FLife

WKLIFE Life History Relationships

19 agosto, 2017

FLife package

```
library(ggplot2)
library(FLife)
library(plyr)
library(reshape)
```

Life history parameters

```
data(wklife)
wklife
```

	name					common		area	stock	
1	Clupea harengus						ring	Celtic Seas	her-nis	F
2	Pollachius pollachius					Pollack		North Sea	pol-nsea	С
3	Molva molva					Ling		•	lin-comb	C
4	Sebastes norvegicus					Rose fish		Northern	smn-con	C
5	Mullus surmuletus					Red mullet		Celtic Seas	$\mathtt{mut-comb}$	F
6	Scopthalmus maximus					Τι	ırbot	North Sea	tur-nsea	F
7	Microstomus kitt					Lemon	sole	North Sea	lem-nsea	C
8	Lepidorhombus whiffiagonis					Me	egrim	North Sea	meg-4a6a	C
9	Ammodytes spp.					Sand	deels	North Sea	san-ns4	C
10	Pleuronectes platessa					P	Laice	Celtic Seas	ple-celt	F
11	Merlangius merlangus					Wha	iting	Celtic Seas	whg-7e-k	F
12	Melanogrammus aeglefinus					Had	ddock	Celtic Seas	had-iris	C
13	Lophius piscatorius White					angle	fish	Celtic Seas	ang-78ab	C
14	Lophius piscatorius Wh				nite	anglerfish		North Sea	ang-ivvi	C
15	Nephrops			ps		Shellfish		Biscay-Iberia	nep-2829	F
	a	b lmax	linf	150	a50	t0	k			
1	0.00480 3.	.20 NA	33.0	23.0	NA	NA	0.606			
2	0.00760 3.	.07 NA	85.6	47.1	NA	NA	0.190			
3	0.00360 3.	.11 NA	119.0	74.0	7.2	NA	0.140			
4	0.01780 2.	.97 NA	50.2	40.3	NA	0.08	0.110			
5	0.00570 3.	.24 NA	47.5	16.9	NA	NA	0.210			
6	0.01490 3.	.08 NA	66.7	34.2	2.2	0.29	0.320			
7	0.01230 2.	.97 NA	37.0	27.0	NA	NA	0.420			
8	0.00220 3.	.34 NA	54.0	23.0	3.0	NA	0.120			
9	0.00490 2.	.78 NA	24.0	12.0	NA	NA	1.000			
10	0.01100 2.	.96 NA	48.0	22.9	NA	NA	0.230			
11	0.01030 2.	.40 NA	38.0	28.0	NA	-1.01	0.380			
12	0.01130 2.	.96 NA	79.9	NA	2.0	-0.36	0.200			
13	0.01980 2.	.90 133	105.6			-0.38				
14	0.02970 2.	.84 NA	106.0	61.0	NA	NA	0.180			

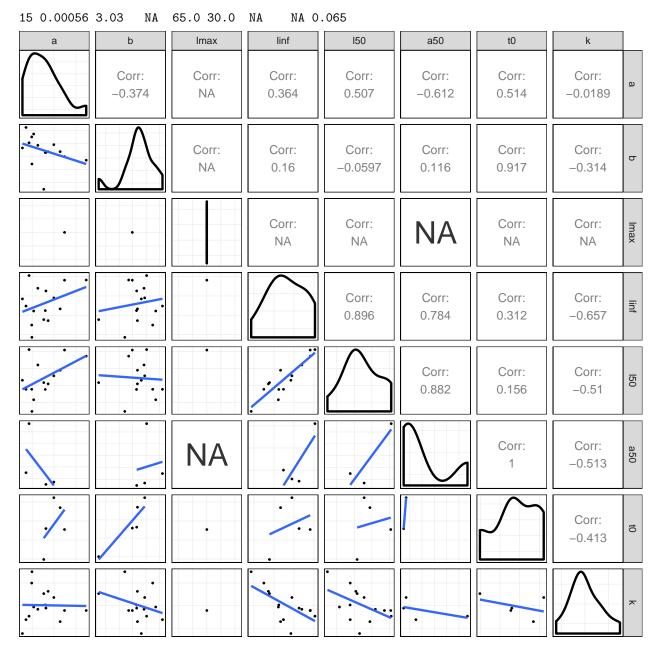


Figure 1 Pairwise scatter plots of life history parameters.

Equilibrium Dynamics

```
Create an FLPar
wkpar=as(wklife[,6:13],"FLPar")
attributes(wkpar)[names(wklife)[1:5]]=wklife[,1:5]
```

Then use life history relationships to estimate missing values

par=lhPar(wkpar)

and then to derive vectors for processes such as natural mortality

```
library(FLBRP)
eql=lhEql(par)
ggplot(FLQuants(eq1,"m","catch.sel","mat","catch.wt"))+
  geom_line(aes(age,data,col=attributes(wkpar)$name[iter]))+
  facet_wrap(~qname,scale="free")+
  scale_x_continuous(limits=c(0,15))+
  guides(colour=guide_legend(title="Species",title.position="top"))
                      m
                                                          catch.sel
    20
                                          1.0
    15
                                                                                 Species
                                                                                     Ammodytes spp.
    10 -
                                                                                     Clupea harengus
                                          0.5
                                                                                     Lepidorhombus whiffiagonis
     5
                                                                                     Lophius piscatorius
                                                                                     Melanogrammus aeglefinus
     0
                                          0.0
                                                                                     Merlangius merlangus
                 5
                           10
                                                                  10
data
                                     15
                                                                            15
                                                                                     Microstomus kitt
                     mat
                                                           catch.wt
                                                                                     Molva molva
  1.00
                                                                                     Mullus surmuletus
                                        15000
                                                                                     Nephrops
```

Figure 2 Vectors of m, selection pattern, maturity and weight-at-age.

and estimate equilibrium dynamics and reference points, e.g. for lemon sole

plot(iter(eq1,7))

10

15

10000

5000

age

15

10

Pleuronectes platessa Pollachius pollachius

Scopthalmus maximus

Sebastes norvegicus

0.75

0.50

0.25

0.00

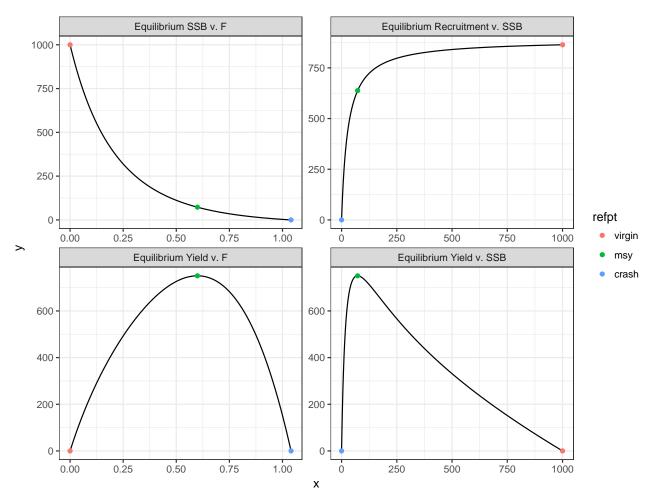


Figure 3 Equilibrium curves for lemon sole.

Simulation

Create a forward projection, i.e. an FLStock from an equilibrium object

```
lmsl=as(iter(eq1,7),"FLStock")
plot(lmsl)
```

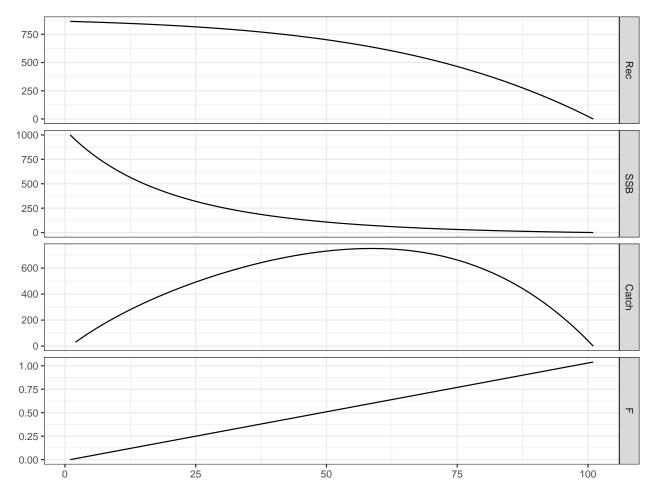


Figure 4 Simulate a stock with increasing F