Modeling Abundance in Relation to Covariates

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Introduction

This tutorial builds on the introductory material presented in the **Rdistance Tutorial for Beginners**. The goal of a distance-sampling study is often to relate abundance to some other variable, e.g., a covariate describing the habitat at each survey site. Rdistance can provide you with site-level estimates of abundance that are corrected for imperfect detection. You can then use these estimates as the response variable in one of the many modeling frameworks available in Program R that you're likely already familiar with (e.g., lm, glm, lme4, etc.). Here, we make use of the example datasets already contained within Rdistance (i.e., line transect surveys of sparrows), to demonstrate an example modeling workflow. This tutorial is current as of version 1.3.2 of Rdistance.

Estimate abundance at each site

The process of generating an overall, study-area estimate of abundance is outlined in the **Rdistance Tutorial** for **Beginners**. Use the by.id argument (by.id=TRUE) within F.abund.estim or F.automated.CDA to estimate abundance for each site surveyed.

```
## Likelihood
                Series
                        Expans
                               Converged? Scale? AIC
                                    3263.4429
## halfnorm cosine 0
                        Yes
                                0k
## halfnorm cosine 1
                                0k
                                    3261.585
                        Yes
## halfnorm hermite 1
                        Yes
                                0k 3262.8066
## halfnorm simple 1
                                    3262.1526
                        Yes
                                Πk
## halfnorm cosine
                        Yes
                                0k
                                    3263.3155
## halfnorm hermite 2
                                Ok
                                    3263.593
                        Yes
## halfnorm simple 2
                        Yes
                                0k
                                    3263.4741
                                    3265.1285
## halfnorm cosine
                        Yes
                                0k
## halfnorm hermite 3
                        Yes
                                0k
                                    3265.5499
## halfnorm simple 3
                                Ok
                                    3265.216
                        Yes
## hazrate
                cosine
                            Yes
                                    0k 3267.6246
                                        3263.3092
## hazrate
                cosine
                            Yes
                                    0k
                       1
                hermite 1
## hazrate
                            No
                                    NΑ
                                        NΑ
## hazrate
                simple 1
                            No
                                    NA NA
                                    0k 3265.0704
## hazrate
                cosine 2
                            Yes
## hazrate
               hermite 2
                            Yes
                                    Ok 3268.9682
## hazrate
                simple 2
                            Yes
                                    0k 3265.5796
                cosine 3
## hazrate
                            Yes
                                    0k 3267.117
```

```
## hazrate
              hermite 3
                          Bad
                                 NA NA
## hazrate
                                 NA NA
              simple 3
                         No
## uniform
              cosine 0
                         Yes
                                 0k 3260.7318
## uniform
              cosine 1
                         Bad
                                 NA NA
## uniform
              hermite 1
                         Yes
                                 0k 3262.736
                                 Ok 3262.2602
## uniform
              simple 1
                         Yes
## uniform
              cosine 2
                         Bad
                                 NA NA
## uniform
              hermite 2
                         No
                                 NA NA
## uniform
              simple 2
                         Bad
                                 NA NA
## uniform
              cosine 3
                         Bad
                                 NA NA
## uniform
              hermite 3
                         No
                                 NA NA
## uniform
                                 NA NA
              simple 3
                         Bad
                                 0k 3263.8153
## negexp
              cosine 0
                         Yes
## negexp
                                 Ok 3260.0985
              cosine 1
                         Yes
## negexp
                         Yes
                                 Ok 3260.3108
              hermite 1
## negexp
              simple 1
                         Yes
                                 0k 3261.0097
                                 0k 3262.0306
## negexp
              cosine 2
                         Yes
## negexp
              hermite 2
                         No
                                 NA NA
                                 Ok 3262.2878
## negexp
              simple 2
                         Yes
## negexp
              cosine 3
                         Yes
                                 0k 3263.4898
## negexp
              hermite 3
                         No
                                 NA NA
## negexp
              simple 3
                         Yes
                                 0k 3264.2055
## Gamma
                             Ok 3586.8919
                  0
                      Yes
## Note: Some models did not converge or had parameters at their boundaries.
## Computing bootstrap confidence interval on \mathbb{N}\dots
##
## ----- Final Automated CDS Abundance Estimate ------ Final Automated CDS Abundance
## Call: F.dfunc.estim(dist = dist, likelihood = fit.table$like[1], w.lo = w.lo, w.hi = w.hi, expan
##
## Coefficients:
##
         Beta
  0.02754839 -0.26542176
##
## Convergence: Success
## Function: NEGEXP with 1 expansion(s) of COSINE series
## Strip: 0 to 150
## Effective strip width: 44.72749
## Scaling: g(0) = 1
## Log likelihood: 1628.032
## AIC: 3260.098
## Abundance estimate: 1.152038; 95% CI=(0.7236135 to 1.411445)
```

Because by.id=TRUE, a data.frame of the site-level abundances is stored in the auto object under the name nhat.df:

```
head(auto$nhat.df)
```

```
## siteID rawcount nhat
## 1 A1 15 3.3536424
## 2 A2 14 3.1300662
```

```
## 3 A3 10 2.2357616
## 4 A4 15 3.3536424
## 5 B1 1 0.2235762
## 6 B2 0 0.0000000
```

Because distances were measured in meters and area=10000, abundance (density) estimates are given as the number of sparrows per hectare.

Model abundance estimates

The original transect dataset contained a variable named **sagemean** that represents the average sagebrush cover at each transect. Suppose we want to examine the relationship between sagebrush cover and sparrow abundance. First, merge the new data.frame of site-level abundance estimates back to the original transect data.frame that includes **sagemean**.

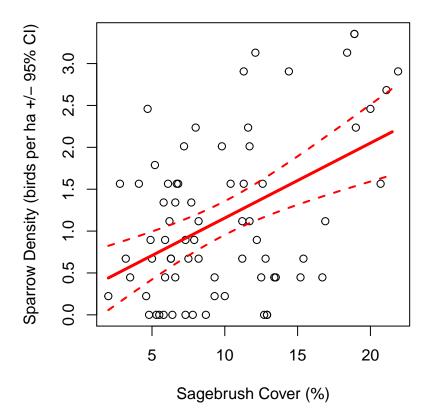
```
mydata <- merge(sparrow.transects, auto$nhat.df, by="siteID")
head(mydata)</pre>
```

```
##
     siteID length observer sagemean sage rawcount
                                                           nhat
## 1
         Α1
               500
                        obs4
                                 18.9 High
                                                   15 3.3536424
## 2
         A2
               500
                        obs4
                                  18.4 High
                                                   14 3.1300662
               500
         ΑЗ
                                                   10 2.2357616
## 3
                        obs5
                                  19.0 High
               500
## 4
         Α4
                        obs5
                                  18.9 High
                                                   15 3.3536424
               500
## 5
         В1
                        obs3
                                   4.6 Low
                                                    1 0.2235762
## 6
         B2
               500
                        obs3
                                   4.8 Low
                                                    0 0.0000000
```

We now have both variables in the same data.frame. For the sake of demonstration, we will fit a simple linear regression model using the 1m function to assess the relationship between sagebrush cover and sparrow density (per ha). We see that sparrow density increases with sagebrush cover, and density nearly quadruples across the range of sagebrush cover surveyed.

```
mod <- lm(nhat ~ sagemean, data=mydata)
summary(mod)</pre>
```

```
##
## lm(formula = nhat ~ sagemean, data = mydata)
##
## Residuals:
##
      Min
               1Q Median
                                3Q
                                       Max
## -1.4157 -0.7356 -0.1005 0.6878 1.7860
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                0.2611
                            0.2286
                                     1.142
                                              0.257
                            0.0206
                 0.0895
                                     4.344 4.64e-05 ***
## sagemean
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.8569 on 70 degrees of freedom
## Multiple R-squared: 0.2123, Adjusted R-squared:
## F-statistic: 18.87 on 1 and 70 DF, p-value: 4.639e-05
```



Conclusion

By using Rdistance to estimate site-level abundance, you can account for imperfect detection, and use the modeling framework you choose to then relate site-level covariates to abundance. We demonstrated this workflow using Rdistance's F.automated.CDA function to fit a detection function and provide abundance estimates for each site, and the standard lm function to model the relationship between a habitat covariate and sparrow density.