MAOM and POM NEON assignment

2022-10-18

R. Markdown

This first code chunk formats your data - be sure to change for your file path and your site.

```
#loading packages
library(tidyr)

## Warning: package 'tidyr' was built under R version 4.1.3

library(ggplot2)

## Warning: package 'ggplot2' was built under R version 4.1.3
```

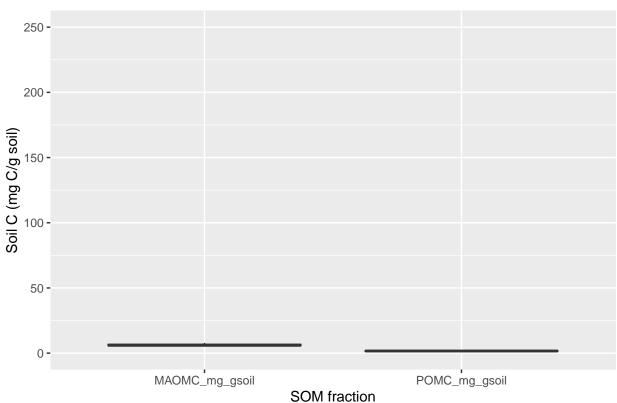
```
#formatting data
#import data
NEON_MAOMPOM <- read.csv(".../data/NEON_POMMAOMdata.csv")

#select your site and just the top soil layer (0-15cm)
N_MP_CPER <- NEON_MAOMPOM[NEON_MAOMPOM$Site== "CPER" & NEON_MAOMPOM$layer== "M1", ]</pre>
```

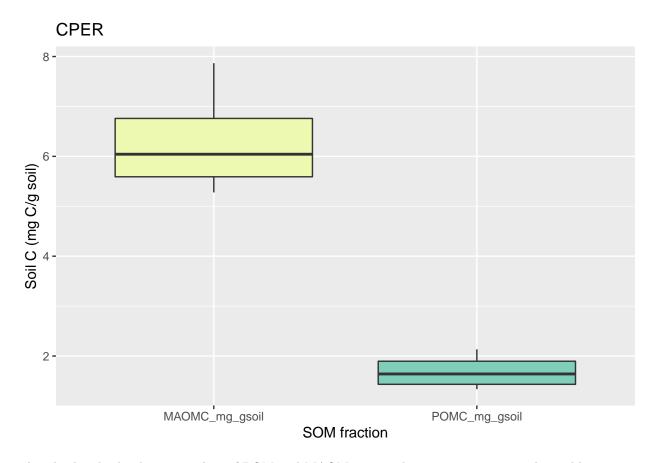
This second code chunk creates box plots with fixed axes and unfixed axes

```
#boxplot of MAOM and POM with y axis as 0-250 mg C/g soil
#rename variables
N_MP_CPER$POM <- N_MP_CPER$POMC_mg_gsoil
N_MP_CPER$MAOM <- N_MP_CPER$MAOMC_mg_gsoil
#stack to have MAOM and POM on top of one another
NMP_CPER_stacked <-gather(N_MP_CPER, "SOM_frac", "C_conc", 5:6)
#plot with specific axes
ggplot(NMP_CPER_stacked, aes(y=C_conc, x=SOM_frac, fill=SOM_frac)) +
    geom_boxplot()+ scale_fill_brewer(palette="YIGnBu") + ggtitle("CPER") + ylim(0,250) +
    xlab("SOM fraction") + ylab("Soil C (mg C/g soil)") + theme(legend.position="none")</pre>
```

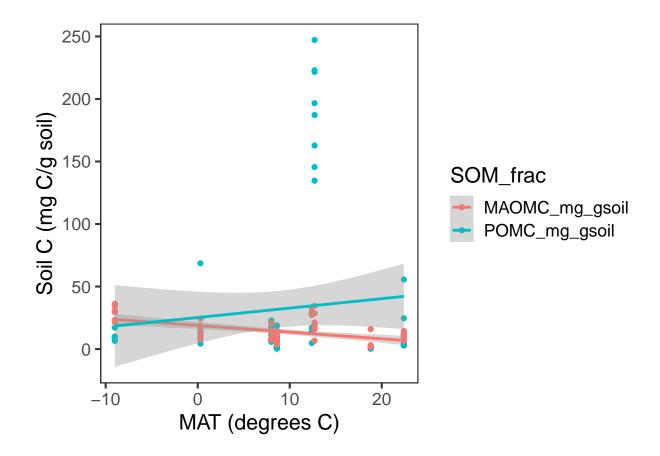


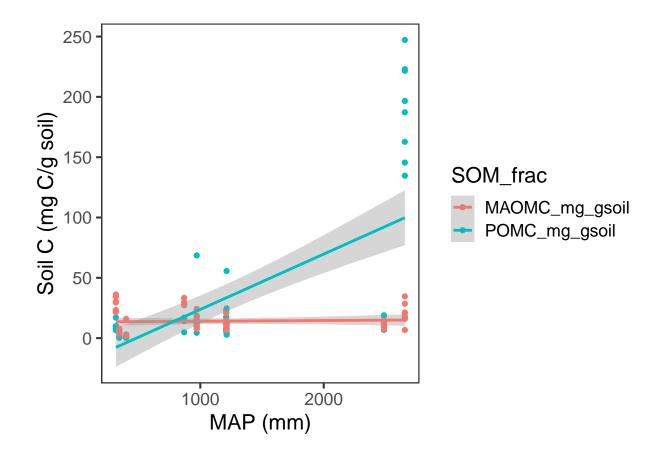


```
#plot without specific axes - remove ylim code
ggplot(NMP_CPER_stacked, aes(y=C_conc, x=SOM_frac, fill=SOM_frac)) +
  geom_boxplot()+ scale_fill_brewer(palette="YlGnBu") + ggtitle("CPER") +
  xlab("SOM fraction") + ylab("Soil C (mg C/g soil)") + theme(legend.position="none")
```



This third code chunk creates plots of POM and MAOM regressed against environmental variables





```
#initial_pH
ggplot(NMP_stacked, aes(x=initial_pH,y=C_conc, color=SOM_frac)) + geom_point() + geom_smooth(method="lm
    xlab("Soil pH") + ylab("Soil C (mg C/g soil)") +
    theme_bw(base_size = 16) + theme(panel.grid.major = element_blank(), panel.grid.minor = element_blank()
```

