# Online Analytical Chemistry notes: data analysis part II

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# Overview

These notes are on analysis of an intervention in time series data.

## Main ideas

- 1. Autocorrelation makes statistical analysis more difficult
- 2. Lurking variables can cause problems
- 3. We can make inferences without statistical models (at least about one experimental unit)
- 4. Time series methods exist but are not simple and can (should?) be avoided
- 5. Repetition at experimental unit level is (still) important for inferences
- 6. Response feature analysis works well for assessing interventions

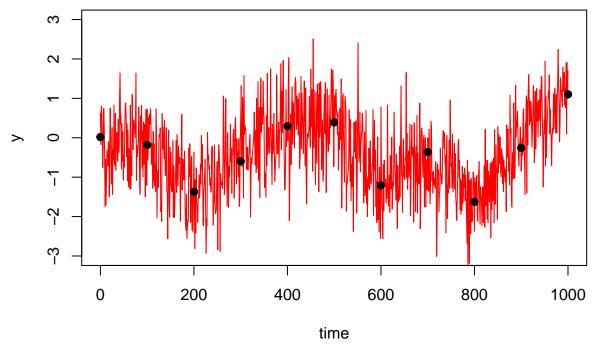
## Autocorrelation

Autocorrelation is just correlation between adjacent measurements. It is commonly present in time series type data. Autocorrelation could have multiple sources, e.g., instrument drift, reaction or mass transfer causing changes over time, change in ambient temperature or other conditions, ...

Here are some generated autocorrelated data.

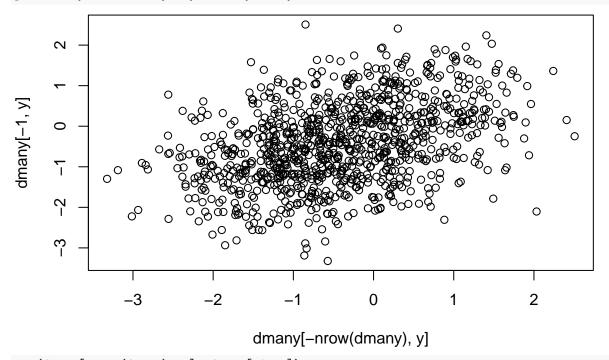
```
set.seed(10)
dfew <- data.table(time = 0:10 * 100, y = rnorm(11))
plot(y ~ time, data = dfew, pch = 19, ylim = c(-3, 3))
dmany <- data.table(time = 0:1000, y = approx(dfew[, time], dfew[, y], xout = 0:1000)$y + rnorm(1001, scheme)</pre>
```

```
lines(y ~ time, data = dmany, col = 'red')
points(y ~ time, data = dfew, pch = 19)
```



See the correlation? Find it by

plot(dmany[-nrow(dmany), y], dmany[-1, y])



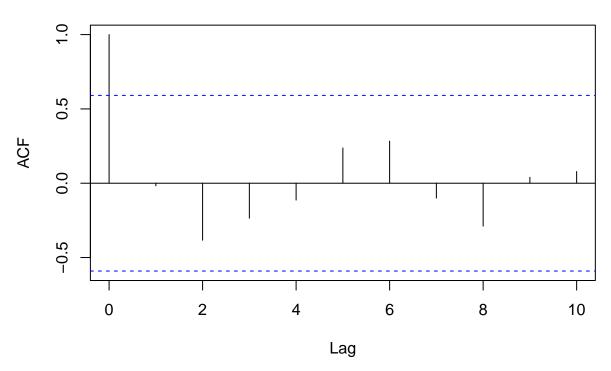
cor(dmany[-nrow(dmany), y], dmany[-1, y])

## [1] 0.4059568

In R the acf() function can be used to check. It returns the correlation between the data and lagged versions.

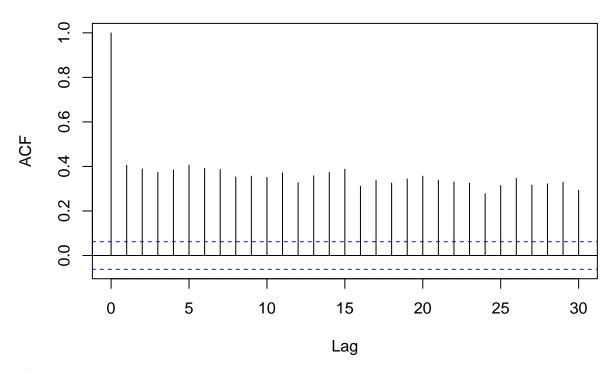
(acf(dfew\$y))

# Series dfew\$y



```
##
## Autocorrelations of series 'dfew$y', by lag
##
## 0 1 2 3 4 5 6 7 8 9 10
## 1.000 -0.018 -0.383 -0.236 -0.113 0.237 0.283 -0.100 -0.289 0.040 0.079
(acf(dmany$y))
```

# Series dmany\$y



```
##
  Autocorrelations of series 'dmany$y', by lag
##
                    2
                           3
                                                     7
##
              1
                                 4
                                        5
                                              6
                                                           8
                                                                  9
                                                                       10
                                                                              11
                                                                                    12
  1.000 0.405 0.389 0.374 0.384 0.406 0.391 0.386 0.353 0.356 0.351 0.371
##
                                                                                 0.327
##
      13
             14
                   15
                          16
                                             19
                                                    20
                                                          21
                                                                 22
                                                                       23
                                                                              24
                                                                                    25
                                17
                                       18
## 0.358 0.373 0.388 0.310 0.337 0.326 0.344 0.356 0.338 0.330 0.325 0.277 0.314
             27
                   28
                          29
                                30
##
      26
## 0.346 0.316 0.322 0.329 0.293
```

The presence of autocorrelation violates the assumption of independent observations. So it is important that measurements are not treated as independent.

# Lurking variables

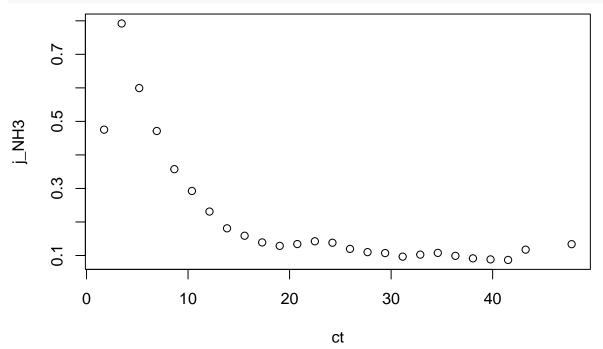
Lurking variables are some variables that affect the response but are not measured or considered in an analysis. They are probably more of a problem for observational data than experimental data. But a combination of a lurking variable and autocorrelation could make for very wrong inferences! Here is an example.

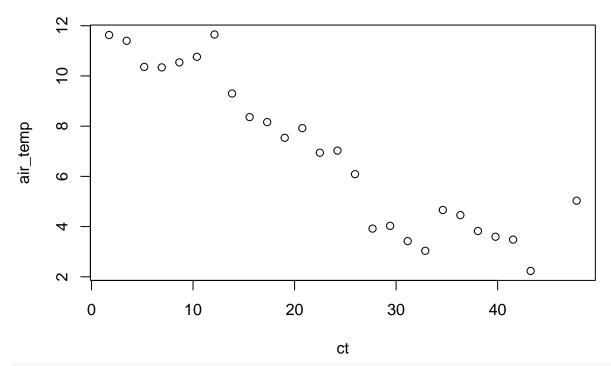
```
library(data.table)
amm_int <- fread('../data/NH3_emis_acid_interval.csv')</pre>
amm_int
##
         pmid
                   ct
                                 dt
                                                 t start
                                                                         t end
      1: 1947
##
                 1.73
                        1.7333 1.73 2020-11-18 13:40:00 2020-11-18 15:24:00
      2: 1947
                        3.4667 1.73 2020-11-18 15:24:00 2020-11-18 17:08:00
##
                 3.46
##
      3: 1947
                 5.19
                        5.2000 1.73 2020-11-18 17:08:00 2020-11-18 18:52:00
##
      4: 1947
                 6.92
                        6.9333 1.73 2020-11-18 18:52:00 2020-11-18 20:36:00
      5: 1947
                 8.65
                        8.6667 1.73 2020-11-18 20:36:00 2020-11-18 22:20:00
##
```

```
##
## 3485: 1982 178.19 178.5300 1.73 2020-12-16 23:49:00 2020-12-17 01:33:00
## 3486: 1982 179.92 180.2700 1.73 2020-12-17 01:33:00 2020-12-17 03:17:00
## 3487: 1982 181.65 182.0000 1.73 2020-12-17 03:17:00 2020-12-17 05:01:00
  3488: 1982 183.38 183.7300 1.73 2020-12-17 05:01:00 2020-12-17 06:45:00
  3489: 1982 185.11 185.4700 1.73 2020-12-17 06:45:00 2020-12-17 08:29:00
##
             j_NH3 air_temp
      1: 0.0088216
                     11.470
##
##
      2: 0.0000000
                     11.600
##
      3: 0.0061700
                     10.510
##
      4: 0.0136090
                     10.240
##
      5: 0.0154260
                     10.560
##
## 3485: 0.0100490
                      5.818
## 3486: 0.0098460
                      5.763
## 3487: 0.0095709
                      5.917
## 3488: 0.0099536
                      6.172
## 3489: 0.0116350
                      6.791
amm_sub <- amm_int[ct < 48 & pmid == 1951, ]
```

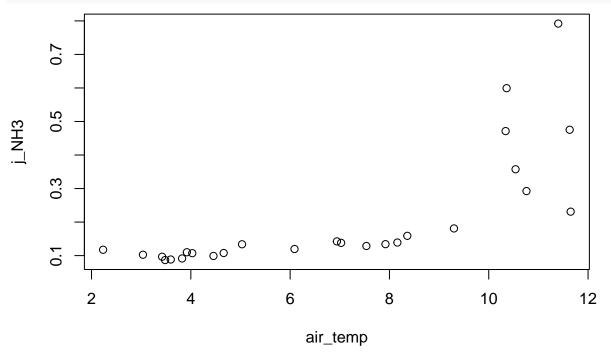
Emission decreased over the first couple days and so did temperature. Did the change in temperature cause the drop in emission?

```
plot(j_NH3 ~ ct, data = amm_sub)
```





plot(j\_NH3 ~ air\_temp, data = amm\_sub)



Let's fit a (stupid!) model.

```
mod1 <- lm(j_NH3 ~ air_temp, data = amm_sub)
summary(mod1)</pre>
```

```
##
## Call:
## lm(formula = j_NH3 ~ air_temp, data = amm_sub)
##
## Residuals:
```

```
##
                 1Q
                     Median
## -0.18560 -0.08390 -0.00142 0.03396 0.38613
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.087489
                          0.060474 -1.447
                                               0.161
                           0.008007
                                     5.405 1.5e-05 ***
## air_temp
               0.043275
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.1243 on 24 degrees of freedom
## Multiple R-squared: 0.549, Adjusted R-squared: 0.5302
## F-statistic: 29.21 on 1 and 24 DF, p-value: 1.496e-05
mod2 <- lm(log10(j_NH3) ~ air_temp, data = amm_sub)</pre>
summary(mod2)
##
## Call:
## lm(formula = log10(j_NH3) ~ air_temp, data = amm_sub)
## Residuals:
##
       Min
                  1Q
                      Median
                                    30
## -0.22435 -0.08768 -0.01825 0.05138 0.32997
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.31326
                          0.07009 -18.736 7.84e-16 ***
                           0.00928
                                   8.337 1.51e-08 ***
## air_temp
               0.07737
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.1441 on 24 degrees of freedom
## Multiple R-squared: 0.7433, Adjusted R-squared: 0.7326
## F-statistic: 69.51 on 1 and 24 DF, p-value: 1.508e-08
That's a very low p value. This is bad bad practice!
```

# Inferences without statistics

Here are some data from a recent paper.

### **ENVIRONMENTAL RESEARCH**

**LETTERS** 



#### LETTER

#### **OPEN ACCESS**

# A high efficiency gas phase photoreactor for eradication of methane from low-concentration sources

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E-mail: msj@chem.ku.dk 18 December 2023

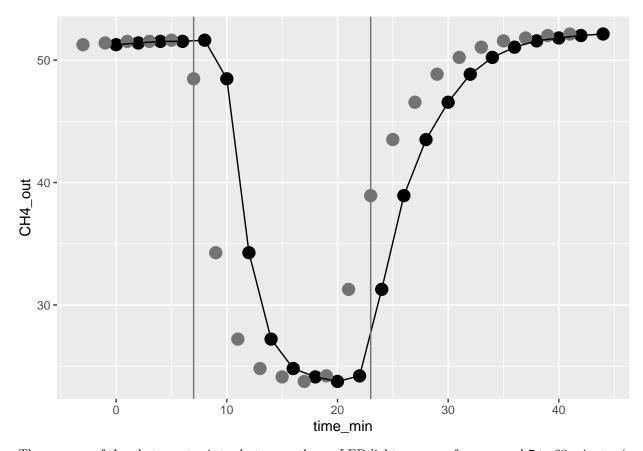
```
library(data.table)
```

```
pr <- fread('../data/photo_reactor.csv')</pre>
pr[, led := factor(led)]
head(pr)
```

```
##
                date_time Cl2_out
                                     Cl2_in CH4_out
                                                        CH4_in time_min led
## 1: 2023-06-09 12:15:00 102.2350 102.06809 51.25882 51.30022
## 2: 2023-06-09 12:17:00 101.4895 101.29243 51.39887 51.78323
                                                                      2
                                                                          0
## 3: 2023-06-09 12:19:00 101.1526 99.82636 51.52996 51.88457
                                                                          0
## 4: 2023-06-09 12:21:00 100.6248 100.54421 51.53150 51.87321
                                                                      6
                                                                          0
## 5: 2023-06-09 12:23:00 100.0695 100.10595 51.62850 51.97208
                                                                      8
                                                                          1
## 6: 2023-06-09 12:25:00 89.7600 100.15390 48.47417 52.33748
                                                                     10
```

### library(ggplot2)

```
ggplot(pr, aes(time_min, CH4_out)) +
  geom point(size = 4) +
  geom_point(aes(x = I(time_min - 3)), colour = 'gray45', size = 4) +
  geom_line() +
  geom_vline(xintercept = c(7, 23), colour = 'gray45')
```



The purpose of the photoreactor is to destroy methane. LED lights were on from around 7 to 23 minutes (see vertical lines). The authors concluded—without application of a statistical model—that these measurements show that the light drive about destruction of about 50% of the methane coming into the reactor.

## Time series methods

There are some statistical methods designed for analyzing time series data. Most seem to have a focus on prediction and perhaps less so on hypothesis testing. ARMA and ARIMA are approaches that include a moving average component. There are approaches for assessing interventions or perturbations. If you are interested, the book by McDowall et al. (2019) is not a bad place to start, but it is not focused on implementation in R or Python.

# Repeat experiments

As we discussed last week, statistical inference with any really useul scope really requires repeated application of a treatment or intervention to individual experimental units, i.e., multiple experimental units. So if you can, repeat. Then response feature analysis can easily be used. Here we will create a larger dataset from the one photoreactor run.

```
pr <- fread('../data/photo_reactor.csv')
pr[, led := factor(led)]
head(pr)</pre>
```

```
## 1: 2023-06-09 12:15:00 102.2350 102.06809 51.25882 51.30022 0 0 ## 2: 2023-06-09 12:17:00 101.4895 101.29243 51.39887 51.78323 2 0 ## 3: 2023-06-09 12:19:00 101.1526 99.82636 51.52996 51.88457 4 0
```

```
## 4: 2023-06-09 12:21:00 100.6248 100.54421 51.53150 51.87321 6 0
## 5: 2023-06-09 12:23:00 100.0695 100.10595 51.62850 51.97208 8 1
## 6: 2023-06-09 12:25:00 89.7600 100.15390 48.47417 52.33748 10 1
```

We'll make a fake dataset here with 5 separate runs, all operated the same way.

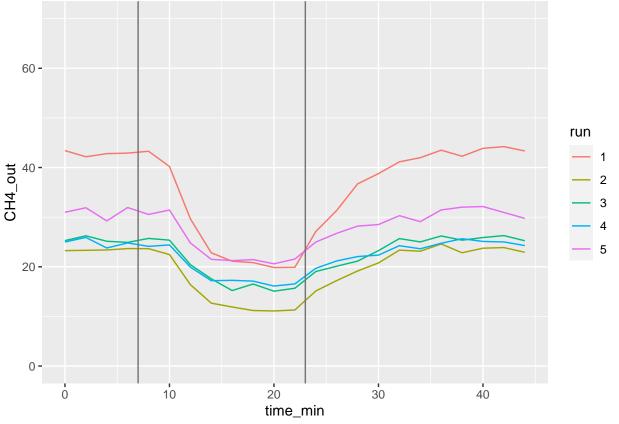
```
prm <- data.table()

n <- 5

set.seed(2)
for (i in 1:n) {
    d <- pr
    d[, run := i]
    d[, CH4_out := CH4_out * rnorm(1, mean = 1, sd = 0.2) + rnorm(1, sd = 3) + rnorm(nrow(pr), sd = 0.5)]
    prm <- rbind(prm, d)
}

prm[, run := factor(run)]

ggplot(prm, aes(time_min, CH4_out, colour = run)) +
    geom_line() +
    geom_vline(xintercept = c(7, 23), colour = 'gray45') +
    ylim(0, 70)</pre>
```



```
table(prm[, .(time_min, led)])
```

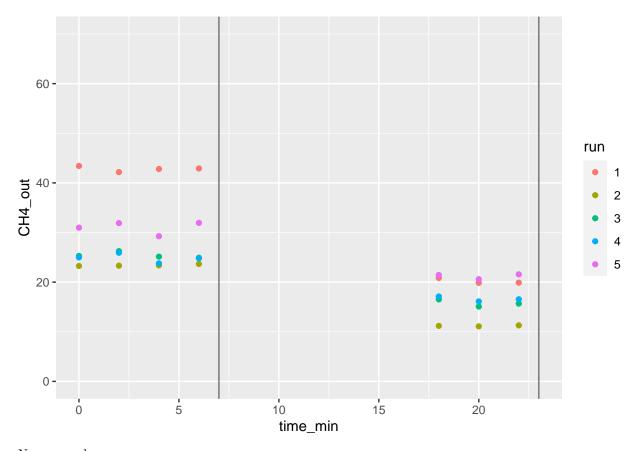
```
## led
## time_min 0 1
```

```
0
            5 0
##
         2
            5 0
##
         4
            5 0
##
##
         6 5 0
            0 5
##
         8
##
         10 0 5
##
         12 0 5
         14 0 5
##
##
         16 0 5
##
         18 0 5
##
         20 0 5
##
         22 0 5
##
         24 5 0
         26 5 0
##
##
         28 5 0
         30 5 0
##
##
         32 5 0
         34 5 0
##
##
         36 5 0
         38 5 0
##
         40 5 0
##
##
         42 5 0
##
         44 5 0
```

Let's try to cut out the decay and build up of methane, i.e., target what seem to be steady-state concentrations (although really we should probably have longer runs for this).

```
pra <- prm[time_min < 8 | (time_min > 16 & time_min < 24), ]

ggplot(pra, aes(time_min, CH4_out, colour = run)) +
  geom_point() +
  geom_vline(xintercept = c(7, 23), colour = 'gray45') +
  ylim(0, 70)</pre>
```



Now mean by run.

```
prm <- pra[, .(CH4_out_mn = mean(CH4_out)), by = .(run, led)]
prm</pre>
```

```
##
       run led CH4_out_mn
                  42.82241
##
    1:
         1
              0
##
    2:
         1
              1
                  20.19788
         2
                  23.40956
##
    3:
              0
##
    4:
         2
                  11.19447
              1
                  25.39618
##
    5:
         3
              0
##
    6:
         3
                  15.76424
              1
##
    7:
         4
              0
                  24.87557
##
    8:
         4
                  16.59396
##
    9:
         5
              0
                  31.01816
## 10:
         5
              1
                  21.20504
```

And fit model, using run as blocking variable.

```
mod1 <- lm(CH4_out_mn ~ led + run, data = prm)
summary(mod1)</pre>
```

```
##
## Call:
## lm(formula = CH4_out_mn ~ led + run, data = prm)
##
## Residuals:
## 1 2 3 4 5 6 7 8 9 10
## 5.0556 -5.0556 -0.1491 0.1491 -1.4407 1.4407 -2.1158 2.1158 -1.3501 1.3501
```

```
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 37.767 3.192 11.833 0.000292 ***
                           2.606 -4.802 0.008638 **
## led1
              -12.513
## run2
              -14.208
                           4.120 -3.448 0.026098 *
## run3
              -10.930
                           4.120 -2.653 0.056828 .
                           4.120 -2.615 0.059104 .
## run4
              -10.775
## run5
               -5.399
                           4.120 -1.310 0.260308
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 4.12 on 4 degrees of freedom
## Multiple R-squared: 0.9043, Adjusted R-squared: 0.7847
## F-statistic: 7.562 on 5 and 4 DF, p-value: 0.03628
anova (mod1)
## Analysis of Variance Table
## Response: CH4_out_mn
            Df Sum Sq Mean Sq F value
                                     Pr(>F)
             1 391.45 391.45 23.0561 0.008638 **
## led
             4 250.50 62.63 3.6886 0.117066
## run
## Residuals 4 67.91
                      16.98
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```