## partial RDA

## Pablo E. Gutiérrez-Fonseca

## 11/17/2021

```
library(ggplot2)
library(dplyr)
library(vegan)
```

Primer paso: cargar las librerias que necesitas.

```
species=read.csv("data/RDA_species.csv", header=T, row.names=NULL, sep=",")
env=read.csv("data/RDA_environmetal_standart.csv", header=T, row.names=NULL, sep=",")
```

Segundo paso: cargar los datos.

```
species_1 <- select(species, -site)
env_1 <- select(env, -site)</pre>
```

Remover la columna de sitos.

**Transformar datos.** Hellinger es una transformacion recomendada por Legendre & Callagher (2001) en datos de abundancia y con una respuesta lineal

```
species_2 <- decostand(species_1, method = "hellinger")</pre>
```

Vamos a construir los modelos: combinados e individuales

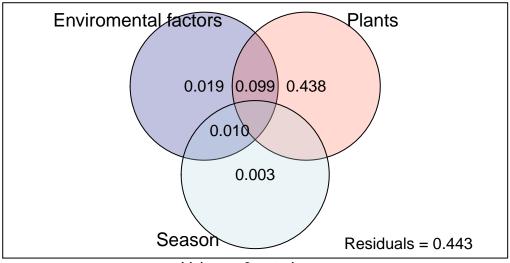
```
RsquareAdj (partial_rda_all)
Ver los R2adj de cada uno
## $r.squared
## [1] 0.6423136
## $adj.r.squared
## [1] 0.5567799
RsquareAdj (partial_rda_env)
## $r.squared
## [1] 0.1834995
## $adj.r.squared
## [1] 0.1218768
RsquareAdj(partial_rda_phy)
## $r.squared
## [1] 0.5749283
## $adj.r.squared
## [1] 0.5249199
RsquareAdj(partial_rda_season)
## $r.squared
## [1] 0.01787828
## $adj.r.squared
## [1] 0.0003403971
varp <- varpart(species_2, ~ temperature + oxygen + pH + conductivity,</pre>
                ~ plants + land_use + margin, ~season, data = env_1)
varp
Vamos a correr un RDA con las matrices (diferentes factores) separadas.
##
## Partition of variance in RDA
## Call: varpart(Y = species_2, X = ~temperature + oxygen + pH +
## conductivity, ~plants + land_use + margin, ~season, data = env_1)
```

##

```
## Explanatory tables:
        ~temperature + oxygen + pH + conductivity
## X1:
        ~plants + land use + margin
## X3:
        ~season
## No. of explanatory tables: 3
## Total variation (SS): 34.967
               Variance: 0.61346
##
## No. of observations: 58
##
## Partition table:
                          Df R.square Adj.R.square Testable
##
                                            0.12188
## [a+d+f+g] = X1
                              0.18350
                                                        TRUE
## [b+d+e+g] = X2
                              0.57493
                                                        TRUE
                                            0.52492
## [c+e+f+g] = X3
                              0.01788
                                            0.00034
                                                        TRUE
                           1
## [a+b+d+e+f+g] = X1+X2 10
                              0.63213
                                            0.55386
                                                        TRUE
## [a+c+d+e+f+g] = X1+X3
                          5
                              0.19602
                                            0.11871
                                                        TRUE
## [b+c+d+e+f+g] = X2+X3
                          7
                              0.59442
                                            0.53764
                                                        TRUE
## [a+b+c+d+e+f+g] = All 11
                                                        TRUE
                              0.64231
                                            0.55678
## Individual fractions
## [a] = X1 | X2+X3
                           4
                                            0.01914
                                                        TRUE
## [b] = X2 | X1+X3
                           6
                                            0.43807
                                                        TRUE
## [c] = X3 | X1+X2
                                                        TRUE
                           1
                                            0.00292
## [d]
                           0
                                                       FALSE
                                            0.09923
## [e]
                           0
                                           -0.00608
                                                       FALSE
## [f]
                           0
                                            0.00980
                                                       FALSE
## [g]
                           0
                                           -0.00630
                                                       FALSE
## [h] = Residuals
                                            0.44322
                                                       FALSE
## Controlling 1 table X
## [a+d] = X1 | X3
                                            0.11837
                                                        TRUE
## [a+f] = X1 | X2
                           4
                                            0.02894
                                                        TRUE
## [b+d] = X2 | X3
                           6
                                            0.53730
                                                        TRUE
                           6
## [b+e] = X2 | X1
                                            0.43198
                                                        TRUE
## [c+e] = X3 | X1
                                                        TRUE
                           1
                                           -0.00316
## [c+f] = X3 | X2
                                            0.01272
                                                        TRUE
## ---
## Use function 'rda' to test significance of fractions of interest
```

Legendre (2008) [doi: 10.1093/jpe/rtm001] argued that "Negative values of Ra2 are interpreted as zeros; they correspond to cases where the explanatory variables explain less variation than random normal variables would."

**Plot the results** The plot shows the adjusted R2 values associated with each partition or for overlapping partitions.



Values <0 not shown

```
## Show values for all partitions by putting 'cutoff' low enough: #plot(varp, cutoff = -Inf, cex = 0.7, bg=2:5)
```

Then we can test the significance of each individual component.

```
## Permutation test for rda under reduced model
## Permutation: free
## Number of permutations: 999
##
## Model: rda(formula = species_2 ~ temperature + oxygen + pH + conductivity + Condition(plants + land_*
## Df Variance F Pr(>F)
## Model 4 0.02938 1.5398 0.027 *
## Residual 46 0.21943
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
# Significance of partition from Physical characteristics
anova(rda(species_2 ~ plants + land_use + margin +
          Condition(temperature + oxygen + pH + conductivity) +
          Condition(season), data=env_1))
## Permutation test for rda under reduced model
## Permutation: free
## Number of permutations: 999
##
## Model: rda(formula = species_2 ~ plants + land_use + margin + Condition(temperature + oxygen + pH +
           Df Variance
                            F Pr(>F)
##
            6 0.27378 9.5659 0.001 ***
## Model
## Residual 46 0.21943
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
\# Significance of partition from Season
anova(rda(species_2 ~ season +
          Condition(plants + land_use + margin) +
          Condition(temperature + oxygen + pH + conductivity),
          data=env_1))
## Permutation test for rda under reduced model
## Permutation: free
## Number of permutations: 999
## Model: rda(formula = species_2 ~ season + Condition(plants + land_use + margin) + Condition(temperat
           Df Variance
##
                            F Pr(>F)
            1 0.006247 1.3096 0.201
## Model
## Residual 46 0.219425
```

Physicochemical and Physical characteriscs are all statistically significant in their contributions to Odonata community composition, even though the amount of variation explained by Physicochemistry is small. There was no statistically significant of season in Odonata composition independent of these other measured drivers.