

BIO311: Population Ecology
Prac 8: Life tables & Population Matrices

Koen van Benthem & Tina Cornioley

`koen.vanbenthem@ieu.uzh.ch`
`tina.cornioley@ieu.uzh.ch`

Spring 2014

Contents

| | | |
|----------|---|-----------|
| 1 | Preparing the data | 2 |
| 2 | Finding transition rates based on linear fits (lm) | 2 |
| 3 | Taking into account that $0 < S_J < 1$ and $0 < S_A < 1$ | 16 |
| 4 | What the students found | 18 |

This is 1 out of 3 methods (so far) for extracting transition rates from the raw data.

1 Preparing the data

```
rot<-read.csv("rdata.csv",header=T,sep=",")
```

Since we aim for transitions, rearrange the data, for the transition from timestep 1 to 2:

```
rot1<-subset(rot,Day==1)
rot2<-subset(rot,Day==2)
rot12<-merge(rot1,rot2,by=c("Population","Copper","Replicate"))
```

And for the transition from 2 to 3:

```
rot3<-subset(rot,Day==3)
rot23<-merge(rot2,rot3,by=c("Population","Copper","Replicate"))
```

And we create one super object that contains all transitions:

```
rot123<-rbind(rot12,rot23)
```

2 Finding transition rates based on linear fits (lm)

First we focus only on the transition from timestep 2 to 3

```
library('xtable')
store<-data.frame(Population=c(NA),Copper=c(NA),R=c(NA),SA=c(NA),SJ=c(NA))
for(i in levels(rot123$Population)){
  temp<-subset(rot23,Population==i)
  for(j in levels(rot123$Copper)){
    # Selecting the correct part of the dataset:
    temp2<-subset(temp,Copper==j)

    cat("\\subsection{",i,j,"}\\n")
    cat("\\subsubsection{Determining reproduction}\\n\\n")
    cat("First we fit a value for the reproductive rate using \\texttt{lm} where we force th

    par(mar=c(6,6,2,2))
    plot(temp2$Alive_Adult.x,temp2$Alive_Juv.y,xlab="Number of adults at time $t$",ylab="Num
```

```

reg1<-lm(temp2$Alive_Juv.y~temp2$Alive_Adult.x+0)
abline(reg1)
cat("\n From this regression we found that R=",reg1$coef[[1]],"\n\n")

cat("\n\\subsection{Determining survival}\n\n")
cat("Now we perform a second regression. We fit a plane in which the number of adults ne
reg2<-lm(temp2$Alive_Adult.y~temp2$Alive_Adult.x+temp2$Alive_Juv.x+0)
cat("\\\\From this we find:\\\\ \n$S_A=$",reg2$coef[[1]],"\n$S_J=$",reg2$coef[[2]],"\n")

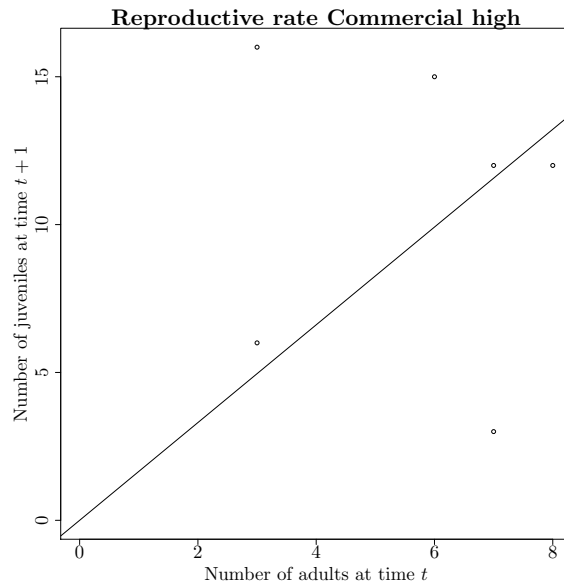
storet<-data.frame(Population=c(i),Copper=c(j),R=c(reg1$coef[[1]]),SA=c(reg2$coef[[1]]),
store<-rbind(store,storet)
}
}

```

2.1 Commercial high

2.1.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 1.653$

2.1.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the

number of adults this year

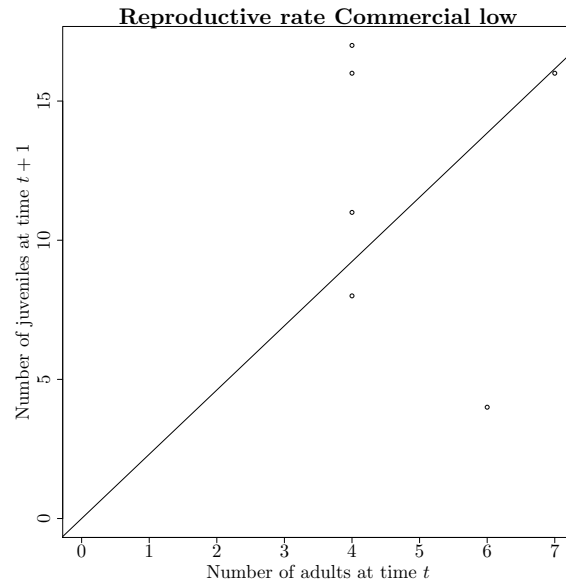
From this we find:

$$S_A = 1.698 \quad S_J = 0.5537$$

2.2 Commercial low

2.2.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 2.309$

2.2.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

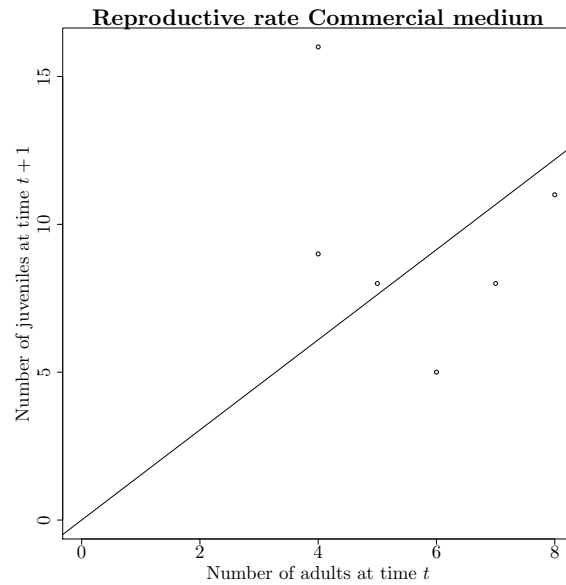
From this we find:

$$S_A = 0.893 \quad S_J = 1.039$$

2.3 Commercial medium

2.3.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 1.524$

2.3.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

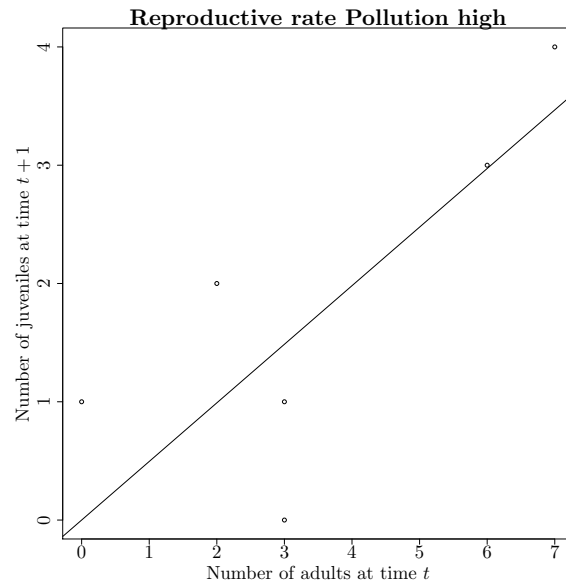
From this we find:

$$S_A = 1.669 \quad S_J = 0.3427$$

2.4 Pollution high

2.4.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.4953$

2.4.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

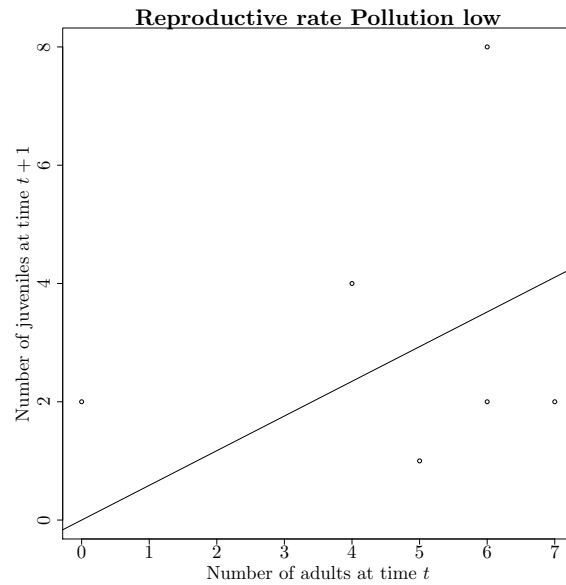
From this we find:

$$S_A = 1.295 \quad S_J = -0.07418$$

2.5 Pollution low

2.5.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.5864$

2.5.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

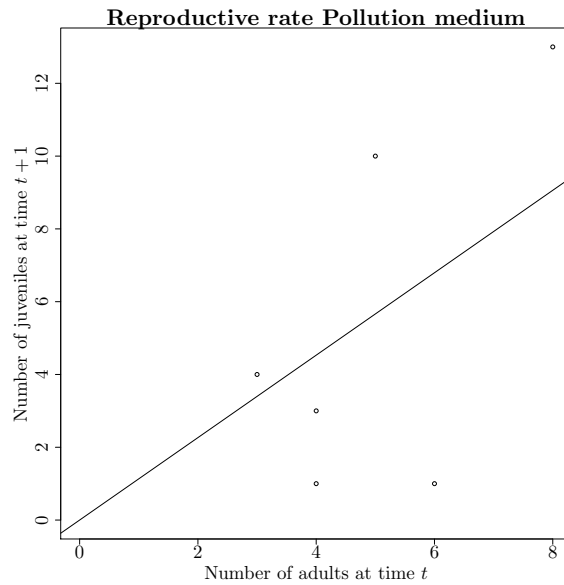
From this we find:

$$S_A = 1.373 \quad S_J = 0.424$$

2.6 Pollution medium

2.6.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 1.133$

2.6.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

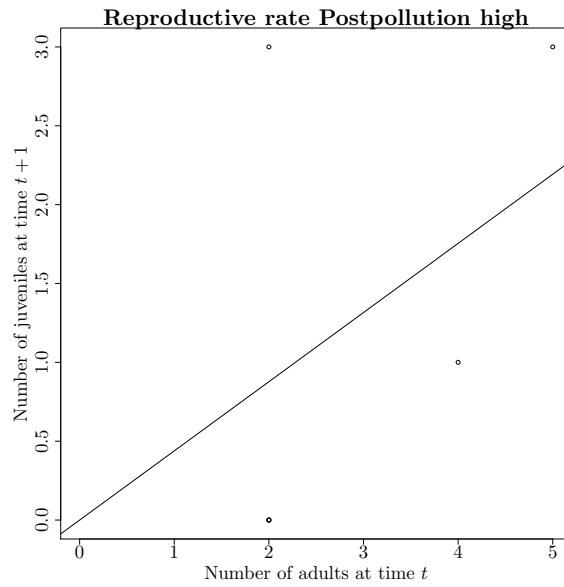
From this we find:

$$S_A = 1.086 \quad S_J = -0.08364$$

2.7 Postpollution high

2.7.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.4386$

2.7.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

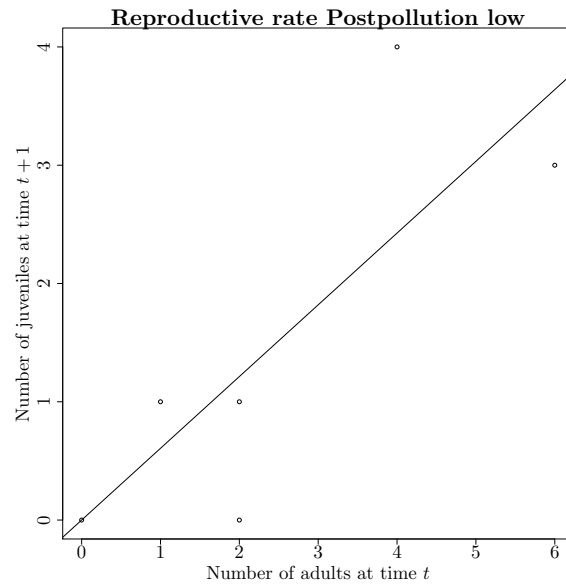
From this we find:

$$S_A = 1.024 \quad S_J = 0.4491$$

2.8 Postpollution low

2.8.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.6066$

2.8.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

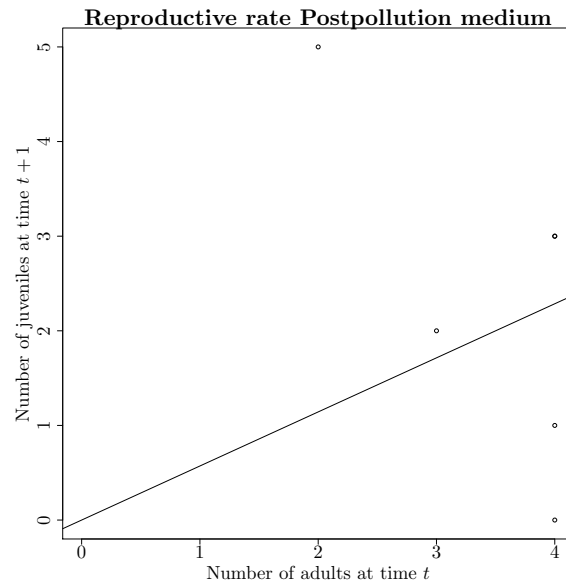
From this we find:

$$S_A = 0.07692 \quad S_J = 4.692$$

2.9 Postpollution medium

2.9.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.5714$

2.9.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

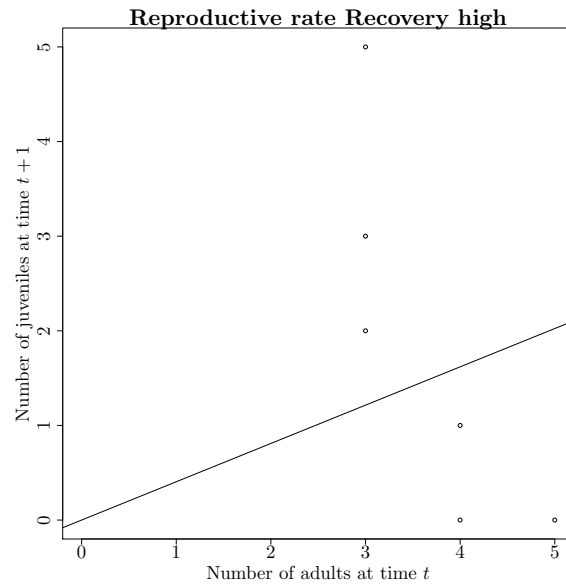
From this we find:

$$S_A = 0.8715 \quad S_J = -0.257$$

2.10 Recovery high

2.10.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.4048$

2.10.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

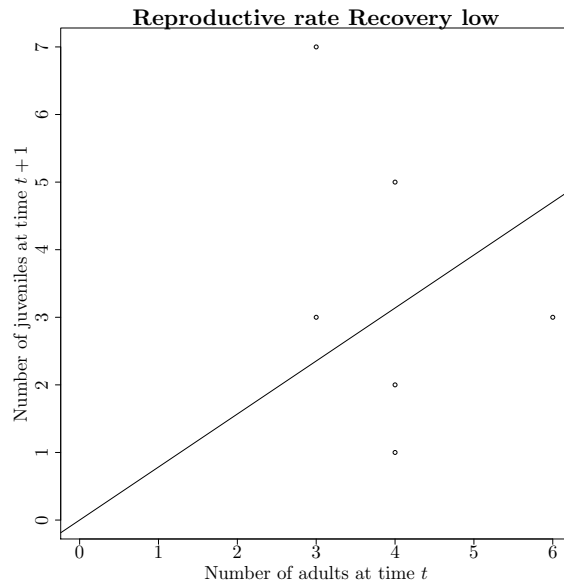
From this we find:

$$S_A = 1.097 \quad S_J = 0.4803$$

2.11 Recovery low

2.11.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.7843$

2.11.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

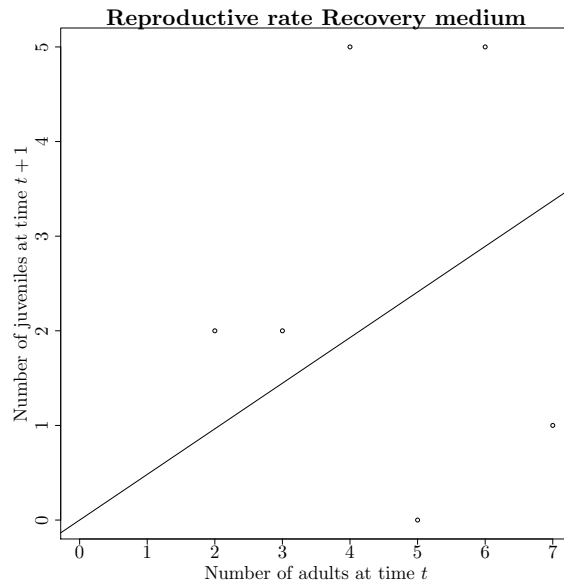
From this we find:

$$S_A = 0.8964 \quad S_J = 0.5193$$

2.12 Recovery medium

2.12.1 Determining reproduction

First we fit a value for the reproductive rate using `lm` where we force the line to go through 0.



From this regression we found that $R = 0.482$

2.12.2 Determining survival

Now we perform a second regression. We fit a plane in which the number of adults next year depends on both the number of juveniles this year and the number of adults this year

From this we find:

$$S_A = 1.09 \quad S_J = -0.1955$$

```
cat("\\subsection{Summary of the data for the 2--3 transition}")
```

2.13 Summary of the data for the 2–3 transition

```
store<-store[-1,]
print(xtable(store, digits=2),
      size="footnotesize", #Change size; useful for bigger tables
      include.rownames=FALSE, #Don't print rownames
      include.colnames=FALSE, #We create them ourselves
      floating=FALSE,
      hline.after=NULL, #We don't need hline; we use booktabs
      add.to.row = list(pos = list(-1,
                                   nrow(store)),
                        command = c(paste("\\toprule \\n",
                                           "$Population$ & $Copper$ & $R$ & $S_A$ & $S_J$ \\\\",
                                           "\\midrule \\n"),
```

```
"\\bottomrule \n")
```

```
))
```

| <i>Population</i> | <i>Copper</i> | <i>R</i> | <i>S_A</i> | <i>S_J</i> |
|-------------------|---------------|----------|----------------------|----------------------|
| Commercial | high | 1.65 | 1.70 | 0.55 |
| Commercial | low | 2.31 | 0.89 | 1.04 |
| Commercial | medium | 1.52 | 1.67 | 0.34 |
| Pollution | high | 0.50 | 1.30 | -0.07 |
| Pollution | low | 0.59 | 1.37 | 0.42 |
| Pollution | medium | 1.13 | 1.09 | -0.08 |
| Postpollution | high | 0.44 | 1.02 | 0.45 |
| Postpollution | low | 0.61 | 0.08 | 4.69 |
| Postpollution | medium | 0.57 | 0.87 | -0.26 |
| Recovery | high | 0.40 | 1.10 | 0.48 |
| Recovery | low | 0.78 | 0.90 | 0.52 |
| Recovery | medium | 0.48 | 1.09 | -0.20 |

```
cat("\\subsection{Summary of the data for both transitions}")
```

2.14 Summary of the data for both transitions

```
store<-data.frame(Population=c(NA),Copper=c(NA),R=c(NA),SA=c(NA),SJ=c(NA))
for(i in levels(rot123$Population)){
  temp<-subset(rot123,Population==i)
  for(j in levels(rot123$Copper)){
    # Selecting the correct part of the dataset:
    temp2<-subset(temp,Copper==j)
    reg1<-lm(temp2$Alive_Juv.y~temp2$Alive_Adult.x+0)
    reg2<-lm(temp2$Alive_Adult.y~temp2$Alive_Adult.x+temp2$Alive_Juv.x+0)

    storet<-data.frame(Population=c(i),Copper=c(j),R=c(reg1$coef[[1]]),SA=c(reg2$coef[[1]]))
    store<-rbind(store,storet)
  }
}
store<-store[-1,]
print(xtable(store, digits=2),
      size="footnotesize", #Change size; useful for bigger tables
      include.rownames=FALSE, #Don't print rownames
      include.colnames=FALSE, #We create them ourselves
      floating=FALSE,
      hline.after=NULL, #We don't need hline; we use booktabs
      add.to.row = list(pos = list(-1,
                                   nrow(store)),
                        command = c(paste("\\toprule \n",
                                           "$Population$ & $Copper$ & $R$ & $S_A$ & $S_J$ \n",
                                           "\\midrule \n"),
```

"\\bottomrule \n")

))

| <i>Population</i> | <i>Copper</i> | <i>R</i> | <i>S_A</i> | <i>S_J</i> |
|-------------------|---------------|----------|----------------------|----------------------|
| Commercial | high | 1.59 | 1.72 | 0.42 |
| Commercial | low | 2.26 | 0.75 | 1.09 |
| Commercial | medium | 1.55 | 1.66 | 0.32 |
| Pollution | high | 0.48 | 1.31 | -0.13 |
| Pollution | low | 0.59 | 1.40 | 0.24 |
| Pollution | medium | 1.09 | 0.79 | 0.61 |
| Postpollution | high | 0.40 | 1.15 | -0.10 |
| Postpollution | low | 0.40 | 0.95 | -0.04 |
| Postpollution | medium | 0.69 | 0.76 | 0.09 |
| Recovery | high | 0.62 | 1.14 | 0.29 |
| Recovery | low | 0.78 | 0.94 | 0.42 |
| Recovery | medium | 0.36 | 1.05 | 0.30 |

3 Taking into account that $0 < S_J < 1$ and $0 < S_A < 1$

We now use `nls()` to ensure that both S_J and S_A stay between 0 and 1. (not sure if this is the best solution, but at least it is a solution.)

```
library('xtable')
store<-data.frame(Population=c(NA),Copper=c(NA),R=c(NA),SA=c(NA),SJ=c(NA))
for(i in levels(rot123$Population)){
  temp<-subset(rot123,Population==i)
  for(j in levels(rot123$Copper)){
    # Selecting the correct part of the dataset:
    temp2<-subset(temp,Copper==j)
    reg1<-lm(temp2$Alive_Juv.y~temp2$Alive_Adult.x+0)
    y<-temp2$Alive_Adult.y
    x1<-temp2$Alive_Adult.x
    x2<-temp2$Alive_Juv.x
    nlsfit<-nls(y~sa*x1+sj*x2,start=c(sa=0.5,sj=0.5),upper=c(sa=1,sj=1),lower=c(sa=0,sj=0),a

    storet<-data.frame(Population=c(i),Copper=c(j),R=c(reg1$coef[[1]]),SA=c(coef(nlsfit)[[1]]),SJ=c(coef(nlsfit)[[2]]))
    store<-rbind(store,storet)
  }
}
store<-store[-1,]
print(xtable(store, digits=2),
      size="footnotesize", #Change size; useful for bigger tables
      include.rownames=FALSE, #Don't print rownames
      include.colnames=FALSE, #We create them ourselves
```



```

floating=FALSE,
hline.after=NULL, #We don't need hline; we use booktabs
add.to.row = list(pos = list(-1,
                             nrow(store)),
                  command = c(paste("\\toprule \n",
                                     "$Population$ & $Copper$ & $R$ & $S_A$ & $S_J$ \\",
                                     "\\midrule \n"),
                              "\\bottomrule \n")
                  ))

```

| <i>Population</i> | <i>Copper</i> | <i>R</i> | <i>S_A</i> | <i>S_J</i> |
|-------------------|---------------|----------|----------------------|----------------------|
| Commercial | high | 1.59 | 1.00 | 1.00 |
| Commercial | low | 2.26 | 0.85 | 1.00 |
| Commercial | medium | 1.55 | 1.00 | 0.96 |
| Pollution | high | 0.48 | 1.00 | 0.25 |
| Pollution | low | 0.59 | 1.00 | 0.72 |
| Pollution | medium | 1.09 | 0.79 | 0.61 |
| Postpollution | high | 0.40 | 1.00 | 0.06 |
| Postpollution | low | 0.40 | 0.92 | 0.00 |
| Postpollution | medium | 0.69 | 0.76 | 0.09 |
| Recovery | high | 0.62 | 1.00 | 0.42 |
| Recovery | low | 0.78 | 0.94 | 0.42 |
| Recovery | medium | 0.36 | 1.00 | 0.35 |

```

store$lambda<-NA
library('popbio')

## Loading required package: quadprog

for(i in 1:length(store$SJ)){
  A<-matrix(c(0,store$SJ[i],store$R[i],store$SA[i]),nrow=2)

  store$lambda[i]<-lambda(A)
}
print(xtable(store, digits=4),
      size="footnotesize", #Change size; useful for bigger tables
      include.rownames=FALSE, #Don't print rownames
      include.colnames=FALSE, #We create them ourselves
      floating=FALSE,
      hline.after=NULL, #We don't need hline; we use booktabs
      add.to.row = list(pos = list(-1,
                                   nrow(store)),
                        command = c(paste("\\toprule \n",
                                           "$Population$ & $Copper$ & $R$ & $S_A$ & $S_J$ & $S",
                                           "\\midrule \n"),
                                   "\\bottomrule \n")
                        ))

```

| <i>Population</i> | <i>Copper</i> | <i>R</i> | <i>S_A</i> | <i>S_J</i> | <i>λ</i> |
|-------------------|---------------|----------|----------------------|----------------------|----------|
| Commercial | high | 1.5889 | 1.0000 | 1.0000 | 1.8561 |
| Commercial | low | 2.2562 | 0.8522 | 1.0000 | 1.9874 |
| Commercial | medium | 1.5538 | 1.0000 | 0.9565 | 1.8177 |
| Pollution | high | 0.4783 | 1.0000 | 0.2500 | 1.1079 |
| Pollution | low | 0.5926 | 1.0000 | 0.7241 | 1.3241 |
| Pollution | medium | 1.0900 | 0.7948 | 0.6130 | 1.3063 |
| Postpollution | high | 0.3962 | 1.0000 | 0.0645 | 1.0249 |
| Postpollution | low | 0.4000 | 0.9217 | 0.0000 | 0.9217 |
| Postpollution | medium | 0.6928 | 0.7583 | 0.0945 | 0.8366 |
| Recovery | high | 0.6165 | 1.0000 | 0.4153 | 1.2114 |
| Recovery | low | 0.7821 | 0.9356 | 0.4224 | 1.2089 |
| Recovery | medium | 0.3591 | 1.0000 | 0.3529 | 1.1138 |

4 What the students found

| <i>Population</i> | <i>Copper</i> | <i>R</i> | <i>S_A</i> | <i>S_J</i> | <i>λ</i> |
|-------------------|---------------|----------|----------------------|----------------------|----------|
| Commercial | high | 2.8557 | 0.9523 | 0.8917 | 2.1414 |
| Commercial | low | 3.3523 | 0.9523 | 0.8555 | 2.2353 |
| Commercial | medium | 2.2175 | 0.9583 | 0.9207 | 1.9862 |
| Pollution | high | 0.5126 | 0.8334 | 0.5833 | 1.1042 |
| Pollution | low | 0.8452 | 0.6695 | 0.5833 | 1.1126 |
| Pollution | medium | 1.2013 | 0.8790 | 0.3195 | 1.1991 |
| Postpollution | high | 0.5750 | 0.5833 | 0.4167 | 0.8614 |
| Postpollution | low | 1.0000 | 0.5833 | 0.2500 | 0.8705 |
| Postpollution | medium | 1.1528 | 0.5833 | 0.4000 | 1.0307 |
| Recovery | high | 0.5973 | 0.9583 | 0.7222 | 1.2922 |
| Recovery | low | 0.8888 | 0.8888 | 0.5833 | 1.2906 |
| Recovery | medium | 0.5912 | 0.7112 | 0.2000 | 0.8502 |