MyDas

Gurnard Life History Relationships

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```
dirMy="/home/laurence/Desktop/sea++/mydas"
#dirMy =getwd()
dirInp=file.path(dirMy,"tasks/inputs")
dirDat=file.path(dirMy,"tasks/data")

library(ggplot2)
library(GGally)

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

Life history parameters

```
load(file.path(dirDat, "gurnards.RData"))
```

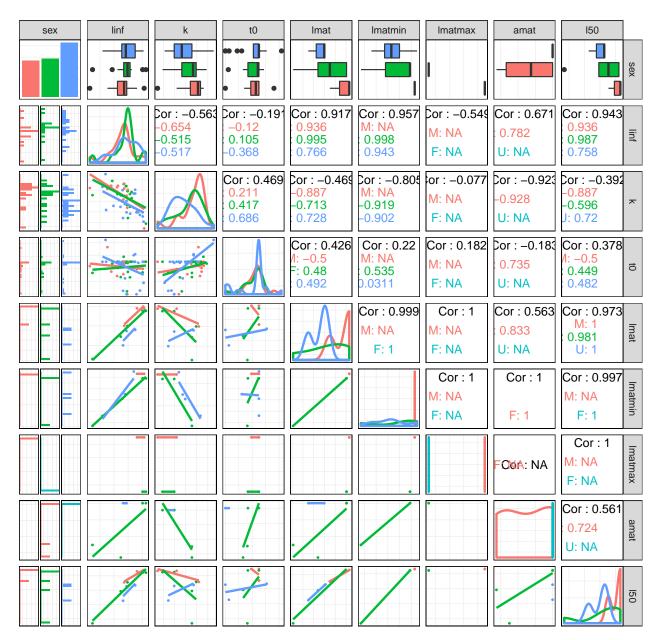


Figure 1 Pairwise scatter plots of life history parameters.

Equilibrium Dynamics

```
Create \ an \ {\tt 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(eql,"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"))
```

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

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

Figure 4 Simulate a stock with increasing F