

# Package: FLCore (via r-universe)

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**Title** Core Package of FLR, Fisheries Modelling in R

**Version** 2.6.20.9322

**Description** Core classes and methods for FLR, a framework for fisheries modelling and management strategy simulation in R. Developed by a team of fisheries scientists in various countries. More information can be found at <http://flr-project.org/>.

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**Suggests** testthat, rlang, hedgehog

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acc

*Catch curve estimates of total mortality at age (Z)***Description**

Catch curve estimates of total mortality at age (Z)

**Usage**

acc(object, ...)

**Examples**

```
data(ple4)
```

---

accessors

*accessor and replacement methods for FLCore classes*

---

**Description**

All S4 classes defined in FLCore have methods for accessing and replacing any of their slots. These methods are named as the slot, and will return the content of the slot, for the accessor method, or modify it with the provided value.

**Usage**

```
name(object, ...)
```

```
desc(object, ...)
```

```
range(x, i) <- value
```

```
catch(object, ...)
```

```
catch.n(object, ...) <- value
```

```
catch.wt(object, ...)
```

```
discards(object, ...)
```

```
discards.n(object, ...)
```

```
discards.wt(object, ...)
```

```
landings(object, ...)
```

```
landings.n(object, ...)
```

```
landings.wt(object, ...)
```

```
m(object, ...)
```

```
stock(object, ...)
```

```
stock.n(object, ...)
```

```
stock.wt(object, ...)
```

m.spwn(object, ...)  
harvest(object, catch, ...)  
harvest.spwn(object, ...)  
mat(object, ...)  
n(object, ...)  
m(object, ...)  
wt(object, ...)  
fec(object, ...)  
spwn(object, ...)  
effort(object, metier, ...)  
type(object, ...)  
distr(object, ...)  
distribution(object, ...)  
index(object, ...)  
index.var(object, ...)  
catch.n(object, ...)  
catch.wt(object, ...)  
sel.pattern(object, ...)  
index.q(object, ...)  
model(object, ...)  
logl(object, ...)  
gr(object, ...)  
initial(object, ...)  
logLik(object, ...)

```
vcov(object, ...) <- value
hessian(object, ...)
logerror(object, ...)
details(object, ...)
residuals(object, ...) <- value
fitted(object, ...)
rec(object, ...)
rec.obs(object, ...)
catch.q(object, ...)
discards.sel(object, ...)
landings.sel(object, ...)
params(object, ...)

## S4 replacement method for signature 'FLS,FLQuants'
catch(object) <- value
```

### Arguments

object	The object from which a slot is to be extracted or replaced
value	Object to be inserted into the relevant slot

### Details

Accessors and replacement methods, with some exception, are created at build time by calls to the `createFLAccessors` function. An accessor method is created for each slot, with simply calls `slot()` on the relevant slot name. For slots of class `FLQuant`, or `FLArray`-based, two methods are created: one if `value` is of class `FLQuant`, and another for `value` being a numeric vector. The later would insert the vector into the slot structure, using R's recycling rules.

Users are encouraged to use the accessor methods, rather than the `'@'` operator or the `slot()` method, to isolate code from the internal structure of the class. If a slot was to be altered or deleted in the future, a method would be provided to return the same value, computed from other slots.

Some of these methods might already not access directly an slot, and instead carry out a calculation to return the requested value, depending on the class being called with. Please refer to the particular method implementation to see if this is the case.

Accessor methods for slots of class `predictModel` behave differently depending on the `compute` argument. Please refer to the relevant help page for further clarification.

An object of class FLQuants, containing three elements named *catch*, *catch.n* and *catch.wt*, as returned by `computeCatch`, can be assigned directly to an object using `catch<-`.

### Value

The required slot, for an accessor method, or invisible modifies the object, for the replacement one.

### Author(s)

The FLR Team

### See Also

[FLQuant](#), [FLStock](#), [FLIndex](#), [FLBiol](#), [predictModel](#)

### Examples

```
data(ple4)

# To access the catch slot in an FLStock, use
catch(ple4)

# while to modify it, do
catch(ple4) <- catch(ple4) * 2

# A number can be used as input, to be recycled
m(ple4) <- 0.3
# same as a longer vector, by age
m(ple4) <- 0.4^(seq(1, 2, length=10))

# To see the methods defined by createFLAccessors, run, for example
getMethod('catch', 'FLS')

# Assign the 3 catch slots
catch(ple4) <- computeCatch(ple4, slot="all")
```

---

adjust,FLStock-method *Recalculate to adjust abundances to F and M*

---

### Description

An FLStock object is projected forward using the initial abundances and the total mortality-at-age per timestep. New values for the stock.n and catch.n slots are calculated, assuming that harvest and m are correct. This calculation provides a test of the internal consistency of the object.

### Usage

```
## S4 method for signature 'FLStock'
adjust(object)
```



**Arguments**

object            an FLStock object

**Value**

FLStock object

**See Also**

[harvest](#)

**Examples**

```
data(ple4)
test <- adjust(ple4)
# Difference in catch due to estimation error
plot(FLStocks(PLE=ple4, TEST=test))
```

---

ageopt	<i>Age at which a cohort reaches its maximum biomass, calculated by year</i>
--------	--

---

**Description**

The optimal (or critical) age is the transition point when a cohort achieves its maximum biomass in the absence of fishing, i.e. losses due to natural mortality are now greater than gains due to increase in individual biomass.

**Usage**

```
## S4 method for signature 'FLStock'
ageopt(object)
```

**Arguments**

object            An object of class 'FLStock'

**Value**

The age at which maximum biomass is reached, an 'FLQuant'.

**Author(s)**

The FLR Team

**See Also**

[FLStock](#)

**Examples**

```
data(ple4)
ageopt(ple4)
```

---

AIC

*Method AIC*

---

**Description**

Akaike's information criterion (AIC) method A method to calculate Akaike's 'An Information Criterion' (AIC) of an [FLModel](#) object from the value of the obtained log-likelihood stored in its logLik slot.

**Usage**

```
## S4 method for signature 'FLModel,numeric'
AIC(object, k = 2)
```

**Arguments**

object            an FLModel object  
k                 the penalty per parameter to be used; the default 'k = 2' is the classical AIC.

**Generic function**

```
AIC(object, k)
```

**Author(s)**

The FLR Team

**See Also**

[AIC](#), [logLik](#), [FLModel](#)

**Examples**

```
data(nsher)
AIC(nsher)
```

---

`append-FLCore`*Append objects along the year dimension*

---

### Description

Method to append objects along the *year* dimensions, by extending, combining and substituting sections of them.

### Usage

```
## S4 method for signature 'FLQuant,FLQuant'  
append(x, values, after = dims(values)$minyear - 1)
```

```
## S4 method for signature 'FLStock,FLStock'  
append(x, values, after = dims(values)$minyear - 1)
```

### Arguments

<code>x</code>	the object to which the values are to be appended to.
<code>values</code>	to be included in the modified object.
<code>after</code>	a year dimname after with the values are to be appended.

### Details

FLR objects are commonly manipulated along the year dimension, and the append method offers a simple interface for substituting parts of an object with another, or combine them into one, extending them when necessary. The object to be included or added to the first will be placed as defined by the *year* dimnames, unless the *after* input argument specifies otherwise.

Attributes like dimnames and *units* will always be taken from the first argument, unless the necessary changes to dimnames\$year

### Value

An object of the same class as `x` with values appended.

### Author(s)

The FLR Team

### See Also

[base::append](#)

**Examples**

```

# append(FLQuant, FLQuant)
fq1 <- FLQuant(1, dimnames=list(age=1:3, year=2000:2010))
fq2 <- FLQuant(2, dimnames=list(age=1:3, year=2011:2012))
fq3 <- FLQuant(2, dimnames=list(age=1:3, year=2014:2016))

# Appends by dimnames$year
append(fq1, fq2)
# Appends by dimnames$year with gap (2011:2013)
append(fq1, fq3)
# Appends inside x
append(fq1, fq2, after=2009)
# Appends after end of x
append(fq1, fq2, after=2013)

# append(FLStock, FLStock)
data(ple4)
fs1 <- window(ple4, end=2001)
fs2 <- window(ple4, start=2002)
fs3 <- window(ple4, start=2005)

# Appends by dimnames$year
stock.n(append(fs1, fs2))

# Appends by dimnames$year with gap (2011:2013)
stock.n(append(fs1, fs3))

# Appends inside x
stock.n(append(fs1, fs3, after=2000))
# Appends after end of x
stock.n(append(fs1, fs3, after=2005))

```

---

apply,FLArray,numeric,function-method  
*apply method for FLCore classes*

---

**Description**

Applies a function over the margins of an array-based FLCore class

**Usage**

```

## S4 method for signature 'FLArray,numeric,function'
apply(X, MARGIN, FUN, ..., simplify = TRUE)

## S4 method for signature 'FLPar,ANY,ANY'
apply(X, MARGIN, FUN, ..., simplify = TRUE)

## S4 method for signature 'FLQuantJK,numeric,function'

```

```

apply(X, MARGIN, FUN, ..., simplify = TRUE)

## S4 method for signature 'FLParJK,numeric,function'
apply(X, MARGIN, FUN, ..., simplify = TRUE)

```

### Details

These methods call R's `base::apply` on an `FLArray` the standard arithmetic operators included in the `Arith` group ("`+`", "`-`", "`*`", "`^`", "`%%`", "`%/`", and "`/`"), so that they return an object of the appropriate class.

When the operation involves objects of two classes (e.g. `FLPar` and `FLQuant`), the class is the returned object is that of the more complex object, in this case `FLQuant`.

### Author(s)

The FLR Team

### See Also

[base::apply](#)

### Examples

```

flq <- FLQuant(rlnorm(90), dim=c(3,10), units='kg')
flp <- FLPar(a=99)

# FLQuant and numeric
flq * 25
# Two FLQuant objects
flq + flq

```

---

ar1rlnorm

*Generates a time series of possible bias-corrected lognormal autocorrelated random values*

---

### Description

Thorston, 2020.

### Usage

```

ar1rlnorm(
  rho,
  years,
  iters = 1,
  meanlog = 0,
  sdlog = 1,

```

```

    bias.correct = TRUE,
    ...
  )

```

### Arguments

rho	Autocorrelation coefficient.
years	Vector of year names.
iters	Number of iterations.
meanlog	Mean of the series in log space.
sdlog	Marginal standard deviation in log space.
bias.correct	Should bias-correction be applied? Defaults to TRUE.

### Value

An FLQuant object

### Author(s)

Iago Mosqueira (WMR), Henning Winker (JRC).

### References

Thorson, J. T. Predicting recruitment density dependence and intrinsic growth rate for all fishes worldwide using a data-integrated life-history model. *Fish Fish.* 2020; 21: 237– 251. <https://doi-org.ezproxy.library.wur.nl/10.1111/faf.12427>

### See Also

[rlnorm](#)

### Examples

```

devs <- ar1rlnorm(rho=0.6, years=2000:2030, iter=500, meanlog=0, sdlog=1)
plot(devs)

```

---

Arith,numeric,FLArray-method

*Arithmetic operators for FLCORE classes*

---

### Description

Overloaded arithmetic operators for FLCORE classes

**Usage**

```
## S4 method for signature 'numeric,FLArray'
Arith(e1, e2)

## S4 method for signature 'FLArray,numeric'
Arith(e1, e2)

## S4 method for signature 'FLArray,FLArray'
Arith(e1, e2)

## S4 method for signature 'FLPar,FLPar'
Arith(e1, e2)

## S4 method for signature 'FLArray,FLPar'
Arith(e1, e2)

## S4 method for signature 'FLPar,FLArray'
Arith(e1, e2)
```

**Details**

These methods apply the standard arithmetic operators included in the [Arith](#) group ("+", "-", "\*", "^", "%%", "%/%", and "/"), so that they return an object of the appropriate class.

When the operation involves objects of two classes (e.g. [FLPar](#) and [FLQuant](#)), the class is the returned object is that of the more complex object, in this case [FLQuant](#).

**Author(s)**

The FLR Team

**See Also**

[methods::Arith](#) [base::Arithmetic](#)

**Examples**

```
flq <- FLQuant(rlnorm(90), dim=c(3,10), units='kg')
flp <- FLPar(a=99)

# FLQuant and numeric
flq * 25
# Two FLQuant objects
flq + flq

# FLQuant and FLPar
flq / flp
```

---

`as.FLSRs`                      *Convert an FLStock into a list of one or FLSR objects.*

---

### Description

A single FLStock can be coerced into a list with one or more objects of class FLSR, each of them typically set to a different stock-recruit model.

### Usage

```
as.FLSRs(x, models = NULL, ...)
```

### Arguments

`x`                      An estimated FLStock object to coerce.  
`models`                Name(s) of model(s) to fit.  
`...`                    Any extra arguments to be passed to *as.FLSR*.

### Value

An object of class FLSRs

### Author(s)

FLR Team, 2023.

### See Also

[FLSRs](#) [FLSRs](#) [as.FLSR\(\)](#)

### Examples

```
data(ple4)
as.FLSRs(ple4, model=c("bevholt", "segreg"))
```

---

`bias`                      *Bias of estimates through jackknife*

---

### Description

Description: Lorem ipsum dolor sit amet, consectetur adipiscing elit. Pellentesque eleifend odio ac rutrum luctus. Aenean placerat porttitor commodo. Pellentesque eget porta libero. Pellentesque molestie mi sed orci feugiat, non mollis enim tristique.



**Usage**

```
## S4 method for signature 'FLQuantJK'  
bias(x)  
  
## S4 method for signature 'FLParJK'  
bias(x)
```

**Arguments**

x                    An object holding estimates obtained through jackknife

**Details**

Details: Aliquam sagittis feugiat felis eget consequat. Praesent eleifend dolor massa, vitae faucibus justo lacinia a. Cras sed erat et magna pharetra bibendum quis in mi. Sed sodales mollis arcu, sit amet venenatis lorem fringilla vel. Vivamus vitae ipsum sem. Donec malesuada purus at libero bibendum accumsan. Donec ipsum sapien, feugiat blandit arcu in, dapibus dictum felis.

$$\widehat{Bias}_{(\theta)} = (n - 1) \left( \frac{1}{n} \sum_{i=1}^n \hat{\theta}_{(i)} - \hat{\theta} \right)$$

**Value**

A value for the mean bias

**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

**Examples**

```
flq <- FLQuant(1:8)  
flj <- jackknife(flq)  
bias(flj)
```

---

**BIC***Method BIC Bayesian information criterion (BIC) method*

---

**Description**

A method to calculate the Bayesian information criterion (BIC), also known as Schwarz's Bayesian criterion of an [FLModel](#) object from the value of the obtained log-likelihood stored in its `logLik` slot.

**Usage**

```
## S4 method for signature 'FLModel'  
BIC(object)
```

**Arguments**

`object` a fitted `FLModel` object for which there exists a `'logLik'` method to extract the corresponding log-likelihood.

**Generic function**

```
BIC(object)
```

**Author(s)**

The FLR Team

**See Also**

[AIC](#), [BIC](#), [FLModel](#), [logLik](#)

**Examples**

```
data(nsher)  
BIC(nsher)
```

---

**bubbles***Method Bubbles plot*

---

### Description

This method plots three dimensional data such as matrices by age and year or age-class, very common in fisheries. The area of each bubble is proportional to the corresponding value in the matrix. Note that `bubbles` accepts an argument `bub.scale` to control the relative size of the bubbles. Positive and negative values have separate colours.

### Usage

```
## S4 method for signature 'formula,FLQuant'  
bubbles(x, data, bub.scale = 2.5, col = c("blue", "red"), ...)  
  
## S4 method for signature 'formula,data.frame'  
bubbles(x, data, bub.scale = 2.5, col = c("blue", "red"), ...)  
  
## S4 method for signature 'formula,FLCohort'  
bubbles(x, data, bub.scale = 2.5, ...)  
  
## S4 method for signature 'formula,FLQuants'  
bubbles(x, data, bub.scale = 2.5, bub.col = gray(c(0.1, 0.1)), ...)
```

### Generic function

```
bubbles(x, data)
```

### Author(s)

The FLR Team

### See Also

[lattice](#), [FLQuant](#), [FLQuants](#), [FLCohort](#)

### Examples

```
data(ple4)  
bubbles(age~year, data=catch.n(ple4))  
bubbles(age~year, data=catch.n(ple4), bub.scale=5)  
bubbles(age~cohort, data=FLCohort(catch.n(ple4)), bub.scale=5)  
  
qt01 <- log(catch.n(ple4)+1)  
qt02 <- qt01+rnorm(length(qt01))  
flqs <- FLQuants(qt01=qt01, qt02=qt02)  
bubbles(age~year|qname, data=flqs, bub.scale=1)
```

```
qt03 <- FLQuant(rnorm(100),dimnames=list(age=as.character(1:10),
  year=as.character(1:10)))
bubbles(age~year, data=qt03, bub.scale=7, col=c("black","red"), pch=16)
```

---

catch.n,FLQuant-method

*catch.n calculation method*

---

## Description

Calculate catch.n (catch-at-age/length) from abundances, F and M using the catch equation

## Usage

```
## S4 method for signature 'FLQuant'
catch.n(object, harvest, m)
```

## Details

The catch-at-age/length, commonly found in the catch.n slot of an FLStock object, can be simply calculated from abundances-at-age/length, and natural and fishing mortalities-at-age/length by applying the catch equation

$$C = N \cdot F \frac{F}{M + F} \cdot (1 - e^{-(M + F)})$$

## Author(s)

The FLR Team

## See Also

[FLStock](#)

## Examples

```
data(ple4)
res <- catch.n(stock.n(ple4), harvest(ple4), m(ple4))
catch.n(ple4) / res
```

---

catchInmature	<i>Proportion of mature and immature fish in the catch</i>
---------------	--

---

**Description**

The proportion in weight of mature and immature fish in the catch can be computed using `catchMature` and `catchInmature`.

**Usage**

```
catchInmature(object)
```

```
catchMature(object)
```

**Arguments**

`object` An age-structured `FLStock` object

**Value**

An `FLQuant` object

**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

**Examples**

```
data(ple4)
catchInmature(ple4)
catchMature(ple4)
```

---

coerce-methods	<i>Convert Objects Between Classes</i>
----------------	--

---

**Description**

Objects of various **FLCore** classes can be converted into other classes, both basic R ones, like `data.frame`, and others defined in the package. For the specifics of the precise calculations carried out for each pair of classes, see below.

**Arguments**

object	Object to be converted.
Class	Name of the class to convert the object to, character.

**Value**

An object of the requested class.

**FLArray to data.frame**

The six dimensions of an `FLArray` are converted into seven columns, named `quant` (or any other name given to the first dimension in the object), `year`, `unit`, `season`, `area`, `iter` and `data`. The last one contains the actual numbers stored in the array. `units` are stored as an attribute to the `data.frame`. The `year` and `data` columns are of type `numeric`, while all others are `factor`.

**FLPar to data.frame**

The two or more dimensions of an `FLPar` objects are converted into three or more columns. For a 2D objects, they are named `params`, `iter` and `data`. The last one contains the actual numbers stored in the array, in a column type `numeric`, while all others are `factor`.

**Author(s)**

The FLR Team

**See Also**

[base::as](#), [base::coerce](#)

**Examples**

```
# from FLQuant to data.frame
as(FLQuant(rnorm(100), dim=c(5, 20)), "data.frame")
# from FLPar to data.frame
as(FLPar(phi=rnorm(10), rho=rlnorm(10)), "data.frame")
```

---

compare

*A method for comparing FLR objects*

---

**Description**

Comparisons of complete objects of FLR classes can be carried out and a report table is generated to better identify differences. Comparisons do not substitute but complement those provided by R's `all.equal` and `identical`.

**Usage**

```
compare(result, target, ...)
```

**Arguments**

result            First element in comparison, result of method or operation.  
 target            Second element, desired output.

**Value**

A table of comparisons, of class data.frame.

**Author(s)**

Iago Mosqueira (WMR)

---

compute                            *Methods to compute quantities*

---

**Description**

Methods to compute total quant-aggregated catch, landings, discards and stock biomass from age or length-structured numbers and mean weights.

Methods to compute total quant-aggregated catch, landings, discards and stock biomass from age or length-structured numbers and mean weights.

**Usage**

```
computeLandings(object, ...)
```

```
computeDiscards(object, ...)
```

```
computeCatch(object, ...)
```

```
computeStock(object, ...)
```

```
computeHarvest(object, catch, ...)
```

```
computeLandings(object, ...)
```

```
computeDiscards(object, ...)
```

```
computeCatch(object, ...)
```

```
computeStock(object, ...)
```

```
## S4 method for signature 'FLS'  
computeLandings(object, na.rm = TRUE)
```

```
## S4 method for signature 'FLS'
```

```

computeDiscards(object, na.rm = TRUE)

## S4 method for signature 'FLS'
computeCatch(object, slot = "catch", na.rm = TRUE)

## S4 method for signature 'FLS'
computeStock(object, na.rm = TRUE)

```

## Details

These methods compute the total catch, landings, discards and stock biomass from the quant-structured values in numbers and weight per individual. The calculation for landings, discards and stock involves the product of the landings/discards/stock in numbers (`landings.n`, `discards.n` or `stock.n`) by the individual weight-at-quant (`landings.wt`, `discards.wt` or `stock.wt`), as in

$$L = L_n * L_{wt}$$

By selecting `slot="catch"`, `computeCatch` can calculate in the same way the total catch from the catch-at-quant and weight in the catch. Those two values (in slots `catch.n` and `catch.wt`) can also be calculated (from landings and discards) by specifying `slot="n"` and `slot="wt"` respectively. Calling `computeCatch` with option `slot="all"` will carry out the three calculations. In this case, the returned object will be of class `FLQuants`, with element names `catch`, `catch.n` and `catch.wt`, which can then be passed directly to the `catch<-` replacement method.

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## Generic function

```

computeCatch(object, ...)
computeLandings(object, ...)
computeDiscards(object, ...)
computeStock(object, ...)
computeCatch(object, ...)
computeLandings(object, ...)
computeDiscards(object, ...)
computeStock(object, ...)

```



**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

[FLComp](#)

**Examples**

```
data(ple4)
summary(computeLandings(ple4))
summary(computeCatch(ple4, slot="all"))
stock(ple4) <- computeStock(ple4)
landings(ple4) <- computeLandings(ple4)
catch.n(ple4) <- computeCatch(ple4, slot="n")
catch(ple4) <- computeCatch(ple4, slot="all")
```

```
data(ple4)
summary(computeLandings(ple4))
summary(computeCatch(ple4, slot="all"))
stock(ple4) <- computeStock(ple4)
landings(ple4) <- computeLandings(ple4)
catch.n(ple4) <- computeCatch(ple4, slot="n")
catch(ple4) <- computeCatch(ple4, slot="all")
```

---

computeHarvest,FLStock,missing-method

*Computes fishing mortality from abundances, catches and natural mortality*

---

**Description**

Objects or class 'FLStock' already contain a 'harvest' slot to store estimates of fishing mortality at age, for example those obtained from a stock assessment method. Fishing mortality at age can be recalculated using two methods:

**Usage**

```
## S4 method for signature 'FLStock,missing'
computeHarvest(object, units = NULL)
```

**Arguments**

units	Harvest to be computed as 'f' or 'hr', 'character'.
x	An object of class 'FLStock'.

**Value**

An 'FLQuant' with the calculated fishing mortalities at age.

**Author(s)**

The FLR Team

**See Also**

[FLStock harvest\(\)](#) [FLQuant](#)

**Examples**

```
data(ple4)
# Compute 'f' from stock.n and Baranov
computeHarvest(ple4)
# Recomputes all F at age by solving catch Baranov
recomputeHarvest(ple4)
```

---

cpue	<i>cpue, a method to generate an observation of a CPUE index of abundance</i>
------	---

---

**Description**

The observation of stock abundance by CPUE series from commercial fleets is an important step in the generation of management advice that needs to be replicated on an Operating Model during any simulation exercise. This method generates an observation of biomass or numbers-at-age from an FLstock being used as OM.

**Usage**

```
cpue(object, index, ...)

## S4 method for signature 'FLStock,missing'
cpue(
  object,
  sel.pattern = harvest(object),
  effort = units(harvest(object)),
  biomass = TRUE
)
```

**Arguments**

object	The object from which to generate the observation.
effort	Units of index to use to mimic effort series in the fishery, "f" or "hr"
sel	The selectivity of the survey, defaults to be 1 for all ages.
mass	Is the index to be in weight at age?

**Value**

An FLQuant for the index of abundance, age-disaggregated

**Author(s)**

Laurie Kell & Iago Mosqueira, FLR Team.

**See Also**

[FLComp](#)

**Examples**

```
data(ple4)

cpue(ple4)
# Am aggregated biomass CPUE
quantSums(cpue(ple4))

## Not run:
plot(FLQuants(om=stock(ple4), cpue=quantSums(cpue(ple4)),
  hr=quantSums(cpue(ple4, effort="hr"))))

## End(Not run)
```

---

createFLAccesors      *Create accesor methods for a given class*

---

**Description**

This function creates a complete set of standard S4 class accessors and replacers. Not intended for direct use.

**Usage**

```
createFLAccesors(class, exclude = character(1), include = missing)
```

**Arguments**

class	name of the class
exclude	Slot names to exclude
include	Slot names to include

**Author(s)**

The FLR Team

---

datasets

*FLCore datasets*

---

## Description

Example datasets for the classes defined in FLCore.

## Details

- ple4, [FLStockA](#) dataset for North Sea (ICES Area IV) plaice. Catch, landings, discards, natural mortality, weight-at-age and maturity, together with the VPA estimated abundances and fishing mortalities.
- ple4sex, [FLStockA](#) dataset of North Sea (ICES Area IV) plaice disaggregated by sex. Catch, yield, landings, discards, natural mortality, weight-at-age and maturity, together with the VPA estimated abundances and fishing mortalities.
- ple4.index, [FLIndexA](#) dataset of North Sea (ICES Area IV) plaice survey catch per unit effort, index and index variance.
- ple4.indices, [FLIndicesA](#) dataset of three North Sea (ICES Area IV) plaice survey catch per unit effort series. Index and index variance.
- ple4.biol, [FLBiolA](#) dataset of the North Sea plaice population. Numbers, natural mortality, mass and fecundity-at-age.
- nsher , [FLSRStock](#) and recruit data and fitted relationship for autumn spawning North Sea herring.

Datasets can be loaded by issuing the data command, like in `data(ple4)`.

## References

ICES.

## See Also

[FLStock](#), [FLSR](#), [FLIndex](#), [FLStock](#), [FLIndex](#), [FLBiol](#)

## Examples

```
data(ple4)
summary(ple4)
```

```
data(nsher)
is(nsher)
```

---

 dbind

*Methods for binding objects of array classes along a given dimension*


---

### Description

These methods can bind two or more objects of array-based classes (e.g. FLQuant), along the specified dimension.

### Usage

```
dbind(x, y, ...)

## S4 method for signature 'FLArray,FLArray'
dbind(x, y, ..., dim = 1)

qbind(...)
ybind(...)
ubind(...)
sbind(...)
abind(...)
ibind(...)
```

### Arguments

x	First object to bind
y	Second object to bind
...	Other objects to bind
dim	Dimension to bind on, <i>numeric</i> or <i>character</i> .

### Details

The objects to bind must contain the same dimnames in all dimensions other than that used to bind, while dimnames in the selected one must differ. See the examples below for correct and incorrect uses.

Object are bound in the order they are provided, with no attempt to sort according to the dimnames of the chosen dimension.

The implementation is based around a single method (*dbind*), that operates along the dimension position or name indicated by the *dim* argument. A series of shortcut functions call the method for specific dimensions, with names related to the dimensions name they operate on (e.g. *ybind* for *year*).

**Value**

An object of the same class as the inputs

**Author(s)**

Iago Mosqueira (EC JRC)

**See Also**

[FLQuant](#) [FLArray](#)

**Examples**

```
# By iter
x <- FLQuant(rnorm(80000), dim=c(4,20,1,1,1,1000))
y <- FLQuant(rnorm(80000), dim=c(4,20,1,1,1,1000))
dimnames(y) <- list(iter=1001:2000)
ibind(x,y)

# By quant (age)
x <- FLQuant(1, dimnames=list(age=1:3, year=1:10))
y <- FLQuant(2, dimnames=list(age=4:12, year=1:10))
qbind(x, y)

# By year
x <- FLQuant(1, dimnames=list(age=1:3, year=1:10))
y <- FLQuant(2, dimnames=list(age=1:3, year=11:20))
z <- FLQuant(3, dimnames=list(age=1:3, year=21:30))
ybind(x, y, z)

# By season
x <- FLQuant(1, dimnames=list(year=1:10, season=1:2))
y <- FLQuant(2, dimnames=list(year=1:10, season=3:4))
sbind(x, y)
```

---

dims

*Method dims*

---

**Description**

List with information on object dimensions

List with information on object dimensions

**Usage**

```
dims(obj, ...)
```

```
dims(obj, ...)
```

```
## S4 method for signature 'FLQuant'
dims(obj, element, ...)
```

### Details

Method `dims` returns a named list with information on the dimensions and dimension names of a given object. The list returned could be extended in the future and currently contains, depending on the class of the object, some of the following:

- quant** Length of the first dimension
- min** First quant
- max** Last quant
- year** Number of years
- minyear** First year in series
- maxyear** Last year in series
- cohort** Number of cohorts
- mincohort** First cohort in series
- maxcohort** Last cohort in series
- unit** Length of the third (unit) dimension
- season** Length of the fourth (season) dimension
- area** Length of the fifth (area) dimension
- iter** Length of the sixth (iter) dimension

Values in the returned list are of class `numeric`, unless `dimnames` are strings with no numeric translation, in which case the result is `NA`.

Please note that the name of the first element in the returned list changes with the name of the first dimension in the input object. Use `quant` to obtain the name and extract the relevant element from the result list.

Method `dims` returns a named list with information on the dimensions and dimension names of a given object. The list returned could be extended in the future and currently contains, depending on the class of the object, some of the following:

- quant** Length of the first dimension
- min** First quant
- max** Last quant
- year** Number of years
- minyear** First year in series
- maxyear** Last year in series
- cohort** Number of cohorts
- mincohort** First cohort in series
- maxcohort** Last cohort in series
- unit** Length of the third (unit) dimension

**season** Length of the fourth (season) dimension

**area** Length of the fifth (area) dimension

**iter** Length of the sixth (iter) dimension

Values in the returned list are of class `numeric`, unless `dimnames` are strings with no numeric translation, in which case the result is `NA`.

Please note that the name of the first element in the returned list changes with the name of the first dimension in the input object. Use `quant` to obtain the name and extract the relevant element from the result list.

### Generic function

```
dims(obj)
```

```
dims(obj)
```

### Author(s)

The FLR Team

### See Also

[dimnames, FLQuant](#)

[dimnames, FLQuant](#)

### Examples

```
flq <- FLQuant(rnorm(96), dim=c(3,8,1,4), quant='age')
dims(flq)
```

```
# Number of seasons
dims(flq)$season
```

```
# Length of first dimension
dims(flq)[[quant(flq)]]
```

---

dimSummaries

*Summaries by dimension*

---

### Description

Methods to compute various summary calculations (sum, mean, variance) over selected dimensions of objects from any array-based classes (e.g. `FLQuant`). These methods return an object of the same dimensions as the input but with length one in the dimension chosen to operate along.



**Usage**`quantSums(x, ...)``yearSums(x, ...)``unitSums(x, ...)``seasonSums(x, ...)``areaSums(x, ...)``iterSums(x, ...)``dimSums(x, ...)``quantMeans(x, ...)``yearMedians(x, ...)``yearMeans(x, ...)``unitMeans(x, ...)``seasonMeans(x, ...)``areaMeans(x, ...)``iterMeans(x, ...)``dimMeans(x, ...)``quantVars(x, ...)``yearVars(x, ...)``unitVars(x, ...)``seasonVars(x, ...)``areaVars(x, ...)``iterVars(x, ...)``dimVars(x, ...)``iterMedians(x, ...)``iterCVs(x, ...)`

```
iterProb(x, ...)  
  
## S4 method for signature 'FLQuant'  
quantSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
yearSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
unitSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
seasonSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
areaSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
iterSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
quantMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
yearMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
unitMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
seasonMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
areaMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
iterMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
yearMedians(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
iterMedians(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
quantVars(x, na.rm = TRUE)
```

```
## S4 method for signature 'FLQuant'  
yearVars(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
unitVars(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
seasonVars(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
areaVars(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
iterVars(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
iterCVs(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuant'  
iterProb(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
yearSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
unitSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
seasonSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
areaSums(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
yearMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
unitMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
seasonMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
areaMeans(x, na.rm = TRUE)  
  
## S4 method for signature 'FLQuantDistr'  
iterMeans(x, na.rm = TRUE)
```

```

## S4 method for signature 'FLQuantDistr'
iterMedians(x, na.rm = TRUE)

## S4 method for signature 'FLQuantDistr'
quantVars(x, na.rm = TRUE)

## S4 method for signature 'FLQuantDistr'
yearVars(x, na.rm = TRUE)

## S4 method for signature 'FLQuantDistr'
unitVars(x, na.rm = TRUE)

## S4 method for signature 'FLQuantDistr'
seasonVars(x, na.rm = TRUE)

## S4 method for signature 'FLQuantDistr'
areaVars(x, na.rm = TRUE)

## S4 method for signature 'FLQuantDistr'
iterVars(x, na.rm = TRUE)

## S4 method for signature 'FLPar'
iterMeans(x, na.rm = TRUE)

## S4 method for signature 'FLPar'
iterMedians(x, na.rm = TRUE)

## S4 method for signature 'FLPar'
iterVars(x, na.rm = TRUE)

## S4 method for signature 'FLPar'
iterSums(x, na.rm = TRUE)

```

### Arguments

<code>x</code>	An object.
<code>na.rm</code>	Should NAs be removed before calculation? Defaults to TRUE.

### Details

This set of methods computes three different summaries (sum, mean and variance) of an `FLQuant` object along each of the six dimensions (quant, year, unit, season, area, or iter). Medians and CVs can also be computed along the sixth dimension, `iter`.

These methods encapsulate a call to `apply` with the corresponding dimensions and function: `mean`, `median`, `var`, and `sum`, while `iterCVs` are computed as `sqrt(iterVars) / iterMeans`.

In contrast with R standard behaviour, the sum of a dimension where all elements are NA will be NA and not 0. See example below.

Methods working along the `iter` dimension are also defined for objects of class `FLPar`.

Methods to operate over the first dimension refer to it as the quant dimension, regardless of the actual name used in the object.

### Generic methods

quantSums(x), quantMeans(x), quantVars(x) yearSums(x), yearMeans(x), yearVars(x) unitSums(x), unitMeans(x), unitVars(x) seasonSums(x), seasonMeans(x), seasonVars(x) areaSums(x), areaMeans(x), areaVars(x) iterMeans(x), iterVars(x), iterMedians(x), iterSums(x) dimSums(x), dimMeans(x), dimVars(x)

### Author(s)

The FLR Team

### See Also

[FLQuant](#), [sum](#), [mean](#), [var](#)

### Examples

```
flq <- FLQuant(rnorm(4000), dim=c(5,10,2,2,2,10), quant='age')

quantSums(flq)
quantMeans(flq)
yearSums(flq)
iterMeans(flq)
dim(quantSums(flq))

# NA dims stay as NA when summed along
x <- FLQuant(c(NA, NA, NA, rnorm(6)), dim=c(3, 3))
quantSums(x)
# although in fact a sum of no elements (as na.rm=TRUE) is zero
apply(x, 2:6, sum, na.rm=TRUE)
```

---

discardsRatio

*Compute the ratio of discards to total catch in numbers or weight*

---

### Description

A calculation is made of the proportion of discards over total catch at age, either as numbers (value = 'numbers') or weight (value = 'weight'), or for the total discards and catch in biomass (value = 'total').

### Usage

```
discardsRatio(object, value = c("numbers", "weight", "total"))
```

**Arguments**

object            An object of class 'FLStock'  
value            One of 'numbers' (default), 'weight' or 'total'.

**Value**

The discards ratio (between 0 and 1), 'FLQuant'

**Author(s)**

The FLR Team

**See Also**

[FLStock](#)

**Examples**

```
data(ple4)
# Discards ratio at age in numbers
discardsRatio(ple4)
# Total proportion of discards by year
discardsRatio(ple4, value="total")
```

---

drop,FLArray-method    *drop method for FLCORE array-based classes*

---

**Description**

Delete the dimensions of an array which have only one level.

**Usage**

```
## S4 method for signature 'FLArray'
drop(x)
```

**Details**

This method calls R's [base::drop](#) on the @.Data slot of an [FLArray](#). Dimensions of length one are thus dropped, as is the class attribute and the units slot, and an array of equal or less dimensions, a matrix or a vector is returned.

On an FLQuant object with

**Author(s)**

The FLR Team

**See Also**[base::drop](#)**Examples**

```
x <- FLQuant(1:3, dim=c(3,3))
drop(x)
is(drop(x))
dim(drop(x))

# Result of drop can be used for matrix algebra
# for example to calculate aging error

data(ple4)
aging.error <- diag(0.8, 10)
diag(aging.error[-1,]) <- c(rep(0.1, 8), 0.2)
diag(aging.error[, -1]) <- c(0.2, rep(0.1, 8))
t(aging.error) %**% drop(catch.n(ple4))
```

---

**evalPredictModel***Evaluates a predictModel slot inside the object containing it*

---

**Description**

Models in objects of the [predictModel](#) class can make use of slots and methods of the FLR class in which it is contained as a slot. This function can be used by methods wishing to evaluate a single predictModel slot in the context of the class it is part of.

**Usage**

```
evalPredictModel(object, slot, ...)
```

**Arguments**

object	The FLR S4 over which the predictModel evaluation should take place
slot	The predictModel object to be evaluated

**Value**

The result of evaluating the model, usually an FLQuant

**Author(s)**

The FLR Team

**See Also**[predictModel](#)

---

exp, FLQuant-method      *exp and log methods FLCore array-based classes*

---

### Description

Compute the exponential and logarithmic functions

### Usage

```
## S4 method for signature 'FLQuant'
exp(x)

## S4 method for signature 'FLQuant'
log(x, ...)
```

### Details

This method simply calls R's [base::exp](#) and [base::drop](#), but take care of returning the right units of measurement, that is "" or `character(1)`.

### Author(s)

The FLR Team

### See Also

[base::exp](#) [base::log](#)

### Examples

```
x <- FLQuant(c(4,2,7,4,2,9), units="1000")
log(x)
units(log(x))
```

---

Extract

*Extract*

---

### Description

Extract or replace parts of an FLR Object



**Usage**

```
## S4 method for signature 'FLArray,ANY,ANY,ANY'
x[i, j, k, l, m, n, ..., drop = FALSE]

## S4 method for signature 'FLArray,array,missing,missing'
x[i]

## S4 replacement method for signature 'FLArray,ANY,ANY,ANY'
x[i, j, k, l, m, n, ...] <- value

## S4 replacement method for signature 'FLArray,ANY,ANY,FLArray'
x[i, j, k, l, m, n, ...] <- value

## S4 method for signature 'FLQuant'
x$name

## S4 method for signature 'FLQuantDistr,ANY,ANY,ANY'
x[i, j, k, l, m, n, drop = FALSE]

## S4 method for signature 'FLQuantDistr,array,missing,missing'
x[i]

## S4 method for signature 'FLPar,ANY,ANY,ANY'
x[i, j, k, l, m, n, ..., drop = FALSE]

## S4 method for signature 'FLPar,array,missing,missing'
x[i]

## S4 replacement method for signature 'FLPar,ANY,ANY,ANY'
x[i, j, k, l, m, n, ...] <- value

## S4 method for signature 'FLPar'
x$name

## S4 replacement method for signature 'FLPar'
x$name <- value

## S4 method for signature 'FLComp,ANY,ANY,ANY'
x[i, j, k, l, m, n, ..., drop = FALSE]

## S4 replacement method for signature 'FLComp,ANY,ANY,ANY'
x[i, j, k, l, m, n, ...] <- value

## S4 method for signature 'FLStock,ANY,ANY,ANY'
x[i, j, k, l, m, n, ..., drop = FALSE]

## S4 replacement method for signature 'FLStock,ANY,ANY,FLStock'
x[i, j, k, l, m, n, ...] <- value
```

```

## S4 method for signature 'FLI,ANY,ANY,ANY'
x[i, j, k, l, m, n, ..., drop = FALSE]

## S4 method for signature 'predictModel,ANY,missing,ANY'
x[i, k, l, m, n, ..., drop = FALSE]

## S4 replacement method for signature 'FLlst,ANY,missing'
x[[i, j]] <- value

## S4 replacement method for signature 'FLlst'
x$name <- value

## S4 replacement method for signature 'FLlst,ANY,missing,ANY'
x[i, j] <- value

## S4 method for signature 'FLlst,ANY,missing,ANY'
x[i, drop]

```

### Arguments

x	object from which to extract or replace element(s)
i, j, k, l, m, n	indices specifying elements to extract or replace on any of the six dimensions.
...	indices specifying elements to extract or replace by dimension name.
drop	If 'TRUE' the result is coerced to the lowest possible dimension, and so might change class (e.g. drop='TRUE' on an FLQuant might return an array of less dimensions, a matrix or a vector).
value	An object of the same class, or simpler if drop=TRUE, than 'x'.
name	See <a href="#">Extract</a> for further details.

### Details

Operators acting on FLQuant, FLCohort, FLPar, FLComp, and derived classes to extract or replace sections of an object.

Please note the differences between referencing sections of an object by position using values of class numeric, or by using dimnames of class character. See examples below.

All classes that are derived from FLComp (for example, FLStock and FLBio1) can be subset along the six dimensions of their FLQuant slots.

Classes that are derived from FLlst (for example, FLStocks and FLBio1s) can be subset in a similar way to ordinary list objects.

'\$' for the FLPar and FLQuant classes operate only along the first dimension ('params' or 'quant'), and are provided to be used specially in formulas.

### Generic function

[x,i,j,drop](#)

```

[<-(x,i,j,value)
[[<-(x,i,j,value)
\${<-(x,name,value)

```

### Author(s)

The FLR Team

### See Also

[Extract](#)

### Examples

```

flq <- FLQuant(rnorm(200), dimnames=list(age=0:4, year=1991:2000,
season=1:4))

# Extracting by position...
flq[1,]
flq[,1:5]
flq[1:2,,,c(1,3)]

# ...by dimnames
flq['0',]
flq[, '1991']
flq[,as.character(1991:1995),, '1']

# Dimensions of length one can be drop
flq[1, drop=TRUE]

# Replacing part of the object
flq['0',,,1]<-0

```

---

ffwd

---

*Project forward an FLStock for a fbar target*


---

### Description

Projection of an FLStock object for a fishing mortality target does not always require the features of fwd().Fast-forward an FLStock object for a fishing mortality yearly target only.

### Usage

```
ffwd(object, sr, fbar = control, control = fbar, deviances = "missing")
```

**Arguments**

object	An <i>FLStock</i>
sr	A stock-recruit relationship, <i>FLSR</i> or <i>predictModel</i> .
fbar	Yearly target for average fishing mortality, <i>FLQuant</i> .
control	Yearly target for average fishing mortality, <i>fwdControl</i> .
deviances	Deviances for the stock-recruit relationship, <i>FLQuant</i> .

**Value**

The projected *FLStock* object.

**Author(s)**

Iago MOSQUEIRA (MWR), Henning WINKEL (JRC).

**See Also**

[fwd](#)

**Examples**

```
data(ple4)
sr <- predictModel(model=bevholt, params=FLPar(a=140.4e4, b=1.448e5))
# Project for fixed Fbar=0.21
run <- ffwd(ple4, sr=sr, fbar=FLQuant(0.21, dimnames=list(year=1958:2017)))
plot(run)
```

---

FLArray

*Class FLArray*

---

**Description**

A basic 6D array class. No objects of this class are created in *FLCore*, as it is used only for method inheritance.

**Slots**

**.Data** Internal S4 data representation, of class *array*.

**Validity**

**Dimensions:** Array must have 6 dimensions

**Content:** Array must be of class *numeric*

**Author(s)**

The FLR Team

**See Also**

[FLQuant](#), [FLCohort](#)

---

 FLBiol

---

 Class *FLBiol*


---

**Description**

A class for modelling age / length or biomass structured populations.

**Usage**

```
FLBiol(object, ...)
```

```
## S4 method for signature 'FLQuant'
FLBiol(object, plusgroup = dims(object)$max, ...)
```

**Arguments**

<code>object</code>	FLQuant object used for sizing
<code>...</code>	Other objects to be assigned by name to the class slots
<code>plusgroup</code>	Plusgroup age, to be stored in range

**Details**

The FLBiol class is a representation of a biological fish population. This includes information on abundances, natural mortality and fecundity.

**Slots**

**n** Numbers in the population. FLQuant.  
**m** Mortality rate of the population. FLQuant.  
**wt** Mean weight of an individual. FLQuant.  
**mat** predictModel.  
**fec** predictModel.  
**rec** predictModel.  
**spwn** Proportion of time step at which spawning occurs. FLQuant.  
**name** Name of the object. character.  
**desc** Brief description of the object. character.  
**range** Named numeric vector describing the range of the object. numeric.

### Accessors

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

### Constructor

A construction method exists for this class that can take named arguments for any of its slots. All slots are then created to match the requirements of the class validity. If an unnamed FLQuant object is provided, this is used for sizing but not stored in any slot.

### Validity

**Dimensions** All FLQuant slots must have iters equal to 1 or 'n'.

**Iters** The dimname for iter1 should be '1'.

**Dimnames** The name of the quant dimension must be the same for all FLQuant slots.

### Author(s)

The FLR Team

### See Also

[as.FLBIol](#), [as.FLSR](#), [coerce](#), [plot](#), [ssb catch.n,FLBIol-method](#)

### Examples

```
# An FLBIol example dataset
data(ple4.biol)

summary(ple4.biol)
```

---

FLBiols

*Class FLBiols*

---

### Description

A list of FLBIol objects.

**Usage**

```
FLBiols(object, ...)  
  
## S4 method for signature 'FLBiol'  
FLBiols(object, ...)  
  
## S4 method for signature 'missing'  
FLBiols(object, ...)  
  
## S4 method for signature 'list'  
FLBiols(object, ...)
```

**Arguments**

object	unnamed object to be added to the list
...	other named or unnamed objects

**Slots**

**.Data** Internal S4 data representation, of class list.  
**desc** As textual description of the object contents  
**lock** Can the object be extended/trimmed? TRUE or FALSE.  
**names** A character vector for the element names

**Constructor**

A constructor method exists for this class that can take named arguments for any of the list elements.

**Author(s)**

The FLR Team

**See Also**

[FLlst](#), [list](#), [vector](#)

---

FLCohort

*Class FLCohort*

---

**Description**

A class for modelling cohorts.

**Usage**

```

FLCohort(object, ...)

## S4 method for signature 'FLQuant'
FLCohort(object, ...)

## S4 method for signature 'FLCohort'
FLCohort(object, units = units(object))

## S4 method for signature 'array'
FLCohort(
  object,
  dim = rep(1, 6),
  dimnames = "missing",
  units = "NA",
  iter = 1,
  fill.iter = TRUE
)

## S4 method for signature 'vector'
FLCohort(
  object,
  dim = c(length(object), rep(1, 5)),
  dimnames = "missing",
  units = "NA",
  iter = 1
)

## S4 method for signature 'missing'
FLCohort(object, dim = rep(1, 6), dimnames = "missing", units = "NA", iter = 1)

```

**Arguments**

<code>object</code>	Input numeric object
<code>...</code>	Additional arguments

**Details**

This class represents cohorts in columns. It simply shifts the typical matrix representation where cohorts are found on the diagonals, into a matrix where cohorts are found in columns. It is very useful for all analysis that want to make use of cohorts instead of years.

**Slots**

**.Data** Internal S4 data representation. array.  
**units** The data units in some understandable metric. character



**Constructor**

Objects of this class are generally constructed from an [FLQuant](#) object.

**Author(s)**

The FLR Team

**See Also**

[\[, as.data.frame, bubbles, ccplot, FLCohort, FLQuant-method, flc2flq, plot, quant, trim, units, units<- , FLCohort, character-method, xyplot, array](#)

**Examples**

```
data(ple4)
flq <- catch.n(ple4)
flc <- FLCohort(flq)
plot(trim(flq, cohort=1960:2000))
```

---

FLCohorts

---

*Class FLCohorts*


---

**Description**

FLCohorts is a class that extends `list` through `FL1st` but implements a set of features that give a little more structure to list objects. The elements of `FLCohorts` must all be of class `FLCohort`. It implements a lock mechanism that, when turned on, does not allow the user to increase or decrease the object length.

**Usage**

```
FLCohorts(object, ...)
```

**Arguments**

<code>object</code>	unnamed object to be added to the list
<code>...</code>	other named or unnamed objects

**Slots**

**.Data** The data. `list`

**names** Names of the list elements. `character`

**desc** Description of the object. `character`

**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. `logical`

**Constructor**

A constructor method exists for this class that can take named arguments for any of the list elements.

**Author(s)**

The FLR Team

**See Also**

[\\*](#), [Arith](#), [as.data.frame](#), [bubbles](#), [catch<-](#), [iter](#), [model.frame](#), [show](#), [summary](#), [xyplot](#), [FLlst](#), [list](#)

---

FLComp

*Class FLComp*

---

**Description**

A virtual class that forms the basis for most FLR classes composed of slots of class [FLQuant](#). No objects of this class can be constructed.

**Validity**

**Dimensions** All FLQuant slots must have iters equal to 1 or 'n'.

**Iters** The dimname for iter1 should be '1'.

**Dimnames** The name of the quant dimension must be the same for all FLQuant slots.

**Slots**

**name** A character vector for the object name.

**desc** A textual description of the object contents.

**range** A named numeric vector with various values of quant and year ranges, plusgroup, fishing mortality ranges, etc. Elements are specific to each child class.

**Author(s)**

The FLR Team

**See Also**

[\[](#), [\[<-](#), [as.data.frame](#), [iter](#), [propagate](#), [qapply](#), [summary](#), [transform](#), [trim](#), [units,FLComp-method](#), [units<-,FLComp,list-method](#), [window](#)

---

FLComps

*Class FLComps*

---

### Description

A virtual class that forms the basis for many FLR list classes. No objects of this class can be constructed.

### Arguments

object	unnamed object to be added to the list
...	other named or unnamed objects

### Validity

**Elements** All elements must be of a class that inherits from FLComp

### Slots

**.Data** The data. list.

**names** Names of the list elements. character.

**desc** Description of the object. character.

**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. logical.

### Constructor

A constructor method exists for this class that can take named arguments for any of the list elements.

### Author(s)

The FLR Team

### See Also

[FLlst](#), [FLComp](#)

FLI

Class *FLI***Description**

A VIRTUAL class that holds data and parameters related to abundance indices.

**Arguments**

object	FLQuant object used for sizing
...	Other objects to be assigned by name to the class slots

**Slots**

**distribution** Statistical distribution of the index values (character).

**index** Index values (FLQuant).

**index.var** Variance of the index (FLQuant).

**catch.n** Catch numbers used to create the index (FLQuant).

**catch.wt** Catch weight of the index (FLQuant).

**effort** Effort used to create the index (FLQuant).

**sel.pattern** Selection pattern for the index (FLQuant).

**index.q** Catchability of the index (FLQuant).

**name** Name of the stock (character).

**desc** General description of the object (character).

**range** Range of the object (numeric)

**Accessors**

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

**Constructor**

A construction method exists for this class that can take named arguments for any of its slots. All slots are then created to match the requirements of the class validity. If an unnamed FLQuant object is provided, this is used for sizing but not stored in any slot.

**Validity**

**Dimensions** All FLQuant slots must have iters equal to 1 or 'n'.

**Iters** The dimname for iter1 should be '1'.

**Dimnames** The name of the quant dimension must be the same for all FLQuant slots.

**Author(s)**

The FLR Team

**See Also**

[computeCatch](#), [dims](#), [iter](#), [plot](#), [propagate](#), [summary](#), [transform](#), [trim](#), [window](#), [FLComp](#)

---

 FLIndex

---

*Class FLIndex*


---

**Description**

A class for modelling abundance indices.

**Usage**

```
FLIndex(object, ...)
```

```
## S4 method for signature 'FLQuant'
```

```
FLIndex(object, plusgroup = dims(object)$max, ...)
```

```
## S4 method for signature 'missing'
```

```
FLIndex(object, ...)
```

**Details**

The FLIndex object holds data and parameters related to abundance indices.

**Slots**

**type** Type of index (character).

**distribution** Statistical distribution of the index values (character).

**index** Index values (FLQuant).

**index.var** Variance of the index (FLQuant).

**catch.n** Catch numbers used to create the index (FLQuant).

**catch.wt** Catch weight of the index (FLQuant).

**effort** Effort used to create the index (FLQuant).

**sel.pattern** Selection pattern for the index (FLQuant).

**index.q** Catchability of the index (FLQuant).

**name** Name of the stock (character).

**desc** General description of the object (character).

**range** Named numeric vector containing the quant and year ranges, the plusgroup, and the period of the year, expressed as proportions of a year, that corresponds to the index (numeric).

**Author(s)**

The FLR Team

**See Also**

[computeCatch](#), [dims](#), [iter](#), [plot](#), [propagate](#), [summary](#), [transform](#), [trim](#), [window](#), [FLComp](#)

**Examples**

```
# Create an FLIndex object.
fli <- FLIndex(index=FLQuant(rnorm(8), dim=c(1,8)), name="myTestFLIndex")
summary(fli)
index(fli)

# Create an FLIndex object using an existing FLQuant object.
data(ple4)
# Create a perfect index of abundance from abundance at age
fli2 <- FLIndex(index=stock.n(ple4))
# Add some noise around the signal
index(fli2) <- index(fli2)*exp(rnorm(1, index(fli2)-index(fli2), 0.1))
```

---

FLIndexBiomass

*Class FLIndexBiomass*


---

**Description**

A class for modelling biomass indices.

**Usage**

```
FLIndexBiomass(object, ...)

## S4 method for signature 'FLQuant'
FLIndexBiomass(object, plusgroup = dims(object)$max, ...)

## S4 method for signature 'missing'
FLIndexBiomass(object, ...)
```

**Arguments**

`object` FLQuant object used for sizing  
`...` Other objects to be assigned by name to the class slots

**Details**

The FLIndexBiomass object holds data and parameters related to biomass indices.

## Slots

**distribution** Statistical distribution of the index values (character).

**index** Index values (FLQuant).

**index.var** Variance of the index (FLQuant).

**catch.n** Catch numbers used to create the index (FLQuant).

**catch.wt** Catch weight of the index (FLQuant).

**effort** Effort used to create the index (FLQuant).

**sel.pattern** Selection pattern for the index (FLQuant).

**index.q** Catchability of the index (FLQuant).

**name** Name of the stock (character).

**desc** General description of the object (character).

**range** Range of the object (numeric)

## Accessors

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

## Constructor

A construction method exists for this class that can take named arguments for any of its slots. All slots are then created to match the requirements of the class validity. If an unnamed FLQuant object is provided, this is used for sizing but not stored in any slot.

## Validity

**Dimensions** All FLQuant slots must have iters equal to 1 or 'n'.

**Iters** The dimname for iter1 should be '1'.

**Dimnames** The name of the quant dimension must be the same for all FLQuant slots.

## Author(s)

The FLR Team

## See Also

[computeCatch](#), [dims](#), [iter](#), [plot](#), [propagate](#), [summary](#), [transform](#), [trim](#), [window](#), [FLComp](#)

**Examples**

```
idx <- FLIndexBiomass(index=FLQuant(1:10, quant='age'))

data(ple4)
ida <- FLIndexBiomass(index=ssb(ple4),
  catch.n=catch.n(ple4))
```

---

FLIndices

---

Class *FLIndices*


---

**Description**

FLIndices is a class that extends `list` through `FL1st` but implements a set of features that give a little more structure to list objects. The elements of `FLIndices` must all be of class `FLIndex`. It implements a lock mechanism that, when turned on, does not allow the user to increase or decrease the object length.

**Usage**

```
FLIndices(object, ...)

## S4 method for signature 'FLI'
FLIndices(object, ...)

## S4 method for signature 'missing'
FLIndices(object, ...)

## S4 method for signature 'list'
FLIndices(object, ...)
```

**Arguments**

<code>object</code>	unnamed object to be added to the list
<code>...</code>	other named or unnamed objects

**Slots**

**.Data** The data. `list`.

**names** Names of the list elements. `character`.

**desc** Description of the object. `character`.

**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. `logical`.

**Constructor**

A constructor method exists for this class that can take named arguments for any of the list elements.



**Author(s)**

The FLR Team

**See Also**

[FLlst, list](#)

**Examples**

```
data(ple4.index)
flis <- FLIndices(INDa=ple4.index, INDb=window(ple4.index, end=2000))
```

---

FLlst

*Class FLlst*


---

**Description**

FLlst is a class that extends `list` but implements a set of features that give a little more structure to list objects. First the elements of FLlst must all be of the same class. Second it implements a lock mechanism that, when turned on, does not allow the user to increase or decrease the object length.

**Usage**

```
FLlst(object, ...)
```

**Arguments**

<code>object</code>	unnamed object to be added to the list
<code>...</code>	other named or unnamed objects

**Slots**

**.Data** The data. `list`.

**names** Names of the list elements. `character`.

**desc** Description of the object. `character`.

**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. `logical`.

**Constructor**

A constructor method exists for this class that can take named arguments for any of the list elements.

**Author(s)**

The FLR Team

**See Also**

[\[, \[ \$<\$ -, \[\[ \$<\$ -, \\$ \$<\$ -, coerce, lapply, window, list](#)

**Examples**

```
f1101 <- new("FLlst", list(a=1:10, b=10:20))
f1102 <- new("FLlst", list(1:10, 10:20), names=c("a", "b"))
f1103 <- FLlst(a=1:10, b=10:20)
f1104 <- FLlst(list(a=1:10, b=10:20))
f1105 <- FLlst(c(1:10), c(10:20))
names(f1105) <- names(f1101)
names(f1101)
```

---

FLModel

*Class FLModel*


---

**Description**

A virtual class for statistical models

**Usage**

```
FLModel(model, ...)
```

**Details**

The FLModel class provides a virtual class that developers of various statistical models can use to implement classes that allow those models to be tested, fitted and presented.

Slots in this class attempt to map all the usual outputs for a modelling exercise, together with the standard inputs. Input data are stored in slots created by a specified class based on FLModel. See for example [FLSR](#) for a class used for stock-recruitment models.

The `initial` slot contains a function used to obtain initial values for the numerical solver. It can also contain two attributes, `upper` and `lower` that limit the search area for each parameter.

Various fitting algorithms, similar to those present in the basic R packages, are currently available for FLModel, including [fmle](#), [nls-FLCore](#) and [glm](#).

**Slots**

**name** Name of the object, character.

**desc** Description of the object, character.

**range** Range, numeric.

**distribution** Associated error probability distribution, factor.

**fitted** Estimated values, FLQuant.

**residuals** Residuals obtained from the model fit, FLQuant.

**model** Model formula, formula.

**gr** Function returning the gradient of the likelihood, function.

**logl** Log-likelihood function. function.

**initial** Function returning initial parameter values for the optimizer, as an object of class FLPar, function.

**params** Estimated parameter values, FLPar.

**logLik** Value of the log-likelihood, logLik.

**vcov** Variance-covariance matrix, array.

**hessian** Hessian matrix obtained from the parameter fitting, array.

**details** extra information on the model fit procedure, list.

### Author(s)

The FLR Team

### See Also

[AIC](#), [BIC](#), [fmle](#), [nls](#), [FLComp](#)

### Examples

```
# Normally, FLModel objects won't be created if "class" is not set
summary(FLModel(length~width*alpha))

# Objects of FLModel-based classes use their own constructor,
# which internally calls FLModel
fsr <- FLModel(rec~ssb*a, class='FLSR')
is(fsr)
summary(fsr)

# An example constructor method for an FLModel-based class
# Create class FLGrowth with a single new slot, 'mass'
setClass('FLGrowth', representation('FLModel', mass='FLArray'))

# Define a creator method based on FLModel
setGeneric("FLGrowth", function(object, ...) standardGeneric("FLGrowth"))
setMethod('FLGrowth', signature(object='ANY'),
  function(object, ...) return(FLModel(object, ..., class='FLGrowth')))
setMethod('FLGrowth', signature(object='missing'),
  function(...) return(FLModel(formula(NULL), ..., class='FLGrowth'))))

# Define an accessor method
setMethod('mass', signature(object='FLGrowth'),
  function(object) return(slot(object, 'mass')))
```

---

`FLModelSim`*Class FLModelSim*

---

### Description

A virtual class for statistical simulation models

### Usage

```
FLModelSim(object, ...)
```

```
## S4 method for signature 'missing'  
FLModelSim(object, ...)
```

### Details

The `FLModelSim` class provides a virtual class that developers of various statistical models can use to implement classes that allow those models to be tested, fitted and presented.

Slots in this class attempt to map all the usual outputs for a modelling exercise, together with the standard inputs. Input data are stored in slots created by a specified class that is based on `FLModelSim`. See for example [FLSR](#) for a class used for stock-recruitment models.

Various fitting algorithms, similar to those present in the basic R packages, are currently available for `FLModelSim`, including [fmle](#), [nls-FLCore](#) and [glm](#).

### Slots

**params** Estimated parameter values. `FLPar`.

**distr** character

**vcov** array

**model** formula

### Author(s)

The FLR Team

### See Also

[AIC](#), [BIC](#), [fmle](#), [nls](#)

---

`FLModelSims`*Class FLModelSims*

---

**Description**

A list of `FLModelSim` objects.

**Usage**

```
FLModelSims(object, ...)
```

```
## S4 method for signature 'ANY'  
FLModelSims(object, ...)
```

```
## S4 method for signature 'missing'  
FLModelSims(object, ...)
```

```
## S4 method for signature 'list'  
FLModelSims(object)
```

```
## S4 method for signature 'FLModelSims'  
FLModelSims(object)
```

**Arguments**

<code>object</code>	unnamed object to be added to the list
<code>...</code>	other named or unnamed objects

**Slots**

**.Data** The data. `list`.

**names** Names of the list elements. `character`.

**desc** Description of the object. `character`.

**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. `logical`.

**Constructor**

A constructor method exists for this class that can take named arguments for any of the list elements.

**Author(s)**

The FLR Team

**See Also**

[FLlst](#), [list](#), [vector](#)

---

FLPar

*Class FLPar*

---

### Description

A class for storing parameters of a model.

### Usage

```
FLPar(object, ...)
```

### Details

The FLPar class is based on the array class which can store Monte Carlo samples and the names of the relevant parameter vectors.

Methods for this class include subsetting and replacement as for the [FLQuant](#) class. There are methods for extracting statistics of the sample (mean, median etc.) and for plotting the parameter samples.

### Slots

**.Data** Describe slot. array.

**units** Units of measurement. character.

### Author(s)

The FLR Team

### See Also

[\[](#), [\[<-](#), [as.data.frame](#), [densityplot](#), [histogram](#), [iter](#), [iter<-](#), [mean](#), [median](#), [plot](#), [splom](#), [summary](#), [units,FLPar-method](#), [units<-,FLPar,character-method](#), [var](#)

### Examples

```
FLPar(rnorm(4), params=c('a','b','c','sigma2'))
```

```
FLPar(rnorm(20), dimnames=list(params=c('a','b'), year=1990:1999, iter=1),  
      units='NA')
```

```
# with iters
```

```
FLPar(rnorm(80), params=c('a', 'b'), iter=1:40)
```

---

FLParJK

*Class FLParJK*

---

### Description

A class for storing parameters of a jackknifed model fit.

### Usage

```
## S4 method for signature 'ANY'
FLParJK(object, orig)

## S4 method for signature 'FLParJK'
orig(object)
```

### Slots

.Data Jackknifed object, FLPar.  
units units of measurement, character.  
orig original object being jackknifed, FLPar.

### Validity

You can inspect the class validity function by using `getValidity(getClassDef('FLParJK'))`

### Accessors

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

### Constructor

Objects of this class are commonly created by calling the `jackknife()` method. A construction method exists for this class that can take named arguments for any of its slots. All slots are then created to match the requirements of the class validity.

### Author(s)

The FLR Team

### See Also

[FLPar](#)

---

 FLPars

 Class *FLPars*


---

### Description

A list of FLPar objects.

### Usage

```
FLPars(object, ...)

## S4 method for signature 'ANY'
FLPars(object, ...)

## S4 method for signature 'missing'
FLPars(object, ...)

## S4 method for signature 'list'
FLPars(object)

## S4 method for signature 'FLPars'
FLPars(object)
```

### Arguments

object	unnamed object to be added to the list
...	other named or unnamed objects

### Slots

**.Data** Internal S4 data representation, of class `list`.  
**desc** As textual description of the object contents  
**lock** Can the object be extended/trimmed? TRUE or FALSE.  
**names** A character vector for the element names

### Constructor

A constructor method exists for this class that can take named arguments for any of the list elements.

### Author(s)

The FLR Team

### See Also

[FLlist](#), [list](#), [vector](#)



---

FLQuant

*FLQuant class for numerical data*

---

### Description

The FLQuant class is a six-dimensional [array](#) designed to store most quantitative data used in fisheries and population modelling.

### Usage

```
FLQuant(object, ...)
```

```
## S4 method for signature 'missing'
```

```
FLQuant(  
  object,  
  dim = rep(1, 6),  
  dimnames = "missing",  
  quant = NULL,  
  units = "NA",  
  iter = 1  
)
```

```
## S4 method for signature 'vector'
```

```
FLQuant(  
  object,  
  dim = rep(1, 6),  
  dimnames = "missing",  
  quant = NULL,  
  units = "NA",  
  iter = 1,  
  fill.iter = TRUE  
)
```

```
## S4 method for signature 'array'
```

```
FLQuant(  
  object,  
  dim = rep(1, 6),  
  dimnames = "missing",  
  quant = NULL,  
  units = "NA",  
  iter = 1,  
  fill.iter = TRUE  
)
```

```
## S4 method for signature 'matrix'
```

```
FLQuant(object, dim = lapply(dimnames, length), dimnames = "missing", ...)
```

```
## S4 method for signature 'FLQuant'
FLQuant(
  object,
  quant = attributes(object)[["quant"]],
  units = attributes(object)[["units"]],
  dimnames = attributes(object)[["dimnames"]],
  iter = dim(object)[6],
  fill.iter = TRUE,
  dim = attributes(object)[["dim"]]
)
```

### Arguments

<code>object</code>	Input numeric object
<code>...</code>	Additional arguments
<code>dim</code>	Vector of dimension lengths
<code>dimnames</code>	List of dimension names
<code>quant</code>	Character vector for name of first dimension
<code>units</code>	Character vctor of units of measurement, see <a href="#">uom</a>
<code>iter</code>	Number of iterations, i.e. length of the 6th dimension
<code>fill.iter</code>	Should iterations be filled with the same content as the first?

### Details

The six dimensions are named. The name of the first dimension can be altered by the user from its default, `quant`. This could typically be `age` or `length` for data related to natural populations. The only name not accepted is `'cohort'`, as data structured along cohort should be stored using the [FLCohort](#) class instead. Other dimensions are always names as follows: `year`, for the calendar year of the datapoint; `unit`, for any kind of division of the population, e.g. by sex; `season`, for any temporal strata shorter than year; `area`, for any kind of spatial stratification; and `iter`, for replicates obtained through bootstrap, simulation or Bayesian analysis.

In addition, `FLQuant` objects contain a `units` attribute, of class [character](#), intended to contain the units of measurement relevant to the data.

### Slots

- .Data** A 6-D array for numeric data. [array](#).
- units** Units of measurement. [character](#).

### Validity

- Dimensions:** Array must have 6 dimensions
- Content:** Array must be of class `numeric`
- Dimnames:** Dimensions 2 to 6 must be named `"year"`, `"unit"`, `"season"`, `"area"` and `"iter"`

## Constructor

The FLQuant method provides a flexible constructor for objects of the class. Inputs can be of class:

**vector:** A numeric vector will be placed along the year dimension by default.

**matrix:** A matrix will be placed along dimensions 1 and 2, unless otherwise specified by 'dim'.  
The matrix dimnames will be used unless overridden by 'dimnames'.

**array:** As above

**missing:** If no input is given, an empty FLQuant (NA) is returned, but dimensions and dimnames can still be specified.

Additional arguments to the constructor:

**units:** The units of measurement, a [character](#) string.

**dim:** The dimensions of the object, a [numeric](#) vector of length 6.

**dimnames:** A [list](#) object providing the dimnames of the array. Only those different from the default ones need to be specified.

**quant:** The name of the first dimension, if different from 'quant', as a [character](#) string.

## Author(s)

The FLR Team

## See Also

[FLQuant](#)

## Examples

```
# creating a new FLQuant
flq <- FLQuant()
flq <- FLQuant(1:10, dim=c(2,5))
summary(flq)

# Vectors are used column first...
dim(FLQuant(1:10))
# ...while matrices go row first.
dim(FLQuant(matrix(1:10)))

FLQuant(matrix(rnorm(100), ncol=20))

FLQuant(array(rnorm(100), dim=c(5,2,1,1,1,10)))
FLQuant(array(rnorm(100), dim=c(5,2)), iter=10)

# working with FLQuant objects
flq <- FLQuant(rnorm(200), dimnames=list(age=1:5, year=2000:2008), units='diff')
summary(flq)

flq[1,]
flq[,1]
flq[1,1] <- 0
```

```

units(flq)
quant(flq)

plot(flq)

FLQuant()
summary(FLQuant())

FLQuant(1:10)

FLQuant(array(rnorm(9), dim=c(3,3,3)))
FLQuant(matrix(rnorm(12), nrow=4, ncol=3))

FLQuant(FLQuant(array(rnorm(9), dim=c(3,3,3)), units='kg'), units='t')
```

---

FLQuantDistr

*A class for samples of a probability distribution*


---

### Description

This extended FLQuant class holds both a measure of central tendency (mean, median) and of dispersion (typically variance), to be later used to generate, for example, random numbers with those mean and variances.

### Usage

```

FLQuantDistr(object, var, ...)

## S4 method for signature 'ANY,ANY'
FLQuantDistr(object, var, ...)

## S4 method for signature 'FLQuant,FLQuant'
FLQuantDistr(object, var, units = object@units, distr = "norm")
```

### Arguments

<code>object</code>	Input numeric object
<code>...</code>	Additional arguments

### Slots

**.Data** Unnamed slot for storing the mean (or other measure of expectation) (FLQuant).  
**var** Variance, or other measure of dispersion, (FLQuant).  
**distr** Name of the probability distribution, see Details (character).

**Validity**

**slot dims** .Data and var slots must have the same dimensions.

**slot dimnames** .Data and var slots must have the same dimnames.

You can inspect the class validity function by using `getValidity(getClassDef('FLQuantDistr'))`

**Accessors**

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

The contents of the unnamed slot (.Data) can be accessed through the `e()` method, see Example below.

**Constructor**

A construction method exists for this class that can take named arguments for any of its slots. All slots are then created to match the requirements of the class validity. If an unnamed FLQuant object is provided, this is used for the .Data slot.

**Methods**

Methods exist for various calculations based on values stored in the class:

**Arith** .

**Arithmetic**

The methods under the *Arith* group have been defined for objects of this class, both for operations between two FLQuantDistr objects and with objects of class FLQuant (FLArray) as follows:

**+**, FLQuantDistr,FLArray .

**-**, FLQuantDistr,FLArray .

**/**, FLQuantDistr,FLArray. *\item*/, FLQuantDistr,FLArray. *\item*+, FLQuantDistr,FLQuantDistr. *\item*-, FLQuantDistr

**Author(s)**

The FLR Team

**See Also**

[FLQuant](#)

**Examples**

```
data(ple4)
fqd <- FLQuantDistr(catch.n(ple4), var=catch.n(ple4) * 10, distr='norm')
```

---

FLQuantJK

*A class for jackknifing fisheries data*


---

**Description**

This extended FLQuant class holds both a jackknifed FLQuant, one in which each iter is missing one element, and the original object, as a separate FLQuant in the orig slot.

**Usage**

```
## S4 method for signature 'ANY'
FLQuantJK(object, orig)

## S4 method for signature 'FLQuantJK'
orig(object)

## S4 method for signature 'FLQuants'
orig(object)
```

**Arguments**

object	Input numeric object
...	Additional arguments

**Slots**

**.Data** Unnamed slot containing the jackknifed object([FLQuant](#)).  
**orig** Original object, ([FLQuant](#)).

**Validity**

**slot dims** .Data and orig slots must have the same dimensions 1-5.  
**slot dimnames** .Data and var slots must have the same dimnames 1-5.  
You can inspect the class validity function by using `getValidity(getClassDef('FLQuantJK'))`

**Accessors**

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

**Constructor**

Objects of this class must be constructed from an [FLQuant](#) that is to be jackknifed, through the [jackknife](#) method.

**Methods**

All methods defined for the [FLQuant](#) class are available, but they will operate only on the jackknifed (.Data) slot. Please use `orig()` to apply them to the original object stored in the class.

**Author(s)**

The FLR Team

**See Also**

[FLQuant](#)

**Examples**

```
data(ple4)
fjk <- jackknife(stock(ple4))
# New object has as many iters as length of jackknifed dimension (defaults to 'year')
dim(fjk)
```

---

FLQuantPoint

*Class FLQuantPoint*

---

**Description**

The FLQuantPoint class summarizes the contents of an FLQuant object with multiple iterations along its sixth dimension using a number of descriptive statistics.

**Usage**

```
FLQuantPoint(object, ...)

## S4 method for signature 'missing'
FLQuantPoint(..., units = "NA", n = 1)

## S4 method for signature 'FLQuant'
FLQuantPoint(object, ..., probs = c(0.25, 0.75))

## S4 method for signature 'FLQuantPoint'
n(object, ...)
```

**Arguments**

object	Input numeric object
...	Additional arguments

**Details**

An object of this class has a set structure along its sixth dimension (*iter*), which will always be of length 5, and with dimnames *mean*, *median*, *var*, *uppq* and *lowq*. They refer, respectively, to the sample mean, sample median, variance, and lower (0.25) and upper (0.75) quantiles.

Objects of this class will be typically created from an FLQuant. The various statistics are calculated along the *iter* dimension of the original FLQuant using [apply](#).

**Slots**

**.Data** The main array holding the computed statistics. array.  
**units** Units of measurement. character.

**Accessors**

**mean,mean<-:** 'mean' element on 6th dimension, arithmetic mean.  
**median,median<-:** 'median' element on 6th dimension, median.  
**var,var<-:** 'var' element on 6th dimension, variance.  
**lowq,lowq<-:** 'lowq' element on 6th dimension, lower quantile (0.25 by default).  
**uppq,uppq<-:** 'uppq' element on 6th dimension, upper quantile (0.75 by default).  
**quantile:** returns the 'lowq' or 'uppq' iter, depending on the value of 'probs' (0.25 or 0.75).

**Constructor**

Inputs can be of class:

FLQuant: An FLQuant object with iters (i.e. `dim6 > 1`)

**Validity**

**iter:** iter dimension is of length 5.

**Dimnames:** iter dimnames are 'mean', 'median', 'var', 'uppq' and 'lowq'

**Author(s)**

The FLR Team

**See Also**

[FLQuant](#)



**Examples**

```

flq <- FLQuant(rlnorm(2000), dim=c(10,20,1,1,1,200), units="kg")
flqp <- FLQuantPoint(flq)
flqp <- FLQuantPoint(flq, probs=c(0.05, 0.95))
summary(flqp)
mean(flqp)
var(flqp)
rnorm(200, flqp)

```

---

FLQuants

*Class FLQuants*


---

**Description**

FLQuants is a list of FLQuant objects. It is very similar to the standard list class. It implements a lock mechanism that, when turned on, does not allow the user to increase or decrease the object length. The elements of FLQuants must all be of class FLQuant.

**Usage**

```

FLQuants(object, ...)

## S4 method for signature 'ANY'
FLQuants(object, ...)

## S4 method for signature 'FLComp'
FLQuants(object, ...)

## S4 method for signature 'missing'
FLQuants(object, ...)

## S4 method for signature 'list'
FLQuants(object, ...)

## S4 method for signature 'FLQuants'
FLQuants(object, ...)

```

**Arguments**

object	unnamed object to be added to the list
...	other named or unnamed objects

**Slots**

**.Data** The data. list.  
**names** Names of the list elements. character.  
**desc** Description of the object. character.

**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. logical.

### Constructor

A constructor method exists for this class that can take named arguments for any of the list elements.

### Author(s)

The FLR Team

### See Also

[\\*](#), [Arith](#), [as.data.frame](#), [bubbles](#), [catch<-](#), [iter](#), [model.frame](#), [show](#), [summary](#), [xyplot](#), [FLlst](#), [list](#)

### Examples

```
# Compute various FLStock indicators
data(ple4)
fqs <- FLQuants(ssb=ssb(ple4), catch=catch(ple4), rec=rec(ple4),
  f=fbar(ple4))
summary(fqs)
xyplot(data~year|qname, fqs, type='b', scales=list(relation='free'))
```

---

FLS

*Class FLS*

---

### Description

A virtual class that forms the basis for the [FLStock](#) and [FLStockLen](#) classes. No objects of this class can be constructed.

### Validity

**None** No particular validity checks

### Slots

**catch** Total catch weight (FLQuant).  
**catch.n** Catch numbers (FLQuant).  
**catch.wt** Mean catch weights (FLQuant).  
**desc** Description of the stock (character).  
**discards** Total discards weight (FLQuant).  
**discards.n** Discard numbers (FLQuant).  
**discards.wt** Mean discard weights (FLQuant).  
**landings** Total landings weight (FLQuant).

**landings.n** Landing numbers (FLQuant).  
**landings.wt** Landing weights (FLQuant).  
**stock** Total stock weight (FLQuant).  
**stock.n** Stock numbers (FLQuant).  
**stock.wt** Mean stock weights (FLQuant).  
**m** Natural mortality (FLQuant).  
**m.spwn** Proportion of natural mortality before spawning (FLQuant).  
**mat** Proportion mature (FLQuant).  
**harvest** Harvest rate or fishing mortality. The units of this slot should be set to 'harvest' or 'f' accordingly (FLQuant).  
**harvest.spwn** Proportion of harvest/fishing mortality before spawning (FLQuant).  
**name** Name of the stock (character).  
**range** Named numeric vector containing the quant and year ranges, the plusgroup and the quant range that the average fishing mortality should be calculated over (numeric).

**Author(s)**

The FLR Team

**See Also**

[, [`<-`, `as.data.frame`, `iter`, `propagate`, `qapply`, `summary`, `transform`, `trim`, `units,FLComp-method`, `units<-`, `FLComp,list-method`, `window`

---

FLSR

*Class FLSR*

---

**Description**

Class for stock-recruitment models.

**Usage**

```
FLSR(model, ...)

## S4 method for signature 'ANY'
FLSR(model, ...)

## S4 method for signature 'missing'
FLSR(model, ...)
```

**Details**

A series of commonly-used stock-recruitment models are already available, including the corresponding likelihood functions and calculation of initial values. See [SRModels](#) for more details and the exact formulation implemented for each of them.

**Slots**

**name** Name of the object (character).  
**desc** Description of the object (character).  
**range** Range (numeric).  
**rec** Recruitment series (FLQuant).  
**ssb** Index of reproductive potential, e.g. SSB or egg or egg production (FLQuant).  
**fitted** Estimated values for rec (FLQuant).  
**residuals** Residuals obtained from the model fit (FLArray).  
**covar** Covariates for SR model (FLQuants).  
**model** Model formula (formula).  
**gr** Function returning the gradient of the likelihood (function).  
**logl** Log-likelihood function (function).  
**initial** Function returning initial parameter values for the optimizer (function).  
**params** Estimated parameter values (FLPar).  
**logLik** Value of the log-likelihood (logLik).  
**vcov** Variance-covariance matrix (array).  
**details** Extra information on the model fit procedure (list).  
**logerror** Is the error on a log scale (logical).  
**distribution** (factor).  
**hessian** Resulting Hessian matrix from the fit (array).

**Author(s)**

The FLR Team

**See Also**

[FLModel](#), [FLComp](#)

**Examples**

```
# Create an empty FLSR object.
sr1 <- FLSR()

# Create an FLSR object using the existing SR models.
sr2 <- FLSR(model = 'ricker')
sr2@model
sr2@initial
sr2@logl

sr3 <- FLSR(model = 'bevholt')
sr3@model
sr3@initial
sr3@logl
```

```
# Create an FLSR using a function.
mysr1 <- function(){
  model <- rec ~ a*ssb^b
  return(list(model = model))}

sr4 <- FLSR(model = mysr1)

# Create an FLSR using a function and check that it works.
mysr2 <- function(){
  formula <- rec ~ a+ssb*b

  logl <- function(a, b, sigma, rec, ssb) sum(dnorm(rec,
    a + ssb*b, sqrt(sigma), TRUE))

  initial <- structure(function(rec, ssb) {
    a <- mean(rec)
    b <- 1
    sigma <- sqrt(var(rec))

    return(list(a=a, b=b, sigma=sigma)),
    lower = c(0, 1e-04, 1e-04), upper = rep(Inf, 3))

  return(list(model = formula, initial = initial, logl = logl))
}

ssb <- FLQuant(runif(10, 10000, 100000))
rec <- 10000 + 2*ssb + rnorm(10,0,1)
sr5 <- FLSR(model = mysr2, ssb = ssb, rec = rec)

sr5.mle <- fmle(sr5)
sr5.nls <- nls(sr5)

# NS Herring stock-recruitment dataset
data(nsher)

# already fitted with a Ricker SR model
summary(nsher)

plot(nsher)

# change model
model(nsher) <- bevholt()

# fit through MLE
nsher <- fmle(nsher)

plot(nsher)
```

---

FLSRs *FLSRs is a class that extends list through FLlist but implements a set of features that give a little bit more structure to list objects. The elements of FLSRs must all be of class FLSR. It implements a lock mechanism that, when turned on, does not allow the user to increase or decrease the object length.*

---

### Description

FLSRs is a class that extends list through FLlist but implements a set of features that give a little bit more structure to list objects. The elements of FLSRs must all be of class FLSR. It implements a lock mechanism that, when turned on, does not allow the user to increase or decrease the object length.

### Usage

```
FLSRs(object, ...)

## S4 method for signature 'FLSR'
FLSRs(object, ...)

## S4 method for signature 'missing'
FLSRs(object, ...)

## S4 method for signature 'list'
FLSRs(object, ...)
```

### Slots

**.Data** The data. list.  
**names** Names of the list elements. character.  
**desc** Description of the object. character.  
**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. logical.

### Author(s)

The FLR Team

### See Also

[FLlist](#), [list](#), [FLSR](#)

### Examples

```
data(nsher)
bnsher <- nsher
model(bnsher) <- bevholt
```

```

bnsher <- fmle(bnsher)
fls <- FLSRs(Ricker=bnsher, BevHolt=bnsher)
summary(flS)

```

---

FLStock

*Class FLStock*


---

### Description

A class for modelling a fish stock.

### Usage

```

FLStock(object, ...)

## S4 method for signature 'FLQuant'
FLStock(object, plusgroup = dims(object)$max, ...)

## S4 method for signature 'missing'
FLStock(object, ...)

## S4 method for signature 'FLQuants'
FLStock(object, ...)

```

### Arguments

object	FLQuant object used for sizing
...	Other objects to be assigned by name to the class slots
plusgroup	Plusgroup age, to be stored in range

### Details

The FLStock object contains a representation of a fish stock as constructed for the purposes of scientific analysis and advice. This includes information on removals (i.e. catches, landings and discards), maturity, natural mortality and the results of an analytical assessment (i.e. estimates of abundance and removal rates) .

### Slots

**catch** Total catch weight (FLQuant).  
**catch.n** Catch numbers (FLQuant).  
**catch.wt** Mean catch weights (FLQuant).  
**discards** Total discards weight (FLQuant).  
**discards.n** Discard numbers (FLQuant).  
**discards.wt** Mean discard weights (FLQuant).

**landings** Total landings weight (FLQuant).

**landings.n** Landing numbers (FLQuant).

**landings.wt** Landing weights (FLQuant).

**stock** Total stock weight (FLQuant).

**stock.n** Stock numbers (FLQuant).

**stock.wt** Mean stock weights (FLQuant).

**m** Natural mortality (FLQuant).

**mat** Proportion mature (FLQuant).

**harvest** Harvest rate or fishing mortality. The units of this slot should be set to 'hr' or 'f' accordingly (FLQuant).

**harvest.spwn** Proportion of harvest/fishing mortality before spawning (FLQuant).

**m.spwn** Proportion of natural mortality before spawning (FLQuant).

**name** Name of the stock (character).

**desc** Description of the stock (character).

**range** Named numeric vector containing the quant and year ranges, the plusgroup and the quant range that the average fishing mortality should be calculated over (numeric).

### Accessors

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

### Constructor

A construction method exists for this class that can take named arguments for any of its slots. All slots are then created to match the requirements of the class validity. If an unnamed FLQuant object is provided, this is used for sizing but not stored in any slot.

### Author(s)

The FLR Team

### See Also

[\[](#), [\[<-](#), [as.FLBIol](#), [as.FLSR](#), [catch](#), [catch<-](#), [catch.n](#), [catch.n<-](#), [catch.wt](#), [catch.wt<-](#), [coerce](#), [computeCatch](#), [computeDiscards](#), [computeLandings](#), [discards](#), [discards<-](#), [discards.n](#), [discards.n<-](#), [discards.wt](#), [discards.wt<-](#), [harvest](#), [harvest<-](#), [harvest.spwn](#), [landings](#), [landings<-](#), [landings.n](#), [landings.n<-](#), [landings.wt](#), [landings.wt<-](#), [m](#), [m<-](#), [mat](#), [m.spwn](#), [plot](#), [ssb](#), [ssbpurec](#), [stock](#), [stock.n](#), [stock.wt](#), [trim](#), [FLComp](#)



**Examples**

```

data(ple4)
summary(ple4)

# get the landings slot and assign values to it
landings(ple4)
landings(ple4) <- apply(landings.n(ple4)*landings.wt(ple4),2,sum)

# perform similar calculation as the preceding apply function
discards(ple4) <- computeDiscards(ple4)
catch(ple4) <- computeCatch(ple4)
catch(ple4) <- computeCatch(ple4, slot="all")

# set the units of the harvest slot of an FLStock object
harvest(ple4) <- 'f'

# subset and trim the FLStock
ple4[,1]
trim(ple4, age=2:6, year=1980:1990)

# Calculate SSB, and SSB per recruit at zero fishing mortality
ssb(ple4)
ssbpurec(ple4)

# Coerce an FLStock to an FLBiol
biol <- as(ple4, "FLBiol")

# Initialise an FLSR object from an FLStock
flsr <- as.FLSR(ple4)

```

---

FLStockLen

*Class FLStockLen*


---

**Description**

A class for modelling a length-structured fish stock.

**Usage**

```

FLStockLen(object, ...)

## S4 method for signature 'FLQuant'
FLStockLen(object, ...)

## S4 method for signature 'missing'
FLStockLen(object, ...)

```

## Details

The FLStockLen object contains a length based representation of a fish stock. This includes information on removals (i.e. catches, landings and discards), maturity, natural mortality and the results of an analytical assessment (i.e. estimates of abundance and removal rates).

## Slots

**halfwidth** The middle of the length bins (numeric).

**catch** Total catch weight (FLQuant).

**catch.n** Catch numbers (FLQuant).

**catch.wt** Mean catch weights (FLQuant).

**discards** Total discards weight (FLQuant).

**discards.n** Discard numbers (FLQuant).

**discards.wt** Mean discard weights (FLQuant).

**landings** Total landings weight (FLQuant).

**landings.n** Landing numbers (FLQuant).

**landings.wt** Landing weights (FLQuant).

**stock** Total stock weight (FLQuant).

**stock.n** Stock numbers (FLQuant).

**stock.wt** Mean stock weights (FLQuant).

**m** Natural mortality (FLQuant).

**mat** Proportion mature (FLQuant).

**harvest** Harvest rate or fishing mortality. The units of this slot should be set to 'harvest' or 'f' accordingly (FLQuant).

**harvest.spwn** Proportion of harvest/fishing mortality before spawning (FLQuant).

**m.spwn** Proportion of natural mortality before spawning (FLQuant).

**name** Name of the stock (character).

**desc** Description of the stock (character).

**range** Named numeric vector containing the quant and year ranges, the plusgroup and the quant range that the average fishing mortality should be calculated over (numeric).

## Author(s)

The FLR Team

## See Also

[\[](#), [\[<-](#), [as.FLBIol](#), [as.FLSR](#), [computeCatch](#), [computeDiscards](#), [computeLandings](#), [plot](#), [ssb](#), [ssbpurec](#), [trim](#), [FLComp](#)

**Examples**

```

stk1 <- FLStockLen(m=FLQuant(0.2, dimnames=list(len=seq(5, 50, by=2), year=2015:2020)))
summary(stk1)
# Unnamed FLQuant used for sizing
stk1 <- FLStockLen(FLQuant(0.2, dimnames=list(len=seq(5, 50, by=2), year=2015:2020)))
summary(stk1)
m(stk1)

```

---

FLStocks

*Class FLStocks*


---

**Description**

FLStocks is a class that extends `list` through `FLlst` but implements a set of features that give a little bit more structure to list objects. The elements of FLStocks must all be of class `FLStock`. It implements a lock mechanism that, when turned on, does not allow the user to increase or decrease the object length.

**Usage**

```

FLStocks(object, ...)

## S4 method for signature 'FLStock'
FLStocks(object, ...)

## S4 method for signature 'missing'
FLStocks(object, ...)

## S4 method for signature 'list'
FLStocks(object, ...)

```

**Arguments**

<code>object</code>	unnamed object to be added to the list
<code>...</code>	other named or unnamed objects

**Slots**

**.Data** The data. `list`.

**names** Names of the list elements. `character`.

**desc** Description of the object. `character`.

**lock** Lock mechanism, if turned on the length of the list can not be modified by adding or removing elements. `logical`.

**Constructor**

A constructor method exists for this class that can take named arguments for any of the list elements.

**Author(s)**

The FLR Team

**See Also**

[plot](#), [FLlst](#), [list](#)

**Examples**

```
data(ple4)
fls <- FLStocks(sa=ple4, sb=window(ple4, end=1980))
summary(fls)
```

---

FUNCTION

*Extract and modify the recruitment time series*

---

**Description**

Recruitment in number of fish is the first row of the 'stock.n' slot of an age-structured 'FLStock'. These convenience functions allow a clearer syntax when retrieving or altering the content of 'stock.nrec.age', where 'rec.age' is usually the first age in the object.

**Usage**

```
## S4 method for signature 'FLStock'
rec(object, rec.age = as.character(object@range["min"]))
```

**Arguments**

object	An object of class 'FLStock'
rec.age	What age to extract, defaults to first one. As 'character' to select by name or as 'numeric' by position.

**Value**

RETURN Lorem ipsum dolor sit amet

**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

**Examples**

```

data(ple4)
rec(ple4)
# Multiple recruitment by a factor of 2
rec(ple4) <- rec(ple4) * 2

```

---

Funwanted

*Calculate the discards and landings-associated fishing mortalities*


---

**Description**

Computes the fishing mortality at age (harvest) associated with either landings (*Fwanted*) or discards (*Funwanted*) through the respective proportions at age. The function names reflect the convention used in ICES.

**Usage**

```
Funwanted(x, ages = dimnames(x)$age)
```

**Arguments**

`x` An FLStock object, with harvest  
`ages` Ages over which the respective Fbar calculation applies

**Value**

An FLQuant

**Examples**

```

data(ple4)
Fwanted(ple4, ages=2:6)
Funwanted(ple4, ages=1:3)

```

---

fwdWindow

*Extend a FLR object along the year dimension and set future assumed values*


---

**Description**

Objects to be projected into the future are extended until an end year, and the values of certain quantities, usually assume constant, are set following different mechanisms.

**Usage**

```
fwdWindow(x, y, ...)

## S4 method for signature 'FLStock,missing'
fwdWindow(
  x,
  end = dims(x)$maxyear,
  nsq = 3,
  fun = c("mean", "sample"),
  years = list(wt = nsq, mat = nsq, m = nsq, spwn = nsq, discards.ratio = nsq, catch.sel
    = nsq)
)
```

**Arguments**

**x**                    The FLR object to extend.

**y**                    A second object from which information is taken.

**Details**

For 'FLStock'

**Value**

An object of the same class as 'x'.

**Author(s)**

The FLR Team.

**See Also**

[window\(\)](#)

**Examples**

```
data(ple4)
# Use mean of last three years and extend until 2020
fut <- fwdWindow(ple4, end=2020)
# Check values on catch.wt
catch.wt(fut)[, ac(2015:2020)]
# Use mean of the 2010:2015 period
fut <- fwdWindow(ple4, end=2020, years=2010:2015)
# Use last three years mean, but last five for 'wt'
fut <- fwdWindow(ple4, end=2020, nsq=3, years=list(wt=5))
stock.wt(fut)[, ac(2013:2020)]
catch.sel(fut)[, ac(2013:2020)]
# Resample from last years for 'wt'
fut <- fwdWindow(ple4, end=2020, nsq=3, fun=c(wt='sample'))
# Years to resample can be different for 'catch.sel'
```

```
fut <- fwdWindow(ple4, end=2020, nsq=3,
  fun=c(wt='sample', catch.sel='sample'), years=c(wt=10, catch.sel=5))
# 'wt' slot has been resampled,
stock.wt(fut)[, ac(2015:2020)]
# while others have used a 3 year average
catch.sel(fut)[, ac(2015:2020)]
```

---

getSlotNamesClass      *Names of slots of a given class*

---

### Description

This function returns the names, as a character vector, of the slots in an S4 object that are of the class specified by the 'class' argument. Comparison is done using is(), so class inheritance is matched.

### Usage

```
getSlotNamesClass(object, class)
```

### Arguments

object	An S4 object to check slots from.
class	The name of the class to match, 'character'.

### Author(s)

The FLR Team

### Examples

```
data(ple4)
getSlotNamesClass(ple4, 'FLQuant')
```

---

group      *Group objects over some index by applying a function over a single dimension*

---

### Description

Array objects (e.g. FLQuant or FLQuant) are divided along a single dimension following a given index or expression, an aggregating function is applied to each subset, and the results are joined again. Data can be added, for example, by decade or for two age groups.

**Usage**

```
group(x, FUN, ...)

## S4 method for signature 'FLQuant,function'
group(x, FUN = sum, ...)
```

**Arguments**

**x** An object to group.

**FUN** A function to apply along the chosen dimension, defaults to 'sum'.

**...** An expression or indexing vector, named as the chosen dimension. Extra arguments to FUN can also be provided, but cannot match names in x.

**Value**

A single object with reduced dimensionality.

**Author(s)**

Iago Mosqueira (WMR)

**Examples**

```
data(ple4)
# Add catch-at-age along two age groups, 'juv'eniles and 'adu'lts
group(catch.n(ple4), sum, age=c('juv', 'juv', rep('adu', 8)))
# An expression can use based on dimnames
group(catch.n(ple4), sum, age=age < 3)
# Mean by lustrum, by using 'year - year %% 5'
group(catch.n(ple4), mean, year = year - year %% 5)
```

---

iav

---

*Compute the inter-annual variability of a time series*


---

**Description**

The inter-annual variability of a time series stored in an FLQuant object, is computed as  $|x_y - x_{y-1}|/x_{y-1}$ . The resulting object will be one year shorter than the input. The first year will be missing as values are assigned to the final year of each pair.

**Usage**

```
iav(object)
```

**Value**

An object of the same class as object.



**Author(s)**

The FLR Team

**Examples**

```
data(ple4)
# Compute inter-annual variability in catch
iav(catch(ple4))
```

---

indicators.len	<i>Calculate quantile(s) of length distribution</i>
----------------	---

---

**Description**

$$z = (k * (l_{inf} - l_{mean})) / (l_{mean} - l_c)$$

$l_{mean} = \text{sum}(naa * len) / \text{sum}(naa)$   $l_c$ , length at first capture

**Usage**

```
indicators.len(
  object,
  indicators = "lbar",
  model = vonbert,
  params,
  cv = 0.1,
  lmax = 1.25,
  bin = 1,
  n = 500,
  metric = catch.n,
  ...
)

lenquantile(x, quantile = 0.5)

lmax5(x)

l95(x)

l25(x)

lc50(x)

lmode(x)

lbar(x)

lmean(x)
```

```
lmaxy(x, lenwt)

pmeqa(x, linf, lopt = linf * 2/3)

bheqz(x, linf, k, t0, lc = lc50(x))
```

## References

- Kell, L.T., Minto, C., Gerritsen, H.D. 2022. Evaluation of the skill of length-based indicators to identify stock status and trends. ICES Journal of Marine Science. doi: 10.1093/icesjms/fsac043.
- ICES. 2015. Report of the Fifth Workshop on the Development of Quantitative Assessment Methodologies based on Life-history Traits, Exploitation Characteristics and other Relevant Parameters for Data-limited Stocks (WKLIFE V), 5–9 October 2015, Lisbon, Portugal. ICES CM 2015/ACOM:56. 157 pp.
- ICES. 2020. Tenth Workshop on the Development of Quantitative Assessment Methodologies based on LIFE-history traits, exploitation characteristics, and other relevant parameters for data-limited stocks (WKLIFE X). ICES Scientific Reports. 2:98. 72 pp. <http://doi.org/10.17895/ices.pub.5985>

## Examples

```
data(ple4)
indicators.len(ple4, indicators=c('lbar', 'lmaxy'),
  params=FLPar(linf=132, k=0.080, t0=-0.35), metric='catch.n',
  lenwt=FLPar(a=0.01030, b=2.975))
indicators.len(ple4, indicators=c('pmeqa'),
  params=FLPar(linf=60, k=2.29e-01, t0=-1.37), metric='catch.n')
data(ple4.index)
indicators.len(ple4.index, indicators=c('lbar', 'lmean'),
  params=FLPar(linf=132, k=0.080, t0=-0.35), metric='index')
#
ialk <- invALK(params=FLPar(linf = 60, k = 2.29e-01, t0 = -1.37e+00),
  model=vonbert, age=1:10, lmax=1.2)
samps <- lenSamples(catch.n(ple4), invALK=ialk, n=250)
lenquantile(samps, 0.50)
lmax5(samps)
l95(samps)
l25(samps)
lc50(samps)
lmode(samps)
lbar(samps)
lmean(samps)
# Linf(ple4) = 60
lmean(samps) / (0.75 * lc50(samps) + 0.25 * 60) #
lenwt <- FLPar(a=0.01030, b=2.975)
lmaxy(samps, lenwt)
pmeqa(samps, linf=60)
linf <- 60
k <- 2.29e-01
t0 <- -1.37e+00
bheqz(samps, linf = 60, k = 2.29e-01, t0 = -1.37e+00)
```

---

intersect	Returns <i>FLR</i> objects trimmed to their shared dimensions.
-----------	--

---

### Description

Objects sharing certain dimensions, as inferred by their *dimnames*, are subset to the common ones along all dimensions. The returned object is of one of the *FLlst* classes, as corresponds to the input class. The objects in the list can then be, for example, combined or directly compared, as shown in the examples.

### Usage

```
## S4 method for signature 'FLArray,FLArray'  
intersect(x, y)
```

### Arguments

x	First object to be compared and subset
y	Second object to be compared and subset

### Value

And object of the corresponding *FLsdt*-based plural class.

### Author(s)

The FLR Team

### See Also

[base::intersect](#)

### Examples

```
big <- FLQuant(64.39, dimnames=list(age=1:4, year=2001:2012))  
small <- FLQuant(3.52, dimnames=list(age=2:3, year=2001:2005))  
intersect(big, small)  
  
# Two FLQuant objects can be added along their common dimension using Reduce()  
Reduce('+', intersect(big, small))
```

iter

*Methods iter***Description**

Select or modify iterations of an FLR object

**Usage**

```
iter(obj, ...)

## S4 method for signature 'FLArray'
iter(obj, iter)
```

**Details**

To extract or modify a subset of the iterations contained in an FLR object, the `iter` and `iter<-` methods can be used.

In complex objects with various `FLQuant` slots, the `iter` method checks whether individual slots contain more than one iteration, i.e. `dims(object)[6] > 1`. If a particular slot contains a single iteration, that is returned, otherwise the chosen iteration is selected. This is in contrast with the subset operator `[`, which does not carry out this check.

For objects of class [FLModel](#), iters are extracted for slots of classes `FLQuant`, `FLCohort` and `FLPar`.

**Generic function**

```
iter(object) iter<-(object,value)
```

**Author(s)**

The FLR Team

**See Also**

[FLComp](#), [FLQuant](#)

**Examples**

```
# For an FLQuant
flq <- FLQuant(rnorm(800), dim=c(4,10,2), iter=10)
iter(flq, 2)

# For the more complex FLStock object
data(ple4)
fls <- propagate(ple4, 10)
# Extraction using iter...
fls2 <- iter(fls, 2)
summary(fls2)
```

---

`jackknife`*Method jackknife*

---

**Description**

Jackknife resampling

**Usage**

```
## S4 method for signature 'FLQuant'  
jackknife(object, dim = "year", na.rm = TRUE)  
  
## S4 method for signature 'FLQuants'  
jackknife(object, ...)  
  
## S4 method for signature 'FLModel'  
jackknife(object, slot)
```

**Details**

The `jackknife` method sets up objects ready for jackknifing, i.e. to systematically recompute a given statistic leaving out one observation at a time. From this new set of "observations" for the statistic, estimates for the bias and variance of the statistic can be calculated.

Input objects cannot have length > 1 along the `iter` dimension, and the main slot in the resulting object will have as many `iters` as the number of elements in the original object that are not NA.

**Generic function**

```
jackknife(object, ...)
```

**Author(s)**

The FLR Team

**See Also**

[FLQuantJK](#) [FLParJK](#)

**Examples**

```
flq <- FLQuant(1:8)  
flj <- jackknife(flq)  
iters(flj)
```

---

`join`*Joins objects along a dimensions where dimnames differ*

---

**Description**

FLQuant objects are joined along a single dimension, on which dimnames are different. This is the reverse operation to divide.

**Usage**

```
join(x, y, ...)  
  
## S4 method for signature 'FLQuant,FLQuant'  
join(x, y)  
  
## S4 method for signature 'FLQuants,missing'  
join(x, y)
```

**Arguments**

<code>x</code>	An object to join
<code>y</code>	An object to join

**Value**

A single object

**Author(s)**

Iago Mosqueira (WMR)

**Examples**

```
data(ple4)  
# JOIN over age dimension  
x <- catch.n(ple4)[1,]  
y <- catch.n(ple4)[2,]  
join(x, y)  
# JOIN over year dimension  
x <- catch.n(ple4)[,10:20]  
y <- catch.n(ple4)[,21:25]  
join(x, y)  
div <- divide(catch.n(ple4), dim=1)  
is(div)  
length(div)  
join(div)  
all.equal(join(divide(catch.n(ple4), dim=1)), catch.n(ple4))
```

---

lattice

*Lattice methods*

---

## Description

Implementation of Trellis graphics in FLR

## Usage

```
## S4 method for signature 'formula,FLQuant'  
xyplot(x, data, ...)  
  
## S4 method for signature 'formula,FLCohort'  
xyplot(x, data, ...)  
  
## S4 method for signature 'formula,FLQuants'  
xyplot(x, data, ...)  
  
## S4 method for signature 'formula,FLComp'  
xyplot(x, data, ...)  
  
## S4 method for signature 'formula,FLQuant'  
bwplot(x, data, ...)  
  
## S4 method for signature 'formula,FLComp'  
bwplot(x, data, ...)  
  
## S4 method for signature 'formula,FLQuant'  
dotplot(x, data, ...)  
  
## S4 method for signature 'formula,FLComp'  
dotplot(x, data, ...)  
  
## S4 method for signature 'formula,FLQuant'  
barchart(x, data, ...)  
  
## S4 method for signature 'formula,FLComp'  
barchart(x, data, ...)  
  
## S4 method for signature 'formula,FLQuant'  
stripplot(x, data, ...)  
  
## S4 method for signature 'formula,FLComp'  
stripplot(x, data, ...)  
  
## S4 method for signature 'formula,FLQuant'  
histogram(x, data, ...)
```

```
## S4 method for signature 'formula,FLComp'
histogram(x, data, ...)

## S4 method for signature 'formula,FLQuants'
histogram(x, data, ...)

## S4 method for signature 'formula,FLPar'
densityplot(x, data, ...)
```

### Details

Plot methods in the [lattice](#) package are available for an object of classes FLQuant, FLQuants or those derived from FLComp.

See the help page in [lattice](#) for a full description of each plot method and all possible arguments.

Plot methods from lattice are called by passing a [data.frame](#) obtained by converting the FLR objects using [as.data.frame](#). For details on this transformation, see [as.data.frame-FLCore](#).

### Generic function

```
barchart(x, data, ...)
bwplot(x, data, ...)
densityplot(x, data, ...)
dotplot(x, data, ...)
histogram(x, data, ...)
stripplot(x, data, ...)
xyplot(x, data, ...)
```

### Author(s)

The FLR Team

### See Also

[xyplot](#), [barchart](#), [bwplot](#), [densityplot](#), [dotplot](#), [histogram](#), [stripplot](#)

### Examples

```
data(ple4)
# xyplot on FLQuant
xyplot(data~year|age, catch.n(ple4)[, 1:20])
xyplot(data~year|as.factor(age), catch.n(ple4)[, 1:20], type='b', pch=19,
       cex=0.5)

# bwplot on FLQuant with iter...
flq <- rnorm(100, catch.n(ple4)[, 1:20], catch.n(ple4)[,1:20])
bwplot(data~year|as.factor(age), flq)
# ...now with same style modifications
```



```

bwplot(data~year|as.factor(age), flq, scales=list(relation='free',
x=list(at=seq(1, 20, by=5),
labels=dimnames(catch.n(ple4)[,1:20])$year[seq(1, 20, by=5)])),
cex=0.5, strip=strip.custom(strip.names=TRUE, strip.levels=TRUE,
var.name='age'))

```

---

mase

---

*Compute mean absolute scaled error (MASE)*


---

### Description

Franses, PH. "A note on the Mean Absolute Scaled Error". *International Journal of Forecasting*. 32 (1): 20–22. doi:10.1016/j.ijforecast.2015.03.008.

### Usage

```

mase(ref, preds, ...)

## S4 method for signature 'FLQuant,FLQuants'
mase(ref, preds, order = c("inverse", "ahead"))

## S4 method for signature 'FLIndices,list'
mase(ref, preds, order = "inverse", wt = "missing")

```

### Arguments

ref	Reference or naive prediction.
preds	Predictions to compare to reference.
...	Extra arguments.
order	Are predictions in 'inverse' (default) or 'ahead' order.
wt	Mean weights-at-age to use with indices.

### Value

A numeric vector of the same length as 'preds'.

---

meanage	<i>Calculate the mean age in the stock and catch</i>
---------	--

---

**Description**

Average age in the stock numbers or catch-at-age.

**Usage**

```
meanage(object)
```

```
meanageCatch(object)
```

**Arguments**

object            An age-structured FLStock object

**Value**

An FLQuant object

**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

**Examples**

```
data(ple4)
meanage(ple4)
meanageCatch(ple4)
```

---

meanwt	<i>Calculate the mean weight in stock and catch</i>
--------	---

---

**Description**

Average weight in the stock numbers or catch-at-age.

**Usage**

```
meanwt(object)
```

```
meanwtCatch(object)
```

**Arguments**

object            An age-structured FLStock object

**Value**

An FLQuant object

**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

**Examples**

```
data(ple4)
meanwt(ple4)
meanwtCatch(ple4)
```

---

metrics

*Extract simply-defined metrics from complex objects*

---

**Description**

Time series summaries of complex objects are commonly needed, for example for plotting the inputs and outputs of a class like [FLStock](#). These methods allow for simple specification of those metrics by means of function calls and formulas.

**Usage**

```
metrics(object, metrics, ...)

## S4 method for signature 'FLComp,list'
metrics(object, metrics, ...)

## S4 method for signature 'FLS,missing'
metrics(object, metrics, ...)
```

**Arguments**

object            A complex **FLR** object from which to extract time series metrics.

**Value**

An object, generally of class [FLQuants](#).

**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

**Examples**

```
data(ple4)
# missing
metrics(ple4)
# metrics = function
metrics(ple4, metrics=function(x) FLQuants(SSB=ssb(x), REC=rec(x),
  F=fbar(x), SSBREC=ssb(x) / rec(x)))
# metrics = formula
metrics(ple4, metrics=~ssb)
metrics(ple4, metrics=list(SSB=~ssb))
metrics(ple4, metrics=list(SBMSY=~ssb/SBMSY), FLPar(SBMSY=3.87e4))
# metrics = list
metrics(ple4, metrics=list(SSB=ssb, REC=rec, F=fbar))
metrics(ple4, metrics=list(SSB=~ssb, REC=rec, F=fbar))
```

---

mohnMatrix

*Generate a matrix to compute Mohn's rho for a single metric*

---

**Description**

A common measure of the strength of stock assessment retrospective patterns is Mohn's rho. This function does not carry out the calculation but returns a matrix with the metrics value for the  $n$  retrospective runs, in columns, and  $n + 2$  years, in rows.

**Usage**

```
mohnMatrix(stocks, metric = "fbar", ...)
```

**Arguments**

stocks	An FLStocks object from a retrospective analysis
metric	Metric to be computed, as a character vector or function

**Value**

A metrics of  $n + 2 \times n$ , where  $n$  is the numbers of objects in stocks.

---

msy	<i>msy: A series of methods to extract or compute MSY-based reference points</i>
-----	--

---

### Description

Reference points based on equilibrium calculations of Maximum Sustainable Yield (MSY) are computed by various FLR packages. The methods' generics are defined here for convenience. Please refer to the help pages of particular methods for further details

### Usage

```
msy(x, ...)
```

```
bmsy(x, ...)
```

```
sbmsy(x, ...)
```

```
fmsy(x, ...)
```

### Arguments

x                    An input object from which to extract or compute a reference point

### Details

The four methods provide the following parameter estimates:

- `msy` Maximum Sustainable Yield (MSY)
- `fmsy` Fishing mortality level expected to produce on average MSY
- `bmsy` Total biomass that should produce MSY
- `sbmsy` Spawning biomass that should produce MSY

### Value

A value for the requested reference point, 'FLPar'

### Author(s)

The FLR Team

### See Also

[FLPar](#)

---

names

*Method names*

---

### Description

The names method returns the names of the dimnames of an object. For some classes, the names attribute can be modified directly using names<-.

### Usage

```
## S4 method for signature 'FLArray'  
names(x)  
  
## S4 method for signature 'FLPar'  
names(x)  
  
## S4 replacement method for signature 'FLPar,character'  
names(x) <- value
```

### Generic function

```
names(x) names<-(x, value)
```

### Author(s)

The FLR Team

### See Also

[names](#)

### Examples

```
# FLQuant  
data(ple4)  
names(catch.n(ple4))  
  
# Contrast this with  
dimnames(catch.n(ple4))
```

---

plot

*Method plot*

---

### Description

Standard plot methods for every FLCore class. FLR plot methods are based on [lattice](#), and attempt to show a general view of the object contents.

### Usage

```
## S4 method for signature 'FLQuant,missing'
plot(
  x,
  xlab = "year",
  ylab = paste("data (", units(x), ")", sep = ""),
  type = "p",
  ...
)

## S4 method for signature 'FLStock,missing'
plot(x, auto.key = TRUE, ...)

## S4 method for signature 'FLBiol,missing'
plot(x, y, ...)

## S4 method for signature 'FLIndex,missing'
plot(x, type = c("splom"), ...)

## S4 method for signature 'FLSR,missing'
plot(x, main = "Functional form", log.resid = FALSE, cex = 0.8)

## S4 method for signature 'FLPar,missing'
plot(x, y = "missing", ...)
```

### Details

Users are encouraged to write their own plotting code and make use of the overloaded [lattice](#) methods, for example [xyplot](#) or [bwplot](#). See also [lattice-FLCore](#).

### Generic function

```
plot(x,y)
```

### Author(s)

The FLR Team

**See Also**[plot](#)**Examples**

```
data(ple4)

# FLQuant
plot(catch.n(ple4)[, 1:20])
plot(catch.n(ple4)[, 1:20], type='b', pch=19, cex=0.5)

# FLStock
data(ple4sex)
plot(ple4)
plot(ple4sex)

# FLBiol
data(ple4.biol)
plot(ple4.biol)

# FLIndex
data(ple4.index)
plot(ple4.index)

# FLSR
data(nsher)
plot(nsher)

# FLPar
fpa <- FLPar(a=rnorm(100, 1, 20), b=rlnorm(100, 0.5, 0.2))
plot(fpa)
```

---

predictModel

*A class for model prediction*

---

**Description**

Object of the predictModel class are used in various FLR classes to allow flexible modelling of the dynamics of different biological and technological processes.

**Usage**

```
## S4 method for signature 'FLQuants,formula'
predictModel(object, model, params = FLPar())

## S4 method for signature 'FLQuants,missing'
predictModel(object, params = FLPar())
```



```
## S4 method for signature 'FLQuants,character'
predictModel(object, model, params = FLPar())

## S4 method for signature 'FLQuants,function'
predictModel(object, model, params = FLPar())

## S4 method for signature 'FLQuants,list'
predictModel(object, model, params = FLPar())

## S4 method for signature 'missing,ANY'
predictModel(object, model, ...)
```

### Details

The dependency of life history processes, such as maturity and fecundity, to biological and environmental factors, can be represented in objects of this class via a simple model (represented by a formula) and the corresponding parameters (FLPar) and inputs (FLQuants).

### Slots

**.Data** Inputs to the model not found in enclosing class (FLQuants).  
**model** Model representation (formula).  
**params** Model parameters (FLPar).

### Validity

**VALIDITY** Neque porro quisquam est qui dolorem ipsum.

You can inspect the class validity function by using `getValidity(getClassDef('predictModel'))`

### Accessors

All slots in the class have accessor and replacement methods defined that allow retrieving and substituting individual slots.

The values passed for replacement need to be of the class of that slot. A numeric vector can also be used when replacing FLQuant slots, and the vector will be used to substitute the values in the slot, but not its other attributes.

### Constructor

A construction method exists for this class that can take named arguments for any of its slots. All slots are then created to match the requirements of the class validity.

### Methods

Methods exist for various calculations based on values stored in the class:

**METHOD** Neque porro quisquam est qui dolorem ipsum.

**Author(s)**

The FLR Team

**See Also**

[FLQuants](#) [FLPar](#) [FLBiol](#)

**Examples**

```
fec <- FLQuants(fec=FLQuant(rlnorm(10, 20, 5),
  dimnames=list(year=2000:2009), units='1'))
predictModel(fec, model=~fec)
predictModel(fec)
predictModel(fec, model="bevholt")
predictModel(fec, model=bevholt)
predictModel(fec, model=bevholt())
predictModel(model=rec~a*ssb, params=FLPar(a=1.234))
predictModel(model=bevholt, params=FLPar(a=1.234))
predictModel(model="bevholtss3", params=FLPar(a=1.234))
```

---

production

*Returns the computed yearly production*

---

**Description**

Returns the computed yearly production

**Usage**

```
production(object, ...)
```

```
## S4 method for signature 'FLStock'
production(object, what = "ssb", ...)
```

**Arguments**

**object**            An object with biomass and catch data.

**what**             One of the production options: "ssb", "biomass", or "exploitation".

**Details**

Production can be calculated for an FLStock based on the spawning stock biomass ("ssb"), total biomass ("biomass"), or exploitation ("exploitation").

**Value**

The production by year, of class FLQuant.

**Author(s)**

Laurie Kell (Sea++), Iago Mosqueira (WMR)

**Examples**

```
data(ple4)
# For SSB
production(ple4, "ssb")
# For total biomass
production(ple4, "biomass")
```

---

propagate

*Method propagate*

---

**Description**

Methods to extend objects of various FLR classes along the `iter` (6th `FLQuant`) dimension. Objects must generally have a single `iter` to be extended. The new iterations can be filled with copies of the existing, or remain as NA.

**Usage**

```
propagate(object, ...)

## S4 method for signature 'FLQuant'
propagate(object, iter, fill.iter = TRUE)
```

**Arguments**

<code>object</code>	Object to be propagated.
<code>fill.iter</code>	Should first array be copied to others? Defaults to FALSE.
<code>iters</code>	No. of iterations in output.

**Generic function**

```
propagate(object, ...)
```

**Author(s)**

The FLR Team

**See Also**

[FLQuant](#)

**Examples**

```
# An FLQuant with one iter (dim(flq)[6] == 1)
flq <- FLQuant(rnorm(80), dim=c(4,20), quant='age')

# can now be extended along the `iter` dimension, with
# 'copies of the first
propagate(flq, 100)

# or without
iter(propagate(flq, 100, fill.iter=FALSE), 2)
```

---

properties	<i>Returns a series of properties of the fisheries element represented by the class.</i>
------------	--

---

**Description**

Returns a series of properties of the fisheries element represented by the class.

**Usage**

```
properties(object, ...)
```

**Arguments**

object            An object from which properties can be extracted.

**Value**

The corresponding properties, an FLPar.

**Author(s)**

Laurie Kell (Sea++), Iago Mosqueira (WMR)

---

quant	<i>Method quant</i>
-------	---------------------

---

**Description**

Function to get or set the name of the first dimension (quant) in an object of any FLArray-based class, like [FLQuant](#) or [FLCohort](#).

**Usage**

```
quant(object, ...)  
  
## S4 method for signature 'FLArray'  
quant(object)  
  
## S4 replacement method for signature 'FLArray,character'  
quant(object) <- value
```

**Generic function**

```
quant(object) quant<-(object,value)
```

**Author(s)**

The FLR Team

**See Also**

[FLQuant](#), [FLCohort](#)

**Examples**

```
# quant is 'quant' by default  
quant(FLQuant())  
  
flq <- FLQuant(rnorm(80), dim=c(4,20), quant='age')  
quant(flq)  
quant(flq) <- 'length'  
summary(flq)  
  
# quant is 'quant' by default  
quant(FLQuant())  
  
flq <- FLQuant(rnorm(80), dim=c(4,20), quant='age')  
quant(flq)  
quant(flq) <- 'length'  
summary(flq)
```

---

quantTotals

*Methods quantTotals*

---

**Description**

Methods to compute totals over selected dimensions of FLQuant objects. These methods return an object of same dimensions as the input but with the sums along the first (yearTotals) or second dimension (quantTotals). Although the names might appear contradictory, it must be noted that what each method really returns are the totals over the selected dimension.

**Usage**

```
quantTotals(x, ...)
```

**Generic function**

```
quantTotals(x)
```

```
yearTotals(x)
```

**Author(s)**

The FLR Team

**See Also**

[FLQuant](#)

**Examples**

```
flq <- FLQuant(rlnorm(100), dim=c(10,10))
quantTotals(flq)
# See how the values obtained by yearSums are being replicated
yearSums(flq)
# Get the proportions by quant
flq / quantTotals(flq)
# or year
flq / yearTotals(flq)
```

---

readVPAIntercatch	<i>Reads a single file with one year of data in VPA format as output by ICES Intercatch</i>
-------------------	---

---

**Description**

Reads a single file with one year of data in VPA format as output by ICES Intercatch

**Usage**

```
readVPAIntercatch(file)
```

**Arguments**

file                    Intercatch VPA file to load

**Value**

An object of class FLQuant.

---

residuals-FLQuant      *residuals*


---

**Description**

residuals

**Usage**

```
## S4 method for signature 'FLQuant'
residuals(object, fit, type = "log", ...)
```

**Examples**

```
data(ple4)
fit <- rlnorm(1, log(catch(ple4)), 0.1)
residuals(catch(ple4), fit)
residuals(catch(ple4), fit, type="student")
rrow(catch(ple4), fit)
rlogstandard(catch(ple4), fit)
rstandard(catch(ple4), fit)
rstudent(catch(ple4), fit)
```

---

rnoise,numeric,FLQuant-method

*Random noise with different frequencies*


---

**Description**

A noise generator

**Usage**

```
## S4 method for signature 'numeric,FLQuant'
rnoise(
  n = n,
  len = len,
  sd = 1,
  b = 0,
  burn = 0,
  trunc = 0,
  what = c("year", "cohort", "age"),
  seed = NA
)

## S4 method for signature 'numeric,missing'
```

```

rnoise(n = n, sd = 1, b = 0, burn = 0, trunc = 0, seed = NA)

## S4 method for signature 'numeric,FLQuant'
rInoise(
  n = n,
  len = len,
  sd = 1,
  b = 0,
  burn = 0,
  trunc = 0,
  what = c("year", "cohort", "age"),
  seed = NA
)

```

### Arguments

n	number of iterations
len	an FLQuant
sd	standard error for simulated series
b	autocorrelation parameter a real number in 0,1
burn	gets rid of 1st values i series
trunc	get rid of values > abs(trunc)
what	returns time series for year, cohort or age"
...	any

### Value

A FLQuant with autocorrelation equal to B.

### References

Ranta and Kaitala 2001 Proc. R. Soc.  $v_t = b * v_{t-1} + s * \sqrt{1 - b^2}$  s is a normally distributed random variable with mean = 0 b is the autocorrelation parameter

### Examples

```

## Not run:
flq <- FLQuant(1:100, quant="age")
white <- rnoise(100, flq, sd=.3, b=0)
plot(white)
acf(white)

red <- rnoise(100, flq, sd=.3, b=0.7)
plot(red)
acf(red)

res <- rnoise(100, flq, sd=.3, b=0)

```



```

ggplot() +
  geom_point(aes(year, age, size=data),
    data=subset(as.data.frame(res), data>0)) +
  geom_point(aes(year, age, size=-data),
    data=subset(as.data.frame(res), data<=0), colour="red")+
  scale_size_area(max_size=4, guide="none")+
  facet_wrap(~iter)

data(ple4)
res <- rnoise(4,m(ple4),burn=10,b=0.9,what="cohort")
ggplot()+
  geom_point(aes(year, age, size= data),
    data=subset(as.data.frame(res), data>0))+
  geom_point(aes(year, age, size=-data),
    data=subset(as.data.frame(res), data<=0), colour="red")+
  scale_size_area(max_size=4, guide="none")+
  facet_wrap(~iter)

## End(Not run)

```

---

roc

---

*Receiver Operating Characteristic (ROC)*


---

## Description

A receiver operating characteristic (ROC) curve shows the ability of a binary classifier. Here it is applied to compare two sets of values, stored as two FLQuant objects. The first is the result of applying a logical comparison of a given state against a reference value, so it contains a binary (0, 1) label. The second, the score, contains an alternative metric that attempts to measure the absolute value of the first. The examples below compare an observation of stock status, SSB being less than a reference point, and an alternative metric, here the catch curve estimates of total mortality.

## Usage

```
roc(label, ind, direction = c(">=", "<="))
```

```
auc(x = NULL, TPR = x$TPR, FPR = x$FPR)
```

## Examples

```

data(ple4)
# OM 'reality' on stock status (fbar)
state <- fbar(ple4)[, ac(1960:2017)]
# Model estimates of F using catch curves
ind <- acc(catch.n(ple4)[, ac(1960:2017)])
# Compute TSS, returns data.frame
roc(state >= 0.22, ind)
# Needs ggplotFL

```

```
## Not run:
ggplot(roc(state >= 0.22, ind, direction='>='), aes(x=FPR, y=TPR)) +
  geom_line() +
  geom_abline(slope=1, intercept=0, colour="red", linetype=2)

## End(Not run)
# Computes auc using the output of roc()
with(roc(state >= 0.22, ind), auc(TPR=TPR, FPR=FPR))
auc(roc(state >= 0.22, ind))
```

---

ruleset

*Set of verify rules for an FLR class*


---

## Description

Returns a set of standard rules to be used by the verify method for an object of a given class.

## Usage

```
ruleset(object, ...)
```

```
## S4 method for signature 'FLStock'
ruleset(object, ...)
```

## Arguments

object	An object of any FLR class for which the method has been defined.
...	Names of positions in the standard list to subset.

## Details

A standard minimal set of rules to check FLStock objects against using the verify method. The included rules are (with names in italics) check that:

- there are no NAs in any slot, *anyna*.
- *catch.wt*, *landings.wt*, *discards.wt* and *stock.wt* > 0.
- *mat*, *m.spwn* and *harvest.spwn* values are between 0 and 1.
- *harvest* >= 0.
- *cohorts* in the stock.n slot contain decreasing numbers, except for the plusgroup age.

## Value

A named list containing the rules defined for for the class object belongs to.

## Author(s)

The FLR Team

**Examples**

```
data(ple4)
ruleset(ple4)
# Extract single rule by name
ruleset(ple4, 'anyna')
```

---

runstest	<i>Computes Runs Test p-values</i>
----------	------------------------------------

---

**Description**

Computes Runs Test p-values

**Usage**

```
runstest(fit, obs, ...)

## S4 method for signature 'FLQuants,missing'
runstest(fit, combine = TRUE)

## S4 method for signature 'FLQuants,FLQuants'
runstest(fit, obs, combine = TRUE)

## S4 method for signature 'FLQuant,FLQuant'
runstest(fit, obs, combine = TRUE)

## S4 method for signature 'FLQuant,missing'
runstest(fit, combine = TRUE)

## S4 method for signature 'numeric,numeric'
runstest(fit, obs, combine = TRUE)

## S4 method for signature 'numeric,missing'
runstest(fit, obs, combine = TRUE)
```

**Arguments**

fit	The result of a model fit.
obs	The observations used in the fit.
...	Extra arguments.
combine	Should ages be combined by addition, defaults to TRUE.

**Value**

A list with elements 'p.values' and 'pass'.

**Examples**

```

data(nsher)
# Compute 'runstest' for FLSR fit
runstest(fit=fitted(nsher), obs=rec(nsher))
# Example runstest by age
data(ple4)
runstest(catch.n(ple4), landings.n(ple4), combine=FALSE)
runstest(fit=FLQuants(D=residuals(catch(ple4), discards(ple4)),
  L=residuals(catch(ple4), landings(ple4))))
runstest(fit=residuals(fitted(nsher), rec(nsher)))
runstest(FLQuants(residuals(fitted(nsher), rec(nsher))))
# Returns value per iter
runstest(fit=rnorm(25, residuals(fitted(nsher), rec(nsher)), 0.2))
runstest(ssb(nsher))
runstest(rnorm(1, FLQuant(1, dimnames=list(year=1973:2021))))
runstest(rep(0.1, 10), cumsum(rnorm(10, 0.1, 0.01)))
runstest(rnorm(10, 0, 0.1))

```

rwalk

*Generate a random walk time series from a starting point***Description**

The last year of an FLQuant object is used as a starting point to generate a time series following a random walk with drift:

$$z_t = z_{t-1} + \epsilon_t + \delta_t, t = 1, 2, \dots$$

where  $\epsilon$  is  $\mathcal{N}(0, \sigma)$

**Usage**

```
rwalk(x0, end = 1, sd = 0.05, delta = 0)
```

**Arguments**

x0	The initial state of the random walk, 'FLQuant'.
end	The number of years or the final year of the series. numeric.
sd	The standard deviation of the random walk, numeric.
delta	The drift of the random walk.

**Details**

The length of the series is set by argument *end*. This is taken as a number of years, if its value is smaller than the final 'year' of 'x0', or as a final year if larger or of class 'character'.

**Value**

An 'FLQuant' object.

**Author(s)**

Iago Mosqueira, WMR (2023)

**See Also**

[FLQuant rnorm](#)

**Examples**

```
data(ple4)
# Generate random walk recruitment with positive drift
rwalk(rec(ple4), end=5, sd=0.08, delta=0.05)
# Use append() to add the new values at the end
append(rec(ple4), rwalk(rec(ple4), end=10, sd=0.04, delta=0))
# Use end as number of years
rwalk(rec(ple4), end=5)
# or as final year
rwalk(rec(ple4), end=2020)
```

---

show

*Method show*

---

**Description**

Standard complete display of an object contents in an interactive session. Objects of class [FLArray](#) with length > 1 along sixth dimension (*iter*) are output in a summarised form, as `median(mad)`, where `mad` is the median absolute deviation. See [mad](#).

**Usage**

```
## S4 method for signature 'FLArray'
show(object)
```

```
## S4 method for signature 'FLArray'
print(x)
```

**Details**

The same format is used for objects of class [FLPar](#) with length > 1 on the last dimension (*iter*).

**Generic function**

```
show(object)
```

```
show(object)
```

**Author(s)**

The FLR Team

**See Also**[FLComp](#)[FLComp](#)**Examples**

```

# no 'iter'
flq <- FLQuant(rnorm(80), dim=c(4,20), quant='age', units='kg')
flq

# with 'iter'
flq <- FLQuant(rnorm(800), dim=c(4,20,1,1,1,10), quant='age', units='kg')
flq

# no 'iter'
flq <- FLQuant(rnorm(80), dim=c(4,20), quant='age', units='kg')
print(flq)

# with 'iter'
flq <- FLQuant(rnorm(800), dim=c(4,20,1,1,1,10), quant='age', units='kg')
print(flq)

```

simplify

*Aggregate or select along unwanted dimensions***Description**

Objects of many FLR classes might be aggregated along the "unit", "season", and/or "area" dimensions according to the type of data they contain.

**Usage**

```

simplify(object, ...)

## S4 method for signature 'FLStock'
simplify(
  object,
  dims = c("unit", "season", "area")[dim(object)[3:5] > 1],
  spwn.season = 1,
  rec.season = spwn.season,
  harvest = TRUE,
  weighted = FALSE
)

```

**Arguments**

**object** A complex **FLR** object to aggregate.

**Value**

An object of the same class as the input.

**Author(s)**

The FLR Team

---

slim

*Drop unnecessary 'iters'*

---

**Description**

Objects of FLR classes can vary in the length along the sixth dimension in any slot of class [FLQuant](#). This reduces object size and memory usage. If an object has been extended fully, for example by using [propagate](#), we can slim down the object by reducing any slot where all iters are identical and keeping only the first *iter*.

**Usage**

```
slim(object, ...)
```

```
## S4 method for signature 'FLComp'  
slim(object, ...)
```

**Arguments**

object            A complex **FLR** object to slim down.

**Details**

The test for whether a slot can be slimmed is based on checking if the sum of the variance along the 6th dimension is equal to zero.

**Value**

An object of the same class as the input.

**Author(s)**

The FLR Team

**See Also**

[FLQuant propagate](#)

**Examples**

```

data(ple4)
# Extend all of ple4 to 50 iters
ple4 <- propagate(ple4, 50)
# Add variability in catch.n
catch.n(ple4) <- rlnoise(50, log(catch.n(ple4)), log(catch.n(ple4))/10)
summary(ple4)
# slim object by dropping identical iters
spl4 <- slim(ple4)
summary(spl4)

```

---

split-methods

*splits x along the iter dimension into the groups defined by f.*


---

**Description**

Similar to `base::split`, but working along the 6th, *iter*, dimension of any singular FLR object. The object is divided into as many objects as unique values in *f*, and returned as an FLR-derived object, e.g. an FLQuants object when applied to an FLQuant.

**Usage**

```

## S4 method for signature 'FLQuant,vector'
split(x, f)

## S4 method for signature 'FLComp,vector'
split(x, f)

```

**Arguments**

<code>x</code>	The object to be split.
<code>f</code>	The vector of group names.

**Value**

An object of the corresponding plural class (FLQuants from FLQuant).

**Author(s)**

Iago Mosqueira (WMR).

**Examples**

```

# FROM FLQuant to FLQuants
flq <- rlnorm(20, FLQuant(seq(0.1, 0.8, length=10)), 0.2)
split(flq, c(rep(1, 5), rep(2,15)))

```



---

splom	<i>Method splom</i>
-------	---------------------

---

### Description

Draws a conditional scatter plot matrix.

### Usage

```
## S4 method for signature 'FLPar,missing'  
splom(x, data, ...)
```

### Details

See the help page in [lattice](#) for a full description of each plot and all possible arguments.

### Generic function

```
splom(x,data)
```

### Author(s)

The FLR Team

### See Also

[splom](#)

### Examples

```
flp <- FLPar(c(t(mvrnorm(500, mu=c(0, 120, 0.01, 20),  
  Sigma=matrix(.7, nrow=4, ncol=4) + diag(4) * 0.3))),  
  dimnames=list(params=c('a','b','c','d'), iter=1:500), units="NA")  
  
splom(flps)
```

---

spread	<i>A function to make available list elements inside a function or method</i>
--------	---

---

### Description

Inside a function, a call to `spread()` will attach to the function environment, `sys.frame()`, the elements in the list, or of the conversion to list of the object (e.g. named vector or `FLPar`), so that they be called by name. The function environment will be deleted once the function returns, so those variables won't make it to the environment from which the function was called, or further up in the call stack.

### Usage

```
spread(object, FORCE = FALSE)
```

### Arguments

object	A named list or vector whose elements are to be loaded into the calling environment.
FORCE	Should existing variable with matching names be redefined?

### Details

By default, `spread()` will not overwrite variables in the function environment with the same name as any list element, unless `FORCE=TRUE`

### Value

Invisibly the names of the variables loaded into the calling environment.

### Author(s)

The FLR Team

### See Also

[sys.nframe](#)

### Examples

```
# EXAMPLE function
foo <- function (params) {
  a <- spread(params)
  print(a)
  x*y
}
# x and y are accesible to the internal calculation
foo(params=list(x=3.5, y=9))
```

```
# Works with FLPar
foo(params=FLPar(x=3L, y=0.99238))

# Elements in object must be named
## Not run: foo(list(3, y=0.99238))

# If a variable is missing from the spread object, function will fail
## Not run: foo(list(x=4))
# Unless the variable is already defined in the calling environment,
# in this case <environment: R_GlobalEnv>
y <- 45
foo(params=list(x=4))
```

---

SRModels

*Stock-Recruitment models*

---

### **Description**

A range of stock-recruitment (SR) models commonly used in fisheries science are provided in FLCore.

### **Usage**

```
ricker()
bevholt()
bevholtDa()
bevholtss3()
segreg()
segregDa()
geomean()
shepherd()
cushing()
rickerSV()
bevholtSV()
shepherdSV()
```

```

bevholttAR1()
rickerAR1()
segregAR1()
rickerCa()
survRec(ssf, R0, Sfrac, beta, SF0 = ssf[, 1])
bevholtsig()
mixedsr()

```

### Arguments

rho	Autoregression
sigma2	Autoregression
obs	Observed values
hat	estimated values
steepness	Steepness.
vbiomass	Virgin biomass.
spr0	Spawners per recruit at F=0, see <a href="#">spr0</a> .
model	character vector with model name, either 'bevholt' or 'ricker'.

### Details

Each method is defined as a function returning a list with one or more elements as follows:

- `model`: Formula for the model, using the slot names *rec* and *ssb* to refer to the usual inputs
- `logl`: Function to calculate the loglikelihood of the given model when estimated through MLE (See [fmle](#))
- `initial`: Function to provide initial values for all parameters to the minimization algorithms called by [fmle](#) or [nls](#). If required, this function also has two attributes, [lower](#) and [upper](#), that give lower and upper limits for the parameter values, respectively. This is used by some of the methods defined in [optim](#), like "L-BFGS-B".

The `model<-` method for [FLModel](#) can then be called with *value* being a list as described above, the name of the function returning such a list, or the function itself. See the examples below.

Several functions to fit commonly-used SR models are available. They all use maximum likelihood to estimate the parameters through the method [logLAR1](#).

- `ricker`: Ricker stock-recruitment model fit:

$$R = aSe^{-bS}$$

*a* is related to productivity (recruits per stock unit at small stock size) and *b* to density dependence. (*a*, *b* > 0).

- bevholt: Beverton-Holt stock-recruitment model fit:

$$R = \frac{aS}{b + S}$$

$a$  is the maximum recruitment (asymptotically) and  $b$  is the stock level needed to produce the half of maximum recruitment  $\frac{a}{2}$ . ( $a, b > 0$ ).

- segreg: Segmented regression stock-recruitment model fit:

$$R = \text{ifelse}(S \leq b, aS, ab)$$

$a$  is the slope of the recruitment for stock levels below  $b$ , and  $ab$  is the mean recruitment for stock levels above  $b$ . ( $a, b > 0$ ).

- geomean: Constant recruitment model fit, equal to the historical geometric mean recruitment.

$$(R_1 R_2 \dots R_n)^{1/n} = e^{\text{mean}(\log(R_1), \dots, \log(R_n))}$$

- shepherd: Shepherd stock-recruitment model fit:

$$R = \frac{aS}{1 + (\frac{S}{b})^c}$$

$a$  represents density-independent survival (similar to  $a$  in the Ricker stock-recruit model),  $b$  the stock size above which density-dependent processes predominate over density-independent ones (also referred to as the threshold stock size), and  $c$  the degree of compensation.

- cushioning: Cushing stock-recruitment model fit:

$$R = aS e^b$$

This model has been used less often, and is limited by the fact that it is unbounded for  $b \geq 1$  as  $S$  increases. ( $a, b > 0$ ).

Stock recruitment models parameterized for steepness and virgin biomass:

- rickerSV: Fits a ricker stock-recruitment model parameterized for steepness and virgin biomass.

$$a = e^{\frac{b \cdot vbiomass}{spr0}}$$

$$b = \frac{\log(5 \cdot steepness)}{0.8 \cdot vbiomass}$$

- bevholtSV: Fits a Beverton-Holt stock-recruitment model parameterised for steepness and virgin biomass.

$$a = \frac{4 \cdot vbiomass \cdot steepness}{(spr0 \cdot (5 \cdot steepness - 1.0))}$$

$$b = \frac{vbiomass(1.0 - steepness)}{5 \cdot steepness - 1.0}$$

- sheperdSV: Fits a shepher stock-recruitment model parameterized for steepness and virgin biomass.

$$a = \frac{1.0 + (\frac{vbiomass}{b})^c}{spr0}$$

$$b = vbiomass \left( \frac{0.2 - steepness}{steepness(0.2)^c - 0.2} \right) \left( \frac{-1.0}{c} \right)$$

Models fitted using autoregressive residuals of first order:

- bevholtAR1, rickerAR1, segregAR1: Beverton-Holt, Ricker and segmented regression stock-recruitment models with autoregressive normal log residuals of first order. In the model fit, the corresponding stock-recruit model is combined with an autoregressive normal log likelihood of first order for the residuals. If  $R_t$  is the observed recruitment and  $\hat{R}_t$  is the predicted recruitment, an autoregressive model of first order is fitted to the log-residuals,  $x_t = \log(\frac{R_t}{\hat{R}_t})$ .

$$x_t = \rho x_{t-1} + e$$

where  $e$  follows a normal distribution with mean 0:  $e \sim N(0, \sigma_{AR}^2)$ .

Ricker model with one covariate. The covariate can be used, for example, to account for an environmental factor that influences the recruitment dynamics. In the equations,  $c$  is the shape parameter and  $X$  is the covariate.

- rickerCa: Ricker stock-recruitment model with one multiplicative covariate.

$$R = a(1 - cX)S e^{-bS}$$

### Author(s)

The FLR Team

### References

- Beverton, R.J.H. and Holt, S.J. (1957) On the dynamics of exploited fish populations. MAFF Fish. Invest., Ser: II 19, 533.
- Needle, C.L. Recruitment models: diagnosis and prognosis. Reviews in Fish Biology and Fisheries 11: 95-111, 2002.
- Ricker, W.E. (1954) Stock and recruitment. J. Fish. Res. Bd Can. 11, 559-623.
- Shepherd, J.G. (1982) A versatile new stock-recruitment relationship for fisheries and the construction of sustainable yield curves. J. Cons. Int. Explor. Mer 40, 67-75.

### See Also

[FLSR](#), [FLModel](#)

**Examples**

```
# inspect the output of one of the model functions
bevholt()
names(bevholt())
bevholt()$logl

# once an FLSR model is in the workspace ...
data(nsher)

# the three model-definition slots can be modified
# at once by calling 'model<-' with
# (1) a list
model(nsher) <- bevholt()

# (2) the name of the function returning this list
model(nsher) <- 'bevholt'

# or (3) the function itself that returns this list
model(nsher) <- bevholt
```

---

ssb

*Calculate or return the Spawning Stock Biomass*


---

**Description**

The calculated Spawning Stock Biomass (SSB) of a fish population is returned by this method. SSB is the combined weight of all individuals in a fish stock that are capable of reproducing. In some classes this is calculated from information stored in different slots, while in others `ssb()` is simply an slot accessor. When the later is the case, the corresponding replacement method also exists.

**Usage**

```
ssb(object, ...)
```

## S4 method for signature 'FLBiol'

```
ssb(object, ...)
```

**Arguments**

`object`            Object on which `ssb` is calculated or extracted.

**Details**

Objects of the *FLBiol* class do not contain any information on catch or fishing mortality, so a call to `ssb()` will only correct abundances for natural mortality to the moment of spawning. The method can also take information on catches or fishing mortality and use them when calculating abundances at spawning time. An *FLQuant* named either 'catch.n', 'f', 'hr' or 'harvest' can be used. The first

three are self-explanatory, while for the last units must be either 'f' or 'hr'. The quantities should refer to total yearly values, as the value in the 'spwn' slot will be used to calculate what fraction of fishing mortality to apply.

### Value

An object, generally of class [FLQuant](#).

### Author(s)

The FLR Team

### See Also

[FLComp](#)

### Examples

```
data(ple4)
biol <- as(ple4, "FLBiol")
# SSB from FLBiol, abundances corrected only for M
ssb(biol)
# Provide catch-at-age, F or HR to correct N
ssb(biol, catch.n=catch.n(ple4))
ssb(biol, f=harvest(ple4))
ssb(biol, harvest=harvest(ple4))
ssb(biol, hr=catch.n(ple4) / stock.n(ple4))
```

---

ssb\_next

*Calculate next yera's SSB from survivors and Fbar*

---

### Description

The spawning stock biomass (SSB) of the stock gets calculated from the survivors of the previous year. This provides a value for the first year after the end of the object. Weights-at-age, maturity in this extra year are calculated as averages over the last *wts.nyears*.

### Usage

```
ssb_next(x, fbar = 0, wts.nyears = 3, fbar.nyears = 3)
```

### Arguments

x	An FLStock object containing estimates of abundance and harvesting.
fbar	The Fbar rate assumed on the extra year. Defaults to 0.
wts.nyears	Number of years in calculation of mean weight-at-age and maturity for the extra year.
fbar.nyears	Number of years in calculation of mean selectivity, natural mortality and fraction of F abnd M before spawning for the extra year.



**Details**

For stocks spawning later in the year, a value for the average fishing mortality,  $fbar$ , expected in that year can be provided. Mortality until spawning is then calculated, with  $M$  and selectivity assumed in the extra year to be an average of the last  $fbar.nyears$ .

**Value**

An FLQuant.

**Examples**

```
data(ple4)
ssb_next(ple4)
# Compare with ssb()
ssb(ple4)[, ac(2014:2017)] / ssb_next(ple4)[, ac(2014:2017)]
```

---

standardUnits

*Standard units of measurement for a complex class object*

---

**Description**

Returns values for the *units* of each *FLQuant* slot according to the standard adopted by the FLR Team for the supplied class.

**Usage**

```
standardUnits(object, ...)

## S4 method for signature 'character'
standardUnits(object, ...)

## S4 method for signature 'FLS'
standardUnits(object, ...)

## S4 method for signature 'FLBiol'
standardUnits(object, ...)
```

**Arguments**

object                    for which the standard *units* are to be returned

**Details**

For objects derived from class *FLS*, which currently includes *FLStock* and *FLStockLen*, the adopted standard includes: 'kg' for individual weights, '1000' for number of individuals, 't' for biomass, 'f' for harvest, 'm' for natural mortality, and an empty string for proportions (spwn, mat).

For objects derived of class *FLBiol* the adopted standard units are: 'kg' for individual weights, '1000' for number of individuals, 'm' for natural mortality, and an empty string for proportions (spwn, mat).

**Value**

A list with the corresponding *units* value for each slot

**Author(s)**

The FLR Team

**See Also**

[units-FLCore](#)

**Examples**

```
stk <- FLStock(catch=FLQuant(runif(20, 2, 120)))
# FLStock object has no units
summary(stk)
# Obtain standard units for the class as a list
standardUnits(stk)
# which can then be assigned to the object
units(stk) <- standardUnits(stk)
summary(stk)
# units<- method also accepts a function to be called to provide units
units(stk) <- standardUnits
bio <- FLBiol(n=FLQuant(runif(50, 2, 120), dim=c(5, 10)))
# Object has no units
summary(bio)
# Obtain standard units for the class as a list
standardUnits(bio)
# which can then be assigned to the object
units(bio) <- standardUnits(bio)
summary(stk)
```

---

summary,FLArray-method

*Method summary*

---

**Description**

Outputs a general summary of the structure and content of an fwdControl object. The method invisibly returns the data.frame shown on screen.

**Usage**

```
## S4 method for signature 'FLArray'
summary(object, .class = TRUE, ...)

## S4 method for signature 'FLQuantPoint'
summary(object, ...)
```

```
## S4 method for signature 'FLPar'  
summary(object, title = TRUE, ...)  
  
## S4 method for signature 'FLComp'  
summary(object, ...)  
  
## S4 method for signature 'FLQuants'  
summary(object)  
  
## S4 method for signature 'predictModel'  
summary(object)  
  
## S4 method for signature 'FLBiol'  
summary(object)  
  
## S4 method for signature 'FLModel'  
summary(object, ...)  
  
## S4 method for signature 'FLlst'  
summary(object)
```

### Generic function

```
summary(object)
```

### Author(s)

The FLR Team

### See Also

[summary](#)

### Examples

```
flq <- FLQuant(rlnorm(90), dim=c(3,10), units='kg')  
summary(flq)  
  
data(ple4)  
summary(ple4)  
  
data(nsher)  
summary(nsher)
```

---

 survey

*A method to generate observations of abundance at age.*


---

### Description

A method to generate observations of abundance at age.

### Usage

```

survey(object, index, ...)

## S4 method for signature 'FLStock,FLIndex'
survey(
  object,
  index,
  sel = sel.pattern(index),
  ages = dimnames(index)$age,
  timing = mean(range(index, c("startf", "endf"))),
  index.q = index@index.q,
  stability = 1
)

## S4 method for signature 'FLStock,FLIndexBiomass'
survey(
  object,
  index,
  sel = sel.pattern(index),
  ages = ac(seq(range(index, c("min")), range(index, c("max")))),
  timing = mean(range(index, c("startf", "endf"))),
  catch.wt = index@catch.wt,
  index.q = index@index.q,
  stability = 1
)

## S4 method for signature 'FLStock,missing'
survey(
  object,
  sel = catch.sel(object),
  ages = dimnames(sel)$age,
  timing = 0.5,
  index.q = 1,
  biomass = FALSE,
  stability = 1
)

## S4 method for signature 'FLStock,FLIndices'
survey(object, index, ...)

```

**Arguments**

object            The object on which to draw the observation

**Value**

An FLQuant for the index of abundance

**Author(s)**

The FLR Team

**See Also**

[FLComp](#)

**Examples**

```
data(ple4)
data(ple4.index)
# CONSTRUCT a survey from stock and index
survey(ple4, ple4.index)
# Create FLIndexBiomass
ple4.biom <- as(ple4.index, "FLIndexBiomass")
survey(ple4, ple4.biom)
data(ple4)
survey(ple4)
survey(ple4, biomass=TRUE)
```

---

survivors

*Calculate the survivors of a stock to the next year.*

---

**Description**

An FLStock object containing estimates of abundance at age ('stock.n') and harvest level at age ('harvest'), is used to bring forward the population by applying the total mortality at age ('z'). No calculation is made on recruitment, so abundances for the first age will be set as 'NA', unless a value is provided.

**Usage**

```
survivors(object, rec = NA)
```

**Arguments**

object            An FLStock with estimated harvest and abundances  
 rec                Value for recruitment, first age abundance, 'numeric' or 'FLQuant'.

**Value**

The abundances at age of the survivors, 'FLQuant'.

**Examples**

```
data(ple4)
stock.n(ple4[, ac(2002:2006)])
survivors(ple4[, ac(2002:2006)])
```

---

sweep,FLArray-method    *Method sweep for FLCore classes*

---

**Description**

Use R's sweep method on FLCore classes

**Usage**

```
## S4 method for signature 'FLArray'
sweep(x, MARGIN, STATS, FUN = "-", check.margin = TRUE, ...)

## S4 method for signature 'FLPar'
sweep(x, MARGIN, STATS, FUN = "-", check.margin = TRUE, ...)
```

**Details**

These methods call base R [sweep](#) method on **FLCore** classes and then ensure that the returned object is of same class.

**Generic function**

```
sweep(x, MARGIN, STATS, FUN = "-", check.margin = TRUE, ...)
```

**Author(s)**

The FLR Team

**See Also**

[sweep](#)

**Examples**

```
flq <- FLQuant(rlnorm(90), dim=c(3,10), units='kg')
# Get ratio of max value by year
sweep(flq, 2, apply(flq, 2, max), "/")
```

---

tail,FLQuant-method    *Returns the first and last parts of an FLQuant.*

---

### Description

Standard tail and head methods can be applied along any dimension of an FLQuant object.

### Usage

```
## S4 method for signature 'FLQuant'  
tail(x, n = 1, dim = 2, ...)  
  
## S4 method for signature 'FLQuant'  
head(x, n = 1, dim = 2, ...)
```

### Arguments

x	The object to extract from, FLQuant.
n	The number of elements to extract, numeric.
dim	Dimension to extract from, defaults to 2, 'year'.

### Value

An FLQuant with the extracted elements.

### Author(s)

Iago Mosqueira (WMR)

### See Also

[base::tail](#)

### Examples

```
x <- FLQuant(1:10)  
  
# Extract the last 3 years  
tail(x, 3)  
  
# Extract all but the first 3 years  
tail(x, -3)  
  
# Extract the first 3 years  
head(x, 3)  
  
# Extract all but the last 3 years  
head(x, -3)
```

---

trim	<i>Method trim</i>
------	--------------------

---

### Description

Trim FLR objects using named dimensions

### Usage

```
trim(x, ...)  
  
## S4 method for signature 'FLArray'  
trim(x, ...)  
  
## S4 method for signature 'FLComp'  
trim(x, ...)  
  
## S4 method for signature 'FLS'  
trim(x, ...)  
  
## S4 method for signature 'FLBiol1'  
trim(x, ...)
```

### Details

Subsetting of FLR objects can be carried out with dimension names by using `trim`. A number of dimension names and selected dimensions are passed to the method and those are used to subset the input object.

Exceptions are made for those classes where certain slots might differ in one or more dimensions. If `trim` is applied to an `FLQuant` object of length 1 in its first dimension and with dimension name equal to 'all', values to `trim` specified for that dimension will be ignored. For example, `FLStock` objects contain slots with length=1 in their first dimension. Specifying values to trim over the first dimension will have no effect on those slots (catch, landings, discards, and stock). Calculations might need to be carried out to recalculate those slots (e.g. using `computeCatch`, `computeLandings`, `computeDiscards` and `computeStock`) if their quant-structured counterparts are modified along the first dimension.

### Generic function

```
trim(x)
```

### Author(s)

The FLR Team

### See Also

[FLQuant](#), [FLStock](#), [FLCohort](#), [FLIndex](#)



**Examples**

```

flq <- FLQuant(rnorm(90), dimnames=list(age=1:10, year=2000:2016))

trim(flq, year=2000:2005)
# which is equivalent to
window(flq, start=2000, end=2005)

trim(flq, year=2000:2005, age=1:2)

# Now on an FLStock
data(ple4)
summary(trim(ple4, year=1990:1995))

# If 'age' is trimmed in ple4, catch, landings and discards need to be
# recalculated
shp14 <- trim(ple4, age=1:4)
landings(shp14) <- computeLandings(shp14)
discards(shp14) <- computeDiscards(shp14)
catch(shp14) <- computeCatch(shp14)
summary(shp14)

```

---

units-FLCore

*Method units*


---

**Description**

units attribute for FLQuant and FLArray-derived objects

**Usage**

```

## S4 method for signature 'FLArray'
units(x)

## S4 replacement method for signature 'FLArray,character'
units(x) <- value

setunits(x, value)

## S4 method for signature 'FLPar'
units(x)

## S4 replacement method for signature 'FLPar,character'
units(x) <- value

## S4 method for signature 'FLComp'
units(x)

```

```
## S4 replacement method for signature 'FLComp,list'
units(x) <- value

## S4 replacement method for signature 'FLComp,character'
units(x) <- value

## S4 replacement method for signature 'FLComp,function'
units(x) <- value
```

## Details

Objects of FLArray-based classes (e.g. [FLQuant](#)) contain a `units` attribute of class `character`. This should be used to store the corresponding units of measurement. This attribute can be directly accessed and modified using the `units` and `units<-` methods.

For complex objects, `units` will return a named list containing the attributes of all `FLQuant` slots. `units` of a complex object can be modified for all slots or a subset of them, by passing a named list with the new values. See examples below.

The complete set of *units* for a complex object can be obtained as a named *list*.

Assignment of *units* to the *FLQuant* slots of a complex object can be carried out passing a named *list* or *character* vector containing the units for the slots to be modified.

## Generic function

```
units(x)
units<-(x,value)
```

## Author(s)

The FLR Team

## See Also

[FLQuant](#), [FLPar](#), [FLCohort](#)

## Examples

```
flq <- FLQuant(rnorm(100), dim=c(5,20), units='kg')
units(flq)
units(flq) <- 't'
summary(flq)

# units for a complex object
data(ple4)
units(ple4)
units(ple4) <- list(harvest='hr')

data(ple4)
units(ple4) <- list(harvest="hr")
units(ple4) <- c(harvest="hr")
```

---

uom *uom Units of Measurement*

---

### Description

The 'units' attribute of FLQuant objects provides a mechanism for keeping track of the units of measurement of that particular piece of data.

### Usage

```
uom(op, u1, u2)
```

```
uomUnits(unit = missing)
```

### Arguments

op	The arithmetic operator to be used, one of '+', '-', '*' or '/'
u1	The units of measurement string of the first object
u2	The units of measurement string of the second object
unit	A character vector for one or more units to be compared with those known to uom.

### Details

Arithmetic operators for 'FLQuant' objects are aware of a limited set of units of measurement and will output the right unit when two object are arithmetically combined. For example, the product of object with units of 'kg' and '1000' will output an object with 'units' of 't' (for metric tonnes).

Operations involving combinations of units not defined will issue a warning, and the resulting 'units' attribute will simply keep a string indicating the input units of measurement and the operation carried out, as in '10 \* 1000'.

Note that no scaling or modification of the values in the object takes place.

Conversion across units is carried out by the uom() function

The list of units known to uom is stored internally but can be queried by calling uomUnits() with no arguments. If a character vector is provided, a logical is returned telling whether the string is included or not in that table.

### Value

uom returns a string with the corresponding units of measurement, or a character vector, showing the operation carried out, when units are not known to uom or not compatible, e.g. "100 \* d".

uomUnits returns TRUE or FALSE if unit is given, otherwise a character vector with all units known to uom.

### Author(s)

The FLR Team

**See Also**

[FLQuant units, FLArray-method](#)

**Examples**

```
# Conversion between weights
FLQuant(1, units='kg') * FLQuant(1000, units='l')

# Conversion between mortalities
FLQuant(0.2, units='m') + FLQuant(0.34, units='f')

# Check if units are known
uomUnits('kg')
uomUnits('kell')
```

---

uomTable

---

*Table for conversions and operations between units of measurement*


---

**Description**

- uom defaults to NA unless defined below.
- unit +/- itself, returns the same unit (e.g. kg + kg = kg)
- numeric unit \* 1 returns same unit (e.g. 1e4 \* 1 = 1e4)
- numeric unit \* numeric unit returns product (e.g. 10 \* 100 = 1000)
- unit / unit returns "" (e.g. 100 / 100 = "")
- numeric unit / smaller numeric unit returns division (e.g. 100 / 10 = 10)
- 100 times kg returns t
- numeric unit \* 'kg' returns the product in tonnes (e.g. kg \* 1e4 = t \* 10)
- units with divisions are parsed (e.g. days/boat \* boat = days)
- 
- 

**Format**

An object of class array

---

`upperlower`*Extract and modify the lower and upper FLModel attributes.*

---

**Description**

Description: Lorem ipsum dolor sit amet, consectetur adipiscing elit.

**Usage**

```
lower(object, ...)
```

```
## S4 method for signature 'FLModel'  
lower(object)
```

```
## S4 method for signature 'FLModel'  
upper(object)
```

**Arguments**

<code>object</code>	Object to extract from or modify
<code>...</code>	Other arguments
<code>value</code>	New value

**Details**

Details: Aliquam sagittis feugiat felis eget consequat.

**Value**

RETURN Lorem ipsum dolor sit amet

**Author(s)**

The FLR Team

**See Also**

[FLModel](#)

---

verify	Verify FLR objects
--------	--------------------

---

### Description

Verifies the content of FLR objects according to a set of rules

### Usage

```
verify(object, ...)
```

```
## S4 method for signature 'FLComp'
verify(object, ..., report = TRUE)
```

```
## S4 method for signature 'FLStock'
verify(object, rules = ruleset(object), ..., report = TRUE)
```

### Arguments

object	An object of any FLR class for which the method has been defined
...	Additional rules to be tested, as a formula or list. See details
report	Should the standard data.frame report be output (if TRUE) or a single logical value for all tests?
rules	Basic set of rules for a given class, as returned by ruleset().

### Details

Classes' validity functions generally check the structure and dimensions of objects and their component slots. But some checks on the data content of objects is often required. The various verify methods implement both a system to create *rules* that an object is expected to pass, and a minimum standard set of rules for each defined class

The data.frame output by the method when report=TRUE contains one row per rule and the following columns:

- *name*, the rule name
- *items*, number of comparisons carried out
- *passes*, number of *TRUE* values
- *fails*, number of *FALSE* values
- *NAs*, number of logical NAs
- *valid*, are all values *TRUE*?
- *rule*, the expression being evaluated

Additional rules can be specify in a call to *verify*, in one of two forms. Simple rules can be defined as a formula involving methods defined for the class. A rule such as `highm = ~ m < 2` will check if values in the *m* slot are higher than 2 and return a logical *FLQuant*.

Some rules cannot simply use existing methods or functions, for example those operating on all slots of the object, or requiring additional computations. In this case, the argument to *verify* can be a list, with an element named *rule* of class *formula* and where *test* is defined. The test then calls for a new function, defined as another element of the list, and which will be used by *verify* when evaluating the set of rules. See below for examples.

A set of rules has been defined for the *FLStock* class, available by calling the *ruleset* method. The *verify* method for *FLStock* will by default evaluate those rules, as well as any other defined in the call.

### Value

A data.frame with the results of applying those rules, or a single logical value, if `report=FALSE`

### Author(s)

The FLR Team

### See Also

[ruleset](#)

### Examples

```
# Verifying a new rule for an FLSR object
data(nsher)
# rule: are all recruitment values greater than 0?
verify(nsher, rec=~rec > 0)

# Define rule calling its own function
data(ple4)
# rule: ssb is less
verify(ple4, ssbstock = ~ssb < stock)
data(ple4)
# verify for the standard set of rules for FLStock
verify(ple4)
# verify a single rule from set
verify(ple4, rules=ruleset(ple4, 'anyina'), report=FALSE)

# add own rule to set
verify(ple4, m = ~m >=0)
```

---

vonbert

*Growth models*

---

### Description

Growth models

ivonbert

gompertz

richards

### Usage

```
vonbert(linf, k, t0, age)
```

```
ivonbert(linf, k, t0, len)
```

```
gompertz(linf, a, k, age)
```

```
richards(linf, k, b, m, age)
```

### Examples

```
data(ple4)
vonbert
```

```
vonbert(linf=35, k=0.352, t0=-0.26, age=1:14)
ivonbert(35, 0.352, -0.26, 1:34)
gompertz(linf=179.13, k=0.4088, a=1.7268, age=1:12)
richards(linf=178.63, k=0.424, b=-7.185, m=2880.4, age=1:12)
```

---

weighted.mean,FLQuants,FLQuants-method

*Weighted means along a FLQuants.*

---

### Description

Facilitates the calculation of weighted means across a FLQuants object.

### Usage

```
## S4 method for signature 'FLQuants,FLQuants'
weighted.mean(x, w)
```



**Arguments**

x                    Values to be averaged, as an object of class FLQuants.  
w                    weights to be used, as an object of class FLQuants.

**Details**

An object of class FLQuants containing elements over which an average is to be computed, is combined with another one, of the same length, containing values to be used as weights. The overall weighted mean is calculated by computing the product of each element to its corresponding weight, and dividing by the sum of all weights. NAs in the value elements are substituted for zeroes, so do not influence the mean.

**Value**

A single FLQuant object.

**Author(s)**

The FLR Team

**See Also**

[FLCore::FLQuants stats::weighted.mean](#)

**Examples**

```
data(ple4)
# Weighted mean of landings and discards weights-at-age
weighted.mean(FLQuants(L=landings.wt(ple4), D=discards.wt(ple4)),
  FLQuants(L=landings.n(ple4), D=discards.n(ple4)))
```

---

wireframe

*Method wireframe*

---

**Description**

3D plot for FLQuant objects

**Usage**

```
## S4 method for signature 'formula,FLQuant'
wireframe(x, data, ...)
```

**Arguments**

x                    a formula formula for lattice  
data                a FLQuant object with the values  
...                Additional argument list to be passed to wireframe

**Details**

Method to plot 3D representations of FLQuant objects

**Value**

a wireframe plot

**Examples**

```
data(ple4)
wireframe(data~age+year, data=harvest(ple4))
```

---

yearSample	<i>Samples along the year dimension</i>
------------	---

---

**Description**

A resample from an FLQuant object along the 'year' dimension is returned. The 'year' dimnames of the output object can be specified, although that is not needed if the resample is to be assigned in a slot.

**Usage**

```
yearSample(x, size = length(years), years, replace = TRUE, prob = NULL)
```

**Arguments**

x	An FLQuant object.
size	Number of samples (years), non-negative integer.
years	Optional vector to set as 'year' dimnames in output.
replace	should sampling be with replacement? Defaults to TRUE.
prob	a vector of probability weights.

**Value**

RETURN Description, class

**Author(s)**

Iago Mosqueira (WMR)

**See Also**

[FLQuant sample\(\)](#)

## Examples

```
data(ple4)
# Take 20 samples of recent recruitment
yearSample(rec(ple4)[, ac(2013:2017)], 20)
# Providing 'years' sets the output object dimnames
yearSample(rec(ple4)[, ac(2013:2017)], 20, year=2000:2019)
```

---

z

*Total mortality z*

---

## Description

Returns the calculation of total mortality,  $z$ , usually as the sum of fishing mortality,  $f$ , and natural mortality,  $m$ .

## Usage

```
z(object, ...)
```

## S4 method for signature 'FLS'

```
z(object, ...)
```

## Arguments

object	Object to calculate on.
...	Any extra arguments.

## Value

An object of the corresponding class, usually *FLQuant*.

## Author(s)

The FLR Team

## See Also

[FLQuant](#)

## Examples

```
data(ple4)

z(ple4)
```

---

 %+%

---

*FLQuant arithmetic operators that extend objects*


---

### Description

Arithmetic operations between two [FLQuant](#) objects using the standard operators (+, -, \*, /, ^, see [Arith](#)) need all dimensions in both objects to match. This requirement is relaxed by using the percent version of those five operators: %+%, %-%, %\*%, %/% and %^%.

### Usage

```
e1 %+% e2
```

```
x %-% y
```

```
x %^% y
```

```
## S4 method for signature 'FLQuant,FLQuant'
x %*% y
```

```
## S4 method for signature 'FLQuant,FLQuant'
e1 %/% e2
```

```
## S4 method for signature 'FLQuant,FLQuant'
e1 %+% e2
```

```
## S4 method for signature 'FLQuant,FLQuant'
x %-% y
```

```
## S4 method for signature 'FLQuant,FLQuant'
x %^% y
```

```
## S4 method for signature 'FLPar,FLQuant'
x %*% y
```

```
## S4 method for signature 'FLPar,FLQuant'
e1 %/% e2
```

```
## S4 method for signature 'FLPar,FLQuant'
e1 %+% e2
```

```
## S4 method for signature 'FLPar,FLQuant'
x %-% y
```

```
## S4 method for signature 'FLPar,FLQuant'
x %^% y
```

```
## S4 method for signature 'FLQuant,FLPar'  
x %*% y  
  
## S4 method for signature 'FLQuant,FLPar'  
e1 %/% e2  
  
## S4 method for signature 'FLQuant,FLPar'  
e1 +% e2  
  
## S4 method for signature 'FLQuant,FLPar'  
x %-% y  
  
## S4 method for signature 'FLQuant,FLPar'  
x %^% y  
  
## S4 method for signature 'FLPar,FLPar'  
x %*% y  
  
## S4 method for signature 'FLPar,FLPar'  
e1 +% e2  
  
## S4 method for signature 'FLPar,FLPar'  
x %-% y  
  
## S4 method for signature 'FLPar,FLPar'  
e1 %/% e2  
  
## S4 method for signature 'FLPar,FLPar'  
x %^% y  
  
## S4 method for signature 'FLQuants,FLPar'  
e1 / e2  
  
## S4 method for signature 'FLQuants,FLPar'  
e1 * e2  
  
## S4 method for signature 'FLQuants,FLPars'  
e1 / e2  
  
## S4 method for signature 'FLQuants,FLPars'  
e1 * e2  
  
## S4 method for signature 'FLQuants,FLQuants'  
e1 / e2  
  
## S4 method for signature 'FLQuants,FLQuants'  
e1 * e2
```

```
## S4 method for signature 'FLQuants,FLQuants'
e1 + e2
```

```
## S4 method for signature 'FLQuants,FLQuants'
e1 - e2
```

## Details

If any of the objects is of length one in a dimensions where the other is longer, the dimensions will be extended and the element-by-element operation then conducted. Dimensions and dimnames of the output will be those of the larger object. See the examples to observe their behaviour.

Please note that this behaviour is already present on the [Arith](#) methods for [FLArray](#)-derived classes but only on the 6th, `iter`, dimension.

The original use of the `%%` operator, as vector product, is not available for [FLQuant](#) objects, but can be applied to the [array](#) inside them, as in the example below.

Methods for operations between an [FLQuant](#) and an [FLPar](#) object will match dimensions by names of dimnames, regardless of position.

## Generic function

```
x %+% y, x %-% y, x %*% y, e1 %/% e2, x %^% y
```

## Author(s)

The FLR Team

## See Also

[FLQuant](#), [matmult](#)

## Examples

```
a <- FLQuant(2, dim=c(3,3,2))
b <- FLQuant(3, dim=c(3,3,1))

# This should fail
## Not run: a * b

a %*% b
a %+% b
# To use base's %% vector product, apply it to a matrix from @.Data
b@.Data[,,,] %% 1:3
# or
b[, ,drop=TRUE] %% 1:3

# FLPar vs. FLQuant works by dimnames' names
flp <- FLPar(2, dimnames=list(params='a', year=2000:2005, iter=1))
flq <- FLQuant(3, dimnames=list(year=2000:2005))
flp %*% flq
```

```

# Divide each FLQuants element by a 'param' in FLPar, e.g. time series
# divide by reference points
FLQuants(SSB=FLQuant(2303), F=FLQuant(0.8)) / FLPar(SSB=1560, F=0.4)

# Product of each FLQuants element by a 'param' in FLPar
FLQuants(SSB=FLQuant(2303), F=FLQuant(0.8)) * FLPar(SSB=1560, F=0.4)
# Divide each FLQuants element by each in FLPar
FLQuants(A=FLQuant(2303), B=FLQuant(1287)) /
  FLPar(A=FLPar(SBMSY=1560), B=FLPar(SBMSY=1000))
# Divide each FLQuants element by each in FLPar
FLQuants(A=FLQuant(2303), B=FLQuant(1287)) *
  FLPar(A=FLPar(SBMSY=1560), B=FLPar(SBMSY=1000))
# Divide each FLQuants element by each in FLPar
FLQuants(A=FLQuant(300), B=FLQuant(200)) /
  FLQuants(A=FLQuant(3), B=FLQuant(2))
# Divide each FLQuants element by each in FLPar
FLQuants(A=FLQuant(100), B=FLQuant(200)) *
  FLQuants(A=FLQuant(3), B=FLQuant(2))
# Divide each FLQuants element by each in FLPar
FLQuants(A=FLQuant(100), B=FLQuant(200)) *
  FLQuants(A=FLQuant(3), B=FLQuant(2))
# Divide each FLQuants element by each in FLPar
FLQuants(A=FLQuant(100), B=FLQuant(200)) *
  FLQuants(A=FLQuant(3), B=FLQuant(2))

```

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