

Package: ss3om (via r-universe)

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Title Tools for Conditioning Fisheries Operating Models Using Stock Synthesis 3

Version 0.5.2.9016

Description Tools for loading Stock Synthesis (SS3) models into FLR. Used in conditioning of Operating Models based on SS3 by considering structural uncertainty in input parameters and assumptions. A grid of SS3 runs can be created and results loaded on objects of various FLR classes.

X-schema.org-keywords fisheries, ss3, flr, R

Depends R (>= 3.3.2), FLCore (>= 2.6.5), r4ss (>= 1.43.0)

Imports methods, stats, data.table, foreach, FLFishery, mse, mvtnorm

Additional_repositories <https://flr.r-universe.dev>

Remotes <https://r4ss.r-universe.dev>

BugReports <https://github.com/flr/ss3om/issues>

Suggests doParallel, testthat, rlang, knitr, rmarkdown

License EUPL

VignetteBuilder knitr

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Repository <https://flr.r-universe.dev>

RemoteUrl <https://github.com/flr/ss3om>

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extractSSB	<i>Extracts derived quantities from SS output</i>
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Description

A number of derived quantities are available in the *Report.sso* file, and are useful for checking that the generated FLR objects lead to the same values. These functions are called on the list returned by *readOutputss3* or *r4ss::SS_output*, and extract the yearly values from the following rows in the *derived_quants* data.frame:

Usage

`extractSSB(out)`

`extractRec(out)`

`extractFbar(out)`

`extractZatage(out)`

`extractDevs(out)`

Arguments

`out` A list as returned by *r4ss::SS_output*.

Details

- *extractSSB*: *SSB_y* for *y* between *startyr* and *endyr*.
- *extractRec*: *Recr_y* for *y* between *startyr* and *endyr*.
- *extractFbar*: *F_y* for *y* between *startyr* and *endyr*.
- *extractZatage*: *Z_{ay}* for *y* between *startyr* and *endyr* and ages but last.

For 2 sex models (*nsexes*), *extractRec* will return a two-unit *FLQuant*, with the *Recr_y* values split according to the recruitment sex ratio. This is extracted from `recruitment_dist[[1]][, "Frac/sex"]`.

The value returned by *extractFbar* is the actual mean F over the age range, and the value in *derived_quants* is corrected according to *F_report_basis* for models where F reporting basis is F/FMSY.

Value

An *FLQuant* object of the requested quantity.

Examples

```

out <- readOutputss3(system.file("ext-data", "herring", package="ss3om"))

extractSSB(out)

extractFbar(out)

```

nameGrid	<i>nameGrid</i>
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Description

Creates folder names from a 'grid' df of scenarios

Usage

```
nameGrid(df, dir, from = 1)
```

Arguments

df	Model grid data.frame
dir	Folder name
from	Starting number

readFLIBss3	<i>A function to read the CPUE series from an SS3 run into an FLIndex object</i>
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Description

A function to read the CPUE series from an SS3 run into an FLIndex object

Usage

```

readFLIBss3(
  dir,
  fleets,
  birthseas = out$birthseas,
  repfile = "Report.sso",
  compfile = "CompReport.sso",
  ...
)

```

Arguments

dir Directory containing the SS3 output files
birthseas The birthseasons for this stock as a numeric vector.
... Any other argument to be passed to r4ss: :SS_output

Value

An object of class [FLStock](#)

Author(s)

Iago Mosqueira, EC JRC

References

Methot RD Jr, Wetzel CR (2013) Stock Synthesis: A biological and statistical framework for fish stock assessment and fishery management. Fisheries Research 142: 86-99.

See Also

[FLComp](#)

readFLSRss3	<i>A function to read the stock-recruit relationships from an SS3 run into an FLSR object</i>
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Description

A function to read the stock-recruit relationships from an SS3 run into an FLSR object

Usage

```
readFLSRss3(  
  dir,  
  birthseas = out$birthseas,  
  repfile = "Report.sso",  
  compfile = "CompReport.sso",  
  ...  
)
```

Arguments

dir Directory containing the SS3 output files
birthseas The birthseasons for this stock as a numeric vector.
... Any other argument to be passed to r4ss: :SS_output

Value

An object of class [FLStock](#)

Author(s)

Iago Mosqueira, EC JRC

References

Methot RD Jr, Wetzel CR (2013) Stock Synthesis: A biological and statistical framework for fish stock assessment and fishery management. Fisheries Research 142: 86-99.

See Also

[FLComp](#)

readFLSss3

A function to read SS3 results as an FLStock object

Description

Results of a run of the Stock Synthesis software, SS3 (Methot & Wetzel, 2013), can be loaded into an object of class [FLStock](#). The code makes use of the `r4ss::SS_output` function to obtain a list from `Report.sso`. The following elements of that list are used to generate the necessary information for the slots in [FLStock](#): "catage", "natage", "agesex", "endgrowth", "catch_units", "nsexes", "nseasons", "nareas", "IsFishFleet", "fleet_ID", "FleetNames", "spawnseas", "inputs" and "SS_version".

Usage

```
readFLSss3(
  dir,
  repfile = "Report.sso",
  compfile = "CompReport.sso",
  wtatage = out$wtatage_switch,
  ...
)
```

Arguments

<code>dir</code>	Directory holding the SS3 output files
<code>...</code>	Any other argument to be passed to <code>r4ss::SS_output</code>
<code>birthseas</code>	Birth seasons for this stock, defaults to <code>spawnseas</code>
<code>name</code>	Name of the output object to fill the name slot
<code>desc</code>	Description of the output object to fill the desc slot

Value

An object of class `\link{FLStock}`

Author(s)

The FLR Team

References

Methot RD Jr, Wetzel CR (2013) Stock Synthesis: A biological and statistical framework for fish stock assessment and fishery management. *Fisheries Research* 142: 86-99.

See Also

[FLComp](#)

 ss3slot

Functions to convert SS3 output into FLQuant(s)

Description

A series of auxiliary functions that convert one or more elements, typically of class `data.frame`, in the list returned by `r4ss::SS_output` into particular `FLQuant` or `FLQuants` objects.

Usage

```
ss3index(cpue, fleets)
```

```
ss3index.res(cpue, fleets)
```

```
ss3index.var(cpue, fleets)
```

```
ss3index.q(cpue, fleets)
```

```
ss3sel.pattern(selex, years, fleets, morphs, factor = "Ase12")
```

```
ss3wt(endgrowth, dmns, birthseas)
```

```
ss3mat(endgrowth, dmns, birthseas, option = 3)
```

```
ss3m(endgrowth, dmns, birthseas)
```

```
ss3n(n, dmns, birthseas)
```

```
ss3catch(catage, wtatage, dmns, birthseas, idx)
```

```
ss3mat30(endgrowth, dmns, spawnseas, option = 3)
```

```
ss3m30(endgrowth, dmns, birthseas)
```

```
ss3n30(n, dmns)
```

```
ss3catch30(catage, wtatage, dmns, birthseas, idx)
```

Arguments

cpue	A data frame obtained from SS_output\$cpue.
fleets	Named vector of fleets (numeric) codes
selex	A data frame obtained from SS_output\$ageselex.
years	Vector of years for which the index applies
morphs	Vector of morphs to use
endgrowth	A data frame obtained from SS_output\$endgrowth.
dmns	dimnames of the output object, usually obtained using getDimnames.
birthseas	The birthseasons for this stock as a numeric vector.
n	A data frame obtained from SS_output\$natage.
catage	A data frame obtained from SS_output\$catage.
wtatage	A data frame obtained from SS_output\$endgrowth but subset for birthseas and RetWt:_idx.
idx	The fishing fleets, as in SS_output\$fleet_ID[SS_output\$IsFishFleet].

Details

- ss3index returns the index slot of each survey/CPUE fleet.
- ss3index.res returns the index.res slot of each survey/CPUE fleet.
- ss3index.var returns the index.var slot of each survey/CPUE fleet.
- ss3index.q returns the index.q slot of each survey/CPUE fleet.
- ss3sel.pattern returns the sel.pattern slot of each survey/CPUE fleet.
- ss3wt returns the stock.wt slot.
- ss3mat returns the mat slot.
- ss3m returns the m slot.
- ss3m returns the m slot.
- ss3catch currently returns the landings.n slot, equal to catch.n as discards are not being parsed.
- ss3mat30 returns the mat slot.

- `ss3m` returns the `m` slot.
- `ss3n30` returns the `stock.n` slot.
- `ss3catch` currently returns the `landings.n` slot, equal to `catch.n` as discards are not being parsed.

Value

An `FLQuant` or `FLQuants` object, depending on the converted data structure

Author(s)

Iago Mosqueira, EC JRC D02

See Also

[FLQuant readFLSs3](#)

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