

Summarizing Age Data

Derek H. Ogle, Northland College

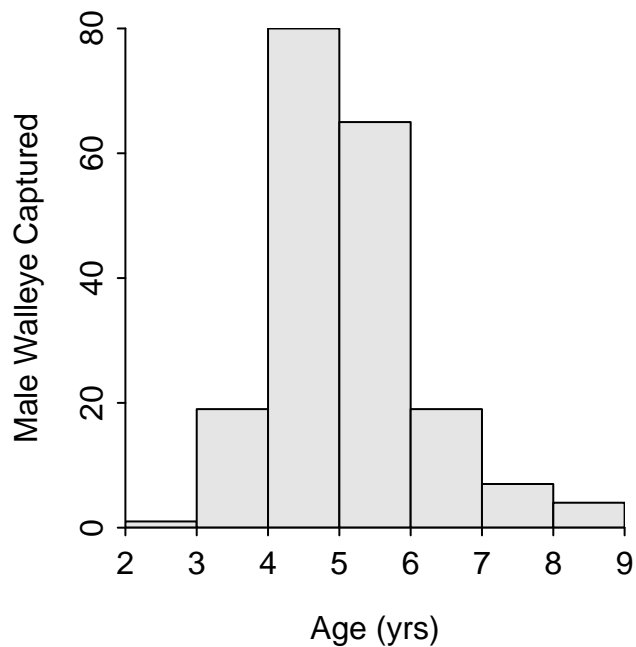
5-Mar-2015

Preliminaries

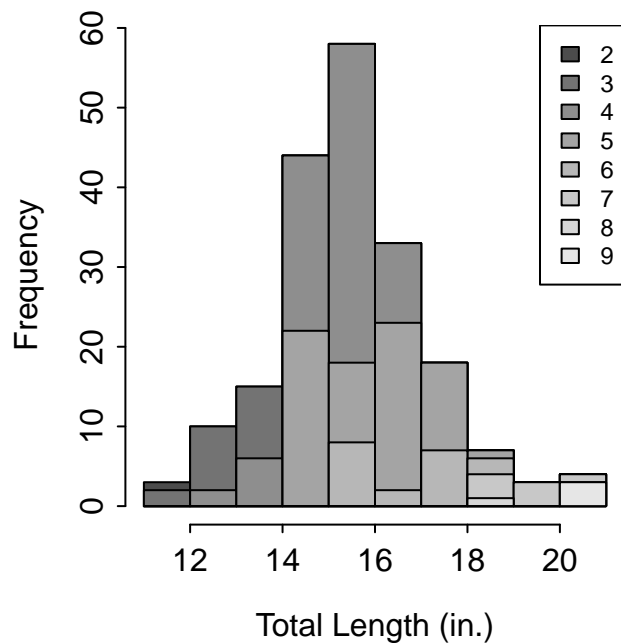
```
> source("02_AgeLengthKey.R")

> ls()
[1] "btxs"      "btys"      "clrs"      "fit1"      "fit2"      "fit3"
[7] "fn"        "hook1"     "lm1"       "mc1"       "ruf"       "ruf2"
[13] "ruf90"     "ruf9000"   "syms"      "tmp"       "tmpx"     "tmpy"
[19] "wae.aged"  "waeF.fn1"  "waeM.fn1"  "waeM.sumlen" "xs"       "ys"

> hist(~Age..observed.annuli.,data=waeM.fn1,xlab="Age (yrs)",ylab="Male Walleye Captured")
```



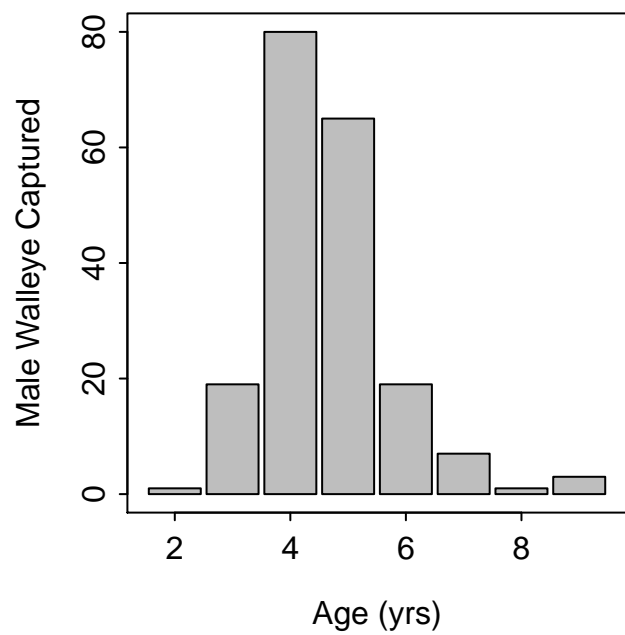
```
> histStack(Length.or.Lower.Length.IN~Age..observed.annuli.,data=waeM.fn1,xlab="Total Length (in.)",
            col="gray.colors",right=FALSE,legend.pos="topright")
Warning in histStack.default(mf[, 1], mf[, 2], breaks = breaks, col = col, : z was converted to a
factor
```



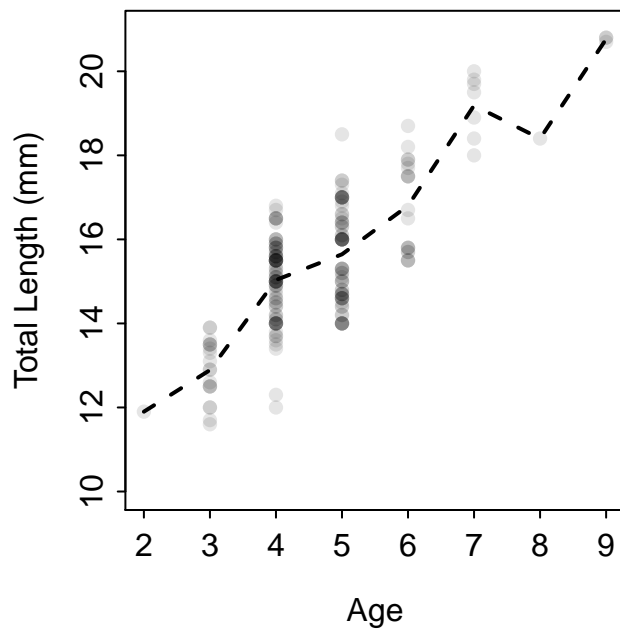
```
> waeM.sumlen <- waeM.fnl %>%
  group_by(Age..observed.annuli.) %>%
  summarize(n=n(),mean=mean(Length.or.Lower.Length.IN),sd=sd(Length.or.Lower.Length.IN),
            min=min(Length.or.Lower.Length.IN),max=max(Length.or.Lower.Length.IN))
> waeM.sumlen
Source: local data frame [8 x 6]
```

	Age..observed.annuli.	n	mean	sd	min	max
1	2	1	11.90000	NaN	11.9	11.9
2	3	19	12.88947	0.72025824	11.6	13.9
3	4	80	15.03375	0.94265230	12.0	16.8
4	5	65	15.64462	1.11313616	14.0	18.5
5	6	19	16.80000	1.10201835	15.5	18.7
6	7	7	19.18571	0.76469726	18.0	20.0
7	8	1	18.40000	NaN	18.4	18.4
8	9	3	20.76667	0.05773503	20.7	20.8

```
> plotH(n~Age..observed.annuli.,data=waem.sumlen,xlab="Age (yrs)",ylab="Male Walleye Captured",
  xlim=c(1.5,9.5),width=0.9)
```



```
> plot(Length.or.Lower.Length.IN~Age..observed.annuli.,data=waeM.fn1,pch=16,col=rgb(0,0,0,1/10),
      xlab="Age",ylab="Total Length (mm)",ylim=c(10,21))
> lines(mean~Age..observed.annuli.,data=waeM.sumlen,lwd=2,lty=2)
```



Construct and Apply an Age-Length Key – Females

Copy the code from above and convert the 'M's to 'F's

Application Assignment

Create a script that performs the following tasks:

1. Continue or `source()` your previous script.
2. Summarize the age distribution from the fish in your sample.
3. Show the mean length-at-age for all fish in your sample in both tabular and graphical forms.
4. (*Time Permitting*) Show the length frequency for all fish in your sample.
5. (*Time Permitting*) Repeat the above for your second sex or species.

Save your script!