

6. Worksheet: Among Site (Beta) Diversity – Part 1

Student Name; Z620: Quantitative Biodiversity, Indiana University

04 February, 2025

OVERVIEW

In this worksheet, we move beyond the investigation of within-site α -diversity. We will explore β -diversity, which is defined as the diversity that occurs among sites. This requires that we examine the compositional similarity of assemblages that vary in space or time.

After completing this exercise you will know how to:

1. formally quantify β -diversity
2. visualize β -diversity with heatmaps, cluster analysis, and ordination
3. test hypotheses about β -diversity using multivariate statistics

Directions:

1. In the Markdown version of this document in your cloned repo, change “Student Name” on line 3 (above) with your name.
2. Complete as much of the worksheet as possible during class.
3. Use the handout as a guide; it contains a more complete description of data sets along with examples of proper scripting needed to carry out the exercises.
4. Answer questions in the worksheet. Space for your answers is provided in this document and is indicated by the “>” character. If you need a second paragraph be sure to start the first line with “>”. You should notice that the answer is highlighted in green by RStudio (color may vary if you changed the editor theme).
5. Before you leave the classroom, **push** this file to your GitHub repo.
6. For the assignment portion of the worksheet, follow the directions at the bottom of this file.
7. When you are done, **Knit** the text and code into a PDF file.
8. After Knitting, submit the completed exercise by creating a **pull request** via GitHub. Your pull request should include this file (**6.BetaDiversity_1_Worksheet.Rmd**) with all code blocks filled out and questions answered) and the PDF output of Knitr (**6.BetaDiversity_1_Worksheet.pdf**).

The completed exercise is due on **Wednesday, February 5th, 2025 before 12:00 PM (noon)**.

1) R SETUP

Typically, the first thing you will do in either an R script or an RMarkdown file is setup your environment. This includes things such as setting the working directory and loading any packages that you will need.

In the R code chunk below, please provide the code to:

- 1) Clear your R environment,
- 2) Print your current working directory,
- 3) Set your working directory to your **Week3-Beta/** folder folder, and
- 4) Load the **vegan** R package (be sure to install first if you have not already).

```
rm(list = ls())
getwd()
```

```
## [1] "/cloud/project/QB2025_Guevara/Week3-Beta"
```

```
setwd("/cloud/project/QB2025_Guevara/Week3-Beta")
```

#If we do ``{r, include = FALSE}, the whole R chunk will not be printed in the knitted file.

2) LOADING DATA

Load dataset

In the R code chunk below, do the following:

1. load the `doubs` dataset from the `ade4` package, and
2. explore the structure of the dataset.

```
# note, please do not print the dataset when submitting
package.list <- c('vegan', 'ade4', 'viridis', 'gplots', 'BiodiversityR', 'indicspecies')
for (package in package.list){
  if (!require(package, character.only = TRUE, quietly = TRUE)) {
    install.packages(package)
    library(package, character.only = TRUE)
  }
}
```

```
## This is vegan 2.6-8
```

```
##
```

```
## Attaching package: 'gplots'
```

```
## The following object is masked from 'package:stats':
```

```
##
```

```
##      lowess
```

```
## Warning in fun(libname, pkgname): couldn't connect to display ":0"
```

```
## BiodiversityR 2.17-1.1: Use command BiodiversityRGUI() to launch the Graphical User Interface;
## to see changes use BiodiversityRGUI(changeLog=TRUE, backward.compatibility.messages=TRUE)
```

```
library(vegan)
```

```
data(doubs)
```

#doubs is our list, fish is our object in this case

```
doubs$fish
```

```
##      Cogo Satr Phph Neba Thth Teso Chna Chto Lele Lece Baba Spbi Gogo Eslu Pefl
## 1      0    3    0    0    0    0    0    0    0    0    0    0    0    0    0
## 2      0    5    4    3    0    0    0    0    0    0    0    0    0    0    0
## 3      0    5    5    5    0    0    0    0    0    0    0    0    0    1    0
## 4      0    4    5    5    0    0    0    0    0    1    0    0    1    2    2
## 5      0    2    3    2    0    0    0    0    5    2    0    0    2    4    4
## 6      0    3    4    5    0    0    0    0    1    2    0    0    1    1    1
## 7      0    5    4    5    0    0    0    0    1    1    0    0    0    0    0
## 8      0    0    0    0    0    0    0    0    0    0    0    0    0    0    0
## 9      0    0    1    3    0    0    0    0    0    5    0    0    0    0    0
## 10     0    1    4    4    0    0    0    0    2    2    0    0    1    0    0
## 11     1    3    4    1    1    0    0    0    0    1    0    0    0    0    0
## 12     2    5    4    4    2    0    0    0    0    1    0    0    0    0    0
```

```

## 13  2  5  5  2  3  2  0  0  0  0  0  0  0  0  0
## 14  3  5  5  4  4  3  0  0  0  1  1  0  1  1  0
## 15  3  4  4  5  2  4  0  0  3  3  2  0  2  0  0
## 16  2  3  3  5  0  5  0  4  5  2  2  1  2  1  1
## 17  1  2  4  4  1  2  1  4  3  2  3  4  1  1  2
## 18  1  1  3  3  1  1  1  3  2  3  3  3  2  1  3
## 19  0  0  3  5  0  1  2  3  2  1  2  2  4  1  1
## 20  0  0  1  2  0  0  2  2  2  3  4  3  4  2  2
## 21  0  0  1  1  0  0  2  2  2  2  4  2  5  3  3
## 22  0  0  0  1  0  0  3  2  3  4  5  1  5  3  4
## 23  0  0  0  0  0  0  0  0  0  1  0  0  0  0  0
## 24  0  0  0  0  0  0  1  0  0  2  0  0  1  0  0
## 25  0  0  0  0  0  0  0  0  1  1  0  0  2  1  0
## 26  0  0  0  1  0  0  1  0  1  2  2  1  3  2  1
## 27  0  0  0  1  0  0  1  1  2  3  4  1  4  4  1
## 28  0  0  0  1  0  0  1  1  2  4  3  1  4  3  2
## 29  0  1  1  1  1  1  2  2  3  4  5  3  5  5  4
## 30  0  0  0  0  0  0  1  2  3  3  3  5  5  4  5
##      Rham Legi Scer Cyca Titi Abbr Icme Acce Ruru Blbj Alal Anan
## 1      0      0      0      0      0      0      0      0      0      0      0      0
## 2      0      0      0      0      0      0      0      0      0      0      0      0
## 3      0      0      0      0      0      0      0      0      0      0      0      0
## 4      0      0      0      0      1      0      0      0      0      0      0      0
## 5      0      0      2      0      3      0      0      0      5      0      0      0
## 6      0      0      0      0      2      0      0      0      1      0      0      0
## 7      0      0      0      0      0      0      0      0      0      0      0      0
## 8      0      0      0      0      0      0      0      0      0      0      0      0
## 9      0      0      0      0      1      0      0      0      4      0      0      0
## 10     0      0      0      0      0      0      0      0      0      0      0      0
## 11     0      0      0      0      0      0      0      0      0      0      0      0
## 12     0      0      0      0      0      0      0      0      0      0      0      0
## 13     0      0      0      0      0      0      0      0      0      0      0      0
## 14     0      0      0      0      0      0      0      0      0      0      0      0
## 15     0      0      0      0      1      0      0      0      0      0      0      0
## 16     0      1      0      1      1      0      0      0      1      0      0      0
## 17     1      1      0      1      1      0      0      0      2      0      2      1
## 18     2      1      0      1      1      0      0      1      2      0      2      1
## 19     2      1      1      1      2      1      0      1      5      1      3      1
## 20     3      2      2      1      4      1      0      2      5      2      5      2
## 21     3      2      2      2      4      3      1      3      5      3      5      2
## 22     3      3      2      3      4      4      2      4      5      4      5      2
## 23     0      0      0      0      0      0      0      0      1      0      2      0
## 24     0      1      0      0      0      0      0      2      2      1      5      0
## 25     0      0      1      0      0      0      0      1      1      0      3      0
## 26     2      2      1      1      3      2      1      4      4      2      5      2
## 27     3      3      1      2      5      3      2      5      5      4      5      3
## 28     4      4      2      4      4      3      3      5      5      5      5      4
## 29     5      5      2      3      3      4      4      5      5      4      5      4
## 30     5      3      5      5      5      5      5      5      5      5      5      5

```

```
str(doubs, max.level = 1)
```

```

## List of 4
## $ env      : 'data.frame': 30 obs. of  11 variables:
## $ fish     : 'data.frame': 30 obs. of 27 variables:

```

```
## $ xy      : 'data.frame': 30 obs. of  2 variables:
## $ species: 'data.frame': 27 obs. of  4 variables:
```

```
head(doubs$env)
```

```
##   dfs alt   slo flo pH har pho nit amm oxy bdo
## 1   3 934 6.176 84 79 45  1 20  0 122 27
## 2  22 932 3.434 100 80 40  2 20 10 103 19
## 3 102 914 3.638 180 83 52  5 22  5 105 35
## 4 185 854 3.497 253 80 72 10 21  0 110 13
## 5 215 849 3.178 264 81 84 38 52 20  80 62
## 6 324 846 3.497 286 79 60 20 15  0 102 53
```

```
head(doubs$fish)
```

```
##   Cogo Satr Phph Neba Thth Teso Chna Chto Lele Lece Baba Spbi Gogo Eslu Pefl
## 1   0   3   0   0   0   0   0   0   0   0   0   0   0   0   0
## 2   0   5   4   3   0   0   0   0   0   0   0   0   0   0   0
## 3   0   5   5   5   0   0   0   0   0   0   0   0   0   1   0
## 4   0   4   5   5   0   0   0   0   0   1   0   0   1   2   2
## 5   0   2   3   2   0   0   0   0   5   2   0   0   2   4   4
## 6   0   3   4   5   0   0   0   0   1   2   0   0   1   1   1
##   Rham Legi Scer Cyca Titi Abbr Icme Acce Ruru Blbj Alal Anan
## 1   0   0   0   0   0   0   0   0   0   0   0   0   0
## 2   0   0   0   0   0   0   0   0   0   0   0   0   0
## 3   0   0   0   0   0   0   0   0   0   0   0   0   0
## 4   0   0   0   0   1   0   0   0   0   0   0   0   0
## 5   0   0   2   0   3   0   0   0   5   0   0   0   0
## 6   0   0   0   0   2   0   0   0   1   0   0   0   0
```

```
? head
```

Question 1: Describe some of the attributes of the `doubs` dataset.

- How many objects are in `doubs`?
- How many fish species are there in the `doubs` dataset?
- How many sites are in the `doubs` dataset?

Answer 1a: There are 4 objects in ‘`doubs`’, where each object is its own `data.frame` (`fish`, `env`, `xy`, and `species`). **Answer 1b:** There are 27 species of fish in the `doubs` dataset **Answer 1c:** There are 30 sites in the `doubs` dataset

Visualizing the Doubs River Dataset

Question 2: Answer the following questions based on the spatial patterns of richness (i.e., α -diversity) and Brown Trout (*Salmo trutta*) abundance in the Doubs River.

- How does fish richness vary along the sampled reach of the Doubs River?
- How does Brown Trout (*Salmo trutta*) abundance vary along the sampled reach of the Doubs River?
- What do these patterns say about the limitations of using richness when examining patterns of biodiversity?

Answer 2a: Fish richness seems to be greatest downstream from Doubs River where it is less rich the further upstream **Answer 2b:** Brown Trout seems to be the opposite of fish richness where there is less abundance of Brown Trout downstream and more Brown Trout upstream **Answer 2c:** These patterns shows the limitation that using richness fails to account for evenness when examining patterns of biodiversity. It is most likely that Brown Trout dominate upstream of this river, which is why upstream fish richness is so low compared to downstream.

3) QUANTIFYING BETA-DIVERSITY

In the R code chunk below, do the following:

1. write a function (`beta.w()`) to calculate Whittaker's β -diversity (i.e., β_w) that accepts a site-by-species matrix with optional arguments to specify pairwise turnover between two sites, and
2. use this function to analyze various aspects of β -diversity in the Doubs River.

```
beta.w <- function(site.by.species = ""){
  SbyS.pa <- decostand(site.byspecies, method = "pa")
  #convert to presence-absence
  S <- ncol(SbyS.pa[,which(colSums(SbyS.pa) > 0)])
  # number of species in the region
  a.bar <- mean(specnumber(SbyS.pa))
  #average richness at each site
  b.w <- round(S/a.bar, 3)
  #round to 3 decimal places
  return(b.w)
}

#Turnover between two sites
beta.w <- function(site.by.species = "", sitenum1 = "", sitenum2 = "", pairwise = FALSE){
  #Only if we specify pairwise as TRUE, do this:
  if (pairwise == TRUE){
    #As a check, let's print an error if we do not provide needed arguments
    if (sitenum1 == "" | sitenum2 == ""){
      print("Error: please specify sites to compare")
      return(NA)}
    #If our function made it this far, let us calculate beta diversity
    site1 = site.by.species[sitenum1,]
    #Select site 1
    site2 = site.by.species[sitenum2,]
    #Select site 2
    site1 = subset(site1, select = site1 > 0)
    #Remove absences
    site2 = subset(site2, select = site2 > 0)
    #Removes absences
    gamma = union(colnames(site1), colnames(site2))
    #Gamma species pool
    s = length(gamma)
    #gamma richness
    a.bar = mean(c(specnumber(site1), specnumber(site2)))
    #Mean sample richness
    b.w = round(s/a.bar - 1, 3)
    return(b.w)
  }
  else{SbyS.pa <- decostand(site.by.species, method = "pa")
  #convert to presence-absence
  S <- ncol(SbyS.pa[,which(colSums(SbyS.pa) > 0)])
  #number of species in region
  a.bar <- mean(specnumber(SbyS.pa))
  #average richness at each site
  b.w <- round(S/a.bar, 3)
  return(b.w)
}
```

```

}

#To answer 3b...We see that
beta.w(site.by.species = doubs$fish, sitenum1 = 1, sitenum2 = 2, pairwise = TRUE)

## [1] 0.5

beta.w(site.by.species = doubs$fish, sitenum1 = 1, sitenum2 = 10, pairwise = TRUE)

## [1] 0.714

```

Question 3: Using your `beta.w()` function above, answer the following questions:

- Describe how local richness (α) and turnover (β) contribute to regional (γ) fish diversity in the Doubs.
- Is the fish assemblage at site 1 more similar to the one at site 2 or site 10?
- Using your understanding of the equation $\beta_w = \gamma/\alpha$, how would your interpretation of β change if we instead defined beta additively (i.e., $\beta = \gamma - \alpha$)?

Answer 3a: According to our beta function, the regional species pool is 2.16x more dense than the average richness at each site within the region (local richness or α). In order to acquire our regional fish diversity in Doubs, we would need to multiply our alpha diversity and beta diversity since the equation is $\text{beta} = \text{gamma}/\alpha$. **Answer 3b:** Site 1 is more similar to site 2 as we see that the Whittaker's beta diversity value smaller indicating more shared species and less turnover.

Answer 3c: From what I understand, $\text{beta}(w)$ is equal to the ratio of regional diversity over the average richness at each site within the specified region subtracted by one to provide the species turnover. If we were to convert this from being a ratio to a subtraction, our values of beta would not reveal how different the total diversity is from the mean site diversity by a given factor, but would instead reveal how many species in the regional pool are not found in the average local site. In sum, if beta was defined as additively rather than multiplicatively, beta would be an difference-based or absolute measure of diversity but would be more difficult to compare values across other regions since the values of diversity are no longer relative.

The Resemblance Matrix

In order to quantify β -diversity for more than two samples, we need to introduce a new primary ecological data structure: the **Resemblance Matrix**.

Question 4: How do incidence- and abundance-based metrics differ in their treatment of rare species?

Answer 4: They differ in their treatment of rare species as incidence-based metrics treat rare species equally to more common/dominant species. Abundance-based metrics take into accountability the abundance (as in the name) of each species, weighing them based on relative abundance within a site. Dominant/common species would affect the similarity/dissimilarity between sites more so than a rare species when using an abundance-based analysis.

In the R code chunk below, do the following:

- make a new object, `fish`, containing the fish abundance data for the Doubs River,
- remove any sites where no fish were observed (i.e., rows with sum of zero),
- construct a resemblance matrix based on Sørensen's Similarity ("fish.ds"), and
- construct a resemblance matrix based on Bray-Curtis Distance ("fish.db").

```

fish <- doubs$fish
fish <- fish[-8, ] #Removes site 8 from data

#Calculate Jaccard
fish.dj <- vegdist(fish, method = "jaccard", binary = TRUE)
new.fish.dj <- vegdist(fish, method = "jaccard", binary = TRUE, upper = TRUE, diag = TRUE)
#Calculate Bray-Curtis

```

```
fish.db <- vegdist(fish, method = "bray")

#Calculate Sørensen
fish.ds <- vegdist(fish, method = "bray", binary = TRUE)
full.fish.ds <- vegdist(fish, method = "bray", binary = TRUE, upper = TRUE, diag = TRUE)
#Printing the Bray-Curtis-based resemblance matrix in the console
fish.db
```

```
##           1           2           3           4           5           6           7
## 2  0.60000000
## 3  0.68421053 0.14285714
## 4  0.75000000 0.33333333 0.18918919
## 5  0.89189189 0.69565217 0.68000000 0.49090909
## 6  0.75000000 0.39393939 0.29729730 0.19047619 0.41818182
## 7  0.68421053 0.14285714 0.12500000 0.24324324 0.64000000 0.24324324
## 9  1.00000000 0.69230769 0.73333333 0.65714286 0.58333333 0.54285714 0.66666667
## 10 0.88235294 0.38461538 0.40000000 0.37142857 0.54166667 0.25714286 0.26666667
## 11 0.57142857 0.30434783 0.40740741 0.43750000 0.68888889 0.43750000 0.33333333
## 12 0.71428571 0.20000000 0.23529412 0.33333333 0.69230769 0.38461538 0.17647059
## 13 0.72727273 0.29032258 0.31428571 0.45000000 0.73584906 0.55000000 0.37142857
## 14 0.80645161 0.40000000 0.31818182 0.34693878 0.67741935 0.42857143 0.36363636
## 15 0.83333333 0.51111111 0.46938776 0.40740741 0.55223881 0.37037037 0.38775510
## 16 0.86046512 0.65384615 0.57142857 0.47540984 0.45945946 0.37704918 0.53571429
## 17 0.91489362 0.67857143 0.63333333 0.50769231 0.51282051 0.44615385 0.60000000
## 18 0.95555556 0.74074074 0.72413793 0.58730159 0.50000000 0.52380952 0.68965517
## 19 1.00000000 0.79310345 0.70967742 0.61194030 0.50000000 0.52238806 0.67741935
## 20 1.00000000 0.91176471 0.88888889 0.74025974 0.48888889 0.68831169 0.86111111
## 21 1.00000000 0.94594595 0.92307692 0.78313253 0.50000000 0.73493976 0.89743590
## 22 1.00000000 0.97619048 0.95454545 0.82795699 0.52830189 0.78494624 0.93181818
## 23 1.00000000 1.00000000 1.00000000 0.92000000 0.89473684 0.84000000 0.90000000
## 24 1.00000000 1.00000000 1.00000000 0.88888889 0.79591837 0.77777778 0.93548387
## 25 1.00000000 1.00000000 0.92592593 0.81250000 0.68888889 0.68750000 0.85185185
## 26 1.00000000 0.96363636 0.93220339 0.78125000 0.55844156 0.68750000 0.89830508
## 27 1.00000000 0.97333333 0.94936709 0.83333333 0.56701031 0.76190476 0.92405063
## 28 1.00000000 0.97560976 0.95348837 0.82417582 0.57692308 0.78021978 0.93023256
## 29 0.97777778 0.93939394 0.92233010 0.81481481 0.53719008 0.77777778 0.90291262
## 30 1.00000000 1.00000000 0.98095238 0.87272727 0.59349593 0.83636364 0.96190476
##           9           10          11          12          13          14          15
## 2
## 3
## 4
## 5
## 6
## 7
## 9
## 10 0.57142857
## 11 0.76000000 0.44000000
## 12 0.68750000 0.37500000 0.24137931
## 13 0.81818182 0.57575758 0.33333333 0.18918919
## 14 0.76190476 0.47619048 0.43589744 0.21739130 0.19148936
## 15 0.65957447 0.40425532 0.50000000 0.33333333 0.38461538 0.24590164
## 16 0.70370370 0.51851852 0.64705882 0.55172414 0.59322034 0.44117647 0.26027397
## 17 0.68965517 0.51724138 0.63636364 0.58064516 0.61904762 0.50000000 0.40259740
## 18 0.64285714 0.57142857 0.69811321 0.66666667 0.70491803 0.60000000 0.46666667
```

```

## 19 0.66666667 0.63333333 0.82456140 0.75000000 0.81538462 0.67567568 0.56962025
## 20 0.68571429 0.77142857 0.91044776 0.89189189 0.92000000 0.83333333 0.70786517
## 21 0.76315789 0.81578947 0.91780822 0.92500000 0.95061728 0.86666667 0.76842105
## 22 0.76744186 0.86046512 0.95180723 0.95555556 0.97802198 0.90000000 0.77142857
## 23 0.77777778 0.88888889 0.86666667 0.90909091 1.00000000 0.93750000 0.94594595
## 24 0.72413793 0.79310345 0.92307692 0.93939394 1.00000000 0.90697674 0.87500000
## 25 0.84000000 0.76000000 0.90909091 0.93103448 1.00000000 0.84615385 0.81818182
## 26 0.71929825 0.82456140 0.92592593 0.93442623 0.96774194 0.85915493 0.76315789
## 27 0.76623377 0.84415584 0.94594595 0.95061728 0.97560976 0.89010989 0.77083333
## 28 0.76190476 0.85714286 0.95061728 0.95454545 0.97752809 0.89795918 0.78640777
## 29 0.78217822 0.84158416 0.89795918 0.90476190 0.90566038 0.84347826 0.73333333
## 30 0.84466019 0.90291262 0.98000000 0.98130841 1.00000000 0.93162393 0.81967213
##          16          17          18          19          20          21          22
## 2
## 3
## 4
## 5
## 6
## 7
## 9
## 10
## 11
## 12
## 13
## 14
## 15
## 16
## 17 0.26190476
## 18 0.34146341 0.13953488
## 19 0.39534884 0.31111111 0.25000000
## 20 0.58333333 0.42000000 0.32653061 0.23529412
## 21 0.62745098 0.49056604 0.40384615 0.29629630 0.10169492
## 22 0.66071429 0.55172414 0.47368421 0.38983051 0.18750000 0.10447761
## 23 0.90909091 0.83333333 0.82608696 0.84000000 0.86666667 0.87878788 0.89473684
## 24 0.81818182 0.69491525 0.64912281 0.63934426 0.57746479 0.61038961 0.65517241
## 25 0.76470588 0.74545455 0.66037736 0.61403509 0.67164179 0.69863014 0.73493976
## 26 0.63855422 0.54022989 0.45882353 0.32584270 0.21212121 0.20000000 0.25217391
## 27 0.66990291 0.57009346 0.48571429 0.37614679 0.19327731 0.13600000 0.12592593
## 28 0.69090909 0.57894737 0.50000000 0.41379310 0.22222222 0.16666667 0.12676056
## 29 0.65354331 0.51145038 0.44186047 0.41353383 0.24475524 0.18120805 0.11949686
## 30 0.72093023 0.57894737 0.52671756 0.48148148 0.29655172 0.23178808 0.18012422
##          23          24          25          26          27          28          29
## 2
## 3
## 4
## 5
## 6
## 7
## 9
## 10
## 11
## 12
## 13
## 14

```



```
## 15
## 16
## 17
## 18
## 19
## 20
## 21
## 22
## 23
## 24 0.57894737
## 25 0.46666667 0.46153846
## 26 0.82978723 0.48275862 0.59259259
## 27 0.88059701 0.61538462 0.70270270 0.18867925
## 28 0.89189189 0.64705882 0.72839506 0.23893805 0.09774436
## 29 0.91208791 0.70588235 0.77551020 0.33846154 0.18666667 0.14649682
## 30 0.91397849 0.71153846 0.78000000 0.36363636 0.19736842 0.15723270 0.14772727
```

```
#Making fish.db into a square matrix from a large diagonal matrix. This doesn't really need to be done.
fish.db <- vegdist(fish, method = "bray", upper = TRUE, diag = TRUE)
fish.db
```

```
##          1          2          3          4          5          6          7
## 1  0.00000000 0.60000000 0.68421053 0.75000000 0.89189189 0.75000000 0.68421053
## 2  0.60000000 0.00000000 0.14285714 0.33333333 0.69565217 0.39393939 0.14285714
## 3  0.68421053 0.14285714 0.00000000 0.18918919 0.68000000 0.29729730 0.12500000
## 4  0.75000000 0.33333333 0.18918919 0.00000000 0.49090909 0.19047619 0.24324324
## 5  0.89189189 0.69565217 0.68000000 0.49090909 0.00000000 0.41818182 0.64000000
## 6  0.75000000 0.39393939 0.29729730 0.19047619 0.41818182 0.00000000 0.24324324
## 7  0.68421053 0.14285714 0.12500000 0.24324324 0.64000000 0.24324324 0.00000000
## 9  1.00000000 0.69230769 0.73333333 0.65714286 0.58333333 0.54285714 0.66666667
## 10 0.88235294 0.38461538 0.40000000 0.37142857 0.54166667 0.25714286 0.26666667
## 11 0.57142857 0.30434783 0.40740741 0.43750000 0.68888889 0.43750000 0.33333333
## 12 0.71428571 0.20000000 0.23529412 0.33333333 0.69230769 0.38461538 0.17647059
## 13 0.72727273 0.29032258 0.31428571 0.45000000 0.73584906 0.55000000 0.37142857
## 14 0.80645161 0.40000000 0.31818182 0.34693878 0.67741935 0.42857143 0.36363636
## 15 0.83333333 0.51111111 0.46938776 0.40740741 0.55223881 0.37037037 0.38775510
## 16 0.86046512 0.65384615 0.57142857 0.47540984 0.45945946 0.37704918 0.53571429
## 17 0.91489362 0.67857143 0.63333333 0.50769231 0.51282051 0.44615385 0.60000000
## 18 0.95555556 0.74074074 0.72413793 0.58730159 0.50000000 0.52380952 0.68965517
## 19 1.00000000 0.79310345 0.70967742 0.61194030 0.50000000 0.52238806 0.67741935
## 20 1.00000000 0.91176471 0.88888889 0.74025974 0.48888889 0.68831169 0.86111111
## 21 1.00000000 0.94594595 0.92307692 0.78313253 0.50000000 0.73493976 0.89743590
## 22 1.00000000 0.97619048 0.95454545 0.82795699 0.52830189 0.78494624 0.93181818
## 23 1.00000000 1.00000000 1.00000000 0.92000000 0.89473684 0.84000000 0.90000000
## 24 1.00000000 1.00000000 1.00000000 0.88888889 0.79591837 0.77777778 0.93548387
## 25 1.00000000 1.00000000 0.92592593 0.81250000 0.68888889 0.68750000 0.85185185
## 26 1.00000000 0.96363636 0.93220339 0.78125000 0.55844156 0.68750000 0.89830508
## 27 1.00000000 0.97333333 0.94936709 0.83333333 0.56701031 0.76190476 0.92405063
## 28 1.00000000 0.97560976 0.95348837 0.82417582 0.57692308 0.78021978 0.93023256
## 29 0.97777778 0.93939394 0.92233010 0.81481481 0.53719008 0.77777778 0.90291262
## 30 1.00000000 1.00000000 0.98095238 0.87272727 0.59349593 0.83636364 0.96190476
##          9          10          11          12          13          14          15
## 1  1.00000000 0.88235294 0.57142857 0.71428571 0.72727273 0.80645161 0.83333333
## 2  0.69230769 0.38461538 0.30434783 0.20000000 0.29032258 0.40000000 0.51111111
## 3  0.73333333 0.40000000 0.40740741 0.23529412 0.31428571 0.31818182 0.46938776
```

## 4	0.65714286	0.37142857	0.43750000	0.33333333	0.45000000	0.34693878	0.40740741
## 5	0.58333333	0.54166667	0.68888889	0.69230769	0.73584906	0.67741935	0.55223881
## 6	0.54285714	0.25714286	0.43750000	0.38461538	0.55000000	0.42857143	0.37037037
## 7	0.66666667	0.26666667	0.33333333	0.17647059	0.37142857	0.36363636	0.38775510
## 9	0.00000000	0.57142857	0.76000000	0.68750000	0.81818182	0.76190476	0.65957447
## 10	0.57142857	0.00000000	0.44000000	0.37500000	0.57575758	0.47619048	0.40425532
## 11	0.76000000	0.44000000	0.00000000	0.24137931	0.33333333	0.43589744	0.50000000
## 12	0.68750000	0.37500000	0.24137931	0.00000000	0.18918919	0.21739130	0.33333333
## 13	0.81818182	0.57575758	0.33333333	0.18918919	0.00000000	0.19148936	0.38461538
## 14	0.76190476	0.47619048	0.43589744	0.21739130	0.19148936	0.00000000	0.24590164
## 15	0.65957447	0.40425532	0.50000000	0.33333333	0.38461538	0.24590164	0.00000000
## 16	0.70370370	0.51851852	0.64705882	0.55172414	0.59322034	0.44117647	0.26027397
## 17	0.68965517	0.51724138	0.63636364	0.58064516	0.61904762	0.50000000	0.40259740
## 18	0.64285714	0.57142857	0.69811321	0.66666667	0.70491803	0.60000000	0.46666667
## 19	0.66666667	0.63333333	0.82456140	0.75000000	0.81538462	0.67567568	0.56962025
## 20	0.68571429	0.77142857	0.91044776	0.89189189	0.92000000	0.83333333	0.70786517
## 21	0.76315789	0.81578947	0.91780822	0.92500000	0.95061728	0.86666667	0.76842105
## 22	0.76744186	0.86046512	0.95180723	0.95555556	0.97802198	0.90000000	0.77142857
## 23	0.77777778	0.88888889	0.86666667	0.90909091	1.00000000	0.93750000	0.94594595
## 24	0.72413793	0.79310345	0.92307692	0.93939394	1.00000000	0.90697674	0.87500000
## 25	0.84000000	0.76000000	0.90909091	0.93103448	1.00000000	0.84615385	0.81818182
## 26	0.71929825	0.82456140	0.92592593	0.93442623	0.96774194	0.85915493	0.76315789
## 27	0.76623377	0.84415584	0.94594595	0.95061728	0.97560976	0.89010989	0.77083333
## 28	0.76190476	0.85714286	0.95061728	0.95454545	0.97752809	0.89795918	0.78640777
## 29	0.78217822	0.84158416	0.89795918	0.90476190	0.90566038	0.84347826	0.73333333
## 30	0.84466019	0.90291262	0.98000000	0.98130841	1.00000000	0.93162393	0.81967213
##	16	17	18	19	20	21	22
## 1	0.86046512	0.91489362	0.95555556	1.00000000	1.00000000	1.00000000	1.00000000
## 2	0.65384615	0.67857143	0.74074074	0.79310345	0.91176471	0.94594595	0.97619048
## 3	0.57142857	0.63333333	0.72413793	0.70967742	0.88888889	0.92307692	0.95454545
## 4	0.47540984	0.50769231	0.58730159	0.61194030	0.74025974	0.78313253	0.82795699
## 5	0.45945946	0.51282051	0.50000000	0.50000000	0.48888889	0.50000000	0.52830189
## 6	0.37704918	0.44615385	0.52380952	0.52238806	0.68831169	0.73493976	0.78494624
## 7	0.53571429	0.60000000	0.68965517	0.67741935	0.86111111	0.89743590	0.93181818
## 9	0.70370370	0.68965517	0.64285714	0.66666667	0.68571429	0.76315789	0.76744186
## 10	0.51851852	0.51724138	0.57142857	0.63333333	0.77142857	0.81578947	0.86046512
## 11	0.64705882	0.63636364	0.69811321	0.82456140	0.91044776	0.91780822	0.95180723
## 12	0.55172414	0.58064516	0.66666667	0.75000000	0.89189189	0.92500000	0.95555556
## 13	0.59322034	0.61904762	0.70491803	0.81538462	0.92000000	0.95061728	0.97802198
## 14	0.44117647	0.50000000	0.60000000	0.67567568	0.83333333	0.86666667	0.90000000
## 15	0.26027397	0.40259740	0.46666667	0.56962025	0.70786517	0.76842105	0.77142857
## 16	0.00000000	0.26190476	0.34146341	0.39534884	0.58333333	0.62745098	0.66071429
## 17	0.26190476	0.00000000	0.13953488	0.31111111	0.42000000	0.49056604	0.55172414
## 18	0.34146341	0.13953488	0.00000000	0.25000000	0.32653061	0.40384615	0.47368421
## 19	0.39534884	0.31111111	0.25000000	0.00000000	0.23529412	0.29629630	0.38983051
## 20	0.58333333	0.42000000	0.32653061	0.23529412	0.00000000	0.10169492	0.18750000
## 21	0.62745098	0.49056604	0.40384615	0.29629630	0.10169492	0.00000000	0.10447761
## 22	0.66071429	0.55172414	0.47368421	0.38983051	0.18750000	0.10447761	0.00000000
## 23	0.90909091	0.83333333	0.82608696	0.84000000	0.86666667	0.87878788	0.89473684
## 24	0.81818182	0.69491525	0.64912281	0.63934426	0.57746479	0.61038961	0.65517241
## 25	0.76470588	0.74545455	0.66037736	0.61403509	0.67164179	0.69863014	0.73493976
## 26	0.63855422	0.54022989	0.45882353	0.32584270	0.21212121	0.20000000	0.25217391
## 27	0.66990291	0.57009346	0.48571429	0.37614679	0.19327731	0.13600000	0.12592593
## 28	0.69090909	0.57894737	0.50000000	0.41379310	0.22222222	0.16666667	0.12676056

##	29	0.65354331	0.51145038	0.44186047	0.41353383	0.24475524	0.18120805	0.11949686
##	30	0.72093023	0.57894737	0.52671756	0.48148148	0.29655172	0.23178808	0.18012422
##		23	24	25	26	27	28	29
##	1	1.00000000	1.00000000	1.00000000	1.00000000	1.00000000	1.00000000	0.97777778
##	2	1.00000000	1.00000000	1.00000000	0.96363636	0.97333333	0.97560976	0.93939394
##	3	1.00000000	1.00000000	0.92592593	0.93220339	0.94936709	0.95348837	0.92233010
##	4	0.92000000	0.88888889	0.81250000	0.78125000	0.83333333	0.82417582	0.81481481
##	5	0.89473684	0.79591837	0.68888889	0.55844156	0.56701031	0.57692308	0.53719008
##	6	0.84000000	0.77777778	0.68750000	0.68750000	0.76190476	0.78021978	0.77777778
##	7	0.90000000	0.93548387	0.85185185	0.89830508	0.92405063	0.93023256	0.90291262
##	9	0.77777778	0.72413793	0.84000000	0.71929825	0.76623377	0.76190476	0.78217822
##	10	0.88888889	0.79310345	0.76000000	0.82456140	0.84415584	0.85714286	0.84158416
##	11	0.86666667	0.92307692	0.90909091	0.92592593	0.94594595	0.95061728	0.89795918
##	12	0.90909091	0.93939394	0.93103448	0.93442623	0.95061728	0.95454545	0.90476190
##	13	1.00000000	1.00000000	1.00000000	0.96774194	0.97560976	0.97752809	0.90566038
##	14	0.93750000	0.90697674	0.84615385	0.85915493	0.89010989	0.89795918	0.84347826
##	15	0.94594595	0.87500000	0.81818182	0.76315789	0.77083333	0.78640777	0.73333333
##	16	0.90909091	0.81818182	0.76470588	0.63855422	0.66990291	0.69090909	0.65354331
##	17	0.83333333	0.69491525	0.74545455	0.54022989	0.57009346	0.57894737	0.51145038
##	18	0.82608696	0.64912281	0.66037736	0.45882353	0.48571429	0.50000000	0.44186047
##	19	0.84000000	0.63934426	0.61403509	0.32584270	0.37614679	0.41379310	0.41353383
##	20	0.86666667	0.57746479	0.67164179	0.21212121	0.19327731	0.22222222	0.24475524
##	21	0.87878788	0.61038961	0.69863014	0.20000000	0.13600000	0.16666667	0.18120805
##	22	0.89473684	0.65517241	0.73493976	0.25217391	0.12592593	0.12676056	0.11949686
##	23	0.00000000	0.57894737	0.46666667	0.82978723	0.88059701	0.89189189	0.91208791
##	24	0.57894737	0.00000000	0.46153846	0.48275862	0.61538462	0.64705882	0.70588235
##	25	0.46666667	0.46153846	0.00000000	0.59259259	0.70270270	0.72839506	0.77551020
##	26	0.82978723	0.48275862	0.59259259	0.00000000	0.18867925	0.23893805	0.33846154
##	27	0.88059701	0.61538462	0.70270270	0.18867925	0.00000000	0.09774436	0.18666667
##	28	0.89189189	0.64705882	0.72839506	0.23893805	0.09774436	0.00000000	0.14649682
##	29	0.91208791	0.70588235	0.77551020	0.33846154	0.18666667	0.14649682	0.00000000
##	30	0.91397849	0.71153846	0.78000000	0.36363636	0.19736842	0.15723270	0.14772727
##		30						
##	1	1.00000000						
##	2	1.00000000						
##	3	0.98095238						
##	4	0.87272727						
##	5	0.59349593						
##	6	0.83636364						
##	7	0.96190476						
##	9	0.84466019						
##	10	0.90291262						
##	11	0.98000000						
##	12	0.98130841						
##	13	1.00000000						
##	14	0.93162393						
##	15	0.81967213						
##	16	0.72093023						
##	17	0.57894737						
##	18	0.52671756						
##	19	0.48148148						
##	20	0.29655172						
##	21	0.23178808						
##	22	0.18012422						

```
## 23 0.91397849
## 24 0.71153846
## 25 0.78000000
## 26 0.36363636
## 27 0.19736842
## 28 0.15723270
## 29 0.14772727
## 30 0.00000000
```

Question 5: Using the distance matrices from above, answer the following questions:

- Does the resemblance matrix (`fish.db`) represent similarity or dissimilarity? What information in the resemblance matrix led you to arrive at your answer?
- Compare the resemblance matrices (`fish.db` or `fish.ds`) you just created. How does the choice of the Sørensen or Bray-Curtis distance influence your interpretation of site (dis)similarity?

Answer 5a: `fish.db` represents dissimilarity as it uses the Bray-Curtis Dissimilarity Abundance-based metric. When comparing each site to itself (diagonal line of zeros), we see that the value is zero indicating that there is no dissimilarity between the sites (makes sense because they are same site). **Answer 5b:** The choice of using Sørensen or Bray-Curtis method differ in their influence on our interpretation of site (dis)similarity as Sørensen is a incidence-based metric so sites will appear more similar if they share lots of species rather than focusing on the abundance of those shared species as does the Bray-Curtis method.

4) VISUALIZING BETA-DIVERSITY

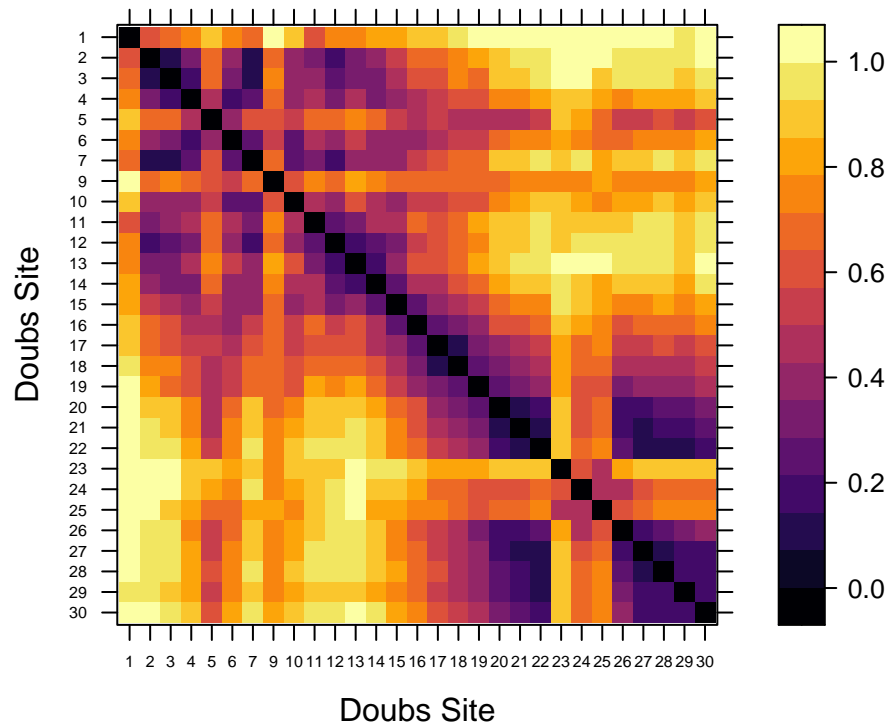
A. Heatmaps

In the R code chunk below, do the following:

- define a color palette,
- define the order of sites in the Doubs River, and
- use the `levelplot()` function to create a heatmap of fish abundances in the Doubs River.

```
#Define Order of Sites
order <- rev(attr(fish.db, "Labels"))
#Plot Heatmap
levelplot(as.matrix(fish.db)[,order], aspect = "iso", col.regions = inferno,
          xlab = "Doubs Site", ylab = "Doubs Site", scales = list(cex = 0.5),
          main = "Bray-Curtis Distance")
```

Bray–Curtis Distance



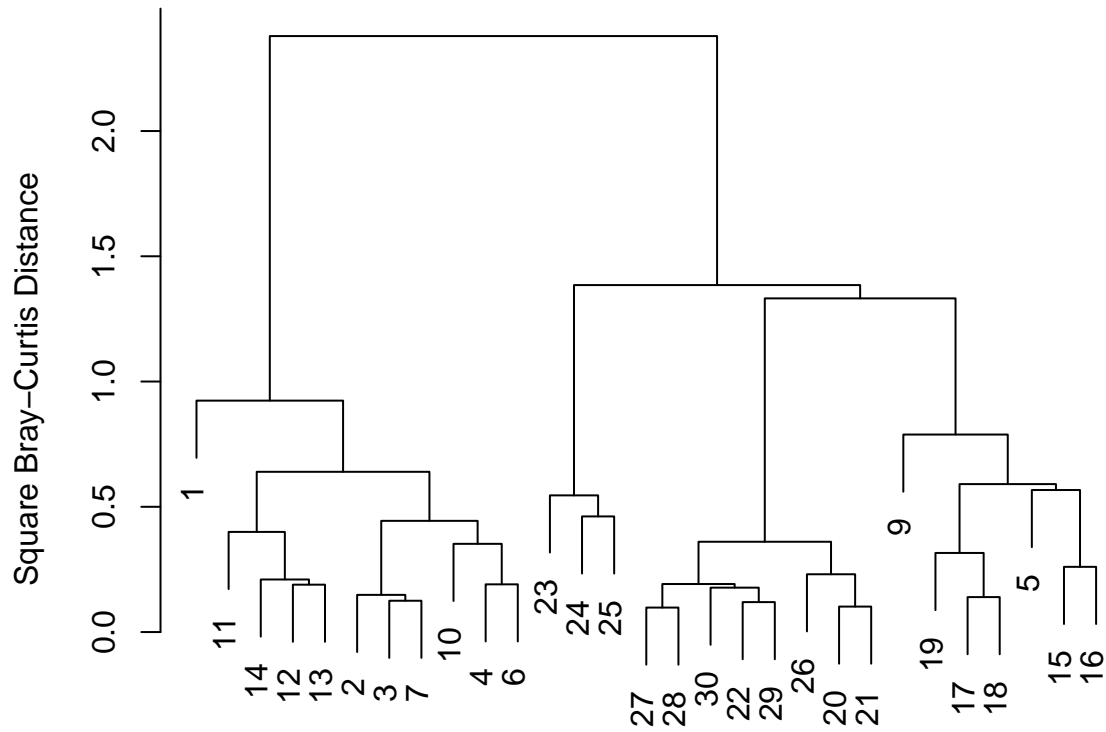
B. Cluster Analysis

In the R code chunk below, do the following:

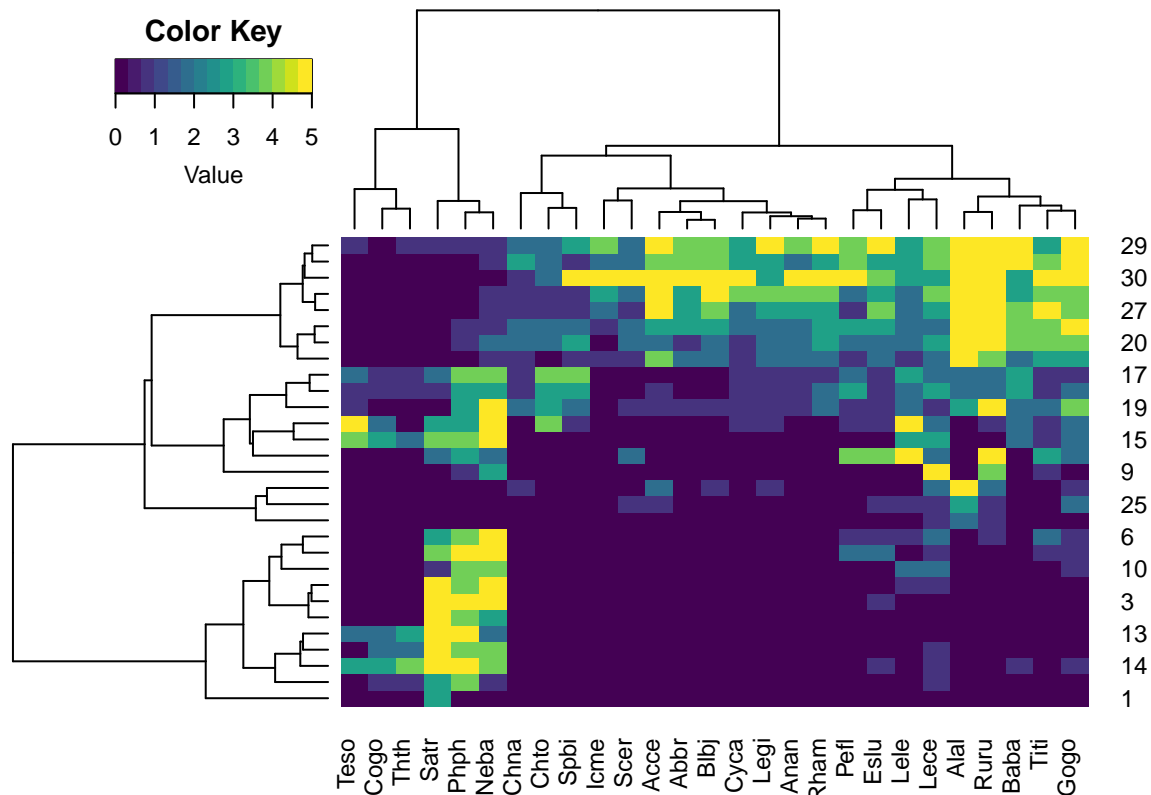
1. perform a cluster analysis using Ward's Clustering, and
2. plot your cluster analysis (use either `hclust` or `heatmap.2`).

```
#Perform Cluster Analysis
fish.ward <- hclust(fish.db, method = "ward.D2")
#Plot Cluster
par(mar = c(1,5,2,2) + 0.1)
plot(fish.ward, main = "Doubs River Fish: Ward's Clustering", ylab = "Square Bray–Curtis Distance")
```

Doubs River Fish: Ward's Clustering



```
gplots::heatmap.2(as.matrix(fish),
  distfun = function(x) vegdist(x, method = "bray"),
  hclustfun = function(x) hclust(x, method = "ward.D2"),
  col = viridis, trace = "none", density.info = "none")
```



Question 6: Based on cluster analyses and the introductory plots that we generated after loading the data, develop an ecological hypothesis for fish diversity the Doubs data set?

Answer 6: Based off of what we are seeing, I am inclined to hypothesized that sites 15 - 30 are close in proximity and sites 1 - 14 are closer in proximity. This could imply that there is a downstream/upstream component in terms of how the fish are being measured where certain species of fish reside either downstream or upstream, or are capable of moving between them. For example, site 5 shows relatively similar similarity across almost all sites. I would hypothesize that sites 1-14 are either downstream/upstream and sites 15 - 30 are the opposite.

C. Ordination

Principal Coordinates Analysis (PCoA)

In the R code chunk below, do the following:

1. perform a Principal Coordinates Analysis to visualize beta-diversity
2. calculate the variation explained by the first three axes in your ordination
3. plot the PCoA ordination,
4. label the sites as points using the Doubs River site number, and
5. identify influential species and add species coordinates to PCoA plot.

```
fish.pcoa <- cmdscale(fish.db, eig = TRUE, k = 3)
explainvar1 <- round(fish.pcoa$eig[1]/sum(fish.pcoa$eig), 3) * 100
explainvar2 <- round(fish.pcoa$eig[2]/sum(fish.pcoa$eig), 3) * 100
explainvar3 <- round(fish.pcoa$eig[3]/sum(fish.pcoa$eig), 3) * 100
sum.eig <- sum(explainvar1, explainvar2, explainvar3)

#Define Plot Parameters
par(mar = c(5, 5, 1, 2) + 0.1)
#Initiate Plot
```

```

plot(fish.pcoa$points[,1], fish.pcoa$points[,2], ylim = c(-0.2, 0.7),
     xlab = paste("PCoA 1 (", explainvar1, "%)", sep = ""),
     ylab = paste("PCoA 2 (", explainvar2, "%)", sep = ""),
     pch = 16, cex = 2.0, type = "n", cex.lab = 1.5,
     cex.axis = 1.2, axes = FALSE)

#Add Axes
axis(side = 1, labels = T, lwd.ticks = 2, cex.axis = 1.2, las = 1)
axis(side = 2, labels = T, lwd.ticks = 2, cex.axis = 1.2, las = 1)
abline(h = 0, v = 0, lty = 3)
box(lwd = 2)

#Add Points and Labels
points(fish.pcoa$points[,1], fish.pcoa$points[,2],
       pch = 19, cex = 3, bg = "gray", col = "gray")
text(fish.pcoa$points[,1], fish.pcoa$points[,2],
     labels = row.names(fish.pcoa$points))

#First we calculate the relative abundances of each species at each site
fishREL <- fish
for(i in 1:nrow(fish)){
  fishREL[i, ] =fish[i, ] / sum(fish[i, ])
}

add.spec.scores.class <- function(ordi,comm,method="cor.scores",multi=1,Rscale=F,scaling="1") {
  ordiscores <- scores(ordi,display="sites")
  n <- ncol(comm)
  p <- ncol(ordiscores)
  specscores <- array(NA,dim=c(n,p))
  rownames(specscores) <- colnames(comm)
  colnames(specscores) <- colnames(ordiscores)
  if (method == "cor.scores") {
    for (i in 1:n) {
      for (j in 1:p) {specscores[i,j] <- cor(comm[,i],ordiscores[,j],method="pearson")}
    }
  }
  if (method == "wa.scores") {specscores <- wascores(ordiscores,comm)}
  if (method == "pcoa.scores") {
    rownames(ordiscores) <- rownames(comm)
    eigenv <- ordi$eig
    accounted <- sum(eigenv)
    tot <- 2*(accounted/ordi$GOF[2])-(accounted/ordi$GOF[1])
    eigen.var <- eigenv/(nrow(comm)-1)
    neg <- length(eigenv[eigenv<0])
    pos <- length(eigenv[eigenv>0])
    tot <- tot/(nrow(comm)-1)
    eigen.percen <- 100*eigen.var/tot
    eigen.cumpercen <- cumsum(eigen.percen)
    constant <- ((nrow(comm)-1)*tot)^0.25
    ordiscores <- ordiscores * (nrow(comm)-1)^-0.5 * tot^-0.5 * constant
    p1 <- min(p, pos)
    for (i in 1:n) {
      for (j in 1:p1) {

```



```

    specscores[i,j] <- cor(comm[,i],ordiscores[,j])*sd(comm[,i])/sd(ordiscores[,j])
    if(is.na(specscores[i,j])) {specscores[i,j]<-0}
  }
}
if (Rscale==T && scaling=="2") {
  percen <- eigen.var/tot
  percen <- percen^0.5
  ordiscores <- sweep(ordiscores,2,percen,"/")
  specscores <- sweep(specscores,2,percen,"*")
}
if (Rscale==F) {
  specscores <- specscores / constant
  ordiscores <- ordi$points
}
ordi$points <- ordiscores
ordi$eig <- eigen.var
ordi$eig.percen <- eigen.percen
ordi$eig.cumpercen <- eigen.cumpercen
ordi$eigen.total <- tot
ordi$R.constant <- constant
ordi$Rscale <- Rscale
ordi$scaling <- scaling
}
specscores <- specscores * multi
ordi$cproj <- specscores
return(ordi)
}

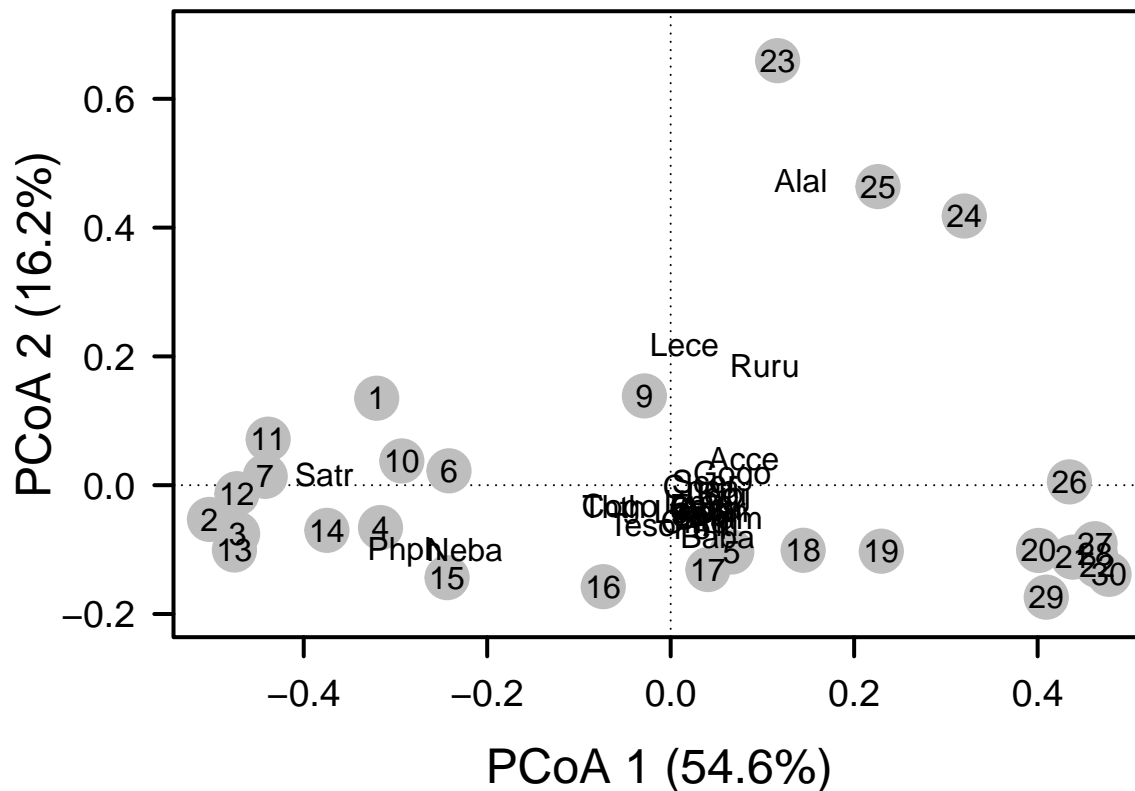
```

#Now we will use the relative abundances of each species at each site to calculate and add species scores

```

library(vegan)
fish.pcoa <- add.spec.scores.class(fish.pcoa,fishREL, method = "pcoa.scores")
text(fish.pcoa$cproj[,1], fish.pcoa$cproj[,2],
     labels = row.names(fish.pcoa$cproj), col = "black")

```



In the R code chunk below, do the following:

1. identify influential species based on correlations along each PCoA axis (use a cutoff of 0.70), and
2. use a permutation test (999 permutations) to test the correlations of each species along each axis.

```
spe.corr <- add.spec.scores.class(fish.pcoa, fishREL, method = "cor.scores")$cproj
corrcut <- 0.7 #user defined cutoff
imp.spp <- spe.corr[abs(spe.corr[,1]) >= corrcut | abs(spe.corr[,2]) >= corrcut, ]

#Permutation test for Species Abundances Across Axes
fit <- envfit(fish.pcoa, fishREL, perm = 999)

#Create "Factors" vector
quality <- c(rep("HQ", 13), rep("MQ", 5), rep("LQ", 6), rep("MQ", 5))
#Run PERMANOVA with adonis function
adonis2(fish ~ quality, method = "bray", permutations = 999)

## Permutation test for adonis under reduced model
## Permutation: free
## Number of permutations: 999
##
## adonis2(formula = fish ~ quality, permutations = 999, method = "bray")
##          Df SumOfSqs      R2      F Pr(>F)
## Model      2   3.0947 0.45765 10.97 0.001 ***
## Residual  26   3.6674 0.54235
## Total     28   6.7621 1.00000
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Question 7: Address the following questions about the ordination results of the Doubs data set:

- Describe the grouping of sites in the Doubs River based on fish community composition.
- Generate a hypothesis about which fish species are potential indicators of river quality.

Answer 7a: Looking at fish community composition, we see a clear break between almost all of sites 1 - 14 and 15-30 with the exception 5, 17, and 16 with a wider distribution of variance across the latter half of the total sites as explained by PCoA 2. **Answer 7b:** I hypothesize that species Alal, Lece, and Ruru are indicators of river quality as they seem to appear quite distinguished from the rest of the species clustering.

SYNTHESIS

Load the dataset from that you and your partner are using for the team project. Use one of the tools introduced in the beta diversity module to visualize your data. Describe any interesting patterns and identify a hypothesis is relevant to the principles of biodiversity.

```
Projdata <- read.csv("/cloud/project/QB2025_Guevara/Week2-Alpha/MAT_fungal_abundances.csv")
str(Projdata)
```

```
## 'data.frame':    704 obs. of  12 variables:
## $ Plot          : chr  "MAT.B1.C" "MAT.B1.C" "MAT.B1.C" "MAT.B1.C" ...
## $ Block         : chr  "B1" "B1" "B1" "B1" ...
## $ Treatment     : chr  "Control" "Control" "Control" "Control" ...
## $ OTU           : chr  "X1" "X2" "X3" "X4" ...
## $ Species       : chr  "Alternaria_alternata" "Articulospora_tetracladia" "Cadophora_finlandica"
## $ Genus         : chr  "Alternaria" "Articulospora" "Cadophora" "Cenococcum" ...
## $ Family        : chr  "Pleosporaceae" "Helotiaceae" "Helotiaceae" "Gloniaceae" ...
## $ Order         : chr  "Pleosporales" "Helotiales" "Helotiales" "Mytilinidiales" ...
## $ Phylum      : chr  "Ascomycota" "Ascomycota" "Ascomycota" "Ascomycota" ...
## $ Relative.Abandance: num  0.0345 0 0 0.0345 0 ...
## $ EcM.density    : num  0.679 0.679 0.679 0.679 0.679 ...
## $ Absolute.Abandance: num  0.0234 0 0 0.0234 0 ...
```

```
#To slim down the data to only columns I will need
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
slim <- Projdata %>%
  select(Plot, Treatment, Species, Relative.Abandance)
slim
```

	Plot	Treatment	Species	Relative.Abandance
## 1	MAT.B1.C	Control	Alternaria_alternata	0.03448276
## 2	MAT.B1.C	Control	Articulospora_tetracladia	0.00000000
## 3	MAT.B1.C	Control	Cadophora_finlandica	0.00000000
## 4	MAT.B1.C	Control	Cenococcum_geophilum	0.03448276
## 5	MAT.B1.C	Control	Chalara_sp._1	0.00000000
## 6	MAT.B1.C	Control	Cortinarius_sp._1	0.37931034

## 7	MAT.B1.C	Control	Cortinarius_croceus_1	0.03448276
## 8	MAT.B1.C	Control	Cortinarius_glandicolor	0.03448276
## 9	MAT.B1.C	Control	Cortinarius_casimiri	0.00000000
## 10	MAT.B1.C	Control	Cortinarius_croceus_2	0.00000000
## 11	MAT.B1.C	Control	Cortinarius_herpeticus	0.00000000
## 12	MAT.B1.C	Control	Cortinarius_tabularis	0.00000000
## 13	MAT.B1.C	Control	Cortinarius_delibutus	0.00000000
## 14	MAT.B1.C	Control	Cortinarius_sp._2	0.00000000
## 15	MAT.B1.C	Control	Cortinarius_paragaudis	0.00000000
## 16	MAT.B1.C	Control	Cortinarius_sp._3	0.00000000
## 17	MAT.B1.C	Control	Helotiales_sp._1	0.00000000
## 18	MAT.B1.C	Control	Laccaria_laccata	0.00000000
## 19	MAT.B1.C	Control	Lactarius_vietus_1	0.10344828
## 20	MAT.B1.C	Control	Lactarius_glyciosmus	0.00000000
## 21	MAT.B1.C	Control	Lactarius_vietus_2	0.00000000
## 22	MAT.B1.C	Control	Leccinum_holopus	0.00000000
## 23	MAT.B1.C	Control	Leccinum_scabrum	0.00000000
## 24	MAT.B1.C	Control	Leccinum_variicolor_1	0.00000000
## 25	MAT.B1.C	Control	Leccinum_variicolor_2	0.00000000
## 26	MAT.B1.C	Control	Leccinum_variicolor_3	0.00000000
## 27	MAT.B1.C	Control	Leptodontidium_elatius	0.00000000
## 28	MAT.B1.C	Control	Meliniomyces_variabilis	0.00000000
## 29	MAT.B1.C	Control	Meliniomyces_bicolor	0.00000000
## 30	MAT.B1.C	Control	Phialocephala_fortinii	0.00000000
## 31	MAT.B1.C	Control	Pseudotomentella_sp._1	0.00000000
## 32	MAT.B1.C	Control	Rhizoscyphus_ericae	0.00000000
## 33	MAT.B1.C	Control	Thelephora_terrestris	0.00000000
## 34	MAT.B1.C	Control	Tomentella_sublilacina	0.00000000
## 35	MAT.B1.C	Control	Tomentella_sp._1	0.00000000
## 36	MAT.B1.C	Control	Tomentellopsis_submollis	0.00000000
## 37	MAT.B1.C	Control	Russula_chamiteae	0.00000000
## 38	MAT.B1.C	Control	Russula_decolorans	0.00000000
## 39	MAT.B1.C	Control	Russula_nitida/sphagnicola	0.00000000
## 40	MAT.B1.C	Control	Russula_sp._1	0.00000000
## 41	MAT.B1.C	Control	Russula_sp._2	0.00000000
## 42	MAT.B1.C	Control	Russula_sp._3	0.37931034
## 43	MAT.B1.C	Control	Russula_sp._4	0.00000000
## 44	MAT.B1.C	Control	Russula_vinosa	0.00000000
## 45	MAT.B1.F	Fertilized	Alternaria_alternata	0.00000000
## 46	MAT.B1.F	Fertilized	Articulospora_tetracladia	0.00000000
## 47	MAT.B1.F	Fertilized	Cadophora_finlandica	0.00000000
## 48	MAT.B1.F	Fertilized	Cenococcum_geophilum	0.00000000
## 49	MAT.B1.F	Fertilized	Chalara_sp._1	0.00000000
## 50	MAT.B1.F	Fertilized	Cortinarius_sp._1	0.00000000
## 51	MAT.B1.F	Fertilized	Cortinarius_croceus_1	0.00000000
## 52	MAT.B1.F	Fertilized	Cortinarius_glandicolor	0.00000000
## 53	MAT.B1.F	Fertilized	Cortinarius_casimiri	0.00000000
## 54	MAT.B1.F	Fertilized	Cortinarius_croceus_2	0.00000000
## 55	MAT.B1.F	Fertilized	Cortinarius_herpeticus	0.00000000
## 56	MAT.B1.F	Fertilized	Cortinarius_tabularis	0.00000000
## 57	MAT.B1.F	Fertilized	Cortinarius_delibutus	0.00000000
## 58	MAT.B1.F	Fertilized	Cortinarius_sp._2	0.00000000
## 59	MAT.B1.F	Fertilized	Cortinarius_paragaudis	0.00000000
## 60	MAT.B1.F	Fertilized	Cortinarius_sp._3	0.00000000

## 61	MAT.B1.F	Fertilized	Helotiales_sp._1	0.00000000
## 62	MAT.B1.F	Fertilized	Laccaria_laccata	0.00000000
## 63	MAT.B1.F	Fertilized	Lactarius_vietus_1	0.00000000
## 64	MAT.B1.F	Fertilized	Lactarius_glyciosmus	0.00000000
## 65	MAT.B1.F	Fertilized	Lactarius_vietus_2	0.00000000
## 66	MAT.B1.F	Fertilized	Leccinum_holopus	0.00000000
## 67	MAT.B1.F	Fertilized	Leccinum_scabrum	0.00000000
## 68	MAT.B1.F	Fertilized	Leccinum_variicolor_1	0.00000000
## 69	MAT.B1.F	Fertilized	Leccinum_variicolor_2	0.00000000
## 70	MAT.B1.F	Fertilized	Leccinum_variicolor_3	0.00000000
## 71	MAT.B1.F	Fertilized	Leptodontidium_elatius	0.00000000
## 72	MAT.B1.F	Fertilized	Meliniomyces_variabilis	0.00000000
## 73	MAT.B1.F	Fertilized	Meliniomyces_bicolor	0.00000000
## 74	MAT.B1.F	Fertilized	Phialocephala_fortinii	0.00000000
## 75	MAT.B1.F	Fertilized	Pseudotomentella_sp._1	0.00000000
## 76	MAT.B1.F	Fertilized	Rhizoscyphus_ericae	0.00000000
## 77	MAT.B1.F	Fertilized	Thelephora_terrestris	0.00000000
## 78	MAT.B1.F	Fertilized	Tomentella_sublilacina	0.00000000
## 79	MAT.B1.F	Fertilized	Tomentella_sp._1	0.00000000
## 80	MAT.B1.F	Fertilized	Tomentellopsis_submollis	0.02777778
## 81	MAT.B1.F	Fertilized	Russula_chamiteae	0.08333333
## 82	MAT.B1.F	Fertilized	Russula_decolorans	0.08333333
## 83	MAT.B1.F	Fertilized	Russula_nitida/sphagnicola	0.77777778
## 84	MAT.B1.F	Fertilized	Russula_sp._1	0.02777778
## 85	MAT.B1.F	Fertilized	Russula_sp._2	0.00000000
## 86	MAT.B1.F	Fertilized	Russula_sp._3	0.00000000
## 87	MAT.B1.F	Fertilized	Russula_sp._4	0.00000000
## 88	MAT.B1.F	Fertilized	Russula_vinosa	0.00000000
## 89	MAT.B1.W	Warmed	Alternaria_alternata	0.00000000
## 90	MAT.B1.W	Warmed	Articulospora_tetracladia	0.00000000
## 91	MAT.B1.W	Warmed	Cadophora_finlandica	0.00000000
## 92	MAT.B1.W	Warmed	Cenococcum_geophilum	0.00000000
## 93	MAT.B1.W	Warmed	Chalara_sp._1	0.00000000
## 94	MAT.B1.W	Warmed	Cortinarius_sp._1	0.00000000
## 95	MAT.B1.W	Warmed	Cortinarius_croceus_1	0.00000000
## 96	MAT.B1.W	Warmed	Cortinarius_glandicolor	0.00000000
## 97	MAT.B1.W	Warmed	Cortinarius_casimiri	0.00000000
## 98	MAT.B1.W	Warmed	Cortinarius_croceus_2	0.00000000
## 99	MAT.B1.W	Warmed	Cortinarius_herpeticus	0.00000000
## 100	MAT.B1.W	Warmed	Cortinarius_tabularis	0.00000000
## 101	MAT.B1.W	Warmed	Cortinarius_delibutus	0.00000000
## 102	MAT.B1.W	Warmed	Cortinarius_sp._2	0.00000000
## 103	MAT.B1.W	Warmed	Cortinarius_paragaudis	0.00000000
## 104	MAT.B1.W	Warmed	Cortinarius_sp._3	0.00000000
## 105	MAT.B1.W	Warmed	Helotiales_sp._1	0.00000000
## 106	MAT.B1.W	Warmed	Laccaria_laccata	0.00000000
## 107	MAT.B1.W	Warmed	Lactarius_vietus_1	0.00000000
## 108	MAT.B1.W	Warmed	Lactarius_glyciosmus	0.03225807
## 109	MAT.B1.W	Warmed	Lactarius_vietus_2	0.00000000
## 110	MAT.B1.W	Warmed	Leccinum_holopus	0.00000000
## 111	MAT.B1.W	Warmed	Leccinum_scabrum	0.00000000
## 112	MAT.B1.W	Warmed	Leccinum_variicolor_1	0.00000000
## 113	MAT.B1.W	Warmed	Leccinum_variicolor_2	0.00000000
## 114	MAT.B1.W	Warmed	Leccinum_variicolor_3	0.00000000

## 115	MAT.B1.W	Warmed	Leptodontidium_elatius	0.00000000
## 116	MAT.B1.W	Warmed	Meliniomyces_variabilis	0.00000000
## 117	MAT.B1.W	Warmed	Meliniomyces_bicolor	0.00000000
## 118	MAT.B1.W	Warmed	Phialocephala_fortinii	0.00000000
## 119	MAT.B1.W	Warmed	Pseudotomentella_sp._1	0.00000000
## 120	MAT.B1.W	Warmed	Rhizoscyphus_ericae	0.00000000
## 121	MAT.B1.W	Warmed	Thelephora_terrestris	0.00000000
## 122	MAT.B1.W	Warmed	Tomentella_sublilacina	0.00000000
## 123	MAT.B1.W	Warmed	Tomentella_sp._1	0.00000000
## 124	MAT.B1.W	Warmed	Tomentellopsis_submollis	0.06451613
## 125	MAT.B1.W	Warmed	Russula_chamiteae	0.16129032
## 126	MAT.B1.W	Warmed	Russula_decolorans	0.38709677
## 127	MAT.B1.W	Warmed	Russula_nitida/sphagnicola	0.16129032
## 128	MAT.B1.W	Warmed	Russula_sp._1	0.09677419
## 129	MAT.B1.W	Warmed	Russula_sp._2	0.00000000
## 130	MAT.B1.W	Warmed	Russula_sp._3	0.09677419
## 131	MAT.B1.W	Warmed	Russula_sp._4	0.00000000
## 132	MAT.B1.W	Warmed	Russula_vinosa	0.00000000
## 133	MAT.B1.WF	Warmed.Fertilized	Alternaria_alternata	0.00000000
## 134	MAT.B1.WF	Warmed.Fertilized	Articulospora_tetracladia	0.00000000
## 135	MAT.B1.WF	Warmed.Fertilized	Cadophora_finlandica	0.00000000
## 136	MAT.B1.WF	Warmed.Fertilized	Cenococcum_geophilum	0.00000000
## 137	MAT.B1.WF	Warmed.Fertilized	Chalara_sp._1	0.02857143
## 138	MAT.B1.WF	Warmed.Fertilized	Cortinarius_sp._1	0.00000000
## 139	MAT.B1.WF	Warmed.Fertilized	Cortinarius_croceus_1	0.00000000
## 140	MAT.B1.WF	Warmed.Fertilized	Cortinarius_glandicolor	0.00000000
## 141	MAT.B1.WF	Warmed.Fertilized	Cortinarius_casimiri	0.00000000
## 142	MAT.B1.WF	Warmed.Fertilized	Cortinarius_croceus_2	0.00000000
## 143	MAT.B1.WF	Warmed.Fertilized	Cortinarius_herpeticus	0.00000000
## 144	MAT.B1.WF	Warmed.Fertilized	Cortinarius_tabularis	0.00000000
## 145	MAT.B1.WF	Warmed.Fertilized	Cortinarius_delibutus	0.00000000
## 146	MAT.B1.WF	Warmed.Fertilized	Cortinarius_sp._2	0.00000000
## 147	MAT.B1.WF	Warmed.Fertilized	Cortinarius_paragaudis	0.00000000
## 148	MAT.B1.WF	Warmed.Fertilized	Cortinarius_sp._3	0.00000000
## 149	MAT.B1.WF	Warmed.Fertilized	Helotiales_sp._1	0.02857143
## 150	MAT.B1.WF	Warmed.Fertilized	Laccaria_laccata	0.40000000
## 151	MAT.B1.WF	Warmed.Fertilized	Lactarius_vietus_1	0.00000000
## 152	MAT.B1.WF	Warmed.Fertilized	Lactarius_glyciosmus	0.00000000
## 153	MAT.B1.WF	Warmed.Fertilized	Lactarius_vietus_2	0.00000000
## 154	MAT.B1.WF	Warmed.Fertilized	Leccinum_holopus	0.00000000
## 155	MAT.B1.WF	Warmed.Fertilized	Leccinum_scabrum	0.00000000
## 156	MAT.B1.WF	Warmed.Fertilized	Leccinum_variicolor_1	0.00000000
## 157	MAT.B1.WF	Warmed.Fertilized	Leccinum_variicolor_2	0.00000000
## 158	MAT.B1.WF	Warmed.Fertilized	Leccinum_variicolor_3	0.00000000
## 159	MAT.B1.WF	Warmed.Fertilized	Leptodontidium_elatius	0.00000000
## 160	MAT.B1.WF	Warmed.Fertilized	Meliniomyces_variabilis	0.00000000
## 161	MAT.B1.WF	Warmed.Fertilized	Meliniomyces_bicolor	0.00000000
## 162	MAT.B1.WF	Warmed.Fertilized	Phialocephala_fortinii	0.00000000
## 163	MAT.B1.WF	Warmed.Fertilized	Pseudotomentella_sp._1	0.00000000
## 164	MAT.B1.WF	Warmed.Fertilized	Rhizoscyphus_ericae	0.00000000
## 165	MAT.B1.WF	Warmed.Fertilized	Thelephora_terrestris	0.14285714
## 166	MAT.B1.WF	Warmed.Fertilized	Tomentella_sublilacina	0.00000000
## 167	MAT.B1.WF	Warmed.Fertilized	Tomentella_sp._1	0.00000000
## 168	MAT.B1.WF	Warmed.Fertilized	Tomentellopsis_submollis	0.20000000

## 169	MAT.B1.WF	Warmed.Fertilized	Russula_chamiteae	0.00000000
## 170	MAT.B1.WF	Warmed.Fertilized	Russula_decolorans	0.00000000
## 171	MAT.B1.WF	Warmed.Fertilized	Russula_nitida/sphagnicola	0.20000000
## 172	MAT.B1.WF	Warmed.Fertilized	Russula_sp._1	0.00000000
## 173	MAT.B1.WF	Warmed.Fertilized	Russula_sp._2	0.00000000
## 174	MAT.B1.WF	Warmed.Fertilized	Russula_sp._3	0.00000000
## 175	MAT.B1.WF	Warmed.Fertilized	Russula_sp._4	0.00000000
## 176	MAT.B1.WF	Warmed.Fertilized	Russula_vinosa	0.00000000
## 177	MAT.B2.C	Control	Alternaria_alternata	0.00000000
## 178	MAT.B2.C	Control	Articulospora_tetracladia	0.00000000
## 179	MAT.B2.C	Control	Cadophora_finlandica	0.00000000
## 180	MAT.B2.C	Control	Cenococcum_geophilum	0.09677419
## 181	MAT.B2.C	Control	Chalara_sp._1	0.00000000
## 182	MAT.B2.C	Control	Cortinarius_sp._1	0.00000000
## 183	MAT.B2.C	Control	Cortinarius_croceus_1	0.00000000
## 184	MAT.B2.C	Control	Cortinarius_glandicolor	0.00000000
## 185	MAT.B2.C	Control	Cortinarius_casimiri	0.00000000
## 186	MAT.B2.C	Control	Cortinarius_croceus_2	0.00000000
## 187	MAT.B2.C	Control	Cortinarius_herpeticus	0.00000000
## 188	MAT.B2.C	Control	Cortinarius_tabularis	0.00000000
## 189	MAT.B2.C	Control	Cortinarius_delibutus	0.00000000
## 190	MAT.B2.C	Control	Cortinarius_sp._2	0.00000000
## 191	MAT.B2.C	Control	Cortinarius_paragaudis	0.00000000
## 192	MAT.B2.C	Control	Cortinarius_sp._3	0.00000000
## 193	MAT.B2.C	Control	Helotiales_sp._1	0.00000000
## 194	MAT.B2.C	Control	Laccaria_laccata	0.00000000
## 195	MAT.B2.C	Control	Lactarius_vietus_1	0.29032258
## 196	MAT.B2.C	Control	Lactarius_glyciosmus	0.06451613
## 197	MAT.B2.C	Control	Lactarius_vietus_2	0.00000000
## 198	MAT.B2.C	Control	Leccinum_holopus	0.03225807
## 199	MAT.B2.C	Control	Leccinum_scabrum	0.00000000
## 200	MAT.B2.C	Control	Leccinum_variicolor_1	0.03225807
## 201	MAT.B2.C	Control	Leccinum_variicolor_2	0.00000000
## 202	MAT.B2.C	Control	Leccinum_variicolor_3	0.03225807
## 203	MAT.B2.C	Control	Leptodontidium_elatius	0.00000000
## 204	MAT.B2.C	Control	Meliniomyces_variabilis	0.03225807
## 205	MAT.B2.C	Control	Meliniomyces_bicolor	0.00000000
## 206	MAT.B2.C	Control	Phialocephala_fortinii	0.03225807
## 207	MAT.B2.C	Control	Pseudotomentella_sp._1	0.03225807
## 208	MAT.B2.C	Control	Rhizoscyphus_ericae	0.00000000
## 209	MAT.B2.C	Control	Thelephora_terrestris	0.00000000
## 210	MAT.B2.C	Control	Tomentella_sublilacina	0.00000000
## 211	MAT.B2.C	Control	Tomentella_sp._1	0.00000000
## 212	MAT.B2.C	Control	Tomentellopsis_submollis	0.00000000
## 213	MAT.B2.C	Control	Russula_chamiteae	0.03225807
## 214	MAT.B2.C	Control	Russula_decolorans	0.25806452
## 215	MAT.B2.C	Control	Russula_nitida/sphagnicola	0.00000000
## 216	MAT.B2.C	Control	Russula_sp._1	0.00000000
## 217	MAT.B2.C	Control	Russula_sp._2	0.00000000
## 218	MAT.B2.C	Control	Russula_sp._3	0.06451613
## 219	MAT.B2.C	Control	Russula_sp._4	0.00000000
## 220	MAT.B2.C	Control	Russula_vinosa	0.00000000
## 221	MAT.B2.F	Fertilized	Alternaria_alternata	0.00000000
## 222	MAT.B2.F	Fertilized	Articulospora_tetracladia	0.03571429

## 223	MAT.B2.F	Fertilized	Cadophora_finlandica	0.00000000
## 224	MAT.B2.F	Fertilized	Cenococcum_geophilum	0.00000000
## 225	MAT.B2.F	Fertilized	Chalara_sp._1	0.00000000
## 226	MAT.B2.F	Fertilized	Cortinarius_sp._1	0.00000000
## 227	MAT.B2.F	Fertilized	Cortinarius_croceus_1	0.00000000
## 228	MAT.B2.F	Fertilized	Cortinarius_glandicolor	0.00000000
## 229	MAT.B2.F	Fertilized	Cortinarius_casimiri	0.00000000
## 230	MAT.B2.F	Fertilized	Cortinarius_croceus_2	0.00000000
## 231	MAT.B2.F	Fertilized	Cortinarius_herpeticus	0.00000000
## 232	MAT.B2.F	Fertilized	Cortinarius_tabularis	0.00000000
## 233	MAT.B2.F	Fertilized	Cortinarius_delibutus	0.00000000
## 234	MAT.B2.F	Fertilized	Cortinarius_sp._2	0.00000000
## 235	MAT.B2.F	Fertilized	Cortinarius_paragaudis	0.00000000
## 236	MAT.B2.F	Fertilized	Cortinarius_sp._3	0.00000000
## 237	MAT.B2.F	Fertilized	Helotiales_sp._1	0.00000000
## 238	MAT.B2.F	Fertilized	Laccaria_laccata	0.00000000
## 239	MAT.B2.F	Fertilized	Lactarius_vietus_1	0.03571429
## 240	MAT.B2.F	Fertilized	Lactarius_glyciosmus	0.00000000
## 241	MAT.B2.F	Fertilized	Lactarius_vietus_2	0.00000000
## 242	MAT.B2.F	Fertilized	Leccinum_holopus	0.00000000
## 243	MAT.B2.F	Fertilized	Leccinum_scabrum	0.00000000
## 244	MAT.B2.F	Fertilized	Leccinum_variicolor_1	0.00000000
## 245	MAT.B2.F	Fertilized	Leccinum_variicolor_2	0.00000000
## 246	MAT.B2.F	Fertilized	Leccinum_variicolor_3	0.00000000
## 247	MAT.B2.F	Fertilized	Leptodontidium_elatius	0.00000000
## 248	MAT.B2.F	Fertilized	Meliniomyces_variabilis	0.03571429
## 249	MAT.B2.F	Fertilized	Meliniomyces_bicolor	0.00000000
## 250	MAT.B2.F	Fertilized	Phialocephala_fortinii	0.00000000
## 251	MAT.B2.F	Fertilized	Pseudotomentella_sp._1	0.03571429
## 252	MAT.B2.F	Fertilized	Rhizoscyphus_ericae	0.03571429
## 253	MAT.B2.F	Fertilized	Thelephora_terrestris	0.67857143
## 254	MAT.B2.F	Fertilized	Tomentella_sublilacina	0.00000000
## 255	MAT.B2.F	Fertilized	Tomentella_sp._1	0.00000000
## 256	MAT.B2.F	Fertilized	Tomentellopsis_submollis	0.14285714
## 257	MAT.B2.F	Fertilized	Russula_chamiteae	0.00000000
## 258	MAT.B2.F	Fertilized	Russula_decolorans	0.00000000
## 259	MAT.B2.F	Fertilized	Russula_nitida/sphagnicola	0.00000000
## 260	MAT.B2.F	Fertilized	Russula_sp._1	0.00000000
## 261	MAT.B2.F	Fertilized	Russula_sp._2	0.00000000
## 262	MAT.B2.F	Fertilized	Russula_sp._3	0.00000000
## 263	MAT.B2.F	Fertilized	Russula_sp._4	0.00000000
## 264	MAT.B2.F	Fertilized	Russula_vinosa	0.00000000
## 265	MAT.B2.W	Warmed	Alternaria_alternata	0.00000000
## 266	MAT.B2.W	Warmed	Articulospora_tetracladia	0.00000000
## 267	MAT.B2.W	Warmed	Cadophora_finlandica	0.00000000
## 268	MAT.B2.W	Warmed	Cenococcum_geophilum	0.15151515
## 269	MAT.B2.W	Warmed	Chalara_sp._1	0.00000000
## 270	MAT.B2.W	Warmed	Cortinarius_sp._1	0.00000000
## 271	MAT.B2.W	Warmed	Cortinarius_croceus_1	0.03030303
## 272	MAT.B2.W	Warmed	Cortinarius_glandicolor	0.00000000
## 273	MAT.B2.W	Warmed	Cortinarius_casimiri	0.00000000
## 274	MAT.B2.W	Warmed	Cortinarius_croceus_2	0.00000000
## 275	MAT.B2.W	Warmed	Cortinarius_herpeticus	0.00000000
## 276	MAT.B2.W	Warmed	Cortinarius_tabularis	0.00000000

## 277	MAT.B2.W	Warmed	Cortinarius_delibutus	0.00000000
## 278	MAT.B2.W	Warmed	Cortinarius_sp._2	0.00000000
## 279	MAT.B2.W	Warmed	Cortinarius_paragaudis	0.00000000
## 280	MAT.B2.W	Warmed	Cortinarius_sp._3	0.00000000
## 281	MAT.B2.W	Warmed	Helotiales_sp._1	0.00000000
## 282	MAT.B2.W	Warmed	Laccaria_laccata	0.00000000
## 283	MAT.B2.W	Warmed	Lactarius_vietus_1	0.03030303
## 284	MAT.B2.W	Warmed	Lactarius_glyciosmus	0.03030303
## 285	MAT.B2.W	Warmed	Lactarius_vietus_2	0.00000000
## 286	MAT.B2.W	Warmed	Leccinum_holopus	0.00000000
## 287	MAT.B2.W	Warmed	Leccinum_scabrum	0.00000000
## 288	MAT.B2.W	Warmed	Leccinum_variicolor_1	0.00000000
## 289	MAT.B2.W	Warmed	Leccinum_variicolor_2	0.00000000
## 290	MAT.B2.W	Warmed	Leccinum_variicolor_3	0.00000000
## 291	MAT.B2.W	Warmed	Leptodontidium_elatius	0.00000000
## 292	MAT.B2.W	Warmed	Meliniomyces_variabilis	0.03030303
## 293	MAT.B2.W	Warmed	Meliniomyces_bicolor	0.00000000
## 294	MAT.B2.W	Warmed	Phialocephala_fortinii	0.00000000
## 295	MAT.B2.W	Warmed	Pseudotomentella_sp._1	0.00000000
## 296	MAT.B2.W	Warmed	Rhizoscyphus_ericae	0.03030303
## 297	MAT.B2.W	Warmed	Thelephora_terrestris	0.00000000
## 298	MAT.B2.W	Warmed	Tomentella_sublilacina	0.00000000
## 299	MAT.B2.W	Warmed	Tomentella_sp._1	0.00000000
## 300	MAT.B2.W	Warmed	Tomentellopsis_submollis	0.00000000
## 301	MAT.B2.W	Warmed	Russula_chamiteae	0.00000000
## 302	MAT.B2.W	Warmed	Russula_decolorans	0.42424242
## 303	MAT.B2.W	Warmed	Russula_nitida/sphagnicola	0.03030303
## 304	MAT.B2.W	Warmed	Russula_sp._1	0.00000000
## 305	MAT.B2.W	Warmed	Russula_sp._2	0.00000000
## 306	MAT.B2.W	Warmed	Russula_sp._3	0.24242424
## 307	MAT.B2.W	Warmed	Russula_sp._4	0.00000000
## 308	MAT.B2.W	Warmed	Russula_vinosa	0.00000000
## 309	MAT.B2.WF	Warmed.Fertilized	Alternaria_alternata	0.00000000
## 310	MAT.B2.WF	Warmed.Fertilized	Articulospora_tetracladia	0.00000000
## 311	MAT.B2.WF	Warmed.Fertilized	Cadophora_finlandica	0.04761905
## 312	MAT.B2.WF	Warmed.Fertilized	Cenococcum_geophilum	0.00000000
## 313	MAT.B2.WF	Warmed.Fertilized	Chalara_sp._1	0.04761905
## 314	MAT.B2.WF	Warmed.Fertilized	Cortinarius_sp._1	0.00000000
## 315	MAT.B2.WF	Warmed.Fertilized	Cortinarius_croceus_1	0.00000000
## 316	MAT.B2.WF	Warmed.Fertilized	Cortinarius_glandicolor	0.00000000
## 317	MAT.B2.WF	Warmed.Fertilized	Cortinarius_casimiri	0.04761905
## 318	MAT.B2.WF	Warmed.Fertilized	Cortinarius_croceus_2	0.00000000
## 319	MAT.B2.WF	Warmed.Fertilized	Cortinarius_herpeticus	0.00000000
## 320	MAT.B2.WF	Warmed.Fertilized	Cortinarius_tabularis	0.00000000
## 321	MAT.B2.WF	Warmed.Fertilized	Cortinarius_delibutus	0.00000000
## 322	MAT.B2.WF	Warmed.Fertilized	Cortinarius_sp._2	0.00000000
## 323	MAT.B2.WF	Warmed.Fertilized	Cortinarius_paragaudis	0.00000000
## 324	MAT.B2.WF	Warmed.Fertilized	Cortinarius_sp._3	0.00000000
## 325	MAT.B2.WF	Warmed.Fertilized	Helotiales_sp._1	0.04761905
## 326	MAT.B2.WF	Warmed.Fertilized	Laccaria_laccata	0.04761905
## 327	MAT.B2.WF	Warmed.Fertilized	Lactarius_vietus_1	0.04761905
## 328	MAT.B2.WF	Warmed.Fertilized	Lactarius_glyciosmus	0.14285714
## 329	MAT.B2.WF	Warmed.Fertilized	Lactarius_vietus_2	0.00000000
## 330	MAT.B2.WF	Warmed.Fertilized	Leccinum_holopus	0.00000000

## 331	MAT.B2.WF	Warmed.Fertilized	Leccinum_scabrum	0.00000000
## 332	MAT.B2.WF	Warmed.Fertilized	Leccinum_variicolor_1	0.00000000
## 333	MAT.B2.WF	Warmed.Fertilized	Leccinum_variicolor_2	0.00000000
## 334	MAT.B2.WF	Warmed.Fertilized	Leccinum_variicolor_3	0.00000000
## 335	MAT.B2.WF	Warmed.Fertilized	Leptodontidium_elatius	0.00000000
## 336	MAT.B2.WF	Warmed.Fertilized	Meliniomyces_variabilis	0.00000000
## 337	MAT.B2.WF	Warmed.Fertilized	Meliniomyces_bicolor	0.04761905
## 338	MAT.B2.WF	Warmed.Fertilized	Phialocephala_fortinii	0.00000000
## 339	MAT.B2.WF	Warmed.Fertilized	Pseudotomentella_sp._1	0.14285714
## 340	MAT.B2.WF	Warmed.Fertilized	Rhizoscyphus_ericae	0.04761905
## 341	MAT.B2.WF	Warmed.Fertilized	Thelephora_terrestris	0.09523809
## 342	MAT.B2.WF	Warmed.Fertilized	Tomentella_sublilacina	0.00000000
## 343	MAT.B2.WF	Warmed.Fertilized	Tomentella_sp._1	0.00000000
## 344	MAT.B2.WF	Warmed.Fertilized	Tomentellopsis_submollis	0.09523809
## 345	MAT.B2.WF	Warmed.Fertilized	Russula_chamiteae	0.00000000
## 346	MAT.B2.WF	Warmed.Fertilized	Russula_decolorans	0.04761905
## 347	MAT.B2.WF	Warmed.Fertilized	Russula_nitida/sphagnicola	0.09523809
## 348	MAT.B2.WF	Warmed.Fertilized	Russula_sp._1	0.00000000
## 349	MAT.B2.WF	Warmed.Fertilized	Russula_sp._2	0.00000000
## 350	MAT.B2.WF	Warmed.Fertilized	Russula_sp._3	0.00000000
## 351	MAT.B2.WF	Warmed.Fertilized	Russula_sp._4	0.00000000
## 352	MAT.B2.WF	Warmed.Fertilized	Russula_vinosa	0.00000000
## 353	MAT.B3.C	Control	Alternaria_alternata	0.00000000
## 354	MAT.B3.C	Control	Articulospora_tetracladia	0.00000000
## 355	MAT.B3.C	Control	Cadophora_finlandica	0.00000000
## 356	MAT.B3.C	Control	Cenococcum_geophilum	0.00000000
## 357	MAT.B3.C	Control	Chalara_sp._1	0.00000000
## 358	MAT.B3.C	Control	Cortinarius_sp._1	0.11764706
## 359	MAT.B3.C	Control	Cortinarius_croceus_1	0.00000000
## 360	MAT.B3.C	Control	Cortinarius_glandicolor	0.00000000
## 361	MAT.B3.C	Control	Cortinarius_casimiri	0.00000000
## 362	MAT.B3.C	Control	Cortinarius_croceus_2	0.17647059
## 363	MAT.B3.C	Control	Cortinarius_herpeticus	0.00000000
## 364	MAT.B3.C	Control	Cortinarius_tabularis	0.00000000
## 365	MAT.B3.C	Control	Cortinarius_delibutus	0.00000000
## 366	MAT.B3.C	Control	Cortinarius_sp._2	0.00000000
## 367	MAT.B3.C	Control	Cortinarius_paragaudis	0.00000000
## 368	MAT.B3.C	Control	Cortinarius_sp._3	0.00000000
## 369	MAT.B3.C	Control	Helotiales_sp._1	0.00000000
## 370	MAT.B3.C	Control	Laccaria_laccata	0.00000000
## 371	MAT.B3.C	Control	Lactarius_vietus_1	0.17647059
## 372	MAT.B3.C	Control	Lactarius_glyciosmus	0.00000000
## 373	MAT.B3.C	Control	Lactarius_vietus_2	0.00000000
## 374	MAT.B3.C	Control	Leccinum_holopus	0.00000000
## 375	MAT.B3.C	Control	Leccinum_scabrum	0.00000000
## 376	MAT.B3.C	Control	Leccinum_variicolor_1	0.00000000
## 377	MAT.B3.C	Control	Leccinum_variicolor_2	0.00000000
## 378	MAT.B3.C	Control	Leccinum_variicolor_3	0.00000000
## 379	MAT.B3.C	Control	Leptodontidium_elatius	0.00000000
## 380	MAT.B3.C	Control	Meliniomyces_variabilis	0.00000000
## 381	MAT.B3.C	Control	Meliniomyces_bicolor	0.00000000
## 382	MAT.B3.C	Control	Phialocephala_fortinii	0.00000000
## 383	MAT.B3.C	Control	Pseudotomentella_sp._1	0.00000000
## 384	MAT.B3.C	Control	Rhizoscyphus_ericae	0.00000000

## 385	MAT.B3.C	Control	Thelephora_terrestris	0.00000000
## 386	MAT.B3.C	Control	Tomentella_sublilacina	0.00000000
## 387	MAT.B3.C	Control	Tomentella_sp._1	0.00000000
## 388	MAT.B3.C	Control	Tomentellopsis_submollis	0.00000000
## 389	MAT.B3.C	Control	Russula_chamiteae	0.00000000
## 390	MAT.B3.C	Control	Russula_decolorans	0.29411765
## 391	MAT.B3.C	Control	Russula_nitida/sphagnicola	0.05882353
## 392	MAT.B3.C	Control	Russula_sp._1	0.17647059
## 393	MAT.B3.C	Control	Russula_sp._2	0.00000000
## 394	MAT.B3.C	Control	Russula_sp._3	0.00000000
## 395	MAT.B3.C	Control	Russula_sp._4	0.00000000
## 396	MAT.B3.C	Control	Russula_vinosa	0.00000000
## 397	MAT.B3.F	Fertilized	Alternaria_alternata	0.00000000
## 398	MAT.B3.F	Fertilized	Articulospora_tetracladia	0.00000000
## 399	MAT.B3.F	Fertilized	Cadophora_finlandica	0.00000000
## 400	MAT.B3.F	Fertilized	Cenococcum_geophilum	0.00000000
## 401	MAT.B3.F	Fertilized	Chalara_sp._1	0.00000000
## 402	MAT.B3.F	Fertilized	Cortinarius_sp._1	0.00000000
## 403	MAT.B3.F	Fertilized	Cortinarius_croceus_1	0.00000000
## 404	MAT.B3.F	Fertilized	Cortinarius_glandicolor	0.00000000
## 405	MAT.B3.F	Fertilized	Cortinarius_casimiri	0.00000000
## 406	MAT.B3.F	Fertilized	Cortinarius_croceus_2	0.00000000
## 407	MAT.B3.F	Fertilized	Cortinarius_herpeticus	0.00000000
## 408	MAT.B3.F	Fertilized	Cortinarius_tabularis	0.00000000
## 409	MAT.B3.F	Fertilized	Cortinarius_delibutus	0.00000000
## 410	MAT.B3.F	Fertilized	Cortinarius_sp._2	0.00000000
## 411	MAT.B3.F	Fertilized	Cortinarius_paragaudis	0.00000000
## 412	MAT.B3.F	Fertilized	Cortinarius_sp._3	0.00000000
## 413	MAT.B3.F	Fertilized	Helotiales_sp._1	0.00000000
## 414	MAT.B3.F	Fertilized	Laccaria_laccata	0.06666667
## 415	MAT.B3.F	Fertilized	Lactarius_vietus_1	0.06666667
## 416	MAT.B3.F	Fertilized	Lactarius_glyciosmus	0.13333333
## 417	MAT.B3.F	Fertilized	Lactarius_vietus_2	0.00000000
## 418	MAT.B3.F	Fertilized	Leccinum_holopus	0.00000000
## 419	MAT.B3.F	Fertilized	Leccinum_scabrum	0.00000000
## 420	MAT.B3.F	Fertilized	Leccinum_variicolor_1	0.00000000
## 421	MAT.B3.F	Fertilized	Leccinum_variicolor_2	0.00000000
## 422	MAT.B3.F	Fertilized	Leccinum_variicolor_3	0.00000000
## 423	MAT.B3.F	Fertilized	Leptodontidium_elatius	0.00000000
## 424	MAT.B3.F	Fertilized	Meliniomyces_variabilis	0.00000000
## 425	MAT.B3.F	Fertilized	Meliniomyces_bicolor	0.00000000
## 426	MAT.B3.F	Fertilized	Phialocephala_fortinii	0.00000000
## 427	MAT.B3.F	Fertilized	Pseudotomentella_sp._1	0.00000000
## 428	MAT.B3.F	Fertilized	Rhizoscyphus_ericae	0.00000000
## 429	MAT.B3.F	Fertilized	Thelephora_terrestris	0.00000000
## 430	MAT.B3.F	Fertilized	Tomentella_sublilacina	0.00000000
## 431	MAT.B3.F	Fertilized	Tomentella_sp._1	0.00000000
## 432	MAT.B3.F	Fertilized	Tomentellopsis_submollis	0.06666667
## 433	MAT.B3.F	Fertilized	Russula_chamiteae	0.00000000
## 434	MAT.B3.F	Fertilized	Russula_decolorans	0.00000000
## 435	MAT.B3.F	Fertilized	Russula_nitida/sphagnicola	0.66666667
## 436	MAT.B3.F	Fertilized	Russula_sp._1	0.00000000
## 437	MAT.B3.F	Fertilized	Russula_sp._2	0.00000000
## 438	MAT.B3.F	Fertilized	Russula_sp._3	0.00000000

## 439	MAT.B3.F	Fertilized	Russula_sp._4	0.00000000
## 440	MAT.B3.F	Fertilized	Russula_vinosa	0.00000000
## 441	MAT.B3.W	Warmed	Alternaria_alternata	0.00000000
## 442	MAT.B3.W	Warmed	Articulospora_tetracladia	0.00000000
## 443	MAT.B3.W	Warmed	Cadophora_finlandica	0.00000000
## 444	MAT.B3.W	Warmed	Cenococcum_geophilum	0.00000000
## 445	MAT.B3.W	Warmed	Chalara_sp._1	0.00000000
## 446	MAT.B3.W	Warmed	Cortinarius_sp._1	0.00000000
## 447	MAT.B3.W	Warmed	Cortinarius_croceus_1	0.00000000
## 448	MAT.B3.W	Warmed	Cortinarius_glandicolor	0.00000000
## 449	MAT.B3.W	Warmed	Cortinarius_casimiri	0.00000000
## 450	MAT.B3.W	Warmed	Cortinarius_croceus_2	0.00000000
## 451	MAT.B3.W	Warmed	Cortinarius_herpeticus	0.00000000
## 452	MAT.B3.W	Warmed	Cortinarius_tabularis	0.00000000
## 453	MAT.B3.W	Warmed	Cortinarius_delibutus	0.00000000
## 454	MAT.B3.W	Warmed	Cortinarius_sp._2	0.00000000
## 455	MAT.B3.W	Warmed	Cortinarius_paragaudis	0.00000000
## 456	MAT.B3.W	Warmed	Cortinarius_sp._3	0.00000000
## 457	MAT.B3.W	Warmed	Helotiales_sp._1	0.00000000
## 458	MAT.B3.W	Warmed	Laccaria_laccata	0.00000000
## 459	MAT.B3.W	Warmed	Lactarius_vietus_1	0.00000000
## 460	MAT.B3.W	Warmed	Lactarius_glyciosmus	0.11111111
## 461	MAT.B3.W	Warmed	Lactarius_vietus_2	0.00000000
## 462	MAT.B3.W	Warmed	Leccinum_holopus	0.00000000
## 463	MAT.B3.W	Warmed	Leccinum_scabrum	0.03703704
## 464	MAT.B3.W	Warmed	Leccinum_variicolor_1	0.00000000
## 465	MAT.B3.W	Warmed	Leccinum_variicolor_2	0.00000000
## 466	MAT.B3.W	Warmed	Leccinum_variicolor_3	0.00000000
## 467	MAT.B3.W	Warmed	Leptodontidium_elatius	0.07407407
## 468	MAT.B3.W	Warmed	Meliniomyces_variabilis	0.00000000
## 469	MAT.B3.W	Warmed	Meliniomyces_bicolor	0.07407407
## 470	MAT.B3.W	Warmed	Phialocephala_fortinii	0.00000000
## 471	MAT.B3.W	Warmed	Pseudotomentella_sp._1	0.00000000
## 472	MAT.B3.W	Warmed	Rhizoscyphus_ericae	0.00000000
## 473	MAT.B3.W	Warmed	Thelephora_terrestris	0.00000000
## 474	MAT.B3.W	Warmed	Tomentella_sublilacina	0.00000000
## 475	MAT.B3.W	Warmed	Tomentella_sp._1	0.00000000
## 476	MAT.B3.W	Warmed	Tomentellopsis_submollis	0.03703704
## 477	MAT.B3.W	Warmed	Russula_chamiteae	0.00000000
## 478	MAT.B3.W	Warmed	Russula_decolorans	0.00000000
## 479	MAT.B3.W	Warmed	Russula_nitida/sphagnicola	0.62962963
## 480	MAT.B3.W	Warmed	Russula_sp._1	0.00000000
## 481	MAT.B3.W	Warmed	Russula_sp._2	0.00000000
## 482	MAT.B3.W	Warmed	Russula_sp._3	0.00000000
## 483	MAT.B3.W	Warmed	Russula_sp._4	0.00000000
## 484	MAT.B3.W	Warmed	Russula_vinosa	0.03703704
## 485	MAT.B3.WF	Warmed.Fertilized	Alternaria_alternata	0.00000000
## 486	MAT.B3.WF	Warmed.Fertilized	Articulospora_tetracladia	0.00000000
## 487	MAT.B3.WF	Warmed.Fertilized	Cadophora_finlandica	0.00000000
## 488	MAT.B3.WF	Warmed.Fertilized	Cenococcum_geophilum	0.00000000
## 489	MAT.B3.WF	Warmed.Fertilized	Chalara_sp._1	0.00000000
## 490	MAT.B3.WF	Warmed.Fertilized	Cortinarius_sp._1	0.00000000
## 491	MAT.B3.WF	Warmed.Fertilized	Cortinarius_croceus_1	0.00000000
## 492	MAT.B3.WF	Warmed.Fertilized	Cortinarius_glandicolor	0.00000000

## 493	MAT.B3.WF	Warmed.Fertilized	Cortinarius_casimiri	0.00000000
## 494	MAT.B3.WF	Warmed.Fertilized	Cortinarius_croceus_2	0.00000000
## 495	MAT.B3.WF	Warmed.Fertilized	Cortinarius_herpeticus	0.00000000
## 496	MAT.B3.WF	Warmed.Fertilized	Cortinarius_tabularis	0.00000000
## 497	MAT.B3.WF	Warmed.Fertilized	Cortinarius_delibutus	0.00000000
## 498	MAT.B3.WF	Warmed.Fertilized	Cortinarius_sp._2	0.00000000
## 499	MAT.B3.WF	Warmed.Fertilized	Cortinarius_paragaudis	0.00000000
## 500	MAT.B3.WF	Warmed.Fertilized	Cortinarius_sp._3	0.00000000
## 501	MAT.B3.WF	Warmed.Fertilized	Helotiales_sp._1	0.00000000
## 502	MAT.B3.WF	Warmed.Fertilized	Laccaria_laccata	0.14285714
## 503	MAT.B3.WF	Warmed.Fertilized	Lactarius_vietus_1	0.00000000
## 504	MAT.B3.WF	Warmed.Fertilized	Lactarius_glyciosmus	0.25000000
## 505	MAT.B3.WF	Warmed.Fertilized	Lactarius_vietus_2	0.00000000
## 506	MAT.B3.WF	Warmed.Fertilized	Leccinum_holopus	0.00000000
## 507	MAT.B3.WF	Warmed.Fertilized	Leccinum_scabrum	0.00000000
## 508	MAT.B3.WF	Warmed.Fertilized	Leccinum_variicolor_1	0.00000000
## 509	MAT.B3.WF	Warmed.Fertilized	Leccinum_variicolor_2	0.00000000
## 510	MAT.B3.WF	Warmed.Fertilized	Leccinum_variicolor_3	0.00000000
## 511	MAT.B3.WF	Warmed.Fertilized	Leptodontidium_elatius	0.00000000
## 512	MAT.B3.WF	Warmed.Fertilized	Meliniomyces_variabilis	0.00000000
## 513	MAT.B3.WF	Warmed.Fertilized	Meliniomyces_bicolor	0.00000000
## 514	MAT.B3.WF	Warmed.Fertilized	Phialocephala_fortinii	0.00000000
## 515	MAT.B3.WF	Warmed.Fertilized	Pseudotomentella_sp._1	0.00000000
## 516	MAT.B3.WF	Warmed.Fertilized	Rhizoscyphus_ericae	0.00000000
## 517	MAT.B3.WF	Warmed.Fertilized	Thelephora_terrestris	0.00000000
## 518	MAT.B3.WF	Warmed.Fertilized	Tomentella_sublilacina	0.00000000
## 519	MAT.B3.WF	Warmed.Fertilized	Tomentella_sp._1	0.00000000
## 520	MAT.B3.WF	Warmed.Fertilized	Tomentellopsis_submollis	0.07142857
## 521	MAT.B3.WF	Warmed.Fertilized	Russula_chamiteae	0.00000000
## 522	MAT.B3.WF	Warmed.Fertilized	Russula_decolorans	0.00000000
## 523	MAT.B3.WF	Warmed.Fertilized	Russula_nitida/sphagnicola	0.17857143
## 524	MAT.B3.WF	Warmed.Fertilized	Russula_sp._1	0.03571429
## 525	MAT.B3.WF	Warmed.Fertilized	Russula_sp._2	0.07142857
## 526	MAT.B3.WF	Warmed.Fertilized	Russula_sp._3	0.25000000
## 527	MAT.B3.WF	Warmed.Fertilized	Russula_sp._4	0.00000000
## 528	MAT.B3.WF	Warmed.Fertilized	Russula_vinosa	0.00000000
## 529	MAT.B4.C	Control	Alternaria_alternata	0.00000000
## 530	MAT.B4.C	Control	Articulospora_tetracladia	0.00000000
## 531	MAT.B4.C	Control	Cadophora_finlandica	0.00000000
## 532	MAT.B4.C	Control	Cenococcum_geophilum	0.10000000
## 533	MAT.B4.C	Control	Chalara_sp._1	0.00000000
## 534	MAT.B4.C	Control	Cortinarius_sp._1	0.16666667
## 535	MAT.B4.C	Control	Cortinarius_croceus_1	0.00000000
## 536	MAT.B4.C	Control	Cortinarius_glandicolor	0.00000000
## 537	MAT.B4.C	Control	Cortinarius_casimiri	0.00000000
## 538	MAT.B4.C	Control	Cortinarius_croceus_2	0.00000000
## 539	MAT.B4.C	Control	Cortinarius_herpeticus	0.10000000
## 540	MAT.B4.C	Control	Cortinarius_tabularis	0.03333333
## 541	MAT.B4.C	Control	Cortinarius_delibutus	0.00000000
## 542	MAT.B4.C	Control	Cortinarius_sp._2	0.00000000
## 543	MAT.B4.C	Control	Cortinarius_paragaudis	0.00000000
## 544	MAT.B4.C	Control	Cortinarius_sp._3	0.00000000
## 545	MAT.B4.C	Control	Helotiales_sp._1	0.00000000
## 546	MAT.B4.C	Control	Laccaria_laccata	0.03333333

## 547	MAT.B4.C	Control	Lactarius_vietus_1	0.00000000
## 548	MAT.B4.C	Control	Lactarius_glyciosmus	0.03333333
## 549	MAT.B4.C	Control	Lactarius_vietus_2	0.00000000
## 550	MAT.B4.C	Control	Leccinum_holopus	0.00000000
## 551	MAT.B4.C	Control	Leccinum_scabrum	0.00000000
## 552	MAT.B4.C	Control	Leccinum_variicolor_1	0.00000000
## 553	MAT.B4.C	Control	Leccinum_variicolor_2	0.03333333
## 554	MAT.B4.C	Control	Leccinum_variicolor_3	0.00000000
## 555	MAT.B4.C	Control	Leptodontidium_elatius	0.00000000
## 556	MAT.B4.C	Control	Meliniomyces_variabilis	0.00000000
## 557	MAT.B4.C	Control	Meliniomyces_bicolor	0.00000000
## 558	MAT.B4.C	Control	Phialocephala_fortinii	0.00000000
## 559	MAT.B4.C	Control	Pseudotomentella_sp._1	0.00000000
## 560	MAT.B4.C	Control	Rhizoscyphus_ericae	0.00000000
## 561	MAT.B4.C	Control	Thelephora_terrestris	0.00000000
## 562	MAT.B4.C	Control	Tomentella_sublilacina	0.00000000
## 563	MAT.B4.C	Control	Tomentella_sp._1	0.00000000
## 564	MAT.B4.C	Control	Tomentellopsis_submollis	0.03333333
## 565	MAT.B4.C	Control	Russula_chamiteae	0.13333333
## 566	MAT.B4.C	Control	Russula_decolorans	0.23333333
## 567	MAT.B4.C	Control	Russula_nitida/sphagnicola	0.03333333
## 568	MAT.B4.C	Control	Russula_sp._1	0.06666667
## 569	MAT.B4.C	Control	Russula_sp._2	0.00000000
## 570	MAT.B4.C	Control	Russula_sp._3	0.00000000
## 571	MAT.B4.C	Control	Russula_sp._4	0.00000000
## 572	MAT.B4.C	Control	Russula_vinosa	0.00000000
## 573	MAT.B4.F	Fertilized	Alternaria_alternata	0.00000000
## 574	MAT.B4.F	Fertilized	Articulospora_tetracladia	0.00000000
## 575	MAT.B4.F	Fertilized	Cadophora_finlandica	0.00000000
## 576	MAT.B4.F	Fertilized	Cenococcum_geophilum	0.00000000
## 577	MAT.B4.F	Fertilized	Chalara_sp._1	0.00000000
## 578	MAT.B4.F	Fertilized	Cortinarius_sp._1	0.00000000
## 579	MAT.B4.F	Fertilized	Cortinarius_croceus_1	0.00000000
## 580	MAT.B4.F	Fertilized	Cortinarius_glandicolor	0.00000000
## 581	MAT.B4.F	Fertilized	Cortinarius_casimiri	0.00000000
## 582	MAT.B4.F	Fertilized	Cortinarius_croceus_2	0.00000000
## 583	MAT.B4.F	Fertilized	Cortinarius_herpeticus	0.00000000
## 584	MAT.B4.F	Fertilized	Cortinarius_tabularis	0.00000000
## 585	MAT.B4.F	Fertilized	Cortinarius_delibutus	0.10000000
## 586	MAT.B4.F	Fertilized	Cortinarius_sp._2	0.00000000
## 587	MAT.B4.F	Fertilized	Cortinarius_paragaudis	0.00000000
## 588	MAT.B4.F	Fertilized	Cortinarius_sp._3	0.00000000
## 589	MAT.B4.F	Fertilized	Helotiales_sp._1	0.00000000
## 590	MAT.B4.F	Fertilized	Laccaria_laccata	0.26666667
## 591	MAT.B4.F	Fertilized	Lactarius_vietus_1	0.00000000
## 592	MAT.B4.F	Fertilized	Lactarius_glyciosmus	0.06666667
## 593	MAT.B4.F	Fertilized	Lactarius_vietus_2	0.03333333
## 594	MAT.B4.F	Fertilized	Leccinum_holopus	0.00000000
## 595	MAT.B4.F	Fertilized	Leccinum_scabrum	0.00000000
## 596	MAT.B4.F	Fertilized	Leccinum_variicolor_1	0.00000000
## 597	MAT.B4.F	Fertilized	Leccinum_variicolor_2	0.00000000
## 598	MAT.B4.F	Fertilized	Leccinum_variicolor_3	0.00000000
## 599	MAT.B4.F	Fertilized	Leptodontidium_elatius	0.00000000
## 600	MAT.B4.F	Fertilized	Meliniomyces_variabilis	0.00000000

## 601	MAT.B4.F	Fertilized	Meliniomyces_bicolor	0.00000000
## 602	MAT.B4.F	Fertilized	Phialocephala_fortinii	0.00000000
## 603	MAT.B4.F	Fertilized	Pseudotomentella_sp._1	0.00000000
## 604	MAT.B4.F	Fertilized	Rhizoscyphus_ericae	0.03333333
## 605	MAT.B4.F	Fertilized	Thelephora_terrestris	0.10000000
## 606	MAT.B4.F	Fertilized	Tomentella_sublilacina	0.00000000
## 607	MAT.B4.F	Fertilized	Tomentella_sp._1	0.00000000
## 608	MAT.B4.F	Fertilized	Tomentellopsis_submollis	0.00000000
## 609	MAT.B4.F	Fertilized	Russula_chamiteae	0.00000000
## 610	MAT.B4.F	Fertilized	Russula_decolorans	0.00000000
## 611	MAT.B4.F	Fertilized	Russula_nitida/sphagnicola	0.40000000
## 612	MAT.B4.F	Fertilized	Russula_sp._1	0.00000000
## 613	MAT.B4.F	Fertilized	Russula_sp._2	0.00000000
## 614	MAT.B4.F	Fertilized	Russula_sp._3	0.00000000
## 615	MAT.B4.F	Fertilized	Russula_sp._4	0.00000000
## 616	MAT.B4.F	Fertilized	Russula_vinosa	0.00000000
## 617	MAT.B4.W	Warmed	Alternaria_alternata	0.00000000
## 618	MAT.B4.W	Warmed	Articulospora_tetracladia	0.00000000
## 619	MAT.B4.W	Warmed	Cadophora_finlandica	0.00000000
## 620	MAT.B4.W	Warmed	Cenococcum_geophilum	0.03389830
## 621	MAT.B4.W	Warmed	Chalara_sp._1	0.00000000
## 622	MAT.B4.W	Warmed	Cortinarius_sp._1	0.00000000
## 623	MAT.B4.W	Warmed	Cortinarius_croceus_1	0.00000000
## 624	MAT.B4.W	Warmed	Cortinarius_glandicolor	0.00000000
## 625	MAT.B4.W	Warmed	Cortinarius_casimiri	0.00000000
## 626	MAT.B4.W	Warmed	Cortinarius_croceus_2	0.01694915
## 627	MAT.B4.W	Warmed	Cortinarius_herpeticus	0.00000000
## 628	MAT.B4.W	Warmed	Cortinarius_tabularis	0.00000000
## 629	MAT.B4.W	Warmed	Cortinarius_delibutus	0.00000000
## 630	MAT.B4.W	Warmed	Cortinarius_sp._2	0.08474576
## 631	MAT.B4.W	Warmed	Cortinarius_paragaudis	0.01694915
## 632	MAT.B4.W	Warmed	Cortinarius_sp._3	0.00000000
## 633	MAT.B4.W	Warmed	Helotiales_sp._1	0.00000000
## 634	MAT.B4.W	Warmed	Laccaria_laccata	0.00000000
## 635	MAT.B4.W	Warmed	Lactarius_vietus_1	0.10169491
## 636	MAT.B4.W	Warmed	Lactarius_glyciosmus	0.01694915
## 637	MAT.B4.W	Warmed	Lactarius_vietus_2	0.00000000
## 638	MAT.B4.W	Warmed	Leccinum_holopus	0.00000000
## 639	MAT.B4.W	Warmed	Leccinum_scabrum	0.00000000
## 640	MAT.B4.W	Warmed	Leccinum_variicolor_1	0.00000000
## 641	MAT.B4.W	Warmed	Leccinum_variicolor_2	0.00000000
## 642	MAT.B4.W	Warmed	Leccinum_variicolor_3	0.00000000
## 643	MAT.B4.W	Warmed	Leptodontidium_elatius	0.00000000
## 644	MAT.B4.W	Warmed	Meliniomyces_variabilis	0.00000000
## 645	MAT.B4.W	Warmed	Meliniomyces_bicolor	0.00000000
## 646	MAT.B4.W	Warmed	Phialocephala_fortinii	0.00000000
## 647	MAT.B4.W	Warmed	Pseudotomentella_sp._1	0.00000000
## 648	MAT.B4.W	Warmed	Rhizoscyphus_ericae	0.00000000
## 649	MAT.B4.W	Warmed	Thelephora_terrestris	0.00000000
## 650	MAT.B4.W	Warmed	Tomentella_sublilacina	0.08474576
## 651	MAT.B4.W	Warmed	Tomentella_sp._1	0.05084746
## 652	MAT.B4.W	Warmed	Tomentellopsis_submollis	0.00000000
## 653	MAT.B4.W	Warmed	Russula_chamiteae	0.08474576
## 654	MAT.B4.W	Warmed	Russula_decolorans	0.49152542

## 655	MAT.B4.W	Warmed	Russula_nitida/sphagnicola	0.00000000
## 656	MAT.B4.W	Warmed	Russula_sp._1	0.00000000
## 657	MAT.B4.W	Warmed	Russula_sp._2	0.00000000
## 658	MAT.B4.W	Warmed	Russula_sp._3	0.00000000
## 659	MAT.B4.W	Warmed	Russula_sp._4	0.01694915
## 660	MAT.B4.W	Warmed	Russula_vinosa	0.00000000
## 661	MAT.B4.WF	Warmed.Fertilized	Alternaria_alternata	0.00000000
## 662	MAT.B4.WF	Warmed.Fertilized	Articulospora_tetracladia	0.00000000
## 663	MAT.B4.WF	Warmed.Fertilized	Cadophora_finlandica	0.00000000
## 664	MAT.B4.WF	Warmed.Fertilized	Cenococcum_geophilum	0.25000000
## 665	MAT.B4.WF	Warmed.Fertilized	Chalara_sp._1	0.00000000
## 666	MAT.B4.WF	Warmed.Fertilized	Cortinarius_sp._1	0.00000000
## 667	MAT.B4.WF	Warmed.Fertilized	Cortinarius_croceus_1	0.00000000
## 668	MAT.B4.WF	Warmed.Fertilized	Cortinarius_glandicolor	0.00000000
## 669	MAT.B4.WF	Warmed.Fertilized	Cortinarius_casimiri	0.06250000
## 670	MAT.B4.WF	Warmed.Fertilized	Cortinarius_croceus_2	0.00000000
## 671	MAT.B4.WF	Warmed.Fertilized	Cortinarius_herpeticus	0.00000000
## 672	MAT.B4.WF	Warmed.Fertilized	Cortinarius_tabularis	0.00000000
## 673	MAT.B4.WF	Warmed.Fertilized	Cortinarius_delibutus	0.00000000
## 674	MAT.B4.WF	Warmed.Fertilized	Cortinarius_sp._2	0.00000000
## 675	MAT.B4.WF	Warmed.Fertilized	Cortinarius_paragaudis	0.00000000
## 676	MAT.B4.WF	Warmed.Fertilized	Cortinarius_sp._3	0.09375000
## 677	MAT.B4.WF	Warmed.Fertilized	Helotiales_sp._1	0.00000000
## 678	MAT.B4.WF	Warmed.Fertilized	Laccaria_laccata	0.00000000
## 679	MAT.B4.WF	Warmed.Fertilized	Lactarius_vietus_1	0.25000000
## 680	MAT.B4.WF	Warmed.Fertilized	Lactarius_glyciosmus	0.06250000
## 681	MAT.B4.WF	Warmed.Fertilized	Lactarius_vietus_2	0.00000000
## 682	MAT.B4.WF	Warmed.Fertilized	Leccinum_holopus	0.00000000
## 683	MAT.B4.WF	Warmed.Fertilized	Leccinum_scabrum	0.00000000
## 684	MAT.B4.WF	Warmed.Fertilized	Leccinum_variicolor_1	0.00000000
## 685	MAT.B4.WF	Warmed.Fertilized	Leccinum_variicolor_2	0.00000000
## 686	MAT.B4.WF	Warmed.Fertilized	Leccinum_variicolor_3	0.00000000
## 687	MAT.B4.WF	Warmed.Fertilized	Leptodontidium_elatius	0.00000000
## 688	MAT.B4.WF	Warmed.Fertilized	Meliniomyces_variabilis	0.00000000
## 689	MAT.B4.WF	Warmed.Fertilized	Meliniomyces_bicolor	0.06250000
## 690	MAT.B4.WF	Warmed.Fertilized	Phialocephala_fortinii	0.00000000
## 691	MAT.B4.WF	Warmed.Fertilized	Pseudotomentella_sp._1	0.00000000
## 692	MAT.B4.WF	Warmed.Fertilized	Rhizoscyphus_ericae	0.00000000
## 693	MAT.B4.WF	Warmed.Fertilized	Thelephora_terrestris	0.00000000
## 694	MAT.B4.WF	Warmed.Fertilized	Tomentella_sublilacina	0.00000000
## 695	MAT.B4.WF	Warmed.Fertilized	Tomentella_sp._1	0.00000000
## 696	MAT.B4.WF	Warmed.Fertilized	Tomentellopsis_submollis	0.03125000
## 697	MAT.B4.WF	Warmed.Fertilized	Russula_chamiteae	0.03125000
## 698	MAT.B4.WF	Warmed.Fertilized	Russula_decolorans	0.00000000
## 699	MAT.B4.WF	Warmed.Fertilized	Russula_nitida/sphagnicola	0.12500000
## 700	MAT.B4.WF	Warmed.Fertilized	Russula_sp._1	0.00000000
## 701	MAT.B4.WF	Warmed.Fertilized	Russula_sp._2	0.00000000
## 702	MAT.B4.WF	Warmed.Fertilized	Russula_sp._3	0.03125000
## 703	MAT.B4.WF	Warmed.Fertilized	Russula_sp._4	0.00000000
## 704	MAT.B4.WF	Warmed.Fertilized	Russula_vinosa	0.00000000

```
#Changing data so that each species is a column, values are the relative.abundance, and rows are each p
library(tidyr)
ExpandProjdata <- slim %>%
```



```

  pivot_wider(names_from = Species, values_from = Relative.Abandance, values_fill = 0)
ExpandProjdata <- ExpandProjdata %>%
  arrange(Treatment)
ExpandProjdata.db <- vegdist(Sorted_ExpandProjdata[, -c(1,2)], method = "bray", upper = TRUE, diag = TRUE)
#Define Order of Sites
order.projdata <- rev(attr(ExpandProjdata.db, "Labels"))

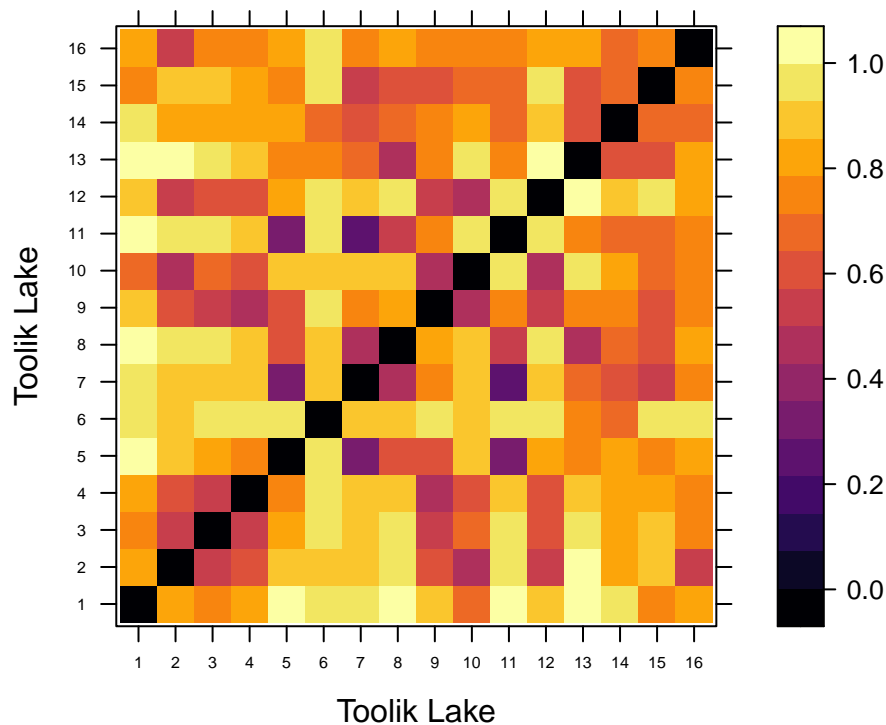
heatmap_matrix <- as.matrix(ExpandProjdata.db)

# Ensure rows and columns match the order
if (!is.null(order.projdata)) {
  heatmap_matrix <- heatmap_matrix[order.projdata, order.projdata]
}

# Now plot
levelplot(heatmap_matrix, aspect = "iso", col.regions = inferno,
  xlab = "Toolik Lake", ylab = "Toolik Lake", scales = list(cex = 0.5),
  main = "Bray-Curtis Distance")

```

Bray-Curtis Distance



#After having reorganized my data so that sites are sorted by treatment group, it is difficult to detect