Package ‘AMPLE’

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Title Shiny Apps to Support Capacity Building on Harvest Control Rules

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Description Three Shiny apps are provided that introduce Harvest Control Rules (HCR) for fisheries management.

'Introduction to HCRs' provides a simple overview to how HCRs work. Users are able to select their own HCR and step through its performance, year by year. Biological variability and estimation uncertainty are introduced.

'Measuring performance' builds on the previous app and introduces the idea of using performance indicators to measure HCR performance.

'Comparing performance' allows multiple HCRs to be created and tested, and their performance compared so that the preferred HCR can be selected.

License GPL (>= 3)

URL https://github.com/PacificCommunity/ofp-sam-ample

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**Description**

Function used by `get_hcr_ip()` to generate input data for an assessment based HCR. The input to the HCR is depletion (i.e. Biomass / K).

**Usage**

```r
assessment(stock, mp_params, yr, iters = 1:dim(stock$biomass)[1])
```

**Arguments**

- `stock` The stock object
- `mp_params` A named list of MP parameters (with `est_sigma` and `est_bias` elements)
- `yr` The timestep that the biomass is taken from.
- `iters` Numeric vector of iters. Default is all of them.
comparing_performance  'Comparing HCR Performance' app launcher

Description

Launches the Comparing Performance Shiny app. See the 'Information' tab in the app for more information. Also see the package vignette(vignette("comparing_performance", package="AMPLE")) for a tutorial.

Usage

comparing_performance(...)

Arguments

... Not used

Examples

## Not run: comparing_performance()

constant  Evaluates a constant harvest control rule

Description

Evaluates a constant harvest control rule, i.e. one that ignores the stock status and just returns the constant level (catch or effort). Used by the hcr_op function.

Usage

constant(mp_params, ...)

Arguments

mp_params The HCR / management procedure parameters used to evaluate the HCR (as a list).
... Unused
estimation_error

Description

Estimation error applied to the 'true' stock status to generate an 'observed' stock status used in the HCR. The error is a combination of bias and lognormally distributed noise.

Usage

estimation_error(input, sigma, bias)

Arguments

input  A vector of the 'true' stock status
sigma  Observation error standard deviation
bias   Observation error bias

get_hcr_ip

Get the input to the HCR

Description

Run the MP analyses function to generate the input to the HCR i.e. observed stock status. For example, estimated biomass from an assessment.

Usage

get_hcr_ip(stock, mp_params, yr, ...)

Arguments

stock  The stock object
mp_params  The HCR / management procedure parameters used to evaluate the HCR (as a list).
yr  The time step of the true stock status used to generate the HCR IP .
...  Other arguments, including iters
get_hcr_op

Evaluates the harvest control rule.

Description
Evaluates the harvest control rule in a single year (timestep).

Usage
get_hcr_op(stock, mp_params, yr, iters = 1:dim(stock$biomass)[1])

Arguments
stock The stock object
mp_params The HCR / management procedure parameters used to evaluate the HCR (as a list).
yr The timestep.
iters A numeric vector of iters.

Value
A vector of outputs from the HCR.

intro_hcr
Introduction to HCRs app launcher

Description
Launches the introduction to HCRs Shiny app. See the ‘Information’ tab in the app for more information. Also see the package vignette (vignette("intro_hcr", package="AMPLE").)

Usage
intro_hcr(...)

Arguments
... Not used

Examples
## Not run: intro_hcr()
measuring_performance  Measuring performance app launcher

Description

Launches the 'Measuring Performance' Shiny app. See the 'Information' tab in the app for more information. Also see the package vignette (vignette("measuring_performance", package="AMPLE")) for a tutorial.

Usage

measuring_performance(...)

Arguments

...  Not used

Examples

## Not run: measuring_performance()

MP modules  mpParamsSetterUI

Description

The interface for the HCR options. The parameter selection inputs shown in the app are conditional on the selected type of HCR. Some of the inputs have initial values that can be set using the function arguments.

Does the setting part of the MP params module. Returns a list of MP params based on the MP inputs.

 Creates the MP params list based on the MP selection from the Shiny UI. Defined outside of a reactive environment above so we can use it non-reactively (helpful for testing).

Usage

mpParamsSetterUI(
  id,
  mp_visible = NULL,
  title = "Select the type of HCR you want to test.",
  init_thresh_max_catch = 140,
  init_thresh_belbow = 0.5,
  init_constant_catch = 50,
  init_constant_effort = 1
)
Stochasticity module

mpParamsSetterServer(id, get_stoch_params = NULL)

mp_params_switcheroo(input, est_sigma = 0, est_bias = 0)

Arguments

id The id (shiny magic)
mp_visible Which HCR types to show.
title The title.
init_thresh_max_catch Initial value of the maximum catch for the catch threshold HCR.
init_thresh_bellow Initial value of the bellow for the catch threshold HCR.
init_constant_catch Initial value of constant catch for the constant catch HCR.
init_constant_effort Initial value of constant effort for the constant effort HCR.
get_stoch_params Reactive expression that gets the parameters from the stochasticity setter. Otherwise est_sigma and est_bias are set to 0.
input List of information taken from the Shiny UI (mpParamsSetterUI)
est_sigma Standard deviation of the estimation variability (default = 0).
est_bias Estimation bias as a proportion. Can be negative (default = 0).

Value

A taglist
A list of HCR options.

Stochasticity module  stochParamsSetterUI

Description

stochParamsSetterUI() is the UI part for the stochasticity options. Stochasticity is included in the projections in two areas: biological variability (e.g. recruitment) and estimation error (to represent the difference between the 'true' status of the stock and the estimated status that is used by the HCR). Estimation error includes bias and variability. The arguments to this function allow only some of these elements to be shown.

stochParamSetterServer() does the server side stuff for the stochasticity options.

set_stoch_params() sets up default values for the stochasticity parameters. Defined as a separate function so it can be used for testing outside of a reactive environment.
Usage

```
stockParamsSetterUI(
   id,
   show_var = FALSE,
   show_biol_sigma = TRUE,
   show_est_sigma = TRUE,
   show_est_bias = TRUE,
   init_biol_sigma = 0,
   init_est_sigma = 0,
   init_est_bias = 0
)
```

```
stockParamsSetterServer(id)
```

```
set_stoch_params(input)
```

Arguments

- **id**
  - The id (shiny magic)

- **show_var**
  - Show the variability options when app opens (default is FALSE).

- **show_biol_sigma**
  - Show the biological productivity variability option (default is TRUE).

- **show_est_sigma**
  - Show the estimation variability option (default is TRUE).

- **show_est_bias**
  - Show the estimation bias option (default is TRUE).

- **init_biol_sigma**
  - Default value for biological productivity variability (ignored if not shown).

- **init_est_sigma**
  - Default value for estimation variability (ignored if not shown).

- **init_est_bias**
  - Default value for estimation bias (ignored if not shown).

- **input**
  - A list of stochasticity parameters.

Value

A taglist

A list of stochasticity options.

---

**Stock**

*R6 Class representing a stock*

**Description**

A stock object has life history parameters, fields and methods for a biomass dynamic model.
Details

A stock has biomass, effort, catch and hcr_ip and hcr_op fields as well as the life history parameters. The population dynamics are a simple biomass dynamic model. The Stock class is used for the Shiny apps in the AMPLE package.

Public fields

biomass Array of biomass
catch Array of catches
effort Array of fishing effort
hcr_ip Array of HCR input signals
hcr_op Array of HCR output signals
msy MSY (default = 100).
r Growth rate (default = 0.6). Set by the user in the app.
k Carrying capacity (default = NULL - set by msy and r when object is initialised).
p Shape of the production curve (default = 1).
q Catchability (default = 1).
lrp Limit reference point, expressed as depletion (default = 0.2).
trp Target reference point, expressed as depletion (default = 0.5).
b0 Virgin biomass (default = NULL - set by msy and r when object is initialised).
current_corrnoise Stores the current values of the correlated noise (by iteration).
biol_sigma Standard deviation of biological variability (default = 0).
last_historical_timestep The last historical timestep of catch and effort data.

Methods

Public methods:

- `Stock$new()`
- `Stock$reset()`
- `Stock$reactive()`
- `Stock$fill_history()`
- `Stock$fill_catch_history()`
- `Stock$fill_biomass()`
- `Stock$as_data_frame()`
- `Stock$project()`
- `Stock$relative_cpue()`
- `Stock$relative_effort()`
- `Stock$replicate_table()`
- `Stock$time_periods()`
- `Stock$performance_indicators()`
- `Stock$pi_table()`
• Stock$clone()

Method new(): Create a new stock object, with fields of the right dimension and NA values (by calling the reset() method. See the reset() method for more details.

Usage:
Stock$new(stock_params, mp_params, niters = 1)

Arguments:
stock_params A list of stock parameters with essential elements: r (growth rate, numeric), stock_history (string: "fully", "over", "under") initial_year (integer), last_historical_timestep (integer), nyears (integer), biol_sigma (numeric).
mp_params A list of the MP parameters. Used to fill HCR ip and op.
niters The number of iters in the stock (default = 1).

Returns: A new Stock object.

Method reset(): Resets an existing stock object, by remaking all fields (possibly with different dimensions for the array fields). Fills up the catch, effort and biomass fields in the historical period based on the stock history and life history parameters in the stock_params argument. This is a reactive method which invalidates a reactive instance of this class after it is called.

Usage:
Stock$reset(stock_params, mp_params, niters)

Arguments:
stock_params A list with essential elements: r (growth rate, numeric, default=6), stock_history (string: "fully", "over", "under", default="fully") initial_year (integer, default=2000), last_historical_timestep (integer, default=10), nyears (integer, default=30), biol_sigma (numeric, default = 0).
mp_params A list of the MP parameters. Used to fill HCR ip and op.
niters The number of iters in the stock (default = 1).

Returns: A new Stock object.

Method reactive(): Method to create a reactive instance of a Stock.

Usage:
Stock$reactive()

Returns: a reactiveExpr.

Method fill_history(): Fills the historical period of the stock

Usage:
Stock$fill_history(stock_params, mp_params)

Arguments:
stock_params Named list with last_historical_timestep and stock_history elements.
mp_params A list of the MP parameters. Used to fill HCR ip and op.

Method fill_catch_history(): Fill up the historical period of catches with random values to simulate a catch history

Usage:
Stock$fill_caListen_history(stock_params)

Arguments:
stock_params A list with essential elements: r (growth rate, numeric), stock_history (string: "fully", "over", "under") initial_year (integer), last_historical_timestep (integer), nyears (integer).
stock_history Character string of the exploitation history (default = "fully", alternatives are "under" or "over").

Method fill_biomass(): Fills the biomass in the next timestep based on current biomass and catches. The surplus production model has the general form: $B_{t+1} = B_t + f(B_t) - Ct$ Where the production function $f()$ is a Pella & Tomlinson model with shape $f(B_t) = \frac{r}{p} B_t * (1 - \left(\frac{B_t}{k}\right)^p)$ Here $p$ is fixed at 1 to give a Schaefer model $cpue = \frac{Ct}{Et} = qB_t$

Usage:
Stock$fill_biomass(ts, iters = 1:dim(self$biomass)[1])

Arguments:
 ts The biomass time step to be filled (required catch etc in ts - 1).
 iters The iterations to calculate the biomass for (optional - default is all of them).

Method as_data_frame(): Produces a data.frame of some of the array-based fields, like biomass. Just used for testing purposes.

Usage:
Stock$as_data_frame()

Method project(): Projects the stock over the time steps given and updates the biomass, HCR ip / op and catches. It uses a simple biomass dynamic model where the catches or fishing effort are set every time step by the harvest control rule.

Usage:
Stock$project(timesteps, mp_params, iters = 1:dim(self$biomass)[1])

Arguments:
 timesteps The timesteps to project over. A vector of length 2 (start and end).
 mp_params A vector of management procedure parameters.
 iters A vector of iterations to be projected. Default is all the iterations in the stock
 Returns: A stock object (a reactiveValues object with bits for the stock)

Method relative_cpue(): The catch per unit effort (CPUE, or catch rate) relative to the CPUE in the last historical period.

Usage:
Stock$relative_cpue()

Returns: An array of same dims as the catch and effort fields.

Method relative_effort(): The effort relative to the effort in the last historical period.

Usage:
Stock$relative_effort()

Returns: An array of same dims as the effort field.
Method replicate_table(): Summarises the final year of each iteration. Only used for the Measuring Performance app.

Usage:
Stock$replicate_table(iters = 1, quantiles = c(0.05, 0.95))

Arguments:
itersthe iterations to calculate the table values for (default is iteration 1).
quantiles Numeric vector of the quantile range. Default values are 0.05 and 0.95.

Method time_periods(): Calculates the short, medium and long term periods to calculate the performance indicators over, based on the last historic year of data and the number of years in the projection.

Usage:
Stock$time_periods()


Usage:
Stock$performance_indicators(
  iters = 1:dim(self$biomass)[1],
  quantiles = c(0.05, 0.95)
)

Arguments:
itersthe iterations to calculate the table values for (default is all of them).
quantiles Numeric vector of the quantile range. Default values are 0.05 and 0.95.

Returns: A data.frame

Method pi_table(): Makes a table of the performance indicators.

Usage:
Stock$pi_table(iters = 1:dim(self$biomass)[1], quantiles = c(0.05, 0.95))

Arguments:
itersthe iterations to calculate the table values for (default is all of them).
quantiles Numeric vector, length 2, of the low and high quantiles.

Method clone(): The objects of this class are cloneable with this method.

Usage:
Stock$clone(deep = FALSE)

Arguments:
deep Whether to make a deep clone.
**Arguments**

- **id**  
  Shiny magic
- **get_stoch_params**  
  Reactive expression that accesses the stochasticity module server.
- **input**  
  List of stock parameters taken from the shiny UI (stockParamsSetterUI()).
- **biol_sigma**  
  Standard deviation of the biological variability (default = 0).

**Value**

A taglist  
A list of stock options.

---

**threshold**  
*Evaluates a threshold harvest control rule*

**Description**

Evaluates a threshold (i.e. hockey stick) harvest control rule. Used by the hcr_op function.

**Usage**

threshold(input, mp_params, ...)

---

**Stock module**  
*stockParamsSetterUI*

**Description**

stockParamsSetterUI() is the interface for the stock options (e.g. life history and exploitation status). stockParamsSetterServer() does the setting of the stock parameters in the server.  
get_stock_params() Sets up default values for the stock, including year range. It’s a separate function so it can be used and tested outside of a reactive environment.

**Usage**

stockParamsSetterUI(id)

stockParamsSetterServer(id, get_stoch_params = NULL)

get_stock_params(input, biol_sigma = 0)
Arguments

input A vector of the 'true' stock status
mp_params The HCR / management procedure parameters used to evaluate the HCR (as a list).
... Unused

Value

A vector of the same dimension as the input.
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