

USER MANUAL FOR
FISHASSESS
0.1-1

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1 Theory and methods

2 Functions

2.1 Growth parameters

VBGF_l

Length based von Bertalanffy growth function

Usage

`VBGFl(t, Linf, K, t0)`

Where

t age

Linf asymptotic length

K growth rate

t0 age at zero length

→ Return the length

References

von Bertalanffy (1938); Sparre and Venema (1998)

VBGF_w

Weight based von Bertalanffy growth function

Usage

`VBGFw(t, Winf, K, t0, b=3)`

Where

t age

Winf asymptotic weight

K growth rate

t0 age at zero length

b allometric coefficient

→ Return the weight

References

von Bertalanffy (1938); Sparre and Venema (1998)

VBGFinv

Inverted von Bertalanffy growth function

Usage

`VBGFinv(l, Linf, K, t0)`

Where

l length

Linf asymptotic length

K growth rate

t0 age at zero length

→ Return the age

References

von Bertalanffy (1938); Sparre and Venema (1998)

NEWk

Adjust growth rate (K) according new L_∞ hold the same ϕ' (phi prime)

Usage

`NEWk(k, Linf, Linf2)`

Where

k growth rate

Linf asymptotic length

Linf2 new asymptotic length

→ Return the new value of K.

References

Munro and Pauly (1983)

phi.prime**Usage**

`phi.prime(k, Linf)`

Where

k growth rate

Linf asymptotic length

→ Return the value of phi prime.

References

Munro and Pauly (1983)

phi**Usage**

`phi(k, Winf)`

Where

k growth rate

Winf asymptotic weight

→ Return the value of phi.

References

Munro and Pauly (1983)

2.2 Mortality rates

Z.hoenig

Usage

Z.hoenig(Tmax)

Where

Tmax maximum age observed

→ Return total mortality estimate (Z)

References

Hoenig (1983)

Z.joneszalinge

Cumulative plot of Jones and van Zalinge to estimate total mortality rate (Z)

Usage

Z.joneszalinge(obj, dlower=0, dupper=0, plot.it=T, ...)

Where

obj object of the class **lengthdata**

dlower number of lower points to be deleted (close to L_∞)

dupper number of upper points to be deleted (not fully recruited)

plot.it TRUE for plot graph

... parameters to be passed to plot method

→ Return total mortality estimate (Z)

References

Sparre and Venema (1998)

M.gunderson

Natural mortality (M) estimation by gonad-somatic index (GSI)

Usage

`M.gunderson(GSI)`

Where

GSI gonad-somatic index

→ Return the estimated natural mortality.

References

Gunderson and Dygert (1988)

M.jensen1

Natural mortality (M) estimation by age at first maturity

Usage

`M.jensen1(Tmat)`

Where

Tmat age at first maturity

→ Return the estimated natural mortality.

References

Jensen (1996)

M.jensen2

Natural mortality (M) estimation by growth rate (K)

Usage

`M.jensen2(K)`

Where

K growth rate

→ Return the estimated natural mortality.

References

Jensen (1996)

M.jensen3

Natural mortality (M) estimation by growth rate (K)

Usage

M.jensen3(K)

Where

K growth rate

→ Return the estimated natural mortality.

References

Jensen (1996)

M.pauly

Natural mortality estimation by Pauly's relationship of von Bertalanffy growth parameters and mean water temperature.

Usage

M.pauly(Linf, K, Temp)

Where

Linf asymptotic length

K growth rate

Temp mean water temperature experienced by the stock

→ Return the estimated natural mortality.

References

Pauly (1980)

M.pauly.w

Natural mortality estimation by Pauly's relationship of von Bertalanffy growth parameters and mean water temperature.

Usage

`M.pauly.w(Winf, k, Temp)`

Where

Winf asymptotic weight

K growth rate

Temp mean water temperature experienced by the stock

→ Return the estimated natural mortality.

References

Pauly (1980)

M.ralston

Natural mortality estimation by Ralston's relationship for snappers and groupers.

Usage

`M.ralston(K)`

Where

K growth rate

→ Return the estimated natural mortality.

References

Ralston (1987)

M.rikhter

Natural mortality (M) estimation by age at first maturity

Usage

`M.rikhter(Tmat)`

Where

Tmat age at first maturity

→ Return the estimated natural mortality.

References

Rikhter and Efanov (1976)

2.3 Virtual populations methods

vpa.age

Performs a virtual population analysis on catch data grouped into age classes by backward solve the nonlinear catch equation

Usage

```
vpa.age(agesdf, catches=NULL, M=NULL, species=NULL)
```

Where

agesdf a vector with lower limit of age classes or an object of class `agedata` containing all required arguments

catches a vector with catch in numbers for each age class

M natural mortality rate

species identification of the species analyzed

→ Return an object of the class `vpaage` with the results.

References

Jones (1984); Quinn and Deriso (1999); Lassen and Medley (2001)

solveCatchEq

Solves catch equation using Newton-Raphson iteration

Usage

```
solveCatchEq(catch, number2, M, approxeps=0.01)
```

Where

catch catch in number in t

number2 number at sea in t+1

M natural mortality rate

approxeps precision for approximation

→ Return a list with number at sea and fishing mortality rate.

References

Press et al. (1992); Lassen and Medley (2001)

vpa.length

Performs a cohort analysis on catch data grouped into length classes by converting lengths to ages using an assumed von Bertalanffy growth curve.

Usage

```
vpa.length(lengthsdf, catches=NULL, Linf=NULL,  
           k=NULL, M=NULL, a=NULL, b=NULL, species=NULL)
```

Where

lengthsdf a numeric vector with lower limit of length classes or an object of class length-data containing all required arguments

catches a vector with catch in numbers for each length class

Linf asymptotic length of the von Bertalanffy growth curve

K Brody growth parameter of the von Bertalanffy growth curve

M natural mortality rate

a intercept of the length-weight relationship

b exponent of the length-weight relationship

species identification of the species analyzed

→ Return an object of the class vpalength with the results.

References

Jones (1984); Quinn and Deriso (1999); Lassen and Medley (2001)

summary.vpalength

Print the summary of results from length based cohort analysis

Usage

```
summary.vpalength(obj)
```

Where

obj object of the class **vpalength**

plot.vpalength

Plot profile of fishing mortalities over lengths and relative ages from length based cohort analysis

Usage

```
plot.vpalength(obj, xlab="Relative age", ylab="Fdt", toplab="Length", ...)
```

Where

obj object of the class **vpalength**

xlab label of age axis

ylab label of fishing mortality axis

toplab label of length axis

... parameters to be passed to plot method

print.vpalength

Print results from length based cohort analysis

Usage

```
print.vpalength(obj)
```

Where

obj object of the class **vpalength**

ThompsonBell

Perform the predictive Thompson & Bell model

Usage

```
ThompsonBell(obj, factors, prices=rep(0, length(obj$lengths)))
```

Where

obj object of the class **vpaage** or **vpalength**

factors vector of multiplication factors of F

prices vector of prices

→ Return an object of the class **TB** with results of the model

References

Sparre and Venema (1998); Quinn and Deriso (1999)

ThompsonBell.MaxY

Find the factor that maximize the yield

Usage

```
ThompsonBell.MaxY(obj, interval=c(0, 10))
```

Where

obj object of the class **TB**

interval interval of factors to be searched

→ Return the factor

ThompsonBell.MaxE

Find the factor that maximize the economic yield

Usage

```
ThompsonBell.MaxE(obj, interval=c(0, 10))
```

Where

obj object of the class **TB**

interval interval of factors to be searched

→ Return the factor

2.4 Beverton and Holt yield per recruit model

Beverton.Holt

Beverton and Holt biomass and yield per recruit calculation for a given set of parameters. Default method for integration is numerical, but also Gulland approximation and relative yield concept are available.

usage

```
Beverton.Holt(F=NULL, FZ=NULL, Tr=NULL, Tc=NULL, Lc=NULL, Tmax=NULL, M,  
             Wt=NULL, Winf=NULL, Linf=NULL, K=NULL, t0=NULL, b=3,  
             approxim=c("integration", "gulland","relative"))
```

Where

F fishing mortality rate

FZ exploitation rate F/Z

Tr age at recruitment

Tc age at first capture

Lc length at first capture

Tmax maximum age attained

M natural mortality rate

Wt weight at age (function)

Winf asymptotic weight

Linf asymptotic length

K growth rate

t0 age at zero length

b constant in length-weight relationship ($W = aL^b$)

approxim method for calculation

→ Return a object of class **BH** with the components:

YR yield per recruit

BR biomass per recruit

... input parameters

References

Beverton and Holt (1957, 1966); Sparre and Venema (1998); Quinn and Deriso (1999)

plot.BH

Plot the yield/recruit curve. The points on the curve for $F_{0.1}$ and F_{max} are shown.

Usage

```
plot.BH(obj, Flab="Fishing mortality",  
        YRlab="Yield/Recruit", BRlab="Biomass/Recruit")
```

Where

obj object of the class **BH**

Flab label for fishing mortality axis

YRlab label for yield/recruit axis

BRlab label for biomass/recruit axis

profile.BH

Profile yield/recruit of Beverton & Holt model over fishing mortalities

Usage

```
profile.BH(obj, Fs=seq(0, round(Fmax(obj) * 3, 0), (round(Fmax(obj) * 3, 0)/100)))
```

Where

obj object of the class **BH**

Fs vector of fishing mortalities

plot.profileBH

Plot the yield/recruit curve as function of fishing mortality

Usage

```
plot.profileBH(obj, Flab="Fishing mortality",  
              YRlab="Yield/Recruit", BRlab="Biomass/Recruit")
```

Where

obj object of the class **BH**

Flab label for fishing mortality axis

YRlab label for yield/recruit axis

BRIab label for biomass/recruit axis

isopleths

Plot yield/recruit isopleth diagram

Usage

`isopleths(obj,F,T)`

Where

obj object of the class **BH**

F vector of fishing mortalities

T vector of ages or lengths at first capture

References

Beverton and Holt (1957, 1966); Quinn and Deriso (1999)

F0.1

Fishing mortality rate at which the slope of the yield per recruit curve as a function of fishing mortality is 10% of its value at the origin. The fishing mortality level $F_{0.1}$ was proposed as a conservative target reference point.

Usage

`F0.1(obj)`

Where

obj object of the class **BH**

→ Return the $F_{0.1}$ value.

References

Gulland (1968); Caddy and Mahon (1995); Quinn and Deriso (1999)

Fmax

Fishing mortality for a given size at first capture, which maximizes the average yield from each recruit entering the fishery. It could be considered as an limit reference point for the stock.

Usage

`Fmax(obj)`

Where

obj object of the class **BH**

→ Return the F_{max} value.

References

Caddy and Mahon (1995); Sparre and Venema (1998); Quinn and Deriso (1999)

2.5 Selectivity

selection

Estimate selection curve from VPA or cohort analysis

Usage

`selection(obj)`

Where

obj object of the class **vpalength** or **vpaage**

→ Return an object of the class `selection` with a vector of selection over lengths or ages.

References

Sparre and Venema (1998)

S.f2logistic2

Fit a function of the product of two logistic equations on selection data

Usage

```
S.f2logistic2(obj, plot.it=T, maxiter=50)
```

Where

obj object of the class **selection**

plot.it TRUE for plot the function and data

maxiter maximum number of iterations (non-linear approximation)

→ Return an object of the class list with the coefficients of the function.

References

Sparre and Venema (1998)

S.f2logistic

Fit a function of the product of two logistic equations on selection data

Usage

```
S.f2logistic(obj, plot.it=T, ...)
```

Where

obj object of the class **selection**

plot.it TRUE for plot the function and data

... parameters to be passed to plot method

→ Return an object of the class list with the coefficients of the function.

References

Sparre and Venema (1998)

S.flogistic

Fit a logistic function on selection data

Usage

`S.flogistic(obj, plot.it=T)`

Where

obj object of the class **selection**

plot.it TRUE for plot the function and data

→ Return an object of the class list with the coefficients of the function.

References

Sparre and Venema (1998)

Plotlogistic2**Usage**

Plot gill-net like selection curve

`Plotlogistic2(lengths, S1, S2, D1, D2)`

Where

lengths vector of lengths

S1 $L_{50\%}$ value

S2 $L_{75\%}$ value

D1 $D_{50\%}$ value

D2 $D_{75\%}$ value

References

Sparre and Venema (1998)

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