Assignment: Local (alpha) Diversity

Evgeniya Polezhaeva Z620: Quantitative Biodiversity, Indiana University

OVERVIEW

In this exercise, we will explore aspects of local or site-specific diversity, also known as alpha (α) diversity. First we will quantify two of the fundamental components of (α) diversity: **richness** and **evenness**. From there, we will then discuss ways to integrate richness and evenness, which will include univariate metrics of diversity along with an investigation of the **species abundance distribution (SAD)**.

Directions:

- 1. Change "Student Name" on line 3 (above) with your name.
- 2. Complete as much of the exercise as possible during class; what you do not complete in class will need to be done on your own outside of class.
- 3. Use the handout as a guide; it contains a more complete description of data sets along with the proper scripting needed to carry out the exercise.
- 4. Be sure to **answer the questions** in this exercise document; they also correspond to the handout. Space for your answer is provided in this document and indicated by the ">" character. If you need a second paragraph be sure to start the first line with ">".
- 5. Before you leave the classroom, **push** this file to your GitHub repo.
- 6. For homework, 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, please submit the completed exercise by creating a **pull request** via GitHub. Your pull request should include this file alpha_assignment.Rmd and the PDF output of Knitr (alpha_assignment.pdf).

1) R SETUP

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 /Week2-Alpha folder, and 4) Load the vegan R package (be sure to install if needed).

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

## [1] "D:/Jane/GitHub/QB2017_Polezhaeva/Week2-Alpha"
setwd("D:/Jane/GitHub/QB2017_Polezhaeva/Week2-Alpha")
getwd()

## [1] "D:/Jane/GitHub/QB2017_Polezhaeva/Week2-Alpha"
install.packages("vegan", repos="http://cran.rstudio.com/")

## Installing package into 'C:/Users/Jane/Documents/R/win-library/3.3'
## (as 'lib' is unspecified)

## package 'vegan' successfully unpacked and MD5 sums checked
##
## The downloaded binary packages are in
## C:\Users\Jane\AppData\Local\Temp\RtmpyopVQl\downloaded_packages
```

```
require("vegan")

## Loading required package: vegan

## Loading required package: permute

## Loading required package: lattice

## This is vegan 2.4-2
```

2) LOADING DATA

In the R code chunk below, do the following: 1) Load your dataset, and 2) Display the structure of the dataset (if the structure is long, use max.level=0 to show just basic information).

```
dataset (if the structure is long, use max.level=0 to show just basic information).
require("vegan")
data(BCI)
typeof(BCI)
## [1] "list"
str(BCI)
                     50 obs. of 225 variables:
  'data.frame':
##
    $ Abarema.macradenia
                                        : int
                                                0 0 0 0 0 0 0 0 0 1 ...
    $ Vachellia.melanoceras
                                        : int
                                                0 0 0 0 0 0 0 0 0 0 ...
##
```

```
$ Acalypha.diversifolia
                                    : int
                                           0 0 0 0 0 0 0 0 0 0 ...
##
  $ Acalypha.macrostachya
                                    : int
                                           0 0 0 0 0 0 0 0 0 0 ...
##
   $ Adelia.triloba
                                    : int
                                           0 0 0 3 1 0 0 0 5 0 ...
                                           0 0 0 0 1 0 1 0 0 1 ...
##
   $ Aegiphila.panamensis
                                    : int
##
  $ Alchornea.costaricensis
                                           2 1 2 18 3 2 0 2 2 2 ...
## $ Alchornea.latifolia
                                           0 0 0 0 0 1 0 0 0 0 ...
                                    : int
##
   $ Alibertia.edulis
                                    : int
                                           0 0 0 0 0 0 0 0 0 0 ...
##
   $ Allophylus.psilospermus
                                    : int
                                           0 0 0 0 1 0 0 0 0 0 ...
## $ Alseis.blackiana
                                           25 26 18 23 16 14 18 14 16 14 ...
                                    : int
## $ Amaioua.corymbosa
                                           0 0 0 0 0 0 0 0 0 0 ...
                                    : int
   $ Anacardium.excelsum
                                           0 0 0 0 0 0 0 1 0 0 ...
                                    : int
## $ Andira.inermis
                                    : int
                                          0 0 0 0 1 1 0 0 1 0 ...
##
  $ Annona.spraguei
                                    : int
                                           1010000110...
                                           13 12 6 3 4 10 5 4 5 5 ...
##
   $ Apeiba.glabra
                                    : int
                                           2 0 1 1 0 0 0 1 0 0 ...
##
   $ Apeiba.tibourbou
                                    : int
##
   $ Aspidosperma.desmanthum
                                    : int
                                           0 0 0 1 1 1 0 0 0 1 ...
   $ Astrocaryum.standleyanum
                                    : int
                                           0 2 1 5 6 2 2 0 2 1 ...
                                           6 0 1 3 0 1 2 2 0 0 ...
##
   $ Astronium.graveolens
                                    : int
##
   $ Attalea.butyracea
                                    : int
                                           0 1 0 0 0 1 1 0 0 0 ...
##
  $ Banara.guianensis
                                    : int
                                           0 0 0 0 0 0 0 0 0 0 ...
                                           4 5 7 5 8 6 5 9 11 14 ...
##
  $ Beilschmiedia.pendula
                                    : int
##
   $ Brosimum.alicastrum
                                    : int
                                           5 2 4 3 2 2 6 4 3 6 ...
##
   $ Brosimum.guianense
                                    : int
                                           0000000000...
##
  $ Calophyllum.longifolium
                                    : int
                                           0 2 0 2 1 2 2 2 2 0 ...
## $ Casearia.aculeata
                                    : int
                                           0 0 0 0 0 0 0 1 0 0 ...
   $ Casearia.arborea
                                           1 1 3 2 4 1 2 3 9 7 ...
##
                                    : int
   $ Casearia.commersoniana
                                    : int 0010100010...
##
  $ Casearia.guianensis
                                    : int 0000000000...
                                           2 1 0 0 0 3 1 0 1 1 ...
##
   $ Casearia.sylvestris
                                    : int
                                    : int 2 0 1 1 3 4 4 0 2 1 ...
   $ Cassipourea.guianensis
```

```
$ Cavanillesia.platanifolia
                                      : int 0000000000...
                                            12 5 7 17 21 4 0 7 2 16 ...
##
   $ Cecropia.insignis
                                      : int
##
   $ Cecropia.obtusifolia
                                      : int
                                             0 0 0 0 1 0 0 2 0 2 ...
##
  $ Cedrela.odorata
                                      : int
                                            0000000000...
##
   $ Ceiba.pentandra
                                      : int
                                             0 1 1 0 1 0 0 1 0 1 ...
##
                                            0 0 0 2 2 0 1 0 0 0 ...
   $ Celtis.schippii
                                      : int
   $ Cespedesia.spathulata
                                             0 0 0 0 0 0 0 0 0 0 ...
                                      : int
                                      : int
##
   $ Chamguava.schippii
                                             0 0 0 0 0 0 0 0 0 0 ...
##
   $ Chimarrhis.parviflora
                                      : int
                                             0 0 0 0 0 0 0 0 0 0 ...
##
   $ Maclura.tinctoria
                                      : int
                                             0 0 0 0 0 0 0 0 0 0 ...
   $ Chrysochlamys.eclipes
                                      : int
                                             0 0 0 0 0 0 0 0 0 0 ...
##
   $ Chrysophyllum.argenteum
                                             4 1 2 2 6 2 3 2 4 2 ...
                                      : int
##
   $ Chrysophyllum.cainito
                                            0 0 0 0 0 0 1 0 0 0 ...
                                      : int
##
                                            0 0 0 1 2 0 0 1 2 1 ...
   $ Coccoloba.coronata
                                      : int
##
   $ Coccoloba.manzinellensis
                                             0 0 0 0 0 0 0 2 0 0 ...
                                      : int
##
   $ Colubrina.glandulosa
                                      : int
                                             0 0 0 0 0 0 0 0 0 0 ...
##
                                            2 3 3 7 1 1 2 0 0 2 ...
   $ Cordia.alliodora
                                      : int
##
   $ Cordia.bicolor
                                            12 14 35 23 13 7 5 10 7 13 ...
                                      : int
   $ Cordia.lasiocalyx
##
                                            8 6 6 11 7 6 6 3 0 4 ...
                                      : int
##
   $ Coussarea.curvigemma
                                      : int
                                            0 0 0 1 0 2 1 0 1 1 ...
##
   $ Croton.billbergianus
                                      : int
                                            2 2 0 11 6 0 0 4 2 0 ...
   $ Cupania.cinerea
                                            0 0 0 0 0 0 0 0 0 0 ...
                                      : int
##
   $ Cupania.latifolia
                                             0 0 0 1 0 0 0 0 0 0 ...
                                      : int
   $ Cupania.rufescens
                                             0 0 0 0 0 0 0 0 0 0 ...
##
                                      : int
##
                                            2 2 1 0 3 0 1 2 2 0 ...
   $ Cupania.seemannii
                                      : int
   $ Dendropanax.arboreus
                                      : int
                                            0 3 6 0 5 2 1 6 1 3 ...
##
   $ Desmopsis.panamensis
                                             0 0 4 0 0 0 0 0 0 1 ...
                                      : int
   $ Diospyros.artanthifolia
                                      : int
                                            1 1 1 1 0 0 0 0 0 1 ...
##
                                            1 1 3 0 0 0 0 2 1 2 ...
   $ Dipteryx.oleifera
                                      : int
##
   $ Drypetes.standleyi
                                      : int
                                             2 1 2 0 0 0 0 0 0 0 ...
##
   $ Elaeis.oleifera
                                      : int
                                             0 0 0 0 0 0 0 0 0 0 ...
##
   $ Enterolobium.schomburgkii
                                      · int
                                             0 0 0 0 0 0 0 0 0 0 ...
##
   $ Erythrina.costaricensis
                                      : int
                                            0 0 0 0 0 3 0 0 1 0 ...
##
                                            0 1 0 0 0 0 0 1 1 1 ...
   $ Erythroxylum.macrophyllum
                                      : int
##
   $ Eugenia.florida
                                             0 1 0 7 2 0 0 1 1 3 ...
                                      : int
##
                                            0 0 0 0 0 0 0 1 0 0 ...
   $ Eugenia.galalonensis
                                      : int
##
   $ Eugenia.nesiotica
                                      : int
                                            0 0 1 0 0 0 5 4 3 0 ...
##
   $ Eugenia.oerstediana
                                      : int
                                             3 2 5 1 5 2 2 3 3 3 ...
##
   $ Faramea.occidentalis
                                             14 36 39 39 22 16 38 41 33 42 ...
                                      : int
                                            0 1 0 0 0 0 0 0 0 0 ...
##
   $ Ficus.colubrinae
                                      : int
                                            0 0 0 0 0 0 0 0 0 0 ...
   $ Ficus.costaricana
                                      : int
##
   $ Ficus.insipida
                                      : int
                                            0000000000...
                                            1000000000...
##
   $ Ficus.maxima
                                      : int
##
   $ Ficus.obtusifolia
                                      : int
                                            0 0 0 0 0 0 0 0 0 0 ...
   $ Ficus.popenoei
                                      : int
                                             0 0 0 0 0 0 1 0 0 0 ...
                                             0 0 1 2 1 0 0 0 0 0 ...
##
   $ Ficus.tonduzii
                                      : int
##
   $ Ficus.trigonata
                                      : int
                                            0 0 0 0 0 0 0 0 0 0 ...
                                            1 0 0 0 0 1 1 0 0 0 ...
##
   $ Ficus.yoponensis
                                      : int
##
   $ Garcinia.intermedia
                                      : int
                                            0 1 1 3 2 1 2 2 1 0 ...
##
   $ Garcinia.madruno
                                      : int
                                             4 0 0 0 1 0 0 0 0 1 ...
##
                                            0 0 1 0 0 0 1 0 1 1 ...
   $ Genipa.americana
                                      : int
## $ Guapira.myrtiflora
                                      : int 3 1 0 1 1 7 3 1 1 1 ...
## $ Guarea.fuzzy
                                     : int 1101300203...
## $ Guarea.grandifolia
                                      : int 000000100...
```

```
$ Guatteria.dumetorum
                                             6 16 6 3 9 7 8 6 2 2 ...
##
                                      : int
##
   $ Guazuma.ulmifolia
                                             0 0 0 1 0 0 0 0 0 0 ...
  $ Guettarda.foliacea
                                             1 5 1 2 1 0 0 4 1 3 ...
##
                                       int
##
   $ Gustavia.superba
                                       int
                                             10 5 0 1 3 1 8 4 4 4 ...
##
                                             0 0 1 0 0 0 0 0 2 1 ...
   $ Hampea.appendiculata
                                      : int
   $ Hasseltia.floribunda
                                             5 9 4 11 9 2 7 6 3 4 ...
                                      : int
##
   $ Heisteria.acuminata
                                      : int
                                             0 0 0 0 1 1 0 0 0 0 ...
##
   $ Heisteria.concinna
                                        int
                                             4546482515...
                                             0 0 0 0 0 0 0 0 0 0 ...
##
   $ Hirtella.americana
                                      : int
## $ Hirtella.triandra
                                             21 14 5 4 6 6 7 14 8 7 ...
                                      : int
                                             0 0 0 0 0 2 1 1 0 0 ...
## $ Hura.crepitans
                                      : int
                                      : int
## $ Hieronyma.alchorneoides
                                             0 2 0 0 0 0 0 0 1 0 ...
     [list output truncated]
  - attr(*, "original.names")= chr "Abarema.macradenium" "Acacia.melanoceras" "Acalypha.diversifolia
```

2 6 2 5 3 4 4 0 1 5 ...

3) SPECIES RICHNESS

\$ Guarea.guidonia

Species richness (S) is simply the number of species in a system or the number of species observed in a sample.

: int

Observed Richness

In the R code chunk below, do the following:

S.obs <- function(x = "") $\{rowSums(x>0) * 1\}$

- 1. Write a function called S.obs to calculate observed richness
- 2. Use your function to determine the number of species in site1, and
- 3. Compare the output of your function to the output of the specnumber() function in vegan.

```
site1 <- BCI[1,]
site1
##
     Abarema.macradenia Vachellia.melanoceras Acalypha.diversifolia
## 1
##
     Acalypha.macrostachya Adelia.triloba Aegiphila.panamensis
## 1
##
     Alchornea.costaricensis Alchornea.latifolia Alibertia.edulis
## 1
##
     Allophylus.psilospermus Alseis.blackiana Amaioua.corymbosa
## 1
##
     Anacardium.excelsum Andira.inermis Annona.spraguei Apeiba.glabra
## 1
##
     Apeiba.tibourbou Aspidosperma.desmanthum Astrocaryum.standleyanum
## 1
##
     Astronium.graveolens Attalea.butyracea Banara.guianensis
## 1
     Beilschmiedia.pendula Brosimum.alicastrum Brosimum.guianense
##
## 1
                                              5
     Calophyllum.longifolium Casearia.aculeata Casearia.arborea
##
## 1
##
     Casearia.commersoniana Casearia.guianensis Casearia.sylvestris
## 1
```

```
Cassipourea.guianensis Cavanillesia.platanifolia Cecropia.insignis
## 1
     Cecropia.obtusifolia Cedrela.odorata Ceiba.pentandra Celtis.schippii
##
## 1
     Cespedesia.spathulata Chamguava.schippii Chimarrhis.parviflora
##
## 1
     Maclura.tinctoria Chrysochlamys.eclipes Chrysophyllum.argenteum
##
## 1
##
     Chrysophyllum.cainito Coccoloba.coronata Coccoloba.manzinellensis
## 1
##
     Colubrina.glandulosa Cordia.alliodora Cordia.bicolor Cordia.lasiocalyx
##
     Coussarea.curvigemma Croton.billbergianus Cupania.cinerea
##
## 1
##
     Cupania.latifolia Cupania.rufescens Cupania.seemannii
## 1
     Dendropanax.arboreus Desmopsis.panamensis Diospyros.artanthifolia
##
     Dipteryx.oleifera Drypetes.standleyi Elaeis.oleifera
##
## 1
##
     Enterolobium.schomburgkii Erythrina.costaricensis
## 1
     Erythroxylum.macrophyllum Eugenia.florida Eugenia.galalonensis
##
## 1
     Eugenia.nesiotica Eugenia.oerstediana Faramea.occidentalis
##
##
     Ficus.colubrinae Ficus.costaricana Ficus.insipida Ficus.maxima
## 1
     Ficus.obtusifolia Ficus.popenoei Ficus.tonduzii Ficus.trigonata
##
## 1
##
     Ficus.yoponensis Garcinia.intermedia Garcinia.madruno Genipa.americana
## 1
     Guapira.myrtiflora Guarea.fuzzy Guarea.grandifolia Guarea.guidonia
## 1
     Guatteria.dumetorum Guazuma.ulmifolia Guettarda.foliacea
##
## 1
##
     Gustavia.superba Hampea.appendiculata Hasseltia.floribunda
## 1
     Heisteria.acuminata Heisteria.concinna Hirtella.americana
##
## 1
     Hirtella.triandra Hura.crepitans Hieronyma.alchorneoides Inga.acuminata
##
     Inga.cocleensis Inga.goldmanii Inga.laurina Inga.semialata Inga.nobilis
##
## 1
     Inga.oerstediana Inga.pezizifera Inga.punctata Inga.ruiziana
##
## 1
     Inga.sapindoides Inga.spectabilis Inga.umbellifera Jacaranda.copaia
##
## 1
     Lacistema.aggregatum Lacmellea.panamensis Laetia.procera Laetia.thamnia
## 1
     Lafoensia.punicifolia Licania.hypoleuca Licania.platypus
##
     Lindackeria.laurina Lonchocarpus.heptaphyllus Luehea.seemannii
## 1
```

```
Macrocnemum.roseum Maquira.guianensis.costaricana Margaritaria.nobilis
## 1
     Marila.laxiflora Maytenus.schippii Miconia.affinis Miconia.argentea
##
##
##
     Miconia.elata Miconia.hondurensis Mosannona.garwoodii Myrcia.gatunensis
##
     Myrospermum.frutescens Nectandra.cissiflora Nectandra.lineata
## 1
##
     Nectandra.purpurea Ochroma.pyramidale Ocotea.cernua Ocotea.oblonga
## 1
     Ocotea.puberula Ocotea.whitei Oenocarpus.mapora Ormosia.amazonica
##
     Ormosia.coccinea Ormosia.macrocalyx Pachira.quinata Pachira.sessilis
##
## 1
     Perebea.xanthochyma Cinnamomum.triplinerve Picramnia.latifolia
##
## 1
     Piper.reticulatum Platymiscium.pinnatum Platypodium.elegans
##
     Posoqueria.latifolia Poulsenia.armata Pourouma.bicolor
##
## 1
##
     Pouteria.fossicola Pouteria.reticulata Pouteria.stipitata
## 1
     Prioria.copaifera Protium.costaricense Protium.panamense
##
## 1
     Protium.tenuifolium Pseudobombax.septenatum Psidium.friedrichsthalianum
##
##
     Psychotria.grandis Pterocarpus.rohrii Quararibea.asterolepis
## 1
     Quassia.amara Randia.armata Sapium.broadleaf Sapium.glandulosum
##
## 1
##
     Schizolobium.parahyba Senna.dariensis Simarouba.amara
## 1
     Siparuna.guianensis Siparuna.pauciflora Sloanea.terniflora
## 1
##
     Socratea.exorrhiza Solanum.hayesii Sorocea.affinis Spachea.membranacea
## 1
     Spondias.mombin Spondias.radlkoferi Sterculia.apetala
##
## 1
##
     Swartzia.simplex.var.grandiflora Swartzia.simplex.continentalis
## 1
     Symphonia.globulifera Handroanthus.guayacan Tabebuia.rosea
## 1
     Tabernaemontana.arborea Tachigali.versicolor Talisia.nervosa
##
## 1
     Talisia.princeps Terminalia.amazonia Terminalia.oblonga
##
## 1
     Tetragastris.panamensis Tetrathylacium.johansenii Theobroma.cacao
##
     Thevetia.ahouai Tocoyena.pittieri Trattinnickia.aspera Trema.micrantha
     Trichanthera.gigantea Trichilia.pallida Trichilia.tuberculata
##
     Trichospermum.galeottii Triplaris.cumingiana Trophis.caucana
## 1
```

```
##
     Trophis.racemosa Turpinia.occidentalis Unonopsis.pittieri
## 1
##
     Virola.multiflora Virola.sebifera Virola.surinamensis Vismia.baccifera
## 1
##
     Vochysia.ferruginea Xylopia.macrantha Zanthoxylum.ekmanii
## 1
     Zanthoxylum.juniperinum Zanthoxylum.panamense Zanthoxylum.setulosum
##
## 1
                                                  2
##
     Zuelania.guidonia
## 1
S.obs(site1)
##
   1
## 93
require("vegan")
specnumber(site1)
## 1
## 93
specnumber(BCI[1:4,])
  1 2 3 4
## 93 84 90 94
# S.obs <- function( ){
    rowSums(
```

Question 1: Does specnumber() from vegan return the same value for observed richness in site1 as our function S.obs? What is the species richness of the first 4 sites (i.e., rows) of the BCI matrix?

Answer 1: Yes, it does. The species richness of the first 4 sites: 93, 84, 90, 94

Coverage. How Well Did You Sample Your Site?

In the R code chunk below, do the following:

- 1. Write a function to calculate Good's Coverage, and
- 2. Use that function to calculate coverage for all sites in the BCI matrix.

[8] 0.9443155 0.9095355 0.9275362 0.9152120 0.9071038 0.9242054 0.9132420

```
## [15] 0.9350649 0.9267735 0.8950131 0.9193084 0.8891455 0.9114219 0.8946078 ## [22] 0.9066986 0.8705882 0.9030612 0.9095023 0.9115479 0.9088729 0.9198966 ## [29] 0.8983516 0.9221053 0.9382423 0.9411765 0.9220183 0.9239374 0.9267887 ## [36] 0.9186047 0.9379310 0.9306488 0.9268868 0.9386503 0.8880597 0.9299517 ## [43] 0.9140049 0.9168704 0.9234234 0.9348837 0.8847059 0.9228916 0.9086651 ## [50] 0.9143519
```

Question 2: Answer the following questions about coverage:

- a. What is the range of values that can be generated by Good's Coverage?
- b. What would we conclude from Good's Coverage if n_i equaled N?
- c. What portion of taxa in site1 were represented by singletons?
- d. Make some observations about coverage at the BCI plots.

Answer 2d: Good's coverage index is about 90% for all sites. It means that

Estimated Richness

In the R code chunk below, do the following:

- 1. Load the microbial dataset (located in the /Week2-Alpha/data folder),
- 2. Transform and transpose the data as needed (see handout),
- 3. Create a vector (soilbac1) with the bacterial OTU abundances at any site in the dataset,
- 4. Calculate the observed richness at that particular site, and
- 5. Calculate the coverage at that particular site

```
soilbac <- read.table("data/soilbac.txt", sep = "\t", header = TRUE, row.names = 1)
soilbac.t <- as.data.frame(t(soilbac))
soilbac1 <- soilbac.t[1,]
soilbac1</pre>
```

```
5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24
##
             2 3
## T1_1 31 11 2 25 124 5 4 0 1
                                   2
                                      0 28
                                             0
                                                0
                                                    4
                                                       0 13
                                                              1
                                                                 0
                                                                    1
                                                                        8
                                                                           0
                                                                              2
        25 26 27 28 29 30 31 32 33
##
                                      34 35 36 37 38 39 40 41 42 43
             4 11
                                       0
                                                           3
                                                               0
                                                                  2
   T1 1
         2
                       1
                          0
                             1
                                 6
                                    4
                                         14
                                              0
                                                13 22
                                                      10
                                                                      0
                                                                         8
##
        48 49
               50 51 52 53 54 55 56 57
                                          58
                                             59
                                                60
                                                   61
                                                       62
                                                          63 64
                                                                 65
                                                                    66
                                                                       67
                                                                                 70
             2
                                                 7
##
         2
                7
                  12
                       2
                          0
                            10
                                 6
                                       8
                                           6
                                              0
                                                     0
                                                        1
                                                          11
                                                               4
                                                                  3
                                                                         4
                                                                               0
  T1_1
                                    1
##
        71 72 73 74
                     75 76
                            77
                                78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93
             2
                       2
                                 0
                                       2
                                           0
                                              0
                                                     2
                                                        0
                                                            0
                                                                  0
                                                                      2
                                                                         2
## T1 1
                0
                    4
                             6
                                    1
                                                 1
                                                               0
                          1
##
        94
            95 96 97
                     98
                         99
                                 101 102 103 104 105 106
                                                           107 108 109
                                                                         110 111
                            100
## T1_1
        2
             0
                0 11
                       1
                               0
                                   0
                                        0
                                            0
                                                0
                                                     0
                                                         1
                                                             18
                                                                  0
                                                                       0
                                                                           0
                                                                                0
        113 114 115 116 117 118 119 120 121 122 123 124
                                                             125
                                                                  126
                                                                      127
                                                       0
##
   T1_1
         12
                   3
                        0
                            0
                                 0
                                     3
                                          0
                                              2
                                                   0
                                                            9
                                                                0
                                                                    0
                                                                         0
##
        130 131 132 133 134 135 136 137 138
                                                139 140 141 142
                                                                  143
                                                                      144 145
                                                                               146
                        0
                            0
                                     0
                                                       2
                                                            0
                                                                0
                                                                    0
                                                                         4
##
  T1 1
           0
               0
                    0
                                 0
                                          1
                                              1
                                                   0
##
        147 148 149 150 151 152 153 154 155 156 157 158 159 160 161 162 163
## T1 1
           0
               0
                  14
                        1
                            0
                                 0
                                     2
                                          2
                                             14
                                                   4
                                                       7
                                                            1
                                                                0
                                                                   13
                                                                             5
                                                                                  0
        164 165 166 167 168 169 170 171 172 173 174 175 176
##
                                                                  177 178
                                                                           179 180
## T1_1
               0
                    0
                            0
                                11
                                     0
                                          2
                                              6
                                                   0
                                                       0
                                                            1
                                                                0
        181 182 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197
```

```
## T1 1 0 0 0 0 1 0 0 0 0 0 0 0 0 0 0
     657 658 659 660 661 662 663 664 665 666 667 668 669 670 671 672 673
## T1 1 0 0 2 0 0 0 0 1 0 0 0 0 0
     674 675 676 677 678 679 680 681 682 683 684 685 686 687 688 689 690
## T1 1 0 4 0 0 5 0 0 0 1 0 0
                                        0 0 0 0
     691 692 693 694 695 696 697 698 699 700 701 702 703 704 705 706 707
## T1 1 1 0 0 0 2 0 0 0 0 0
                                        0 2 0 0 0
     708 709 710 711 712 713 714 715 716 717 718 719 720 721 722 723 724
## T1_1 0 0 0 0 0 0 0 0 0 0 0 4 2 0 0 0
     725 726 727 728 729 730 731 732 733 734 735 736 737 738 739 740 741
## T1_1 2 0 0
               2 0 0 0 4 2 0 1
                                      2
                                        0 0 0
     742 743 744 745 746 747 748 749 750 751 752 753 754 755 756 757 758
## T1_1 0 0 0 0 0 0 1 0 0 1 0 0 1
     759 760 761 762 763 764 765 766 767 768 769 770 771 772 773 774 775
## T1_1 0 0 1 0 0 0 0 0 0 1 0 0 3 0 0 0
##
     776 777 778 779 780 781 782 783 784 785 786 787 788 789 790 791 792
## T1_1 0 0 0 1 0 0 2 0 0 4 1 0 0 0 0 4
     793 794 795 796 797 798 799 800 801 802 803 804 805 806 807 808 809
## T1 1 0 0 3 1 1 0 0 0 0 0 0 0 0 0 0 0
     810 811 812 813 814 815 816 817 818 819 820 821 822 823 824 825 826
## T1_1 0 3 2
              1 0 2 0 0 0 0 2 0 0 0 0 0
     827 828 829 830 831 832 833 834 835 836 837 838 839 840 841 842 843
## T1_1 0 2 0 0 0 0 0 0 0 0 0 0 1
                                                  0
     844 845 846 847 848 849 850 851 852 853 854 855 856 857 858 859 860
## T1_1 0 0
            0 0
                  0 0 0 1 0 0 0 1 0 0
     861 862 863 864 865 866 867 868 869 870 871 872 873 874 875 876 877
## T1_1 1 7 0 0 0 0 0 0 0 2 1 0 0 0 0
     878 879 880 881 882 883 884 885 886 887 888 889 890 891 892 893 894
## T1_1 0 0 0 0 0 0 1 1 0 0 0 0 0 1 1
     895 896 897 898 899 900 901 902 903 904 905 906 907 908 909 910 911
## T1_1 1 0 2 0 0 0 7 0 0 0 0 0 0 0 0 0
     912 913 914 915 916 917 918 919 920 921 922 923 924 925 926 927 928
## T1_1 1 0 0 1 0 0 0 0 1 0 0 0 0 0
     929 930 931 932 933 934 935 936 937 938 939 940 941 942 943 944 945
     0 0 4 1 1 0 1 0 0 1 0 0 0 3 0 0 0
     946 947 948 949 950 951 952 953 954 955 956 957 958 959 960 961 962
     963 964 965 966 967 968 969 970 971 972 973 974 975 976 977 978 979
##
980 981 982 983 984 985 986 987 988 989 990 991 992 993 994 995 996
## T1 1 0 0 3 0 0 0 0 0 0 0 0 0 0 0 0
     997 998 999 1000 1001 1002 1003 1004 1005 1006 1007 1008 1009 1010
## T1 1 0 0 0 1 0 0 2 3 1 1 0 0 0
     1011 1012 1013 1014 1015 1016 1017 1018 1019 1020 1021 1022 1023 1024
## T1_1
     0 1 0 0 0 0 3 0 0 0 0
     1025 1026 1027 1028 1029 1030 1031 1032 1033 1034 1035 1036 1037 1038
##
## T1_1
     0 0 0 0 0 3 0 0 0 0 2
     1039 1040 1041 1042 1043 1044 1045 1046 1047 1048 1049 1050 1051 1052
     0 0 0 0 0 0 0 0 0 0 1 0
     1053 1054 1055 1056 1057 1058 1059 1060 1061 1062 1063 1064 1065 1066
     0 0 1 0 0 0 1 0 0 0 0 0
     1067 1068 1069 1070 1071 1072 1073 1074 1075 1076 1077 1078 1079 1080
## T1_1 4 0 0 0 0 0 0 0 0 3 0 0
     1081 1082 1083 1084 1085 1086 1087 1088 1089 1090 1091 1092 1093 1094
```

```
## T1 1 2 0 0 1 0 0 0 1 0 2 2 1 0
     1473 1474 1475 1476 1477 1478 1479 1480 1481 1482 1483 1484 1485 1486
## T1 1 0 0 4 0 0 0 0 0 0 0 0 1
     1487 1488 1489 1490 1491 1492 1493 1494 1495 1496 1497 1498 1499 1500
## T1 1 0 0 0 1 0 0 0 0 1 0 0 0 1
     1501 1502 1503 1504 1505 1506 1507 1508 1509 1510 1511 1512 1513 1514
## T1 1 0 0 0 1 0 0 0 0 0 0 0 0
     1515 1516 1517 1518 1519 1520 1521 1522 1523 1524 1525 1526 1527 1528
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     1529 1530 1531 1532 1533 1534 1535 1536 1537 1538 1539 1540 1541 1542
## T1_1 1 0 0 0 0 0 0 0 0 0 0 0 0
     1543 1544 1545 1546 1547 1548 1549 1550 1551 1552 1553 1554 1555 1556
## T1_1 0 0 0 0 0 0 1 0 1 0 0 0 0
     1557 1558 1559 1560 1561 1562 1563 1564 1565 1566 1567 1568 1569 1570
## T1_1
     0 0 0 0 0 0 0 1 0 0 1 0 0
##
     1571 1572 1573 1574 1575 1576 1577 1578 1579 1580 1581 1582 1583 1584
     0 0 0 0 1 0 0 0 0 0 0 0
## T1_1
     1585 1586 1587 1588 1589 1590 1591 1592 1593 1594 1595 1596 1597 1598
## T1 1 0 0 0 0 0 0 0 0 0 0 0 1 0
     1599 1600 1601 1602 1603 1604 1605 1606 1607 1608 1609 1610 1611 1612
## T1_1 0 0 0 0 0 0 0 1 0 0 0 0
     1613 1614 1615 1616 1617 1618 1619 1620 1621 1622 1623 1624 1625 1626
## T1_1 0 2 0 0 0 0 0 0 0 0 0
     1627 1628 1629 1630 1631 1632 1633 1634 1635 1636 1637 1638 1639 1640
## T1 1 0 0 0 0 3 1 0 0 0 0 0 0
     1641 1642 1643 1644 1645 1646 1647 1648 1649 1650 1651 1652 1653 1654
## T1_1 0 0 0 0 0 0 2 0 2 0 0 0 0
     1655 1656 1657 1658 1659 1660 1661 1662 1663 1664 1665 1666 1667 1668
## T1_1 0 0 0 0 0 0 0 0 1 0 0 0
     1669 1670 1671 1672 1673 1674 1675 1676 1677 1678 1679 1680 1681 1682
## T1_1
     1683 1684 1685 1686 1687 1688 1689 1690 1691 1692 1693 1694 1695 1696
## T1_1 2 0 0 0 0 0 0 0 0 0 0 0 0
     1697 1698 1699 1700 1701 1702 1703 1704 1705 1706 1707 1708 1709 1710
## T1 1 0 0 0 1 0 0 0 0 0 0 0 0 0
     1711 1712 1713 1714 1715 1716 1717 1718 1719 1720 1721 1722 1723 1724
1725 1726 1727 1728 1729 1730 1731 1732 1733 1734 1735 1736 1737 1738
## T1_1 0 0 0 0 0 0 0 0 0 0 1 0 0
     1739 1740 1741 1742 1743 1744 1745 1746 1747 1748 1749 1750 1751 1752
## T1 1 0 0 0 0 0 0 0 2 0 0 0 0 0
     1753 1754 1755 1756 1757 1758 1759 1760 1761 1762 1763 1764 1765 1766
## T1 1 0 0 0 0 0 0 0 10 0 0 1 0
     1767 1768 1769 1770 1771 1772 1773 1774 1775 1776 1777 1778 1779 1780
## T1_1
     0 0 0 0 0 0 0 1 0 0 0
     1781 1782 1783 1784 1785 1786 1787 1788 1789 1790 1791 1792 1793 1794
##
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0 0
     1795 1796 1797 1798 1799 1800 1801 1802 1803 1804 1805 1806 1807 1808
     0 0 0 0 0 0 0 0 0 0 0 0 0
     1809 1810 1811 1812 1813 1814 1815 1816 1817 1818 1819 1820 1821 1822
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 0
     1823 1824 1825 1826 1827 1828 1829 1830 1831 1832 1833 1834 1835 1836
## T1_1 0 3 0 0 0 0 0 0 0 0 0 0
     1837 1838 1839 1840 1841 1842 1843 1844 1845 1846 1847 1848 1849 1850
```

```
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
    4497 4498 4499 4500 4501 4502 4503 4504 4505 4506 4507 4508 4509 4510
## T1 1 0 0 1 0 0 0 0 1 0 0 0 0
    4511 4512 4513 4514 4515 4516 4517 4518 4519 4520 4521 4522 4523 4524
     4525 4526 4527 4528 4529 4530 4531 4532 4533 4534 4535 4536 4537 4538
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
    4539 4540 4541 4542 4543 4544 4545 4546 4547 4548 4549 4550 4551 4552
## T1_1 0 0 0 0 0 0 0 0 0 0 0
    4553 4554 4555 4556 4557 4558 4559 4560 4561 4562 4563 4564 4565 4566
     0 0 0 0 0 0 0 0 0 0 0 0
    4567 4568 4569 4570 4571 4572 4573 4574 4575 4576 4577 4578 4579 4580
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0 0
    4581 4582 4583 4584 4585 4586 4587 4588 4589 4590 4591 4592 4593 4594
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0
     4595 4596 4597 4598 4599 4600 4601 4602 4603 4604 4605 4606 4607 4608
## T1_1
     4609 4610 4611 4612 4613 4614 4615 4616 4617 4618 4619 4620 4621 4622
     4623 4624 4625 4626 4627 4628 4629 4630 4631 4632 4633 4634 4635 4636
## T1_1
     0 0 0 0 0 0 0 0 0 0 1 1 0
    4637 4638 4639 4640 4641 4642 4643 4644 4645 4646 4647 4648 4649 4650
     0 0 0 0 0 0 0 0 0 0 0
## T1_1
    4651 4652 4653 4654 4655 4656 4657 4658 4659 4660 4661 4662 4663 4664
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
    4665 4666 4667 4668 4669 4670 4671 4672 4673 4674 4675 4676 4677 4678
     0 0 0 0 0 0 1 0 0 0 0
    4679 4680 4681 4682 4683 4684 4685 4686 4687 4688 4689 4690 4691 4692
     4693 4694 4695 4696 4697 4698 4699 4700 4701 4702 4703 4704 4705 4706
     0 0 0 1 0 0 0 2 0 0 0 0 0
## T1_1
     4707 4708 4709 4710 4711 4712 4713 4714 4715 4716 4717 4718 4719 4720
     4721 4722 4723 4724 4725 4726 4727 4728 4729 4730 4731 4732 4733 4734
     0 0 0 0 0 0 0 0 0 0 0 0 1
    4735 4736 4737 4738 4739 4740 4741 4742 4743 4744 4745 4746 4747 4748
     f 0 \ \ 0 \ \ 0 \ \ 0 \ \ 0 \ \ 0 \ \ 0 \ \ 1 \ \ 0 \ \ 0
    4749 4750 4751 4752 4753 4754 4755 4756 4757 4758 4759 4760 4761 4762
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 1
    4763 4764 4765 4766 4767 4768 4769 4770 4771 4772 4773 4774 4775 4776
4777 4778 4779 4780 4781 4782 4783 4784 4785 4786 4787 4788 4789 4790
     0 0 0 0 0 1 0 0 0 0 0
## T1 1
    4791 4792 4793 4794 4795 4796 4797 4798 4799 4800 4801 4802 4803 4804
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0
     4805 4806 4807 4808 4809 4810 4811 4812 4813 4814 4815 4816 4817 4818
## T1_1
     4819 4820 4821 4822 4823 4824 4825 4826 4827 4828 4829 4830 4831 4832
     4833 4834 4835 4836 4837 4838 4839 4840 4841 4842 4843 4844 4845 4846
     4847 4848 4849 4850 4851 4852 4853 4854 4855 4856 4857 4858 4859 4860
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
    4861 4862 4863 4864 4865 4866 4867 4868 4869 4870 4871 4872 4873 4874
```

```
## T1 1 0 0 0 0 0 0 0 1 0 0 0 0
     5253 5254 5255 5256 5257 5258 5259 5260 5261 5262 5263 5264 5265 5266
## T1 1 0 1 0 0 0 0 0 0 0 0 0 0
     5267 5268 5269 5270 5271 5272 5273 5274 5275 5276 5277 5278 5279 5280
## T1 1 0 0 0 0 0 1 0 0 0 0 0 0
     5281 5282 5283 5284 5285 5286 5287 5288 5289 5290 5291 5292 5293 5294
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     5295 5296 5297 5298 5299 5300 5301 5302 5303 5304 5305 5306 5307 5308
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     5309 5310 5311 5312 5313 5314 5315 5316 5317 5318 5319 5320 5321 5322
## T1_1 0 0 0 0 0 0 0 0 0 0 1 0
     5323 5324 5325 5326 5327 5328 5329 5330 5331 5332 5333 5334 5335 5336
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 0
     5337 5338 5339 5340 5341 5342 5343 5344 5345 5346 5347 5348 5349 5350
## T1_1
     0 0 0 0 1 0 0 0 0 0 0
     5351 5352 5353 5354 5355 5356 5357 5358 5359 5360 5361 5362 5363 5364
     0 0 1 0 0 0 0 0 0 0 0 0
## T1_1
     5365 5366 5367 5368 5369 5370 5371 5372 5373 5374 5375 5376 5377 5378
## T1 1 0 0 1 0 0 0 0 0 0 0 0 0 0
     5379 5380 5381 5382 5383 5384 5385 5386 5387 5388 5389 5390 5391 5392
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     5393 5394 5395 5396 5397 5398 5399 5400 5401 5402 5403 5404 5405 5406
## T1_1 0 0 0 1 1 0 0 0 0 0 0 0
     5407 5408 5409 5410 5411 5412 5413 5414 5415 5416 5417 5418 5419 5420
5421 5422 5423 5424 5425 5426 5427 5428 5429 5430 5431 5432 5433 5434
     0 0 0 0 0 0 0 0 0 0 0 1 1
     5435 5436 5437 5438 5439 5440 5441 5442 5443 5444 5445 5446 5447 5448
     5449 5450 5451 5452 5453 5454 5455 5456 5457 5458 5459 5460 5461 5462
     0 0 0 0 0 0 0 0 0 1 0 1 0
## T1_1
     5463 5464 5465 5466 5467 5468 5469 5470 5471 5472 5473 5474 5475 5476
     0 0 0 0 0 0 0 0 0 0 0 0
     5477 5478 5479 5480 5481 5482 5483 5484 5485 5486 5487 5488 5489 5490
     5491 5492 5493 5494 5495 5496 5497 5498 5499 5500 5501 5502 5503 5504
     0 0 0 0 0 0 0 0 0 0 0 0
     5505 5506 5507 5508 5509 5510 5511 5512 5513 5514 5515 5516 5517 5518
## T1_1 0 0 0 0 1 0 0 0 0 0 1
     5519 5520 5521 5522 5523 5524 5525 5526 5527 5528 5529 5530 5531 5532
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     5533 5534 5535 5536 5537 5538 5539 5540 5541 5542 5543 5544 5545 5546
     1 0 0 0 0 0 0 0 0 0 0 0
## T1 1
     5547 5548 5549 5550 5551 5552 5553 5554 5555 5556 5557 5558 5559 5560
## T1_1
     \begin{smallmatrix} 0 & & 0 & & 1 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 1 \\ \end{smallmatrix}
     5561 5562 5563 5564 5565 5566 5567 5568 5569 5570 5571 5572 5573 5574
##
## T1_1
     5575 5576 5577 5578 5579 5580 5581 5582 5583 5584 5585 5586 5587 5588
     0 0 0 0 0 0 0 0 0 0 0 0 0
     5589 5590 5591 5592 5593 5594 5595 5596 5597 5598 5599 5600 5601 5602
     5603 5604 5605 5606 5607 5608 5609 5610 5611 5612 5613 5614 5615 5616
## T1_1 0 0 0 1 0 0 0 0 0 0 0 0
     5617 5618 5619 5620 5621 5622 5623 5624 5625 5626 5627 5628 5629 5630
```

```
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     5631 5632 5633 5634 5635 5636 5637 5638 5639 5640 5641 5642 5643 5644
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     5645 5646 5647 5648 5649 5650 5651 5652 5653 5654 5655 5656 5657 5658
     0 0 0 0 0 0 0 0 0 0 0 1
     5659 5660 5661 5662 5663 5664 5665 5666 5667 5668 5669 5670 5671 5672
## T1_1
     0
            0 0 1 1 0 0 0 0 0 0
     5673 5674 5675 5676 5677 5678 5679 5680 5681 5682 5683 5684 5685 5686
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     5687 5688 5689 5690 5691 5692 5693 5694 5695 5696 5697 5698 5699 5700
     5701 5702 5703 5704 5705 5706 5707 5708 5709 5710 5711 5712 5713 5714
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 0
     5715 5716 5717 5718 5719 5720 5721 5722 5723 5724 5725 5726 5727 5728
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0
     5729 5730 5731 5732 5733 5734 5735 5736 5737 5738 5739 5740 5741 5742
     0 0 0 0 0 0 0 0 0 0 0
## T1_1
     5743 5744 5745 5746 5747 5748 5749 5750 5751 5752 5753 5754 5755 5756
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     5757 5758 5759 5760 5761 5762 5763 5764 5765 5766 5767 5768 5769 5770
## T1_1 0 0 0 1 0 0 0 0 0 0 0 0 0
     5771 5772 5773 5774 5775 5776 5777 5778 5779 5780 5781 5782 5783 5784
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     5785 5786 5787 5788 5789 5790 5791 5792 5793 5794 5795 5796 5797 5798
## T1 1 0 0 0 1 0 0 0 0 0 0 0 0
     5799 5800 5801 5802 5803 5804 5805 5806 5807 5808 5809 5810 5811 5812
     0 0 0 0 0 0 0 0 0 1 0 0
     5813 5814 5815 5816 5817 5818 5819 5820 5821 5822 5823 5824 5825 5826
     5827 5828 5829 5830 5831 5832 5833 5834 5835 5836 5837 5838 5839 5840
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0
     5841 5842 5843 5844 5845 5846 5847 5848 5849 5850 5851 5852 5853 5854
     0 0 0 0 0 0 0 1 0 0 0
     5855 5856 5857 5858 5859 5860 5861 5862 5863 5864 5865 5866 5867 5868
     5869 5870 5871 5872 5873 5874 5875 5876 5877 5878 5879 5880 5881 5882
     0 0 0 0 0 0 0 0 0 0 0 0
     5883 5884 5885 5886 5887 5888 5889 5890 5891 5892 5893 5894 5895 5896
##
     0 0 0 0 0 0 1 0 0 0 0
## T1_1
     5897 5898 5899 5900 5901 5902 5903 5904 5905 5906 5907 5908 5909 5910
## T1 1 0 0 0 0 0 0 0 0 0 0 1 0
     5911 5912 5913 5914 5915 5916 5917 5918 5919 5920 5921 5922 5923 5924
     5925 5926 5927 5928 5929 5930 5931 5932 5933 5934 5935 5936 5937 5938
## T1_1
     0 0 0 0 0 0 1 0 0 0 0 0
     5939 5940 5941 5942 5943 5944 5945 5946 5947 5948 5949 5950 5951 5952
## T1_1
     0 0 0 0 0 0 0 1 0 0 0
     5953 5954 5955 5956 5957 5958 5959 5960 5961 5962 5963 5964 5965 5966
     0 0 0 0 0 0 0 0 0 0 0 0
     5967 5968 5969 5970 5971 5972 5973 5974 5975 5976 5977 5978 5979 5980
     0 0 1 1 0 0 0 1 0 0 0 1
     5981 5982 5983 5984 5985 5986 5987 5988 5989 5990 5991 5992 5993 5994
## T1_1 0 0
             0 0 0 0 0 0 0 0
                                          0
                                               0
     5995 5996 5997 5998 5999 6000 6001 6002 6003 6004 6005 6006 6007 6008
```

```
## T1_1 0 1 0 1 0 0 0 0 0 0 0 0
     6765 6766 6767 6768 6769 6770 6771 6772 6773 6774 6775 6776 6777 6778
## T1 1 0 0 0 0 0 1 0 0 0 1 0 0
     6779 6780 6781 6782 6783 6784 6785 6786 6787 6788 6789 6790 6791 6792
## T1 1 0 0 0 0 0 0 0 2 0 0 0 0
     6793 6794 6795 6796 6797 6798 6799 6800 6801 6802 6803 6804 6805 6806
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     6807 6808 6809 6810 6811 6812 6813 6814 6815 6816 6817 6818 6819 6820
## T1_1 0 0 1 0 0 0 0 0 0 0 0 0
     6821 6822 6823 6824 6825 6826 6827 6828 6829 6830 6831 6832 6833 6834
## T1_1 0 0 0 0 0 0 0 1 0 1 0 0
     6835 6836 6837 6838 6839 6840 6841 6842 6843 6844 6845 6846 6847 6848
## T1_1 0 0 0 0 0 1 0 0 0 1 0 0
     6849 6850 6851 6852 6853 6854 6855 6856 6857 6858 6859 6860 6861 6862
## T1_1
     0 1 0 0 0 0 0 0 0 1 0 0
     6863 6864 6865 6866 6867 6868 6869 6870 6871 6872 6873 6874 6875 6876
## T1_1
     6877 6878 6879 6880 6881 6882 6883 6884 6885 6886 6887 6888 6889 6890
## T1 1 0 0 0 0 0 0 0 0 0 1 0 0 0
     6891 6892 6893 6894 6895 6896 6897 6898 6899 6900 6901 6902 6903 6904
## T1_1 0 0 0 0 0 0 0 1 0 0 0 0
     6905 6906 6907 6908 6909 6910 6911 6912 6913 6914 6915 6916 6917 6918
## T1_1 1 0 0 0 0 0 0 0 0 0 1 0
     6919 6920 6921 6922 6923 6924 6925 6926 6927 6928 6929 6930 6931 6932
## T1 1 0 0 0 0 1 0 0 0 0 0 0 0
     6933 6934 6935 6936 6937 6938 6939 6940 6941 6942 6943 6944 6945 6946
## T1_1 0 0 0 0 0 1 0 0 0 0 0
     6947 6948 6949 6950 6951 6952 6953 6954 6955 6956 6957 6958 6959 6960
     6961 6962 6963 6964 6965 6966 6967 6968 6969 6970 6971 6972 6973 6974
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0 0
     6975 6976 6977 6978 6979 6980 6981 6982 6983 6984 6985 6986 6987 6988
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 0
     6989 6990 6991 6992 6993 6994 6995 6996 6997 6998 6999 7000 7001 7002
     0 0 0 0 0 0 0 0 0 0 1 1 0
     7003 7004 7005 7006 7007 7008 7009 7010 7011 7012 7013 7014 7015 7016
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     7017 7018 7019 7020 7021 7022 7023 7024 7025 7026 7027 7028 7029 7030
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     7031 7032 7033 7034 7035 7036 7037 7038 7039 7040 7041 7042 7043 7044
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     7045 7046 7047 7048 7049 7050 7051 7052 7053 7054 7055 7056 7057 7058
     7059 7060 7061 7062 7063 7064 7065 7066 7067 7068 7069 7070 7071 7072
     0 0 0 0 0 1 0 0 0 0 0
     7073 7074 7075 7076 7077 7078 7079 7080 7081 7082 7083 7084 7085 7086
## T1_1
     0 0 0 0 1 0 0 0 0 0 0 0
     7087 7088 7089 7090 7091 7092 7093 7094 7095 7096 7097 7098 7099 7100
     0 0 0 0 0 0 0 0 0 0 0 0 0
     7101 7102 7103 7104 7105 7106 7107 7108 7109 7110 7111 7112 7113 7114
     0 0 0 0 1 0 0 1 0 0 0 0
     7115 7116 7117 7118 7119 7120 7121 7122 7123 7124 7125 7126 7127 7128
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     7129 7130 7131 7132 7133 7134 7135 7136 7137 7138 7139 7140 7141 7142
```

```
7143 7144 7145 7146 7147 7148 7149 7150 7151 7152 7153 7154 7155 7156
## T1 1 0 1 1 0 0 0 0 0 0 0 0 0 0
        7157 7158 7159 7160 7161 7162 7163 7164 7165 7166 7167 7168 7169 7170
## T1 1 0 0 0 0 0 0 1 0 0 0 0 0
        7171 7172 7173 7174 7175 7176 7177 7178 7179 7180 7181 7182 7183 7184
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
        7185 7186 7187 7188 7189 7190 7191 7192 7193 7194 7195 7196 7197 7198
## T1_1 0 0 0 0 0 0 0 0 0 0 0
        7199 7200 7201 7202 7203 7204 7205 7206 7207 7208 7209 7210 7211 7212
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 0
        7213 7214 7215 7216 7217 7218 7219 7220 7221 7222 7223 7224 7225 7226
## T1_1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
        7227 7228 7229 7230 7231 7232 7233 7234 7235 7236 7237 7238 7239 7240
## T1_1
         0 0 0 1 0 2 0 0 0 0 0
         7241 7242 7243 7244 7245 7246 7247 7248 7249 7250 7251 7252 7253 7254
## T1_1
         7255 7256 7257 7258 7259 7260 7261 7262 7263 7264 7265 7266 7267 7268
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
        7269 7270 7271 7272 7273 7274 7275 7276 7277 7278 7279 7280 7281 7282
7283 7284 7285 7286 7287 7288 7289 7290 7291 7292 7293 7294 7295 7296
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
        7297 7298 7299 7300 7301 7302 7303 7304 7305 7306 7307 7308 7309 7310
7311 7312 7313 7314 7315 7316 7317 7318 7319 7320 7321 7322 7323 7324
         7325 7326 7327 7328 7329 7330 7331 7332 7333 7334 7335 7336 7337 7338
## T1_1 0 0 0 0 0 0 0 0 1 1 0 0
        7339 7340 7341 7342 7343 7344 7345 7346 7347 7348 7349 7350 7351 7352
## T1_1
         7353 7354 7355 7356 7357 7358 7359 7360 7361 7362 7363 7364 7365 7366
## T1_1 0 0 0 0 2 0 0 0 0 0 0 0
        7367 7368 7369 7370 7371 7372 7373 7374 7375 7376 7377 7378 7379 7380
         \begin{smallmatrix} 0 & & 0 & & 1 & & 1 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & & 0 & 
        7381 7382 7383 7384 7385 7386 7387 7388 7389 7390 7391 7392 7393 7394
## T1 1 0 0 0 1 0 0 0 0 1 0 0 0
        7395 7396 7397 7398 7399 7400 7401 7402 7403 7404 7405 7406 7407 7408
## T1_1 0 0 0 0 0 1 0 0 0 0 1 0
        7409 7410 7411 7412 7413 7414 7415 7416 7417 7418 7419 7420 7421 7422
## T1 1 0 0 0 0 0 0 0 0 0 0 1
        7423 7424 7425 7426 7427 7428 7429 7430 7431 7432 7433 7434 7435 7436
         7437 7438 7439 7440 7441 7442 7443 7444 7445 7446 7447 7448 7449 7450
         0 0 0 0 0 0 0 0 0 0 0 0
         7451 7452 7453 7454 7455 7456 7457 7458 7459 7460 7461 7462 7463 7464
## T1_1
         7465 7466 7467 7468 7469 7470 7471 7472 7473 7474 7475 7476 7477 7478
         7479 7480 7481 7482 7483 7484 7485 7486 7487 7488 7489 7490 7491 7492
         7493 7494 7495 7496 7497 7498 7499 7500 7501 7502 7503 7504 7505 7506
## T1_1 0 0 0 0 0 0 0 0 0 0 0
        7507 7508 7509 7510 7511 7512 7513 7514 7515 7516 7517 7518 7519 7520
```

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## T1 1 0 0 0 0 1 1 0 0 0 0 1 1
     7521 7522 7523 7524 7525 7526 7527 7528 7529 7530 7531 7532 7533 7534
7535 7536 7537 7538 7539 7540 7541 7542 7543 7544 7545 7546 7547 7548
## T1 1 0 0 0 0 0 0 1 0 0 0 0 0
     7549 7550 7551 7552 7553 7554 7555 7556 7557 7558 7559 7560 7561 7562
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     7563 7564 7565 7566 7567 7568 7569 7570 7571 7572 7573 7574 7575 7576
## T1_1 0 0 0 0 0 0 0 0 0 0 1
     7577 7578 7579 7580 7581 7582 7583 7584 7585 7586 7587 7588 7589 7590
7591 7592 7593 7594 7595 7596 7597 7598 7599 7600 7601 7602 7603 7604
## T1_1 0 0 0 0 0 0 0 1 1 0 0 0
     7605 7606 7607 7608 7609 7610 7611 7612 7613 7614 7615 7616 7617 7618
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0 0
     7619 7620 7621 7622 7623 7624 7625 7626 7627 7628 7629 7630 7631 7632
## T1_1
     0 0 0 0 0 0 0 1 0 0 0
     7633 7634 7635 7636 7637 7638 7639 7640 7641 7642 7643 7644 7645 7646
## T1 1 0 0 0 1 0 0 0 0 0 0 0 0 0
     7647 7648 7649 7650 7651 7652 7653 7654 7655 7656 7657 7658 7659 7660
7661 7662 7663 7664 7665 7666 7667 7668 7669 7670 7671 7672 7673 7674
     0 0 0 0 0 0 0 0 0 0 1
## T1_1
     7675 7676 7677 7678 7679 7680 7681 7682 7683 7684 7685 7686 7687 7688
## T1 1 0 0 0 0 0 0 0 1 0 0 0
     7689 7690 7691 7692 7693 7694 7695 7696 7697 7698 7699 7700 7701 7702
     0 0 0 0 0 0 1 0 0 0 0
     7703 7704 7705 7706 7707 7708 7709 7710 7711 7712 7713 7714 7715 7716
7717 7718 7719 7720 7721 7722 7723 7724 7725 7726 7727 7728 7729 7730
## T1_1
     0 0 0 0 0 0 0 0 0 0 1 1 0
     7731 7732 7733 7734 7735 7736 7737 7738 7739 7740 7741 7742 7743 7744
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     7745 7746 7747 7748 7749 7750 7751 7752 7753 7754 7755 7756 7757 7758
     0 0 0 0 0 0 0 0 0 0 0 1 0
     7759 7760 7761 7762 7763 7764 7765 7766 7767 7768 7769 7770 7771 7772
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     7773 7774 7775 7776 7777 7778 7779 7780 7781 7782 7783 7784 7785 7786
## T1_1 0 0 0 0 0 1 0 0 1 0 0 0
     7787 7788 7789 7790 7791 7792 7793 7794 7795 7796 7797 7798 7799 7800
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     7801 7802 7803 7804 7805 7806 7807 7808 7809 7810 7811 7812 7813 7814
     0 0 0 0 0 0 0 0 0 0 1 0
     7815 7816 7817 7818 7819 7820 7821 7822 7823 7824 7825 7826 7827 7828
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 1
     7829 7830 7831 7832 7833 7834 7835 7836 7837 7838 7839 7840 7841 7842
## T1_1
     7843 7844 7845 7846 7847 7848 7849 7850 7851 7852 7853 7854 7855 7856
     7857 7858 7859 7860 7861 7862 7863 7864 7865 7866 7867 7868 7869 7870
     0 0 0 0 1 0 0 0 0 0 0 0
     7871 7872 7873 7874 7875 7876 7877 7878 7879 7880 7881 7882 7883 7884
## T1_1 0 0 0 0 0 0 0 0 0 0 1
     7885 7886 7887 7888 7889 7890 7891 7892 7893 7894 7895 7896 7897 7898
```

```
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     7899 7900 7901 7902 7903 7904 7905 7906 7907 7908 7909 7910 7911 7912
## T1 1 0 0 0 1 0 0 0 0 0 0 0 0
     7913 7914 7915 7916 7917 7918 7919 7920 7921 7922 7923 7924 7925 7926
## T1 1 0 1 0 0 0 0 0 0 0 0 0 0
     7927 7928 7929 7930 7931 7932 7933 7934 7935 7936 7937 7938 7939 7940
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     7941 7942 7943 7944 7945 7946 7947 7948 7949 7950 7951 7952 7953 7954
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     7955 7956 7957 7958 7959 7960 7961 7962 7963 7964 7965 7966 7967 7968
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     7969 7970 7971 7972 7973 7974 7975 7976 7977 7978 7979 7980 7981 7982
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0 0
     7983 7984 7985 7986 7987 7988 7989 7990 7991 7992 7993 7994 7995 7996
## T1_1
     7997 7998 7999 8000 8001 8002 8003 8004 8005 8006 8007 8008 8009 8010
## T1_1
     0 0 0 0 0 1 0 0 1 0 0 0
     8011 8012 8013 8014 8015 8016 8017 8018 8019 8020 8021 8022 8023 8024
8025 8026 8027 8028 8029 8030 8031 8032 8033 8034 8035 8036 8037 8038
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 0
     8039 8040 8041 8042 8043 8044 8045 8046 8047 8048 8049 8050 8051 8052
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     8053 8054 8055 8056 8057 8058 8059 8060 8061 8062 8063 8064 8065 8066
8067 8068 8069 8070 8071 8072 8073 8074 8075 8076 8077 8078 8079 8080
## T1_1 0 1 1 0 0 0 0 0 1 0 0 0
     8081 8082 8083 8084 8085 8086 8087 8088 8089 8090 8091 8092 8093 8094
## T1_1 0 0 1 0 0 0 0 0 0 0 0 0
     8095 8096 8097 8098 8099 8100 8101 8102 8103 8104 8105 8106 8107 8108
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0 0
     8109 8110 8111 8112 8113 8114 8115 8116 8117 8118 8119 8120 8121 8122
## T1_1 0 0 0 0 0 0 0 0 1 0 0 0
     8123 8124 8125 8126 8127 8128 8129 8130 8131 8132 8133 8134 8135 8136
## T1 1 0 0 0 0 0 1 0 0 0 0 0 0
     8137 8138 8139 8140 8141 8142 8143 8144 8145 8146 8147 8148 8149 8150
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     8151 8152 8153 8154 8155 8156 8157 8158 8159 8160 8161 8162 8163 8164
## T1 1 0 0 1 0 0 0 1 0 0 0 0 0
     8165 8166 8167 8168 8169 8170 8171 8172 8173 8174 8175 8176 8177 8178
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     8179 8180 8181 8182 8183 8184 8185 8186 8187 8188 8189 8190 8191 8192
## T1 1 0 0 0 0 0 0 0 0 0 1 0 0
     8193 8194 8195 8196 8197 8198 8199 8200 8201 8202 8203 8204 8205 8206
## T1_1
     0 0 0 0 1 0 0 0 0 0 0
     8207 8208 8209 8210 8211 8212 8213 8214 8215 8216 8217 8218 8219 8220
## T1_1
     1 0 0 0 0 0 0 1 0 0 0 0 0
     8221 8222 8223 8224 8225 8226 8227 8228 8229 8230 8231 8232 8233 8234
## T1_1 1 0 0 0 0 0 0 0 0 0 0 0 0
     8235 8236 8237 8238 8239 8240 8241 8242 8243 8244 8245 8246 8247 8248
     8249 8250 8251 8252 8253 8254 8255 8256 8257 8258 8259 8260 8261 8262
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     8263 8264 8265 8266 8267 8268 8269 8270 8271 8272 8273 8274 8275 8276
```

```
## T1 1 0 1 0 0 0 0 0 0 0 0 0 0
     9789 9790 9791 9792 9793 9794 9795 9796 9797 9798 9799 9800 9801 9802
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     9803 9804 9805 9806 9807 9808 9809 9810 9811 9812 9813 9814 9815 9816
## T1 1 0 0 0 0 0 0 0 0 1 0 0 0 0
     9817 9818 9819 9820 9821 9822 9823 9824 9825 9826 9827 9828 9829 9830
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0
     9831 9832 9833 9834 9835 9836 9837 9838 9839 9840 9841 9842 9843 9844
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     9845 9846 9847 9848 9849 9850 9851 9852 9853 9854 9855 9856 9857 9858
## T1_1 0 0 0 0 0 0 1 0 0 0 0
     9859 9860 9861 9862 9863 9864 9865 9866 9867 9868 9869 9870 9871 9872
## T1 1 0 0 0 0 0 0 0 1 0 0 0 0
     9873 9874 9875 9876 9877 9878 9879 9880 9881 9882 9883 9884 9885 9886
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 1 0
     9887 9888 9889 9890 9891 9892 9893 9894 9895 9896 9897 9898 9899 9900
## T1_1
     0 0 0 0 0 0 0 0 0 0 0 0 0
     9901 9902 9903 9904 9905 9906 9907 9908 9909 9910 9911 9912 9913 9914
## T1 1 0 0 0 0 0 0 0 0 0 0 0 0 0
     9915 9916 9917 9918 9919 9920 9921 9922 9923 9924 9925 9926 9927 9928
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0 0
     9929 9930 9931 9932 9933 9934 9935 9936 9937 9938 9939 9940 9941 9942
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     9943 9944 9945 9946 9947 9948 9949 9950 9951 9952 9953 9954 9955 9956
## T1 1 0 0 0 0 0 1 0 0 0 0 0
     9957 9958 9959 9960 9961 9962 9963 9964 9965 9966 9967 9968 9969 9970
9971 9972 9973 9974 9975 9976 9977 9978 9979 9980 9981 9982 9983 9984
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     9985 9986 9987 9988 9989 9990 9991 9992 9993 9994 9995 9996 9997 9998
     0 0 0 0 0 0 0 0 0 0 0 0
## T1_1
     9999 10000 10001 10002 10003 10004 10005 10006 10007 10008 10009
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     10010 10011 10012 10013 10014 10015 10016 10017 10018 10019 10020
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10021 10022 10023 10024 10025 10026 10027 10028 10029 10030 10031
     0 0 0 0 0 0 0 0 0 0
     10032 10033 10034 10035 10036 10037 10038 10039 10040 10041 10042
## T1 1 0 0 0 0 0 0 0 0 0 0
     10043 10044 10045 10046 10047 10048 10049 10050 10051 10052 10053
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10054 10055 10056 10057 10058 10059 10060 10061 10062 10063 10064
## T1 1 0 0 0 0 0 0 0 0 0 0
     10065 10066 10067 10068 10069 10070 10071 10072 10073 10074 10075
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10076 10077 10078 10079 10080 10081 10082 10083 10084 10085 10086
##
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10087 10088 10089 10090 10091 10092 10093 10094 10095 10096 10097
     0 0 0 0 0 0 0 0 0 0
     10098 10099 10100 10101 10102 10103 10104 10105 10106 10107 10108
## T1_1 0 0 0 0 0 0 0 0 0 0
     10109 10110 10111 10112 10113 10114 10115 10116 10117 10118 10119
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     10120 10121 10122 10123 10124 10125 10126 10127 10128 10129 10130
```

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## T1 1 0 0 0 0 0 0 0 0 0 0
## 10131 10132 10133 10134 10135 10136 10137 10138 10139 10140 10141
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10142 10143 10144 10145 10146 10147 10148 10149 10150 10151 10152
10153 10154 10155 10156 10157 10158 10159 10160 10161 10162 10163
## T1 1 0 0 0 0 0 0 1 1 0 0 0
     10164 10165 10166 10167 10168 10169 10170 10171 10172 10173 10174
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10175 10176 10177 10178 10179 10180 10181 10182 10183 10184 10185
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     10186 10187 10188 10189 10190 10191 10192 10193 10194 10195 10196
## T1 1 0 0 0 0 0 0 0 0 0 0
     10197 10198 10199 10200 10201 10202 10203 10204 10205 10206 10207
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10208 10209 10210 10211 10212 10213 10214 10215 10216 10217 10218
## T1_1
     0 0 0 1 0 0 0 0 0 0
     10219 10220 10221 10222 10223 10224 10225 10226 10227 10228 10229
## T1 1 0 0 0 0 0 0 0 0 0 0
     10230 10231 10232 10233 10234 10235 10236 10237 10238 10239 10240
## T1_1 0 0 0 0 0 0 0 0 0 0
     10241 10242 10243 10244 10245 10246 10247 10248 10249 10250 10251
## T1_1 0 0 0 0 0 0 0 0 0 0
     10252 10253 10254 10255 10256 10257 10258 10259 10260 10261 10262
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10263 10264 10265 10266 10267 10268 10269 10270 10271 10272 10273
## T1_1 0 1 0 0 0 0 0 0 0 0 0
     10274 10275 10276 10277 10278 10279 10280 10281 10282 10283 10284
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     10285 10286 10287 10288 10289 10290 10291 10292 10293 10294 10295
     0 0 0 0 0 0 0 0 0 0
## T1_1
     10296 10297 10298 10299 10300 10301 10302 10303 10304 10305 10306
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     10307 10308 10309 10310 10311 10312 10313 10314 10315 10316 10317
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10318 10319 10320 10321 10322 10323 10324 10325 10326 10327 10328
## T1 1 0 0 0 0 0 0 1 0 0 0
     10329 10330 10331 10332 10333 10334 10335 10336 10337 10338 10339
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10340 10341 10342 10343 10344 10345 10346 10347 10348 10349 10350
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10351 10352 10353 10354 10355 10356 10357 10358 10359 10360 10361
## T1 1 0 0 1 0 0 0 0 2 0 0
     10362 10363 10364 10365 10366 10367 10368 10369 10370 10371 10372
## T1_1
     0 1 0 1 0 0 0 0 0 0
     10373 10374 10375 10376 10377 10378 10379 10380 10381 10382 10383
##
## T1_1
     2 0 0 0 0 0 0 0 0 0 0
     10384 10385 10386 10387 10388 10389 10390 10391 10392 10393 10394
## T1_1 0 0 0 0 0 0 0 0 0 0
     10395 10396 10397 10398 10399 10400 10401 10402 10403 10404 10405
## T1_1 0 0 0 0 0 0 0 0 0 0
     10406 10407 10408 10409 10410 10411 10412 10413 10414 10415 10416
## T1_1 0 0 0 0 0 0 0 0 0 0 0
   10417 10418 10419 10420 10421 10422 10423 10424 10425 10426 10427
```

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## T1 1 0 0 0 0 1 0 0 1 0 0
     10428 10429 10430 10431 10432 10433 10434 10435 10436 10437 10438
## T1 1 0 0 0 0 0 0 0 0 1 0 0
     10439 10440 10441 10442 10443 10444 10445 10446 10447 10448 10449
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10450 10451 10452 10453 10454 10455 10456 10457 10458 10459 10460
## T1 1 0 0 0 0 0 0 0 0 1 0
     10461 10462 10463 10464 10465 10466 10467 10468 10469 10470 10471
## T1_1 1 0 0 0 0 0 0 0 0 0
     10472 10473 10474 10475 10476 10477 10478 10479 10480 10481 10482
## T1_1 0 0 0 0 0 0 0 0 0 0
     10483 10484 10485 10486 10487 10488 10489 10490 10491 10492 10493
## T1_1 0 0 0 0 0 1 0 0 0
     10494 10495 10496 10497 10498 10499 10500 10501 10502 10503 10504
## T1_1
     0 0 0 0 0 0 0 1 0 0
     10505 10506 10507 10508 10509 10510 10511 10512 10513 10514 10515
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10516 10517 10518 10519 10520 10521 10522 10523 10524 10525 10526
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10527 10528 10529 10530 10531 10532 10533 10534 10535 10536 10537
## T1_1 0 0 0 0 0 0 0 0 0 0
     10538 10539 10540 10541 10542 10543 10544 10545 10546 10547 10548
     0 0 0 0 0 0 0 0 0 0
## T1_1
     10549 10550 10551 10552 10553 10554 10555 10556 10557 10558 10559
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10560 10561 10562 10563 10564 10565 10566 10567 10568 10569 10570
## T1_1 0 0 0 0 0 1 0 0 0
     10571 10572 10573 10574 10575 10576 10577 10578 10579 10580 10581
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     10582 10583 10584 10585 10586 10587 10588 10589 10590 10591 10592
## T1_1
     0 0 0 1 1 1 0 0 0 0
     10593 10594 10595 10596 10597 10598 10599 10600 10601 10602 10603
## T1_1 0 0 0 0 0 0 0 1 0 0
     10604 10605 10606 10607 10608 10609 10610 10611 10612 10613 10614
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10615 10616 10617 10618 10619 10620 10621 10622 10623 10624 10625
## T1 1 0 0 0 0 0 2 0 0 0 0
     10626 10627 10628 10629 10630 10631 10632 10633 10634 10635 10636
10637 10638 10639 10640 10641 10642 10643 10644 10645 10646 10647
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10648 10649 10650 10651 10652 10653 10654 10655 10656 10657 10658
10659 10660 10661 10662 10663 10664 10665 10666 10667 10668 10669
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10670 10671 10672 10673 10674 10675 10676 10677 10678 10679 10680
##
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10681 10682 10683 10684 10685 10686 10687 10688 10689 10690 10691
## T1_1 0 0 0 0 1 0 0 0 1 0
     10692 10693 10694 10695 10696 10697 10698 10699 10700 10701 10702
## T1_1 0 0 0 0 0 0 0 0 0 0
     10703 10704 10705 10706 10707 10708 10709 10710 10711 10712 10713
## T1_1 0 0 0 0 0 0 0 0 0 0
    10714 10715 10716 10717 10718 10719 10720 10721 10722 10723 10724
```

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## T1 1 0 0 0 0 0 0 0 0 0 0
     10725 10726 10727 10728 10729 10730 10731 10732 10733 10734 10735
## T1 1 0 0 0 0 1 1 0 0 0 0
     10736 10737 10738 10739 10740 10741 10742 10743 10744 10745 10746
## T1 1 0 0 0 1 0 0 0 0 0 0
     10747 10748 10749 10750 10751 10752 10753 10754 10755 10756 10757
## T1 1 0 2 1 0 0 0 1 0 0 0 1
     10758 10759 10760 10761 10762 10763 10764 10765 10766 10767 10768
10769 10770 10771 10772 10773 10774 10775 10776 10777 10778 10779
## T1_1 0 0 0 0 0 0 0 0 0 1
     10780 10781 10782 10783 10784 10785 10786 10787 10788 10789 10790
## T1 1 0 1 0 0 0 0 0 0 0 0
     10791 10792 10793 10794 10795 10796 10797 10798 10799 10800 10801
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10802 10803 10804 10805 10806 10807 10808 10809 10810 10811 10812
## T1_1
     0 0 0 0 0 0 0 1 0 0
     10813 10814 10815 10816 10817 10818 10819 10820 10821 10822 10823
## T1 1 0 0 0 0 0 0 0 0 1 0 0
     10824 10825 10826 10827 10828 10829 10830 10831 10832 10833 10834
10835 10836 10837 10838 10839 10840 10841 10842 10843 10844 10845
## T1_1 0 0 0 0 0 0 0 0 0 0
     10846 10847 10848 10849 10850 10851 10852 10853 10854 10855 10856
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10857 10858 10859 10860 10861 10862 10863 10864 10865 10866 10867
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     10868 10869 10870 10871 10872 10873 10874 10875 10876 10877 10878
## T1_1 0 0 0 0 0 1 0 0 1
     10879 10880 10881 10882 10883 10884 10885 10886 10887 10888 10889
     0 0 0 0 0 0 0 0 0 1
## T1_1
     10890 10891 10892 10893 10894 10895 10896 10897 10898 10899 10900
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     10901 10902 10903 10904 10905 10906 10907 10908 10909 10910 10911
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10912 10913 10914 10915 10916 10917 10918 10919 10920 10921 10922
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10923 10924 10925 10926 10927 10928 10929 10930 10931 10932 10933
## T1 1 0 0 0 0 0 0 0 0 0 0
     10934 10935 10936 10937 10938 10939 10940 10941 10942 10943 10944
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     10945 10946 10947 10948 10949 10950 10951 10952 10953 10954 10955
## T1 1 0 0 0 0 0 0 0 0 0 0
     10956 10957 10958 10959 10960 10961 10962 10963 10964 10965 10966
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10967 10968 10969 10970 10971 10972 10973 10974 10975 10976 10977
##
## T1_1
     0 0 0 0 0 0 0 0 0 0
     10978 10979 10980 10981 10982 10983 10984 10985 10986 10987 10988
## T1_1 0 0 0 0 0 0 0 0 0 0
     10989 10990 10991 10992 10993 10994 10995 10996 10997 10998 10999
## T1_1 0 0 0 0 0 0 0 0 0 0
     11000 11001 11002 11003 11004 11005 11006 11007 11008 11009 11010
## T1_1 0 0 0 0 0 0 0 0 0 0
     11011 11012 11013 11014 11015 11016 11017 11018 11019 11020 11021
```

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## T1_1 0 0 0 0 0 0 0 0 0 0
     11616 11617 11618 11619 11620 11621 11622 11623 11624 11625 11626
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11627 11628 11629 11630 11631 11632 11633 11634 11635 11636 11637
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11638 11639 11640 11641 11642 11643 11644 11645 11646 11647 11648
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11649 11650 11651 11652 11653 11654 11655 11656 11657 11658 11659
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11660 11661 11662 11663 11664 11665 11666 11667 11668 11669 11670
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     11671 11672 11673 11674 11675 11676 11677 11678 11679 11680 11681
## T1_1 0 0 0 0 0 0 0 0 1 1
     11682 11683 11684 11685 11686 11687 11688 11689 11690 11691 11692
## T1_1
     0 0 0 0 0 0 0 0 1 0
##
     11693 11694 11695 11696 11697 11698 11699 11700 11701 11702 11703
## T1_1
     0 0 0 0 0 0 0 0 0 0
     11704 11705 11706 11707 11708 11709 11710 11711 11712 11713 11714
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11715 11716 11717 11718 11719 11720 11721 11722 11723 11724 11725
## T1_1 0 0 0 0 0 0 0 0 0 0
     11726 11727 11728 11729 11730 11731 11732 11733 11734 11735 11736
## T1_1
     0 0 0 0 0 0 0 0 0 0
     11737 11738 11739 11740 11741 11742 11743 11744 11745 11746 11747
## T1 1 0 0 0 0 0 1 0 0 0 0
     11748 11749 11750 11751 11752 11753 11754 11755 11756 11757 11758
## T1_1 0 0 0 0 1 0 0 0 0 0
     11759 11760 11761 11762 11763 11764 11765 11766 11767 11768 11769
## T1_1 0 0 0 1 0 0 0 0 0 0
     11770 11771 11772 11773 11774 11775 11776 11777 11778 11779 11780
## T1_1
     0 0 0 0 0 0 0 0 0 0
     11781 11782 11783 11784 11785 11786 11787 11788 11789 11790 11791
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     11792 11793 11794 11795 11796 11797 11798 11799 11800 11801 11802
## T1 1 0 0 0 1 0 1 0 0 0
     11803 11804 11805 11806 11807 11808 11809 11810 11811 11812 11813
## T1 1 0 0 0 0 0 0 0 0 2 0 0
     11814 11815 11816 11817 11818 11819 11820 11821 11822 11823 11824
## T1 1 0 0 0 0 0 0 0 0 0 1 0
     11825 11826 11827 11828 11829 11830 11831 11832 11833 11834 11835
## T1 1 0 0 0 1 0 0 0 0 0 0
     11836 11837 11838 11839 11840 11841 11842 11843 11844 11845 11846
## T1 1 0 0 0 1 0 0 0 0 0 0
     11847 11848 11849 11850 11851 11852 11853 11854 11855 11856 11857
## T1_1
     0 0 0 0 0 0 0 0 0 0
     11858 11859 11860 11861 11862 11863 11864 11865 11866 11867 11868
##
## T1_1
     0 0 0 0 1 0 0 0 0 0
     11869 11870 11871 11872 11873 11874 11875 11876 11877 11878 11879
## T1_1 0 0 0 0 0 0 1 0 0
     11880 11881 11882 11883 11884 11885 11886 11887 11888 11889 11890
## T1_1 0 0 0 0 0 0 0 0 0 0
     11891 11892 11893 11894 11895 11896 11897 11898 11899 11900 11901
## T1_1 0 0 0 0 0 0 0 0 0 0
    11902 11903 11904 11905 11906 11907 11908 11909 11910 11911 11912
```

```
## T1 1 0 0 0 0 0 0 1 0 0
    11913 11914 11915 11916 11917 11918 11919 11920 11921 11922 11923
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11924 11925 11926 11927 11928 11929 11930 11931 11932 11933 11934
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11935 11936 11937 11938 11939 11940 11941 11942 11943 11944 11945
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11946 11947 11948 11949 11950 11951 11952 11953 11954 11955 11956
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     11957 11958 11959 11960 11961 11962 11963 11964 11965 11966 11967
## T1_1 0 0 0 0 0 0 0 0 0 0
     11968 11969 11970 11971 11972 11973 11974 11975 11976 11977 11978
## T1 1 0 0 0 0 0 0 1 0 0 0
     11979 11980 11981 11982 11983 11984 11985 11986 11987 11988 11989
## T1_1
     0 0 0 0 0 0 0 0 0 0
##
     11990 11991 11992 11993 11994 11995 11996 11997 11998 11999 12000
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12001 12002 12003 12004 12005 12006 12007 12008 12009 12010 12011
## T1 1 0 0 0 0 0 1 0 0 0 0
     12012 12013 12014 12015 12016 12017 12018 12019 12020 12021 12022
## T1_1 0 0 0 0 0 0 0 0 0 0
     12023 12024 12025 12026 12027 12028 12029 12030 12031 12032 12033
## T1_1 0 0 0 0 0 0 0 0 0 1 0
     12034 12035 12036 12037 12038 12039 12040 12041 12042 12043 12044
## T1 1 0 0 0 0 0 1 0 0 0 0
     12045 12046 12047 12048 12049 12050 12051 12052 12053 12054 12055
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     12056 12057 12058 12059 12060 12061 12062 12063 12064 12065 12066
## T1_1 1 0 0 0 0 0 0 0 0 0 0
     12067 12068 12069 12070 12071 12072 12073 12074 12075 12076 12077
     0 0 0 0 0 0 0 0 0 0
## T1_1
##
     12078 12079 12080 12081 12082 12083 12084 12085 12086 12087 12088
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     12089 12090 12091 12092 12093 12094 12095 12096 12097 12098 12099
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12100 12101 12102 12103 12104 12105 12106 12107 12108 12109 12110
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12111 12112 12113 12114 12115 12116 12117 12118 12119 12120 12121
## T1 1 0 0 0 1 0 0 0 0 0 0
     12122 12123 12124 12125 12126 12127 12128 12129 12130 12131 12132
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12133 12134 12135 12136 12137 12138 12139 12140 12141 12142 12143
## T1 1 0 0 0 0 0 0 0 1 1 0
     12144 12145 12146 12147 12148 12149 12150 12151 12152 12153 12154
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12155 12156 12157 12158 12159 12160 12161 12162 12163 12164 12165
##
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12166 12167 12168 12169 12170 12171 12172 12173 12174 12175 12176
     0 0 0 0 0 1 0 0 0 0
     12177 12178 12179 12180 12181 12182 12183 12184 12185 12186 12187
## T1_1 0 0 1 0 0 0 0 0 0 0
     12188 12189 12190 12191 12192 12193 12194 12195 12196 12197 12198
## T1_1 0 0 0 0 0 0 0 0 0 0
    12199 12200 12201 12202 12203 12204 12205 12206 12207 12208 12209
```

```
## T1_1 1 0 0 0 0 0 0 0 0 0
    12210 12211 12212 12213 12214 12215 12216 12217 12218 12219 12220
## T1 1 0 1 0 0 0 0 0 0 0 0 0
     12221 12222 12223 12224 12225 12226 12227 12228 12229 12230 12231
## T1 1 0 1 0 0 0 0 0 0 0 0 0
     12232 12233 12234 12235 12236 12237 12238 12239 12240 12241 12242
## T1 1 0 0 0 1 0 0 0 0 0 0
     12243 12244 12245 12246 12247 12248 12249 12250 12251 12252 12253
## T1 1 0 0 0 0 0 0 0 0 0 0
     12254 12255 12256 12257 12258 12259 12260 12261 12262 12263 12264
     1 0 0 0 0 0 0 0 0 0
     12265 12266 12267 12268 12269 12270 12271 12272 12273 12274 12275
## T1_1
     0 0 0 0 0 0 0 0 0 1
     12276 12277 12278 12279 12280 12281 12282 12283 12284 12285 12286
## T1_1
     0 0 0 0 0 0 1 1 0 0 0
##
     12287 12288 12289 12290 12291 12292 12293 12294 12295 12296 12297
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12298 12299 12300 12301 12302 12303 12304 12305 12306 12307 12308
## T1 1 0 0 0 0 0 0 0 1 0 0
     12309 12310 12311 12312 12313 12314 12315 12316 12317 12318 12319
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     12320 12321 12322 12323 12324 12325 12326 12327 12328 12329 12330
     0 0 0 0 0 0 0 0 0 0
## T1_1
     12331 12332 12333 12334 12335 12336 12337 12338 12339 12340 12341
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12342 12343 12344 12345 12346 12347 12348 12349 12350 12351 12352
## T1_1 0 0 0 0 1 0 0 0 0 0
     12353 12354 12355 12356 12357 12358 12359 12360 12361 12362 12363
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12364 12365 12366 12367 12368 12369 12370 12371 12372 12373 12374
     0 0 1 0 0 0 0 0 0 0
## T1_1
##
     12375 12376 12377 12378 12379 12380 12381 12382 12383 12384 12385
## T1_1 0 0 1 0 0 0 0 0 0 0 0
     12386 12387 12388 12389 12390 12391 12392 12393 12394 12395 12396
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12397 12398 12399 12400 12401 12402 12403 12404 12405 12406 12407
## T1 1 0 0 0 0 1 0 0 0 0 0
     12408 12409 12410 12411 12412 12413 12414 12415 12416 12417 12418
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12419 12420 12421 12422 12423 12424 12425 12426 12427 12428 12429
## T1 1 0 0 0 1 0 0 0 1 0 0
     12430 12431 12432 12433 12434 12435 12436 12437 12438 12439 12440
## T1 1 0 0 0 0 0 0 0 0 0 0
     12441 12442 12443 12444 12445 12446 12447 12448 12449 12450 12451
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12452 12453 12454 12455 12456 12457 12458 12459 12460 12461 12462
##
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12463 12464 12465 12466 12467 12468 12469 12470 12471 12472 12473
     0 0 0 0 0 0 0 0 0 0
     12474 12475 12476 12477 12478 12479 12480 12481 12482 12483 12484
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12485 12486 12487 12488 12489 12490 12491 12492 12493 12494 12495
## T1_1 0 0 0 0 0 0 0 0 0 0
    12496 12497 12498 12499 12500 12501 12502 12503 12504 12505 12506
```

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## T1_1 0 0 0 0 0 0 0 0 0 0
     12507 12508 12509 12510 12511 12512 12513 12514 12515 12516 12517
## T1 1 0 0 1 0 0 0 0 1 1 0 0
     12518 12519 12520 12521 12522 12523 12524 12525 12526 12527 12528
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12529 12530 12531 12532 12533 12534 12535 12536 12537 12538 12539
## T1 1 0 0 0 0 0 0 0 0 0 0
     12540 12541 12542 12543 12544 12545 12546 12547 12548 12549 12550
## T1 1 0 0 0 0 0 0 0 0 0 0
     12551 12552 12553 12554 12555 12556 12557 12558 12559 12560 12561
     0 0 0 0 0 0 0 0 0 0
     12562 12563 12564 12565 12566 12567 12568 12569 12570 12571 12572
## T1_1 0 0 0 0 0 0 0 0 0 0
     12573 12574 12575 12576 12577 12578 12579 12580 12581 12582 12583
## T1_1
     0 0 1 0 0 0 0 0 0 0
##
     12584 12585 12586 12587 12588 12589 12590 12591 12592 12593 12594
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12595 12596 12597 12598 12599 12600 12601 12602 12603 12604 12605
## T1 1 0 0 0 0 0 0 0 0 0 0
     12606 12607 12608 12609 12610 12611 12612 12613 12614 12615 12616
## T1_1 0 0 0 0 0 0 0 0 0 0
     12617 12618 12619 12620 12621 12622 12623 12624 12625 12626 12627
     0 0 0 0 0 0 0 0 0 0
## T1_1
     12628 12629 12630 12631 12632 12633 12634 12635 12636 12637 12638
## T1 1 0 0 0 0 0 0 1 0 0 0
     12639 12640 12641 12642 12643 12644 12645 12646 12647 12648 12649
## T1_1 0 0 0 0 0 0 0 1 0 0
     12650 12651 12652 12653 12654 12655 12656 12657 12658 12659 12660
## T1_1 0 0 0 0 0 0 0 0 0 0 0 0
     12661 12662 12663 12664 12665 12666 12667 12668 12669 12670 12671
## T1_1
     0 0 0 0 0 0 0 0 0 0
##
     12672 12673 12674 12675 12676 12677 12678 12679 12680 12681 12682
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     12683 12684 12685 12686 12687 12688 12689 12690 12691 12692 12693
## T1_1 0 0 0 0 0 0 0 1 0 1
     12694 12695 12696 12697 12698 12699 12700 12701 12702 12703 12704
     0 0 0 0 0 0 0 0 0 0 0
     12705 12706 12707 12708 12709 12710 12711 12712 12713 12714 12715
     0 0 0 0 0 0 0 0 0 0
## T1 1
     12716 12717 12718 12719 12720 12721 12722 12723 12724 12725 12726
     0 0 0 0 0 0 0 0 0 0
     12727 12728 12729 12730 12731 12732 12733 12734 12735 12736 12737
## T1 1 0 0 0 0 0 0 0 0 0 0
     12738 12739 12740 12741 12742 12743 12744 12745 12746 12747 12748
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12749 12750 12751 12752 12753 12754 12755 12756 12757 12758 12759
##
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12760 12761 12762 12763 12764 12765 12766 12767 12768 12769 12770
     0 0 0 0 0 0 0 0 0 0
     12771 12772 12773 12774 12775 12776 12777 12778 12779 12780 12781
     0 0 0 0 0 0 0 1 0 0
     12782 12783 12784 12785 12786 12787 12788 12789 12790 12791 12792
## T1_1 0 0 0 0 0 0 0 0 0 0
    12793 12794 12795 12796 12797 12798 12799 12800 12801 12802 12803
```

```
## T1 1 0 0 0 0 0 1 0 0 0
     12804 12805 12806 12807 12808 12809 12810 12811 12812 12813 12814
## T1 1 0 0 0 0 0 1 0 0 0 0
     12815 12816 12817 12818 12819 12820 12821 12822 12823 12824 12825
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12826 12827 12828 12829 12830 12831 12832 12833 12834 12835 12836
## T1 1 0 0 0 0 1 0 0 0 0 0
     12837 12838 12839 12840 12841 12842 12843 12844 12845 12846 12847
## T1 1 0 1 1 0 0 0 0 0 0 0
     12848 12849 12850 12851 12852 12853 12854 12855 12856 12857 12858
     0 0 0 0 0 0 0 1 0 0
     12859 12860 12861 12862 12863 12864 12865 12866 12867 12868 12869
## T1_1 0 0 0 0 0 0 0 0 0 0
     12870 12871 12872 12873 12874 12875 12876 12877 12878 12879 12880
## T1_1
     0 0 0 0 0 0 0 0 0 0
##
     12881 12882 12883 12884 12885 12886 12887 12888 12889 12890 12891
## T1_1
     0 0 0 0 0 0 0 0 0 0
     12892 12893 12894 12895 12896 12897 12898 12899 12900 12901 12902
## T1 1 0 0 0 0 0 1 0 0 0
     12903 12904 12905 12906 12907 12908 12909 12910 12911 12912 12913
## T1_1 0 0 0 0 0 0 0 0 0 0
     12914 12915 12916 12917 12918 12919 12920 12921 12922 12923 12924
     0 0 0 0 0 0 0 0 0 0
## T1_1
     12925 12926 12927 12928 12929 12930 12931 12932 12933 12934 12935
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12936 12937 12938 12939 12940 12941 12942 12943 12944 12945 12946
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     12947 12948 12949 12950 12951 12952 12953 12954 12955 12956 12957
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     12958 12959 12960 12961 12962 12963 12964 12965 12966 12967 12968
     0 0 0 0 0 0 0 0 0 0
## T1_1
     12969 12970 12971 12972 12973 12974 12975 12976 12977 12978 12979
## T1_1 0 0 0 0 0 0 0 0 0 0 0
     12980 12981 12982 12983 12984 12985 12986 12987 12988 12989 12990
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     12991 12992 12993 12994 12995 12996 12997 12998 12999 13000 13001
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     13002 13003 13004 13005 13006 13007 13008 13009 13010 13011 13012
## T1 1 0 0 0 1 0 0 0 0 0 0
     13013 13014 13015 13016 13017 13018 13019 13020 13021 13022 13023
## T1 1 0 0 0 0 0 0 0 0 0 0 0
     13024 13025 13026 13027 13028 13029 13030 13031 13032 13033 13034
## T1 1 0 0 0 0 0 0 0 0 0 0
     13035 13036 13037 13038 13039 13040 13041 13042 13043 13044 13045
## T1_1
     0 0 0 0 0 0 0 0 0 0
     13046 13047 13048 13049 13050 13051 13052 13053 13054 13055 13056
##
## T1_1
     0 0 0 0 0 0 0 1 0 0
     13057 13058 13059 13060 13061 13062 13063 13064 13065 13066 13067
## T1_1 0 0 0 0 0 0 0 0 0 0
     13068 13069 13070 13071 13072 13073 13074 13075 13076 13077 13078
## T1_1 0 0 0 0 0 0 0 0 0 0
     13079 13080 13081 13082 13083 13084 13085 13086 13087 13088 13089
## T1_1 0 0 0 0 0 0 0 0 0 0
    13090 13091 13092 13093 13094 13095 13096 13097 13098 13099 13100
```

```
1
                         1
                                1
                                      0
##
        13101 13102 13103 13104 13105 13106 13107 13108 13109 13110 13111
## T1_1
                          1
                                2
                                       0
                                                          0
        13112 13113 13114 13115 13116 13117 13118 13119 13120 13121 13122
##
##
  T1 1
             0
                   0
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
                                                                 0
        13123 13124 13125 13126 13127 13128 13129 13130 13131 13132 13133
##
## T1_1
                   0
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
        13134 13135 13136 13137 13138 13139 13140 13141 13142 13143 13144
##
## T1_1
                   0
                          0
                                0
                                      0
                                             0
                                                    0
                                                          0
                                                                 0
                                                                       0
             0
                                                                              0
##
        13145 13146 13147 13148 13149 13150 13151 13152 13153 13154 13155
##
                   0
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
                                                                 0
  T1_1
##
        13156 13157 13158 13159 13160 13161 13162 13163 13164 13165 13166
## T1_1
                          0
                                0
                                      0
                                             0
                                                    0
                                                          0
             0
                   0
                                                                 0
                                                                       0
##
        13167 13168 13169 13170 13171 13172 13173 13174 13175 13176 13177
## T1_1
                          0
                                0
                                      0
                                             0
                                                    1
                                                          0
                                                                 0
##
        13178 13179 13180 13181 13182 13183 13184 13185 13186 13187 13188
## T1_1
                   0
                                      0
                                                    0
                                                          0
             0
                          0
                                1
                                             0
                                                                 0
##
        13189 13190 13191 13192 13193 13194 13195 13196 13197 13198 13199
##
                                0
                                      0
                                                    0
  T1 1
                   0
                         0
                                             0
                                                          1
                                                                0
##
        13200 13201 13202 13203 13204 13205 13206 13207 13208 13209 13210
## T1_1
            0
                   0
                          0
                                0
                                      0
                                             0
                                                    0
                                                          0
                                                                 0
                                                                       0
        13211 13212 13213 13214 13215 13216 13217 13218 13219 13220 13221
##
## T1_1
                   0
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
                                                                 0
##
        13222 13223 13224 13225 13226 13227 13228 13229 13230 13231 13232
## T1 1
                   0
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
                                                                 0
##
        13233 13234 13235 13236 13237 13238 13239 13240 13241 13242 13243
##
  T1_1
                   0
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
                                                                 0
        13244 13245 13246 13247 13248 13249 13250 13251 13252 13253 13254
##
## T1_1
                   0
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
                                                                 0
                                                                       0
##
        13255 13256 13257 13258 13259 13260 13261 13262 13263 13264 13265
## T1_1
                          0
                                0
                                       0
                                             0
                                                    0
                                                          0
##
        13266 13267 13268 13269 13270 13271 13272 13273 13274 13275 13276
## T1_1
                   0
                          0
                                       0
                                             0
                                                    0
                                                          0
                                0
        13277 13278 13279 13280 13281 13282 13283 13284 13285 13286 13287
##
                                      0
                                             0
                                                    0
                                                          0
##
  T1 1
                   0
                         0
                                0
                                                                 0
        13288 13289 13290 13291 13292 13293 13294 13295 13296 13297 13298
##
## T1 1
                   0
                                0
                                       0
                                                          0
        13299 13300 13301 13302 13303 13304 13305 13306 13307 13308 13309
##
                                0
                                       0
                                             1
                                                    0
## T1_1
             1
                   0
                          0
##
        13310
## T1 1
S.obs(soilbac1)
## T1 1
## 1074
```

Question 3: Answer the following questions about the soil bacterial dataset.

- a. How many sequences did we recover from the sample soilbac1, i.e. N?
- b. What is the observed richness of soilbac1?

C(soilbac1)

0.6479471

T1 1

c. How does coverage compare between the BCI sample (site1) and the KBS sample (soilbac1)?

```
Answer 3a: 13310 
Answer 3b: 1074 
Answer 3c: C(soilbac1) = 64.8\%; C(site1) = 93.1\% Coverage of BCI sample is better than one of KBS sample.
```

Richness Estimators

In the R code chunk below, do the following:

- 1. Write a function to calculate **Chao1**,
- 2. Write a function to calculate **Chao2**,
- 3. Write a function to calculate **ACE**, and
- 4. Use these functions to estimate richness at both site1 and soilbac1.

```
S.chao1 <- function(x = ""){S.obs(x) + (sum(x==1)^2) / (2 * sum(x==2))}
S.chao2 <- function(site = "", SbyS = ""){</pre>
  SbyS = as.data.frame(SbyS)
  x = SbyS[site, ]
  SbyS.pa <- (SbyS > 0) * 1 # convert the SbyS to presence/absence
  Q1 = sum(colSums(SbyS.pa) ==1)
  Q2 = sum(colSums(SbyS.pa) == 2)
  S.chao2 = S.obs(x) + (Q1^2)/(2 * Q2)
  return(S.chao2)}
S.ace <- function(x = "", thresh = 10){
  x \leftarrow x[x>0] # excludes zero-abundance taxa
  S.abund <- length(which(x > thresh)) # richness of abundant taxa
  S.rare <- length(which(x <= thresh)) # richness of rare taxa
  singlt <- length(which(x == 1)) # number of singleton taxa</pre>
  N.rare <- sum(x[which(x <= thresh)]) # abundance of rare individuals</pre>
  C.ace <- 1 - (singlt / N.rare) # coverage (prop non-singlt rare inds)</pre>
  i <- c(1:thresh) # threshold abundance range
  count <- function(i, y){ # counter to go through i range</pre>
  length(y[y == i])}
  a.1 <- sapply(i, count, x) # number of individuals in richness i richness classes
  f.1 \leftarrow (i * (i - 1)) * a.1 # k(k-1)kf sensu Gotelli
  G.ace <- (S.rare/C.ace)*(sum(f.1)/(N.rare*(N.rare-1)))</pre>
  S.ace <- S.abund + (S.rare/C.ace) + (singlt/C.ace) * max(G.ace,0)
  return(S.ace)
}
S.chao1(site1)
##
## 119.6944
S.chao2(1, BCI)
## 104.6053
S.ace(site1)
## [1] 159.3404
```

```
S.chao1(soilbac1)

## T1_1
## 2628.514

S.chao2(1, soilbac.t)

## T1_1
## 21055.39

S.ace(soilbac1)

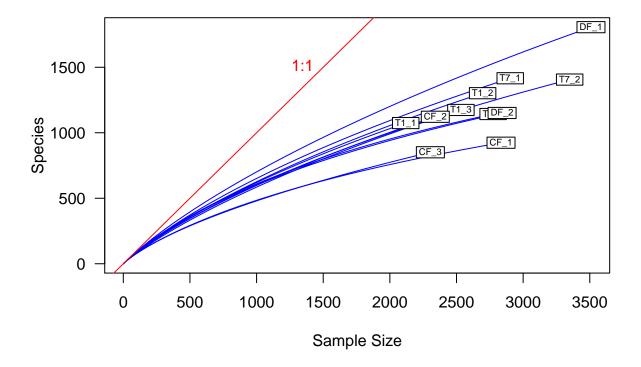
## [1] 4465.983
```

Rarefaction

In the R code chunk below, please do the following:

- 1. Calculate observed richness for all samples in soilbac,
- 2. Determine the size of the smallest sample,
- 3. Use the rarefy() function to rarefy each sample to this level,
- 4. Plot the rarefaction results, and
- 5. Add the 1:1 line and label.

```
soilbac.S <- S.obs(soilbac.t)
min.N <- min(rowSums(soilbac.t))
S.rarefy <- rarefy(x = soilbac.t, sample = min.N, se = TRUE)
rarecurve(x = soilbac.t, step = 20, col = "blue", cex = 0.6, las=1)
abline(0, 1, col = 'red')
text(1500, 1500, "1:1", pos = 2, col = 'red')</pre>
```



Question 4: What is the difference between ACE and the Chao estimators?

Answer 4: Chao estimaters take into account only singltons and doubletons while ACE estimator considers abundance of rare species (species detected less than 'threshold' times).

4) SPECIES EVENNESS

Here, we consider how abundance varies among species, that is, species evenness.

Visualizing Evenness: The Rank Abundance Curve (RAC)

One of the most common ways to visualize evenness is in a **rank-abundance curve** (sometime referred to as a rank-abundance distribution or Whittaker plot). An RAC can be constructed by ranking species from the most abundant to the least abundant without respect to species labels (and hence no worries about 'ties' in abundance).

In the R code chunk below, do the following:

- 1. Write a function to construct a RAC,
- 2. Be sure your function removes species that have zero abundances,
- 3. Order the vector (RAC) from greatest (most abundant) to least (least abundant), and
- 4. Return the ranked vector

```
RAC <- function(x = ""){
  x = as.vector(x)</pre>
```

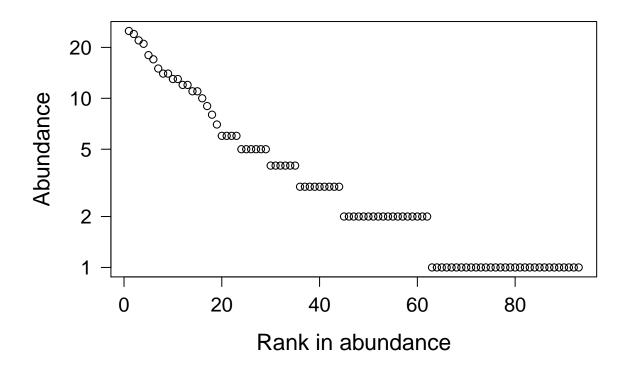
```
x.ab = x[x > 0]
x.ab.ranked = x.ab[order(x.ab, decreasing = TRUE)]
return(x.ab.ranked)}
```

Now, let's examine the RAC for site1 of the BCI data set.

In the R code chunk below, do the following:

- 1. Create a sequence of ranks and plot the RAC with natural-log-transformed abundances,
- 2. Label the x-axis "Rank in abundance" and the y-axis "log(abundance)"

```
rac \leftarrow RAC(x = site1)
ranks <- as.vector(seq(1, length(rac)))
opar <- par(no.readonly = TRUE)</pre>
                                                        # Saves default plot parameters
par(mar = c(5.1, 5.1, 4.1, 2.1))
                                                        # New settings for par
plot(ranks, log(rac), type = 'p', axes = F,
                                                        # Plots w/o axes
 xlab = "Rank in abundance", ylab = "Abundance",
 las = 1, cex.lab = 1.4, cex.axis = 1.25)
box()
                                                         # Manually adds border
axis(side = 1, labels = T, cex.axis = 1.25)
                                                         # Manually adds X-axis
axis(side = 2, las = 1, cex.axis = 1.25,
                                                         # Manually adds Log-Scaled Y-axis
labels = c(1, 2, 5, 10, 20), at = log(c(1, 2, 5, 10, 20)))
```



Question 5: What effect does visualizing species abundance data on a log-scaled axis have on how we interpret evenness in the RAC?

Answer 5: Species abundance data are more convenient to perceive if log-scaled axis is used.

Now that we have visualized unevennes, it is time to quantify it using Simpson's evenness $(E_{1/D})$ and Smith and Wilson's evenness index (E_{var}) .

Simpson's evenness $(E_{1/D})$

In the R code chunk below, do the following:

- 1. Write the function to calculate $E_{1/D}$, and
- 2. Calculate $E_{1/D}$ for site1.

```
SimpE <- function(x = ""){
    S <- S.obs(x)
    x = as.data.frame(x)
    D <- diversity(x, "inv")
    E <- (D)/S
    return(E)}
SimpE(site1)
## 1
## 0.4238232</pre>
```

Smith and Wilson's evenness index (E_{var})

In the R code chunk below, please do the following:

- 1. Write the function to calculate E_{var} ,
- 2. Calculate E_{var} for site1, and
- 3. Compare $E_{1/D}$ and E_{var} .

```
Evar <- function(x){
    x <- as.vector(x [x > 0])
    1 - (2/pi)*atan(var(log(x)))}
Evar(site1)
```

```
## [1] 0.5067211
```

```
DeltaEvenness = Evar(site1) - SimpE(site1)
DeltaEvenness
```

```
## 1
## 0.08289795
```

Question 6: Compare estimates of evenness for site1 of BCI using $E_{1/D}$ and E_{var} . Do they agree? If so, why? If not, why? What can you infer from the results.

Answer 6: $E_{1/D}$ is close but not equal to E_{var} (lower by 20%). E_{var} decreases input of more abundant species comparing to $E_{1/D}$.

5) INTEGRATING RICHNESS AND EVENNESS: DIVERSITY METRICS

So far, we have introduced two primary aspects of diversity, i.e., richness and evenness. Here, we will use popular indices to estimate diversity, which explicitly incorporate richness and evenness We will write our own diversity functions and compare them against the functions in vegan.

Shannon's diversity (a.k.a., Shannon's entropy)

In the R code chunk below, please do the following:

- 1. Provide the code for calculating H' (Shannon's diversity),
- 2. Compare this estimate with the output of vegan's diversity function using method = "shannon".

```
ShanH <- function(x = ""){
    H = 0
    for (n_i in x){
        if(n_i > 0) {
            p = n_i / sum(x)
            H = H - p*log(p)}}
    return(H)}

ShanH(site1)

## [1] 4.018412
diversity(site1, index = "shannon")
```

Simpson's diversity (or dominance)

[1] 4.018412

In the R code chunk below, please do the following:

- 1. Provide the code for calculating D (Simpson's diversity),
- 2. Calculate both the inverse (1/D) and 1 D,
- 3. Compare this estimate with the output of vegan's diversity function using method = "simp".

```
SimpD \leftarrow function(x = ""){
  D = 0
  N = sum(x)
  for (n_i in x){
    D = D + (n_i^2)/(N^2)
  return(D)}
D.inv <- 1/SimpD(site1)</pre>
D.sub <- 1-SimpD(site1)</pre>
print(D.inv)
## [1] 39.41555
print(D.sub)
## [1] 0.9746293
diversity(site1, "inv")
## [1] 39.41555
diversity(site1, "simp")
## [1] 0.9746293
```

Question 7: Compare estimates of evenness for site1 of BCI using $E_{H'}$ and E_{var} . Do they agree? If so, why? If not, why? What can you infer from the results.

Answer 7: It is Question 6. **Answer 7**: "vegan"s diversity functions return the same values as functions SimpD

Fisher's α

In the R code chunk below, please do the following:

- 1. Provide the code for calculating Fisher's α ,
- 2. Calculate Fisher's α for site1 of BCI.

```
rac <- as.vector(site1[site1 > 0])
invD <- diversity(rac, "inv")
invD
## [1] 39.41555
Fisher <- fisher.alpha(rac)
Fisher</pre>
```

[1] 35.67297

Question 8: How is Fisher's α different from $E_{H'}$ and E_{var} ? What does Fisher's α take into account that $E_{H'}$ and E_{var} do not?

Answer 8: Fisher's alpha is fitted parameter in model for the RAC, diversity metrics (takes into account both richness and evenness), while E(H') and Evar are measures of evenness.

6) MOVING BEYOND UNIVARIATE METRICS OF α DIVERSITY

The diversity metrics that we just learned about attempt to integrate richness and evenness into a single, univariate metric. Although useful, information is invariably lost in this process. If we go back to the rank-abundance curve, we can retrieve additional information – and in some cases – make inferences about the processes influencing the structure of an ecological system.

Species abundance models

The RAC is a simple data structure that is both a vector of abundances. It is also a row in the site-by-species matrix (minus the zeros, i.e., absences).

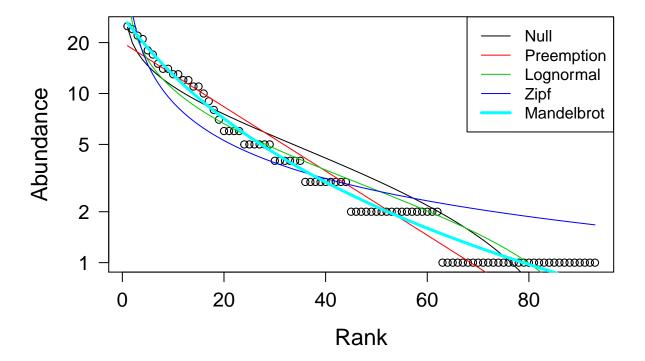
Predicting the form of the RAC is the first test that any biodiversity theory must pass and there are no less than 20 models that have attempted to explain the uneven form of the RAC across ecological systems.

In the R code chunk below, please do the following:

- 1. Use the radfit() function in the vegan package to fit the predictions of various species abundance models to the RAC of site1 in BCI,
- 2. Display the results of the radfit() function, and
- 3. Plot the results of the radfit() function using the code provided in the handout.

```
RACresults <- radfit(site1)
print(RACresults)</pre>
```

```
##
## RAD models, family poisson
  No. of species 93, total abundance 448
##
##
              par1
                         par2
                                  par3
                                           Deviance AIC
## Null
                                            39.5261 315.4362 315.4362
## Preemption
               0.042797
                                            21.8939 299.8041 302.3367
                                            25.1528 305.0629 310.1281
## Lognormal
               1.0687
                          1.0186
## Zipf
               0.11033
                         -0.74705
                                            61.0465 340.9567 346.0219
## Mandelbrot
               100.52
                         -2.312
                                   24.084
                                             4.2271 286.1372 293.7350
plot.new()
plot(RACresults, las = 1, cex.lab = 1.4, cex.axis = 1.25)
```



Question 9: Answer the following questions about the rank abundance curves: a) Based on the output of radfit() and plotting above, discuss which model best fits our rank-abundance curve for site1? b) Can we make any inferences about the forces, processes, and/or mechanisms influencing the structure of our system, e.g., an ecological community?

Answer 9a: From the outpur of radfit() function we can see that 5 models were fitted to the data. Deviance of Mandelbrot model is the lowest in the list, as well as AIc and BIC, which means that this model best fits the rank-abundance curve (Mandelbrot model is the only model in the list that uses three parameters). The second best model is one parametric Preemption model. Answer 9b

Question 10: Answer the following questions about the preemption model: a. What does the preemption model assume about the relationship between total abundance (N) and total resources that can be preempted? b. Why does the niche preemption model look like a straight line in the RAD plot?

Answer 10a: The more resources that can be preemted (the the same alpha) the higher is total abundance. **Answer 10b:** It represents exponential equation

Question 11: Why is it important to account for the number of parameters a model uses when judging how well it explains a given set of data?

Answer 11: The higher number of parameters the easier (usually) to find those which will better describe data.

SYNTHESIS

1. As stated by Magurran (2004) the $D = \sum p_i^2$ derivation of Simpson's Diversity only applies to communities of infinite size. For anything but an infinitely large community, Simpson's Diversity index is calculated as $D = \sum \frac{n_i(n_i-1)}{N(N-1)}$. Assuming a finite community, calculate Simpson's D, 1 - D, and Simpson's inverse (i.e. 1/D) for site 1 of the BCI site-by-species matrix.

2. Along with the rank-abundance curve (RAC), another way to visualize the distribution of abundance among species is with a histogram (a.k.a., frequency distribution) that shows the frequency of different abundance classes.

For example, in a given sample, there may be 10 species represented by a single individual, 8 species with two individuals, 4 species with three individuals, and so on. In fact, the rank-abundance curve and the frequency distribution are the two most common ways to visualize the species-abundance distribution (SAD) and to test species abundance models and biodiversity theories. To address this homework question, use the R function hist() to plot the frequency distribution for site 1 of the BCI site-by-species matrix, and describe the general pattern you see.

plot.new() hist(rac, main="Frequency distribution of species in site1", xlab="Species", col="yellow", freq = TRUE, breaks=length(rac))

3. We asked you to find a biodiversity dataset with your partner. This data could be one of your own or it could be something that you obtained from the literature. Load that dataset.

frogtable <- read.table("data/Frogdata.txt", quote = "", sep = ",", header = TRUE, fill = TRUE) str(frogtable)

How many sites are there? 15480 How many species are there in the entire site-by-species matrix? 122 Any other interesting observations based on what you learned this week?

SUBMITTING YOUR ASSIGNMENT

Use Knitr to create a PDF of your completed alpha_assignment.Rmd document, push it to GitHub, and create a pull request. Please make sure your updated repo include both the HTML and RMarkdown files.

Unless otherwise noted, this assignment is due on Wednesday, January 25th, 2015 at 12:00 PM (noon).