

# Figure3A

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## Intro

Compares the functional diversity and richness between the communities associated to the semesters Summer-Autumn and Winter-Spring, to reproduce the analyses and the Figure 3A from the original publication Seasonal dynamics of the coastal microbiome and its association with environmental factors.

### 1. Set the environment

```
library(tidyverse)
library(vegan)
```

### 2. Load data

opus\_workable.tsv contains the rarefied OPUs abundance profiles, with samples as rows and OPUs as columns.

date2season2community.tsv is table mapping the date, season, and community columns.

```
ABUND <- read_tsv("../data/opus_workable.tsv.gz", show_col_types = FALSE)
DATE2SEASON2COMMUNITY <- read_tsv("../data/date2season2community.tsv", show_col_types = FALSE)
```

### 3. Format ABUND as wide

```
ABUND <- ABUND %>%
  pivot_wider(values_fill = 0,
              values_from = abundance,
              names_from = opu_id) %>%
  column_to_rownames("Date")
```

### 4. Compute diversity

Compute Shannon diversity, observed richness and Chao1 estimator.

```
div <- diversity(ABUND)
rich_obs <- estimateR(ABUND) ["S.obs",]
rich_chao1 <- estimateR(ABUND) ["S.chao1",]

output_long_df <- data.frame(Shannon = div,
                             Richness_obs = rich_obs,
                             Richness_chao1 = rich_chao1) %>%
  rownames_to_column(var = "Date") %>%
  pivot_longer(names_to = "Index", values_to = "value",
               cols = -Date) %>%
  left_join(DATE2SEASON2COMMUNITY %>%
```

```
select(Date, Community) %>%
  mutate(Date= as.character(Date)),
  by = "Date")
```

## 5.Run ANOVA Welch test

Perform ANOVA Welch test to evaluate if the diversity and richness differences are significant.

```
anova_welch_shannon <- output_long_df %>%
  filter(Index == "Shannon") %>%
  oneway.test(value ~ Community,
              data = ., var.equal = FALSE)

anova_welch_shannon
```

```
##
## One-way analysis of means (not assuming equal variances)
##
## data: value and Community
## F = 2.731, num df = 1.000, denom df = 10.878, p-value = 0.127
```

```
anova_welch_rich_obs <- output_long_df %>%
  filter(Index == "Richness_obs") %>%
  oneway.test(value ~ Community,
              data = ., var.equal = FALSE)

anova_welch_rich_obs
```

```
##
## One-way analysis of means (not assuming equal variances)
##
## data: value and Community
## F = 2.1394, num df = 1.000, denom df = 12.207, p-value = 0.1688
```

```
anova_welch_rich_chao <- output_long_df %>%
  filter(Index == "Richness_chao1") %>%
  oneway.test(value ~ Community,
              data = ., var.equal = FALSE)

anova_welch_rich_chao
```

```
##
## One-way analysis of means (not assuming equal variances)
##
## data: value and Community
## F = 1.4454, num df = 1.000, denom df = 14.699, p-value = 0.2483
```

## 5. Plot diversity and richness value distributions

```
text_size <- 12

plot_labels <- c("Shannon" = "Shannon diversity",
                 "Richness_obs" = "Observed richness",
                 "Richness_chao1" = "Chao1 estimator")

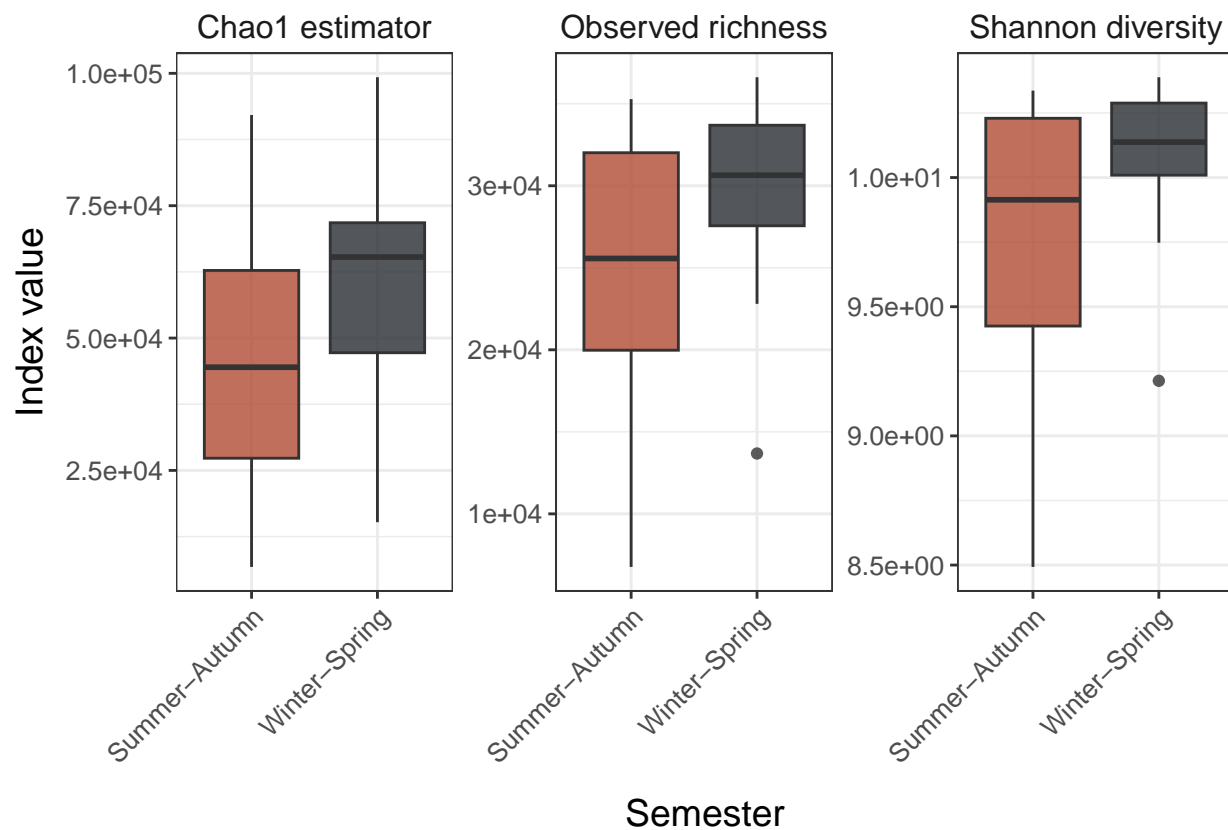
div_plot <- output_long_df %>%
```

```

filter(Index %in% c("Shannon", "Richness_obs", "Richness_chao1")) %>%
ggplot(aes(x = Community, y = value, fill = Community)) +
facet_wrap(~Index, scales = "free", labeller = as_labeller(plot_labels)) +
xlab("Semester") +
ylab("Index value") +
scale_y_continuous(labels = scales::scientific_format()) +
scale_x_discrete(labels = c("S1" = "Summer-Autumn", "S2" = "Winter-Spring")) +
# scale_fill_manual(values = c("#e28743", "#063970")) +
scale_fill_manual(values = c("#b14b34", "#292c33")) +
geom_boxplot(alpha = 0.8) +
theme_bw() +
theme(
  axis.text.x = element_text(size = text_size-2, angle = 45, hjust = 1),
  axis.text.y = element_text(size = text_size-2),
  axis.title.x = element_text(size = text_size+2, margin = unit(c(4,0,0,0), "mm")),
  axis.title.y = element_text(size = text_size+2, margin = unit(c(0,2,0,0), "mm")),
  strip.text = element_text(size = text_size),
  strip.background = element_blank()
) +
guides(fill = "none")

```

div\_plot



## 6. Print session info

```
sessionInfo()
```

```
## R version 4.4.2 (2024-10-31)
## Platform: x86_64-pc-linux-gnu
## Running under: Ubuntu 20.04.6 LTS
##
## Matrix products: default
## BLAS: /usr/lib/x86_64-linux-gnu/blas/libblas.so.3.9.0
## LAPACK: /usr/lib/x86_64-linux-gnu/lapack/liblapack.so.3.9.0
##
## locale:
##  [1] LC_CTYPE=en_US.UTF-8      LC_NUMERIC=C              LC_TIME=en_US.UTF-8      LC_COLLATE=en_US.UTF-8
##  [5] LC_MONETARY=en_US.UTF-8   LC_MESSAGES=en_US.UTF-8   LC_PAPER=en_US.UTF-8     LC_NAME=C
##  [9] LC_ADDRESS=C              LC_TELEPHONE=C            LC_MEASUREMENT=en_US.UTF-8 LC_IDENTIFICATION=C
##
## time zone: Etc/UTC
## tzcode source: system (glibc)
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods    base
##
## other attached packages:
##  [1] vegan_2.6-8      lattice_0.22-6  permute_0.9-7   lubridate_1.9.3 forcats_1.0.0   stringr_1.5.1
##  [8] purrr_1.0.2      readr_2.1.5     tidyr_1.3.1     tibble_3.2.1    ggplot2_3.5.1   tidyverse_2.0.0
##
## loaded via a namespace (and not attached):
##  [1] utf8_1.2.4      generics_0.1.3   stringi_1.8.4    digest_0.6.37    hms_1.1.3        magrittr_2.0.3
##  [7] evaluate_1.0.1  grid_4.4.2       timechange_0.3.0 fastmap_1.2.0    Matrix_1.7-0     tinytex_0.49
## [13] mgcv_1.9-1      fansi_1.0.6       scales_1.3.0     cli_3.6.3        rlang_1.1.4      crayon_1.5.2
## [19] bit64_4.5.2     munsell_0.5.1    splines_4.4.2    yaml_2.3.10      withr_3.0.1      tools_4.0.2
## [25] parallel_4.4.2  tzdb_0.4.0        colorspace_2.1-1 vctrs_0.6.5      R6_2.5.1          lifecycle_1.0.3
## [31] bit_4.5.0       vroom_1.6.5      MASS_7.3-61      cluster_2.1.6    pkgconfig_2.0.3   pillar_1.9.0
## [37] gtable_0.3.5    glue_1.8.0        highr_0.11       xfun_0.48         tidyselect_1.2.1  rstudioapi_0.15
## [43] knitr_1.48      farver_2.1.2      htmltools_0.5.8.1 nlme_3.1-166     labeling_0.4.3    rmarkdown_2.27
## [49] compiler_4.4.2
```