

# Package ‘icesTAF’

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icesTAF-package      *Functions to Support the ICES Transparent Assessment Framework*

---

### Description

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

### Details

*Web services:*

[download](#)    download file in binary mode

*Read and write files:*

[read.dls](#)    read DLS3.2 results from file  
[read.taf](#)    read TAF table from file  
[write.dls](#)    write DLS3.2 results to file  
[write.taf](#)    write TAF table to file

*Run scripts:*

<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

*Other file management:*

<code>clean</code>	clean TAF directories
<code>cp</code>	copy files
<code>dos2unix</code>	convert line endings
<code>mkdir</code>	create directory
<code>taf.skeleton</code>	create empty TAF template
<code>unix2dos</code>	convert line endings

*Table tools:*

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

*Plotting tools:*

<code>lim</code>	compute axis limits
<code>taf.colors</code>	predefined colors
<code>tafpng</code>	open PNG graphics device

*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:*

<code>deps</code>	list dependencies
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**Author(s)**

Arni Magnusson and Colin Millar.

**References**

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

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

---

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. 656.

**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`*Catch at Age in TAF Format*

---

**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. 656.

**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. 656.

### 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 TAF directories: data, input, model, output, report.

**Usage**

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

**Arguments**

dirs                directories to delete.

**Note**

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

**See Also**

[sourceTAF](#) runs a TAF script.

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

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

**Examples**

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

**Arguments**

from	source filenames, e.g. *.csv.
to	destination filenames, or directory.
move	whether to move instead of copy.

**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.

**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

```
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)

## Not run:
deps(dir, available=FALSE) # dependencies that might be unavailable

## 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.

**Note**

TAF uses `unix2dos` to ensure that text files on the server have Dos line endings.

### See Also

[file](#) is used to open a binary connection to the text file.

[writeLines](#) is used to apply CRLF or LF 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 in Binary Mode*

---

### Description

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

### Usage

```
download(url, dir = ".", mode = "wb", chmod = file_ext(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>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.

## See Also

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

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

## Examples

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

## 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")
```

---

 lim

*Axis Limits*


---

**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(rivers, ylim=lim(rivers))
```

---

long2taf	<i>Convert Long Table to TAF Format</i>
----------	---

---

**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 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)
```

---

make	<i>Run R Script If Needed</i>
------	-------------------------------

---

**Description**

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

**Usage**

```
make(recipe, prereq, target, include = TRUE, engine = source,  
      debug = 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.
...	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", "input/input.dat", "model/results.dat")  
  
## End(Not run)
```



---

makeAll	<i>Run All TAF Scripts as Needed</i>
---------	--------------------------------------

---

**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, input.R, model.R, output.R, and report.R.

If a begin.R script exists, it is ignored.

**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 an empty directory.

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

[clean](#) cleans TAF directories.

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

### Examples

```
## Not run:
mkdir("emptydir")
unlink("emptydir", recursive=TRUE)

mkdir("outer/inner")
unlink("outer/inner", 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...")
```

---

`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.

**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.dls	<i>Read DLS3.2 Results from File</i>
----------	--------------------------------------

---

**Description**

Read results from the DLS3.2 advisory method from a file into a list.

**Usage**

```
read.dls(file)
```

**Arguments**

file            a filename.

**Value**

A list containing advice and other elements showing intermediate steps in the calculations.

**See Also**

[write.dls](#) writes DLS3.2 results to a file.

[DLS3.2](#) in the **icesAdvice** package can be used to calculate catch advice for data-limited stocks (DLS).

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

**Examples**

```
## Not run:
survey <- data.frame(year=2001:2010, randu[1:10,])
dls <- icesAdvice::DLS3.2(1000, survey$y)

write.dls(dls, "dls.txt")
read.dls("dls.txt")

file.remove("dls.txt")

## End(Not run)
```

---

read.taf	<i>Read TAF Table from File</i>
----------	---------------------------------

---

**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**

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

**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)
```

---

`rnd`

*Round Columns*

---

## Description

Round column values in a data frame.

## Usage

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

## Arguments

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

---

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: data.R, input.R, model.R, output.R, and report.R.  
If a begin.R script exists, it is ignored.



**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)
```

---

sourceTAF

*Run TAF Script*


---

**Description**

Run a TAF script and return to the original directory.

**Usage**

```
sourceTAF(script, rm = FALSE, clean = TRUE, 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 remove the target directory before running the script.
<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 by creating a new empty directory. The target directory of a TAF script has the same filename prefix as the script: `data.R` creates ‘`data`’ etc. An important exception is that a directory called ‘`begin`’ will never be deleted.

**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 after running a script, 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.colors

*TAF Colors*


---

### Description

Predefined colors for 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  
barplot(5:1, col=c(taf.green, taf.orange, taf.blue, taf.dark, taf.light))
```

---

taf.skeleton

*TAF Skeleton*

---

## Description

Create an empty template for a new TAF analysis: initial directories and R scripts with TAF header comments.

## Usage

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

## Arguments

name	main directory name for the analysis.
path	path to create the analysis directory in.
force	whether to overwrite an existing directory.

## Details

Use name = "." to create initial directories and scripts inside the current working directory.

## 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)
```

---

taf2long	<i>Convert TAF Table to Long Format</i>
----------	---

---

### 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 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.

[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 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)
```

---

tafpng	<i>PNG Device</i>
--------	-------------------

---

### Description

Open PNG graphics device to create an image file inside the TAF report folder.

### Usage

```
tafpng(filename, width = 800, height = 600, pointsize = 22, ...)
```

### Arguments

filename	image filename.
width	image width.
height	image height.
pointsize	text size.
...	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/myplot.png", width=800, height=600, pointsize=22)
```

to

```
tafpng("myplot")
```

To successfully export trellis and ggplot objects to image files in scripts, these objects should be explicitly printed (see examples below).

### See Also

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

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

## Examples

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

library(lattice)
tafpng("mytrellis")
print(xyplot(1~1))
dev.off()

library(ggplot2)
tafpng("myggplot")
print(qplot(1))
dev.off()

## End(Not run)
```

---

tt

*TAF Transpose*

---

## 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**

[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.dls

*Write DLS3.2 Results to File*

---

**Description**

Write results from the DLS3.2 advisory method to a file.

**Usage**

```
write.dls(x, file = "")
```

**Arguments**

x	a list generated by the DLS3.2 function.
file	a filename.

**Note**

The resulting text file has Dos line endings (CRLF).

**See Also**

[read.dls](#) reads DLS3.2 results from a file back into R.

[DLS3.2](#) in the **icesAdvice** package can be used to calculate catch advice for data-limited stocks (DLS).

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

**Examples**

```
## Not run:
survey <- data.frame(year=2001:2010, randu[1:10,])
dls <- icesAdvice::DLS3.2(1000, survey$y)

write.dls(dls, "dls.txt")
read.dls("dls.txt")

file.remove("dls.txt")

## End(Not run)
```

---

write.taf	<i>Write TAF Table to File</i>
-----------	--------------------------------

---

**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**

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

**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.

**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)
```

---

xtab2taf

*Convert Crosstab Table to TAF Format*

---

**Description**

Convert a table from crosstab format to TAF format.

**Usage**

```
xtab2taf(x)
```

**Arguments**

x                    a data frame in crosstab format.

**Value**

A data frame in TAF format.

**Note**

TAF stores tables as data frames 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.

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

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

**Examples**

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

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