

Overview of Unmarked: An R Package for the Analysis of Wildlife Data

Ian Fiske and Richard Chandler

February 25, 2010

Abstract

Unmarked aims to be a complete environment for the statistical analysis of wildlife data. Currently, the focus is on 2-level hierarchical models that separately model a latent state and an observation process. Unmarked uses S4 classes to help the user both explore their data and analyze in a transparent manner.

1 Overview of unmarked

Occupancy and abundance data are often associated with metadata related to the design of the study. For example, in distance sampling, the study design (line- or point-transect), distance class break points, transect lengths, and units of measurement need to be accounted for in the analysis. Unmarked uses S4 classes to store data and metadata in a way that allows for easy data manipulation, summarization, and model specification. Table 1 lists the currently implemented models and their associated fitting functions and data classes.

Each data class can be created with a call to the constructor function of the same name as described in the examples below.

2 Typical unmarked session

The first step is to import the data into R. This can be accomplished with either a call to the appropriate type of `unmarkedFrame`:

```
> library(unmarked)
> wt <- read.csv(system.file("csv", "widewt.csv", package = "unmarked"))
> head(wt)
  site y.1 y.2 y.3      elev      forest   length   date.1
1    1  0  0  0 -1.1729446 -1.156228147 1.824549 -1.761481
2    2  0  0  0 -1.1265010 -0.501483710 1.629241 -2.904339
3    3  0  0  0 -0.1976283 -0.101362109 1.458615 -1.690053
4    4  0  0  0 -0.1047411  0.007761963 1.686399 -2.190053
5    5  0  0  0 -1.0336137 -1.192602838 1.280934 -1.832910
6    6  0  0  0 -0.8478392  0.917129237 1.808289 -2.618624
      date.2   date.3   ivel.1   ivel.2   ivel.3
1  0.3099471 1.3813757 -0.5060353 -0.5060353 -0.5060353
2 -1.0471958 0.5956614 -0.9336151 -0.9907486 -1.1621491
3 -0.4757672 1.4528042 -1.1355754 -1.3388644 -1.6099164
```

Model	Fitting Function	Data	Citation
Occupancy	occu	unmarkedFrameOccu	[2]
Royle-Nichols	occuRN	unmarkedFrameOccu	[5]
Point Count	pcount	unmarkedFramePCount	[4]
Distance-sampling	distsamp	unmarkedFrameDS	[6]
Arbitrary multinomial-Poisson	multinomPois	unmarkedFrameMPois	[3]
Colonization-extinction	colect	unmarkedMultFrame	[1]

Table 1: Models handled by unmarked.

```

4 -0.6900529 1.2385185 -0.8193481 -0.9272669 -1.1970640
5 0.1670899 1.3813757 0.6375563 0.8803737 1.0422520
6 0.1670899 1.3813757 -1.3288666 -1.0422624 -0.8989603
> y <- wt[, 2:4]
> siteCovs <- wt[, c("elev", "forest", "length")]
> obsCovs <- reshape(wt[, c("date.1", "date.2", "date.3",
    "ivel.1", "ivel.2", "ivel.3")], varying = 1:6, direction = "long")
> obsCovs <- obsCovs[order(obsCovs$id, obsCovs$time), c(2:3)]
> wt <- unmarkedFrameOccu(y = y, siteCovs = siteCovs, obsCovs = obsCovs)
> summary(wt)
unmarkedFrame Object

```

```

237 sites
Maximum number of observations per site: 3
Mean number of observations per site: 2.81
Sites with at least one detection: 79

```

Tabulation of y observations:

```

0    1 <NA>
483 182  46

```

Site-level covariates:

elev	forest	length
Min. : -1.436125	Min. : -1.265e+00	Min. : 0.1823
1st Qu.: -0.940726	1st Qu.: -9.744e-01	1st Qu.: 1.4351
Median : -0.166666	Median : -6.499e-02	Median : 1.6094
Mean : 0.007612	Mean : 8.798e-05	Mean : 1.5924
3rd Qu.: 0.994425	3rd Qu.: 8.080e-01	3rd Qu.: 1.7750
Max. : 2.434177	Max. : 2.299e+00	Max. : 2.2407

Observation-level covariates:

date	ivel
Min. : -2.9043386	Min. : -1.753e+00
1st Qu.: -1.1186243	1st Qu.: -6.660e-01
Median : -0.1186243	Median : -1.395e-01
Mean : -0.0002173	Mean : -3.008e-11
3rd Qu.: 1.3099471	3rd Qu.: 5.493e-01
Max. : 3.8099471	Max. : 5.980e+00
NA's : 42.0000000	NA's : 4.600e+01

or by using the convenience function csvToUMF:

```

> wt <- csvToUMF(system.file("csv", "widewt.csv", package = "unmarked"),
    long = FALSE, type = "unmarkedFrameOccu")

```

If not all sites have the same numbers of observations, then manual importation of data in long format can be tricky. csvToUMF seamlessly handles this situation.

```

> pcru <- csvToUMF(system.file("csv", "frog2001pcru.csv",
    package = "unmarked"), long = TRUE, type = "unmarkedFrameOccu")
> summary(pcr)
unmarkedFrame Object

```

```

130 sites
Maximum number of observations per site: 3
Mean number of observations per site: 2.59
Sites with at least one detection: 96

```

Tabulation of y observations:

```

0    1    2    3 <NA>
197  25  28  87  53

```

Observation-level covariates:

MinAfterSunset	Wind	Sky	Temperature
Min. : -21.00	Min. : 0.0000	Min. : 0.0000	Min. : 4.00
1st Qu.: 66.00	1st Qu.: 0.0000	1st Qu.: 0.0000	1st Qu.: 13.00
Median : 97.00	Median : 1.0000	Median : 0.0000	Median : 17.50
Mean : 97.57	Mean : 0.8813	Mean : 0.4837	Mean : 16.61
3rd Qu.: 126.00	3rd Qu.: 2.0000	3rd Qu.: 1.0000	3rd Qu.: 20.60
Max. : 228.00	Max. : 3.0000	Max. : 5.0000	Max. : 28.00
NA's : 53.00	NA's : 53.0000	NA's : 53.0000	NA's : 53.00

JulianDate
Min. : 72.0
1st Qu.: 95.0
Median : 123.0
Mean : 127.4
3rd Qu.: 159.0
Max. : 179.0
NA's : 53.0

To help stabilize the numerical optimization algorithm, we recommend standardizing the covariates.

```
> obsCovs(pcr) <- scale(obsCovs(pcr))
```

Occupancy models can then be fit with the `occu()` function:

```
> fm1 <- occu(~1 ~ 1, pcr)
> fm2 <- occu(~MinAfterSunset + Temperature ~ 1, pcr)
> summary(fm1)
```

Call:

```
occu(formula = ~1 ~ 1, data = pcr)
```

Occupancy (logit-scale):

Estimate	SE	z	P(> z)
2.95	1.44	2.05	0.04

Detection (logit-scale):

Estimate	SE	z	P(> z)
-0.249	0.170	-1.47	0.142

AIC: 461.0986

Sample size: 130

optim convergence code: 0

optim iterations: 22

```
> summary(fm2)
```

Call:

```
occu(formula = ~MinAfterSunset + Temperature ~ 1, data = pcr)
```

Occupancy (logit-scale):

Estimate	SE	z	P(> z)
1.54	0.292	5.26	1.42e-07

Detection (logit-scale):

	Estimate	SE	z	P(> z)
(Intercept)	0.2098	0.206	1.017	3.09e-01
MinAfterSunset	-0.0855	0.160	-0.536	5.92e-01
Temperature	-1.8936	0.291	-6.508	7.60e-11

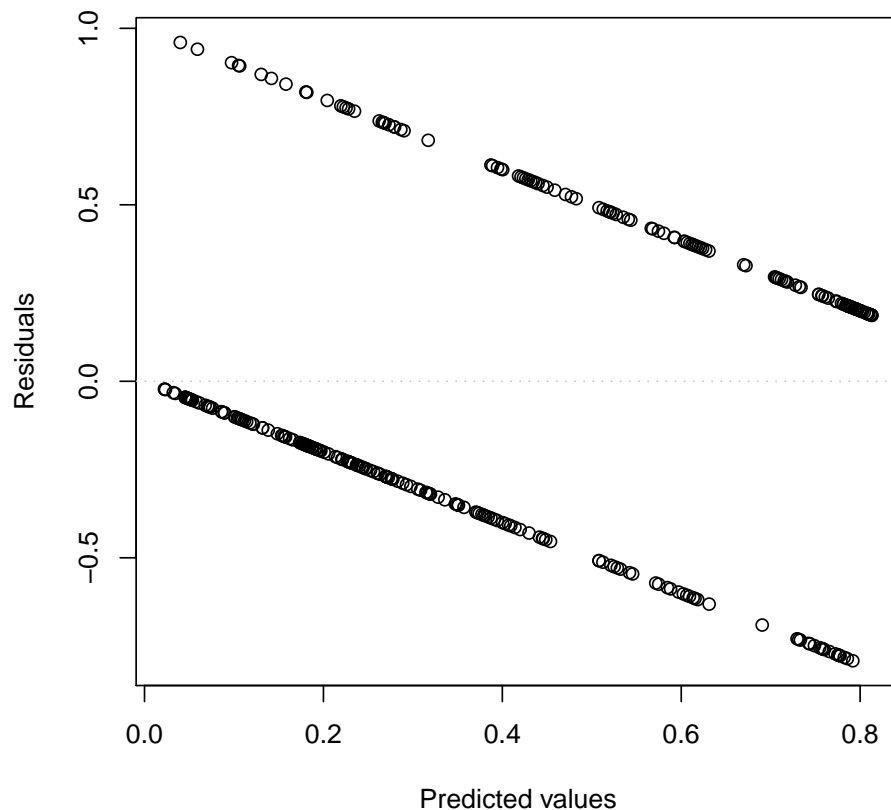
AIC: 357.0791

Sample size: 130

optim convergence code: 0

optim iterations: 21

```
> plot(fm2)
```



Here, we have specified that the detection process is modeled with the MinAfterSunset and Temperature covariates. No covariates are specified for occupancy here. See ?occu for more details.

Unmarked fitting functions return unmarkedFit objects which can be queried to investigate the model fit. Variables can be back-transformed to the unconstrained scale using backTransform. Standard errors are computed using the delta method.

```
> backTransform(fm2, "state")
Backtransformed linear combination(s) of Occupancy estimate(s)
```

Estimate	SE LinComb	(Intercept)
0.823	0.0425	1.54

Transformation: logistic

Because the detection component was modeled with covariates, covariate coefficients must be specified to back-transform. Here, we request the probability of detection given a site is occupied and all covariates are set to 0.

```
> backTransform(linearComb(fm2, coefficients = c(1, 0,
0), type = "det"))
Backtransformed linear combination(s) of Detection estimate(s)
```

Estimate	SE LinComb	(Intercept)	MinAfterSunset	Temperature
0.552	0.051	0.210	1	0

Transformation: logistic

A predict method also exists.

```
> newData <- data.frame(MinAfterSunset = 0, Temperature = -2:2)
> head(predict(fm2, type = "det", newdata = newData, appendData = TRUE))
      Predicted      SE MinAfterSunset Temperature
1 0.98196076 0.01266193           0          -2
2 0.89123189 0.04248804           0          -1
3 0.55225129 0.05102660           0           0
4 0.15658708 0.03298276           0           1
5 0.02718682 0.01326263           0           2
```

Confidence intervals are requested with `confint`, using either the asymptotic normal approximation or profiling.

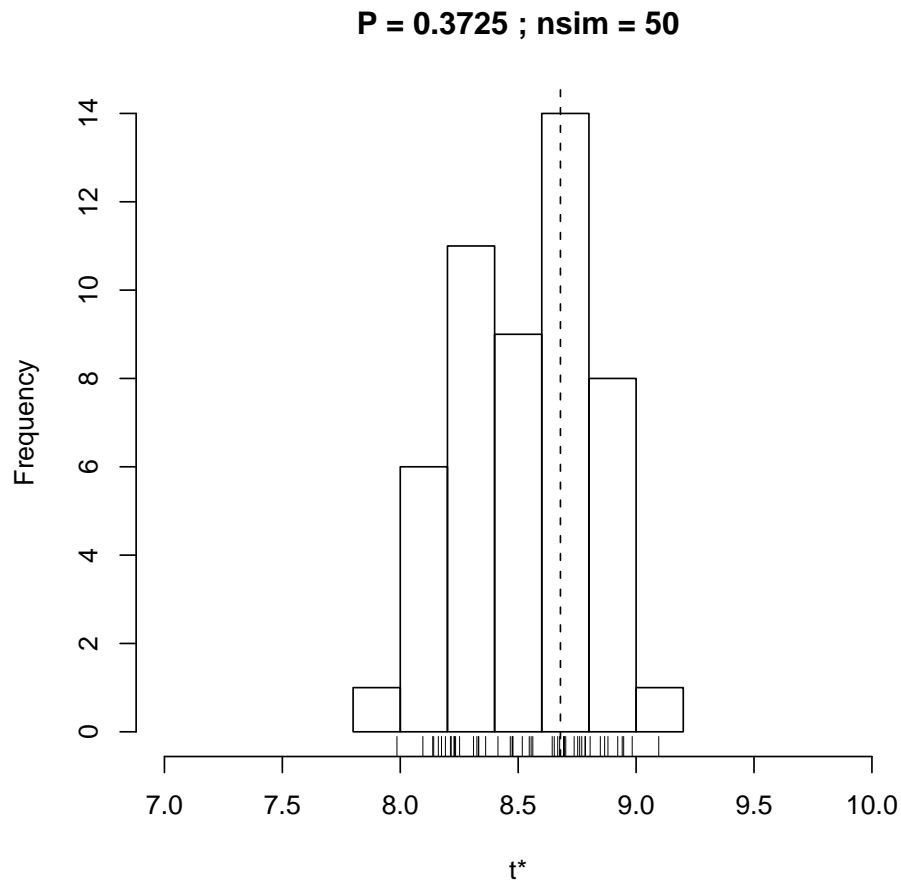
```
> confint(fm2, type = "det")
              0.025      0.975
(Intercept)  -0.1946872  0.6142292
MinAfterSunset -0.3985642  0.2274722
Temperature   -2.4638797 -1.3233511
> confint(fm2, type = "det", method = "profile")
Profiling parameter 1 of 3 ... done.
Profiling parameter 2 of 3 ... done.
Profiling parameter 3 of 3 ... done.
              0.025      0.975
p(Int)          -0.1929210  0.6208837
p(MinAfterSunset) -0.4044794  0.2244221
p(Temperature)   -2.5189984 -1.3789261
```

Model selection and multi-model inference can be implemented after organizing models using the `fitList` function.

```
> fms <- fitList(NULL = fm1, TimeTemp = fm2)
> modSel(fms, nullmod = fm1)
      n nPars    AIC deltaAIC    AICwt    Rsq cumltvAICwt
TimeTemp 130     4 357.08     0.00 1.0000e+00 0.58243      1
Null      130     2 461.10    104.02 2.5849e-23 0.00000      1
> head(predict(fms, type = "det", newdata = newData, appendData = TRUE))
      Predicted      SE MinAfterSunset Temperature
1 0.98196076 0.01266193           0          -2
2 0.89123189 0.04248804           0          -1
3 0.55225129 0.05102660           0           0
4 0.15658708 0.03298276           0           1
5 0.02718682 0.01326263           0           2
```

Parametric bootstrapping can be used to check the adequacy of model fit.

```
> pcru.pb <- parboot(fm2, nsim = 50, report = 5)
t0 = 8.679173
8.6, 8.7, 8.4, 9, 8
8.8, 8.7, 8.4, 8, 8.5
8.5, 8.4, 8.9, 8.5, 9.1
8.7, 8.6, 8.3, 8.8, 8.4
8.3, 8.7, 8.9, 9.1, 8.5
8.8, 8.6, 8.4, 8.9, 8.8
8.4, 8.7, 8.8, 8.4, 8.1
8.2, 8.6, 8.5, 8.8, 8.4
8.9, 9.1, 8, 8.6, 8.3
8.7, 8.6, 8.7, 8.2, 8.4
> plot(pcr.pb)
```



This example suggests an adequate fit.

References

- [1] Darryl~I. MacKenzie, James~D. Nichols, James~E. Hines, Melinda~G. Knutson, and Alan~B. Franklin. Estimating site occupancy, colonization, and local extinction when a species is detected imperfectly. *Ecology*, 84(8):2200–2207, 2003.
- [2] Darryl~I. MacKenzie, James~D. Nichols, G.~B. Lachman, S.~Droege, J.~A. Royle, and C.~A. Langtimm. Estimating site occupancy rates when detection probabilities are less than one. *Ecology*, 83(8):2248–2255, 2002.
- [3] J.~A. Royle. Generalized estimators of avian abundance from count survey data. *Animal Biodiversity and Conservation*, 27(1):375–386, 2004.
- [4] J.~A Royle. N-mixture models for estimating population size from spatially replicated counts. *Biometrics*, 60(1):108–115, 2004.
- [5] J.~A. Royle and J.~D. Nichols. Estimating abundance from repeated presence-absence data or point counts. *Ecology*, 84(3):777–790, 2003.
- [6] JA Royle, DK Dawson, and S.~Bates. Modeling abundance effects in distance sampling. *Ecology*, 85(6):1591–1597, 2004.