

# Observed Data Prep for Model Analysis - Soils

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## Purpose

Generate BEACH soil calibration data with:

- Time.csv (Julian day marker)
- Composite soils (MonitoringScopeSoils\_R.csv)
- Detailed soils (DetailConc.csv & DetailIsotopes.csv)

```
if (MAC) {  
  if (WIN){  
    path = file.path("C:/Users/DayTimeChunks/Documents/PhD/HydrologicalMonitoring")  
  
  } else {  
    # path = file.path("/Users/DayTightChunks/Documents/PhD/HydrologicalMonitoring")  
    path = file.path("/Users/DayTightChunks/Documents/PhD/HydroMonitor/.nosync/HydrologicalMonitoring")  
    time = read.csv2("/Users/DayTightChunks/Documents/PhD/Models/.nosync/pesti-beach16/Analysis/Data/Time.csv")  
    time$DayMoYr = as.POSIXct(strptime(time$Date, "%d/%m/%Y", tz="EST"))  
  }  
} else {  
  path = file.path("D:/Documents/these_pablo/Alteckendorf2016/HydrologicalMonitoring")  
  time = read.csv2("D:/Documents/these_pablo/Models/BEACH2016/Analysis/Data/Time.csv")  
  time$DayMoYr = as.POSIXct(strptime(time$Date, "%d/%m/%Y", tz="EST"))  
}  
  
# Lab parameters and field constants  
source(file.path(path, "global.R"))
```

## Packages

```
# Plotting functions  
library("scales")  
library("tidyr")  
  
library("reshape")  
library("zoo") # na.approx()  
  
library("plyr")  
library("dplyr")
```

## Working directory

```
# setwd("D:/Documents/these_pablo/Alteckendorf2016/R")  
  
# MAC
```

```
# setwd("/Users/DayTightChunks/Documents/PhD/Routput/Alteck/R")

# Mac-WIN
# setwd("C:/Users/DayTightChunks/Documents/Models/pesti-beach16/Analysis/Data")
getwd()

## [1] "/Users/DayTightChunks/Documents/PhD/Models/.nosync/pesti-beach16/Analysis/Data"
```

## Merge Transects with Time (Julian Days)

```
s = read.csv2(file.path(path, "Data/MonitoringScopeSoils_R.csv"))

#q$date = as.POSIXct(strptime(q$dateCheck, "%d/%m/%Y %H:%M", tz="EST"))
s$DayMoYr = as.POSIXct(strptime(s$date.Soil, "%d/%m/%Y", tz="EST"))

north = subset(s, Transect == "N")
valley = subset(s, Transect == "T")
south = subset(s, Transect == "S")

njd = merge(time, north, by = "DayMoYr", all = T)
vjd = merge(time, valley, by = "DayMoYr", all = T)
sjd = merge(time, south, by = "DayMoYr", all = T)

SAVE = F
if (SAVE) {
  write.table(njd, "BEACH_R/north.tss", sep="\t", row.names = F)
  write.table(vjd, "BEACH_R/valley.tss", sep="\t", row.names = F)
  write.table(sjd, "BEACH_R/south.tss", sep="\t", row.names = F)
}
```

## Merge detail concentration and isotopes

### Steps

- group isotope values by ID
- evaluate standard deviations
- compute means
- merge with concentrations

```
co = read.csv("DetailConc.csv")
is = read.csv("DetailIsotopes.csv")
```

### Group

```
# Same procedure as in Book 06_MassSoils_Composite
siso <- is[, c("ID", "d13C")]
sumIS = ddply(siso, c("ID"),
  summarise,
  N_detsoil = length(d13C),
```

```

det.d13C = mean(d13C),
det.d13C.SD = sd(d13C)
)

```

## Merge conc and isotopes

```

ci = merge(co, sumIS, by = "ID", all = T)
ci$DayMoYr = as.POSIXct(strptime(ci$Sampled, "%d/%m/%Y", tz="EST"))
ci = subset(ci, !is.na(ci$DayMoYr))

ci$Plot = paste(ci$Transect, ci$Plot, sep = "-")
ci = ci[, c("DayMoYr", "ID", "Plot", "ug.g", "det.d13C", "det.d13C.SD")]
detailed = merge(time, ci, by = "DayMoYr", all = T)

```

## Create plot subsets

```

N1 = subset(detailed, is.na(Plot) | Plot == "N-1")
N2 = subset(detailed, is.na(Plot) | Plot == "N-2")
N3 = subset(detailed, is.na(Plot) | Plot == "N-3")
N4 = subset(detailed, is.na(Plot) | Plot == "N-4")
N5 = subset(detailed, is.na(Plot) | Plot == "N-5")
N7 = subset(detailed, is.na(Plot) | Plot == "N-7")
N8 = subset(detailed, is.na(Plot) | Plot == "N-8")

T4 = subset(detailed, is.na(Plot) | Plot == "T-4")
T5 = subset(detailed, is.na(Plot) | Plot == "T-5")
T7 = subset(detailed, is.na(Plot) | Plot == "T-7")
T8 = subset(detailed, is.na(Plot) | Plot == "T-8")
T9 = subset(detailed, is.na(Plot) | Plot == "T-9")
T10 = subset(detailed, is.na(Plot) | Plot == "T-10")

S11 = subset(detailed, is.na(Plot) | Plot == "S-11")
S12 = subset(detailed, is.na(Plot) | Plot == "S-12")
S13 = subset(detailed, is.na(Plot) | Plot == "S-13")

nor = list(N1, N2, N3, N4, N5, N7, N8)
val = list(T4, T5, T7, T8, T9, T10)
sou = list(S11, S12, S13)

```

## Save

```

SAVE = F

if (SAVE) {
  counter = 0
  for (d in nor) {
    counter = counter + 1
    transect = "N"
  }
}

```

```

    if (counter < 6) {
      name = paste(transect, as.character(counter), ".tss", sep = "")
    }
    else if (counter == 6) {
      counter = counter + 1
      name = paste(transect, as.character(counter), ".tss", sep = "")
    } else {
      name = paste(transect, as.character(counter), ".tss", sep = "")
    }
    write.table(d, paste("BEACH_R/", name, sep = ""), sep="\t", row.names = F)
  }

  counter = 3
  for (d in val) {
    counter = counter + 1
    transect = "T"
    if (counter < 6) {
      name = paste(transect, as.character(counter), ".tss", sep = "")
    }
    else if (counter == 6) {
      counter = counter + 1
      name = paste(transect, as.character(counter), ".tss", sep = "")
    } else {
      name = paste(transect, as.character(counter), ".tss", sep = "")
    }
    write.table(d, paste("BEACH_R/", name, sep = ""), sep="\t", row.names = F)
  }

  counter = 10
  for (d in sou) {
    counter = counter + 1
    transect = "S"
    name = paste(transect, as.character(counter), ".tss", sep = "")
    write.table(d, paste("BEACH_R/", name, sep = ""), sep="\t", row.names = F)
  }
}

```