

Mark-recapture lab

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Question 1

The Lincoln-Peterson estimate of abundance is $N=200$.

```
n1<-100
n2<-50
m2<-25

N<-n1*n2/m2
N
```

```
## [1] 200
```

Models

```
require(mra)
```

```
## Loading required package: mra
```

```
## mra (version 2.16.11)
```

```
##M0 model
M0 <- F.huggins.estim(capture=-1, recapture=NULL, histories=ch.mat)
M0
```

```
## Call:
## F.huggins.estim(capture = -1, recapture = NULL, histories = ch.mat)
##
## Capture and Recapture model:
## Variable Est SE
## (Intercept) -2.37255 0.49911
##
## Population Size Estimate (se): 41.0364 (16.6626)
## 95% confidence interval for population size: 24.04 to 99.05
## Individuals observed: 17
## Effective sample size: 102
##
## Message = SUCCESS: Convergence criterion met
## Number of estimable coefficients (estimated) = 1
## Log likelihood = -43.9354578026216
## Deviance = 87.8709156052432
## AIC = 89.8709156052432
## AICc = 89.9189156052432
```

```
round(M0$phat, digits=3)
```

```
## [1,] [2,] [3,] [4,] [5,] [6,]
## [1,] 0.085 0.085 0.085 0.085 0.085 0.085
## [2,] 0.085 0.085 0.085 0.085 0.085 0.085
## [3,] 0.085 0.085 0.085 0.085 0.085 0.085
## [4,] 0.085 0.085 0.085 0.085 0.085 0.085
## [5,] 0.085 0.085 0.085 0.085 0.085 0.085
## [6,] 0.085 0.085 0.085 0.085 0.085 0.085
## [7,] 0.085 0.085 0.085 0.085 0.085 0.085
## [8,] 0.085 0.085 0.085 0.085 0.085 0.085
## [9,] 0.085 0.085 0.085 0.085 0.085 0.085
## [10,] 0.085 0.085 0.085 0.085 0.085 0.085
## [11,] 0.085 0.085 0.085 0.085 0.085 0.085
## [12,] 0.085 0.085 0.085 0.085 0.085 0.085
## [13,] 0.085 0.085 0.085 0.085 0.085 0.085
## [14,] 0.085 0.085 0.085 0.085 0.085 0.085
## [15,] 0.085 0.085 0.085 0.085 0.085 0.085
## [16,] 0.085 0.085 0.085 0.085 0.085 0.085
## [17,] 0.085 0.085 0.085 0.085 0.085 0.085
```

```
##Mb model
Mb <- F.huggins.estim(capture=-1, recapture=-1, histories=ch.mat)
Mb
```

```
## Call:
## F.huggins.estim(capture = -1, recapture = -1, histories = ch.mat)
##
## Capture Model Est SE Recapture Model Est SE
## (Intercept) -0.67847 0.48712 C:(Intercept) -0.67847 0.48712(fixed)
## B:(Intercept) -1.97829 0.71051
##
## Population Size Estimate (se): 18.5841 (2.1527)
## 95% confidence interval for population size: 17.21 to 28.76
## Individuals observed: 17
## Effective sample size: 102
##
## Message = SUCCESS: Convergence criterion met
## Number of estimable coefficients (estimated) = 2
## Log likelihood = -41.609252319818
## Deviance = 83.2185064639637
## AIC = 87.2185064639637
## AICc = 87.3397185851758
```

```
round(Mb$phat, digits=3)
```

```
## [1,] [2,] [3,] [4,] [5,] [6,]
## [1,] 0.337 0.337 0.337 0.337 0.337 0.337
## [2,] 0.337 0.337 0.337 0.337 0.337 0.337
## [3,] 0.337 0.337 0.337 0.337 0.337 0.337
## [4,] 0.337 0.337 0.337 0.337 0.337 0.337
## [5,] 0.337 0.337 0.337 0.337 0.337 0.337
## [6,] 0.337 0.337 0.337 0.337 0.337 0.337
## [7,] 0.337 0.337 0.337 0.337 0.337 0.337
## [8,] 0.337 0.337 0.337 0.337 0.337 0.337
## [9,] 0.337 0.337 0.337 0.337 0.337 0.337
## [10,] 0.337 0.337 0.337 0.337 0.337 0.337
## [11,] 0.337 0.337 0.337 0.337 0.337 0.337
## [12,] 0.337 0.337 0.337 0.337 0.337 0.337
## [13,] 0.337 0.337 0.337 0.337 0.337 0.337
## [14,] 0.337 0.337 0.337 0.337 0.337 0.337
## [15,] 0.337 0.337 0.337 0.337 0.337 0.337
## [16,] 0.337 0.337 0.337 0.337 0.337 0.337
## [17,] 0.337 0.337 0.337 0.337 0.337 0.337
```

```
round(Mb$Sc.hat, digits=3)
```

```
## [1,] [2,] [3,] [4,] [5,] [6,]
## [1,] NA 0.066 0.066 0.066 0.066 0.066
## [2,] NA 0.066 0.066 0.066 0.066 0.066
## [3,] NA 0.066 0.066 0.066 0.066 0.066
## [4,] NA 0.066 0.066 0.066 0.066 0.066
## [5,] NA 0.066 0.066 0.066 0.066 0.066
## [6,] NA 0.066 0.066 0.066 0.066 0.066
## [7,] NA 0.066 0.066 0.066 0.066 0.066
## [8,] NA 0.066 0.066 0.066 0.066 0.066
## [9,] NA 0.066 0.066 0.066 0.066 0.066
## [10,] NA 0.066 0.066 0.066 0.066 0.066
## [11,] NA 0.066 0.066 0.066 0.066 0.066
## [12,] NA 0.066 0.066 0.066 0.066 0.066
## [13,] NA 0.066 0.066 0.066 0.066 0.066
## [14,] NA 0.066 0.066 0.066 0.066 0.066
## [15,] NA 0.066 0.066 0.066 0.066 0.066
## [16,] NA 0.066 0.066 0.066 0.066 0.066
## [17,] NA 0.066 0.066 0.066 0.066 0.066
```

```
##Mt model
time <- tvar(factor(1:6), nan=nrow(ch.mat)) ## 6 time periods. 14 animals.
Mt <- F.huggins.estim(capture=time, recapture=NULL, histories=ch.mat)
Mt
```

```
## Call:
## F.huggins.estim(capture = -time, recapture = NULL, histories = ch.mat)
##
## Capture and Recapture model:
## Variable Est SE
## (Intercept) -1.68759 0.64342
## time:2 -0.78331 0.74956
## time:3 0.18611 0.6113
## time:4 -1.21711 0.85368
## time:5 -0.78331 0.74956
## time:6 -31.02057 268608.00057
##
## Population Size Estimate (se): 37.7944 (14.7706)
## 95% confidence interval for population size: 22.94 to 89.78
## Individuals observed: 17
## Effective sample size: 102
##
## Message = SUCCESS: Convergence criterion met
## Number of estimable coefficients (estimated) = 5
## Log likelihood = -37.42450409473
## Deviance = 74.8490081894601
## AIC = 84.8490081894601
## AICc = 85.4740081894601
```

```
round(Mt$phat, digits=3)
```

```
## [1,] [2,] [3,] [4,] [5,] [6,]
## [1,] 0.159 0.079 0.185 0.053 0.079 0
## [2,] 0.159 0.079 0.185 0.053 0.079 0
## [3,] 0.159 0.079 0.185 0.053 0.079 0
## [4,] 0.159 0.079 0.185 0.053 0.079 0
## [5,] 0.159 0.079 0.185 0.053 0.079 0
## [6,] 0.159 0.079 0.185 0.053 0.079 0
## [7,] 0.159 0.079 0.185 0.053 0.079 0
## [8,] 0.159 0.079 0.185 0.053 0.079 0
## [9,] 0.159 0.079 0.185 0.053 0.079 0
## [10,] 0.159 0.079 0.185 0.053 0.079 0
## [11,] 0.159 0.079 0.185 0.053 0.079 0
## [12,] 0.159 0.079 0.185 0.053 0.079 0
## [13,] 0.159 0.079 0.185 0.053 0.079 0
## [14,] 0.159 0.079 0.185 0.053 0.079 0
## [15,] 0.159 0.079 0.185 0.053 0.079 0
## [16,] 0.159 0.079 0.185 0.053 0.079 0
## [17,] 0.159 0.079 0.185 0.053 0.079 0
```

```
round(Mt$Sc.hat, digits=3)
```

```
## [1,] [2,] [3,] [4,] [5,] [6,]
## [1,] NA 0.079 0.185 0.053 0.079 0
## [2,] NA 0.079 0.185 0.053 0.079 0
## [3,] NA 0.079 0.185 0.053 0.079 0
## [4,] NA 0.079 0.185 0.053 0.079 0
## [5,] NA 0.079 0.185 0.053 0.079 0
## [6,] NA 0.079 0.185 0.053 0.079 0
## [7,] NA 0.079 0.185 0.053 0.079 0
## [8,] NA 0.079 0.185 0.053 0.079 0
## [9,] NA 0.079 0.185 0.053 0.079 0
## [10,] NA 0.079 0.185 0.053 0.079 0
## [11,] NA 0.079 0.185 0.053 0.079 0
## [12,] NA 0.079 0.185 0.053 0.079 0
## [13,] NA 0.079 0.185 0.053 0.079 0
## [14,] NA 0.079 0.185 0.053 0.079 0
## [15,] NA 0.079 0.185 0.053 0.079 0
## [16,] NA 0.079 0.185 0.053 0.079 0
## [17,] NA 0.079 0.185 0.053 0.079 0
```

```
##Mtb model
Mtb <- F.huggins.estim(capture=-time, recapture=-1, histories=ch.mat)
Mtb
```

```
## Call:
## F.huggins.estim(capture = -time, recapture = -1, histories = ch.mat)
##
## Capture Model Est SE Recapture Model Est SE
## (Intercept) -0.96169 0.96236 C:(Intercept) -0.96169 0.96236 (fixed)
## time:2 -0.62584 0.84758 C:time:2 -0.62584 0.84758 (fixed)
## time:3 0.66289 1.23518 C:time:3 0.66289 1.23518 (fixed)
## time:4 -0.67604 1.49986 C:time:4 -0.67604 1.49986 (fixed)
## time:5 -0.13542 1.69741 C:time:5 -0.13542 1.69741 (fixed)
## time:6 -31.22411 122955.34396 C:time:6 -31.22411 122955.34396 (fixed)
## B:(Intercept) -1.23793 2.07832
##
## Population Size Estimate (se): 21.6967 (13.1183)
## 95% confidence interval for population size: 17.26 to 101.55
## Individuals observed: 17
## Effective sample size: 102
##
## Message = SUCCESS: Convergence criterion met
## Number of estimable coefficients (estimated) = 6
## Log likelihood = -37.3456782098957
## Deviance = 74.6913564197915
## AIC = 86.6913564197915
## AICc = 87.5755669461072
```

```
round(Mtb$phat, digits=3)
```

```
## [1,] [2,] [3,] [4,] [5,] [6,]
## [1,] 0.277 0.17 0.426 0.163 0.25 0
## [2,] 0.277 0.17 0.426 0.163 0.25 0
## [3,] 0.277 0.17 0.426 0.163 0.25 0
## [4,] 0.277 0.17 0.426 0.163 0.25 0
## [5,] 0.277 0.17 0.426 0.163 0.25 0
## [6,] 0.277 0.17 0.426 0.163 0.25 0
## [7,] 0.277 0.17 0.426 0.163 0.25 0
## [8,] 0.277 0.17 0.426 0.163 0.25 0
## [9,] 0.277 0.17 0.426 0.163 0.25 0
## [10,] 0.277 0.17 0.426 0.163 0.25 0
## [11,] 0.277 0.17 0.426 0.163 0.25 0
## [12,] 0.277 0.17 0.426 0.163 0.25 0
## [13,] 0.277 0.17 0.426 0.163 0.25 0
## [14,] 0.277 0.17 0.426 0.163 0.25 0
## [15,] 0.277 0.17 0.426 0.163 0.25 0
## [16,] 0.277 0.17 0.426 0.163 0.25 0
## [17,] 0.277 0.17 0.426 0.163 0.25 0
```

```
round(Mtb$Sc.hat, digits=3)
```

```
## [1,] [2,] [3,] [4,] [5,] [6,]
## [1,] NA 0.056 0.177 0.053 0.088 0
## [2,] NA 0.056 0.177 0.053 0.088 0
## [3,] NA 0.056 0.177 0.053 0.088 0
## [4,] NA 0.056 0.177 0.053 0.088 0
## [5,] NA 0.056 0.177 0.053 0.088 0
## [6,] NA 0.056 0.177 0.053 0.088 0
## [7,] NA 0.056 0.177 0.053 0.088 0
## [8,] NA 0.056 0.177 0.053 0.088 0
## [9,] NA 0.056 0.177 0.053 0.088 0
## [10,] NA 0.056 0.177 0.053 0.088 0
## [11,] NA 0.056 0.177 0.053 0.088 0
## [12,] NA 0.056 0.177 0.053 0.088 0
## [13,] NA 0.056 0.177 0.053 0.088 0
## [14,] NA 0.056 0.177 0.053 0.088 0
## [15,] NA 0.056 0.177 0.053 0.088 0
## [16,] NA 0.056 0.177 0.053 0.088 0
## [17,] NA 0.056 0.177 0.053 0.088 0
```

```
# create HTML table using kableExtra
library(kableExtra)
# create HTML table using kableExtra
library(kableExtra)
require(knitr)

Model<-c("M0", "Mt", "Mb", "Mtb")
Model_Description<-c("The most basic model in which p and c are constant", "p differs among sampling occasions and pt = ct", "Behavioral response model in which p and c differ.", "Combination of models Mt and Mb")
N<-c(41.03, 37.79,18.58, 21.76))
SE<-c(16.66, 14.77, 2.15, 13.12))
AICc<-c(89.91, 85.47, 87.34, 87.58))
tab<-data.frame(Model,Model_Description,N,SE,AICc)
```

```
#table
options(knitr.kable.NA = "") # leave NA cells empty
knitr::kable(tab, digits = 3,booktabs=T, align="c",
caption = "<center><strong>Table 1. Results for mark-recapture analysis for stink pots.</strong></center>")%>%
kable_styling(bootstrap_options = c("striped", "hover", "condensed", "bordered"))
```

Table 1. Results for mark-recapture analysis for stink pots.				
Model	Model_Description	N	SE	AICc
M0	The most basic model in which p and c are constant	41.03	16.66	89.91
Mt	p differs among sampling occasions and pt = ct	37.79	14.77	85.47
Mb	Behavioral response model in which p and c differ.	18.58	2.15	87.34
Mtb	Combination of models Mt and Mb	21.70	13.12	87.58

```
Time<-c(1, 2, 3, 4, 5, 6))
```

```
pt<-c(0.159, 0.079, 0.185, 0.053 ,0.079,0))
```

```
pdata<-data.frame(Time,pt)
```

```
require(ggplot2)
```

```
## Loading required package: ggplot2
```

```
ggplot(pdata, aes(x=Time, y=pt)) + geom_point() + geom_line()+
scale_x_continuous("Time interval", breaks =c(1,2,3,4,5,6)) +
scale_y_continuous("Capture probability (pt)") +
theme_classic() +
ggtitle(label = "Fig 1. The capture probability of stinkpots over the season.") +
theme(plot.title = element_text(face = "bold", hjust=0.5), plot.subtitle=element_text(hjust=0.5))
```

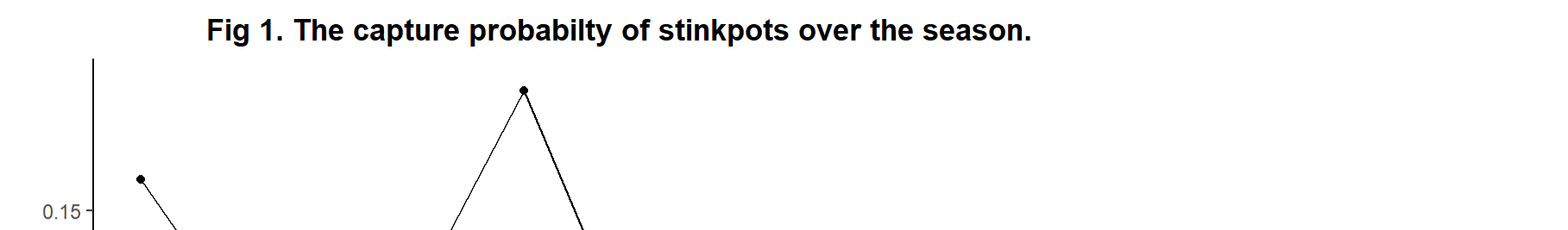


Fig 1. The capture probability of stinkpots over the season.

Question 4

Model M_1 has the lowest AICc value. Model M_1 does not measure behavioural responses but model M_0 and M_0 both do. Model M_0 suggests that recapture is relatively low (0.06) suggesting a trap shy strategy, and model M_0 shows that recapture varies depending on the trapping interval. For interval 2 there is a low recapture (0.05), interval 3 there is a higher recapture rate (0.17), and interval 4 has a lower recapture (0.08). This suggests for this model that there is a trap shy behavior after the first time an individual is caught and marked, a trap happy-ish response for interval 3, and then for interval 4 and 5 trap shyness.

Question 5

Model M_0 was the worst model because captures and recaptures varied overtime and this model was unable to account for this variation since it assumed both were constant over time.