

Package ‘icesSAG’

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Suggests testthat, icesVocab

Description R interface to access the web services of the ICES Stock Assessment
Graphs database <<https://sg.ices.dk>>.

License GPL (>= 2)

URL <https://sg.ices.dk>, <https://github.com/ices-tools-prod/icesSAG>

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icesSAG-package	<i>Stock Assessment Graphs Database Web Services</i>
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Description

R interface to access the web services of the ICES Stock Assessment Graphs database.

Details

Get dataset:

getSummaryTable	summary results
getFishStockReferencePoints	reference points
getSAG	any data

Look up codes:

findAssessmentKey	find assessment key
getListStocks	list of stocks

Author(s)

Colin Millar, Scott Large, Arni Magnusson, Carlos Pinto and Laura Andreea Petre.

References

ICES Stock Assessment Graphs database: <http://sg.ices.dk>.
ICES Stock Assessment Graphs web services: <http://sg.ices.dk/webservices.aspx>.

See Also

Useful links:

- <https://sg.ices.dk>
- <https://github.com/ices-tools-prod/icesSAG>
- Report bugs at <https://github.com/ices-tools-prod/icesSAG/issues>

`convertSAGxml`*Create and read the SAG XML data transfer file*

Description

Convert between R data (a list and a data.frame) and the XML format required for uploading data to the SAG database.

Usage

```
createSAGxml(info, fishdata)

readSAGxml(file)

writeSAGxml(info, fishdata, file)
```

Arguments

<code>info</code>	a list of stock information
<code>fishdata</code>	a data frame of fish data
<code>file</code>	an xml file name

Value

Either a list containing `info` and `fishdata`, or a string containing the xml file.

See Also

[stockInfo](#) creates a list of stock information.
[stockFishdata](#) creates a data frame of fish stock summary data.

Examples

```
info <- stockInfo(StockCode = "cod.27.347d",
                  AssessmentYear = 2017,
                  StockCategory = 1,
                  ModelType = "A",
                  ModelName = "SCA",
                  ContactPerson = "itsme@fisheries.com")
fishdata <- stockFishdata(Year = 1990:2017, Catches = 100)
```

```
xmlfile <- createSAGxml(info, fishdata)

out <- readSAGxml(xmlfile)
```

findAssessmentKey	<i>Find a Key</i>
-------------------	-------------------

Description

Find a lookup key corresponding to a stock in a given assessment year.

Usage

```
findAssessmentKey(
  stock = NULL,
  year = 0,
  published = TRUE,
  regex = TRUE,
  full = FALSE
)
```

Arguments

stock	a stock name, e.g. cod-347d, or cod to find all cod stocks, or NULL (default) to process all stocks.
year	the assessment year, e.g. 2015, or 0 to process all years.
published	whether to include only years where status is "Published" (applies only when non-secure web services are in use, secure web service always returns unpublished stocks).
regex	whether to match the stock name as a regular expression.
full	whether to return a data frame with all stock list columns.

Value

A vector of keys (default) or a data frame if full is TRUE.

Author(s)

Arni Magnusson and Colin Millar.

See Also

[getListStocks](#) gets a list of stocks.
[icesSAG-package](#) gives an overview of the package.

Examples

```
## Not run:
findAssessmentKey("had.27.46a20", 2023, full = TRUE)

## End(Not run)
```

`getCustomColumns`*Get the Custom Columns for SAG records*

Description

Get custom columns, such as alternative biomass series or Fproxy reference points for records in the SAG database.

Usage

```
getCustomColumns(assessmentKey, ...)
```

Arguments

`assessmentKey` the unique identifier of the stock assessment
`...` arguments passed to [sag_get](#).

Value

A data frame.

Author(s)

Colin Millar.

See Also

[getSAG](#) supports querying many years and quarters in one function call.
[getListStocks](#) and [getFishStockReferencePoints](#) get a list of stocks and reference points.
[icesSAG-package](#) gives an overview of the package.

Examples

```
## Not run:
assessmentKey <- findAssessmentKey("bli.27.5a14")
customs <- getCustomColumns(assessmentKey)
head(customs)

## End(Not run)
```

getFishStockReferencePoints
Get Reference Points

Description

Get biological reference points for all stocks in a given assessment year.

Usage

```
getFishStockReferencePoints(assessmentKey, ...)
```

Arguments

assessmentKey the unique identifier of the stock assessment
... arguments passed to [sag_get](#).

Value

A data frame.

Author(s)

Colin Millar.

See Also

[getSAG](#) supports querying many years and quarters in one function call.
[getListStocks](#) and [getSummaryTable](#) get a list of stocks and summary results.
[icesSAG-package](#) gives an overview of the package.

Examples

```
## Not run:
assessmentKey <- findAssessmentKey("cod.27.21", year = 2023)
refpts <- getFishStockReferencePoints(assessmentKey)
refpts

# To get all reference points in a given assessment year:
keys2022 <- findAssessmentKey(year = 2022, full = TRUE)
keys2022 <- keys2022[keys2022$Purpose == "Advice",]
refpts2022 <- getFishStockReferencePoints(keys2022$AssessmentKey)
refpts2022

## End(Not run)
```

`getLatestStockAdviceList`*Get List of Most Recent Advice*

Description

Get a list of the most recent advice for all fish stocks.

Usage

```
getLatestStockAdviceList(...)
```

Arguments

... arguments passed to [sag_get](#).

Value

A data frame.

Author(s)

Colin Millar, Scott Large, and Arni Magnusson.

See Also

[getSummaryTable](#) gets a summary table of historical stock size.

[getFishStockReferencePoints](#) gets biological reference points.

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

Examples

```
## Not run:  
stocks <- getLatestStockAdviceList()  
  
## End(Not run)
```

getListStocks*Get a List of Fish Stocks*

Description

Get a list of fish stocks for a given assessment year.

Usage

```
getListStocks(year, stock = NULL, modifiedAfter = NULL, ...)
```

Arguments

year	the assessment year, e.g. 2015, or 0 to process all years.
stock	a stock name, e.g. lin.27.5a.
modifiedAfter	date-time parameter in the format "YYYY/MM/DD". If set will only return stocks assessments modified after the provided date.
...	arguments passed to sag_get .

Value

A data frame.

Author(s)

Colin Millar.

See Also

[getSummaryTable](#) gets a summary table of historical stock size.
[getFishStockReferencePoints](#) gets biological reference points.
[icesSAG-package](#) gives an overview of the package.

Examples

```
## Not run:  
stocks <- getListStocks(2023)  
nshad <- getListStocks(stock = "had.27.46a20")  
  
## End(Not run)
```

getSAG*Get Any SAG Data*

Description

This function combines the functionality of [getListStocks](#), [getFishStockReferencePoints](#), [getSummaryTable](#) and [getStockDownloadData](#). It supports querying many stocks and years in one function call.

Usage

```
getSAG(stock, year, data = "summary", combine = TRUE, purpose = "Advice")
```

Arguments

stock	a stock name, e.g. cod-347d, or cod to find all cod stocks, or NULL to process all stocks.
year	the assessment year, e.g. 2015, or 0 to process all years.
data	the data of interest, either "summary", "refpts" or "source".
combine	whether to combine the list output to a data frame.
purpose	the purpose of the entry, options are "Advice", "Bench", "InitAdvice", default is "Advice".

Value

A data frame (default) or a list if combine is TRUE.

Note

Only years with "Published" status are returned.

Author(s)

Colin Millar.

See Also

[getListStocks](#), [getSummaryTable](#), [getFishStockReferencePoints](#), and [getStockDownloadData](#) get a list of stocks, summary results, reference points, and all data including custom columns.

[findAssessmentKey](#) finds lookup keys.

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

Examples

```
## Not run:
summary <- getSAG("had.27.46a20", 2022)
refpts <- getSAG("had.27.46a20", 2022, "refpts")

cod_summary <- getSAG("cod", 2022)
cod_refpts <- getSAG("cod", 2015:2016, "refpts")
cod_data <- getSAG("cod", 2017, "source-data")

## End(Not run)
```

getSAGGraphs

*Get Summary Graphs of Stock Assessment Output***Description**

Get summary graphs of catches, recruitment, fishing pressure, and spawning stock biomass.

Usage

```
getSAGGraphs(assessmentKey, ...)
```

Arguments

`assessmentKey` the unique identifier of the stock assessment
`...` to allow scope for back compatibility

Value

An array representing a bitmap.

Author(s)

Colin Millar and Scott Large.

See Also

[getListStocks](#) gets a list of stocks.
[getFishStockReferencePoints](#) gets biological reference points.
[icesSAG-package](#) gives an overview of the package.

Examples

```
## Not run:
assessmentKey <- findAssessmentKey("cod", 2015)
graphs <- getSAGGraphs(assessmentKey[1])
plot(graphs)
# note this stock only has one graph see:
# http://standardgraphs.ices.dk/ViewCharts.aspx?key=8309

## End(Not run)
```

getSAGTypegraphsandSettings

Get Details on SAG Charts and Settings

Description

List all possible chart settings for each chart type (0 = general, 1 = Landings, ...).

Usage

```
getSAGTypeGraphs(...)

getSAGTypeSettings(SAGChartKey, ...)
```

Arguments

...	arguments passed to sag_get .
SAGChartKey	the type identifier of the SAG chart, e.g. 0, 1, 2, ...

Value

a data frame with SAG chart type IDs and settings IDs.

Examples

```
## Not run:
getSAGTypeGraphs()

getSAGTypeSettings(0)[-4]

## End(Not run)
```

getsetSAGSettingsForASTock

Get and Set SAG Chart Settings

Description

details

Usage

```
getSAGSettingsForASTock(assessmentKey, ...)

setSAGSettingForASTock(
  assessmentKey,
  chartKey,
  settingKey,
  settingValue,
  copyNextYear,
  ...
)
```

Arguments

assessmentKey	the unique identifier of the stock assessment
...	arguments passed to sag_get .
chartKey	the type identifier of the SAG chart, e.g. 0, 1, 2, ...
settingKey	the type identifier of the SAG chart setting, e.g. 0, 1, 2, ...
settingValue	the value of the setting
copyNextYear	should the settings be copied to next year (TRUE) or not (FALSE)

Value

A data frame with SAG chart type IDs, settings IDs and setting values.

Examples

```
## Not run:
key <- findAssessmentKey("cod.21.1", 2017)
graphs <- getSAGGraphs(key[1])
plot(graphs)
getSAGSettingsForASTock(key [1])
chart1 <- getLandingsGraph(key [1])
setSAGSettingForASTock(key [2], 1, 1, "Catches of cod.21.1 in 2017",
FALSE)
setSAGSettingForASTock(key [2], 1, 11, 10,
FALSE)
plot(chart1)
```

```
chart2 <- getSpawningStockBiomassGraph(key [1])
plot(chart2)
setSAGSettingForASTock(key [1], 4, 1, "SSB of cod.21.1 in 2017",
FALSE)
plot(chart2)

## End(Not run)
```

`getStandardAssessmentGraphs`*Get a Graph of Stock Assessment Output*

Description

Get a graph of stock assessment output, e.g., historical stock size, recruitment, and fishing pressure.

Usage

```
getLandingsGraph(assessmentKey, ...)
getRecruitmentGraph(assessmentKey, ...)
getFishingMortalityGraph(assessmentKey, ...)
getSpawningStockBiomassGraph(assessmentKey, ...)
getFishMortality(assessmentKey, ...)
getstock_recruitment(assessmentKey, ...)
getYSSB(assessmentKey, ...)
getSSBHistoricalPerformance(assessmentKey, ...)
getFishingMortalityHistoricalPerformance(assessmentKey, ...)
getRecruitmentHistoricalPerformance(assessmentKey, ...)
getStockStatusTable(assessmentKey, ...)
```

Arguments

<code>assessmentKey</code>	the unique identifier of the stock assessment
<code>...</code>	to allow scope for back compatibility

Value

An array representing a bitmap.

See Also

[getListStocks](#) gets a list of stocks.

[getFishStockReferencePoints](#) gets biological reference points.

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

Examples

```
## Not run:
assessmentKeys <- findAssessmentKey("had", 2015)
landings_img <- getLandingsGraph(assessmentKeys[1])
landings_img

landings_plots <- getLandingsGraph(assessmentKeys)
landings_plots

## End(Not run)
```

getStockSourceData	<i>Get Source Data</i>
--------------------	------------------------

Description

Get a copy of the source data for the specified stocks.

Usage

```
getStockDownloadData(assessmentKey, ...)
```

Arguments

assessmentKey	the unique identifier of the stock assessment, can be a vector
...	arguments passed to sag_get .

Value

A data frame.

Author(s)

Colin Millar.

See Also

[getSAG](#) supports querying many years in one function call.

[getListStocks](#) and [getFishStockReferencePoints](#) get a list of stocks and reference points.

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

Examples

```
## Not run:
assessmentKey <- findAssessmentKey("cod-2224", year = 2016)
sourcedat <- getStockDownloadData(assessmentKey)
head(sourcedat[[1]])

## End(Not run)
```

getStockStatusValues	<i>Get the Values in a Stock Status Table</i>
----------------------	---

Description

Get summary results of historical stock size, recruitment, and fishing pressure.

Usage

```
getStockStatusValues(assessmentKey, ...)
```

Arguments

assessmentKey	the unique identifier of the stock assessment
...	arguments passed to sag_get .

Value

A data frame.

Author(s)

Colin Millar.

See Also

[getSAG](#) supports querying many years and quarters in one function call.

[getListStocks](#) and [getFishStockReferencePoints](#) get a list of stocks and reference points.

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

Examples

```
## Not run:
assessmentKey <- findAssessmentKey("had.27.46a20", year = 2022)
status <- getStockStatusValues(assessmentKey)
status

## End(Not run)
```

getSummaryTable	<i>Get a Summary Table of Historical Stock Size</i>
-----------------	---

Description

Get summary results of historical stock size, recruitment, and fishing pressure.

Usage

```
getSummaryTable(assessmentKey, ...)
```

Arguments

assessmentKey the unique identifier of the stock assessment
 ... arguments passed to [sag_get](#).

Value

A data frame.

Author(s)

Colin Millar.

See Also

[getSAG](#) supports querying many years and quarters in one function call.
[getListStocks](#) and [getFishStockReferencePoints](#) get a list of stocks and reference points.
[icesSAG-package](#) gives an overview of the package.

Examples

```
## Not run:
assessmentKey <- findAssessmentKey("had.27.46a20", year = 2022)
sumtab <- getSummaryTable(assessmentKey)
head(sumtab)

## End(Not run)
```

sag_api	<i>Build a SAG web service url</i>
---------	------------------------------------

Description

utility to build a url with optional query arguments

Usage

```
sag_api(service, ...)
```

Arguments

service	the name of the service
...	name arguments will be added as queries

Value

a complete url as a character string

Examples

```
sag_api("hi", bye = 21)
sag_api("StockList", year = 2021)
```

sag_get	<i>Get a url</i>
---------	------------------

Description

Get a url, optionally using an ICES authentication token

Usage

```
sag_get(
  url,
  retry = TRUE,
  quiet = !getOption("icesSAG.messages"),
  verbose = FALSE,
  content = TRUE,
  use_token = getOption("icesSAG.use_token")
)

sag_get_cached(
  url,
```

```

    retry = TRUE,
    quiet = !getOption("icesSAG.messages"),
    verbose = FALSE,
    content = TRUE,
    use_token = getOption("icesSAG.use_token")
  )

```

Arguments

url	the url to get.
retry	should the get request be retried if first attempt fails? default TRUE.
quiet	should all messages be suppressed, default FALSE.
verbose	should verbose output from the http request be returned? default FALSE.
content	should content be returned, or the full http response? default TRUE, i.e. content is returned by default.
use_token	should an authentication token be sent with the request? default is the value of the option icesSAG.use_token.

Value

content or an http response.

Functions

- `sag_get_cached()`: cached version of `sag_get`

See Also

[sag_api](#) builds a SAG web service url.
[icesSAG-package](#) gives an overview of the package.

Examples

```

## Not run:
sag_get(sag_api("StockList", year = 2022))

## End(Not run)

```

sag_post

Post to a url

Description

Post to a url using an ICES authentication token

Usage

```
sag_post(url, body = list(), verbose = FALSE)
```

Arguments

url	the url to get.
body	a list of named arguments to be sent as the body of the post request.
verbose	should verbose output from the http request be returned? default FALSE.

Value

content of the http response.

See Also

[sag_api](#) builds a SAG web service url.
[icesSAG-package](#) gives an overview of the package.

setPackageOptions	<i>Get and Set icesSAG package options</i>
-------------------	--

Description

There are two options of interest, 1) switch on or off the use of authenticated web service calls, and 2) switch on or off the display of messages to the console.

Usage

```
sag_use_token(value)  
  
sag_messages(value)
```

Arguments

value	TRUE or FALSE
-------	---------------

Value

invisible return of the old value.

Examples

```
## Not run:  
sag_use_token(TRUE)  
sag_messages(TRUE)  
  
## End(Not run)
```

stockFishdata	<i>Create a data.frame of fish stock data</i>
---------------	---

Description

This function is a wrapper to `data.frame(...)` in which the names are forced to match with the names required for the SAG database. See <http://dome.ices.dk/datsu/selRep.aspx?Dataset=126> for more details.

Usage

```
stockFishdata(Year, ...)
```

Arguments

Year	a vector of years.
...	additional information, e.g. Recruitment, StockSize, Landings, ...

Value

A data.frame, where all names are valid column names in the SAG database.

Author(s)

Colin Millar.

Examples

```
stockFishdata(Year = 1990:2017, Catches = 100)
```

stockInfo	<i>Create a list of fish stock information</i>
-----------	--

Description

This function is a wrapper to `list(...)` in which the names are forced to match with the names required for the SAG database. See <http://dome.ices.dk/datsu/selRep.aspx?Dataset=126> for more details.

Usage

```
stockInfo(
  StockCode,
  AssessmentYear,
  ContactPerson,
  StockCategory,
  Purpose = "Advice",
  ModelType,
  ModelName,
  ...
)
```

Arguments

StockCode	a stock name, e.g. cod-347d.
AssessmentYear	the assessment year, e.g. 2015.
ContactPerson	the email for the person responsible for uploading the stock data.
StockCategory	Category of the assessment used (see below)
Purpose	the purpose of the entry, options are "Advice", "Bench", "InitAdvice", default is "Advice".
ModelType	the type of the model used (see below for links to more information)
ModelName	the name (acronym) of the model used if available (see below for links to more information)
...	additional information, e.g. BMGT, FMSY, RecruitmentAge, ...

Value

A named `sag.list`, inheriting from a list, where all names are valid column names in the SAG database.

Author(s)

Colin Millar.

See Also

Links to the relevant ICES vocabularies list are here StockCode: <https://vocab.ices.dk/?ref=357> StockCategory: <https://vocab.ices.dk/?ref=1526> Purpose: <https://vocab.ices.dk/?ref=1516> ModelType: <https://vocab.ices.dk/?ref=1524> ModelName: <https://vocab.ices.dk/?ref=1525>

Link to the relevant format description is <https://datsu.ices.dk/web/selfRep.aspx?Dataset=126>

Examples

```

info <-
  stockInfo(StockCode = "cod.27.47d20",
            AssessmentYear = 2017,
            StockCategory = 1,
            ModelType = "A",
            ModelName = "SCA",
            ContactPerson = "itsme@fisheries.com")

info
info$mistake <- "oops"
info
# should have gotten a warning message

## Not run:
# use icesVocab to list valid codes etc.
library(icesVocab)
# print the list of valid stock codes
stock.codes <- getCodeList("ICES_StockCode")
stock.codes[1:10,1:2]

# print the list of assessment model types in the ICES vocabulary
model.types <- getCodeList("AssessmentModelType")
model.types[1:2]

# print the list of assessment model names in the ICES vocabulary
model.names <- getCodeList("AssessmentModelName")
model.names$Key

## End(Not run)

```

uploadStock

Upload New or Updated Fish Stock Assessment Results

Description

Get summary results of historical stock size, recruitment, and fishing pressure.

Usage

```
uploadStock(file, upload = FALSE, verbose = FALSE)
```

Arguments

file	the xml file to check and upload
upload	should the file be uploaded to the database? Default is FALSE.
verbose	if TRUE more verbose messages are reported

Value

The database key of the new / updated stock, or 0 if there was an error.

Author(s)

Colin Millar.

See Also

[stockInfo](#) creates a list of stock information.

[stockFishdata](#) creates a data.frame of fish stock summary data.

Examples

```
## Not run:
info <-
stockInfo(
  StockCode = "whb-comb",
  AssessmentYear = 1996,
  ContactPerson = "its_me@somewhere.gov",
  StockCategory = 3,
  Purpose = "InitAdvice",
  ModelType = "A",
  ModelName = "XSA"
)
fishdata <- stockFishdata(1950:1996)
# simulate some landings for something a bit interesting
set.seed(1232)
fishdata$Landings <- 10^6 * exp(cumsum(cumsum(rnorm(nrow(fishdata), 0, 0.1))))

# You should write out to an xml file and then upload that file
tempfile <- tempfile(fileext = ".xml")
writeSAGxml(info, fishdata, file = tempfile)

# you can check the file is formatted correctly
uploadStock(tempfile, upload = FALSE)

# and if all checks pass, the file can be uploaded
uploadStock(tempfile, upload = TRUE)

## End(Not run)
```

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