	round column values
<code>sam2taf</code>	convert SAM to TAF
<code>taf2long</code>	convert TAF to long format
<code>taf2xtab</code>	convert TAF to crosstab
<code>tt</code>	transpose TAF table
<code>xtab2long</code>	convert crosstab to long format
<code>xtab2taf</code>	convert crosstab to TAF

*Plots:*

<code>lim</code>	compute axis limits
<code>taf.colors</code>	predefined colors
<code>taf.png</code>	open PNG graphics device
<code>zoom</code>	change lattice text size

*Example tables:*

<code>catage.long</code>	long format
<code>catage.taf</code>	TAF format
<code>catage.xtab</code>	crosstab format
<code>summary.taf</code>	summary results

*Administrative tools, rarely used in scripts:*

<code>clean.data</code>	clean bootstrap data
<code>clean.library</code>	clean TAF library
<code>clean.software</code>	clean TAF software
<code>deps</code>	list dependencies
<code>detach.packages</code>	detach all packages
<code>dos2unix</code>	convert line endings

<code>download</code>	download file
<code>download.github</code>	download repository
<code>file.encoding</code>	examine file encoding
<code>get.remote.sha</code>	look up SHA code
<code>is.r.package</code>	check if file is an R package
<code>latin1.to.utf8</code>	convert file encoding
<code>line.endings</code>	examine line endings
<code>rmdir</code>	remove empty directory
<code>taf.install</code>	install package in TAF library
<code>taf.libPaths</code>	add TAF library to search path
<code>taf.session</code>	show session information
<code>unix2dos</code>	convert line endings
<code>utf8.to.latin1</code>	convert file encoding

**Author(s)**

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**References**

ICES Transparent Assessment Framework: <https://taf.ices.dk>.

To explore example TAF stock assessments, see the introductory [video](#) and [tutorial](#).

The [TAF Wiki](#) provides additional help resources.

---

catage.long

*Catch at Age in Long Format*

---

**Description**

Small catch-at-age table to describe a long format data frame to store year-age values.

**Usage**

`catage.long`

**Format**

Data frame containing three columns:

Year	year
Age	age
Catch	catch (millions of individuals)

**Details**

The data are an excerpt (first years and ages) from the catch-at-age table for North Sea cod from the ICES (2016) assessment.

**Source**

ICES (2016) Report of the working group on the assessment of demersal stocks in the North Sea and Skagerrak (WGNSSK). *ICES CM 2016/ACOM:14*, p. 673.

**See Also**

[catage.taf](#) and [catage.xtab](#) describe alternative table formats.

[long2taf](#) converts a long table to TAF format.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
catage.long
long2taf(catage.long)
```

---

catage.taf	<i>Catch at Age in TAF Format</i>
------------	-----------------------------------

---

**Description**

Small catch-at-age table to describe a TAF format data frame to store year-age values.

**Usage**

```
catage.taf
```

**Format**

Data frame containing five columns:

Year	year
1	number of one-year-olds in the catch (millions)
2	number of two-year-olds in the catch (millions)
3	number of three-year-olds in the catch (millions)
4	number of four-year-olds in the catch (millions)

**Details**

The data are an excerpt (first years and ages) from the catch-at-age table for North Sea cod from the ICES (2016) assessment.

**Source**

ICES (2016) Report of the working group on the assessment of demersal stocks in the North Sea and Skagerrak (WGNSSK). *ICES CM 2016/ACOM:14*, p. 673.

**See Also**

[catage.long](#) and [catage.xtab](#) describe alternative table formats.  
[taf2long](#) and [taf2xtab](#) convert a TAF table to alternative formats.  
[icesTAF-package](#) gives an overview of the package.

**Examples**

```
catage.taf
taf2long(catage.taf)
taf2xtab(catage.taf)
```

---

catage.xtab	<i>Catch at Age in Crosstab Format</i>
-------------	--

---

**Description**

Small catch-at-age table to describe a crosstab format data frame to store year-age values.

**Usage**

```
catage.xtab
```

**Format**

Data frame with years as row names and containing four columns:

- 1 number of one-year-olds in the catch (millions)
- 2 number of two-year-olds in the catch (millions)
- 3 number of three-year-olds in the catch (millions)
- 4 number of four-year-olds in the catch (millions)

**Details**

The data are an excerpt (first years and ages) from the catch-at-age table for North Sea cod from the ICES (2016) assessment.

**Source**

ICES (2016) Report of the working group on the assessment of demersal stocks in the North Sea and Skagerrak (WGNSSK). *ICES CM 2016/ACOM:14*, p. 673.

**See Also**

[catage.long](#) and [catage.taf](#) describe alternative table formats.  
[xtab2taf](#) converts a crosstab table to TAF format.  
[icesTAF-package](#) gives an overview of the package.

## Examples

```
catage.xtab  
xtab2taf(catage.xtab)
```

---

clean	<i>Clean TAF Directories</i>
-------	------------------------------

---

## Description

Remove working TAF directories (data, model, output, report), bootstrap, or other directories.

## Usage

```
clean(dirs = c("data", "model", "output", "report"), force = FALSE)
```

## Arguments

dirs	directories to delete.
force	passed to software and clean.library if any of the dirs is "bootstrap".

## Note

The purpose of removing the directories is to make sure that subsequent TAF scripts start by creating new empty directories.

If any of the dirs is "bootstrap" it is treated specially. Instead of completely removing the bootstrap directory, only the subdirectories data is removed, while clean.software and clean.library are used to clean the bootstrap/software and bootstrap/library subdirectories. This protects the subdirectory bootstrap/initial and \*.bib metadata files from being accidentally deleted.

## See Also

[clean.software](#) selectively removes software from bootstrap/software.

[clean.library](#) selectively removes packages from bootstrap/library.

[clean.data](#) selectively removes data from bootstrap/data.

[mkdir](#) and [rmdir](#) create and remove empty directories.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:  
clean()  
  
## End(Not run)
```



---

`clean.data`*Clean Data*

---

**Description**

Selectively remove data from the bootstrap/data folder if not listed in DATA.bib.

**Usage**

```
clean.data(folder = "bootstrap/data", quiet = FALSE, force = FALSE)
```

**Arguments**

folder	location of bootstrap/data.
quiet	whether to suppress messages about removed data.
force	whether to remove folder, regardless of how it compares to DATA.bib entries.

**Note**

For each data file or subfolder, the cleaning procedure selects between two cases:

1. Data entry found in DATA.bib - do nothing.
2. Data entry is not listed in DATA.bib - remove.

The `taf.bootstrap` procedure cleans the bootstrap/data folder, without requiring the user to run `clean.data`.

**See Also**

[taf.bootstrap](#) calls `clean.data` as part of the default bootstrap procedure.

[clean.software](#) cleans the local TAF software folder.

[clean.library](#) cleans the local TAF library.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
clean.data()  
  
## End(Not run)
```

clean.library

*Clean TAF Library*

---

**Description**

Selectively remove packages from the local TAF library if not listed in SOFTWARE.bib.

**Usage**

```
clean.library(folder = "bootstrap/library", quiet = FALSE, force = FALSE)
```

**Arguments**

folder	location of local TAF library.
quiet	whether to suppress messages about removed packages.
force	whether to remove the local TAF library, regardless of how it compares to SOFTWARE.bib entries.

**Note**

For each package, the cleaning procedure selects between three cases:

1. Installed package matches SOFTWARE.bib - do nothing.
2. Installed package is not the version listed in SOFTWARE.bib - remove.
3. Installed package is not listed in SOFTWARE.bib - remove.

The taf.bootstrap procedure cleans the TAF library, without requiring the user to run clean.library. The main reason for a TAF user to run clean.library directly is to experiment with installing and removing different versions of software without modifying the SOFTWARE.bib file.

**See Also**

[taf.bootstrap](#) calls clean.library as part of the default bootstrap procedure.

[taf.install](#) installs a package in the local TAF library.

[clean.software](#) cleans the local TAF software folder.

[clean.data](#) cleans the bootstrap/data folder.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
clean.library()  
  
## End(Not run)
```

---

clean.software	<i>Clean TAF Software</i>
----------------	---------------------------

---

### Description

Selectively remove software from the local TAF software folder if not listed in SOFTWARE.bib.

### Usage

```
clean.software(folder = "bootstrap/software", quiet = FALSE, force = FALSE)
```

### Arguments

folder	location of local TAF software folder.
quiet	whether to suppress messages about removed software.
force	whether to remove the local TAF software folder, regardless of how it compares to SOFTWARE.bib entries.

### Note

For each file (and subdirectory) in the software folder, the cleaning procedure selects between three cases:

1. File and version matches SOFTWARE.bib - do nothing.
2. Filename does not contain the version listed in SOFTWARE.bib - remove.
3. File is not listed in SOFTWARE.bib - remove.

The taf.bootstrap procedure cleans the TAF software folder, without requiring the user to run clean.software. The main reason for a TAF user to run clean.software directly is to experiment with installing and removing different versions of software without modifying the SOFTWARE.bib file.

### See Also

[taf.bootstrap](#) calls clean.software as part of the default bootstrap procedure.

[download.github](#) downloads a GitHub repository.

[clean.library](#) cleans the local TAF library.

[clean.data](#) cleans the bootstrap/data folder.

[icesTAF-package](#) gives an overview of the package.

### Examples

```
## Not run:  
clean.software()  
  
## End(Not run)
```

---

convert.spaces	<i>Convert Spaces</i>
----------------	-----------------------

---

### Description

Convert spaces in filenames.

### Usage

```
convert.spaces(file, sep = "_")
```

### Arguments

file	filename, e.g. "file name.csv", "*.csv", or "dir/*".
sep	character to use instead of spaces.

### Value

TRUE for success, FALSE for failure, invisibly.

### Note

This function treats '%20' in filenames as a space and converts to sep.

### See Also

[file.rename](#) is the base function to rename files.  
[icesTAF-package](#) gives an overview of the package.

### Examples

```
## Not run:  
write(pi, "A B.txt")  
convert.spaces("A B.txt")  
  
## Many files  
convert.spaces("bootstrap/initial/data/*")  
  
## End(Not run)
```

---

cp	<i>Copy Files</i>
----	-------------------

---

### Description

Copy or move files, overwriting existing files if necessary, and returning the result invisibly.

### Usage

```
cp(from, to, move = FALSE, ignore = FALSE, overwrite = TRUE, quiet = TRUE)
```

### Arguments

from	source filenames, e.g. *.csv.
to	destination filenames, or directory.
move	whether to move instead of copy.
ignore	whether to suppress error if source file does not exist.
overwrite	whether to overwrite if destination file exists.
quiet	whether to suppress messages.

### Value

TRUE for success, FALSE for failure, invisibly.

### Note

To prevent accidental loss of files, two safeguards are enforced when `move = TRUE`:

1. When moving files, the `to` argument must either have a filename extension or be an existing directory.
2. When moving many files to one destination, the `to` argument must be an existing directory.

If these conditions do not hold, no files are changed and an error is returned.

### See Also

[file.copy](#) and [unlink](#) are the underlying functions used to copy and (if `move = TRUE`) delete files.

[file.rename](#) is the base function to rename files.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:
write(pi, "A.txt")
cp("A.txt", "B.txt")
cp("A.txt", "B.txt", move=TRUE)

## Copy directory tree
cp(system.file(package="datasets"), ".")
mkdir("everything")
cp("datasets/*", "everything")

## End(Not run)
```

---

 deps

*List Dependencies*


---

**Description**

Search R scripts for packages that are required.

**Usage**

```
deps(path = ".", base = FALSE, installed = TRUE, available = TRUE,
      list = FALSE)
```

**Arguments**

path	a directory or file containing R scripts.
base	whether to include base packages in the output.
installed	whether to include installed packages in the output.
available	whether to include available packages in the output.
list	whether to return packages in list format, split by script.

**Value**

Names of packages as a vector, or in list format if `list=TRUE`. If no dependencies are found, the return value is `NULL`.

**Note**

Package names are matched based on four patterns:

```
library(*)
require(*)
*::object
*:::object
```

The search algorithm may return false-positive dependencies if these patterns occur inside if-clauses, strings, comments, etc.

**See Also**

[installed.packages](#), [available.packages](#).

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:
dir <- system.file(package="MASS", "scripts")
script <- system.file(package="MASS", "scripts/ch08.R")

deps(script)           # dependencies
deps(script, base=TRUE) # including base packages
deps(script, installed=FALSE) # not (yet) installed

deps(dir)
deps(dir, list=TRUE)

deps(dir, available=FALSE) # dependencies that might be unavailable

## End(Not run)
```

---

detach.packages

*Detach Packages*

---

**Description**

Detach all non-base packages that have been attached using `library` or `taf.library`.

**Usage**

```
detach.packages(quiet = FALSE)
```

**Arguments**

`quiet` whether to suppress messages.

**See Also**

`detach` is the underlying base function to detach a package.

`taf.library` loads a package from bootstrap/library.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:  
detach.packages()  
  
## End(Not run)
```

---

div

*Divide Columns*

---

## Description

Divide column values in a data frame with a common number.

## Usage

```
div(x, cols, by = 1000, grep = FALSE, ...)
```

## Arguments

x	a data frame.
cols	column names, or column indices.
by	a number to divide with.
grep	whether cols is a regular expression.
...	passed to grep().

## Value

A data frame similar to x, after dividing columns cols by the number by.

## Note

Provides notation that is convenient for modifying many columns at once.

## See Also

[transform](#) can also be used to recalculate column values, using a more general and verbose syntax.

[grep](#) is the underlying function used to match column names if grep is TRUE.

[rnd](#) is a similar function that rounds columns.

[icesTAF-package](#) gives an overview of the package.



## Examples

```
# These are equivalent:

x1 <- div(summary.taf, c("Rec", "Rec_lo", "Rec_hi",
                        "TSB", "TSB_lo", "TSB_hi",
                        "SSB", "SSB_lo", "SSB_hi",
                        "Removals", "Removals_lo", "Removals_hi"))

x2 <- div(summary.taf, "Rec|TSB|SSB|Removals", grep=TRUE)

x3 <- div(summary.taf, "Year|Fbar", grep=TRUE, invert=TRUE)

# Less reliable in scripts if columns have been added/deleted/reordered:

x4 <- div(summary.taf, 2:13)
```

---

dos2unix

*Convert Line Endings*

---

## Description

Convert line endings in a text file between Dos (CRLF) and Unix (LF) format.

## Usage

```
dos2unix(file)
```

```
unix2dos(file)
```

## Arguments

file            a filename.

## See Also

[line.endings](#) examines line endings.

[write.taf](#) uses unix2dos to ensure that the resulting files have Dos line endings.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:
file <- "test.txt"
write("123", file)

dos2unix(file)
file.size(file)
```

```

unix2dos(file)
file.size(file)

file.remove(file)

## End(Not run)

```

---

download

*Download File*


---

### Description

Download a file in binary mode, e.g. a model executable.

### Usage

```

download(url, dir = ".", mode = "wb", chmod = file_ext(url) == "",
  destfile = file.path(dir, basename(url)), quiet = TRUE, ...)

```

### Arguments

<code>url</code>	URL of file to download.
<code>dir</code>	directory to download to.
<code>mode</code>	download mode, see details.
<code>chmod</code>	whether to set execute permission (default is TRUE if file has no filename extension).
<code>destfile</code>	destination path and filename (optional, overrides <code>dir</code> ).
<code>quiet</code>	whether to suppress messages.
<code>...</code>	passed to <code>download.file</code> .

### Details

With the default mode "wb" the file is downloaded in binary mode (see [download.file](#)), to prevent R from adding ^M at line ends. This is particularly relevant for Windows model executables, while the `chmod` switch is useful when downloading Linux executables.

This function can be convenient for downloading any file, including text files. Data files in CSV or other text format can also be read directly into memory using `read.table`, `read.taf` or similar functions, without writing to the file system.

**Note**

If `destfile` contains a question mark it is removed from the `destfile` filename. Similarly, if `destfile` contains spaces or ‘%20’ sequences, those are converted to underscores.

In general, TAF scripts do not access the internet using `download` or similar functions. Instead, data and software are declared in `DATA.bib` and `SOFTWARE.bib` and then downloaded using `taf.bootstrap`. The exception is when a bootstrap script is used to fetch files from a web service (see [TAF Wiki](#)).

**See Also**

`download.file` is the underlying base function to download files.

`download.github` downloads a GitHub repository.

`icesTAF-package` gives an overview of the package.

**Examples**

```
## Not run:
url <- paste0("https://github.com/ices-taf/2015_had-iceg/raw/master/",
             "bootstrap/initial/software/catageysa.exe")
download(url)

## End(Not run)
```

---

download.github	<i>Download GitHub Repository</i>
-----------------	-----------------------------------

---

**Description**

Download a repository from GitHub in ‘tar.gz’ format.

**Usage**

```
download.github(repo, dir = ".", quiet = FALSE)
```

**Arguments**

repo	GitHub reference of the form owner/repo[/subdir]@ref.
dir	directory to download to.
quiet	whether to suppress messages.

**Value**

Name of downloaded tar.gz file.

**Note**

In general, TAF scripts do not access the internet using `download.github` or similar functions. Instead, data and software are declared in `DATA.bib` and `SOFTWARE.bib` and then downloaded using `taf.bootstrap`. The exception is when a bootstrap script is used to fetch files from a web service (see [TAF Wiki](#)).

**See Also**

`taf.bootstrap` uses `download.github` to fetch software and data repositories.

`download` downloads a file.

`untar` extracts a tar.gz archive.

`taf.install` installs a package in tar.gz format.

`icesTAF-package` gives an overview of the package.

**Examples**

```
## Not run:
# Specify release tag
download.github("ices-tools-prod/icesTAF@2.0-0")

# Specify SHA reference code
download.github("ices-tools-prod/icesTAF@d5a8947")

## End(Not run)
```

---

draft.data

*Draft DATA.bib*

---

**Description**

Create an initial draft version of a ‘DATA.bib’ metadata file.

**Usage**

```
draft.data(originator = NULL, year = format(Sys.time(), "%Y"),
  title = NULL, period = NULL, access = "Public", source = NULL,
  file = "", append = FALSE, data.files = dir("bootstrap/initial/data"),
  data.scripts = dir("bootstrap", pattern = "\\R$"))
```

**Arguments**

originator	who prepared the data, e.g. a working group acronym.
year	year of the analysis when the data were used. The default is the current year.
title	description of the data, including survey names or the like.

period	a string of the form "1990-2000", indicating the first and last year that the data cover, separated by a simple dash. Alternatively, a single number if the data cover only one year. If the data do not cover specific years, this metadata field can be suppressed using period = FALSE.
access	data access code: "OSPAR", "Public", or "Restricted".
source	where the data are copied/downloaded from. This can be a URL, filename, special value "file", or special value "script".
file	optional filename to save the draft metadata to a file. The value TRUE can be used as shorthand for "bootstrap/DATA.bib".
append	whether to append metadata entries to an existing file.
data.files	data files to consider. The default is all folders and files inside bootstrap/initial/data.
data.scripts	bootstrap data scripts to consider. The default is all *.R files in the bootstrap folder.

### Details

Typical usage is to specify originator, while using the default values for the other arguments. Most data files have the same originator, which can be specified to facilitate completing the entries after creating the initial draft.

The data access codes come from <https://vocab.ices.dk/?ref=1435>.

The special values source = "file" and source = "script" are described on the [TAF Wiki](#), along with other metadata information.

The default value file = "" prints the initial draft in the console, instead of writing it to a file. The output can then be pasted into a file to edit further, without accidentally overwriting an existing metadata file.

### Value

Object of class Bibtex.

### Note

This function is intended to be called from the top directory of a TAF analysis. It looks for data files inside bootstrap/initial/data folder and data scripts inside bootstrap.

After creating the initial draft, the user can complete the description of each data entry inside the title field and look into each file to specify the period that the data cover.

### See Also

[period](#) pastes two years to form a period string.

[draft.software](#) creates an initial draft version of a SOFTWARE.bib metadata file.

[taf.bootstrap](#) reads and processes metadata entries.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:
# Print in console
draft.data("WGEF", 2015)

# Export to file
draft.data("WGEF", 2015, file=TRUE)

## End(Not run)
```

---

draft.data.script      *Draft or create a bootstrap data script*

---

## Description

Create an 'R' file to fetch data including adding metadata via roxygen2 fields to the top of the file.

## Usage

```
draft.data.script(name, title, description, format, originator, year, period,
  access, content)
```

## Arguments

name	the name of the dataset and the file name that will be created.
title	description of the data, including survey names or the like.
description	a more involved description if required.
format	the format of the data produced, e.g. "csv"
originator	who prepared the data, e.g. a working group acronym.
year	year of the analysis when the data were used. The default is the current year.
period	a numeric vector of the form <code>c(1990, 2000)</code> , indicating the first and last year that the data cover. Alternatively, a single number if the data cover only one year.
access	data access code: "OSPAR", "Public", or "Restricted".
content	the r code that fetches and saves the data

---

draft.software	<i>Draft SOFTWARE.bib</i>
----------------	---------------------------

---

## Description

Create an initial draft version of a ‘SOFTWARE.bib’ metadata file.

## Usage

```
draft.software(package, author = NULL, year = NULL, title = NULL,
  version = NULL, source = NULL, file = "", append = FALSE)
```

## Arguments

package	name of one or more R packages, or files/folders starting with the path bootstrap/initial/software.
author	author(s) of the software.
year	year when this version of the software was released, or the publication year of the cited manual/article/etc.
title	title or short description of the software.
version	string to specify details about the version, e.g. GitHub branch and commit date.
source	string to specify where the software are copied/downloaded from. This can be a GitHub reference of the form owner/repo[/subdir]@ref, URL, or a filename.
file	optional filename to save the draft metadata to a file. The value TRUE can be used as shorthand for "bootstrap/SOFTWARE.bib".
append	whether to append metadata entries to an existing file.

## Details

Typical usage is to specify package, while using the default values for the other arguments.

If package is an R package, it can either be a package that is already installed ("icesAdvice") or a GitHub reference ("ices-tools-prod/icesAdvice@4271797").

With the default version = NULL, the function will automatically suggest an appropriate version entry for CRAN packages, but for GitHub packages it is left to the user to add further information about the GitHub branch (if different from master) and the commit date.

With the default source = NULL, the function will automatically suggest an appropriate source entry for CRAN and GitHub packages, but for other R packages it is left to the user to add information about where the software can be accessed.

The default value file = "" prints the initial draft in the console, instead of writing it to a file. The output can then be pasted into a file to edit further, without accidentally overwriting an existing metadata file.

## Value

Object of class Bibtex.

**Note**

After creating the initial draft, the user can complete the version, source, and other fields as required. This function is especially useful for citing exact versions of R packages on GitHub. To prepare metadata for software other than R packages, see the [TAF Wiki](#) for an example.

**See Also**

[citation](#) and [packageDescription](#) are the underlying functions to access information about installed R packages.

[draft.data](#) creates an initial draft version of a DATA.bib metadata file.

[taf.bootstrap](#) reads and processes metadata entries.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
# Print in console
draft.software("icesTAF")

## Not run:
# Export to file
draft.software("icesTAF", file=TRUE)

## End(Not run)
```

---

file.encoding

*File Encoding*

---

**Description**

Examine file encoding.

**Usage**

```
file.encoding(file)
```

**Arguments**

file            a filename.

**Value**

"latin1", "UTF-8", "unknown", or NA.

This function requires the file shell command. If the file utility is not found in the path, this function looks for it inside c:/Rtools/bin. If the required software is not installed, this function returns NA.



**Note**

The encoding "unknown" indicates that the file is an ASCII text file or a binary file.  
 In TAF, text files that have non-ASCII characters should be encoded as UTF-8.  
 If this function fails in Windows, the `guess_encoding` function in the **readr** package may help.

**See Also**

[Encoding](#) examines the encoding of a string.  
[latin1.to.utf8](#) converts files from latin1 to UTF-8 encoding.  
[line.endings](#) examines line endings.  
[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:
file.base <- system.file(package="base", "DESCRIPTION")
file.nlme <- system.file(package="nlme", "DESCRIPTION")
file.encoding(file.base) # ASCII
file.encoding(file.nlme)

## End(Not run)
```

---

 flr2taf

---

*Convert FLR Table to TAF Format*


---

**Description**

Convert a table from FLR format to TAF format.

**Usage**

```
flr2taf(x, colname = "Value")
```

**Arguments**

`x` a table of class FLQuant.  
`colname` a column name to use if the FLR table contains only one row.

**Value**

A data frame in TAF format.

**Note**

FLR uses the FLQuant class to store tables as 6-dimensional arrays, while TAF tables are stored as data frames with a year column.

**See Also**

[catage.taf](#) describes the TAF format.

[as.data.frame](#) is a method provided by the **FLCore** package to convert FLQuant tables to a 7-column long format.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
x <- array(t(catage.xtab), dim=c(4,8,1,1,1,1))
dimnames(x) <- list(age=1:4, year=1963:1970,
                   unit="unique", season="all", area="unique", iter=1)
flr2taf(x)

x1 <- x[1,,,,,drop=FALSE]
flr2taf(x1)
flr2taf(x1, "Juveniles")
```

---

get.remote.sha

*Get Remote SHA*

---

**Description**

Look up SHA reference code on GitHub.

**Usage**

```
get.remote.sha(owner, repo, ref, seven = TRUE)
```

**Arguments**

owner	repository owner.
repo	repository name.
ref	reference.
seven	whether to truncate SHA reference code to seven characters.

**Value**

SHA reference code as a string.

**See Also**

[taf.bootstrap](#) uses `get.remote.sha` to determine whether it is necessary to remove or download files, via [clean.library](#), [clean.software](#), and [download.github](#).

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:
get.remote.sha("ices-tools-prod", "icesTAF", "master")
get.remote.sha("ices-tools-prod", "icesTAF", "3.0-0")
get.remote.sha("ices-tools-prod", "icesTAF", "3.0-0", seven=FALSE)

## End(Not run)
```

---

is.r.package

*Is R Package*

---

## Description

Check if '.tar.gz' file is an R package.

## Usage

```
is.r.package(targz, spec = NULL, warn = TRUE)
```

## Arguments

targz	a filename ending with tar.gz.
spec	an optional list generated with parse.repo.
warn	whether to warn if the file contents look like an R package nested inside a repository.

## Details

The only purpose of passing spec is to get a more helpful warning message if the file contents look like an R package nested inside a repository.

## Value

Logical indicating whether targz is an R package.

## Examples

```
## Not run:
is.r.package("bootstrap/software/SAM.tar.gz")
is.r.package("bootstrap/software/stockassessment.tar.gz")

## End(Not run)
```

---

latin1.to.utf8	<i>Convert File Encoding</i>
----------------	------------------------------

---

**Description**

Convert file encoding between "latin1" and "UTF-8".

**Usage**

```
latin1.to.utf8(file, force = FALSE)
```

```
utf8.to.latin1(file, force = FALSE)
```

**Arguments**

file	a filename.
force	whether to perform the conversion even if the current file encoding cannot be verified with <a href="#">file.encoding</a> . Not recommended.

**Note**

In TAF, text files that have non-ASCII characters must be encoded as UTF-8.

**See Also**

[iconv](#) converts the encoding of a string.

[file.encoding](#) examines the encoding of a file.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
utf8.to.latin1("data.txt")  
latin1.to.utf8("data.txt")  
  
## End(Not run)
```

---

lim	<i>Axis Limits</i>
-----	--------------------

---

**Description**

Compute axis limits. The lower limit is 0 and the upper limit is determined by the highest data value, times a multiplier.

**Usage**

```
lim(x, mult = 1.1)
```

**Arguments**

x	a vector of data values.
mult	a number to multiply with the highest data value.

**Value**

A vector of length two, which can be used as axis limits.

**See Also**

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
plot(precip)
plot(precip, ylim=lim(precip))
plot(precip, ylim=lim(precip), yaxs="i")
```

---

line.endings	<i>Line Endings</i>
--------------	---------------------

---

**Description**

Examine whether file has Dos or Unix line endings.

**Usage**

```
line.endings(file)
```

**Arguments**

file	a filename.
------	-------------

**Value**

String indicating the line endings: "Dos" or "Unix".

**See Also**

[file.encoding](#) examines the encoding of a file.  
[dos2unix](#) and [unix2dos](#) convert line endings.  
[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
file <- system.file(package="icesTAF", "DESCRIPTION")  
line.endings(file)  
  
## End(Not run)
```

---

long2taf

*Convert Long Table to TAF Format*

---

**Description**

Convert a table from long format to TAF format.

**Usage**

```
long2taf(x)
```

**Arguments**

x                    a data frame in long format.

**Value**

A data frame in TAF format.

**Note**

TAF stores tables as data frames, usually with a year column as seen in stock assessment reports. The long format is more convenient for analysis and producing plots.

**See Also**

[catage.long](#) and [catage.taf](#) describe the long and TAF formats.  
[taf2long](#) converts a TAF table to long format.  
[icesTAF-package](#) gives an overview of the package.

**Examples**

```
long2taf(catage.long)
```

---

long2xtab

*Convert Long Table to Crosstab Format*

---

**Description**

Convert a table from long format to crosstab format.

**Usage**

```
long2xtab(x)
```

**Arguments**

x                    a data frame in long format.

**Value**

A data frame with years as row names.

**See Also**

[catage.long](#) and [catage.xtab](#) describe the long and crosstab formats.

[long2taf](#) and [taf2xtab](#) are the underlying functions that perform the conversion.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
long2xtab(catage.long)
```

---

 make

*Run R Script If Needed*


---

### Description

Run an R script if underlying files have changed, otherwise do nothing.

### Usage

```
make(recipe, prereq, target, include = TRUE, engine = source,
      debug = FALSE, force = FALSE, recon = FALSE, ...)
```

### Arguments

recipe	script filename.
prereq	one or more underlying files, required by the script. For example, data files and/or scripts.
target	one or more output files, produced by the script. Directory names can also be used.
include	whether to automatically include the script itself as a prerequisite file.
engine	function to source the script.
debug	whether to show a diagnostic table of files and time last modified.
force	whether to run the R script unconditionally.
recon	whether to return TRUE or FALSE, without actually running the R script.
...	passed to engine.

### Value

TRUE or FALSE, indicating whether the script was run.

### Note

This function provides functionality similar to makefile rules, to determine whether a script should be (re)run or not.

If any target is missing or older than any prereq, then the script is run.

### References

Stallman, R. M. *et al.* An introduction to makefiles. Chapter 2 in the *GNU Make manual*.

### See Also

[source](#) runs any R script, [sourceTAF](#) is more convenient for running a TAF script, and [sourceAll](#) runs all TAF scripts.

[make](#), [makeTAF](#), and [makeAll](#) are similar to the source functions, except they avoid repeating tasks that have already been run.

[icesTAF-package](#) gives an overview of the package.



**Examples**

```
## Not run:  
make("model.R", "data/input.dat", "model/results.dat")  
  
## End(Not run)
```

---

`makeAll`*Run All TAF Scripts as Needed*

---

**Description**

Run core TAF scripts that have changed, or if previous steps were rerun.

**Usage**

```
makeAll(...)
```

**Arguments**

... passed to `makeTAF`.

**Value**

Logical vector indicating which scripts were run.

**Note**

TAF scripts that will be run as needed: `data.R`, `model.R`, `output.R`, and `report.R`.

**See Also**

`source` runs any R script, `sourceTAF` is more convenient for running a TAF script, and `sourceAll` runs all TAF scripts.

`make`, `makeTAF`, and `makeAll` are similar to the source functions, except they avoid repeating tasks that have already been run.

`icesTAF-package` gives an overview of the package.

**Examples**

```
## Not run:  
makeAll()  
  
## End(Not run)
```

---

`makeTAF`*Run TAF Script If Needed*

---

**Description**

Run a TAF script if the target directory is either older than the script, or older than the directory of the previous TAF step.

**Usage**

```
makeTAF(script, ...)
```

**Arguments**

<code>script</code>	TAF script filename.
<code>...</code>	passed to <a href="#">make</a> and <a href="#">sourceTAF</a> .

**Value**

TRUE or FALSE, indicating whether the script was run.

**Note**

Any underlying scripts are automatically included if they share the same filename prefix, followed by an underscore. For example, when determining whether a script `data.R` should be run, this function checks whether `data_foo.R` and `data_bar.R` have been recently modified.

**See Also**

[source](#) runs any R script, [sourceTAF](#) is more convenient for running a TAF script, and [sourceAll](#) runs all TAF scripts.

[make](#), [makeTAF](#), and [makeAll](#) are similar to the source functions, except they avoid repeating tasks that have already been run.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
makeTAF("model.R")  
  
## End(Not run)
```

---

mkdir	<i>Create Directory</i>
-------	-------------------------

---

### Description

Create directory, including parent directories if necessary, without generating a warning if the directory already exists.

### Usage

```
mkdir(path)
```

### Arguments

path            a directory name.

### Value

TRUE for success, FALSE for failure, invisibly.

### See Also

[dir.create](#) is the base function to create a new directory.

[rmdir](#) removes an empty directory.

[clean](#) can be used to remove non-empty directories.

[icesTAF-package](#) gives an overview of the package.

### Examples

```
## Not run:  
mkdir("emptydir")  
rmdir("emptydir")  
  
mkdir("outer/inner")  
rmdir("outer", recursive=TRUE)  
  
## End(Not run)
```

msg *Show Message*

---

**Description**

Show a message, as well as the current time.

**Usage**

```
msg(...)
```

**Arguments**

... passed to message.

**See Also**

[message](#) is the base function to show messages, without the current time.

[sourceTAF](#) reports progress using msg.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
msg("script.R running..")
```

---

os *Operating System*

---

**Description**

Determine operating system name.

**Usage**

```
os()
```

```
os.linux()
```

```
os.macos()
```

```
os.windows()
```

```
os.unix()
```

**Value**

os returns the name of the operating system, typically "Linux", "Darwin", or "Windows".  
 os.linux, os.macos, os.unix, and os.windows return TRUE or FALSE.

**Note**

The macOS operating system identifies itself as "Darwin".

Both Linux and macOS are os.unix.

These shorthand functions can be useful when writing workaround solutions in platform-independent scripts.

**See Also**

[Sys.info](#) is the underlying function used to extract the operating system name.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
os()
os.linux()
os.macos()
os.unix()
os.windows()
```

---

period

*Period*

---

**Description**

Paste two years to form a period string.

**Usage**

```
period(x, y = NULL)
```

**Arguments**

x                    the first year, vector of years, matrix, or data frame.  
 y                    the last year, if x is only the first year.

**Details**

If x is a vector or a data frame, then the lowest and highest years are used, and y is ignored.

If x is a matrix or data frame, this function looks for years in the first column. If the values of the first column do not look like years (four digits), then it looks for years in the row names.

**Value**

A string of the form "1990-2000".

**Note**

This function can be useful when working with `draft.data`.

**See Also**

`paste` is the underlying function to paste strings.

`draft.data` has an argument called `period`.

`icesTAF-package` gives an overview of the package.

**Examples**

```
period(1963, 1970)
period(c(1963, 1970))
period(1963:1970)

period(range(catage.taf$Year))
period(catage.taf$Year)
period(catage.taf)
period(catage.xtab)
period(catage.long)
```

---

plus

*Rename Plus Group Column*

---

**Description**

Rename the last column in a data frame, by appending a "+" character. This is useful if the last column is a plus group.

**Usage**

```
plus(x)
```

**Arguments**

x                    a data frame.

**Value**

A data frame similar to x, after renaming the last column.

**Note**

If the last column name already ends with a "+", the original data frame is returned without modifications.

**See Also**

[names](#) is the underlying function to rename columns.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
catage <- catage.taf

# Rename last column
catage <- plus(catage)

# Shorter and less error-prone than
names(catage)[names(catage)=="4"] <- "4+"
```

---

read.bib

*bibtex parser*

---

**Description**

Parser for bibliography databases written in the bib format.

**Usage**

```
read.bib(file)
```

**Arguments**

file            bib file to parse

**Details**

Upload a table of F at age to the ICES TAF results database. The data and assessment\_info will be checked against a schema, and any errors reported back to the user as attributes to the FALSE return value.

**Value**

TRUE if successfull, FALSE otherwise

**Note**

The data argument expects a data.frame with the first column named "year" as described in the help for [xtab2taf](#).

**See Also**

The [xstab2taf](#) function is to create a data.frame

[taf.png](#) opens a PNG graphics device.

[icesTAF-package](#) gives an overview of the package.

---

 read.taf

*Read TAF Table from File*


---

**Description**

Read a TAF table from a file into a data frame.

**Usage**

```
read.taf(file, check.names = FALSE, stringsAsFactors = FALSE,
         fileEncoding = "UTF-8", ...)
```

**Arguments**

<code>file</code>	a filename.
<code>check.names</code>	whether to enforce regular column names, e.g. convert column name <code>"3"</code> to <code>"X3"</code> .
<code>stringsAsFactors</code>	whether to import strings as factors.
<code>fileEncoding</code>	character encoding of input file.
<code>...</code>	passed to <code>read.csv</code> .

**Details**

Alternatively, `file` can be a directory or a vector of filenames, to read many tables in one call.

**Value**

A data frame in TAF format, or a list of data frames if `file` is a directory or a vector of filenames.

**See Also**

[read.csv](#) is the underlying function used to read a table from a file.

[write.taf](#) writes a TAF table to a file.

[icesTAF-package](#) gives an overview of the package.



## Examples

```
## Not run:
write.taf(catage.taf, "catage.csv")
catage <- read.taf("catage.csv")

write.taf(catage)
file.remove("catage.csv")

## End(Not run)
```

---

rmdir	<i>Remove Empty Directory</i>
-------	-------------------------------

---

## Description

Remove empty directory under any operating system.

## Usage

```
rmdir(path, recursive = FALSE)
```

## Arguments

path	a directory name.
recursive	whether to remove empty subdirectories as well.

## Value

TRUE for success, FALSE for failure, invisibly.

## Note

The base function `unlink(dir,recursive=FALSE)` does not remove empty directories in Windows and `unlink(dir,recursive=TRUE)` removes non-empty directories, making it unsuitable for tidying up empty ones.

## See Also

[unlink](#) with `recursive = TRUE` removes non-empty directories.

[mkdir](#) creates a new directory.

[clean](#) can be used to remove non-empty directories.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:
mkdir("emptydir")
rmdir("emptydir")

mkdir("outer/inner")
rmdir("outer", recursive=TRUE)

## End(Not run)
```

---

rnd

*Round Columns*

---

## Description

Round column values in a data frame.

## Usage

```
rnd(x, cols, digits = 0, grep = FALSE, ...)
```

## Arguments

x	a data frame.
cols	column names, or column indices.
digits	number of decimal places.
grep	whether cols is a regular expression.
...	passed to <code>grep()</code> .

## Value

A data frame similar to `x`, after rounding columns `cols` to the number of `digits`.

## Note

Provides notation that is convenient for modifying many columns at once.

## See Also

[round](#) is the underlying function used to round numbers.

[grep](#) is the underlying function used to match column names if `grep` is `TRUE`.

[div](#) is a similar function that divides columns with a common number.

[icesTAF-package](#) gives an overview of the package.

The **icesAdvice** package provides the [icesRound](#) function to round values for ICES advice sheets.

## Examples

```
# With rnd() we no longer need to repeat the column names:

m <- mtcars
m[c("mpg", "disp", "qsec")] <- round(m[c("mpg", "disp", "qsec")])
m <- rnd(m, c("mpg", "disp", "qsec"))

# The x1/x2/x3/x4 approaches are equivalent:

x1 <- rnd(summary.taf, c("Rec", "Rec_lo", "Rec_hi",
                        "TSB", "TSB_lo", "TSB_hi",
                        "SSB", "SSB_lo", "SSB_hi",
                        "Removals", "Removals_lo", "Removals_hi"))
x1 <- rnd(x1, c("Fbar", "Fbar_lo", "Fbar_hi"), 3)

x2 <- rnd(summary.taf, "Rec|TSB|SSB|Removals", grep=TRUE)
x2 <- rnd(x2, "Fbar", 3, grep=TRUE)

x3 <- rnd(summary.taf, "Fbar", grep=TRUE, invert=TRUE)
x3 <- rnd(x3, "Fbar", 3, grep=TRUE)

# Less reliable in scripts if columns have been added/deleted/reordered:

x4 <- rnd(summary.taf, 2:13)
x4 <- rnd(x4, 14:16, 3)
```

---

sam2taf

*Convert SAM Table to TAF Format*

---

## Description

Convert a table from SAM format to TAF format.

## Usage

```
sam2taf(x, colname = NULL, year = TRUE)
```

## Arguments

x	a matrix containing columns Estimate, Low, and High.
colname	a descriptive column name for the output.
year	whether to include a year column.

**Details**

The default when `colname = NULL` is to try to infer a column name from the `x` argument. For example,

```
sam2taf(ssbtable(fit))
sam2taf(ssb)
sam2taf(SSB)
```

will recognize `ssbtable` calls and `ssb` object names, implicitly setting `colname = "SSB"` if the user does not pass an explicit value for `colname`.

**Value**

A data frame in TAF format.

**Note**

The **stockassessment** package provides accessor functions that return a matrix with columns Estimate, Low, and High, while TAF tables are stored as data frames with a year column.

**See Also**

[summary.taf](#) describes the TAF format.

`catchtable`, `fbartable`, `rectable`, `ssbtable`, and `tsbtable` (in the **stockassessment** package) return matrices with SAM estimates and confidence limits.

The `summary` method for `sam` objects produces a summary table with some key quantities of interest, containing duplicated column names (Low, High) and rounded values.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Example objects
x <- as.matrix(summary.taf[grep("SSB", names(summary.taf))])
rec <- as.matrix(summary.taf[grep("Rec", names(summary.taf))])
tsb <- as.matrix(summary.taf[grep("TSB", names(summary.taf))])
dimnames(x) <- list(summary.taf$Year, c("Estimate", "Low", "High"))
dimnames(rec) <- dimnames(tsb) <- dimnames(x)

## One SAM table, arbitrary object name
sam2taf(x)
sam2taf(x, "SSB")
sam2taf(x, "SSB", year=FALSE)

## Many SAM tables, recognized names
sam2taf(rec)
data.frame(sam2taf(rec), sam2taf(tsb, year=FALSE))

## Not run:
```

```
## Accessing tables from SAM fit object
data.frame(sam2taf(rectable(fit)), sam2taf(tsbtable(fit), year=FALSE))

## End(Not run)
```

---

sourceAll

*Run All TAF Scripts*

---

### Description

Run core TAF scripts in current directory.

### Usage

```
sourceAll(...)
```

### Arguments

... passed to [sourceTAF](#).

### Value

Logical vector, indicating which scripts ran without errors.

### Note

TAF scripts that will be run if they exist: utilities.R data.R, model.R, output.R, and report.R.

### See Also

[sourceTAF](#) runs a TAF script.

[makeAll](#) runs all TAF scripts as needed.

[clean](#) cleans TAF directories.

[icesTAF-package](#) gives an overview of the package.

### Examples

```
## Not run:
sourceAll()

## End(Not run)
```

---

sourceDir	<i>Source Directory</i>
-----------	-------------------------

---

### Description

Read all \*.R files from a directory containing R functions.

### Usage

```
sourceDir(dir, pattern = "\\.[r|R]$", all.files = FALSE,  
          recursive = FALSE, quiet = TRUE, ...)
```

### Arguments

dir	a directory containing R source files.
pattern	passed to <code>dir</code> when selecting files.
all.files	passed to <code>dir</code> when selecting files.
recursive	passed to <code>dir</code> when selecting files.
quiet	whether to suppress messages.
...	passed to <code>source</code> when sourcing files.

### Details

The `dir` argument can also be a vector of filenames, instead of a directory name. This can be useful to specify certain files while avoiding others.

### Value

Names of sourced files.

### Note

This function is convenient in TAF analyses when many R utility functions are stored in a directory, see example below.

### See Also

[source](#) is the base function to read R code from a file.  
[icesTAF-package](#) gives an overview of the package.

### Examples

```
## Not run:  
sourceDir("bootstrap/software/utilities")  
  
## End(Not run)
```

---

`sourceTAF`*Run TAF Script*

---

**Description**

Run a TAF script and return to the original directory.

**Usage**

```
sourceTAF(script, rm = FALSE, clean = TRUE, detach = FALSE, taf = NULL,  
          quiet = FALSE)
```

**Arguments**

<code>script</code>	script filename.
<code>rm</code>	whether to remove all objects from the global environment before and after the script is run.
<code>clean</code>	whether to <a href="#">clean</a> the target directory before running the script.
<code>detach</code>	whether to detach all non-base packages before running the script, to ensure that the script is not affected by packages that may have been attached outside the script.
<code>taf</code>	a convenience flag where <code>taf = TRUE</code> sets <code>rm</code> , <code>clean</code> , and <code>detach</code> to <code>TRUE</code> , as is done on the TAF server. Any other value of <code>taf</code> is ignored.
<code>quiet</code>	whether to suppress messages reporting progress.

**Details**

The default value of `rm = FALSE` is to protect users from accidental loss of work, but the TAF server always runs with `rm = TRUE` to make sure that only files, not objects, are carried over between scripts.

Likewise, the TAF server runs with `clean = TRUE` to make sure that the script starts with a clean directory. The target directory of a TAF script has the same filename prefix as the script: `data.R` creates `'data'` etc.

**Value**

TRUE or FALSE, indicating whether the script ran without errors.

**Note**

Commands within a script (such as `setwd`) may change the working directory, but `sourceTAF` guarantees that the working directory reported by `getwd()` is the same before and after running a script.

**See Also**

[source](#) is the base function to run R scripts.

[makeTAF](#) runs a TAF script if needed.

[sourceAll](#) runs all TAF scripts in a directory.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:
write("print(pi)", "script.R")
source("script.R")
sourceTAF("script.R")
file.remove("script.R")

## End(Not run)
```

---

summary.taf

*Summary Results in TAF Format*


---

**Description**

Small summary results table to describe a TAF format data frame to store values by year.

**Usage**

```
summary.taf
```

**Format**

Data frame containing 16 columns:

Year	year
Rec	recruitment, numbers at age 1 in this year (thousands)
Rec_lo	lower 95% confidence limit
Rec_hi	upper 95% confidence limit
TSB	total stock biomass (tonnes)
TSB_lo	lower 95% confidence limit
TSB_hi	upper 95% confidence limit
SSB	spawning stock biomass (tonnes)
SSB_lo	lower 95% confidence limit
SSB_hi	upper 95% confidence limit
Removals	total removals, including catches due to unaccounted mortality
Removals_lo	lower 95% confidence limit
Removals_hi	upper 95% confidence limit
Fbar	average fishing mortality (ages 2-4)
Fbar_lo	lower 95% confidence limit
Fbar_hi	upper 95% confidence limit



**Details**

The data are an excerpt (first years) from the summary results table for North Sea cod from the ICES (2016) assessment.

**Source**

ICES (2016) Report of the working group on the assessment of demersal stocks in the North Sea and Skagerrak (WGNSSK). *ICES CM 2016/ACOM:14*, p. 673.

**See Also**

[div](#) and [rnd](#) can modify a large number of columns.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
summary.taf
x <- div(summary.taf, "Rec|TSB|SSB|Removals", grep=TRUE)
x <- rnd(x, "Rec|TSB|SSB|Removals", grep=TRUE)
x <- rnd(x, "Fbar", 3, grep=TRUE)
```

---

taf.boot.path

*Construct Path to a TAF bootstrap folder*


---

**Description**

Construct the path to a file in the TAF bootstrap data folder from components in a platform-independent way. This function checks to see if R is running in the bootstrap folder - i.e. 'taf.bootstrap()' is running, and adjusts the path accordingly.

**Usage**

```
taf.boot.path(..., fsep = .Platform$file.sep)
```

**Arguments**

... character vectors. Long vectors are not supported.  
fsep the path separator to use (assumed to be ASCII).

**Details**

This function, simplifies the construction of file paths to the boot (bootstrap) folder.

**Value**

character

**See Also**[file.path](#)

taf.bootstrap

*Bootstrap TAF Analysis***Description**

Process metadata files ‘SOFTWARE.bib’ and ‘DATA.bib’ to set up software and data files required for the analysis.

**Usage**

```
taf.bootstrap(software = TRUE, data = TRUE, clean = TRUE,
              force = FALSE, taf = NULL, quiet = FALSE)
```

**Arguments**

software	whether to process SOFTWARE.bib.
data	whether to process DATA.bib.
clean	whether to <a href="#">clean</a> directories during the bootstrap procedure.
force	whether to remove existing bootstrap/data, bootstrap/library, and bootstrap/software directories before the bootstrap procedure.
taf	a convenience flag where taf = TRUE sets software, data, clean, and force to TRUE, as is done on the TAF server. Any other value of taf is ignored.
quiet	whether to suppress messages reporting progress.

**Details**

If clean = TRUE then:

1. [clean.software](#) and [clean.library](#) are run if ‘SOFTWARE.bib’ is processed.
2. [clean.data](#) is run if ‘DATA.bib’ is processed.

The default behavior of taf.bootstrap is to skip downloading of remote files (GitHub resources, URLs, bootstrap scripts) and also skip installing R packages from GitHub if the files seem to be already in place. This is done to speed up the bootstrap procedure as much as possible. To override this and guarantee that all data and software files are updated, pass force = TRUE to download and install everything declared in SOFTWARE.bib and DATA.bib.

**Value**

Logical vector indicating which metadata files were processed.

**Note**

This function should be called from the top directory of a TAF analysis. It looks for a directory called 'bootstrap' and prepares data files and software according to metadata specifications.

The bootstrap procedure consists of the following steps:

1. If a bootstrap/SOFTWARE.bib metadata file exists, it is processed.
2. If a bootstrap/DATA.bib metadata file exists, it is processed.

After the bootstrap procedure, software and data have been documented and are ready to be used in the subsequent analysis. Specifically, the procedure populates up to three new directories:

- bootstrap/data with data files.
- bootstrap/library with R packages compiled for the local platform.
- bootstrap/software with software files, such as R packages in tar.gz source code format.

Model settings and configuration files can be set up within DATA.bib, see [TAF Wiki](#).

**See Also**

[draft.data](#) and [draft.software](#) can be used to create initial draft versions of 'DATA.bib' and 'SOFTWARE.bib' metadata files.

[taf.library](#) loads a package from bootstrap/library.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
taf.bootstrap()  
  
## End(Not run)
```

---

taf.colors

*TAF Colors*

---

**Description**

Predefined colors that can be useful in TAF plots.

**Usage**

```
taf.green  
taf.orange  
taf.blue  
taf.dark  
taf.light
```

**See Also**

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
taf.green

par(mfrow=c(3,1))

barplot(5:1, main="Five",
        col=c(taf.green, taf.orange, taf.blue, taf.dark, taf.light))

barplot(6:1, main="Six", col=c(taf.green, taf.orange, taf.blue,
                              taf.dark, taf.light, "white"))

barplot(7:1, main="Seven", col=c("black", taf.dark, taf.light,
                                 taf.green, taf.orange, taf.blue, "white"))
```

---

taf.data.path	<i>Construct Path to a TAF bootstrap data file</i>
---------------	--

---

**Description**

Construct the path to a file in the TAF bootstrap data folder from components in a platform-independent way. This function checks to see if R is running in the bootstrap folder - i.e. ‘taf.bootstrap()’ is running, and adjusts the path accordingly.

**Usage**

```
taf.data.path(..., fsep = .Platform$file.sep)
```

**Arguments**

...            character vectors. Long vectors are not supported.  
fsep            the path separator to use (assumed to be ASCII).

**Details**

This function, simplifies the construction of file paths to initial data files gathered during the TAF bootstrapping step. In addition, this function is useful when developing scripts used in the bootstrap procedure, as these scripts are run with the working directory set to the bootstrap folder, and hence make it to develop and debug.

**Value**

character

**See Also**[file.path](#)

---

`taf.install`*TAF Install*

---

**Description**

Install packages in ‘tar.gz’ format in local TAF library.

**Usage**

```
taf.install(targz = NULL, lib = "bootstrap/library", quiet = FALSE)
```

**Arguments**

<code>targz</code>	a package filename, vector of filenames, or NULL.
<code>lib</code>	location of local TAF library.
<code>quiet</code>	whether to suppress messages.

**Details**

If `targz = NULL`, all packages found in `bootstrap/software` are installed, as long as they have filenames of the form `package_sha.tar.gz` containing a 7-character SHA reference code.

The default behavior of `taf.install` is to install packages in alphabetical order. When the installation order matters because of dependencies, the user can specify a vector of package filenames to install.

**Note**

The `taf.bootstrap` procedure downloads and installs R packages, without requiring the user to run `taf.install`. The main reason for a TAF user to run `taf.install` directly is to initialize and run a TAF analysis without running the bootstrap procedure, e.g. to avoid updating the underlying datasets and software.

After installing the package, this function writes the remote SHA reference code into the package files `DESCRIPTION` and `Meta/package.rds`.

**See Also**

[taf.bootstrap](#) calls [download.github](#) and `taf.install` to download and install R packages.

[taf.library](#) loads a package from `bootstrap/library`.

[clean.library](#) selectively removes packages from the local TAF library.

[install.packages](#) is the underlying base function to install a package.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:  
# Install one package  
taf.install("bootstrap/software/FLAssess_f1e5acb.tar.gz")  
  
# Install all packages found in bootstrap/software  
taf.install()  
  
## End(Not run)
```

---

taf.libPaths	<i>Add TAF Library Path</i>
--------------	-----------------------------

---

## Description

Add TAF library to the search path for R packages.

## Usage

```
taf.libPaths(remove = FALSE)
```

## Arguments

remove                    whether to remove TAF library from the search path, instead of adding it.

## Value

The resulting vector of file paths.

## Warning

An unwanted side effect of having the TAF library as the first element in the search path is that `install.packages` will then install packages inside `bootstrap/library`. This is not a serious side effect, since a subsequent call to `taf.bootstrap` or `clean.library` will remove packages from the TAF library that are not declared in the `'SOFTWARE.bib'` file.

## Note

Specifically, this function sets `"bootstrap/library"` as the first element of `.libPaths()`. This is rarely beneficial in TAF scripts, but can be useful when using the `sessioninfo` package, for example.

## See Also

[.libPaths](#) is the underlying function to modify the search path for R packages.  
[taf.library](#) loads a package from `bootstrap/library`.  
[icesTAF-package](#) gives an overview of the package.

## Examples

```
taf.libPaths()  
taf.libPaths(remove=TRUE)
```

---

taf.library	<i>TAF Library</i>
-------------	--------------------

---

## Description

Load and attach package from local TAF library.

## Usage

```
taf.library(package, messages = FALSE, warnings = FALSE)
```

## Arguments

package	name of a package found in bootstrap/library.
messages	whether to show messages when package loads.
warnings	whether to show warnings when package loads.

## Value

The names of packages currently installed in the TAF library.

## Note

The purpose of the TAF library is to retain R packages that are not commonly used (and not on CRAN), to support long-term reproducibility of TAF analyses.

## See Also

[library](#) is the underlying base function to load and attach a package.

[taf.bootstrap](#) is the procedure to install packages into a local TAF library, via the SOFTWARE.bib metadata file.

[detach.packages](#) detaches all packages.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:  
  
# Show packages in TAF library  
taf.library()  
  
# Load packages  
taf.library(this)  
taf.library(that)  
  
## End(Not run)
```

---

taf.png

*PNG Device*

---

## Description

Open PNG graphics device to export a plot into the TAF report folder.

## Usage

```
taf.png(filename, width = 1600, height = 1200, res = 200, ...)
```

## Arguments

filename	plot filename.
width	image width.
height	image height.
res	resolution determining the text size, line width, plot symbol size, etc.
...	passed to png.

## Details

The filename can be passed without the preceding "report/", and without the ".png" filename extension.

Specifically, the function prepends "report/" to the filename if (1) the filename does not contain a "/" separator, (2) the working directory is not report, and (3) the directory report exists. The function also appends ".png" to the filename if it does not already have that filename extension.

This automatic filename manipulation can be bypassed by using the png function directly.



**Note**

A simple convenience function to shorten

```
png("report/plot.png", width=1600, height=1200, res=200)
```

to

```
taf.png("plot")
```

The `res` argument affects the text size, along with all other plot elements. To change the text size of specific lattice plot elements, the `zoom` function can be helpful.

For consistent image width and text size, it can be useful to keep the default width = 1600 but vary the height to adjust the desired aspect ratio for each plot.

**See Also**

[png](#) is the underlying function used to open a PNG graphics device.

[zoom](#) changes text size in a lattice plot.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:
taf.png("myplot")
plot(1)
dev.off()

library(lattice)
taf.png("mytrellis")
xyplot(1~1)
dev.off()

library(ggplot2)
taf.png("myggplot")
qplot(1, 1)
dev.off()

## End(Not run)
```

**Description**

This function builds documentation for a TAF repository using roxygen syntax headers. It depends on the `roxygen2` package adding some extra functionality to produce citation entries for data sources

**Usage**

```
taf.roxygenise(path = ".", files)
```

**Arguments**

path	location of taf repository top level directory. Default is working directory.
files	a vector of file names to parse for documentation.

---

taf.session	<i>TAF Session</i>
-------------	--------------------

---

**Description**

Show session information about loaded packages, clearly indicating which packages were loaded from the local TAF library.

**Usage**

```
taf.session(sort = FALSE, details = FALSE)
```

**Arguments**

sort	whether to sort packages by name.
details	whether to include more detailed session information.

**Value**

List containing session information about loaded packages.

**See Also**

[sessionInfo](#) and the **sessioninfo** package provide similar information, but do not indicate clearly packages that were loaded from the local TAF library.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
taf.session()  
taf.session(sort=TRUE)  
taf.session(details=TRUE)
```

---

taf.skeleton	<i>TAF Skeleton</i>
--------------	---------------------

---

**Description**

Create initial directories and R scripts for a new TAF analysis.

**Usage**

```
taf.skeleton(path = ".", force = FALSE)
```

**Arguments**

path	where to create initial directories and R scripts. The default is the current working directory.
force	whether to overwrite existing scripts.

**Value**

Full path to analysis directory.

**See Also**

[package.skeleton](#) creates an empty template for a new R package.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
taf.skeleton()  
  
## End(Not run)
```

---

taf.sources	<i>Extract sources from TAF *.bib file</i>
-------------	--

---

**Description**

Extract a list of sources from a TAF \*.bib file (i.e. DATA.bib or SOFTWARE.bib). This allows the user to print the list of sources but also to process them individually, giving more flexibility when developing larger projects.

**Usage**

```
taf.sources(type)
```

**Arguments**

type                    one of "data", "software" or "both"

**See Also**

[process.entry](#) to process one of the entries returned by 'taf.sources'.

---

taf.unzip

*Unzip File*


---

**Description**

Extract files from a zip archive, retaining executable file permissions.

**Usage**

```
taf.unzip(zipfile, files = NULL, exdir = ".", unzip = NULL, ...)
```

**Arguments**

zipfile                zip archive filename.  
files                    files to extract, default is all files.  
exdir                    directory to extract to, will be created if necessary.  
unzip                    extraction method to use, see details below.  
...                      passed to [unzip](#).

**Details**

The default method unzip = NULL uses the external unzip program in Unix-compatible operating systems, but an internal method in Windows. For additional information, see the [unzip](#) help page.

**Note**

One shortcoming of the base unzip function is that the default "internal" method resets file permissions, so Linux and macOS executables will return a 'Permission denied' error when run.

This function is identical to the base unzip function, except the default value unzip = NULL chooses an appropriate extraction method in all operating systems, making it useful when writing platform-independent scripts.

**See Also**

[unzip](#) is the base function to unzip files.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
## Not run:
exefile <- if(os.unix()) "run" else "run.exe"
taf.unzip("bootstrap/software/archive.zip", files=exefile, exdir="model")

## End(Not run)
```

---

taf2long

*Convert TAF Table to Long Format*

---

## Description

Convert a table from TAF format to long format.

## Usage

```
taf2long(x, names = c("Year", "Age", "Value"))
```

## Arguments

`x` a data frame in TAF format.  
`names` a vector of three column names for the resulting data frame.

## Value

A data frame with three columns.

## Note

TAF stores tables as data frames, usually with a year column as seen in stock assessment reports. The long format is more convenient for analysis and producing plots.

## See Also

[catage.taf](#) and [catage.long](#) describe the TAF and long formats.

[long2taf](#) converts a long table to TAF format.

[icesTAF-package](#) gives an overview of the package.

## Examples

```
taf2long(catage.taf, names=c("Year", "Age", "Catch"))
```

---

taf2xtab	<i>Convert TAF Table to Crosstab Format</i>
----------	---

---

**Description**

Convert a table from TAF format to crosstab format.

**Usage**

```
taf2xtab(x)
```

**Arguments**

`x` a data frame in TAF format.

**Value**

A data frame with years as row names.

**Note**

TAF stores tables as data frames, usually with a year column as seen in stock assessment reports. The crosstab format can be more convenient for analysis and producing plots.

**See Also**

[catage.taf](#) and [catage.xtab](#) describe the TAF and crosstab formats.

[tt](#) converts a TAF table to transposed crosstab format.

[xtab2taf](#) converts a crosstab table to TAF format.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
taf2xtab(catage.taf)
```

---

tt	<i>TAF Transpose</i>
----	----------------------

---

**Description**

Convert a table from TAF format to transposed crosstab format.

**Usage**

```
tt(x, column = FALSE)
```

**Arguments**

x	a data frame in TAF format.
column	a logical indicating whether the group names should be stored in a column called 'Age' instead of in row names. Alternatively, column can be a string supplying another name for that first column.

**Value**

A data frame with years as column names.

**Note**

Transposing can be useful when comparing TAF tables to stock assessment reports.

**See Also**

[t](#) transposes a matrix.  
[catage.taf](#) describes the TAF format.  
[taf2xtab](#) converts a TAF table to crosstab format, without transposing.  
[icesTAF-package](#) gives an overview of the package.

**Examples**

```
taf2xtab(catage.taf)  
tt(catage.taf)  
tt(catage.taf, TRUE)  
tt(catage.taf, "Custom")
```

---

`write.taf`*Write TAF Table to File*

---

### Description

Write a TAF table to a file.

### Usage

```
write.taf(x, file = NULL, dir = NULL, quote = FALSE, row.names = FALSE,  
          fileEncoding = "UTF-8", underscore = TRUE, ...)
```

### Arguments

<code>x</code>	a data frame in TAF format.
<code>file</code>	a filename.
<code>dir</code>	an optional directory name.
<code>quote</code>	whether to quote strings.
<code>row.names</code>	whether to include row names.
<code>fileEncoding</code>	character encoding for output file.
<code>underscore</code>	whether automatically generated filenames (when <code>file = NULL</code> ) should use underscore separators instead of dots.
<code>...</code>	passed to <code>write.csv</code> .

### Details

Alternatively, `x` can be a list of data frames or a string vector of object names, to write many tables in one call. The resulting files are named automatically, similar to `file = NULL`.

The default value `file = NULL` uses the name of `x` as a filename, so a data frame called `survey.uk` will be written to a file called `'survey_uk.csv'` (when `underscore = TRUE`) or `'survey.uk.csv'` (when `underscore = FALSE`).

The special value `file = ""` prints the data frame in the console, similar to `write.csv`.

### Note

The resulting CSV file has Dos line endings, as specified in the RFC 4180 standard (IETF 2005).

This function gives a warning when column names are duplicated, unless the target directory name is `report`.

### References

IETF (2005) Common format and Mime type for Comma-Separated Values (CSV) files. *IETF RFC 4180*.



**See Also**

[write.csv](#) is the underlying function used to write a table to a file.

[read.taf](#) reads a TAF table from a file into a data frame.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
## Not run:
write.taf(catage.taf, "catage.csv")
catage <- read.taf("catage.csv")

write.taf(catage)
file.remove("catage.csv")

## End(Not run)
```

---

xstab2long

*Convert Crosstab Table to Long Format*

---

**Description**

Convert a table from crosstab format to long format.

**Usage**

```
xstab2long(x, names = c("Year", "Age", "Value"))
```

**Arguments**

x                    a data frame in crosstab format.  
names                a vector of three column names for the resulting data frame.

**Value**

A data frame with three columns.

**See Also**

[catage.xtab](#) and [catage.long](#) describe the crosstab and long formats.

[xtab2taf](#) and [taf2long](#) are the underlying functions that perform the conversion.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
xstab2long(catage.xtab, names=c("Year", "Age", "Catch"))
```

---

xtab2taf	<i>Convert Crosstab Table to TAF Format</i>
----------	---

---

**Description**

Convert a table from crosstab format to TAF format.

**Usage**

```
xtab2taf(x, colname = "Year")
```

**Arguments**

x	a data frame in crosstab format.
colname	name for first column.

**Value**

A data frame in TAF format.

**Note**

TAF stores tables as data frames, usually with a year column as seen in stock assessment reports. The crosstab format can be more convenient for analysis and producing plots.

**See Also**

[catage.xtab](#) and [catage.taf](#) describe the crosstab and TAF formats.

[taf2xtab](#) converts a TAF table to crosstab format.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
xtab2taf(catage.xtab)
```

---

 zoom

*Zoom*


---

### Description

Change text size in a lattice plot.

### Usage

```
zoom(x, ...)
```

```
## S3 method for class 'trellis'
zoom(x, size = 1, main = 1.2 * size, lab = size,
      axis = size, strip = size, sub = 0.9 * size, legend = 0.9 * size,
      splom = 0.9 * size, ...)
```

### Arguments

x	a lattice plot of class "trellis".
...	further arguments, currently ignored.
size	text size multiplier.
main	size of main title (default is 1.2 * size).
lab	size of axis labels (default is size).
axis	size of tick labels (default is size).
strip	size of strip labels (default is size).
sub	size of subtitle (default is 0.9 * size).
legend	size of legend labels (default is 0.9 * size).
splom	size of scatterplot matrix diagonal labels (default is 0.9 * size).

### Details

Pass NULL for any argument to avoid changing the size of that text component.

The legend component of a lattice plot can be somewhat fickle, as the object structure varies between plots. One solution is to pass `legend = NULL` and tweak the legend before or after calling the `zoom` function.

### Value

The same lattice object, but with altered text size.

### Note

The default values result in lattice plots that have similar text size as base plots, when using `taf.png`.

This function ends with a `print` call, to make it easy to export the lattice plot to a file, without the need of an explicit `print`.

**See Also**

[Lattice](#) plots are created using [xyplot](#) or related functions.

[taf.png](#) opens a PNG graphics device.

[icesTAF-package](#) gives an overview of the package.

**Examples**

```
library(lattice)

xyplot(1~1)
zoom(xyplot(1~1))
zoom(xyplot(1~1), size=1.2)
zoom(xyplot(1~1), axis=0.8)
zoom(xyplot(1~1), axis=NULL)

## Not run:
taf.png("myplot")
plot(1)
dev.off()

taf.png("mytrellis")
xyplot(1~1)
dev.off()

taf.png("mytrellis_zoom")
zoom(xyplot(1~1))
dev.off()

## End(Not run)
```

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