Mixed-effects models

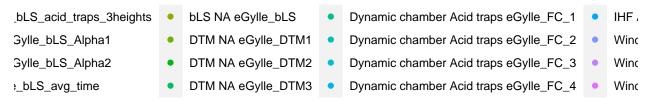
Sasha D. Hafner

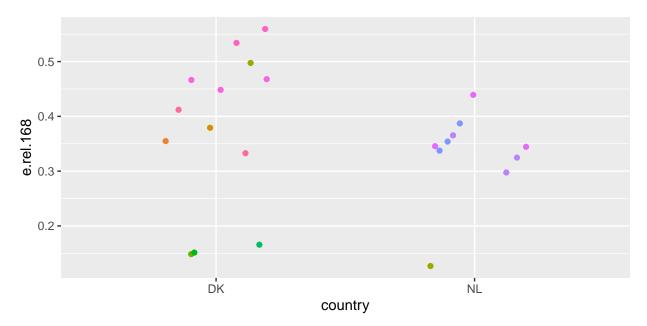
22 January, 2024 14:26

1. Take a look

```
pdat[, meas.tech.3 := paste(meas.tech, meas.tech.det, treat)]
ggplot(pdat, aes(country, e.rel.168, colour = meas.tech.3)) +
  geom_jitter(height = 0, width = 0.2) +
  theme(legend.position = 'top')
```

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





2. Mixed-effects models

```
pdat[, meas.level := paste(meas.tech, meas.tech.det, treat)]
pdat[, meas.level := gsub('bLS Alpha samplers.+$', 'bLS-alpha', meas.level)]
pdat[, meas.level := gsub('DTM.+$', 'DTM', meas.level)]
```

```
pdat[, meas.level := gsub('Wind tunnel NA', 'WT', meas.level)]
pdat[, meas.level := gsub('bLS NA eGylle_bLS', 'bLS-CRDS', meas.level)]
pdat[, meas.level := gsub('.+_FC_.+$', 'FC', meas.level)]
pdat[, meas.level := gsub('diluted$', '', meas.level)]
d1 <- pdat[country == 'DK', ]
d2 <- pdat[country == 'NL', ]</pre>
```

I-AU micromet

```
d1mm <- d1[meas.tech2 == 'micro met', ]</pre>
m1 <- lmer(e.rel.168 ~ 1 meas.level, data = d1mm)
summary(m1)
## Linear mixed model fit by REML ['lmerMod']
## Formula: e.rel.168 ~ 1 | meas.level
##
     Data: d1mm
##
## REML criterion at convergence: -5.8
##
## Scaled residuals:
      Min
           1Q Median
                                       Max
## -0.7730 -0.3206 0.1318 0.3865 0.6412
## Random effects:
## Groups
                           Variance Std.Dev.
              Name
## meas.level (Intercept) 0.0082889 0.09104
## Residual
                           0.0002955 0.01719
## Number of obs: 3, groups: meas.level, 2
## Fixed effects:
              Estimate Std. Error t value
## (Intercept) 0.43161
                          0.06523
                                     6.617
VarCorr(m1)
                           Std.Dev.
## Groups
              Name
## meas.level (Intercept) 0.091044
## Residual
                           0.017190
m2 <- lmer(log10(e.rel.168) ~ 1 meas.level, data = d1mm)
summary(m2)
## Linear mixed model fit by REML ['lmerMod']
## Formula: log10(e.rel.168) ~ 1 | meas.level
##
     Data: d1mm
##
## REML criterion at convergence: -5.5
##
## Scaled residuals:
               1Q Median
                                3Q
      Min
                                       Max
## -0.7840 -0.3151 0.1538 0.3920 0.6302
## Random effects:
```

```
## Groups
               Name
                           Variance Std.Dev.
## meas.level (Intercept) 0.0084479 0.09191
                           0.0004142 0.02035
## Number of obs: 3, groups: meas.level, 2
## Fixed effects:
               Estimate Std. Error t value
## (Intercept) -0.37024
                        0.06617 -5.595
VarCorr(m2)
## Groups
               Name
                           Std.Dev.
## meas.level (Intercept) 0.091912
## Residual
                           0.020352
100 * (10^(as.data.frame(VarCorr(m2))[, 5]) - 1)
## [1] 23.569771 4.797817
Total.
sqrt(sum(as.data.frame(VarCorr(m2))[, 5]^2))
## [1] 0.09413859
I-AU enclosure
d1en <- d1[meas.tech2 != 'micro met', ]</pre>
m1 <- lmer(e.rel.168 ~ 1 meas.level, data = d1en)
summary(m1)
## Linear mixed model fit by REML ['lmerMod']
## Formula: e.rel.168 ~ 1 | meas.level
      Data: d1en
##
## REML criterion at convergence: -24.3
##
## Scaled residuals:
##
       \mathtt{Min}
                1Q
                      Median
                                    3Q
## -1.56605 -0.40378 0.01123 0.33245 1.55594
##
## Random effects:
## Groups
               Name
                           Variance Std.Dev.
## meas.level (Intercept) 0.028064 0.1675
                           0.000645 0.0254
## Number of obs: 10, groups: meas.level, 4
## Fixed effects:
               Estimate Std. Error t value
## (Intercept) 0.38367
                           0.08416
VarCorr(m1)
## Groups
               Name
                           Std.Dev.
## meas.level (Intercept) 0.167524
```

0.025397

Residual

```
m2 <- lmer(log10(e.rel.168) ~ 1 meas.level, data = d1en)
summary(m2)
## Linear mixed model fit by REML ['lmerMod']
## Formula: log10(e.rel.168) ~ 1 | meas.level
##
      Data: d1en
##
## REML criterion at convergence: -19.4
##
## Scaled residuals:
##
       Min
                  1Q
                      Median
                                    30
                                            Max
## -1.45940 -0.37882 -0.03926 0.33695 1.47468
##
## Random effects:
## Groups
                           Variance Std.Dev.
## meas.level (Intercept) 0.058782 0.24245
## Residual
                           0.001001 0.03164
## Number of obs: 10, groups: meas.level, 4
##
## Fixed effects:
              Estimate Std. Error t value
##
## (Intercept) -0.4601
                            0.1217 -3.782
VarCorr(m2)
## Groups
                           Std.Dev.
              Name
## meas.level (Intercept) 0.242450
## Residual
                           0.031637
100 * (10^(as.data.frame(VarCorr(m2))[, 5]) - 1)
## [1] 74.763381 7.556652
```

II-WUR micromet

```
d2mm <- d2[meas.tech2 == 'micro met', ]</pre>
d2mm[, .(meas.level, e.rel.168)]
##
                                           meas.level e.rel.168
## 1:
                                            bLS-CRDS
                                                       0.12683
## 2: bLS Acid traps eGylle_bLS_acid_traps_3heights
                                                             NA
## 3:
                  bLS CRDS avg. eGylle_bLS_avg_time
                                                             NA
## 4:
                           IHF Acid traps eGylle_IHF
                                                             NΑ
d2mm[, .(meas.level, e.rel.final)]
##
                                           meas.level e.rel.final
## 1:
                                            bLS-CRDS
                                                          0.12844
## 2: bLS Acid traps eGylle_bLS_acid_traps_3heights
                                                          0.14881
## 3:
                  bLS CRDS avg. eGylle_bLS_avg_time
                                                          0.12428
## 4:
                           IHF Acid traps eGylle_IHF
                                                          0.09831
```

Drop 3 heights and add replicate data (or some representation of it) manually. **Note: IHF value does not match Table 3–update ALFAM2 data!**

```
dimpinger <- data.table(meas.level = 'bLS-impinger', e.rel.final = c(0.125, 0.148, 0.171))
sd(dimpinger[, e.rel.final])
## [1] 0.023
mean(dimpinger[, e.rel.final])
## [1] 0.148
d2mm <- d2mm[meas.level != 'bLS Acid traps eGylle_bLS_acid_traps_3heights', ]</pre>
d2mm <- rbind(d2mm, dimpinger, fill = TRUE)</pre>
d2mm[, .(meas.level, e.rel.final)]
##
                             meas.level e.rel.final
                                bLS-CRDS
## 1:
                                             0.12844
## 2: bLS CRDS avg. eGylle_bLS_avg_time
                                             0.12428
              IHF Acid traps eGylle_IHF
## 3:
                                             0.09831
## 4:
                           bLS-impinger
                                             0.12500
## 5:
                           bLS-impinger
                                             0.14800
                           bLS-impinger
                                             0.17100
m1 <- lmer(e.rel.final ~ 1 meas.level, data = d2mm)</pre>
summary(m1)
## Linear mixed model fit by REML ['lmerMod']
## Formula: e.rel.final ~ 1 | meas.level
##
      Data: d2mm
## REML criterion at convergence: -21.3
##
## Scaled residuals:
        \mathtt{Min}
                 1Q
                      Median
                                     3Q
                                             Max
## -1.01716 -0.56392 -0.08467 0.29274 1.49161
##
## Random effects:
## Groups
             Name
                           Variance Std.Dev.
## meas.level (Intercept) 0.0001909 0.01382
                           0.0004403 0.02098
## Residual
## Number of obs: 6, groups: meas.level, 4
##
## Fixed effects:
               Estimate Std. Error t value
## (Intercept) 0.12891
                           0.01139
                                      11.32
VarCorr(m1)
## Groups
               Name
                           Std.Dev.
## meas.level (Intercept) 0.013817
## Residual
                           0.020983
m2 <- lmer(log10(e.rel.final) ~ 1 meas.level, data = d2mm)</pre>
summary(m2)
## Linear mixed model fit by REML ['lmerMod']
## Formula: log10(e.rel.final) ~ 1 | meas.level
##
      Data: d2mm
##
```

```
## REML criterion at convergence: -9.6
##
## Scaled residuals:
##
       Min
            1Q
                     Median
                                   3Q
                                           Max
## -1.00957 -0.58008 0.00164 0.30991 1.36761
##
## Random effects:
## Groups
              Name
                          Variance Std.Dev.
## meas.level (Intercept) 0.002795 0.05287
## Residual
                          0.004123 0.06421
## Number of obs: 6, groups: meas.level, 4
##
## Fixed effects:
##
              Estimate Std. Error t value
## (Intercept) -0.89863
                          0.03853 -23.32
VarCorr(m2)
## Groups
              Name
                          Std.Dev.
## meas.level (Intercept) 0.052872
## Residual
                          0.064207
100 * (10^(as.data.frame(VarCorr(m2))[, 5]) - 1)
## [1] 12.94623 15.93303
sqrt(sum(as.data.frame(VarCorr(m2))[, 5]^2))
## [1] 0.08317441
II-WUR enclosure
d2en <- d2[meas.tech2 != 'micro met' & meas.level %in% c('WT 20', 'FC') & !grep1('diluted', treat), ]
d2en[, .(meas.tech.orig, e.rel.168)]
##
      meas.tech.orig e.rel.168
## 1:
         Wind tunnel
                       0.35403
## 2:
         Wind tunnel
                       0.38712
## 3:
         Wind tunnel
                       0.33742
## 4: Dynamic chamber
## 5: Dynamic chamber
                            NA
```

```
##
      meas.level e.rel.168
           WT 20
                   0.35403
## 1:
## 2:
           WT 20
                   0.38712
                  0.33742
## 3:
           WT 20
## 4:
              FC
              FC
## 5:
                        NA
## 6:
              FC
                        NA
            FC
## 7:
                        NA
```

d2en[, .(meas.level, e.rel.168)]

6: Dynamic chamber

7: Dynamic chamber

NA

```
d2en[, .(meas.level, e.rel.final)]
      meas.level e.rel.final
## 1:
          WT 20
                    0.35426
          WT 20
                     0.38741
## 2:
## 3:
          WT 20
                     0.33770
## 4:
             FC
                     0.15707
## 5:
             FC
                     0.16987
## 6:
             FC
                     0.16513
## 7:
             FC
                     0.22263
m1 <- lmer(e.rel.final ~ 1 meas.level, data = d2en)
summary(m1)
## Linear mixed model fit by REML ['lmerMod']
## Formula: e.rel.final ~ 1 | meas.level
     Data: d2en
##
##
## REML criterion at convergence: -19.6
##
## Scaled residuals:
##
      Min
            1Q Median
                                ЗQ
                                       Max
## -0.8083 -0.6282 -0.3524 0.4451 1.5268
##
## Random effects:
## Groups
            Name
                          Variance Std.Dev.
## meas.level (Intercept) 0.0161714 0.12717
## Residual
                           0.0007882 0.02808
## Number of obs: 7, groups: meas.level, 2
##
## Fixed effects:
              Estimate Std. Error t value
## (Intercept) 0.26905
                           0.09056
                                     2.971
VarCorr(m1)
## Groups
               Name
                           Std.Dev.
## meas.level (Intercept) 0.127167
## Residual
                           0.028076
m2 <- lmer(log10(e.rel.final) ~ 1 meas.level, data = d2en)
summary(m2)
## Linear mixed model fit by REML ['lmerMod']
## Formula: log10(e.rel.final) ~ 1 | meas.level
##
      Data: d2en
##
## REML criterion at convergence: -11.7
##
## Scaled residuals:
               1Q Median
       Min
                                3Q
                                       Max
## -0.9709 -0.5004 -0.3632 0.3001 1.7348
##
## Random effects:
                           Variance Std.Dev.
## Groups
               Name
## meas.level (Intercept) 0.046344 0.21528
```