

# Package: icesSAG (via r-universe)

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**Title** Stock Assessment Graphs Database Web Services

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**Suggests** testthat

**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>

**BugReports** <https://github.com/ices-tools-prod/icesSAG/issues>

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icesSAG-package      *Stock Assessment Graphs Database Web Services*

---

### Description

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

### Details

*Get dataset:*

<code>getSummaryTable</code>	summary results
<code>getFishStockReferencePoints</code>	reference points
<code>getSAG</code>	any data

*Look up codes:*

<code>findAssessmentKey</code>	find assessment key
<code>getListStocks</code>	list of stocks

### Author(s)

Colin Millar, Scott Large, and Arni Magnusson.

## References

ICES Stock Assessment Graphs database: <http://sg.ices.dk>.

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

---

convertSAGxml	<i>Create and read the SAG XML data transfer file</i>
---------------	---

---

## 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)
```

## Arguments

info	a list of stock information
fishdata	a data frame of fish data
file	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      *Find a Key*

---

### 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

[StockList](#) 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)
```

---

FishStockReferencePoints

*Get Reference Points*

---

## Description

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

## Usage

```
FishStockReferencePoints(assessmentKey, ...)
```

## Arguments

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

## Value

A data frame.

## Author(s)

Colin Millar.

## See Also

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

[StockList](#) and [SummaryTable](#) 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 <- FishStockReferencePoints(assessmentKey)
refpts

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

## 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 [ices\\_get](#).

### Value

A data frame.

### Author(s)

Colin Millar.

### See Also

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

[StockList](#) and [FishStockReferencePoints](#) 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  
... to allow scope for back compatibility

**Value**

A data frame.

**Author(s)**

Colin Millar and Scott Large.

**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-2224", year = 2016)  
refpts <- getFishStockReferencePoints(assessmentKey)  
refpts  
  
#To get all reference points in a given assessment year:  
keys2016 <- findAssessmentKey(year = 2016)  
refpts2016 <- getFishStockReferencePoints(keys2016)  
refpts2016  
  
## 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()
```

**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)
```



**Arguments**

year                    the assessment year, e.g. 2015, or 0 to process all years.

**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 <- getListStocks(2015)

## End(Not run)
```

---

getSAG

*Get Any SAG Data*


---

**Description**

This function combines the functionality of `getListStocks`, `getFishStockReferencePoints`, and `getSummaryTable`. 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**

[StockList](#), [SummaryTable](#), and [FishStockReferencePoints](#) get a list of stocks, summary results, and reference points.

[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 etc.).

**Usage**

```
getSAGTypeGraphs()
```

```
getSAGTypeSettings(SAGChartKey)
```

**Arguments**

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
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)
```

---

getsetStockChartSettings

*Get and Set SAG Chart Settings*


---

**Description**

details

**Usage**

```
StockSettings(assessmentKey, ...)
```

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

**Arguments**

assessmentKey the unique identifier of the stock assessment  
... arguments passed to [ices\\_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:
assessmentKey <- findAssessmentKey("had.27.46a20", year = 2022)

StockSettings(assessmentKey)

## 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**

assessmentKey the unique identifier of the stock assessment  
... 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])  
plot(landings_img)  
  
landings_plots <- getLandingsGraph(assessmentKeys)  
plot(landings_plots)  
  
## End(Not run)
```

---

getStockSourceData      *Get Source Data*

---

**Description**

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

**Usage**

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

**Arguments**

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

**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 *Get the Values in a Stock Status Table*

---

**Description**

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

**Usage**

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

**Arguments**

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

**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("cod-2224", year = 2016)
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
...	to allow scope for back compatibility

**Value**

A data frame.

**Author(s)**

Colin Millar and Scott Large.

**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("cod-2224", year = 2016)
sumtab <- getSummaryTable(assessmentKey)
head(sumtab)
attributes(sumtab)$notes

## End(Not run)
```



**Value**

A data frame.

**Author(s)**

Colin Millar and Scott Large.

**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("cod-2224", year = 2015)
sumtab <- getYSBRSummaryTable(assessmentKey)
head(sumtab)

## End(Not run)
```

---

get_custom_plot	<i>Get a Custom Graph of Stock Assessment Output</i>
-----------------	--

---

**Description**

Get a custom graph of stock assessment output.

**Usage**

```
get_custom_plot(assessmentKey, type = c(15, 16, 17, 18), width = 800, ...)
```

**Arguments**

assessmentKey	the unique identifier of the stock assessment
type	The type of plot: values can be 15, 16, 17, or 18.
width	width of the image in pixels
...	to allow scope for back compatibility

**Value**

An array representing a bitmap.

**See Also**

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

[FishStockReferencePoints](#) gets biological reference points.

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

**Examples**

```
## Not run:
assessmentKey <- findAssessmentKey("ane.27.9a", year = 2022)
landings_img <- get_custom_plot(assessmentKey, "15")

plot(landings_img)

landings_plots <- get_image(assessmentKey, "landings")
plot(landings_plots)

## End(Not run)
```

---

get\_image

*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**

```
get_image(
  assessmentKey,
  type = c("landings", "recruitment", "ssb", "mortality", "historical_mortality",
           "historical_ssb", "historical_recruitment"),
  width = 800,
  ...
)
```

**Arguments**

assessmentKey	the unique identifier of the stock assessment
type	The type of plot: values can be landings, recruitment ssb, mortality, historical_mortality, historical_ssb, historical_recruitment.
width	width of the image in pixels
...	to allow scope for back compatibility

**Value**

An array representing a bitmap.

**See Also**

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

[FishStockReferencePoints](#) gets biological reference points.

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

**Examples**

```
## Not run:
assessmentKey <- findAssessmentKey("cod.27.21", year = 2022:2023)
landings_img <- get_image(assessmentKey[1], "landings")

plot(landings_img)

landings_plots <- get_image(assessmentKey, "landings")
plot(landings_plots)

## End(Not run)
```

---

ices\_get

*Get a url*

---

**Description**

Get a url, optionally using an ICES authentication token

**Usage**

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

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

**Arguments**

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

**Value**

content or an http response.

**Functions**

- `ices_get_cached()`: cached version of `ices_get`

**See Also**

[sag\\_api](#) builds a SAG web service url.

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

**Examples**

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

## End(Not run)
```

---

LatestStockAdviceList *Get a List of the Latest Fish Stock Assessments*

---

**Description**

Get a list of the latest fish stock assessments. Only published stocks are returned.

**Usage**

```
LatestStockAdviceList(...)
```

**Arguments**

... arguments passed to [ices\\_get](#).

**Value**

A data frame.

**See Also**

[StockList](#) Get a list of fish stock assessments.  
[getFishStockReferencePoints](#) gets biological reference points.  
[icesSAG-package](#) gives an overview of the package.

**Examples**

```
## Not run:  
stock_list <- LatestStockAdviceList()  
  
## End(Not run)
```

---

SAGTypeChartsandSettings

*Get Details on SAG Charts and Settings*

---

**Description**

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

**Usage**

```
SAGTypeCharts(...)  
  
SettingsForChartType(SAGChartKey, ...)
```

**Arguments**

... arguments passed to [ices\\_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:  
SAGTypeCharts()  
  
SettingsForChartType(0)[-4]  
  
## 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)
```

---

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

---

**Description**

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

**Usage**

```
StockDownload(assessmentKey, ...)
```

**Arguments**

assessmentKey	the unique identifier of the stock assessment, can be a vector
...	arguments passed to <a href="#">ices_get</a> .

**Value**

A data frame.



**Author(s)**

Colin Millar.

**See Also**

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

[StockList](#) 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)
sourcedat <- StockDownload(assessmentKey)
head(sourcedat)

## End(Not run)
```

---

stockFishdata

*Create a data.frame of fish stock data*

---

**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

*Create a list of fish stock information*


---

### 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/se1Rep.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)
```

---

 StockList

---

*Get a List of Fish Stock Assessments*


---

**Description**

Get a list of fish stock assessments for a given assessment year. If an authentication token is not provided, only published stocks are returned.

**Usage**

```
StockList(year, modifiedAfter = NULL, ...)
```

**Arguments**

`year` the assessment year, e.g. 2022, or 0 to process all years.  
`modifiedAfter` date-time parameter in the format "". If set will only return stocks assessments modified after the provided date.  
`...` arguments passed to [ices\\_get](#).

**Value**

A data frame.

**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:
stock_list <- StockList(year = 2022)

## End(Not run)
```

---

StockStatusValues      *Get the Values in a Stock Status Table*

---

**Description**

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

**Usage**

```
StockStatusValues(assessmentKey, ...)
```

**Arguments**

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

**Value**

A data frame.

**Author(s)**

Colin Millar.

**See Also**

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

[StockList](#) and [FishStockReferencePoints](#) 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 <- StockStatusValues(assessmentKey)
status

## End(Not run)
```

---

SummaryTable

*Get a Summary Table of Historical Stock Size*

---

**Description**

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

**Usage**

```
SummaryTable(assessmentKey, ...)
```

**Arguments**

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

**Value**

A data frame.

**Author(s)**

Colin Millar.

**See Also**

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

[StockList](#) and [FishStockReferencePoints](#) 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 <- SummaryTable(assessmentKey)
head(sumtab)

## 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(info, fishdata, verbose = FALSE)
```

**Arguments**

<code>info</code>	a list of stock information
<code>fishdata</code>	a data.frame of fish data
<code>verbose</code>	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.

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