eda-final.R

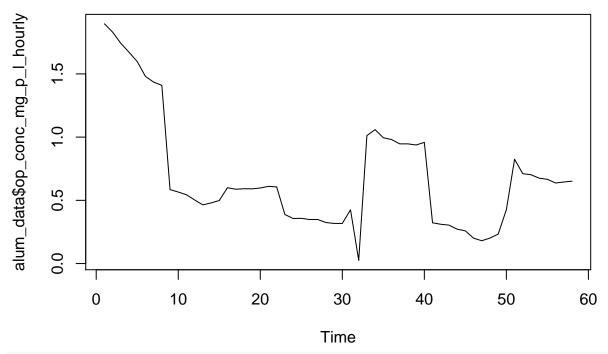
niko

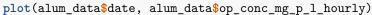
2020-06-09

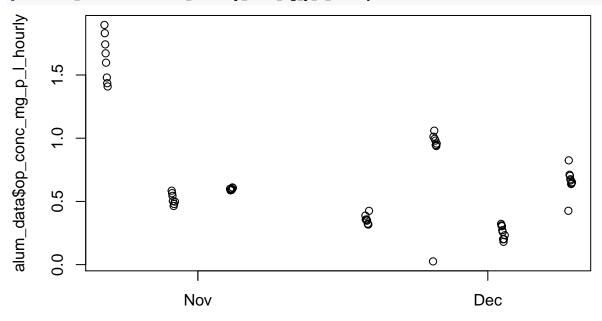
```
rm( list = ls() )
library(tidyverse)
## -- Attaching packages ----
## v ggplot2 3.3.0
                     v purrr
                               0.3.3
## v tibble 2.1.3
                     v dplyr
                               0.8.5
## v tidyr
           1.0.2
                    v stringr 1.4.0
## v readr
           1.3.1
                     v forcats 0.5.0
## -- Conflicts ------ tidyvers
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
library(fields)
## Loading required package: spam
## Loading required package: dotCall64
## Loading required package: grid
## Spam version 2.5-1 (2019-12-12) is loaded.
## Type 'help( Spam)' or 'demo( spam)' for a short introduction
## and overview of this package.
## Help for individual functions is also obtained by adding the
## suffix '.spam' to the function name, e.g. 'help( chol.spam)'.
##
## Attaching package: 'spam'
## The following objects are masked from 'package:base':
##
##
      backsolve, forwardsolve
## Loading required package: maps
## Attaching package: 'maps'
## The following object is masked from 'package:purrr':
##
##
      map
## See https://github.com/NCAR/Fields for
```

an extensive vignette, other supplements and source code

```
library(lubridate)
## Attaching package: 'lubridate'
## The following object is masked from 'package:base':
##
       date
library(latex2exp)
library(reshape2)
## Attaching package: 'reshape2'
## The following object is masked from 'package:tidyr':
##
##
       smiths
# make sure src is the current working directory
load("../data/final-data.rda")
load("../data/boulderMoWater.rda")
# display the number of observations for each type of coagulant
table(final_data$coagulant)
##
##
            Alum Ferric
                          None
       0
             58
                     45
# splitting the data by coagulant
ferr_data <- final_data %>% filter(coagulant == "Ferric")
alum_data <- final_data %>% filter(coagulant == "Alum")
none_data <- final_data %>% filter(coagulant == "None")
# average was found by hour maybe do a moving average instead?
ts.plot(alum_data$op_conc_mg_p_l_hourly)
```



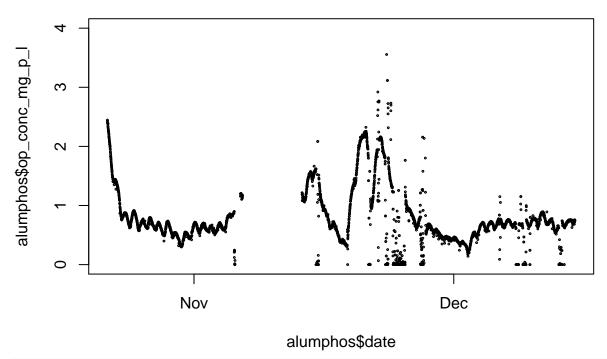




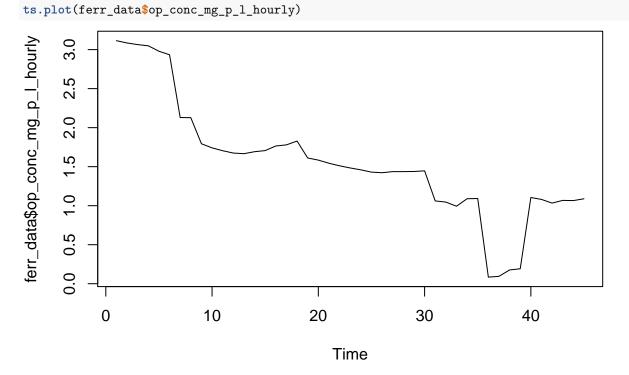
alum_data\$date

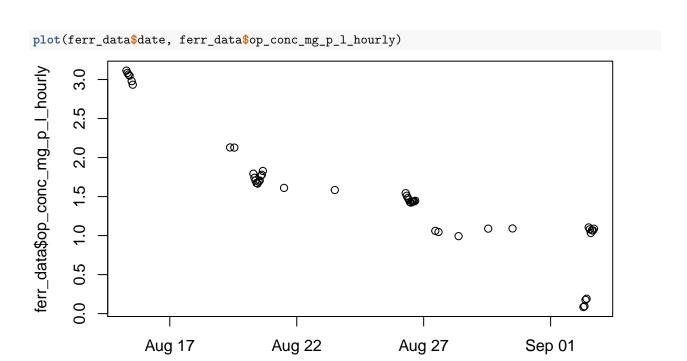
```
# gap in november looking at this variable at original time scale no average
alumphos <- phosfax_10m %>%
  filter(date >= ymd("2019-10-22")) %>%
  filter(date <= ymd("2019-12-15"))
plot(alumphos$date, alumphos$op_conc_mg_p_l,
    cex = .25,
    main = "Alum effluent OP",
    ylim = c(0,4))</pre>
```

Alum effluent OP



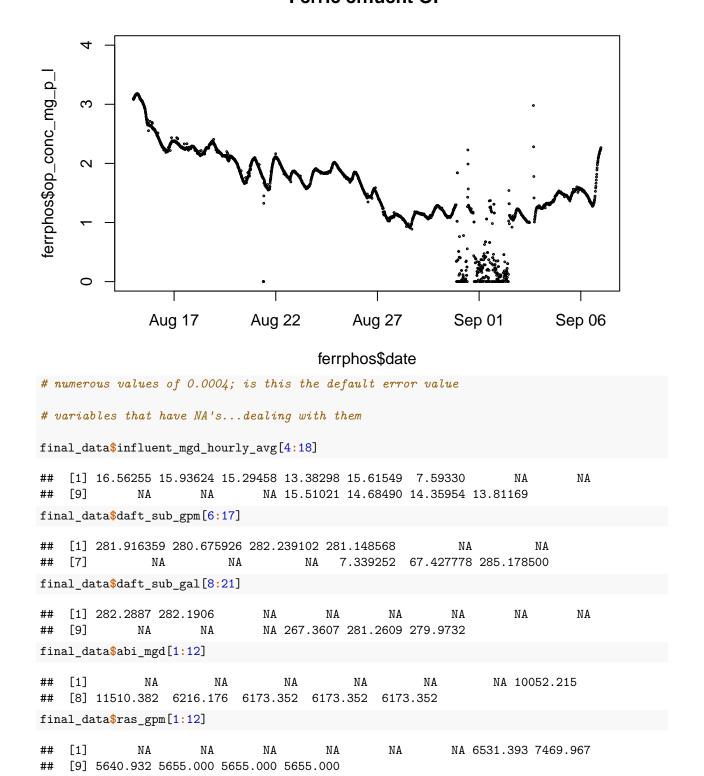
comparing this to above we see 11/18 - 11/25 our merged data doesnt exist. a lot of outliers in this
what occurred between 08/28 - 09/02





ferr_data\$date

Ferric effluent OP



NA 12.80994 67.42778 295.33417 295.47741

NA

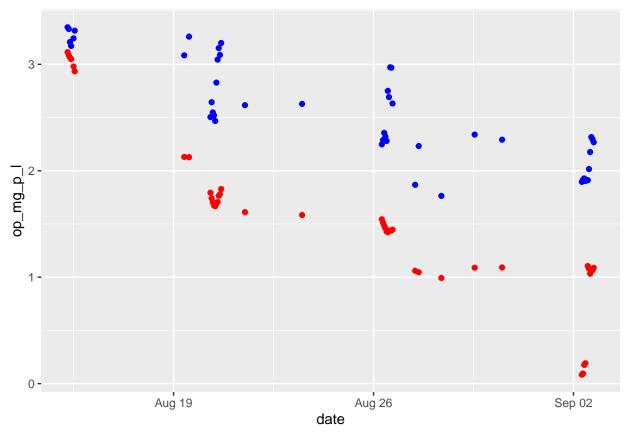
NA

final_data\$mlws_flow_gpm[7:18]

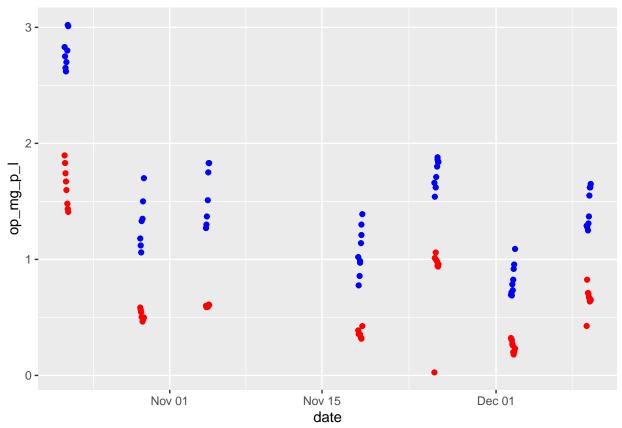
[1] 294.45824 294.68749 295.53401

```
# remove? fill with average, moving average?

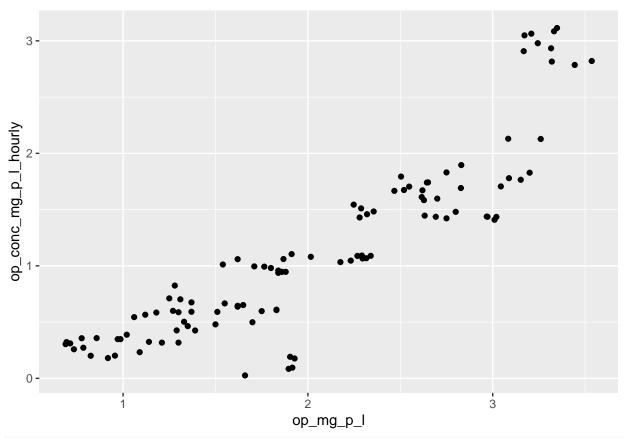
ferr_data %>% ggplot() +
   geom_point(aes(date, op_mg_p_l), col = "blue") +
   geom_point(aes(date, op_conc_mg_p_l_hourly), col = "red")
```



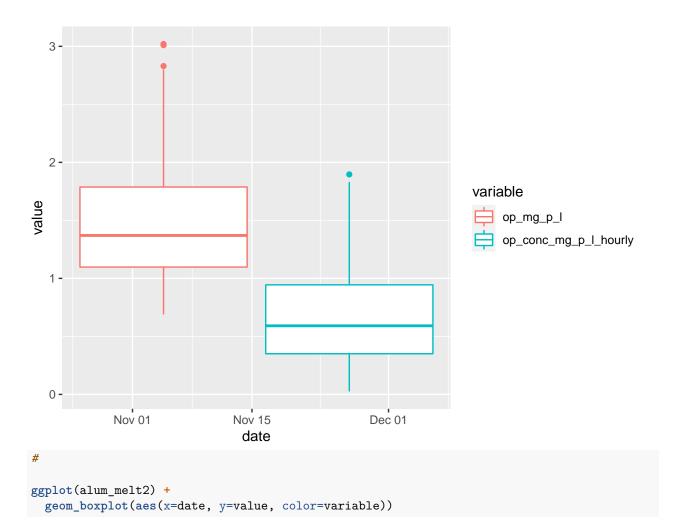
```
alum_data %>% ggplot() +
  geom_point(aes(date, op_mg_p_1), col = "blue") +
  geom_point(aes(date, op_conc_mg_p_1_hourly), col = "red")
```

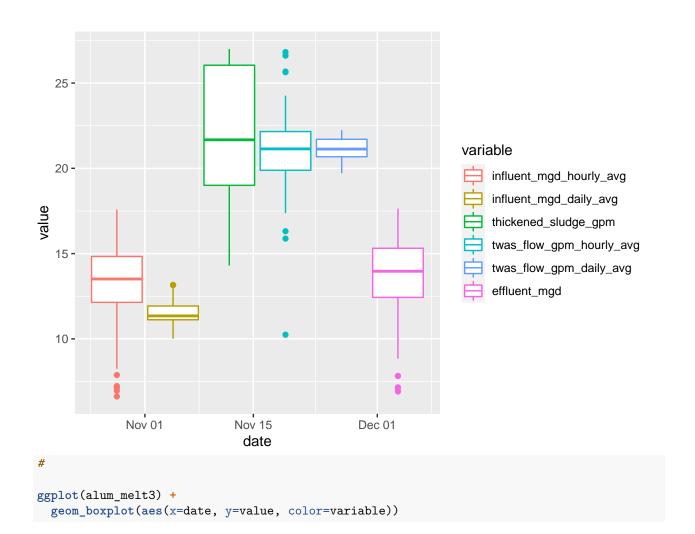


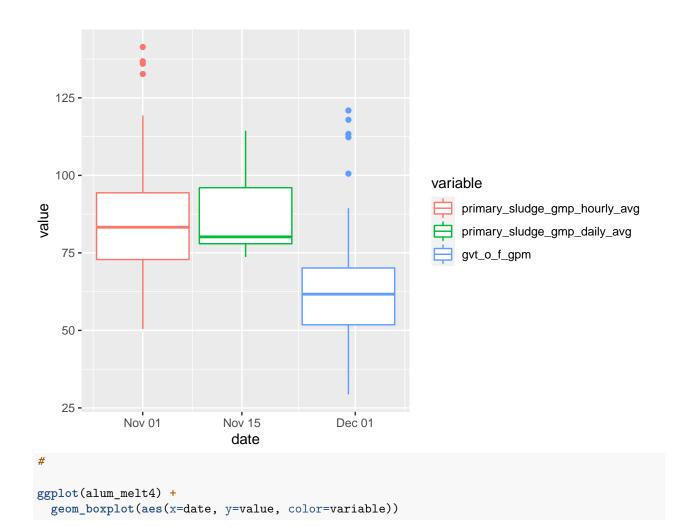
shows linear relationship shifted from effluent being less due to dosing
final_data %>% ggplot() +
 geom_point(aes(op_mg_p_l, op_conc_mg_p_l_hourly))

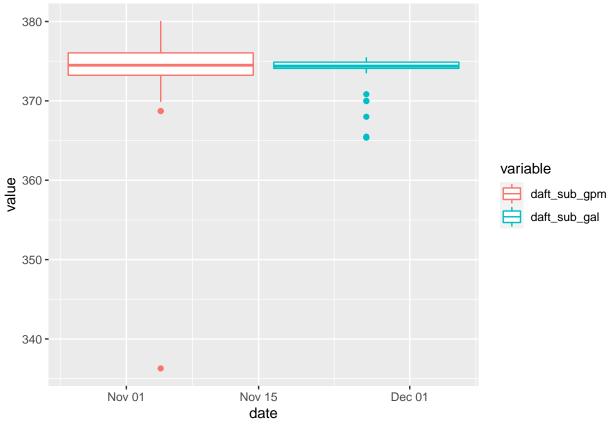


```
alum_melt1 <- melt(alum_data[,c(1,3:4)],id.vars='date', measure.vars=colnames(alum_data[,c(1,3:4)])[-1]
alum_melt2 <- melt(alum_data[,c(1,5:6,10,19:20,22)],id.vars='date', measure.vars=colnames(alum_data[,c(alum_melt3)],id.vars='date', measure.vars=colnames(alum_data[,c(1,7:8,12)])
alum_melt3 <- melt(alum_data[,c(1,14:15)],id.vars='date', measure.vars=colnames(alum_data[,c(1,14:15)])
alum_melt5 <- melt(alum_data[,c(1,18)],id.vars='date', measure.vars=colnames(alum_data[,c(1,18)])[-1])
alum_melt6 <- melt(alum_data[,c(1,21)],id.vars='date', measure.vars=colnames(alum_data[,c(1,21)])[-1])
alum_melt7 <- melt(alum_data[,c(1,9,11,13)],id.vars='date', measure.vars=colnames(alum_data[,c(1,9,11,13)])
##
ggplot(alum_melt1) +
geom_boxplot(aes(x=date, y=value, color=variable))</pre>
```



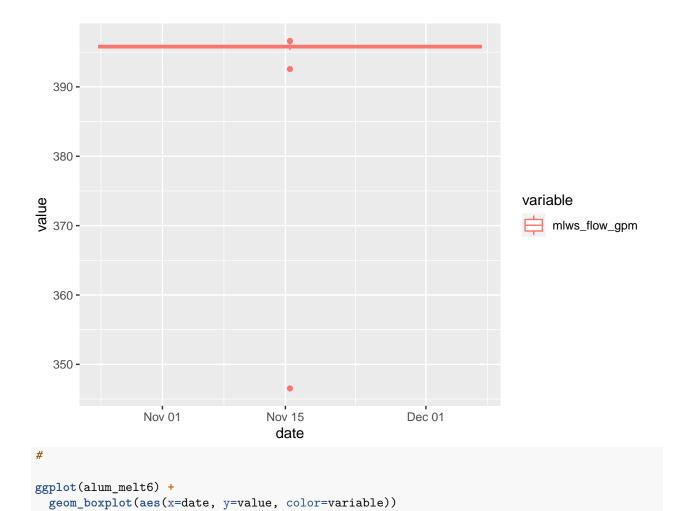


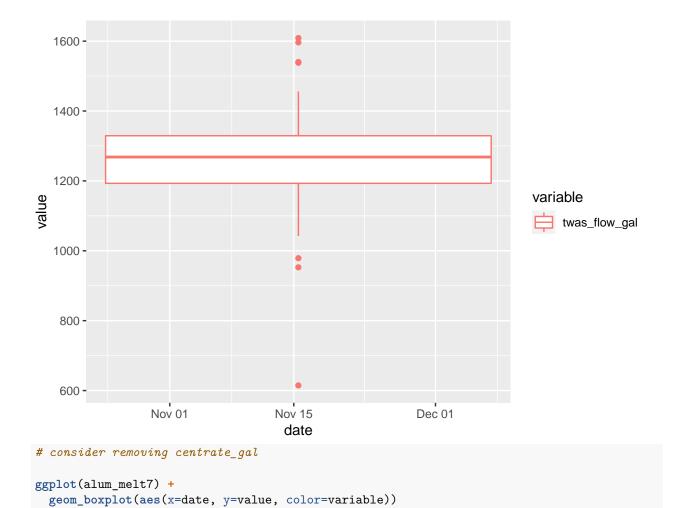




```
# outlier in melt 5. determine how to approach and then replot

ggplot(alum_melt5) +
  geom_boxplot(aes(x=date, y=value, color=variable))
```





Warning: Removed 2 rows containing non-finite values (stat_boxplot).

