warmXtrophic Project: CN Plots

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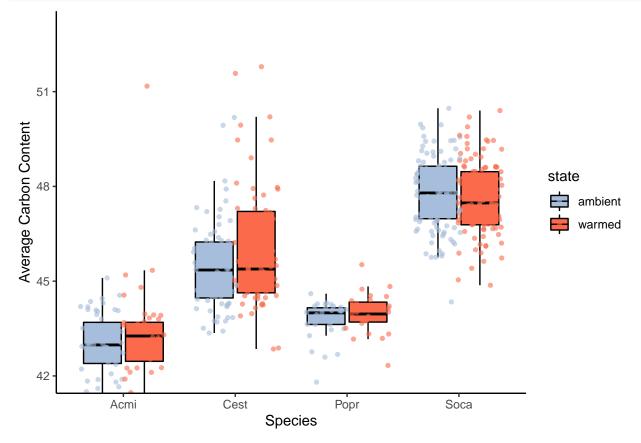
Load in data

```
# Clear all existing data
rm(list=ls())
#Load packages
library(tidyverse)
library(plotrix)
# Set working directory to Google Drive
setwd("/Volumes/GoogleDrive/Shared drives/SpaCE_Lab_warmXtrophic/data/")
# Read in data
cn <- read.csv("L1/CN/CN_L1.csv")</pre>
# Summary of data
with(cn,table(cn$site,cn$species))
##
##
          Acmi Cest Popr Soca
##
            99
                  0
                       0
                          232
    kbs
             0 138
                      58
with(cn,table(cn$year,cn$species))
##
##
          Acmi Cest Popr Soca
     2017
                          161
##
            99
                 69
     2019
                 69
                      58
with(cn,table(cn$year,cn$site))
##
##
          kbs umbs
     2017 260
     2019 71 127
```

Carbon data: all yrs, sites

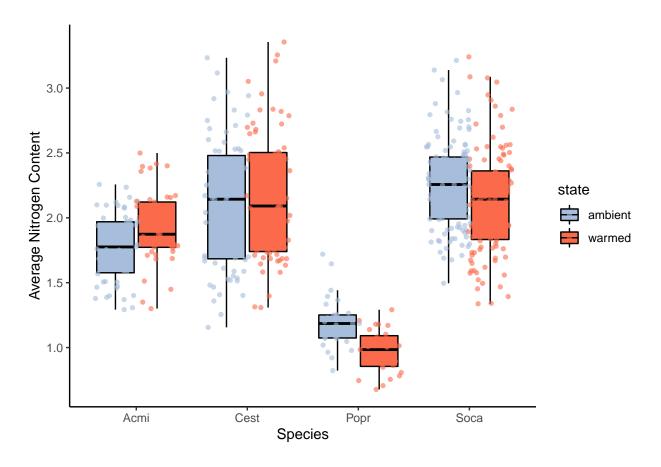
Cest and Popr = UMBS, Soca = KBS (working on faceting this)

```
ggplot(cn, aes(x = species, y = carbon, fill = state)) +
    #facet_grid(.~site) +
    geom_boxplot(color = "black", outlier.shape = NA) +
    labs(x = "Species", y = "Average Carbon Content") +
    scale_fill_manual(values = c("#a6bddb", "#fb6a4a")) +
    scale_x_discrete(labels=c("ambient" = "A", "warmed" = "W")) +
    geom_jitter(shape=16, position=position_jitterdodge(), alpha = 0.6, aes(colour = state)) +
    scale_color_manual(values = c("ambient" = "#a6bddb", "warmed" = "#fb6a4a")) +
    coord_cartesian(ylim = c(42, 53)) +
    theme_classic()
```



Nitrogen data: all yrs, sites

```
ggplot(cn, aes(x = species, y = nitrogen, fill = state)) +
    #facet_grid(.~site) +
    geom_boxplot(color = "black", outlier.shape = NA) +
    labs(x = "Species", y = "Average Nitrogen Content") +
    scale_fill_manual(values = c("#a6bddb", "#fb6a4a")) +
    scale_x_discrete(labels=c("ambient" = "A", "warmed" = "W")) +
    geom_jitter(shape=16, position=position_jitterdodge(), alpha = 0.6, aes(colour = state)) +
    scale_color_manual(values = c("ambient" = "#a6bddb", "warmed" = "#fb6a4a")) +
    theme_classic()
```



Nitrogen data: all yrs, by species - ignore the single boxplots where there should be 1 per yr (not sure why this is happening)

```
ggplot(cn, aes(x = year, y = nitrogen, fill = state)) +
    geom_boxplot(color = "black", outlier.shape = NA) +
    geom_jitter(shape=16, position=position_jitterdodge(), alpha = 0.6, aes(colour = state)) +
    facet_grid(.~species) +
    labs(x = "Year", y = "Average Nitrogen Content") +
    scale_fill_manual(values = c("#a6bddb", "#fb6a4a")) +
    #scale_x_discrete(labels=c("ambient" = "A", "warmed" = "W")) +
    scale_color_manual(values = c("ambient" = "#a6bddb", "warmed" = "#fb6a4a")) +
    facet_grid(.~species) +
    theme_classic()
```

