NxWater - 16S

November 20, 2023

Contents

1 PCoA

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## Analysis of Variance Table
##
## Response: Distances
             Df Sum Sq Mean Sq F value
                                           Pr(>F)
             9 0.43831 0.048702 3.5636 0.0003632 ***
## Residuals 231 3.15694 0.013666
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Analysis of Variance Table
## Response: Distances
             Df Sum Sq Mean Sq F value Pr(>F)
## Groups
             1 0.04924 0.049238
                                   4.291 0.03939 *
## Residuals 239 2.74246 0.011475
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Analysis of Variance Table
##
## Response: Distances
             Df Sum Sq Mean Sq F value Pr(>F)
              2 0.07907 0.039533 3.4424 0.03359 *
## Groups
## Residuals 238 2.73322 0.011484
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Permutation test for adonis under reduced model
## Terms added sequentially (first to last)
## Permutation: free
## Number of permutations: 999
## adonis2(formula = dist ~ Nitrogen * Water * Sample.date, data = d)
##
                              Df SumOfSqs
                                              R2
                                                       F Pr(>F)
## Nitrogen
                                   0.636 0.01464 2.4872 0.006 **
```

```
0.426 0.00981 3.3313 0.006 **
## Water
                              1
## Sample.date
                              9
                                   12.545 0.28874 10.8992 0.001 ***
                             2
                                 0.452 0.01041 1.7690 0.044 *
## Nitrogen:Water
## Nitrogen:Sample.date
                                   2.651 0.06101 1.1516 0.115
                             18
## Water:Sample.date
                              8
                                   1.355 0.03119 1.3245 0.068
## Nitrogen:Water:Sample.date 16 1.851 0.04259 0.9044 0.775
## Residual
                             184
                                   23.532 0.54160
## Total
                                   43.449 1.00000
                             240
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## [1] 0.014642254 0.009805738 0.288736017 0.010414291 0.061012791 0.031189213
## [7] 0.042594729 0.541604969 1.000000000
## [1] 0.458395
## Permutation test for dbrda under reduced model
## Permutation: free
## Number of permutations: 999
## Model: dbrda(formula = dist ~ Nitrogen * Water * Sample.date, data = d)
            Df SumOfSqs
                            F Pr(>F)
                 19.917 2.7809 0.001 ***
            56
## Model
## Residual 184
                 23.532
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
```

Bacteria (Bray–Curtis dbRDA)

