

SUPPLEMENTARY ELECTRONIC MATERIAL

ESTIMATING THE ABUNDANCE AND HABITAT
SELECTION OF CONSERVATION PRIORITY MARSH-
DWELLING PASSERINES WITH A DOUBLE-OBSERVER
APPROACH

ESTIMA DE LA ABUNDANCIA Y SELECCIÓN DE HÁBITAT DE
PASERIFORMES PALUSTRES DE CONSERVACIÓN
PRIORITARITARIA A TRAVÉS DE UNA APROXIMACIÓN DE
DOBLE OBSERVADOR

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Simulating data and WinBUGS analysis.
[Simulación de datos y análisis con WinBUGS.]

```
library("R2WinBUGS")
```

```
setwd('.')
```

```
# We simulate data for double observer
```

```
nSites <- 750
```

```
psi <- runif(1, 0.05, 0.1)
```

```
p1 <- 0.4
```

```
p2 <- 0.7
```

```
cp <- c(p1*(1-p2), p2*(1-p1), p1*p2)
```

```
set.seed(9023)
```

```
N <- rpois(nSites, psi)
```

```
y <- matrix(NA, nSites, 3)
```

```
for(i in 1:nSites) {
```

```
  y[i,] <- rmultinom(1, N[i], c(cp, 1-sum(cp)))[1:3]
```

```
}
```

```
x <- apply(y, 2, sum)
```

```
x01 <- x[1]
```

```
x10 <- x[2]
```

```
x11 <- x[3]
```

```
# This code (Royle & Dorazio, 2008) creates the multinomial trial representation
```

```

# of the data

ncap2<-matrix(0,nrow=sum(x),ncol=4)

rid<-c(1:x[1], (x[1]+1):(x[1]+x[2]),(x[1]+x[2]+1):nrow(ncap2))

cid<-c( rep(1,x[1]),rep(2,x[2]),rep(3,x[3]) )

ncap2[cbind(rid,cid)]<-1

# We augment the encounter histories with histories of the last type
# which are those not seen by either observer (0,0,0,1).

nz<-nSites-dim(ncap2)[1]

ncap<-rbind(ncap2,matrix(rep(c(0,0,0,1),nz),ncol=4,byrow=TRUE))

nind<-dim(ncap2)[1]

# Real population size:

cat("Real population", sum(N), "individuals", "\n")

# Naive estimation:

cat("We detect", nind, "individuals", "\n")

sink("model.txt")

cat("

model {

# Priors

psi~dunif(0,1)

```

```

tau~dunif(0,10)

mu1.p~dunif(-5,5)

mu2.p~dunif(-5,5)

# First nind observations
for(i in 1:nind){
  z[i]~dbin(psi,1)
}

# Data augmentation
for(i in (nind+1):(nind+nz)){
  z[i]~dbin(psi,1)
}

for(i in 1:(nind+nz)){
  e[i]~dnorm(0,tau)I(-5,5)

  # Logit for probability covariates
  logit(p1[i])<- mu1.p + e[i]
  logit(p2[i])<- mu2.p + e[i]
  cp1[i]<- (1-p2[i])*p1[i]
  cp2[i]<- p2[i]*(1-p1[i])
  cp3[i]<- p1[i]*p2[i]
  cp4[i]<- (1-p1[i])*(1-p2[i])
  mu[i,1]<-z[i]*cp1[i]
  mu[i,2]<-z[i]*cp2[i]
  mu[i,3]<-z[i]*cp3[i]
  mu[i,4]<-z[i]*cp4[i] + (1-z[i])

```

```

ncap[i,1:4]~dmulti(mu[i,1:4],1)
}
# We transform back to obtain observation probabilities
logit(p1bar)<-mu1.p
logit(p2bar)<-mu2.p
Nind<-sum(z[1:(nind+nz)])
}
",fill=TRUE)
sink()

data<-list("ncap"=ncap, "nind"=nind, "nz"=nz)
zst<-c(rep(1, nind),rbinom(nz, 1, 0.5))
inits<-function(){
  list(mu1.p=0, mu2.p=0, z=zst)
}
parameters <- c('Nind', 'mu1.p', 'mu2.p', 'p1bar', 'p2bar', 'psi')

ni=5000
nb=2500
nthin=1
nc=3

library(R2WinBUGS)
out <- bugs(data, inits, parameters, "model.txt", n.thin=nthin,n.chains=nc,
  n.burnin=nb,n.iter=ni, debug=TRUE)
print(out, dig=3)

```

```

estima_moda <- function(x) {
  d <- density(x)
  d$x[which.max(d$y)]
}

moda<-estima_moda(out$sims.list$N); moda

hist(out$sims.list$Nind, col="lightgray", border="white", main="",
  xlab="Number of individuals", ylab="Frecuency", xlim=c(20,170), breaks=20)

abline(v=sum(N), lwd=2, col='red')
abline(v=moda, lwd=2, lty=2, col='blue')
abline(v=nind, lwd=2, lty=2, col='darkgreen')

legend('topright',c("Real population", "Estimation (Mode)", "Naïve") , lwd=2,
  lty=c(1,2,2), col=c('red', 'blue','darkgreen' ), bty='n', cex=0.9)

```