

FLife

WKLIFE Life History Relationships

Laurence Kell

10 enero, 2018

FLife package

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

Life history parameters

```
data(wklife)
```

```
wklife
```

		name	common	area	stock	sex		
1		Clupea harengus	Herring	Celtic Seas	her-nis	F		
2		Pollachius pollachius	Pollack	North Sea	pol-nsea	C		
3		Molva molva	Ling	Widely	lin-comb	C		
4		Sebastes norvegicus	Rose fish	Northern	smn-con	C		
5		Mullus surmuletus	Red mullet	Celtic Seas	mut-comb	F		
6		Scophthalmus maximus	Turbot	North Sea	tur-nsea	F		
7		Microstomus kitt	Lemon sole	North Sea	lem-nsea	C		
8		Lepidorhombus whiffiagonis	Megrim	North Sea	meg-4a6a	C		
9		Ammodytes spp.	Sandeels	North Sea	san-ns4	C		
10		Pleuronectes platessa	Plaice	Celtic Seas	ple-celt	F		
11		Merlangius merlangus	Whiting	Celtic Seas	whg-7e-k	F		
12		Melanogrammus aeglefinus	Haddock	Celtic Seas	had-iris	C		
13		Lophius piscatorius	White anglerfish	Celtic Seas	ang-78ab	C		
14		Lophius piscatorius	White anglerfish	North Sea	ang-ivvi	C		
15		Nephrops	Shellfish	Biscay-Iberia	nep-2829	F		
	a	b	lmax	linf	l50	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 73.0 NA -0.38 0.180
14 0.02970 2.84 NA 106.0 61.0 NA NA 0.180
15 0.00056 3.03 NA 65.0 30.0 NA NA 0.065

```

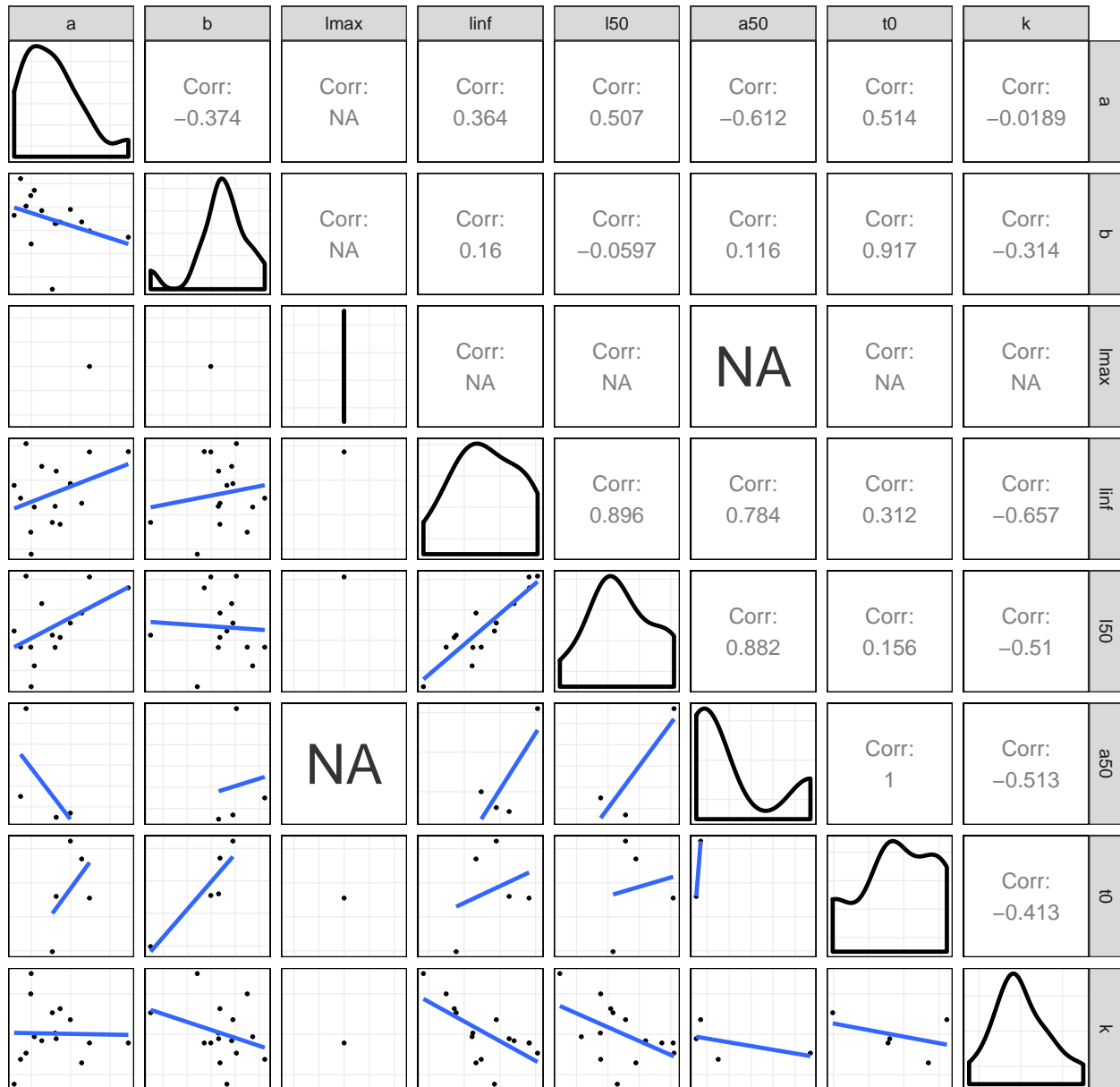


Figure 1 Pairwise scatter plots of life history parameters.

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)
```

```
sel<-function(x)
  catch.sel(x)/%fapex(catch.sel(x))

ggplot(FLQuants(eql,"m","catch.sel"=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"))
```

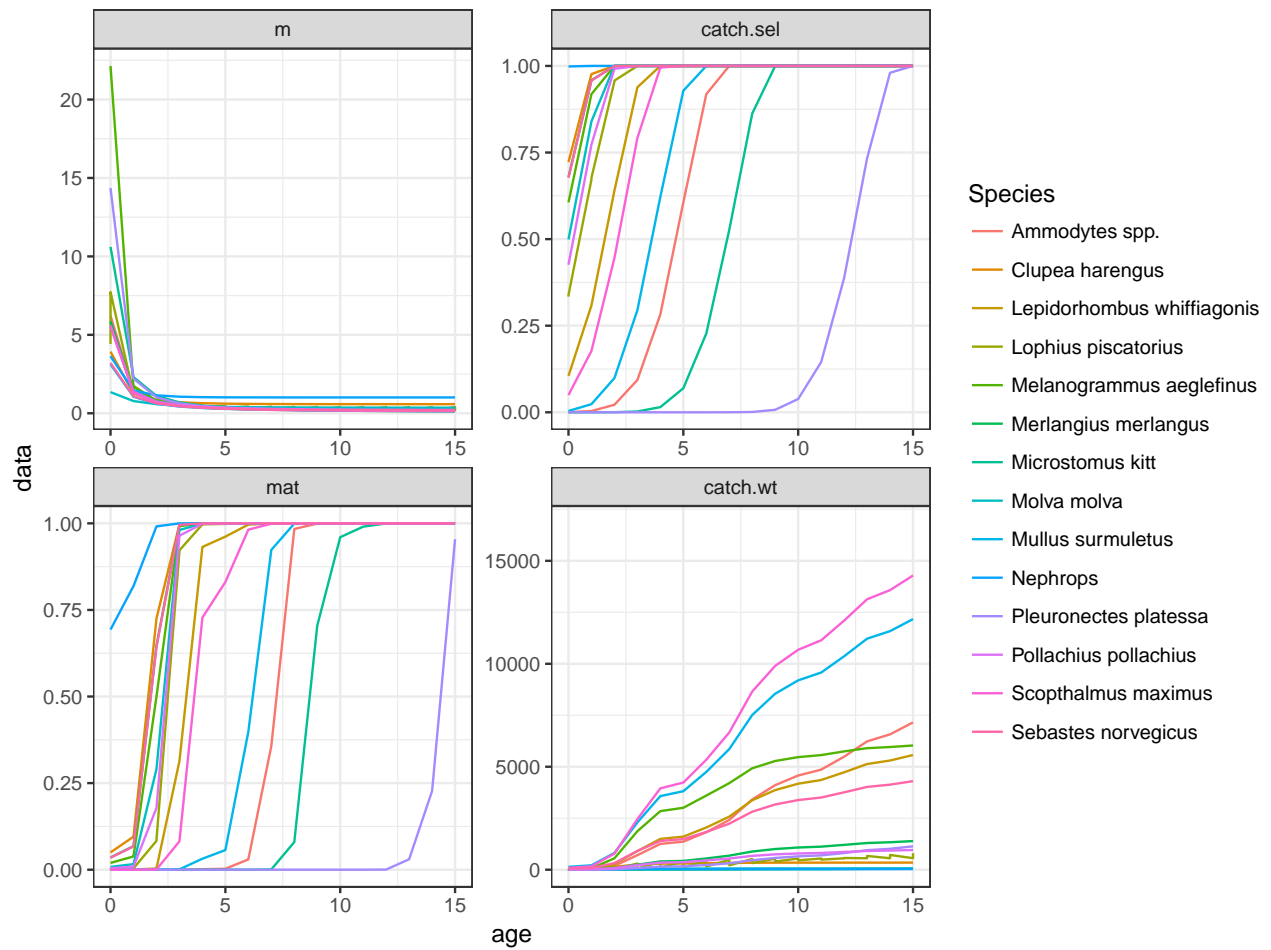


Figure 2 Vectors of m , selection pattern, maturity and weight-at-age. and estimate equilibrium dynamics and reference points, e.g. for lemon sole

```
plot(iter(eq1,7))
```

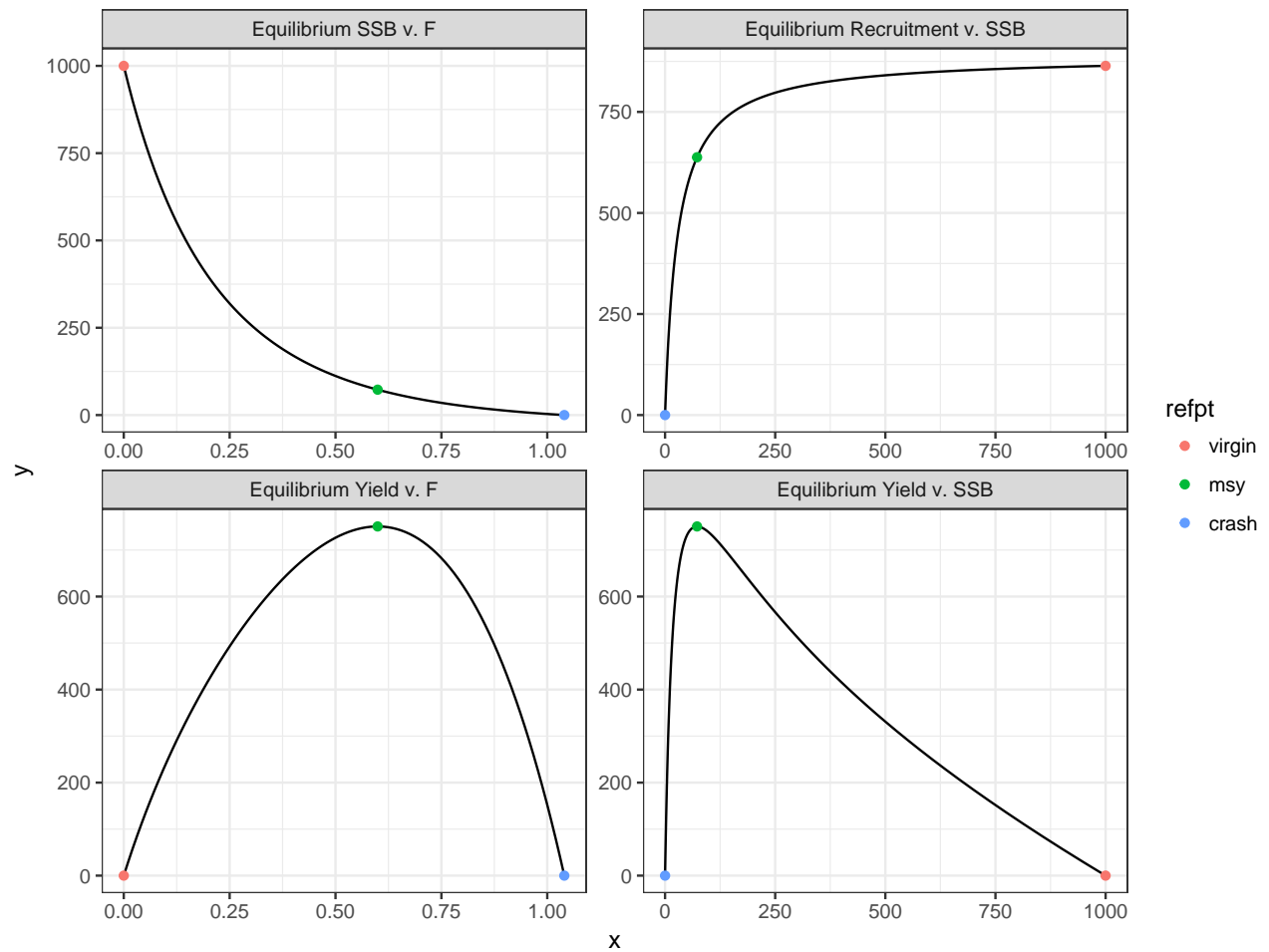


Figure 3 Equilibrium curves for lemon sole.

Simulation

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

```
lms1=as(iter(eq1,7),"FLStock")
```

```
plot(lms1)
```

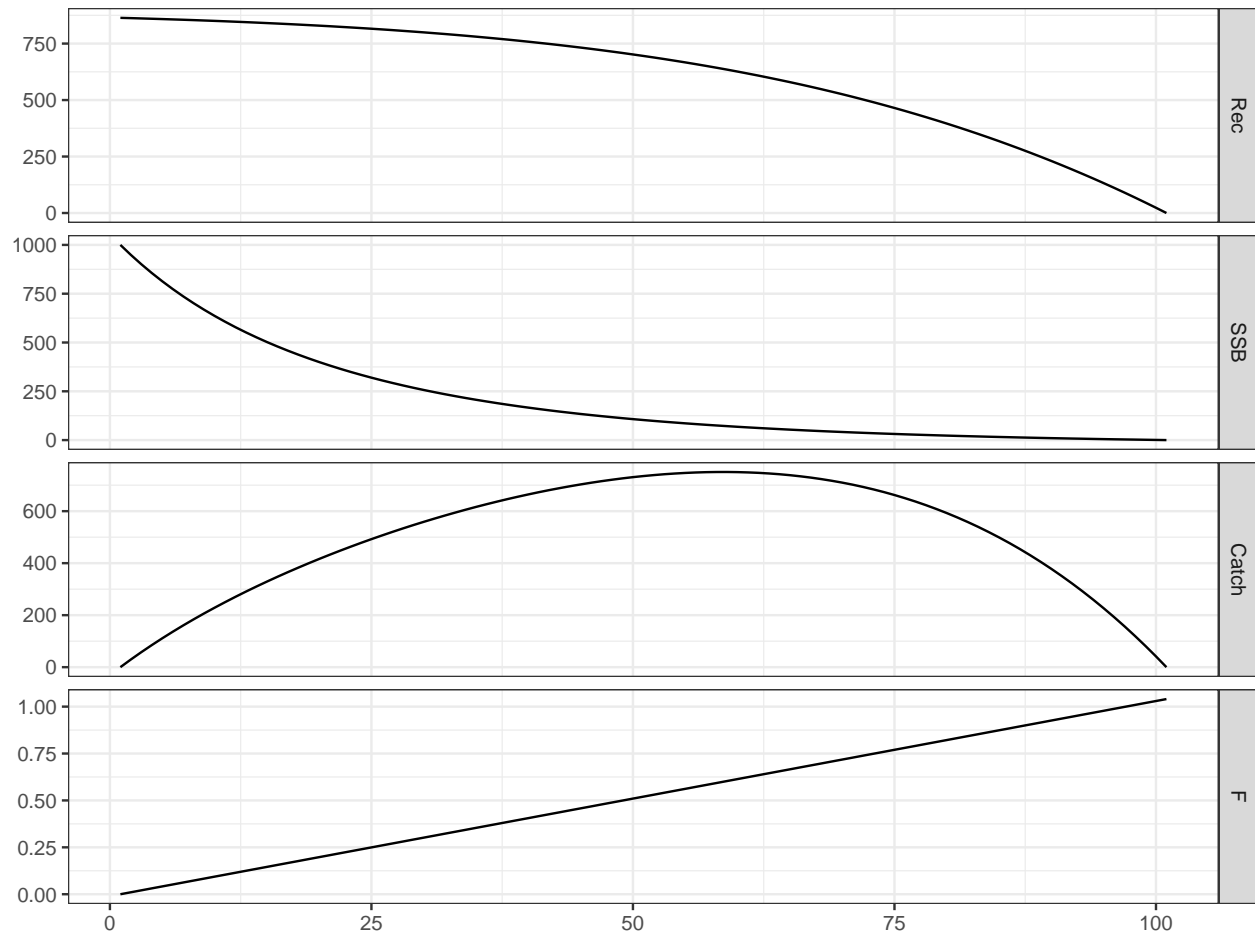


Figure 4 Simulate a stock with increasing F

Software Versions

- R version 3.4.1 (2017-06-30)
- FLCore: 2.6.5
- FLPKG:
- **Compiled:** Wed Jan 10 19:45:17 2018
- **Git Hash:** c6526f5

Author information

Laurence KELL. laurie.kell.es

Acknowledgements

This vignette and many of the methods documented in it were developed under the MyDas project funded by the Irish exchequer and EMFF 2014-2020. The overall aim of MyDas is to develop and test a range of assessment models and methods to establish Maximum Sustainable Yield (MSY) reference points (or proxy MSY reference points) across the spectrum of data-limited stocks.

References