

Package ‘icesTAF’

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Title Functions to Support the ICES Transparent Assessment Framework

Depends TAF

Imports purrr, roxygen2

Suggests git2r

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

URL <https://taf.ices.dk>

Encoding UTF-8

RoxygenNote 7.2.1

NeedsCompilation no

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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.script</code>	create bootstrap script
<code>taf.roxygenise</code>	create DATA.bib entry from tags

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.

add.data.script*Import a bootstrap data script from ICES datasets repo***Description**

Download an ‘R’ file from the ICES datasets repo to fetch data including adding metadata via roxygen2 fields to the top of the file.

Usage

```
add.data.script(name, install.deps = TRUE, commit = FALSE)
```

Arguments

<code>name</code>	the name of the dataset.
<code>install.deps</code>	install packages used in the script if not already installed.
<code>commit</code>	should the bootstrap file be added and committed to the analysis.

Examples

```
## Not run:  
  
# Create bootstrap folder  
mkdir(taf.boot.path())  
  
# Create bootstrap script, bootstrap/mydata.R  
add.data.script(name = "vms")  
  
# Create metadata, bootstrap/DATA.bib  
taf.roxygenise(files = "vms.R")  
  
# Run bootstrap script, creating bootstrap/data/vms/...  
taf.bootstrap()  
  
## 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

<code>name</code>	the name of the dataset and the file name that will be created.
<code>title</code>	description of the data, including survey names or the like.
<code>description</code>	a more involved description if required.
<code>format</code>	the format of the data produced, e.g. “csv”
<code>originator</code>	who prepared the data, e.g. a working group acronym.
<code>year</code>	year of the analysis when the data were used. The default is the current year.
<code>period</code>	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.
<code>access</code>	data access code: “OSPAR”, “Public”, or “Restricted”.
<code>content</code>	the R code that fetches and saves the data

Examples

```
## Not run:

# Create bootstrap folder
mkdir("bootstrap")

# Create bootstrap script, bootstrap/mydata.R
draft.data.script(name="mydata", title="Title", description="Description",
                   format="txt", originator="Me", year="2022",
                   period=c(2000,2020), access="Public",
                   content='write(pi, file="pi.txt")')

# Create metadata, bootstrap/DATA.bib
taf.roxygenise(files="mydata.R")

# Run bootstrap script, creating bootstrap/data/mydata/pi.txt
taf.bootstrap()

## End(Not run)
```

install.deps

Install packages dependencies of a TAF analysis

Description

Search R scripts for packages that are required and install them.

Usage

```
install.deps(...)
```

Arguments

...	arguments passed on to install.packages
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See Also

[deps](#)

Examples

```
## Not run:

library(icesTAF)

# Download a TAF analysis
download("https://github.com/ices-taf/2019_san.sa.6/archive/refs/heads/master.zip")
unzip("master.zip")
```

```
# move into analysis folder  
setwd("2019_san.sa.6-master")  
  
# list dependencies  
deps()  
  
# install dependencies  
install.deps()  
  
## End(Not run)
```

taf.roxygenise *Process a TAF repo with the taf roclt*

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.

Examples

```
## Not run:  
  
# Create bootstrap folder  
mkdir("bootstrap")  
  
# Create bootstrap script, bootstrap/mydata.R  
draft.data.script(name="mydata", title="Title", description="Description",  
    format="txt", originator="Me", year="2022",  
    period=c(2000,2020), access="Public",  
    content='write(pi, file="pi.txt")')  
  
# Create metadata, bootstrap/DATA.bib  
taf.roxygenise(files="mydata.R")  
  
# Run bootstrap script, creating bootstrap/data/mydata/pi.txt  
taf.bootstrap()  
  
## End(Not run)
```

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