Soils & Discharge Merged

PAZ

17 novembre 2016

Purpose

This file merges outlet data with soil data to plot cumulative exported and remaining S-metolachlor mass. The nearest soil sample date for each transect is used to match the initial time ("ti") of the sampling discharge period. This is most adequate merging location given that samples took place shortly before relaunching the automatic sampler.

Note that week numbers for water and soils are offset by one. I.e. Week 1 soils influence/regulate Week 2's water sample results.

Imports:

- **WeeklyHydroContam_R.csv**
- **WeeklySoils_R.csv**

Generates:

- **WeekSoilHydroCont_R.csv**

Required R-packages:

```
library("plyr")
library("dplyr")
```

Working directory

```
# setwd("D:/Documents/these_pablo/Alteckendorf2016/R")
# setwd("/Users/DayTightChunks/Documents/PhD/Routput/Alteck/R")
# setwd("D:/Documents/these_pablo/Alteckendorf2016/00_TransparencyFolder")
getwd()
```

```
## [1] "D:/Documents/these_pablo/Alteckendorf2016/HydrologicalMonitoring"
```

Import files

```
outlet = read.csv2("Data/WeeklyHydroContam_R.csv", header = T)
outlet$ti <- as.POSIXct(outlet$ti, "%Y-%m-%d %H:%M", tz = "EST")
sum(is.na(outlet$ti))
```

```
## [1] 0
```

```
# Select variables from Water dataset
outlet <- outlet[, c("ti", "WeekSubWeek", "Event", "B.diss", "B.filt", "CumOutDiss.g", "CumOutFilt.g",
print("Water")
```

```
## [1] "Water"
```

```r
str(outlet)
```

```
## 'data.frame':    51 obs. of  9 variables:
##  $ ti           : POSIXct, format: "2016-03-25 00:04:00" "2016-03-25 12:04:00" ...
##  $ WeekSubWeek  : Factor w/ 51 levels "W0-0x","W0-1",..: 1 2 3 4 5 6 26 27 28 29 ...
##  $ Event        : int  0 0 0 1 1 1 2 2 2 3 ...
##  $ B.diss       : num  NA 93.1 NA 35.4 29.4 ...
##  $ B.filt       : num  NA NA NA NA NA ...
##  $ CumOutDiss.g : num  0.00354 0.02815 0.19818 2.84809 5.2051 ...
##  $ CumOutFilt.g : num  0.00345 0.0069 0.01263 0.01571 0.01923 ...
##  $ CumAppMass.g : num  6369 6369 6369 6369 6369 ...
##  $ CumOutMELsm.g: num  0.302 2.38 4.76 35.001 62.009 ...
```

```r
soils = read.csv2("Data/WeeklySoils_Rng.csv", header =T) # Corrected with only ngC > 2.0
soils$Date.ti <- as.POSIXct(soils$Date.ti, "%Y-%m-%d %H:%M", tz = "EST")
#soils$Date.ti <- as.POSIXct(soils$Date.ti, "%d/%m/%Y %H:%M", tz = "EST")
sum(is.na(soils$Date.ti))
```

```
## [1] 0
```

```r
print("Soils")
```

```
## [1] "Soils"
```

```r
str(soils)
```

```
## 'data.frame':    51 obs. of  26 variables:
##  $ ID               : Factor w/ 51 levels "AW-N-0","AW-N-0x",..: 2 19 36 1 18 35 3 20 37 10 ...
##  $ Transect         : Factor w/ 3 levels "N","S","T": 1 2 3 1 2 3 1 2 3 1 ...
##  $ Wnum             : int  -1 -1 -1 0 0 0 1 1 1 2 ...
##  $ Date.Soil        : Factor w/ 17 levels "03/05/2016 13:10",..: 13 13 13 16 16 16 3 3 3 7 ...
##  $ Date.ti          : POSIXct, format: "2016-03-25 00:04:00" "2016-03-25 00:04:00" ...
##  $ Conc.mug.g.dry.soil: num  0.0183 0.0285 0.0205 0.8893 3.204 ...
##  $ Conc.ComSoil.SD  : num  NA NA NA 1.46 2.77 ...
##  $ N_compsoil       : int  NA NA NA NA NA NA NA 2 NA 3 ...
##  $ comp.d13C        : num  NA NA NA NA NA ...
##  $ comp.d13C.SD     : num  NA NA NA NA NA ...
##  $ comp.d13C.SE     : num  NA NA NA NA NA ...
##  $ N_ngC            : int  NA NA NA NA NA NA NA 2 NA 3 ...
##  $ ngC.mean         : num  NA NA NA NA NA ...
##  $ ngC.SD           : num  NA NA NA NA NA ...
##  $ ngC.SE           : num  NA NA NA NA NA ...
##  $ DD13C.comp       : num  NA NA NA NA NA ...
##  $ f.max.comp       : num  NA NA NA NA NA ...
##  $ B.max.comp       : num  NA NA NA NA NA ...
##  $ f.min.comp       : num  NA NA NA NA NA ...
##  $ B.min.comp       : num  NA NA NA NA NA ...
##  $ f.mean.comp      : num  NA NA NA NA NA ...
##  $ B.mean.comp      : num  NA NA NA NA NA ...
##  $ MassSoil.g       : num  12.61 18.8 4.44 613.08 2112.12 ...
##  $ Area.N           : num  139266 139266 139266 139266 139266 ...
##  $ Area.T           : num  43713 43713 43713 43713 43713 ...
##  $ Area.S           : num  133175 133175 133175 133175 133175 ...
```

Get soil concentrations for each transect for merging horizontally

```r
# North
soils.N <- subset(soils, soils$Transect == "N")
soils.N <- soils.N[, c("Date.ti", "B.mean.comp",  "B.max.comp",  "B.min.comp",  "MassSoil.g",
                       "comp.d13C", "comp.d13C.SD", "comp.d13C.SE",
                       "ID", "Area.N", "Area.T", "Area.S" )]

colnames(soils.N) <- c("ti",  "B.mean.comp.North", "B.max.comp.North", "B.min.comp.North", "MassSoil.g.
                       "comp.d13C.North", "comp.d13C.SD.North", "comp.d13C.SE.North",
                       "ID.N", "Area.N", "Area.T", "Area.S" )

# Talweg
soils.T <- subset(soils, soils$Transect == "T")
soils.T <- soils.T[, c("Date.ti", "B.mean.comp",  "B.max.comp",  "B.min.comp", "MassSoil.g",
                       "comp.d13C", "comp.d13C.SD", "comp.d13C.SE",
                       "ID")]

colnames(soils.T) <- c("ti",  "B.mean.comp.Talweg", "B.max.comp.Talweg", "B.min.comp.Talweg", "MassSoil
                       "comp.d13C.Talweg", "comp.d13C.SD.Talweg", "comp.d13C.SE.Talweg",
                       "ID.T" )

# South
soils.S <- subset(soils, soils$Transect == "S")
soils.S <- soils.S[, c("Date.ti", "B.mean.comp",  "B.max.comp",  "B.min.comp", "MassSoil.g",
                       "comp.d13C", "comp.d13C.SD", "comp.d13C.SE",
                       "ID" )]
colnames(soils.S) <- c("ti",  "B.mean.comp.South", "B.max.comp.South", "B.min.comp.South", "MassSoil.g.S
                       "comp.d13C.South", "comp.d13C.SD.South", "comp.d13C.SE.South",
                       "ID.S" )
```

Total Catchment Mass, Bulk Mass and Bulk Isotopes

Bulk isotopes are calculated based on the following:

$$\delta_{bulk} = \frac{M_{North}}{M_{tot}}\delta_{North} + \frac{M_{Talweg}}{M_{tot}}\delta_{Talweg} + \frac{M_{South}}{M_{tot}}\delta_{South}$$

Bulk mass is a proxy for the potential mass that can be discharged at a give time. It is calculated base don the pondered mass from each transect area and the proportion of that area in relation to the entire catchment such that:

$$M_{bulk} = \frac{A_{North}}{A_{tot}}M_{North} + \frac{A_{Talweg}}{A_{tot}}M_{Talweg} + \frac{A_{South}}{A_{tot}}M_{South}$$

```r
library(zoo)
```

```
##
## Attaching package: 'zoo'

## The following objects are masked from 'package:base':
##
##     as.Date, as.Date.numeric
```

```r
class(outlet$ti)
```

```
## [1] "POSIXct" "POSIXt"
```

```r
class(soils.T$ti)
```

```
## [1] "POSIXct" "POSIXt"
```

```r
soilsOut <- merge(outlet, soils.N, by = "ti", all = T)
soilsOut <- merge(soilsOut, soils.T, by = "ti", all = T)
soilsOut <- merge(soilsOut, soils.S, by = "ti", all = T)

# Total mass in catchment
soilsOut$CatchMassSoil.g <-
  soilsOut$MassSoil.g.North +
  soilsOut$MassSoil.g.Talweg +
  soilsOut$MassSoil.g.South

# Approximate initial mass for first 6 observation (before recording composite soils)
#soilsOut$CatchMassSoil.g[1:6] <- ifelse(is.na(soilsOut$CatchMassSoil.g),
#                                        soilsOut$CumAppMass.g-soilsOut$CumOutMELsm.g,
#                                        soilsOut$CatchMassSoil.g)



soilsOut$BulkMass.g <-
  soilsOut$MassSoil.g.North*(soilsOut$Area.N/(soilsOut$Area.N+soilsOut$Area.T+soilsOut$Area.S)) +
  soilsOut$MassSoil.g.Talweg*(soilsOut$Area.T/(soilsOut$Area.N+soilsOut$Area.T+soilsOut$Area.S)) +
  soilsOut$MassSoil.g.South*(soilsOut$Area.S/(soilsOut$Area.N+soilsOut$Area.T+soilsOut$Area.S))

# Bulk catchment isotopes
soilsOut$BulkCatch.d13 <-
  (soilsOut$MassSoil.g.North/soilsOut$CatchMassSoil.g)*soilsOut$comp.d13C.North +
  (soilsOut$MassSoil.g.Talweg/soilsOut$CatchMassSoil.g)*soilsOut$comp.d13C.Talweg +
  (soilsOut$MassSoil.g.South/soilsOut$CatchMassSoil.g)*soilsOut$comp.d13C.South

print("Merged Soils and Outlet by ti")
```

```
## [1] "Merged Soils and Outlet by ti"
```

```r
str(soilsOut)
```

```
## 'data.frame':    52 obs. of  39 variables:
##  $ ti                : POSIXct, format: "2016-03-25 00:04:00" "2016-03-25 12:04:00" ...
##  $ WeekSubWeek       : Factor w/ 51 levels "W0-0x","W0-1",..: 1 2 3 4 5 6 26 27 28 29 ...
##  $ Event             : int  0 0 0 1 1 1 2 2 2 3 ...
##  $ B.diss            : num  NA 93.1 NA 35.4 29.4 ...
##  $ B.filt            : num  NA NA NA NA NA ...
##  $ CumOutDiss.g      : num  0.00354 0.02815 0.19818 2.84809 5.2051 ...
##  $ CumOutFilt.g      : num  0.00345 0.0069 0.01263 0.01571 0.01923 ...
##  $ CumAppMass.g      : num  6369 6369 6369 6369 6369 ...
##  $ CumOutMELsm.g     : num  0.302 2.38 4.76 35.001 62.009 ...
##  $ B.mean.comp.North : num  NA NA NA NA NA ...
##  $ B.max.comp.North  : num  NA NA NA NA NA ...
##  $ B.min.comp.North  : num  NA NA NA NA NA ...
##  $ MassSoil.g.North  : num  12.6 NA NA 613.1 NA ...
```

4

```
## $ comp.d13C.North    : num  NA NA NA NA NA ...
## $ comp.d13C.SD.North : num  NA NA NA NA NA ...
## $ comp.d13C.SE.North : num  NA NA NA NA NA ...
## $ ID.N               : Factor w/ 51 levels "AW-N-0","AW-N-0x",..: 2 NA NA 1 NA NA 3 NA NA 10 ...
## $ Area.N             : num  139266 NA NA 139266 NA ...
## $ Area.T             : num  43713 NA NA 43713 NA ...
## $ Area.S             : num  133175 NA NA 133175 NA ...
## $ B.mean.comp.Talweg : num  NA NA NA NA NA ...
## $ B.max.comp.Talweg  : num  NA NA NA NA NA ...
## $ B.min.comp.Talweg  : num  NA NA NA NA NA ...
## $ MassSoil.g.Talweg  : num  4.44 NA NA 173.27 NA ...
## $ comp.d13C.Talweg   : num  NA NA NA NA NA ...
## $ comp.d13C.SD.Talweg: num  NA NA NA NA NA ...
## $ comp.d13C.SE.Talweg: num  NA NA NA NA NA ...
## $ ID.T               : Factor w/ 51 levels "AW-N-0","AW-N-0x",..: 36 NA NA 35 NA NA 37 NA NA 44 ...
## $ B.mean.comp.South  : num  NA NA NA NA NA ...
## $ B.max.comp.South   : num  NA NA NA NA NA ...
## $ B.min.comp.South   : num  NA NA NA NA NA ...
## $ MassSoil.g.South   : num  18.8 NA NA 2112.1 NA ...
## $ comp.d13C.South    : num  NA NA NA NA NA ...
## $ comp.d13C.SD.South : num  NA NA NA NA NA ...
## $ comp.d13C.SE.South : num  NA NA NA NA NA ...
## $ ID.S               : Factor w/ 51 levels "AW-N-0","AW-N-0x",..: 19 NA NA 18 NA NA 20 NA NA 27 ...
## $ CatchMassSoil.g    : num  35.8 NA NA 2898.5 NA ...
## $ BulkMass.g         : num  14.1 NA NA 1183.7 NA ...
## $ BulkCatch.d13      : num  NA NA NA NA NA ...
```

```r
write.csv2(soilsOut,
           'Data/MassBalance_R.csv', row.names = F)
```

Plot

```r
library("ggplot2")
library("scales")
library("reshape2")
library("cowplot")
```

```
##
## Attaching package: 'cowplot'

## The following object is masked from 'package:ggplot2':
##
##     ggsave
```

```r
# Melt data set
##Subset the necessary columns
soilsRemainMass <- soilsOut[, c("ti" ,"CumAppMass.g", "CumOutDiss.g", "CumOutFilt.g", "CumOutMELsm.g" ,
soilsRemainMass$CumAppMass.g[1:3] <- NA

# Replace Catchment Mass's NA with the most recent non-NA prior to it (assumes no degradation).
# Purpose: To match continuous outlet time array
soilsRemainMass$CatchMassSoil.g <- na.locf(soilsRemainMass$CatchMassSoil.g)
# soilsRemainMass <- na.omit(soilsRemainMass)
```

```r
##Then rearrange your data frame
remainMassMolten = melt(soilsRemainMass, id=c("ti"))

# View(remainMassMolten)

pg <- remainMassMolten

# Change variable names:
levels(pg$variable)[levels(pg$variable)=="CumAppMass.g"] <- "Applied SM Cum. (Survey)"
levels(pg$variable)[levels(pg$variable)=="CumOutMELsm.g"] <- "MEL-SM Cum. (Outlet)"
levels(pg$variable)[levels(pg$variable)=="CatchMassSoil.g"] <- "Remaining SM (1cm Soil)"

levels(pg$variable)[levels(pg$variable)=="CumOutDiss.g"] <- "Dissolved SM Cum. (Outlet)"
levels(pg$variable)[levels(pg$variable)=="CumOutFilt.g"] <- "Sediment SM Cum. (Outlet)"

# Change the order:
levels(pg$variable)
```

```
## [1] "Applied SM Cum. (Survey)"   "Dissolved SM Cum. (Outlet)"
## [3] "Sediment SM Cum. (Outlet)"  "MEL-SM Cum. (Outlet)"
## [5] "Remaining SM (1cm Soil)"
```

```r
pg$variable <- factor(pg$variable, levels = c("Applied SM Cum. (Survey)",  "Remaining SM (1cm Soil)", "

pgSimple <- pg[which(pg$variable != ("Dissolved SM Cum. (Outlet)") & pg$variable != ("Sediment SM Cum.
# names(pg)[names(pg)=="variable"]  <- "Estimated Mass"

massBalTop <- ggplot(pg) +
  geom_line(aes(x=ti, y=value, group = variable, color=variable)) +

  # Themes and axes
  theme_bw() +
  theme(# axis.text.x=element_text(angle = 45, hjust = 1),
        axis.text.x=element_blank(),
        axis.title.x=element_blank(),
        legend.position="top"

        )+
  labs(color = "Estimated Mass") +
  guides(col = guide_legend(ncol = 1)) +  # Sets legend parameters

  # xlab("Date") +
  scale_x_datetime(breaks = date_breaks("2 weeks"), labels = date_format("%b %d")) +
  ylab(expression(paste("Mass ", {(g)}))) +
  # scale_y_continuous(breaks = c(100, 5000, 10000, 20000), limits = c(100, 20000) )
  scale_y_continuous(trans=log_trans(), breaks=c(1,5,10,50,100,500,1000,5000, 10000))
massBalTop
```
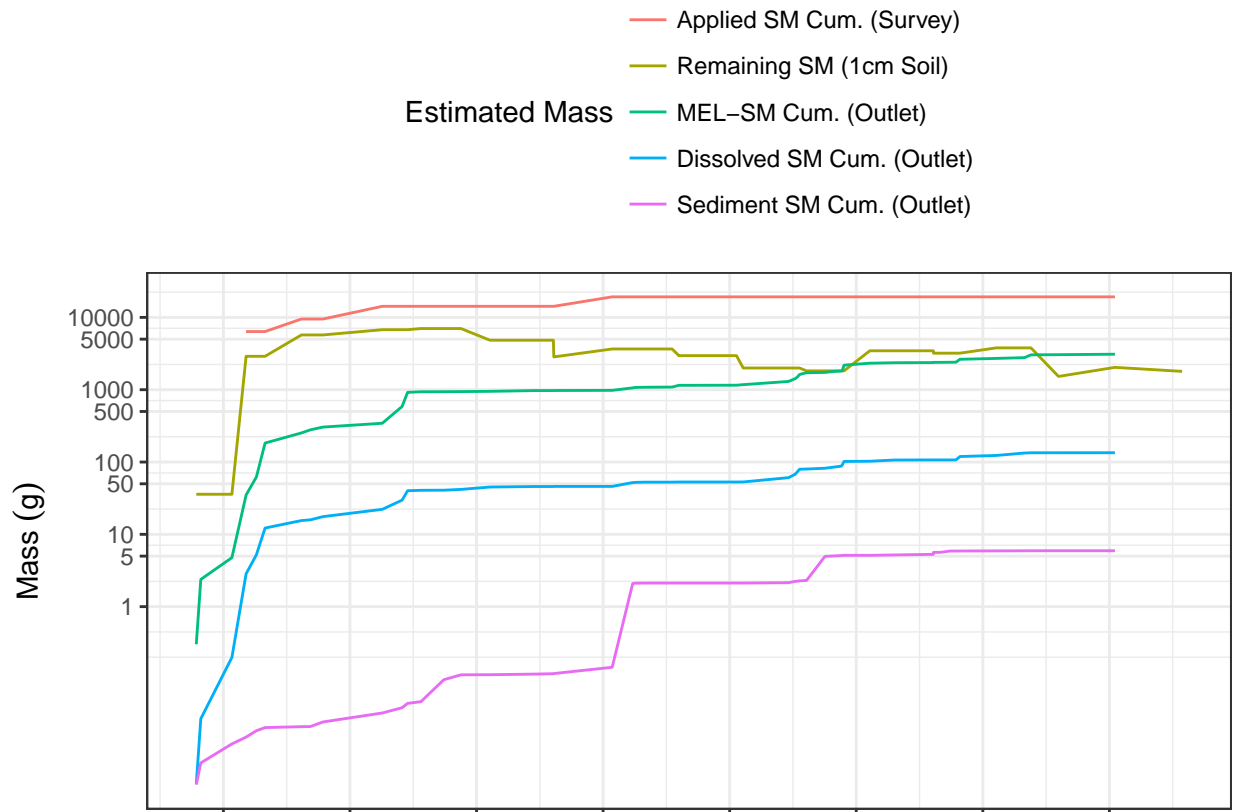
```
## Warning: Removed 7 rows containing missing values (geom_path).
```

```
massBalBottom <- ggplot(pg) +
  geom_line(aes(x=ti, y=value, color=variable)) +

  # Themes and axes
  theme_bw() +
  theme(axis.text.x=element_text(angle = 45, hjust = 1),
        #axis.text.x=element_blank(),
        #axis.title.x=element_blank(),
        legend.position="none"
        )+
  # guides(col = guide_legend(nrows = 2)) +  # Sets legend parameters
  xlab("Date") +
  scale_x_datetime(breaks = date_breaks("2 weeks"), labels = date_format("%b %d")) +
  ylab(expression(paste("Mass. S-Meto. ", {(g)}))) +
  scale_y_continuous(breaks = c(1, 25, 50, 100), limits = c(0, 100) )


massBal = plot_grid(massBalTop, massBalBottom, ncol = 1, nrow = 2, align = "v")
```

```
## Warning: Removed 7 rows containing missing values (geom_path).
```

```
## Warning: Removed 112 rows containing missing values (geom_path).
```

```
massBal_MEL <- ggplot(pgSimple) +
  geom_line(aes(x=ti, y=value, group = variable, color=variable)) +

  # Themes and axes
```

```
    theme_bw() +
    theme(# axis.text.x=element_text(angle = 45, hjust = 1),
          axis.text.x=element_blank(),
          axis.title.x=element_blank(),
          legend.position="top"

          )+
    labs(color = "Estimated Mass") +
    guides(col = guide_legend(ncol = 3)) +  # Sets legend parameters

    # xlab("Date") +
    scale_x_datetime(breaks = date_breaks("2 weeks"), labels = date_format("%b %d")) +
    ylab(expression(paste("Mass ", {(g)})))

massBal_MEL
```

Warning: Removed 5 rows containing missing values (geom_path).