

Climate Water Loss Experiment - Capture Hydration Analysis

Savannah Weaver

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Packages

```
if (!require("tidyverse")) install.packages("tidyverse")
library("tidyverse") # workflow and plots
if (!require("lme4")) install.packages("lme4")
library("lme4") # for LMMs
if (!require("lmerTest")) install.packages("lmerTest")
library("lmerTest") # for p-values
if (!require("UsingR")) install.packages("UsingR")
library("UsingR") # simple.eda model assumption checker
if (!require("ggpubr")) install.packages("ggpubr")
library("ggpubr") # for multi-ggplot figs
if (!require("broom.mixed")) install.packages("broom.mixed")
library("broom.mixed") # lmer model export
if (!require("AICcmodavg")) install.packages("AICcmodavg")
library("AICcmodavg") # model selection
if (!require("car")) install.packages("car")
library("car") # VIFs
if (!require("AICcmodavg")) install.packages("AICcmodavg")
library("AICcmodavg") # model selection
if (!require("RColorBrewer")) install.packages("RColorBrewer")
library("RColorBrewer") # color
```

Background and Goals

This data was collected June - August by Master's student Savannah Weaver, advisor Dr. Emily Taylor, and research assistants Tess McIntyre and Taylor Van Rossum. Adult male *Sceloporus occidentalis* were caught across the Cal Poly campus then acclimated to 4 different climate treatments. **This R file analyzes the state and variation of osmotic balance and regulation at the time of capture.** Please refer to the published scientific journal article for full details.

Load Data

```
dat <- read_rds("./data/analysis_data_capture.RDS")
summary(dat)
```

```
## individual_ID      mass_g      hematocrit_percent      SVL_mm
## 201      : 1      Min.      : 8.80      Min.      :27.00      Min.      :60.00
## 202      : 1      1st Qu.:10.60      1st Qu.:34.25      1st Qu.:66.00
## 203      : 1      Median :11.65      Median :39.00      Median :67.00
## 204      : 1      Mean      :11.73      Mean      :38.93      Mean      :67.71
## 205      : 1      3rd Qu.:12.70      3rd Qu.:43.00      3rd Qu.:70.00
## 206      : 1      Max.      :17.40      Max.      :52.00      Max.      :77.00
## (Other):132
## capture_date      osmolality_mmol_kg_mean      CEWL_g_m2h_mean      msmt_temp_C
## Min.      :2021-06-16      Min.      :305.0      Min.      : 7.152      Min.      :25.90
## 1st Qu.:2021-06-26      1st Qu.:334.3      1st Qu.:17.255      1st Qu.:26.72
## Median :2021-07-20      Median :344.6      Median :21.030      Median :26.96
## Mean      :2021-07-16      Mean      :348.3      Mean      :20.760      Mean      :27.20
```

```

## 3rd Qu.:2021-08-08 3rd Qu.:361.9 3rd Qu.:24.416 3rd Qu.:27.50
## Max. :2021-08-22 Max. :395.0 Max. :34.660 Max. :29.20
##
## msmt_RH_percent cloacal_temp_C date_time msmt_temp_K
## Min. :25.52 Min. :25.00 Min. :2021-06-16 09:54:00 Min. :299.1
## 1st Qu.:45.77 1st Qu.:26.00 1st Qu.:2021-06-26 12:59:30 1st Qu.:299.9
## Median :47.09 Median :26.00 Median :2021-07-20 13:17:00 Median :300.1
## Mean :44.08 Mean :26.45 Mean :2021-07-17 06:56:12 Mean :300.3
## 3rd Qu.:48.44 3rd Qu.:27.00 3rd Qu.:2021-08-08 13:39:00 3rd Qu.:300.6
## Max. :53.15 Max. :30.00 Max. :2021-08-22 15:19:00 Max. :302.4
##
## e_s_kPa_m e_a_kPa_m msmt_VPD_kPa SMI
## Min. :3.441 Min. :0.9894 Min. :1.612 Min. : 9.122
## 1st Qu.:3.616 1st Qu.:1.6913 1st Qu.:1.846 1st Qu.:10.926
## Median :3.669 Median :1.7342 Median :1.942 Median :11.687
## Mean :3.724 Mean :1.6312 Mean :2.093 Mean :11.690
## 3rd Qu.:3.790 3rd Qu.:1.7865 3rd Qu.:2.053 3rd Qu.:12.347
## Max. :4.194 Max. :1.8502 Max. :3.021 Max. :14.263
##
## capture_date_time hold_time_sec hold_time_min
## Min. :2021-06-16 08:28:00 Length:138 Length:138
## 1st Qu.:2021-06-26 09:44:45 Class :difftime Class :difftime
## Median :2021-07-20 09:52:00 Mode :numeric Mode :numeric
## Mean :2021-07-14 14:50:11
## 3rd Qu.:2021-08-08 09:56:45
## Max. :2021-08-22 13:25:00
## NA's :14
## hold_time_hr temp_C_interpol RH_percent_interpol VPD_kPa_int
## Length:138 Min. :15.11 Min. : 19.73 Min. :0.0000
## Class :difftime 1st Qu.:19.91 1st Qu.: 59.20 1st Qu.:0.5420
## Mode :numeric Median :21.91 Median : 69.33 Median :0.8284
## Mean :23.41 Mean : 62.27 Mean :1.4295
## 3rd Qu.:23.91 3rd Qu.: 77.29 3rd Qu.:1.2321
## Max. :35.83 Max. :100.00 Max. :4.9400
## NA's :14 NA's :14 NA's :14
## wind_mph_interpol solar_rad_W_sqm_interpol
## Min. : 0.100 Min. : 294.7
## 1st Qu.: 2.025 1st Qu.: 682.9
## Median : 3.100 Median : 759.9
## Mean : 4.406 Mean : 762.9
## 3rd Qu.: 5.880 3rd Qu.: 873.2
## Max. :12.720 Max. :1007.0
## NA's :14 NA's :14

```

note IDs I do not have data for: 254, 284, 304

LMMs

We use simple linear models here because date and individual ID do not need to be random effects: we want to know how/why values varied across dates, and each individual only has one measurement.

Hematocrit

Models

First, start with a full model with every probable potential predictor in it, then check for multicollinearity.

```
hct_mod1 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # body size options
               mass_g + SVL_mm + SMI +
               # weather at the time of capture
               temp_C_interpol * VPD_kPa_int +
               wind_mph_interpol + solar_rad_W_sqm_interpol
               )
hct_mod1_VIFs <- data.frame(VIF = car::vif(hct_mod1)) %>%
  arrange(desc(VIF))
hct_mod1_VIFs
```

```
##                                VIF
## VPD_kPa_int                    776.842577
## temp_C_interpol:VPD_kPa_int  259.587300
## temp_C_interpol               180.895804
## mass_g                        147.299036
## SVL_mm                       137.309907
## SMI                           69.984055
## wind_mph_interpol             5.202791
## solar_rad_W_sqm_interpol      3.988220
```

remove VPD*temp interaction:

```
hct_mod2 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # body size
               mass_g + SVL_mm + SMI +
               # weather at the time of capture
               temp_C_interpol + VPD_kPa_int +
               wind_mph_interpol + solar_rad_W_sqm_interpol
               )
hct_mod2_VIFs <- data.frame(VIF = car::vif(hct_mod2)) %>%
  arrange(desc(VIF))
hct_mod2_VIFs
```

```
##                                VIF
## mass_g                        144.935446
## SVL_mm                       134.828923
## temp_C_interpol               89.583462
## VPD_kPa_int                   88.715132
## SMI                           68.677230
## wind_mph_interpol             4.086594
## solar_rad_W_sqm_interpol      2.293753
```

```
drop1(hct_mod2)
```

```
## Single term deletions
```

```
##
```

```
## Model:
```

```
## hematocrit_percent ~ mass_g + SVL_mm + SMI + temp_C_interpol +
##      VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol
##                Df Sum of Sq    RSS    AIC
## <none>                        2998.3 411.00
## mass_g                1      0.970 2999.3 409.05
## SVL_mm                1      0.725 2999.0 409.03
## SMI                   1      0.685 2999.0 409.03
## temp_C_interpol       1      4.349 3002.7 409.18
## VPD_kPa_int           1      1.631 2999.9 409.07
## wind_mph_interpol     1    250.305 3248.6 418.95
## solar_rad_W_sqm_interpol 1     77.557 3075.9 412.17
```

drop SVL:

```
hct_mod3 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # body size
               mass_g + SMI +
               # weather at the time of capture
               temp_C_interpol + VPD_kPa_int +
               wind_mph_interpol + solar_rad_W_sqm_interpol
               )
hct_mod3_VIFs <- data.frame(VIF = car::vif(hct_mod3)) %>%
  arrange(desc(VIF))
hct_mod3_VIFs
```

```
##                VIF
## temp_C_interpol  84.911535
## VPD_kPa_int     83.181379
## wind_mph_interpol  4.052405
## solar_rad_W_sqm_interpol 2.215093
## SMI             1.405795
## mass_g          1.246836
```

```
drop1(hct_mod3)
```

Single term deletions

##

Model:

```
## hematocrit_percent ~ mass_g + SMI + temp_C_interpol + VPD_kPa_int +
##      wind_mph_interpol + solar_rad_W_sqm_interpol
##                Df Sum of Sq    RSS    AIC
## <none>                        2999.0 409.03
## mass_g                1      2.186 3001.2 407.13
## SMI                   1    136.330 3135.4 412.55
## temp_C_interpol       1      3.773 3002.8 407.19
## VPD_kPa_int           1      1.208 3000.2 407.08
## wind_mph_interpol     1    254.908 3253.9 417.15
## solar_rad_W_sqm_interpol 1     77.461 3076.5 410.20
```

drop temperature:

```
hct_mod4 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # body size
```

```

        mass_g + SMI +
        # weather at the time of capture
        VPD_kPa_int +
        wind_mph_interpol + solar_rad_W_sqm_interpol
    )
hct_mod4_VIFs <- data.frame(VIF = car::vif(hct_mod4)) %>%
  arrange(desc(VIF))
hct_mod4_VIFs

##              VIF
## wind_mph_interpol  4.039979
## VPD_kPa_int        4.010514
## SMI                1.402937
## mass_g             1.246609
## solar_rad_W_sqm_interpol 1.068123
drop1(hct_mod4)

## Single term deletions
##
## Model:
## hematocrit_percent ~ mass_g + SMI + VPD_kPa_int + wind_mph_interpol +
##   solar_rad_W_sqm_interpol
##              Df Sum of Sq    RSS    AIC
## <none>                        3002.8 407.19
## mass_g              1      2.110 3004.9 405.28
## SMI                 1    134.567 3137.4 410.63
## VPD_kPa_int         1     13.135 3015.9 405.73
## wind_mph_interpol    1    252.259 3255.1 415.19
## solar_rad_W_sqm_interpol 1    113.672 3116.5 409.80

VIFs are all below 5 now, so start backwards selection.

Drop mass first:
hct_mod5 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # body size
               SMI +
               # weather at the time of capture
               VPD_kPa_int +
               wind_mph_interpol + solar_rad_W_sqm_interpol
               )
drop1(hct_mod5)

## Single term deletions
##
## Model:
## hematocrit_percent ~ SMI + VPD_kPa_int + wind_mph_interpol +
##   solar_rad_W_sqm_interpol
##              Df Sum of Sq    RSS    AIC
## <none>                        3004.9 405.28
## SMI              1    185.660 3190.6 410.71
## VPD_kPa_int       1     13.221 3018.1 403.82
## wind_mph_interpol  1    262.114 3267.0 413.65

```

```
## solar_rad_W_sqm_interpol 1 116.011 3120.9 407.98
```

Drop VPD:

```
hct_mod6 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # body size
               SMI +
               # weather at the time of capture
               wind_mph_interpol + solar_rad_W_sqm_interpol
               )
summary(hct_mod6)
```

```
##
## Call:
## lm(formula = hematocrit_percent ~ SMI + wind_mph_interpol + solar_rad_W_sqm_interpol,
##     data = dat)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -11.7663  -3.0319  -0.1456   3.4610  11.0669
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    17.030786    6.116508   2.784  0.00623 **
## SMI              1.244585    0.447364   2.782  0.00628 **
## wind_mph_interpol    0.607540    0.123216   4.931 2.66e-06 ***
## solar_rad_W_sqm_interpol  0.006307    0.003098   2.036  0.04398 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.015 on 120 degrees of freedom
## (14 observations deleted due to missingness)
## Multiple R-squared:  0.1945, Adjusted R-squared:  0.1744
## F-statistic: 9.659 on 3 and 120 DF,  p-value: 9.26e-06
```

```
drop1(hct_mod6)
```

```
## Single term deletions
##
## Model:
## hematocrit_percent ~ SMI + wind_mph_interpol + solar_rad_W_sqm_interpol
##              Df Sum of Sq  RSS   AIC
## <none>                 3018.1 403.82
## SMI                   1   194.66 3212.8 409.57
## wind_mph_interpol     1   611.47 3629.6 424.70
## solar_rad_W_sqm_interpol 1   104.23 3122.4 406.03
```

Drop solar:

```
hct_mod7 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # body size
               SMI +
               # weather at the time of capture
```

```

        wind_mph_interpol
      )
drop1(hct_mod7)

## Single term deletions
##
## Model:
## hematocrit_percent ~ SMI + wind_mph_interpol
##           Df Sum of Sq    RSS    AIC
## <none>                 3122.4 406.03
## SMI             1     171.27 3293.6 410.65
## wind_mph_interpol 1     594.00 3716.4 425.63

```

Drop SMI:

```

hct_mod8 <- lm(data = dat,
               # response variable
               hematocrit_percent ~
               # weather at the time of capture
               wind_mph_interpol
               )

```

Finally, null model:

```

hct_mod_null <- lm(data = dat,
                  # response variable
                  hematocrit_percent ~ 1)

```

Selection

Compare models 4-8 and the null model.

```

hct_models <- list(hct_mod4, hct_mod5, hct_mod6, hct_mod7,
                  hct_mod8, hct_mod_null)

#specify model names
hct_mod_names <- c('(model 4) ~ Wind-C, SMI, Solar-C, VPD-C, Mass',
                  '(model 5) ~ Wind-C, SMI, Solar-C, VPD-C',
                  '(model 6) ~ Wind-C, SMI, Solar-C',
                  '(model 7) ~ Wind-C, SMI',
                  '(model 8) ~ Wind-C',
                  'null model')

#calculate AIC of each model
hct_AICc <- data.frame(aictab(cand.set = hct_models,
                             modnames = hct_mod_names))

hct_AICc

##           Modnames K    AICc Delta_AICc
## 3      (model 6) ~ Wind-C, SMI, Solar-C 5 758.2275    0.000000
## 2      (model 5) ~ Wind-C, SMI, Solar-C, VPD-C 6 759.8926    1.665102
## 4      (model 7) ~ Wind-C, SMI 4 760.2651    2.037581
## 1 (model 4) ~ Wind-C, SMI, Solar-C, VPD-C, Mass 7 762.0530    3.825573
## 5      (model 8) ~ Wind-C 3 764.7505    6.523000
## 6      null model 2 861.5454 103.317962
##           Modellik      AICcWt      LL      Cum.Wt
## 3 1.000000e+00 5.045488e-01 -373.8595 0.5045488

```



```
## 2 4.349384e-01 2.194476e-01 -373.5873 0.7239964
## 4 3.610313e-01 1.821579e-01 -375.9645 0.9061543
## 1 1.476683e-01 7.450586e-02 -373.5438 0.9806602
## 5 3.833086e-02 1.933979e-02 -379.2752 1.0000000
## 6 3.671044e-23 1.852221e-23 -428.7283 1.0000000
```

The best models are models 6 and 5.

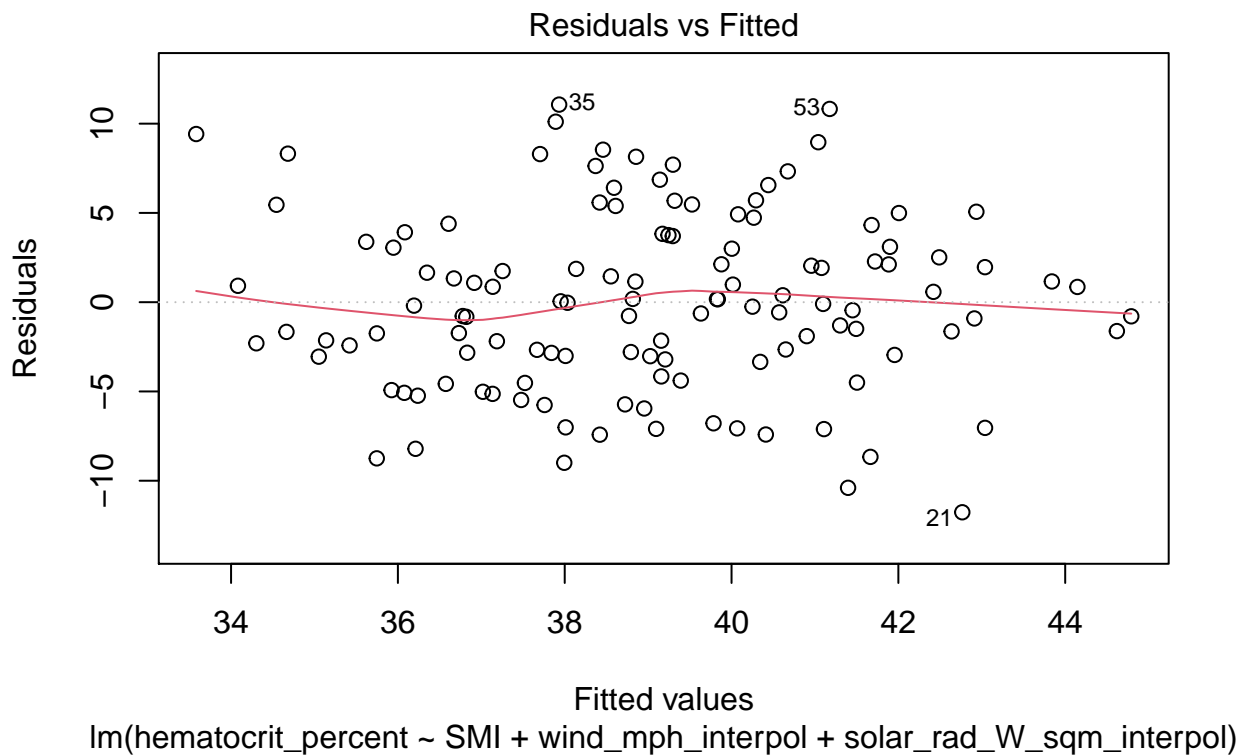
LM Conditions

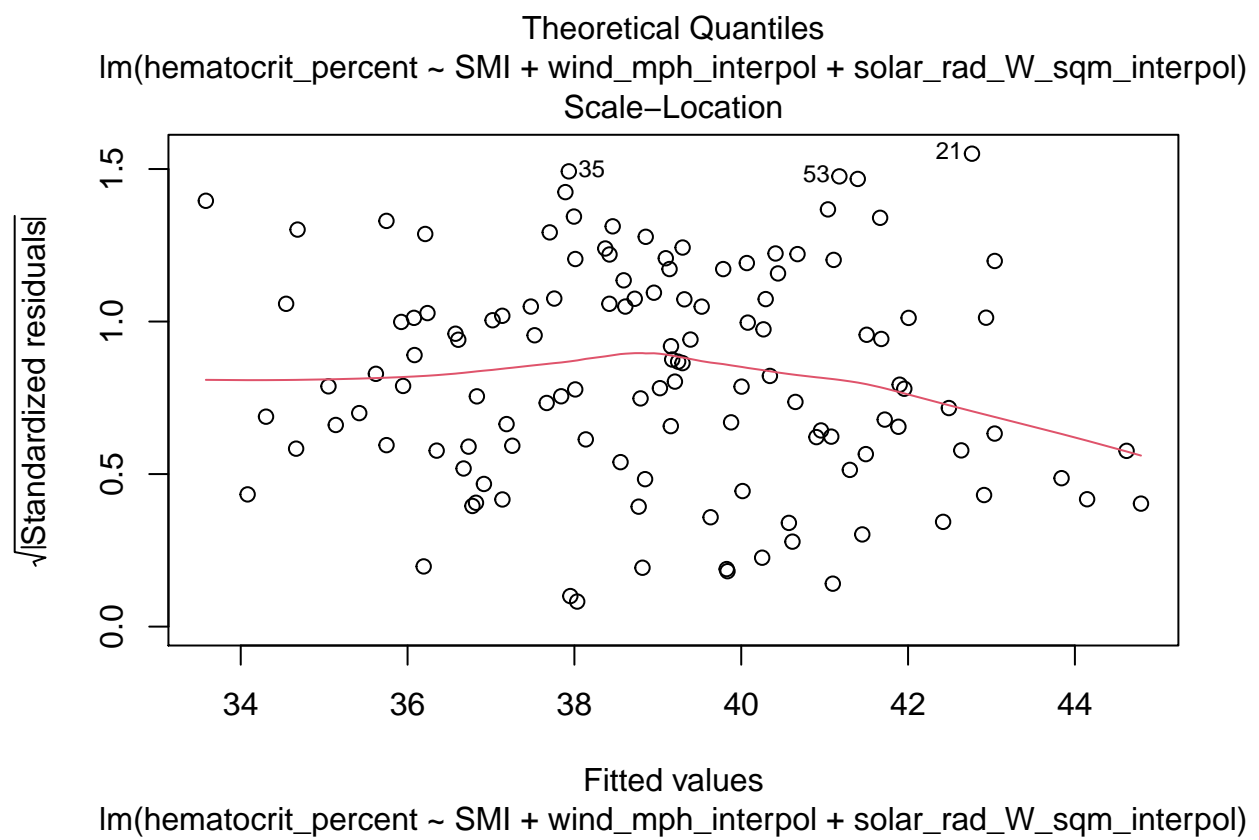
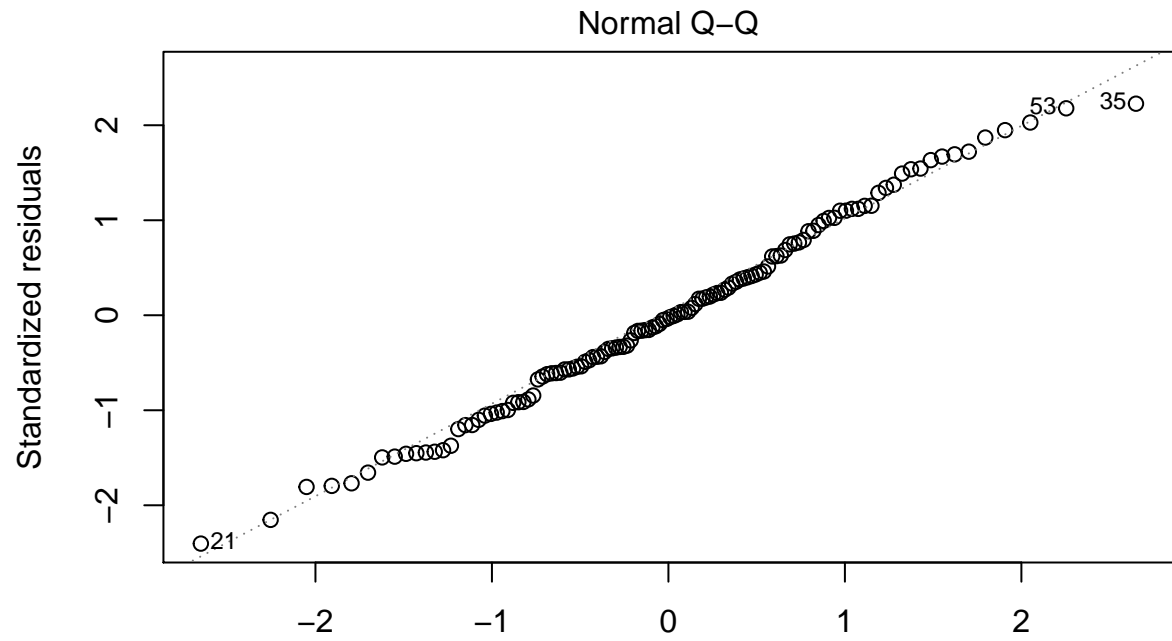
Check that the best model meets the criteria for linear regression and has no collinearity.

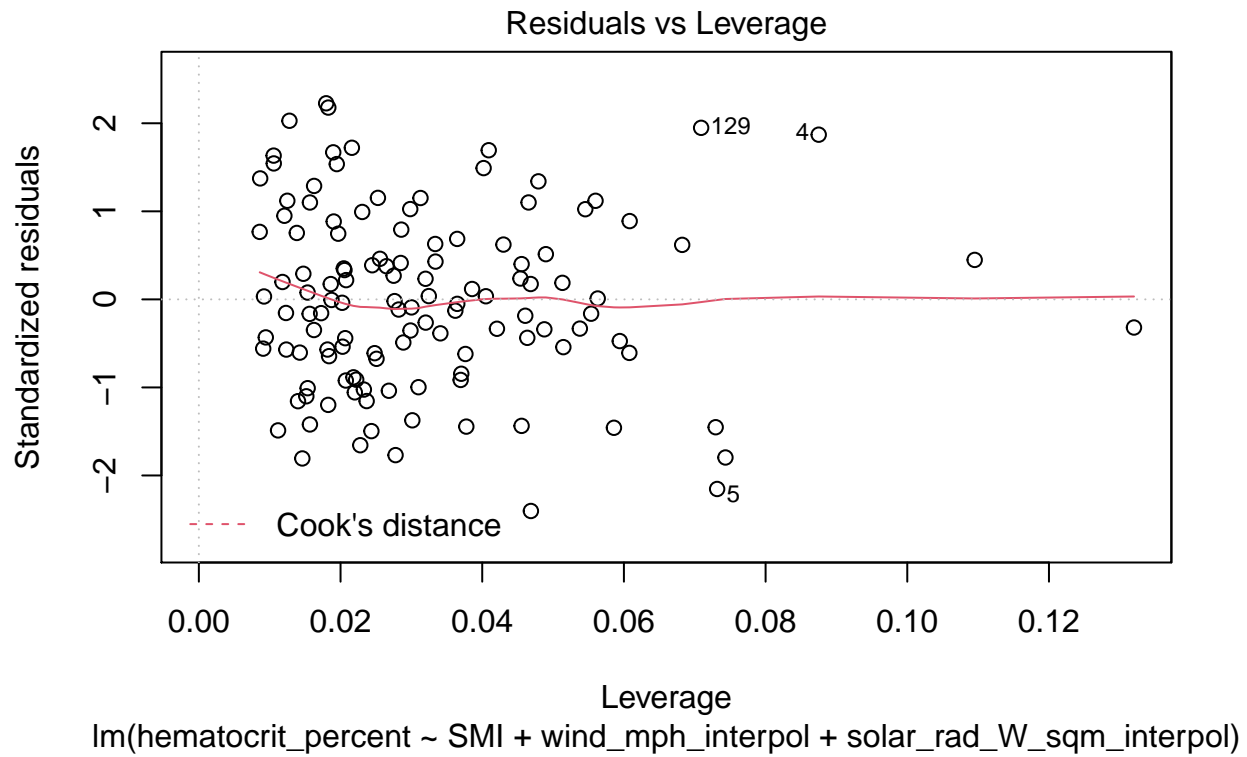
```
vif(hct_mod6)
```

```
##                SMI      wind_mph_interpol solar_rad_W_sqm_interpol
##                1.124966                1.117346                1.008219
```

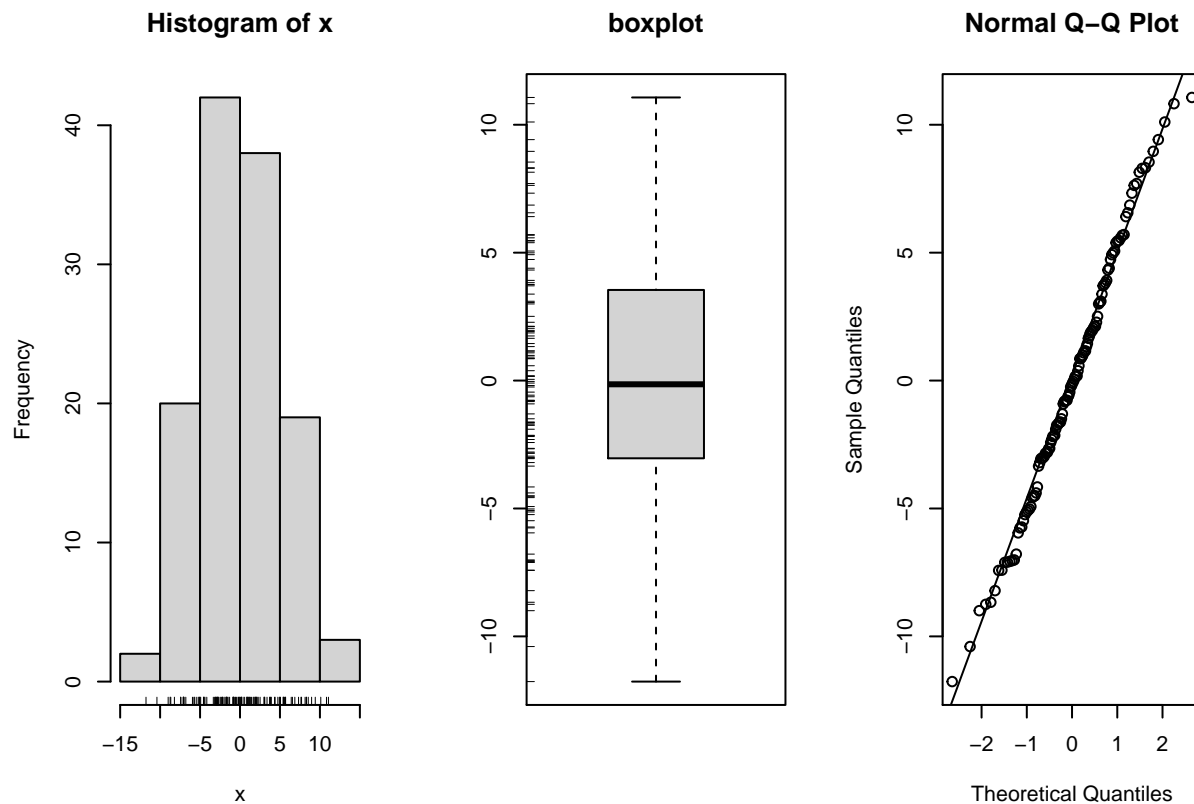
```
plot(hct_mod6)
```







```
simple.eda(residuals(hct_mod6))
```



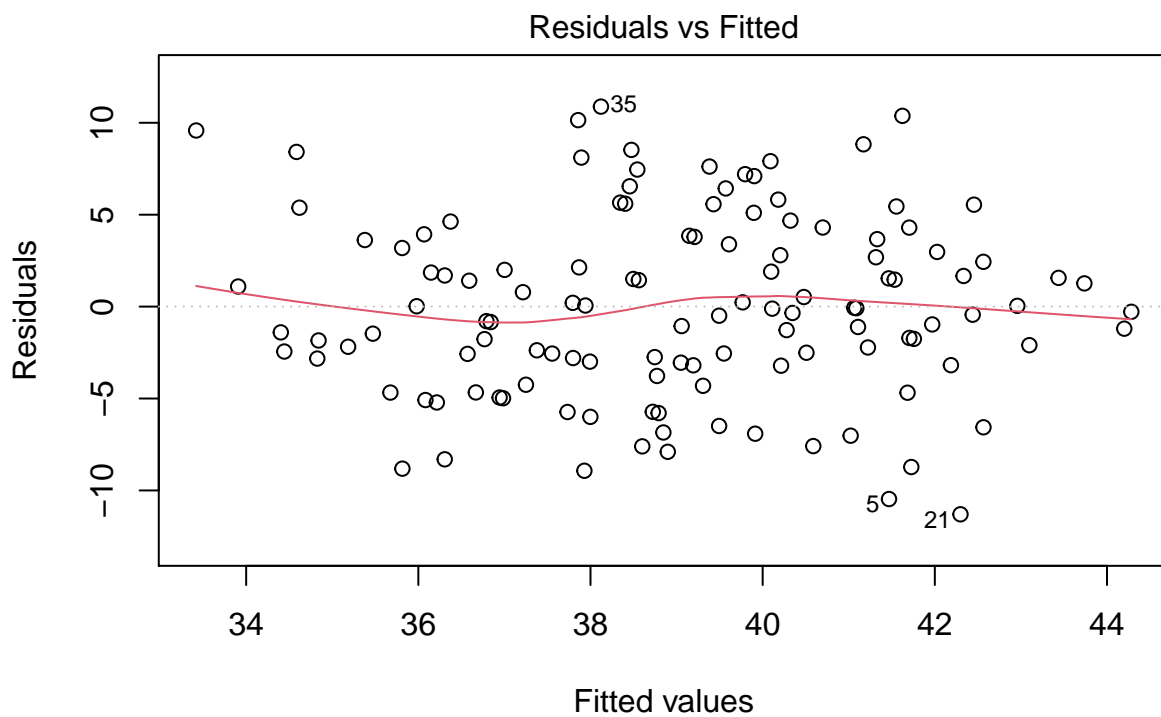
```
shapiro.test(residuals(hct_mod6))
```

```
##  
## Shapiro-Wilk normality test  
##  
## data: residuals(hct_mod6)  
## W = 0.9913, p-value = 0.6322
```

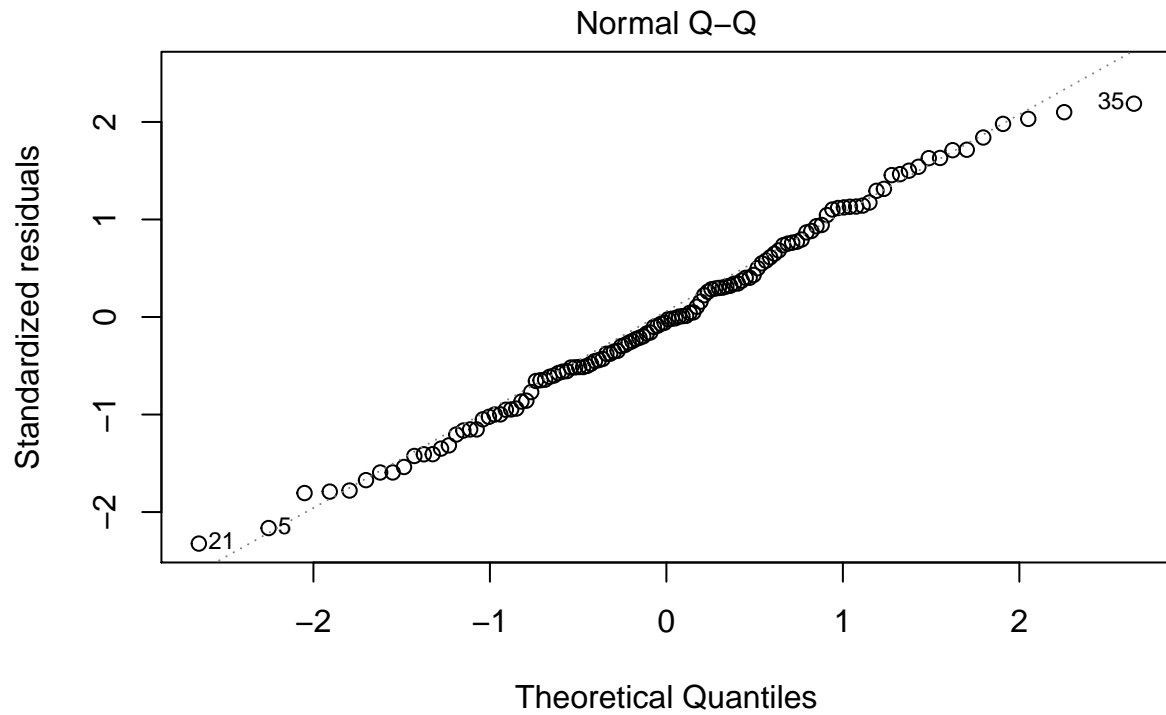
```
vif(hct_mod5)
```

```
##                SMI                VPD_kPa_int        wind_mph_interpol  
##          1.131865          4.010251          3.977224  
## solar_rad_W_sqm_interpol  
##          1.064119
```

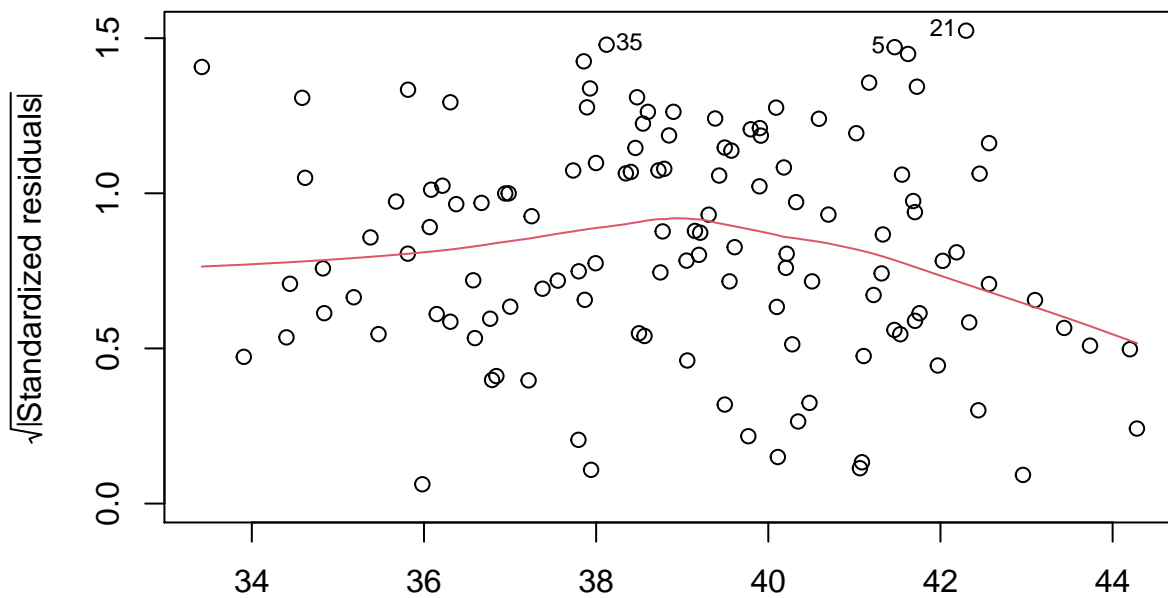
```
plot(hct_mod5)
```



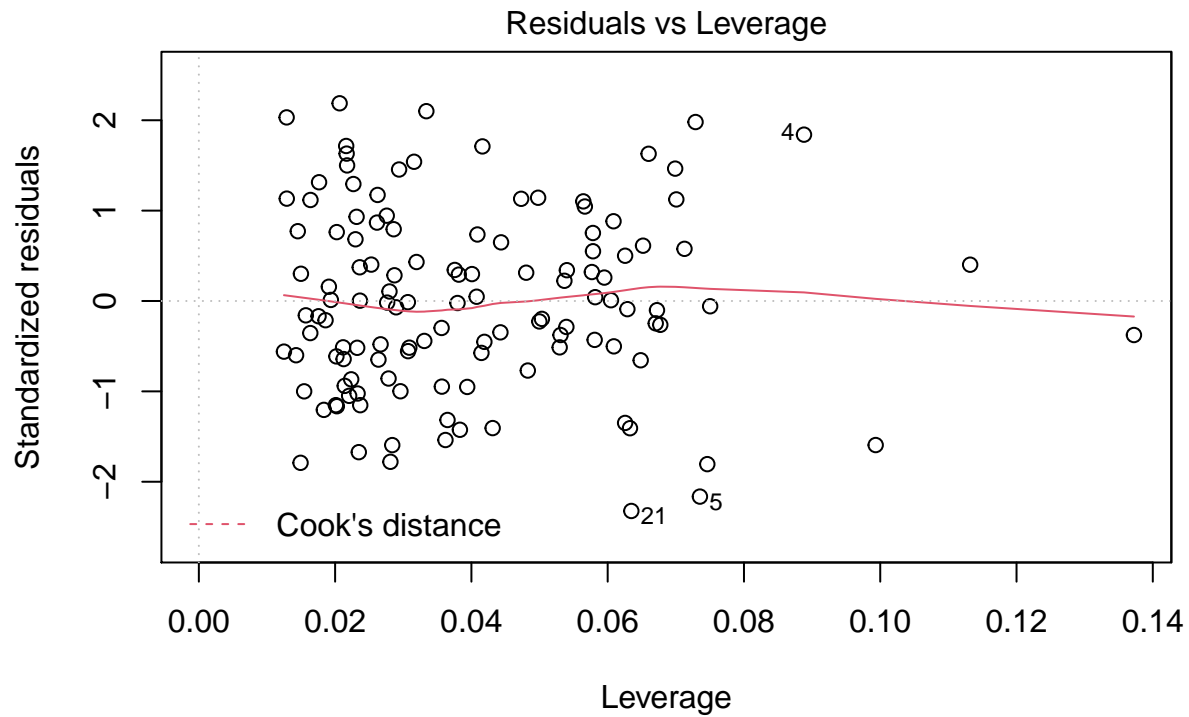
lm(hematocrit_percent ~ SMI + VPD_kPa_int + wind_mph_interpol + solar_rad_W ..



lm(hematocrit_percent ~ SMI + VPD_kPa_int + wind_mph_interpol + solar_rad_W ..
Scale-Location



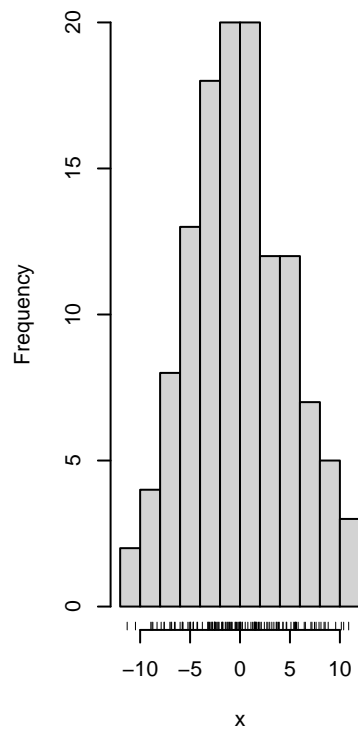
lm(hematocrit_percent ~ SMI + VPD_kPa_int + wind_mph_interpol + solar_rad_W ..



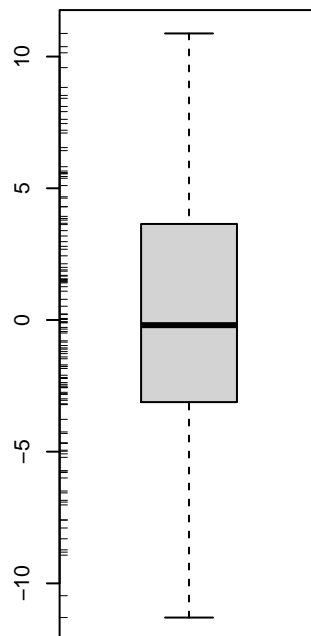
lm(hematocrit_percent ~ SMI + VPD_kPa_int + wind_mph_interpol + solar_rad_W ..

```
simple.eda(residuals(hct_mod5))
```

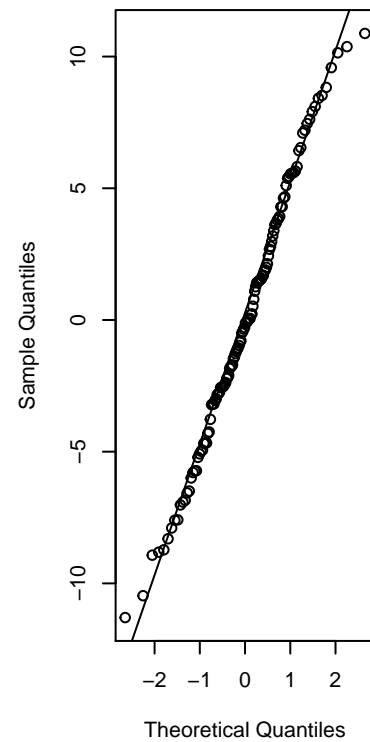
Histogram of x



boxplot



Normal Q-Q Plot



```
shapiro.test(residuals(hct_mod5))
```

```
##  
## Shapiro-Wilk normality test  
##  
## data: residuals(hct_mod5)  
## W = 0.9903, p-value = 0.5368
```

Everything is almost perfect.

Export

```
write.csv(hct_AICc, "./results_statistics/capture_hct_mod_rankings.csv")  
write.csv(broom.mixed::tidy(hct_mod6),  
          "./results_statistics/capture_hct_best_mod1.csv")  
write.csv(broom.mixed::tidy(hct_mod5),  
          "./results_statistics/capture_hct_best_mod2.csv")
```

Osmolality

Models

Since there are large differences in osmolality by date, but we are interested in what's different among dates, rather than the capture date itself, we will include that as a random effect in the model.

We would also include whether or not a blood sample is hemolyzed as a random effect, but only 11 of the almost 150 samples were hemolyzed, so we will assume that any potential effects will be undetectable and/or overshadowed. We do not have concern about using those points.

First, start with a full model with every probable potential predictor in it, then check for multicollinearity.

```
osml_mod1 <- lme4::lmer(data = dat,  
                       # response variable  
                       osmolality_mmol_kg_mean ~  
                       # body size  
                       mass_g + SVL_mm + SMI +  
                       # blood sample traits  
                       hematocrit_percent +  
                       # weather at the time of capture  
                       temp_C_interpol * VPD_kPa_int +  
                       wind_mph_interpol + solar_rad_W_sqm_interpol +  
                       (1|capture_date))  
osml_mod1_VIFs <- data.frame(VIF = car::vif(osml_mod1)) %>%  
  arrange(desc(VIF))  
osml_mod1_VIFs
```

##	VIF
## VPD_kPa_int	230.770824
## mass_g	157.598336
## SVL_mm	134.421136
## temp_C_interpol:VPD_kPa_int	74.795939
## temp_C_interpol	74.150446
## SMI	68.079595
## solar_rad_W_sqm_interpol	4.986184
## wind_mph_interpol	1.733699
## hematocrit_percent	1.144594

VPD and temperature introduce a lot of collinearity, so start by dropping their interaction:

```
osml_mod2 <- lme4::lmer(data = dat,
  # response variable
  osmolality_mmol_kg_mean ~
  # body size
  mass_g + SVL_mm + SMI +
  # blood sample traits
  hematocrit_percent +
  # weather at the time of capture
  temp_C_interpol + VPD_kPa_int +
  wind_mph_interpol + solar_rad_W_sqm_interpol +
  (1|capture_date))
osml_mod2_VIFs <- data.frame(VIF = car::vif(osml_mod2)) %>%
  arrange(desc(VIF))
osml_mod2_VIFs

##              VIF
## mass_g          157.205557
## SVL_mm          133.945843
## SMI              67.851081
## temp_C_interpol  25.018579
## VPD_kPa_int      17.519956
## solar_rad_W_sqm_interpol  3.794844
## wind_mph_interpol  1.287093
## hematocrit_percent  1.135184

drop1(osml_mod2)

## Single term deletions
##
## Model:
## osmolality_mmol_kg_mean ~ mass_g + SVL_mm + SMI + hematocrit_percent +
##   temp_C_interpol + VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##   (1 | capture_date)
##              npar      AIC
## <none>              1026.6
## mass_g              1 1025.3
## SVL_mm              1 1025.5
## SMI                 1 1025.2
## hematocrit_percent  1 1024.7
## temp_C_interpol     1 1026.0
## VPD_kPa_int         1 1026.6
## wind_mph_interpol   1 1025.0
## solar_rad_W_sqm_interpol  1 1026.5
```

Drop mass next, since it's extremely collinear and we get slightly better AIC by dropping mass compared to SVL:

```
osml_mod3 <- lme4::lmer(data = dat,
  # response variable
  osmolality_mmol_kg_mean ~
  # body size
  SVL_mm + SMI +
  # blood sample traits
  hematocrit_percent +
```



```

# weather at the time of capture
temp_C_interpol + VPD_kPa_int +
wind_mph_interpol + solar_rad_W_sqm_interpol +
(1|capture_date))
osml_mod3_VIFs <- data.frame(VIF = car::vif(osml_mod3)) %>%
  arrange(desc(VIF))
osml_mod3_VIFs

##                VIF
## temp_C_interpol  24.145020
## VPD_kPa_int      17.159996
## solar_rad_W_sqm_interpol  3.673673
## wind_mph_interpol  1.277514
## SMI               1.141551
## hematocrit_percent  1.135063
## SVL_mm            1.065997

drop1(osml_mod3)

## Single term deletions
##
## Model:
## osmolality_mmol_kg_mean ~ SVL_mm + SMI + hematocrit_percent +
##   temp_C_interpol + VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##   (1 | capture_date)
##               npar    AIC
## <none>                1025.3
## SVL_mm                1 1026.5
## SMI                   1 1023.5
## hematocrit_percent    1 1023.3
## temp_C_interpol       1 1024.3
## VPD_kPa_int           1 1025.0
## wind_mph_interpol     1 1023.6
## solar_rad_W_sqm_interpol 1 1025.7

Temperature is still introducing a lot of multicollinearity, so drop:
osml_mod4 <- lme4::lmer(data = dat,
  # response variable
  osmolality_mmol_kg_mean ~
  # body size
  SVL_mm + SMI +
  # blood sample traits
  hematocrit_percent +
  # weather at the time of capture
  VPD_kPa_int +
  wind_mph_interpol + solar_rad_W_sqm_interpol +
  (1|capture_date))
osml_mod4_VIFs <- data.frame(VIF = car::vif(osml_mod4)) %>%
  arrange(desc(VIF))
osml_mod4_VIFs

##                VIF
## VPD_kPa_int      2.679148
## solar_rad_W_sqm_interpol 2.423187
## wind_mph_interpol  1.190631

```

```

## SMI                                1.138075
## hematocrit_percent                 1.133355
## SVL_mm                             1.064398
summary(osml_mod4)

## Linear mixed model fit by REML ['lmerMod']
## Formula: osmolality_mmol_kg_mean ~ SVL_mm + SMI + hematocrit_percent +
##          VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##          (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 1000.8
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3276 -0.6763 -0.0915  0.6145  3.1713
##
## Random effects:
##   Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 290.4    17.04
## Residual              179.4    13.39
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   287.94425   34.47818   8.351
## SVL_mm         0.67238    0.38976   1.725
## SMI           -0.49620    1.27272  -0.390
## hematocrit_percent 0.04670    0.24779   0.188
## VPD_kPa_int   -3.55091    3.96947  -0.895
## wind_mph_interpol -0.15522    1.19310  -0.130
## solar_rad_W_sqm_interpol 0.03473    0.01316   2.639
##
## Correlation of Fixed Effects:
##              (Intr) SVL_mm SMI    hmtcr_ VPD_P_ wnd_m_
## SVL_mm       -0.836
## SMI          -0.555  0.223
## hmtcrt_prcn -0.114 -0.025 -0.244
## VPD_kPa_int  0.056  0.043  0.084 -0.104
## wnd_mph_ntr -0.058 -0.035  0.012 -0.145 -0.290
## slr_rd_W_s_ -0.136 -0.072  0.021 -0.014 -0.742  0.093
drop1(osml_mod4)

## Single term deletions
##
## Model:
## osmolality_mmol_kg_mean ~ SVL_mm + SMI + hematocrit_percent +
##          VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##          (1 | capture_date)
##              npar      AIC
## <none>              1024.3
## SVL_mm              1 1025.4
## SMI                  1 1022.5

```

```
## hematocrit_percent      1 1022.4
## VPD_kPa_int             1 1023.2
## wind_mph_interpol       1 1022.4
## solar_rad_W_sqm_interpol 1 1029.8
```

Great, VIFs are well-within acceptable ranges. Now we can start backwards model selection.

Start by dropping wind:

```
osml_mod5 <- lme4::lmer(data = dat,
                        # response variable
                        osmolality_mmol_kg_mean ~
                        # body size
                        SVL_mm + SMI +
                        # blood sample traits
                        hematocrit_percent +
                        # weather at the time of capture
                        VPD_kPa_int + solar_rad_W_sqm_interpol +
                        (1|capture_date))
summary(osml_mod5)

## Linear mixed model fit by REML ['lmerMod']
## Formula: osmolality_mmol_kg_mean ~ SVL_mm + SMI + hematocrit_percent +
##       VPD_kPa_int + solar_rad_W_sqm_interpol + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 1003
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3330 -0.6846 -0.1048  0.6265  3.1859
##
## Random effects:
## Groups      Name                Variance Std.Dev.
## capture_date (Intercept) 285.8      16.91
## Residual              178.0      13.34
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   287.68166   34.27361   8.394
## SVL_mm         0.67062    0.38793   1.729
## SMI           -0.49383    1.26746  -0.390
## hematocrit_percent 0.04192    0.24416   0.172
## VPD_kPa_int   -3.70219    3.77636  -0.980
## solar_rad_W_sqm_interpol 0.03489    0.01303   2.677
##
## Correlation of Fixed Effects:
##              (Intr) SVL_mm SMI    hmtcr_ VPD_P_
## SVL_mm       -0.840
## SMI          -0.555  0.223
## hmtcrt_prcn -0.124 -0.030 -0.245
## VPD_kPa_int  0.041  0.034  0.092 -0.154
## slr_rd_W_s_ -0.131 -0.069  0.020  0.000 -0.749
```

```
drop1(osml_mod5)
```

```
## Single term deletions
##
## Model:
## osmolality_mmol_kg_mean ~ SVL_mm + SMI + hematocrit_percent +
##   VPD_kPa_int + solar_rad_W_sqm_interpol + (1 | capture_date)
##               npar    AIC
## <none>                1022.4
## SVL_mm                1 1023.5
## SMI                   1 1020.6
## hematocrit_percent    1 1020.5
## VPD_kPa_int           1 1021.6
## solar_rad_W_sqm_interpol 1 1027.9
```

Drop hematocrit:

```
osml_mod6 <- lme4::lmer(data = dat,
                        # response variable
                        osmolality_mmol_kg_mean ~
                        # body size
                        SVL_mm + SMI +
                        # weather at the time of capture
                        VPD_kPa_int + solar_rad_W_sqm_interpol +
                        (1|capture_date))
summary(osml_mod6)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula:
## osmolality_mmol_kg_mean ~ SVL_mm + SMI + VPD_kPa_int + solar_rad_W_sqm_interpol +
##   (1 | capture_date)
##   Data: dat
##
## REML criterion at convergence: 1002
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3426 -0.6858 -0.0985  0.6211  3.1839
##
## Random effects:
##   Groups       Name             Variance Std.Dev.
##   capture_date (Intercept) 284.0      16.85
##   Residual                176.5      13.29
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##               Estimate Std. Error t value
## (Intercept)    288.41208   33.87099   8.515
## SVL_mm          0.67263    0.38616   1.742
## SMI            -0.44051    1.22367  -0.360
## VPD_kPa_int    -3.60191    3.71759  -0.969
## solar_rad_W_sqm_interpol 0.03489    0.01298   2.687
##
## Correlation of Fixed Effects:
##              (Intr) SVL_mm SMI      VPD_P_
```

```
## SVL_mm      -0.851
## SMI          -0.609  0.223
## VPD_kPa_int  0.023  0.030  0.056
## slr_rd_W_s_ -0.132 -0.069  0.021 -0.758

drop1(osml_mod6)

## Single term deletions
##
## Model:
## osmolality_mmol_kg_mean ~ SVL_mm + SMI + VPD_kPa_int + solar_rad_W_sqm_interpol +
## (1 | capture_date)
##               npar      AIC
## <none>                1020.5
## SVL_mm                1 1021.5
## SMI                   1 1018.6
## VPD_kPa_int           1 1019.6
## solar_rad_W_sqm_interpol 1 1025.9
```

Drop SMI:

```
osml_mod7 <- lme4::lmer(data = dat,
                        # response variable
                        osmolality_mmol_kg_mean ~
                        # body size
                        SVL_mm +
                        # weather at the time of capture
                        VPD_kPa_int + solar_rad_W_sqm_interpol +
                        (1|capture_date))
summary(osml_mod7)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula:
## osmolality_mmol_kg_mean ~ SVL_mm + VPD_kPa_int + solar_rad_W_sqm_interpol +
## (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 1004.4
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3370 -0.6953 -0.0963  0.6058  3.2280
##
## Random effects:
## Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 282.0    16.79
## Residual          175.2    13.24
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##               Estimate Std. Error t value
## (Intercept)    280.98723   26.77002  10.496
## SVL_mm          0.70358    0.37510   1.876
## VPD_kPa_int     -3.52647    3.69840  -0.954
## solar_rad_W_sqm_interpol 0.03499    0.01293   2.705
##
```

```

## Correlation of Fixed Effects:
##           (Intr) SVL_mm VPD_P_
## SVL_mm      -0.925
## VPD_kPa_int  0.072  0.018
## slr_rd_W_s_ -0.151 -0.075 -0.761
drop1(osml_mod7)

## Single term deletions
##
## Model:
## osmolality_mmol_kg_mean ~ SVL_mm + VPD_kPa_int + solar_rad_W_sqm_interpol +
##      (1 | capture_date)
##               npar      AIC
## <none>                1018.6
## SVL_mm                1 1020.1
## VPD_kPa_int            1 1017.7
## solar_rad_W_sqm_interpol 1 1024.1

Drop VPD:
osml_mod8 <- lme4::lmer(data = dat,
                        # response variable
                        osmolality_mmol_kg_mean ~
                        # body size
                        SVL_mm +
                        # weather at the time of capture
                        solar_rad_W_sqm_interpol +
                        (1|capture_date))
summary(osml_mod8)

## Linear mixed model fit by REML ['lmerMod']
## Formula: osmolality_mmol_kg_mean ~ SVL_mm + solar_rad_W_sqm_interpol +
##      (1 | capture_date)
##      Data: dat
##
## REML criterion at convergence: 1009.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3259 -0.6885 -0.0971  0.5530  3.2419
##
## Random effects:
##      Groups       Name             Variance Std.Dev.
## capture_date (Intercept) 288.1       16.97
## Residual                175.0       13.23
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)    2.828e+02  2.671e+01  10.589
## SVL_mm          7.103e-01  3.748e-01   1.895
## solar_rad_W_sqm_interpol 2.560e-02  8.387e-03   3.053
##
## Correlation of Fixed Effects:
##           (Intr) SVL_mm

```

```
## SVL_mm      -0.928
## slr_rd_W_s_ -0.148 -0.095
```

Drop SVL:

```
osml_mod9 <- lme4::lmer(data = dat,
                        # response variable
                        osmolality_mmol_kg_mean ~
                        # weather at the time of capture
                        solar_rad_W_sqm_interpol +
                        (1|capture_date))
summary(osml_mod9)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula:
## osmolality_mmol_kg_mean ~ solar_rad_W_sqm_interpol + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 1013.2
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.4085 -0.6875 -0.1434  0.5776  3.3124
##
## Random effects:
## Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 277.1    16.65
## Residual                179.0    13.38
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)    3.297e+02  9.915e+00  33.256
## solar_rad_W_sqm_interpol 2.712e-02  8.444e-03   3.212
##
## Correlation of Fixed Effects:
##              (Intr)
## slr_rd_W_s_ -0.648
```

Lastly, compute null model:

```
osml_mod_null <- lme4::lmer(data = dat,
                           osmolality_mmol_kg_mean ~ 1 +
                           (1|capture_date))
summary(osml_mod_null)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: osmolality_mmol_kg_mean ~ 1 + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 1127.8
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.4072 -0.6642 -0.1005  0.5332  3.1645
##
```

```
## Random effects:
##   Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 262.1    16.19
## Residual          190.9    13.82
## Number of obs: 138, groups: capture_date, 5
##
## Fixed effects:
##           Estimate Std. Error t value
## (Intercept) 350.168      7.338  47.72
```

Selection

Compare models 4-9 and null.

```
osml_models <- list(osml_mod4, osml_mod5, osml_mod6,
                    osml_mod7, osml_mod8, osml_mod9,
                    osml_mod_null)
#specify model names
osml_mod_names <- c('(model 4) ~ Solar-C, SVL, VPD-C, SMI, Hct, Wind-C',
                    '(model 5) ~ Solar-C, SVL, VPD-C, SMI, Hct',
                    '(model 6) ~ Solar-C, SVL, VPD-C, SMI',
                    '(model 7) ~ Solar-C, SVL, VPD-C',
                    '(model 8) ~ Solar-C, SVL',
                    '(model 9) ~ Solar-C',
                    'null model')
#calculate AIC of each model
osml_AICc <- data.frame(aictab(cand.set = osml_models,
                              modnames = osml_mod_names))
```

```
## Warning in aictab.AIClmerMod(cand.set = osml_models, modnames = osml_mod_names):
## Model selection for fixed effects is only appropriate with ML estimation:
## REML (default) should only be used to select random effects for a constant set of fixed effects
```

```
osml_AICc
```

	Modnames	K	AICc	Delta_AICc
## 3	(model 6) ~ Solar-C, SVL, VPD-C, SMI	7	1016.982	0.0000000
## 4	(model 7) ~ Solar-C, SVL, VPD-C	6	1017.103	0.1204429
## 2	(model 5) ~ Solar-C, SVL, VPD-C, SMI, Hct	8	1020.225	3.2431270
## 5	(model 8) ~ Solar-C, SVL	5	1020.255	3.2728106
## 1	(model 4) ~ Solar-C, SVL, VPD-C, SMI, Hct, Wind-C	9	1020.348	3.3662143
## 6	(model 9) ~ Solar-C	4	1021.516	4.5333722
## 7	null model	3	1133.960	116.9777599

	ModelLik	AICcWt	Res.LL	Cum.Wt
## 3	1.000000e+00	3.812029e-01	-501.0083	0.3812029
## 4	9.415560e-01	3.589239e-01	-502.1923	0.7401267
## 2	1.975895e-01	7.532170e-02	-501.4865	0.8154484
## 5	1.946786e-01	7.421204e-02	-504.8732	0.8896605
## 1	1.857958e-01	7.082589e-02	-500.3847	0.9604864
## 6	1.036551e-01	3.951363e-02	-506.5897	1.0000000
## 7	3.968279e-26	1.512719e-26	-563.8904	1.0000000

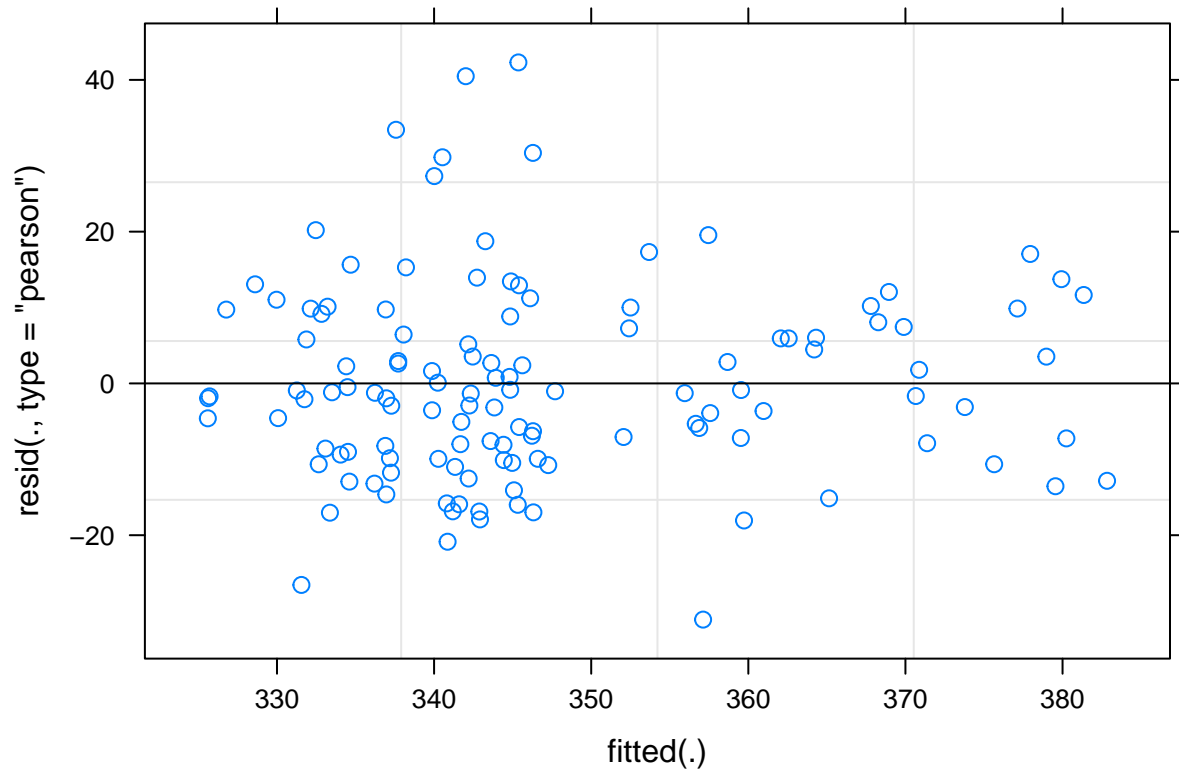
LM Conditions

Check residual plots and VIFs

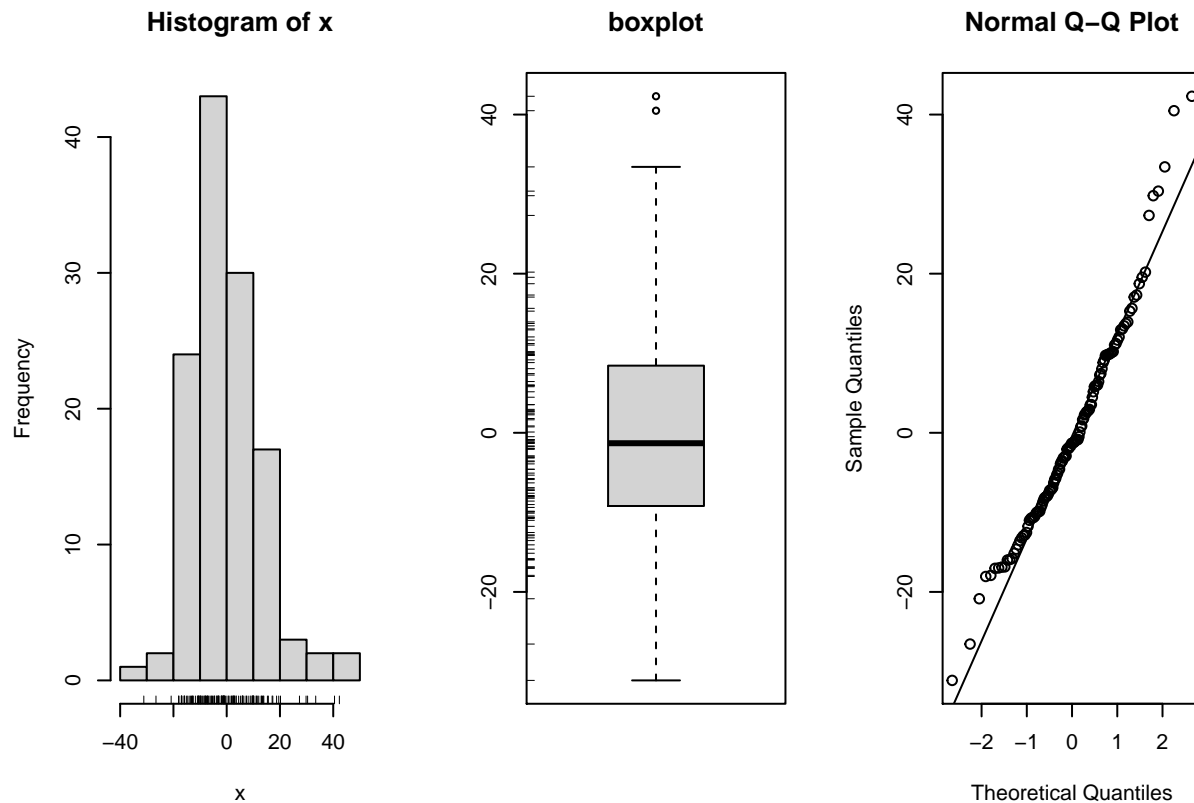

```
vif(osml_mod6)
```

```
##           SVL_mm           SMI           VPD_kPa_int  
##           1.062154        1.069446        2.391413  
## solar_rad_W_sqm_interpol  
##           2.397751
```

```
plot(osml_mod6)
```



```
simple.eda(residuals(osml_mod6))
```



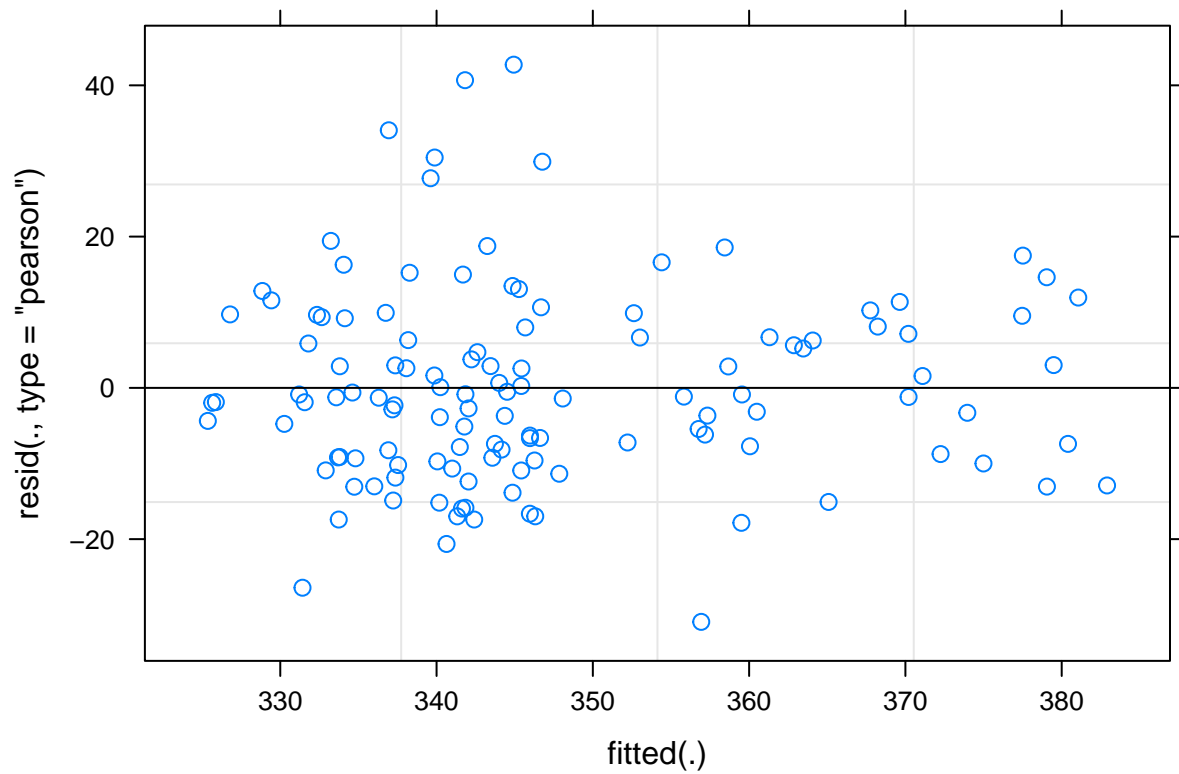
```
shapiro.test(residuals(osml_mod6))
```

```
##
##  Shapiro-Wilk normality test
##
## data:  residuals(osml_mod6)
## W = 0.96768, p-value = 0.004547
```

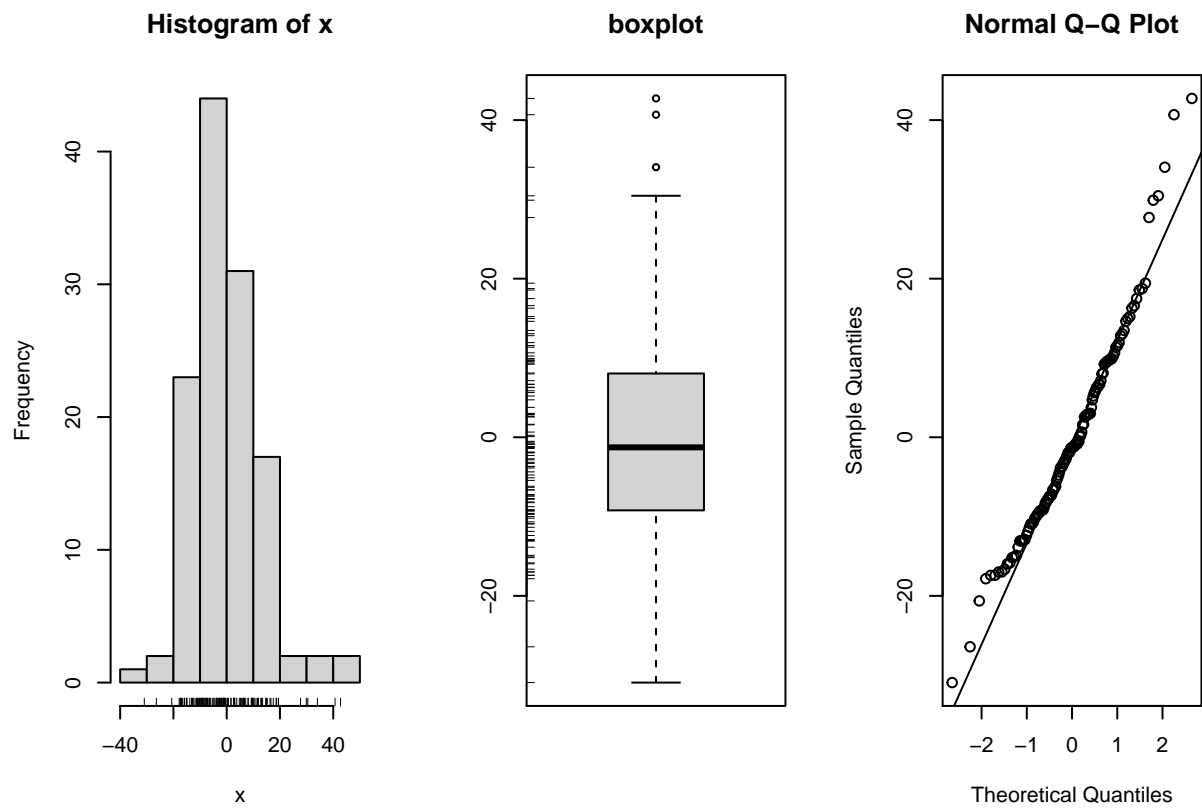
```
vif(osml_mod7)
```

```
##              SVL_mm              VPD_kPa_int solar_rad_W_sqm_interpol
##              1.009504              2.383988              2.396868
```

```
plot(osml_mod7)
```



```
simple.eda(residuals(osml_mod7))
```



```
shapiro.test(residuals(osml_mod7))
```

```
##
##  Shapiro-Wilk normality test
##
## data:  residuals(osml_mod7)
## W = 0.96557, p-value = 0.002973
```

There is no clear pattern in the residuals ~ fitted plot, so linearity seems satisfied. slight fanning, but equal error variance seems fine. Normality seems fine, even though the Shapiro-Wilk normality test is significant. VIFs essentially negligible.

Export

First, re-run for p-values:

```
osml_mod6p <- lmerTest::lmer(data = dat,
                             # response variable
                             osmolality_mmol_kg_mean ~
                             # body size
                             SVL_mm + SMI +
                             # weather at the time of capture
                             VPD_kPa_int + solar_rad_W_sqm_interpol +
                             (1|capture_date))
summary(osml_mod6p)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## osmolality_mmol_kg_mean ~ SVL_mm + SMI + VPD_kPa_int + solar_rad_W_sqm_interpol +
## (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 1002
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3426 -0.6858 -0.0985  0.6211  3.1839
##
## Random effects:
## Groups      Name                Variance Std.Dev.
## capture_date (Intercept) 284.0      16.85
## Residual                176.5      13.29
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)  288.41208   33.87099 117.08658   8.515 6.54e-14 ***
## SVL_mm        0.67263    0.38616 116.54146   1.742  0.0842 .
## SMI          -0.44051    1.22367 115.79045  -0.360  0.7195
## VPD_kPa_int  -3.60191    3.71759  15.08255  -0.969  0.3479
## solar_rad_W_sqm_interpol  0.03489    0.01298  34.46638   2.687  0.0110 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Correlation of Fixed Effects:
##           (Intr) SVL_mm SMI    VPD_P_
## SVL_mm      -0.851
## SMI          -0.609  0.223
## VPD_kPa_int  0.023  0.030  0.056
## slr_rd_W_s_ -0.132 -0.069  0.021 -0.758

osml_mod7p <- lmerTest::lmer(data = dat,
                             # response variable
                             osmolality_mmol_kg_mean ~
                             # body size
                             SVL_mm +
                             # weather at the time of capture
                             VPD_kPa_int + solar_rad_W_sqm_interpol +
                             (1|capture_date))

summary(osml_mod7p)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## osmolality_mmol_kg_mean ~ SVL_mm + VPD_kPa_int + solar_rad_W_sqm_interpol +
## (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 1004.4
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3370 -0.6953 -0.0963  0.6058  3.2280
##
## Random effects:
## Groups      Name                Variance Std.Dev.
## capture_date (Intercept) 282.0      16.79
## Residual              175.2      13.24
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)    280.98723   26.77002 109.79196  10.496  <2e-16 ***
## SVL_mm          0.70358    0.37510 117.63142   1.876   0.0632 .
## VPD_kPa_int     -3.52647    3.69840  14.98489  -0.954   0.3555
## solar_rad_W_sqm_interpol  0.03499    0.01293  34.53408   2.705   0.0105 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##           (Intr) SVL_mm VPD_P_
## SVL_mm      -0.925
## VPD_kPa_int  0.072  0.018
## slr_rd_W_s_ -0.151 -0.075 -0.761
```

Save the model output.

```
write.csv(broom.mixed::tidy(osml_mod6p),
          "./results_statistics/capture_osml_best_model1.csv")
write.csv(broom.mixed::tidy(osml_mod7p),
```

```

      "./results_statistics/capture_osml_best_model2.csv")
write.csv(osml_AICc, "./results_statistics/capture_osml_mod_rankings.csv")

```

To report in paper:

The best models to predict the variation in baseline plasma osmolality included SVL, SMI, VPD, and solar radiation at the time of capture as fixed effects. Date was included as a random effect. The final model had acceptable LM conditions. The full model included mass, SVL, SMI, percent hematocrit, and temperature, VPD, wind speed, and solar radiation at the time of capture, with date as a random effect.

CEWL

It looks like there are meaningful differences in CEWL across individuals/dates (probably confounded), and based on cloacal temp, capture temp, capture VPD, capture wind, and capture solar radiation.

I didn't calculate holding time!!!!!!!!!!!!!!!!!!!!

Models

Start with the full model of all potential predictor variables. We will again include date as a random effect. Individual ID is not included as a random effect bc each lizard only has one set of capture observations.

When we have this many variables, it's extremely important to start with checking for multicollinearity.

```

CEWL_mod1 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # essential covariate
  cloacal_temp_C +
  # body size
  mass_g + SVL_mm + SMI +
  # blood
  osmolality_mmol_kg_mean + hematocrit_percent +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
  # weather at the time of capture
  temp_C_interpol * VPD_kPa_int +
  wind_mph_interpol + solar_rad_W_sqm_interpol +
  # time between capture and measurements
  hold_time_hr +
  (1|capture_date))
CEWL_mod1_VIFs <- data.frame(VIF = car::vif(CEWL_mod1)) %>%
  arrange(desc(VIF))
CEWL_mod1_VIFs

```

```

##                                VIF
## VPD_kPa_int                    856.642445
## temp_C_interpol:VPD_kPa_int    272.795884
## temp_C_interpol                171.003017
## mass_g                        160.346694
## SVL_mm                        138.353857
## SMI                            69.684728
## msmt_VPD_kPa                  30.887901
## msmt_temp_C                   12.625546
## solar_rad_W_sqm_interpol       4.956034
## wind_mph_interpol              3.480824

```

```
## hold_time_hr          1.961889
## hematocrit_percent    1.220159
## osmolality_mmol_kg_mean 1.179014
## cloacal_temp_C        1.152096
```

```
drop1(CEWL_mod1)
```

```
## Single term deletions
```

```
##
```

```
## Model:
```

```
## CEWL_g_m2h_mean ~ cloacal_temp_C + mass_g + SVL_mm + SMI + osmolality_mmol_kg_mean +
##      hematocrit_percent + msmt_temp_C + msmt_VPD_kPa + temp_C_interpol *
##      VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##      hold_time_hr + (1 | capture_date)
```

```
##              npar      AIC
## <none>              653.45
## cloacal_temp_C      1 651.49
## mass_g              1 654.50
## SVL_mm              1 654.36
## SMI                 1 654.13
## osmolality_mmol_kg_mean 1 663.51
## hematocrit_percent  1 651.54
## msmt_temp_C         1 667.57
## msmt_VPD_kPa        1 658.33
## wind_mph_interpol   1 658.51
## solar_rad_W_sqm_interpol 1 652.77
## hold_time_hr        1 658.03
## temp_C_interpol:VPD_kPa_int 1 651.89
```

Just as for osmolality, VPD and temperature introduce a lot of collinearity. Start with dropping their interaction:

```
CEWL_mod2 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # essential covariate
  cloacal_temp_C +
  # body size
  mass_g + SVL_mm + SMI +
  # blood
  osmolality_mmol_kg_mean + hematocrit_percent +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
  # weather at the time of capture
  temp_C_interpol + VPD_kPa_int +
  wind_mph_interpol + solar_rad_W_sqm_interpol +
  # time between capture and measurements
  hold_time_hr +
  (1|capture_date))
```

```
CEWL_mod2_VIFs <- data.frame(VIF = car::vif(CEWL_mod2)) %>%
  arrange(desc(VIF))
CEWL_mod2_VIFs
```

```
##              VIF
## mass_g        160.075943
## SVL_mm        138.388861
```

```
## SMI                                69.706986
## temp_C_interpol                    34.379121
## VPD_kPa_int                        30.735088
## msmt_VPD_kPa                       11.232713
## msmt_temp_C                        5.060439
## solar_rad_W_sqm_interpol           4.618568
## wind_mph_interpol                  3.757447
## hold_time_hr                       1.738348
## hematocrit_percent                 1.213102
## osmolality_mmol_kg_mean            1.182087
## cloacal_temp_C                     1.151022
```

```
drop1(CEWL_mod2)
```

```
## Single term deletions
```

```
##
```

```
## Model:
```

```
## CEWL_g_m2h_mean ~ cloacal_temp_C + mass_g + SVL_mm + SMI + osmolality_mmol_kg_mean +
##      hematocrit_percent + msmt_temp_C + msmt_VPD_kPa + temp_C_interpol +
##      VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##      hold_time_hr + (1 | capture_date)
##               npar      AIC
## <none>                651.89
## cloacal_temp_C        1 649.91
## mass_g                 1 652.94
## SVL_mm                 1 652.82
## SMI                    1 652.60
## osmolality_mmol_kg_mean 1 662.07
## hematocrit_percent     1 650.01
## msmt_temp_C            1 677.25
## msmt_VPD_kPa           1 661.94
## temp_C_interpol        1 658.23
## VPD_kPa_int            1 660.20
## wind_mph_interpol      1 656.88
## solar_rad_W_sqm_interpol 1 651.88
## hold_time_hr           1 656.12
```

MUCH better. Drop SVL next:

```
CEWL_mod3 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # essential covariate
  cloacal_temp_C +
  # body size
  mass_g + SMI +
  # blood
  osmolality_mmol_kg_mean + hematocrit_percent +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
  # weather at the time of capture
  temp_C_interpol + VPD_kPa_int +
  wind_mph_interpol + solar_rad_W_sqm_interpol +
  # time between capture and measurements
  hold_time_hr +
  (1|capture_date))
```



```
CEWL_mod3_VIFs <- data.frame(VIF = car::vif(CEWL_mod3)) %>%
  arrange(desc(VIF))
CEWL_mod3_VIFs
```

```
##                               VIF
## temp_C_interpol              32.331655
## VPD_kPa_int                  29.352386
## msmt_VPD_kPa                 10.735526
## msmt_temp_C                  5.021935
## solar_rad_W_sqm_interpol     4.459750
## wind_mph_interpol            3.641960
## hold_time_hr                 1.716197
## SMI                           1.376741
## mass_g                       1.347833
## hematocrit_percent           1.212158
## osmolality_mmol_kg_mean      1.167705
## cloacal_temp_C               1.143432
```

```
drop1(CEWL_mod3)
```

```
## Single term deletions
```

```
##
```

```
## Model:
```

```
## CEWL_g_m2h_mean ~ cloacal_temp_C + mass_g + SMI + osmolality_mmol_kg_mean +
##   hematocrit_percent + msmt_temp_C + msmt_VPD_kPa + temp_C_interpol +
##   VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##   hold_time_hr + (1 | capture_date)
```

```
##                               npar    AIC
## <none>                          652.82
## cloacal_temp_C                   1 650.82
## mass_g                           1 651.04
## SMI                              1 650.93
## osmolality_mmol_kg_mean          1 663.97
## hematocrit_percent               1 650.93
## msmt_temp_C                      1 677.25
## msmt_VPD_kPa                     1 661.74
## temp_C_interpol                  1 657.40
## VPD_kPa_int                      1 659.57
## wind_mph_interpol                1 657.23
## solar_rad_W_sqm_interpol         1 652.00
## hold_time_hr                     1 656.12
```

```
Next drop temperature at the time of capture:
```

```
CEWL_mod4 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # essential covariate
  cloacal_temp_C +
  # body size
  mass_g + SMI +
  # blood
  osmolality_mmol_kg_mean + hematocrit_percent +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
```

```

# weather at the time of capture
VPD_kPa_int +
wind_mph_interpol + solar_rad_W_sqm_interpol +
# time between capture and measurements
hold_time_hr +
(1|capture_date))
CEWL_mod4_VIFs <- data.frame(VIF = car::vif(CEWL_mod4)) %>%
  arrange(desc(VIF))
CEWL_mod4_VIFs

##              VIF
## msmt_VPD_kPa    8.374622
## VPD_kPa_int     4.949436
## msmt_temp_C     4.856765
## solar_rad_W_sqm_interpol 3.215116
## wind_mph_interpol 2.950819
## hold_time_hr    1.692077
## SMI              1.363902
## mass_g           1.342405
## hematocrit_percent 1.203612
## osmolality_mmol_kg_mean 1.150762
## cloacal_temp_C   1.108965

summary(CEWL_mod4)

## Linear mixed model fit by REML ['lmerMod']
## Formula:
## CEWL_g_m2h_mean ~ cloacal_temp_C + mass_g + SMI + osmolality_mmol_kg_mean +
##   hematocrit_percent + msmt_temp_C + msmt_VPD_kPa + VPD_kPa_int +
##   wind_mph_interpol + solar_rad_W_sqm_interpol + hold_time_hr +
##   (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 644.8
##
## Scaled residuals:
##   Min       1Q   Median       3Q      Max
## -2.1622 -0.6141 -0.0049  0.4957  3.3967
##
## Random effects:
##   Groups       Name             Variance Std.Dev.
##   capture_date (Intercept) 4.956      2.226
##   Residual              9.386      3.064
## Number of obs: 124, groups:  capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   -1.663e+02  3.531e+01  -4.709
## cloacal_temp_C    1.330e-01  4.045e-01   0.329
## mass_g         -1.465e-01  2.012e-01  -0.728
## SMI             -1.918e-02  3.177e-01  -0.060
## osmolality_mmol_kg_mean  7.173e-02  2.080e-02   3.448
## hematocrit_percent  1.522e-02  5.822e-02   0.261
## msmt_temp_C      7.148e+00  1.501e+00   4.763

```

```

## msmt_VPD_kPa          -1.893e+01  6.283e+00  -3.012
## VPD_kPa_int           -1.576e+00  1.057e+00  -1.490
## wind_mph_interpol      8.408e-01  3.434e-01  2.449
## solar_rad_W_sqm_interpol 2.313e-03  3.463e-03  0.668
## hold_time_hr           6.129e-01  3.234e-01  1.895
##
## Correlation of Fixed Effects:
##          (Intr) clc__C mass_g SMI      osm___ hmtcr_ msm__C m_VPD_ VPD_P_
## clocl_tmp_C -0.309
## mass_g      -0.023  0.072
## SMI          -0.100 -0.063 -0.443
## osmlly_m__ -0.299  0.038 -0.125  0.117
## hmtcr_prcn  0.027 -0.012 -0.021 -0.202 -0.019
## msmt_temp_C -0.899  0.043  0.016  0.022  0.095 -0.118
## msmt_VPD_kP  0.494 -0.176 -0.049  0.019 -0.057  0.232 -0.699
## VPD_kPa_int  0.126  0.024  0.011  0.037  0.028 -0.202  0.056 -0.526
## wnd_mph_ntr -0.329  0.193  0.048  0.008  0.110 -0.229  0.391 -0.624  0.035
## slr_rd_W_s_  0.352  0.024  0.002  0.015 -0.221  0.090 -0.505  0.588 -0.636
## hold_tim_hr  0.113  0.245  0.207 -0.085  0.063 -0.031 -0.155 -0.214  0.170
##          wnd_m_ s__W__
## clocl_tmp_C
## mass_g
## SMI
## osmlly_m__
## hmtcr_prcn
## msmt_temp_C
## msmt_VPD_kP
## VPD_kPa_int
## wnd_mph_ntr
## slr_rd_W_s_ -0.174
## hold_tim_hr  0.413  0.067

```

```
drop1(CEWL_mod4)
```

```

## Single term deletions
##
## Model:
## CEWL_g_m2h_mean ~ cloacal_temp_C + mass_g + SMI + osmolality_mmol_kg_mean +
##      hematocrit_percent + msmt_temp_C + msmt_VPD_kPa + VPD_kPa_int +
##      wind_mph_interpol + solar_rad_W_sqm_interpol + hold_time_hr +
##      (1 | capture_date)
##
##          npar      AIC
## <none>          657.40
## cloacal_temp_C      1 655.48
## mass_g              1 655.92
## SMI                 1 655.40
## osmolality_mmol_kg_mean 1 668.42
## hematocrit_percent      1 655.45
## msmt_temp_C            1 678.51
## msmt_VPD_kPa           1 665.68
## VPD_kPa_int            1 657.64
## wind_mph_interpol      1 662.39
## solar_rad_W_sqm_interpol 1 655.71
## hold_time_hr           1 659.19

```

Great, VIFs are minimal and we're ready to start backwards selection!

Start with dropping SMI:

```
CEWL_mod5 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # essential covariate
  cloacal_temp_C +
  # body size
  mass_g +
  # blood
  osmolality_mmol_kg_mean + hematocrit_percent +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
  # weather at the time of capture
  VPD_kPa_int +
  wind_mph_interpol + solar_rad_W_sqm_interpol +
  # time between capture and measurements
  hold_time_hr +
  (1|capture_date))

summary(CEWL_mod5)

## Linear mixed model fit by REML ['lmerMod']
## Formula: CEWL_g_m2h_mean ~ cloacal_temp_C + mass_g + osmolality_mmol_kg_mean +
##      hematocrit_percent + msmt_temp_C + msmt_VPD_kPa + VPD_kPa_int +
##      wind_mph_interpol + solar_rad_W_sqm_interpol + hold_time_hr +
##      (1 | capture_date)
##      Data: dat
##
## REML criterion at convergence: 644.3
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1751 -0.6201 -0.0115  0.4975  3.4051
##
## Random effects:
##      Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 4.952     2.225
## Residual              9.302     3.050
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   -1.665e+02  3.499e+01  -4.758
## cloacal_temp_C    1.316e-01  4.019e-01   0.328
## mass_g         -1.519e-01  1.795e-01  -0.846
## osmolality_mmol_kg_mean  7.186e-02  2.057e-02   3.493
## hematocrit_percent  1.453e-02  5.677e-02   0.256
## msmt_temp_C      7.148e+00  1.494e+00   4.784
## msmt_VPD_kPa    -1.891e+01  6.258e+00  -3.023
## VPD_kPa_int     -1.574e+00  1.052e+00  -1.496
## wind_mph_interpol  8.407e-01  3.420e-01   2.458
## solar_rad_W_sqm_interpol  2.321e-03  3.448e-03   0.673
## hold_time_hr     6.113e-01  3.208e-01   1.905
```

```
##
## Correlation of Fixed Effects:
##      (Intr) clc__C mass_g osm___ hmtcr_ msm__C m_VPD_ VPD_P_ wnd_m_
## clocl_tmp_C -0.318
## mass_g      -0.076  0.050
## osmlly_m__ -0.291  0.046 -0.082
## hmtcr_prcn  0.007 -0.026 -0.126  0.005
## msmt_temp_C -0.902  0.045  0.029  0.093 -0.116
## msmt_VPD_kP  0.499 -0.175 -0.045 -0.060  0.241 -0.700
## VPD_kPa_int  0.131  0.026  0.030  0.024 -0.198  0.055 -0.526
## wnd_mph_ntr -0.329  0.194  0.058  0.110 -0.232  0.390 -0.624  0.034
## slr_rd_W_s_  0.355  0.025  0.009 -0.225  0.095 -0.505  0.587 -0.637 -0.173
## hold_tim_hr  0.105  0.241  0.189  0.074 -0.050 -0.153 -0.213  0.173  0.415
##      s__W__
## clocl_tmp_C
## mass_g
## osmlly_m__
## hmtcr_prcn
## msmt_temp_C
## msmt_VPD_kP
## VPD_kPa_int
## wnd_mph_ntr
## slr_rd_W_s_
## hold_tim_hr  0.069
```

```
drop1(CEWL_mod5)
```

```
## Single term deletions
##
## Model:
## CEWL_g_m2h_mean ~ cloacal_temp_C + mass_g + osmolality_mmol_kg_mean +
##      hematocrit_percent + msmt_temp_C + msmt_VPD_kPa + VPD_kPa_int +
##      wind_mph_interpol + solar_rad_W_sqm_interpol + hold_time_hr +
##      (1 | capture_date)
##      npar      AIC
## <none>          655.40
## cloacal_temp_C    1 653.49
## mass_g            1 654.11
## osmolality_mmol_kg_mean 1 666.64
## hematocrit_percent 1 653.45
## msmt_temp_C       1 676.54
## msmt_VPD_kPa      1 663.68
## VPD_kPa_int       1 655.64
## wind_mph_interpol 1 660.40
## solar_rad_W_sqm_interpol 1 653.72
## hold_time_hr      1 657.19
```

next drop cloacal temperature (What?!):

```
CEWL_mod6 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # body size
  mass_g +
  # blood
  osmolality_mmol_kg_mean + hematocrit_percent +
```

```

# microclimate at the time of msmt
msmt_temp_C + msmt_VPD_kPa +
# weather at the time of capture
VPD_kPa_int +
wind_mph_interpol + solar_rad_W_sqm_interpol +
# time between capture and measurements
hold_time_hr +
(1|capture_date))
summary(CEWL_mod6)

## Linear mixed model fit by REML ['lmerMod']
## Formula:
## CEWL_g_m2h_mean ~ mass_g + osmolality_mmol_kg_mean + hematocrit_percent +
## msmt_temp_C + msmt_VPD_kPa + VPD_kPa_int + wind_mph_interpol +
## solar_rad_W_sqm_interpol + hold_time_hr + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 644.4
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1956 -0.6066 -0.0044  0.5205  3.4154
##
## Random effects:
## Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 4.863    2.205
## Residual          9.231    3.038
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)    -1.629e+02  3.303e+01  -4.931
## mass_g          -1.547e-01  1.786e-01  -0.866
## osmolality_mmol_kg_mean  7.158e-02  2.047e-02   3.497
## hematocrit_percent  1.498e-02  5.653e-02   0.265
## msmt_temp_C      7.129e+00  1.486e+00   4.796
## msmt_VPD_kPa    -1.857e+01  6.133e+00  -3.027
## VPD_kPa_int     -1.582e+00  1.047e+00  -1.511
## wind_mph_interpol  8.193e-01  3.340e-01   2.453
## solar_rad_W_sqm_interpol  2.287e-03  3.433e-03   0.666
## hold_time_hr     5.858e-01  3.101e-01   1.889
##
## Correlation of Fixed Effects:
##              (Intr) mass_g osm___ hmtcr_ msm__C m_VPD_ VPD_P_ wnd_m_ s__W__
## mass_g        -0.064
## osmlly_m__    -0.292 -0.084
## hmtcrt_prcn   -0.001 -0.125  0.007
## msmt_temp_C   -0.937  0.026  0.091 -0.115
## msmt_VPD_kP   0.475 -0.037 -0.053  0.240 -0.703
## VPD_kPa_int   0.147  0.029  0.023 -0.198  0.055 -0.531
## wnd_mph_ntr  -0.288  0.049  0.103 -0.231  0.390 -0.611  0.030
## slr_rd_W_s_   0.383  0.008 -0.226  0.096 -0.507  0.601 -0.638 -0.182
## hold_tim_hr   0.198  0.183  0.065 -0.045 -0.169 -0.179  0.172  0.387  0.065

```

```
drop1(CEWL_mod6)
```

```
## Single term deletions
##
## Model:
## CEWL_g_m2h_mean ~ mass_g + osmolality_mmol_kg_mean + hematocrit_percent +
##   msmt_temp_C + msmt_VPD_kPa + VPD_kPa_int + wind_mph_interpol +
##   solar_rad_W_sqm_interpol + hold_time_hr + (1 | capture_date)
##               npar      AIC
## <none>                653.49
## mass_g                1 652.21
## osmolality_mmol_kg_mean 1 664.67
## hematocrit_percent      1 651.54
## msmt_temp_C            1 674.57
## msmt_VPD_kPa           1 661.72
## VPD_kPa_int            1 653.73
## wind_mph_interpol      1 658.45
## solar_rad_W_sqm_interpol 1 651.79
## hold_time_hr           1 655.24
```

next drop hematocrit:

```
CEWL_mod7 <- lme4::lmer(data = dat,
                        # response variable
                        CEWL_g_m2h_mean ~
                        # body size
                        mass_g +
                        # blood
                        osmolality_mmol_kg_mean +
                        # microclimate at the time of msmt
                        msmt_temp_C + msmt_VPD_kPa +
                        # weather at the time of capture
                        VPD_kPa_int +
                        wind_mph_interpol + solar_rad_W_sqm_interpol +
                        # time between capture and measurements
                        hold_time_hr +
                        (1|capture_date))
summary(CEWL_mod7)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: CEWL_g_m2h_mean ~ mass_g + osmolality_mmol_kg_mean + msmt_temp_C +
##   msmt_VPD_kPa + VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##   hold_time_hr + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 640.6
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1691 -0.6099  0.0036  0.5315  3.4388
##
## Random effects:
##   Groups             Name             Variance Std.Dev.
##   capture_date (Intercept) 4.814         2.194
##   Residual                9.157         3.026
```

```
## Number of obs: 124, groups:  capture_date, 5
##
## Fixed effects:
##
##              Estimate Std. Error t value
## (Intercept)    -1.629e+02  3.290e+01  -4.952
## mass_g          -1.488e-01  1.765e-01  -0.843
## osmolality_mmol_kg_mean  7.154e-02  2.038e-02   3.510
## msmt_temp_C      7.174e+00  1.470e+00   4.880
## msmt_VPD_kPa     -1.896e+01  5.929e+00  -3.198
## VPD_kPa_int      -1.527e+00  1.022e+00  -1.494
## wind_mph_interpol  8.399e-01  3.236e-01   2.596
## solar_rad_W_sqm_interpol  2.199e-03  3.403e-03   0.646
## hold_time_hr      5.895e-01  3.085e-01   1.911
##
## Correlation of Fixed Effects:
##              (Intr) mass_g osm___ msm__C m_VPD_ VPD_P_ wnd_m_ s__W__
## mass_g        -0.064
## osmllty_m__   -0.292 -0.084
## msmt_temp_C   -0.944  0.012  0.092
## msmt_VPD_kP   0.489 -0.007 -0.056 -0.700
## VPD_kPa_int   0.149  0.004  0.025  0.033 -0.509
## wnd_mph_ntr  -0.297  0.021  0.108  0.376 -0.588 -0.017
## slr_rd_W_s_   0.385  0.020 -0.228 -0.502  0.599 -0.634 -0.165
## hold_tim_hr   0.198  0.179  0.065 -0.176 -0.173  0.167  0.387  0.069
drop1(CEWL_mod7)

## Single term deletions
##
## Model:
## CEWL_g_m2h_mean ~ mass_g + osmolality_mmol_kg_mean + msmt_temp_C +
##      msmt_VPD_kPa + VPD_kPa_int + wind_mph_interpol + solar_rad_W_sqm_interpol +
##      hold_time_hr + (1 | capture_date)
##              npar      AIC
## <none>                651.54
## mass_g                1 650.23
## osmolality_mmol_kg_mean  1 662.72
## msmt_temp_C           1 673.17
## msmt_VPD_kPa          1 660.75
## VPD_kPa_int           1 651.74
## wind_mph_interpol      1 657.21
## solar_rad_W_sqm_interpol  1 649.82
## hold_time_hr           1 653.34
```

next drop solar radiation:

```
CEWL_mod8 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # body size
  mass_g +
  # blood
  osmolality_mmol_kg_mean +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
  # weather at the time of capture
```



```

        VPD_kPa_int +
        wind_mph_interpol +
        # time between capture and measurements
        hold_time_hr +
        (1|capture_date))
summary(CEWL_mod8)

## Linear mixed model fit by REML ['lmerMod']
## Formula: CEWL_g_m2h_mean ~ mass_g + osmolality_mmol_kg_mean + msmt_temp_C +
##      msmt_VPD_kPa + VPD_kPa_int + wind_mph_interpol + hold_time_hr +
##      (1 | capture_date)
##      Data: dat
##
## REML criterion at convergence: 631.5
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1289 -0.6630  0.0006  0.5170  3.4722
##
## Random effects:
##      Groups          Name          Variance Std.Dev.
## capture_date (Intercept) 4.368      2.090
## Residual                9.129      3.021
## Number of obs: 124, groups:  capture_date, 5
##
## Fixed effects:
##
##              Estimate Std. Error t value
## (Intercept)    -171.43335    30.15878   -5.684
## mass_g          -0.15039     0.17619   -0.854
## osmolality_mmol_kg_mean  0.07461     0.01977    3.773
## msmt_temp_C      7.66653     1.25869    6.091
## msmt_VPD_kPa    -21.30228     4.66496   -4.566
## VPD_kPa_int     -1.10894     0.78696   -1.409
## wind_mph_interpol  0.87805     0.31656    2.774
## hold_time_hr     0.57564     0.30695    1.875
##
## Correlation of Fixed Effects:
##              (Intr) mass_g osm___ msm__C m_VPD_ VPD_P_ wnd_m_
## mass_g        -0.079
## osmllty_m__   -0.228 -0.081
## msmt_temp_C   -0.940  0.026 -0.026
## msmt_VPD_kP   0.343 -0.025  0.105 -0.571
## VPD_kPa_int   0.549  0.022 -0.161 -0.423 -0.221
## wnd_mph_ntr  -0.259  0.024  0.079  0.346 -0.624 -0.160
## hold_tim_hr   0.186  0.178  0.085 -0.165 -0.270  0.275  0.403
drop1(CEWL_mod8)

## Single term deletions
##
## Model:
## CEWL_g_m2h_mean ~ mass_g + osmolality_mmol_kg_mean + msmt_temp_C +
##      msmt_VPD_kPa + VPD_kPa_int + wind_mph_interpol + hold_time_hr +
##      (1 | capture_date)

```

```
##               npar    AIC
## <none>                649.82
## mass_g                1 648.51
## osmolality_mmol_kg_mean 1 662.57
## msmt_temp_C           1 674.36
## msmt_VPD_kPa          1 664.06
## VPD_kPa_int           1 650.01
## wind_mph_interpol     1 656.43
## hold_time_hr          1 651.50
```

next drop mass:

```
CEWL_mod9 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # blood
  osmolality_mmol_kg_mean +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
  # weather at the time of capture
  VPD_kPa_int +
  wind_mph_interpol +
  # time between capture and measurements
  hold_time_hr +
  (1|capture_date))
summary(CEWL_mod9)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula:
## CEWL_g_m2h_mean ~ osmolality_mmol_kg_mean + msmt_temp_C + msmt_VPD_kPa +
##   VPD_kPa_int + wind_mph_interpol + hold_time_hr + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 630.6
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1586 -0.6440  0.0060  0.5491  3.3550
##
## Random effects:
##   Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 4.231    2.057
## Residual              9.114    3.019
## Number of obs: 124, groups: capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   -173.58299   29.99067  -5.788
## osmolality_mmol_kg_mean    0.07327    0.01968   3.723
## msmt_temp_C         7.70005    1.25371   6.142
## msmt_VPD_kPa       -21.41767    4.63600  -4.620
## VPD_kPa_int       -1.09462    0.78545  -1.394
## wind_mph_interpol    0.88553    0.31553   2.807
## hold_time_hr        0.62206    0.30170   2.062
##
```

```

## Correlation of Fixed Effects:
##          (Intr) osm___ msm__C m_VPD_ VPD_P_ wnd_m_
## osmllty_m__ -0.237
## msmt_temp_C -0.941 -0.024
## msmt_VPD_kP  0.341  0.104 -0.568
## VPD_kPa_int  0.552 -0.161 -0.422 -0.225
## wnd_mph_ntr -0.259  0.083  0.347 -0.625 -0.161
## hold_tim_hr  0.204  0.102 -0.174 -0.270  0.276  0.405
drop1(CEWL_mod9)

## Single term deletions
##
## Model:
## CEWL_g_m2h_mean ~ osmolality_mmol_kg_mean + msmt_temp_C + msmt_VPD_kPa +
##      VPD_kPa_int + wind_mph_interpol + hold_time_hr + (1 | capture_date)
##              npar      AIC
## <none>                648.51
## osmolality_mmol_kg_mean    1 660.83
## msmt_temp_C                1 673.00
## msmt_VPD_kPa               1 662.83
## VPD_kPa_int                1 648.64
## wind_mph_interpol          1 655.26
## hold_time_hr               1 650.87

next drop VPD:
CEWL_mod10 <- lme4::lmer(data = dat,
                        # response variable
                        CEWL_g_m2h_mean ~
                        # blood
                        osmolality_mmol_kg_mean +
                        # microclimate at the time of msmt
                        msmt_temp_C + msmt_VPD_kPa +
                        # weather at the time of capture
                        wind_mph_interpol +
                        # time between capture and measurements
                        hold_time_hr +
                        (1|capture_date))
summary(CEWL_mod10)

## Linear mixed model fit by REML ['lmerMod']
## Formula:
## CEWL_g_m2h_mean ~ osmolality_mmol_kg_mean + msmt_temp_C + msmt_VPD_kPa +
##      wind_mph_interpol + hold_time_hr + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 633.8
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3031 -0.6815  0.0296  0.6037  3.2530
##
## Random effects:
##      Groups      Name      Variance Std.Dev.
## capture_date (Intercept) 4.417      2.102

```

```
## Residual          9.179    3.030
## Number of obs: 124, groups:  capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   -150.31800   25.12710  -5.982
## osmolality_mmol_kg_mean    0.06886    0.01951   3.529
## msmt_temp_C       6.95213    1.14343   6.080
## msmt_VPD_kPa     -22.82527    4.56608  -4.999
## wind_mph_interpol    0.81352    0.31334   2.596
## hold_time_hr      0.73782    0.29121   2.534
##
## Correlation of Fixed Effects:
##              (Intr) osm___ msm__C m_VPD_ wnd_m_
## osmlly_m__ -0.180
## msmt_temp_C -0.937 -0.103
## msmt_VPD_kP  0.572  0.071 -0.751
## wnd_mph_ntr -0.206  0.056  0.311 -0.684
## hold_tim_hr  0.065  0.153 -0.065 -0.223  0.474
```

can't test drop1 bc of NA's

based on t-values, wind should be dropped next:

```
CEWL_mod11 <- lme4::lmer(data = dat,
  # response variable
  CEWL_g_m2h_mean ~
  # blood
  osmolality_mmol_kg_mean +
  # microclimate at the time of msmt
  msmt_temp_C + msmt_VPD_kPa +
  # time between capture and measurements
  hold_time_hr +
  (1|capture_date))
summary(CEWL_mod11)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula:
## CEWL_g_m2h_mean ~ osmolality_mmol_kg_mean + msmt_temp_C + msmt_VPD_kPa +
##   hold_time_hr + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 639.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3700 -0.7378  0.0347  0.5964  3.2260
##
## Random effects:
## Groups      Name                Variance Std.Dev.
## capture_date (Intercept) 6.893      2.625
## Residual          9.510      3.084
## Number of obs: 124, groups:  capture_date, 5
##
## Fixed effects:
```

```
##               Estimate Std. Error t value
## (Intercept)    -136.16631   25.49925  -5.340
## osmolality_mmol_kg_mean    0.06695   0.02004   3.341
## msmt_temp_C         5.96258   1.14748   5.196
## msmt_VPD_kPa       -14.35840   3.88834  -3.693
## hold_time_hr         0.38517   0.26150   1.473
##
## Correlation of Fixed Effects:
##           (Intr) osm___ msm__C m_VPD_
## osmllty_m__ -0.177
## msmt_temp_C -0.936 -0.115
## msmt_VPD_kP  0.598  0.118 -0.779
## hold_tim_hr  0.175  0.144 -0.229  0.110
```

drop hold time:

```
CEWL_mod11a <- lme4::lmer(data = dat,
                          # response variable
                          CEWL_g_m2h_mean ~
                          # blood
                          osmolality_mmol_kg_mean +
                          # microclimate at the time of msmt
                          msmt_temp_C + msmt_VPD_kPa +
                          (1|capture_date))
summary(CEWL_mod11a)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula:
## CEWL_g_m2h_mean ~ osmolality_mmol_kg_mean + msmt_temp_C + msmt_VPD_kPa +
##   (1 | capture_date)
##   Data: dat
##
## REML criterion at convergence: 740.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.0076 -0.6511 -0.0032  0.6792  3.4191
##
## Random effects:
##   Groups       Name             Variance Std.Dev.
## capture_date (Intercept)  2.828     1.682
## Residual                12.064     3.473
## Number of obs: 138, groups: capture_date, 5
##
## Fixed effects:
##               Estimate Std. Error t value
## (Intercept)    -137.99713   26.31861  -5.243
## osmolality_mmol_kg_mean    0.05162   0.02061   2.505
## msmt_temp_C         6.44585   1.11981   5.756
## msmt_VPD_kPa       -16.51862   2.96126  -5.578
##
## Correlation of Fixed Effects:
##           (Intr) osm___ msm__C
## osmllty_m__ -0.195
## msmt_temp_C -0.944 -0.102
```

```
## msmt_VPD_kP 0.624 0.169 -0.783
```

drop osmolality:

```
CEWL_mod12 <- lme4::lmer(data = dat,  
  # response variable  
  CEWL_g_m2h_mean ~  
  # microclimate at the time of msmt  
  msmt_temp_C + msmt_VPD_kPa +  
  (1|capture_date))  
summary(CEWL_mod12)
```

```
## Linear mixed model fit by REML ['lmerMod']  
## Formula: CEWL_g_m2h_mean ~ msmt_temp_C + msmt_VPD_kPa + (1 | capture_date)  
## Data: dat  
##  
## REML criterion at convergence: 740.9  
##  
## Scaled residuals:  
##      Min       1Q   Median       3Q      Max   
## -2.1447 -0.6565 -0.0485  0.6256  3.3559   
##  
## Random effects:  
## Groups      Name      Variance Std.Dev.  
## capture_date (Intercept) 2.385   1.544  
## Residual              12.581   3.547  
## Number of obs: 138, groups: capture_date, 5  
##  
## Fixed effects:  
##              Estimate Std. Error t value  
## (Intercept)  -125.832    26.094  -4.822  
## msmt_temp_C     6.769     1.119   6.048  
## msmt_VPD_kPa  -17.893     2.795  -6.401  
##  
## Correlation of Fixed Effects:  
##              (Intr) msm__C  
## msmt_temp_C  -0.990  
## msmt_VPD_kP  0.692 -0.786
```

```
drop1(CEWL_mod12)
```

```
## Single term deletions
```

```
##
```

```
## Model:
```

```
## CEWL_g_m2h_mean ~ msmt_temp_C + msmt_VPD_kPa + (1 | capture_date)
```

```
##              npar      AIC
```

```
## <none>          756.56
```

```
## msmt_temp_C      1 779.51
```

```
## msmt_VPD_kPa     1 768.96
```

drop VPD at the time of msmt:

```
CEWL_mod13 <- lme4::lmer(data = dat,  
  # response variable  
  CEWL_g_m2h_mean ~  
  # microclimate at the time of msmt  
  msmt_temp_C +
```

```

                                (1|capture_date))
summary(CEWL_mod13)

## Linear mixed model fit by REML ['lmerMod']
## Formula: CEWL_g_m2h_mean ~ msmt_temp_C + (1 | capture_date)
##   Data: dat
##
## REML criterion at convergence: 755.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1664 -0.6915 -0.0549  0.5410  3.2942
##
## Random effects:
##   Groups       Name             Variance Std.Dev.
## capture_date (Intercept) 39.63      6.295
## Residual                12.55      3.543
## Number of obs: 138, groups:  capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept) -72.3114    22.5292  -3.210
## msmt_temp_C   3.4235     0.8222   4.164
##
## Correlation of Fixed Effects:
##              (Intr)
## msmt_temp_C -0.992

```

And finally, null model:

```

CEWL_mod_null <- lme4::lmer(data = dat,
                             # response variable
                             CEWL_g_m2h_mean ~ 1 +
                             (1|capture_date))
summary(CEWL_mod_null)

## Linear mixed model fit by REML ['lmerMod']
## Formula: CEWL_g_m2h_mean ~ 1 + (1 | capture_date)
##   Data: dat
##
## REML criterion at convergence: 772.6
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.4713 -0.6736 -0.0596  0.7187  3.1910
##
## Random effects:
##   Groups       Name             Variance Std.Dev.
## capture_date (Intercept) 19.32      4.396
## Residual                14.29      3.780
## Number of obs: 138, groups:  capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)  20.746     1.993    10.41

```

Selection

compare models 4-13 and the null

```
CEWL_models <- list(CEWL_mod4, CEWL_mod5, CEWL_mod6, CEWL_mod7,
  CEWL_mod8, CEWL_mod9, CEWL_mod10, CEWL_mod11, CEWL_mod11a,
  CEWL_mod12, CEWL_mod13, CEWL_mod_null)

#specify model names
CEWL_mod_names <- c('(model 4) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hct, Temp-Clo, SMI,
  '(model 5) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hct, Temp-Clo, Hold',
  '(model 6) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hct, Hold',
  '(model 7) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hold',
  '(model 8) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Hold',
  '(model 9) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Hold',
  '(model 10) ~ Temp-M, VPD-M, Osml, Wind-C, Hold',
  '(model 11) ~ Temp-M, VPD-M, Osml, Hold',
  '(model 11a) ~ Temp-M, VPD-M, Osml',
  '(model 12) ~ Temp-M, VPD-M',
  '(model 13) ~ Temp-M',
  'null model')

#calculate AIC of each model
CEWL_AICc <- data.frame(aictab(cand.set = CEWL_models,
  modnames = CEWL_mod_names))

## Warning in aictab.AIClmerMod(cand.set = CEWL_models, modnames = CEWL_mod_names):
## Model selection for fixed effects is only appropriate with ML estimation:
## REML (default) should only be used to select random effects for a constant set of fixed effects
CEWL_AICc
```

									Modnames
## 6									(model 9) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Hold
## 7									(model 10) ~ Temp-M, VPD-M, Osml, Wind-C, Hold
## 5									(model 8) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Hold
## 8									(model 11) ~ Temp-M, VPD-M, Osml, Hold
## 4									(model 7) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hold
## 3									(model 6) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hct, Hold
## 2									(model 5) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hct, Temp-Clo, Hold
## 1									(model 4) ~ Temp-M, VPD-M, Osml, Wind-C, VPD-C, Mass, Solar-C, Hct, Temp-Clo, SMI, Hold
## 10									(model 12) ~ Temp-M, VPD-M
## 9									(model 11a) ~ Temp-M, VPD-M, Osml
## 11									(model 13) ~ Temp-M
## 12									null model
##	K	AICc	Delta_AICc	ModelLik	AICcWt	Res.LL	Cum.Wt		
## 6	9	650.1317	0.0000000	1.000000e+00	5.223517e-01	-315.2764	0.5223517		
## 7	8	651.0968	0.9650419	6.172254e-01	3.224088e-01	-316.9223	0.8447604		
## 5	10	653.4071	3.2753338	1.944331e-01	1.015625e-01	-315.7301	0.9463229		
## 8	7	654.6950	4.5632966	1.021158e-01	5.334033e-02	-319.8648	0.9996633		
## 4	11	664.9398	14.8080831	6.087873e-04	3.180011e-04	-320.2913	0.9999813		
## 3	12	671.2351	21.1033836	2.614920e-05	1.365908e-05	-322.2122	0.9999949		
## 2	13	673.6151	23.4834176	7.955009e-06	4.155312e-06	-322.1530	0.9999991		
## 1	14	676.6154	26.4837148	1.774739e-06	9.270377e-07	-322.3811	1.0000000		
## 10	5	751.3494	101.2176916	1.049198e-22	5.480505e-23	-370.4474	1.0000000		
## 9	6	753.3335	103.2017320	3.890708e-23	2.032318e-23	-370.3461	1.0000000		
## 11	4	764.0328	113.9010693	1.847979e-25	9.652950e-26	-377.8660	1.0000000		


```
## 12 3 778.7422 128.6105011 1.181910e-28 6.173728e-29 -386.2816 1.0000000
```

The best two models are 9 & 10, which included Temp-M, VPD-M, Osmol, Wind-C, VPD-C (not in 10), and Hold time.

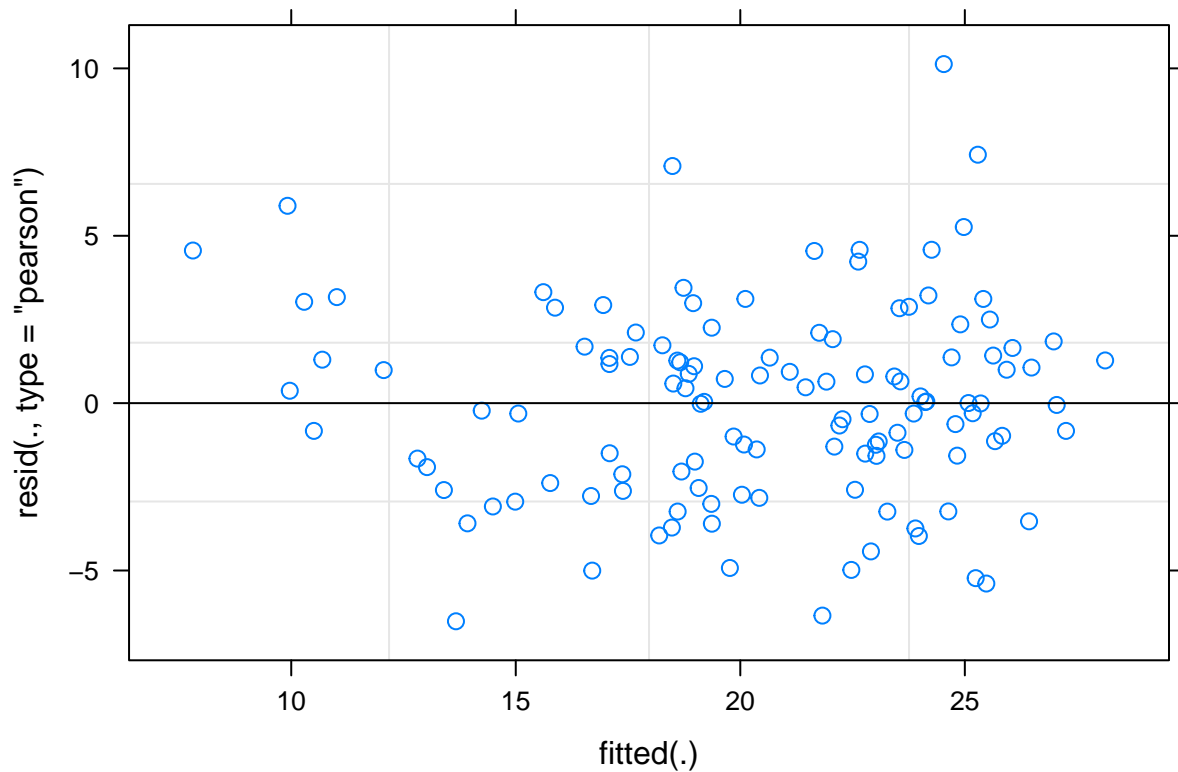
LM Conditions

Check that the best model meets the criteria for linear regression and has no collinearity.

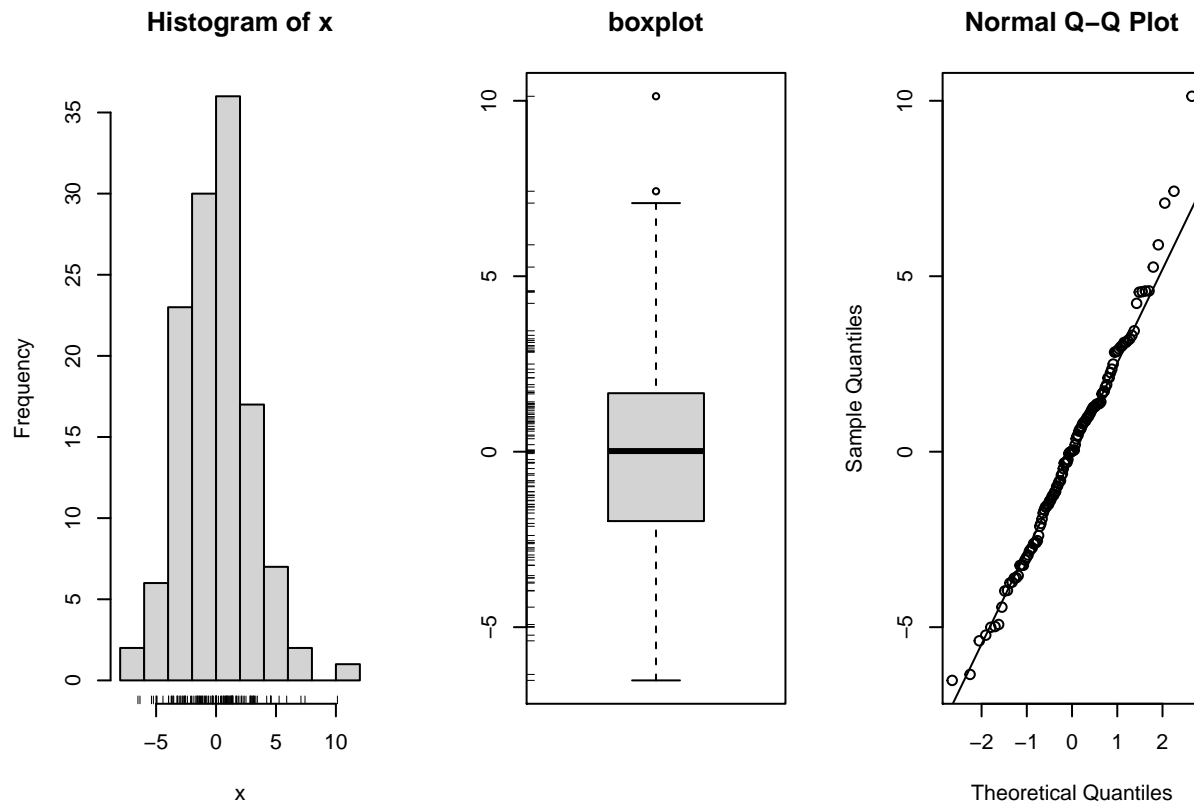
```
vif(CEWL_mod9)
```

```
## osmolality_mmol_kg_mean      msmt_temp_C      msmt_VPD_kPa
##           1.070914           3.562068           5.117716
##           VPD_kPa_int      wind_mph_interpol      hold_time_hr
##           2.957004           2.743307           1.520147
```

```
plot(CEWL_mod9)
```



```
simple.eda(residuals(CEWL_mod9))
```



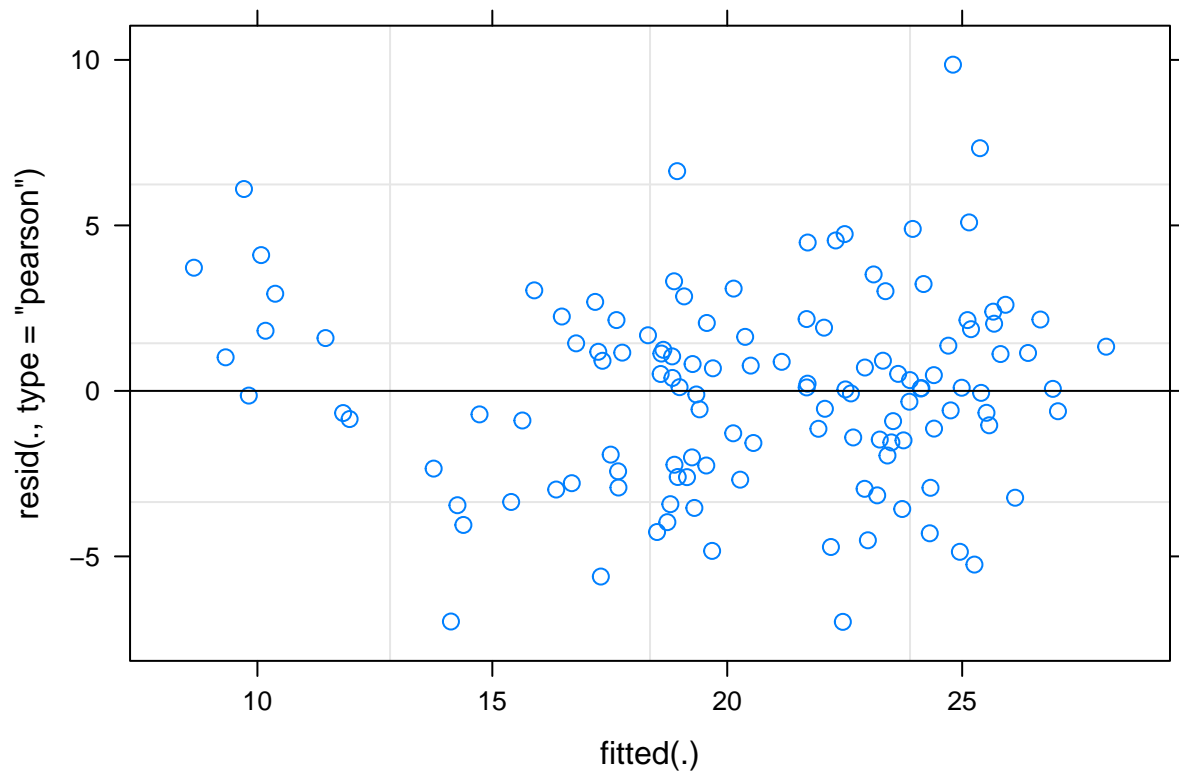
```
shapiro.test(residuals(CEWL_mod9))
```

```
##
##  Shapiro-Wilk normality test
##
## data:  residuals(CEWL_mod9)
## W = 0.98747, p-value = 0.315
```

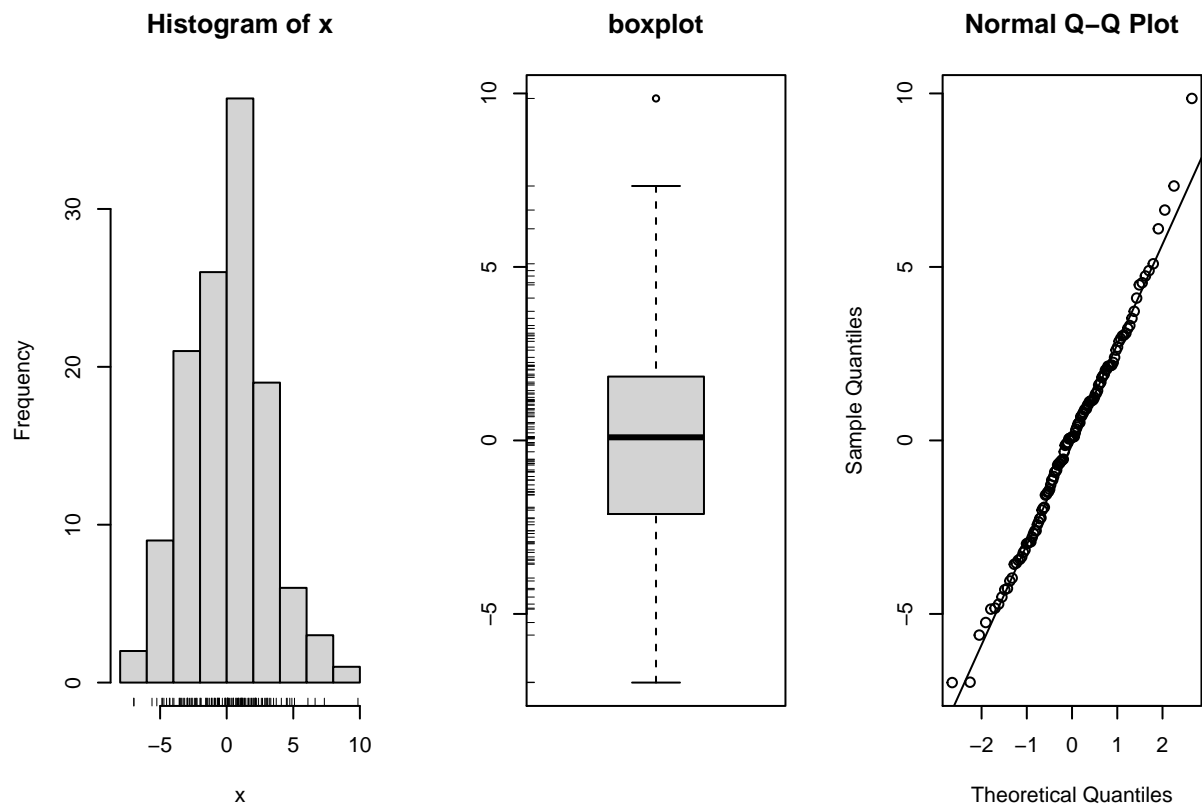
```
vif(CEWL_mod10)
```

```
## osmolality_mmol_kg_mean      msmt_temp_C      msmt_VPD_kPa
##           1.042362           2.924902           4.811710
##      wind_mph_interpol      hold_time_hr
##           2.636065           1.405275
```

```
plot(CEWL_mod10)
```



```
simple.eda(residuals(CEWL_mod10))
```



```
shapiro.test(residuals(CEWL_mod10))
```

```
##
##  Shapiro-Wilk normality test
##
## data:  residuals(CEWL_mod10)
## W = 0.99091, p-value = 0.594
```

There is some slight fanning in the residuals ~ fitted plot, suggesting equal error variance is not perfect, but overall, all LNE conditions appear to be met and VIFs are very low.

Export

First, re-run the best model using lmerTest for p-values.

```
CEWL_mod9p <- lmerTest::lmer(data = dat,
                             # response variable
                             CEWL_g_m2h_mean ~
                             # blood
                             osmolality_mmol_kg_mean +
                             # microclimate at the time of msmt
                             msmt_temp_C + msmt_VPD_kPa +
                             # weather at the time of capture
                             VPD_kPa_int +
                             wind_mph_interpol + hold_time_hr +
                             (1|capture_date))
summary(CEWL_mod9p)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## CEWL_g_m2h_mean ~ osmolality_mmol_kg_mean + msmt_temp_C + msmt_VPD_kPa +
##   VPD_kPa_int + wind_mph_interpol + hold_time_hr + (1 | capture_date)
## Data: dat
##
## REML criterion at convergence: 630.6
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.1586 -0.6440  0.0060  0.5491  3.3550
##
## Random effects:
## Groups           Name          Variance Std.Dev.
## capture_date (Intercept) 4.231      2.057
## Residual              9.114      3.019
## Number of obs: 124, groups:  capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)   -173.58299    29.99067  85.43506  -5.788 1.15e-07 ***
## osmolality_mmol_kg_mean    0.07327    0.01968 113.48079   3.723 0.000308 ***
## msmt_temp_C         7.70005    1.25371  51.02494   6.142 1.23e-07 ***
## msmt_VPD_kPa     -21.41767    4.63600  19.93145  -4.620 0.000167 ***
## VPD_kPa_int       -1.09462    0.78545 110.73114  -1.394 0.166224
## wind_mph_interpol    0.88553    0.31553  66.73483   2.807 0.006558 **
```

```

## hold_time_hr          0.62206    0.30170  116.90288    2.062 0.041437 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) osm___ msm__C m_VPD_ VPD_P_ wnd_m_
## osmllty_m__ -0.237
## msmt_temp_C -0.941 -0.024
## msmt_VPD_kPa 0.341  0.104 -0.568
## VPD_kPa_int  0.552 -0.161 -0.422 -0.225
## wnd_mph_ntr -0.259  0.083  0.347 -0.625 -0.161
## hold_tim_hr  0.204  0.102 -0.174 -0.270  0.276  0.405

CEWL_mod10p <- lmerTest::lmer(data = dat,
                             # response variable
                             CEWL_g_m2h_mean ~
                             # blood
                             osmolality_mmol_kg_mean +
                             # microclimate at the time of msmt
                             msmt_temp_C + msmt_VPD_kPa +
                             # weather at the time of capture
                             wind_mph_interpol + hold_time_hr +
                             (1|capture_date))
summary(CEWL_mod10p)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula:
## CEWL_g_m2h_mean ~ osmolality_mmol_kg_mean + msmt_temp_C + msmt_VPD_kPa +
##   wind_mph_interpol + hold_time_hr + (1 | capture_date)
##   Data: dat
##
## REML criterion at convergence: 633.8
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3031 -0.6815  0.0296  0.6037  3.2530
##
## Random effects:
##   Groups             Name             Variance Std.Dev.
##   capture_date (Intercept) 4.417       2.102
##   Residual                9.179       3.030
## Number of obs: 124, groups:  capture_date, 5
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)   -150.31800   25.12710  106.33080  -5.982 3.00e-08 ***
## osmolality_mmol_kg_mean    0.06886    0.01951  112.56853   3.529 0.000605 ***
## msmt_temp_C         6.95213    1.14343   68.01507   6.080 6.15e-08 ***
## msmt_VPD_kPa     -22.82527    4.56608   14.34413  -4.999 0.000181 ***
## wind_mph_interpol    0.81352    0.31334   67.59866   2.596 0.011553 *
## hold_time_hr      0.73782    0.29121  117.56630   2.534 0.012605 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##

```

```
## Correlation of Fixed Effects:
##           (Intr)  osm___  msm__C  m_VPD_  wnd_m_
## osmllty_m__ -0.180
## msmt_temp_C -0.937 -0.103
## msmt_VPD_kP  0.572  0.071 -0.751
## wnd_mph_ntr -0.206  0.056  0.311 -0.684
## hold_tim_hr  0.065  0.153 -0.065 -0.223  0.474
```

Save the best CEWL model output.

```
write.csv(broom.mixed::tidy(CEWL_mod9p),
          "./results_statistics/capture_CEWL_best_model1.csv")
write.csv(broom.mixed::tidy(CEWL_mod10p),
          "./results_statistics/capture_CEWL_best_model2.csv")
write.csv(CEWL_AICc,
          "./results_statistics/capture_CEWL_mod_rankings.csv")
```

To report in paper:

The best model to predict CEWL included plasma osmolality, temperature and VPD at the time of measurement, and VPD and wind at the time of capture. The final model met all linear regression conditions for linearity, normality, and equal error variance, and there was no multicollinearity.

Pub Figures

Custom Colors

```
lizard_color = "turquoise"
VPD_color = "blue"
temp_color = "gray"
solar_color = "orange"
wind_color = "orange"
date_color = "gray"
osml_color <- c(brewer.pal(6, "Set2")[c(2)])
```

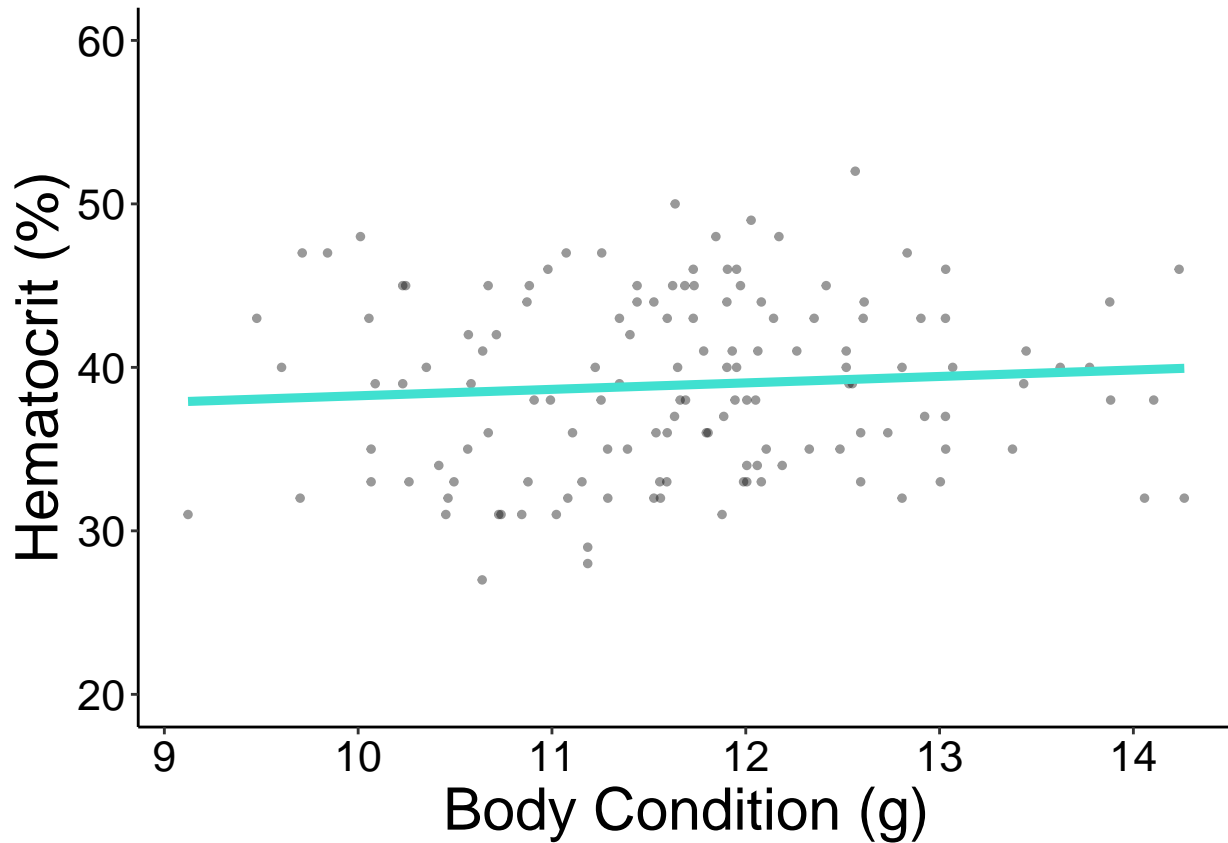
Hct ~ SMI

```
ggplot(dat) +
  aes(x = SMI,
      y = hematocrit_percent) +
  geom_point(size = 1,
             alpha = 0.4) +
  stat_smooth(formula = y ~ x,
              method = "lm",
              se = F,
              size = 1.6,
              color = lizard_color,
              alpha = 1) +
  theme_classic() +
  xlab("Body Condition (g)") +
  ylab("Hematocrit (%)") +
  #ylab("") +
  #xlim() +
  ylim(20,60) +
```

```

theme(text = element_text(color = "black",
                           family = "sans",
                           size = 22),
      axis.text = element_text(color = "black",
                               family = "sans",
                               size = 16),
      #axis.text.y = element_blank(),
      #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
    ) -> cap_hct_SMI_fig
cap_hct_SMI_fig

```



Hct ~ VPD at Capture

```

ggplot(dat) +
  aes(x = VPD_kPa_int,
      y = hematocrit_percent) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
            method = "lm",
            se = F,
            color = VPD_color,
            size = 1.6,
            alpha = 1) +
  theme_classic() +
  xlab("VPD at Capture (kPa)") +

```

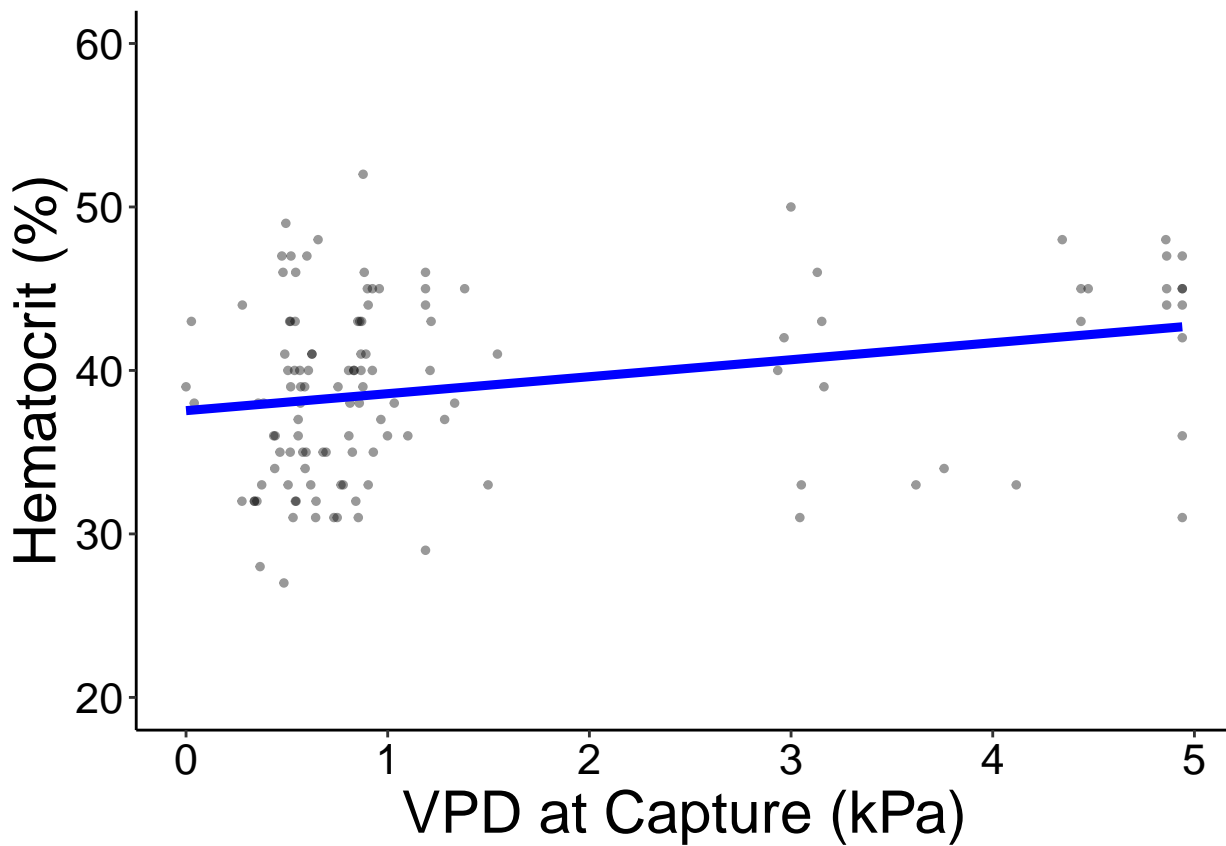
```

ylab("Hematocrit (%)") +
#ylab("") +
#xlim() +
ylim(20,60) +
theme(text = element_text(color = "black",
                             family = "sans",
                             size = 22),
      axis.text = element_text(color = "black",
                                family = "sans",
                                size = 16),
      #axis.text.y = element_blank(),
      #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
    ) -> cap_hct_VPD_fig
cap_hct_VPD_fig

```

Warning: Removed 14 rows containing non-finite values (stat_smooth).

Warning: Removed 14 rows containing missing values (geom_point).



Hct ~ Wind Speed at Capture

```

ggplot(dat) +
  aes(x = wind_mph_interpol,
       y = hematocrit_percent)+
  geom_point(size = 1,
             alpha = 0.4) +
  stat_smooth(formula = y ~ x,

```



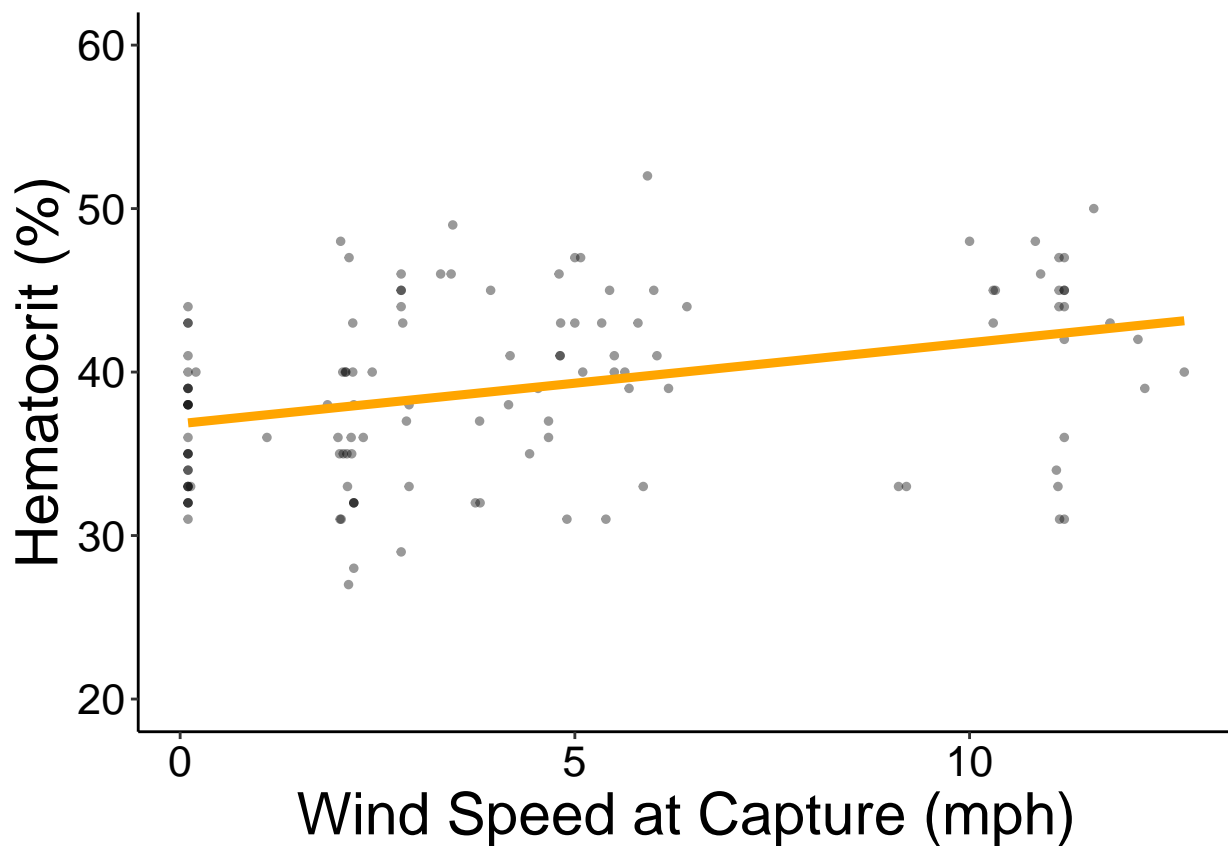
```

      method = "lm",
      se = F,
      color = wind_color,
      size = 1.6,
      alpha = 1 ) +
theme_classic() +
xlab("Wind Speed at Capture (mph)") +
ylab("Hematocrit (%)") +
#ylab("") +
#xlim() +
ylim(20, 60) +
theme(text = element_text(color = "black",
                           family = "sans",
                           size = 22),
      axis.text = element_text(color = "black",
                                family = "sans",
                                size = 16),
      #axis.text.y = element_blank(),
      #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
    ) -> cap_hct_wind_fig
cap_hct_wind_fig

```

Warning: Removed 14 rows containing non-finite values (stat_smooth).

Warning: Removed 14 rows containing missing values (geom_point).

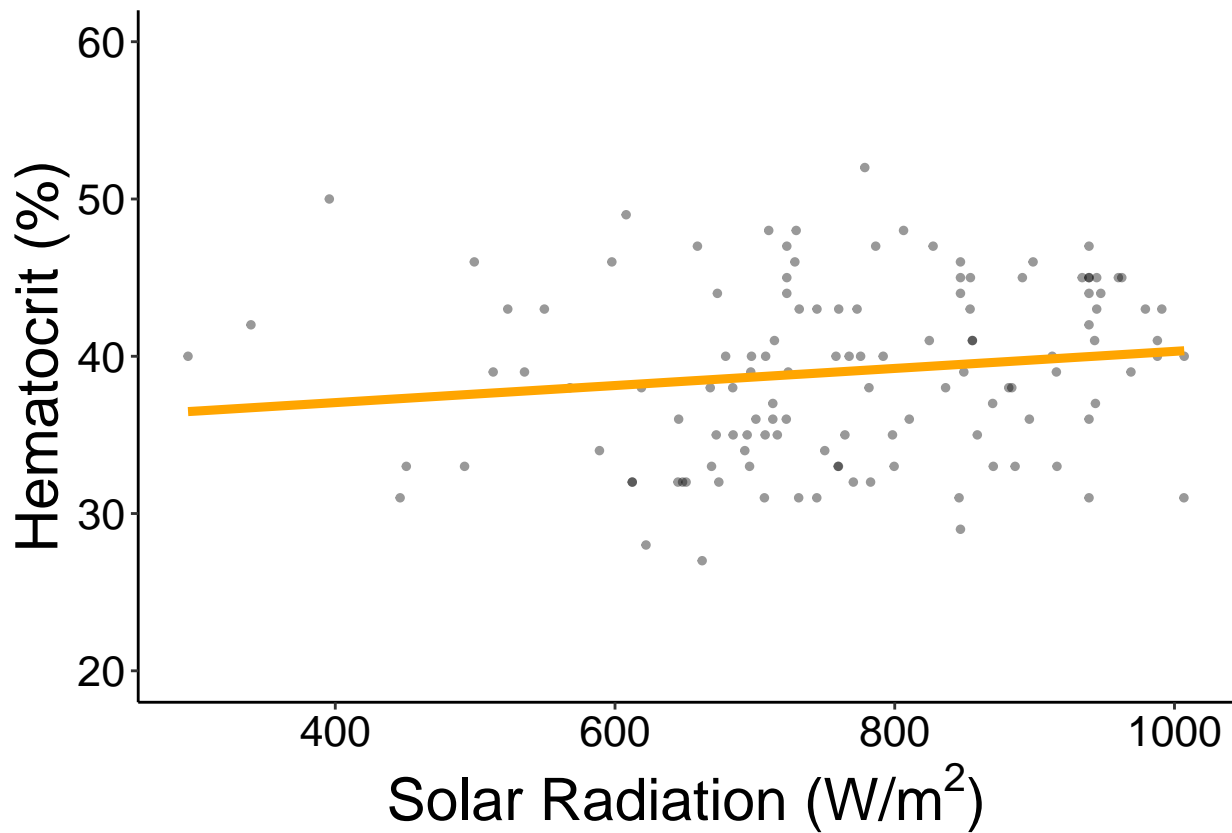


Hct ~ Solar Radiation at Capture

```
ggplot(dat) +
  aes(x = solar_rad_W_sqm_interpol,
      y = hematocrit_percent) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
             method = "lm",
             se = F,
             color = solar_color,
             size = 1.6,
             alpha = 1) +
  theme_classic() +
  xlab(bquote('Solar Radiation (W/*m^2*')')) +
  ylab("Hematocrit (%)") +
  #ylab("") +
  ylim(20, 60) +
  theme(text = element_text(color = "black",
                            family = "sans",
                            size = 22),
        axis.text = element_text(color = "black",
                                  family = "sans",
                                  size = 16),
        #axis.text.y = element_blank(),
        #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
  ) -> cap_hct_sorad_fig
cap_hct_sorad_fig
```

```
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 14 rows containing missing values (geom_point).
```



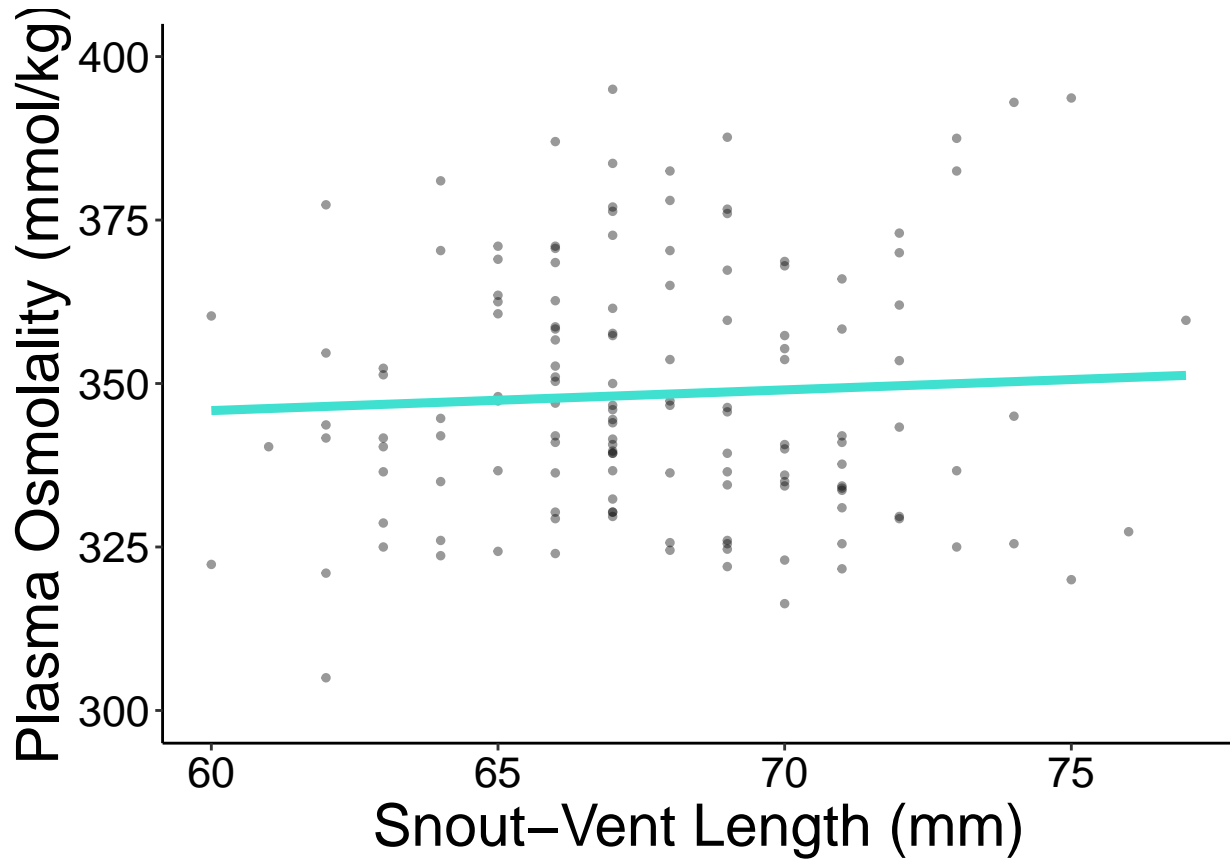
Osmolality ~ SVL

```
ggplot(dat) +
  aes(x = SVL_mm,
      y = osmolality_mmol_kg_mean) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
            method = "lm",
            se = F,
            size = 1.6,
            color = lizard_color,
            alpha = 1) +
  theme_classic() +
  xlab("Snout-Vent Length (mm)") +
  ylab("Plasma Osmolality (mmol/kg)") +
  #ylab("") +
  #xlim() +
  ylim(300,400) +
  theme(text = element_text(color = "black",
                            family = "sans",
                            size = 22),
        axis.text = element_text(color = "black",
                                family = "sans",
                                size = 16),
        #axis.text.y = element_blank(),
```

```

#plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
) -> cap_osml_SVL_fig
cap_osml_SVL_fig

```



Osmolality ~ SMI

```

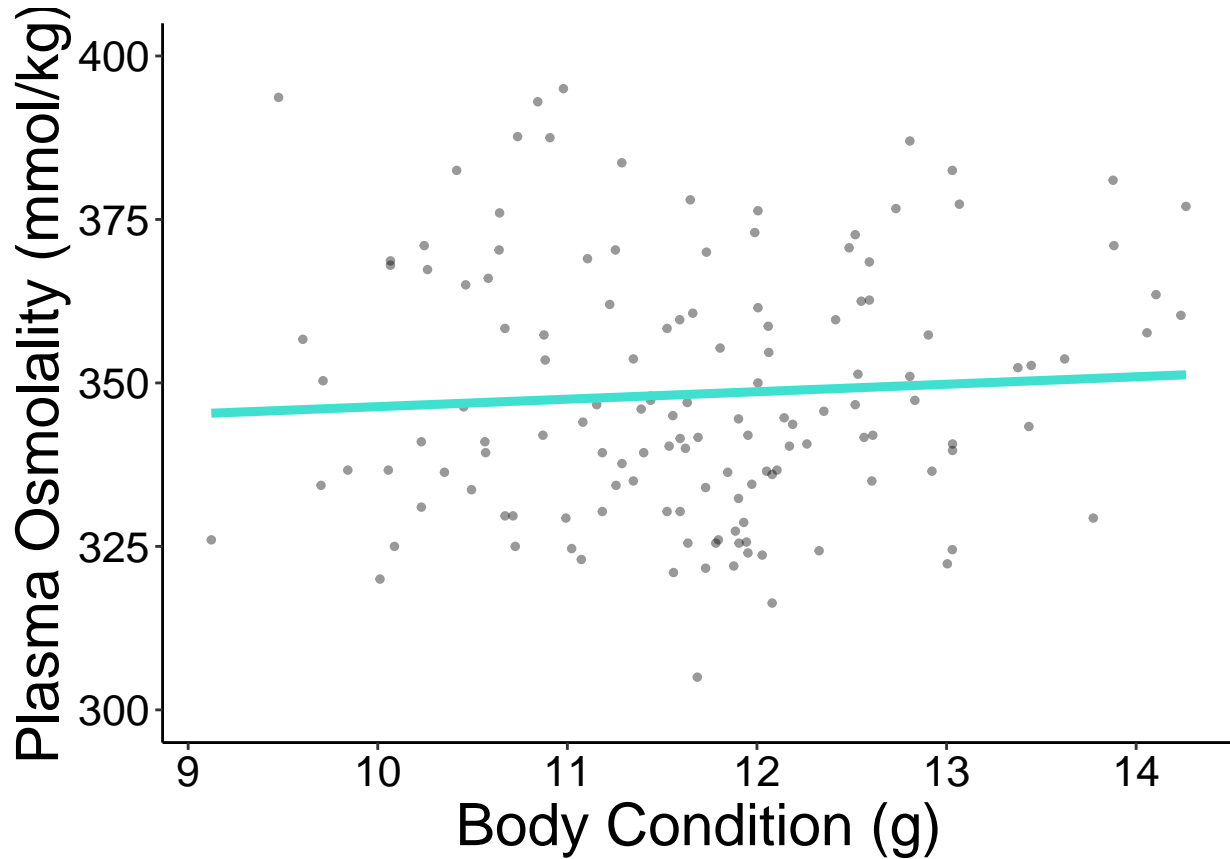
ggplot(dat) +
  aes(x = SMI,
      y = osmolality_mmol_kg_mean) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
            method = "lm",
            se = F,
            size = 1.6,
            color = lizard_color,
            alpha = 1) +
  theme_classic() +
  xlab("Body Condition (g)") +
  ylab("Plasma Osmolality (mmol/kg)") +
  #ylab("") +
  #xlim() +
  ylim(300,400) +
  theme(text = element_text(color = "black",
                            family = "sans",

```

```

        size = 22),
axis.text = element_text(color = "black",
                          family = "sans",
                          size = 16),
#axis.text.y = element_blank(),
#plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
) -> cap_osml_SMI_fig
cap_osml_SMI_fig

```



Osmolality ~ VPD at Capture

```

ggplot(dat) +
  aes(x = VPD_kPa_int,
      y = osmolality_mmol_kg_mean) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
            method = "lm",
            se = F,
            color = VPD_color,
            size = 1.6,
            alpha = 1) +
  theme_classic() +
  xlab("VPD at Capture (kPa)") +
  ylab("Plasma Osmolality (mmol/kg)") +

```

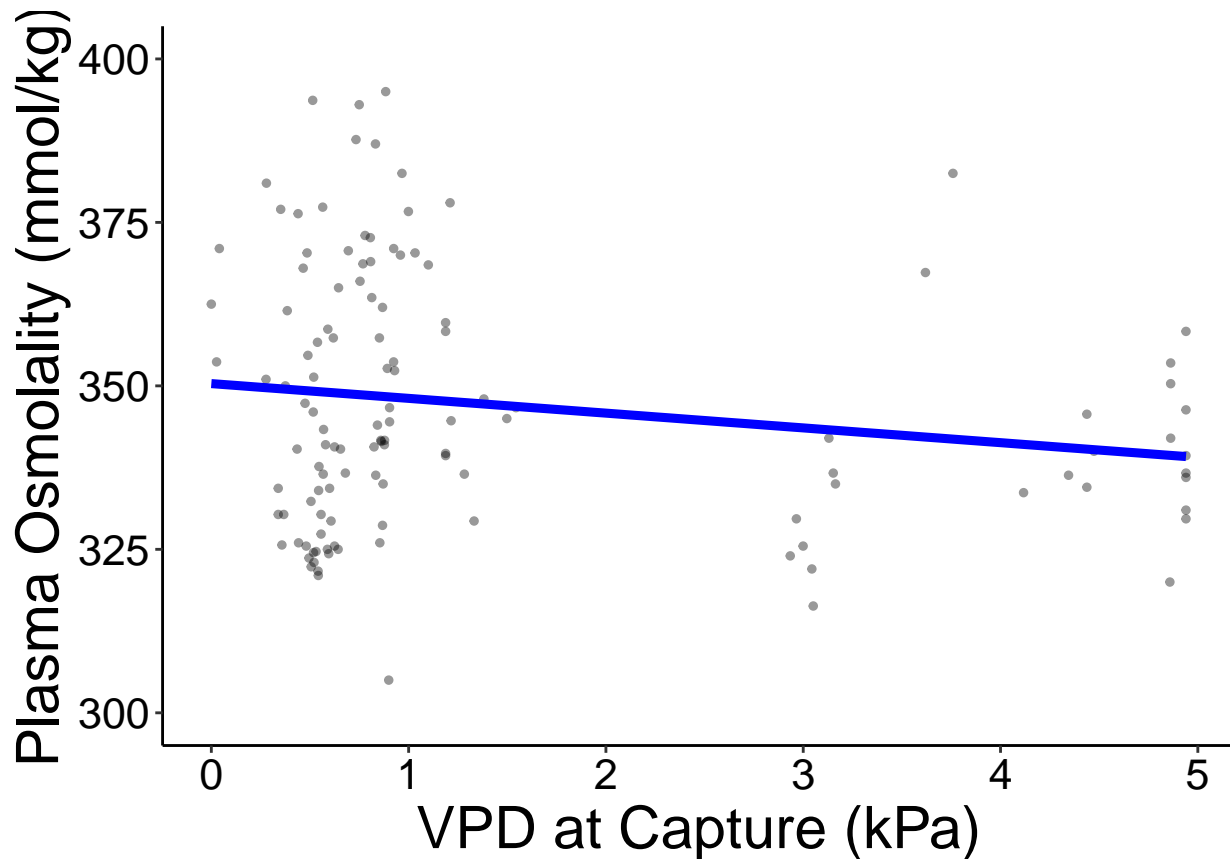
```

#ylab("") +
#xlim() +
ylim(300,400) +
theme(text = element_text(color = "black",
                             family = "sans",
                             size = 22),
      axis.text = element_text(color = "black",
                                family = "sans",
                                size = 16),
      #axis.text.y = element_blank(),
      #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
    ) -> cap_osml_VPD_fig
cap_osml_VPD_fig

```

Warning: Removed 14 rows containing non-finite values (stat_smooth).

Warning: Removed 14 rows containing missing values (geom_point).



Osmolality ~ Solar Radiation at Capture

```

ggplot(dat) +
  aes(x = solar_rad_W_sqm_interpol,
       y = osmolality_mmol_kg_mean) +
  geom_point(size = 1,
             alpha = 0.4) +
  stat_smooth(formula = y ~ x,

```

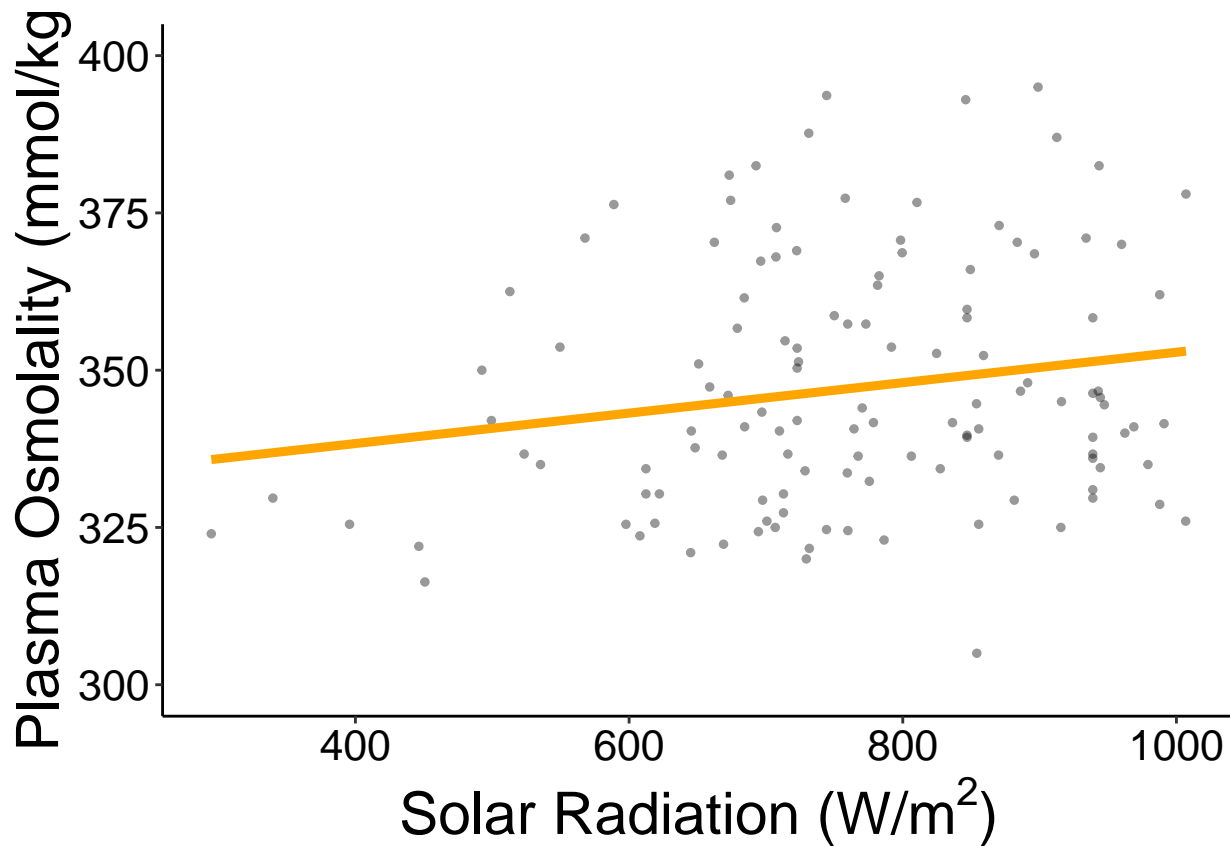
```

      method = "lm",
      se = F,
      color = solar_color,
      size = 1.6,
      alpha = 1 ) +
theme_classic() +
xlab(bquote('Solar Radiation (W/*m^2*')')) +
ylab("Plasma Osmolality (mmol/kg)") +
#ylab("") +
ylim(300,400) +
theme(text = element_text(color = "black",
                           family = "sans",
                           size = 22),
      axis.text = element_text(color = "black",
                                family = "sans",
                                size = 16),
      #axis.text.y = element_blank(),
      #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
    ) -> cap_osml_sorad_fig
cap_osml_sorad_fig

```

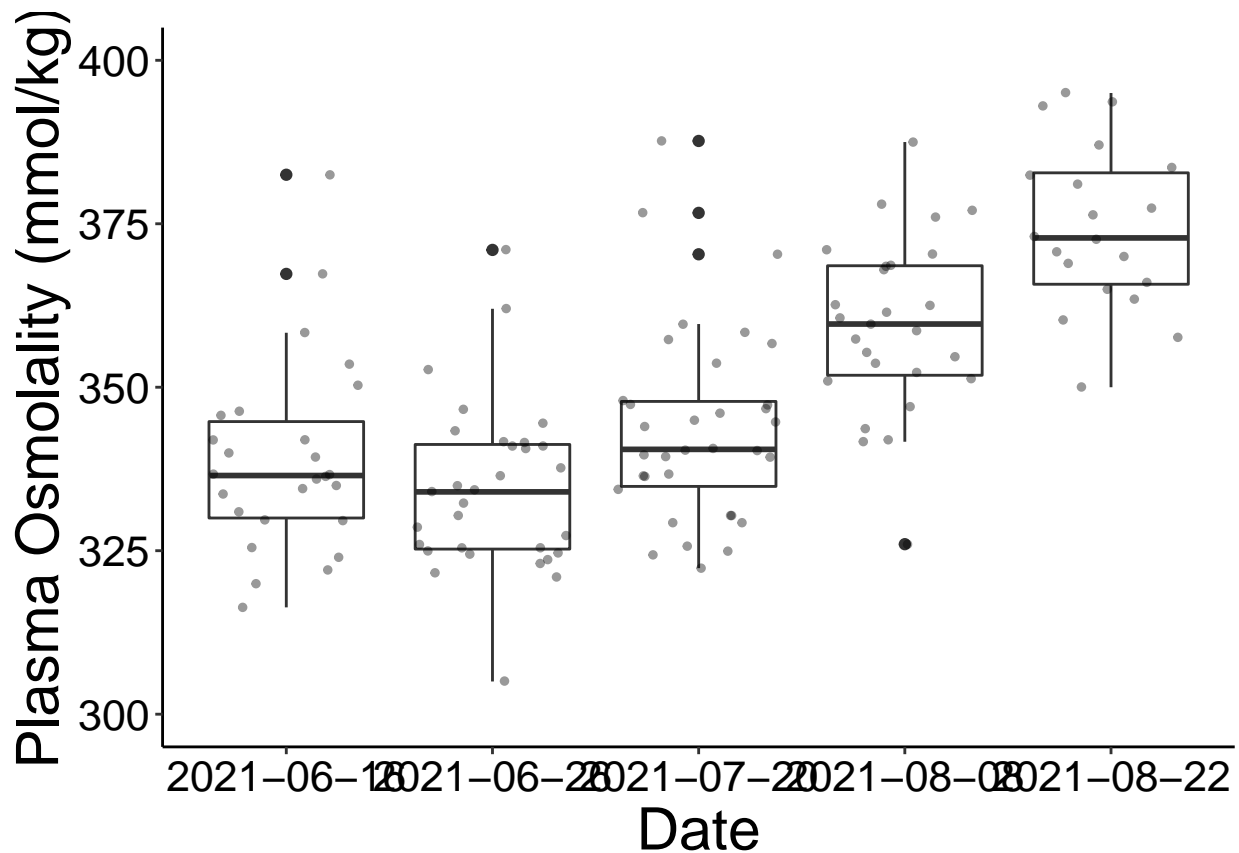
Warning: Removed 14 rows containing non-finite values (stat_smooth).

Warning: Removed 14 rows containing missing values (geom_point).



Osmolality ~ Date

```
ggplot(dat) +
  aes(x = as.factor(capture_date),
       y = osmolality_mmol_kg_mean,
       group = as.factor(capture_date)) +
  geom_boxplot() +
  geom_jitter(size = 1,
              alpha = 0.4) +
  theme_classic() +
  xlab("Date") +
  ylab("Plasma Osmolality (mmol/kg)") +
  #ylab("") +
  #xlim() +
  ylim(300, 400) +
  #annotate("text", x = , y = ,
  #             label = "paste(italic(R) ^ 2, \" = 0.\")",
  #             parse = TRUE,
  #             size = 6) +
  #annotate("text", x = , y = ,
  #             label = "paste(italic(p), \" < 0.0001\")",
  #             parse = TRUE,
  #             size = 6) +
  theme(text = element_text(color = "black",
                             family = "sans",
                             size = 22),
        axis.text = element_text(color = "black",
                                   family = "sans",
                                   size = 16),
        #axis.text.y = element_blank(),
        #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
  ) -> cap_osml_date_fig
cap_osml_date_fig
```

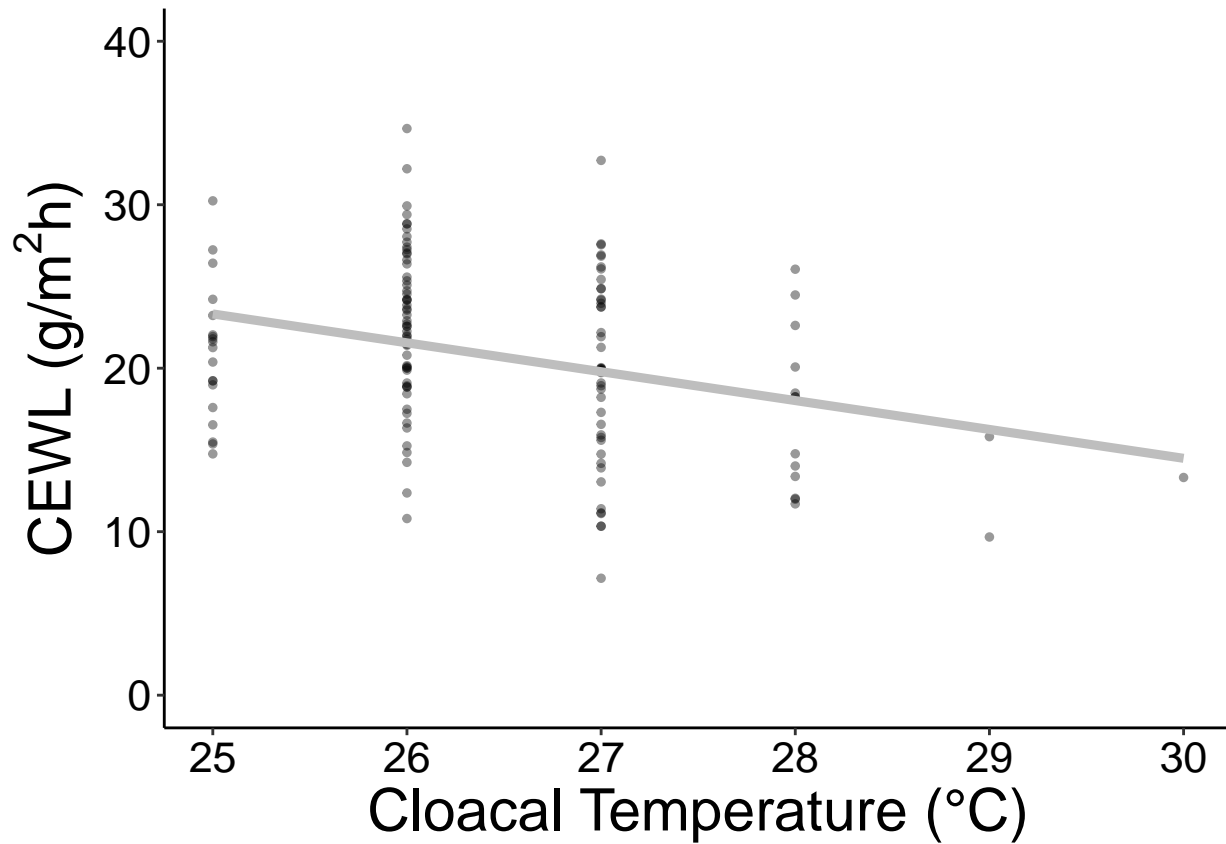
CEWL ~ Cloacal Temperature

```
ggplot(dat) +
  aes(x = cloacal_temp_C,
       y = CEWL_g_m2h_mean) +
  geom_point(size = 1,
             alpha = 0.4) +
  stat_smooth(formula = y ~ x,
              method = "lm",
              se = F,
              color = temp_color,
              size = 1.6,
              alpha = 1) +
  theme_classic() +
  xlab("Cloacal Temperature (°C)") +
  ylab(bquote('CEWL (g/'*m^2*'h)')) +
  #ylab("") +
  #xlim() +
  ylim(0, 40) +
  theme(text = element_text(color = "black",
                             family = "sans",
                             size = 22),
        axis.text = element_text(color = "black",
                                  family = "sans",
                                  size = 16),
        #axis.text.y = element_blank(),
```

```

#plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
) -> cap_CEWL_clotemp_fig
cap_CEWL_clotemp_fig

```



CEWL ~ Plasma Osmolality

```

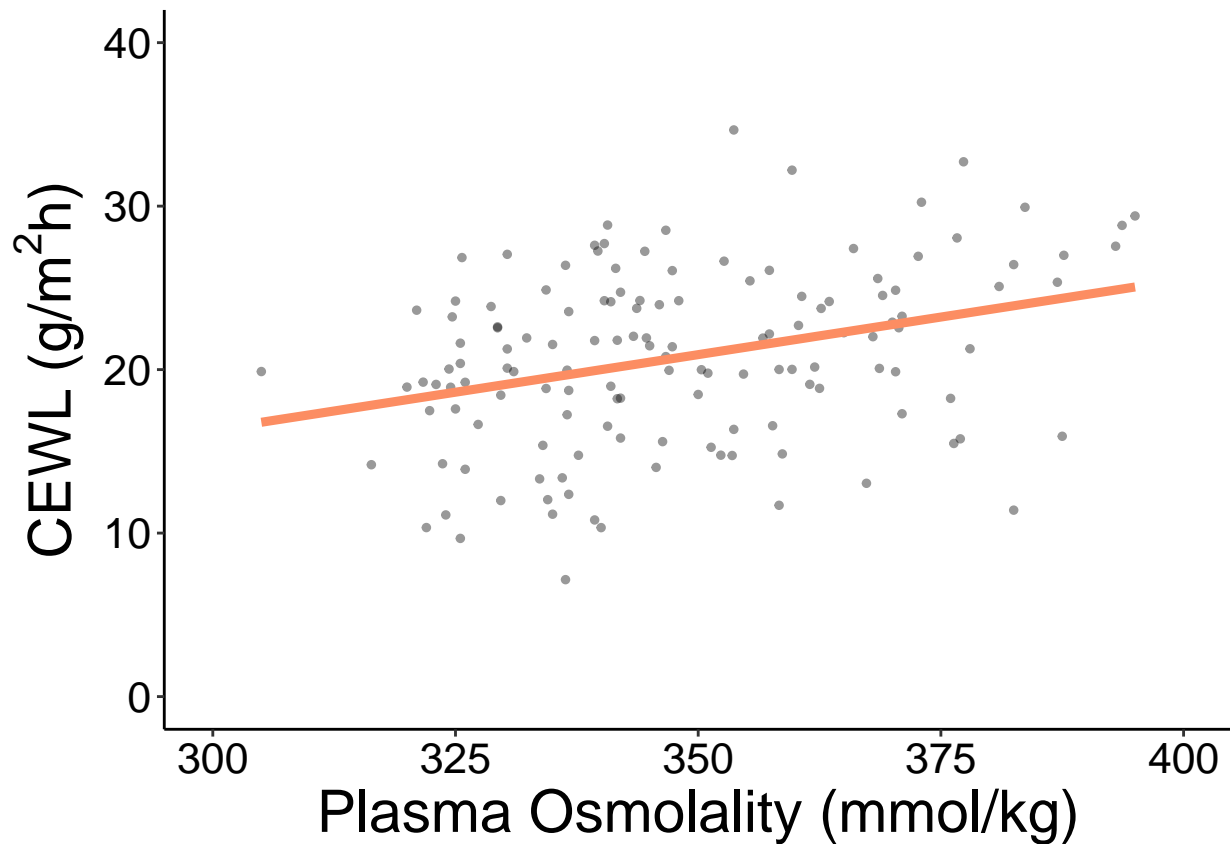
ggplot(dat) +
  aes(x = osmolality_mmol_kg_mean,
      y = CEWL_g_m2h_mean) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
            method = "lm",
            se = F,
            color = osml_color,
            size = 1.6,
            alpha = 1) +
  theme_classic() +
  xlab("Plasma Osmolality (mmol/kg)") +
  ylab(bquote('CEWL (g/'*m2*'h)')) +
  #ylab("") +
  xlim(300, 400) +
  ylim(0, 40) +
  # manual_color_brewer(, name = "") +
  theme(text = element_text(color = "black",
                            family = "sans",

```

```

        size = 22),
axis.text = element_text(color = "black",
                          family = "sans",
                          size = 16),
#axis.text.y = element_blank(),
#plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
) -> cap_CEWL_osml_fig
cap_CEWL_osml_fig

```



```

ggsave(filename = "cap_CEWL_osml_fig.jpeg",
plot = cap_CEWL_osml_fig,
path = "./results_figures",
device = "jpeg",
dpi = 600,
width = 6, height = 4)

```

CEWL ~ Temperature at Measurement

```

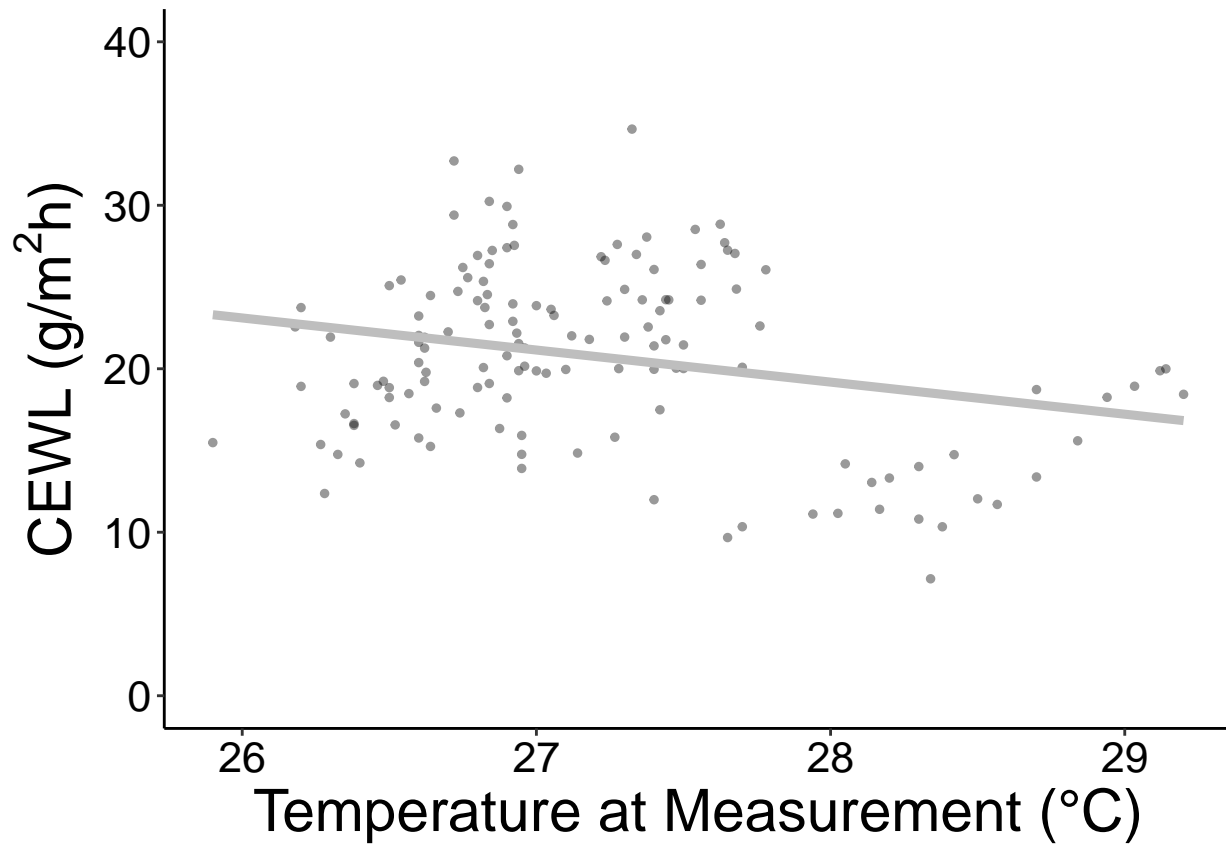
ggplot(dat) +
  aes(x = msmt_temp_C,
      y = CEWL_g_m2h_mean) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
            method = "lm",
            se = F,

```

```

    color = temp_color,
    size = 1.6,
    alpha = 1) +
  theme_classic() +
  xlab("Temperature at Measurement (°C)") +
  ylab(bquote('CEWL (g/'*m2*h)')) +
  #ylab("") +
  #xlim() +
  ylim(0, 40) +
  theme(text = element_text(color = "black",
                             family = "sans",
                             size = 22),
        axis.text = element_text(color = "black",
                                  family = "sans",
                                  size = 16),
        #axis.text.y = element_blank(),
        #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
  ) -> cap_CEWL_temp_fig
cap_CEWL_temp_fig

```



CEWL ~ VPD at Measurement

```

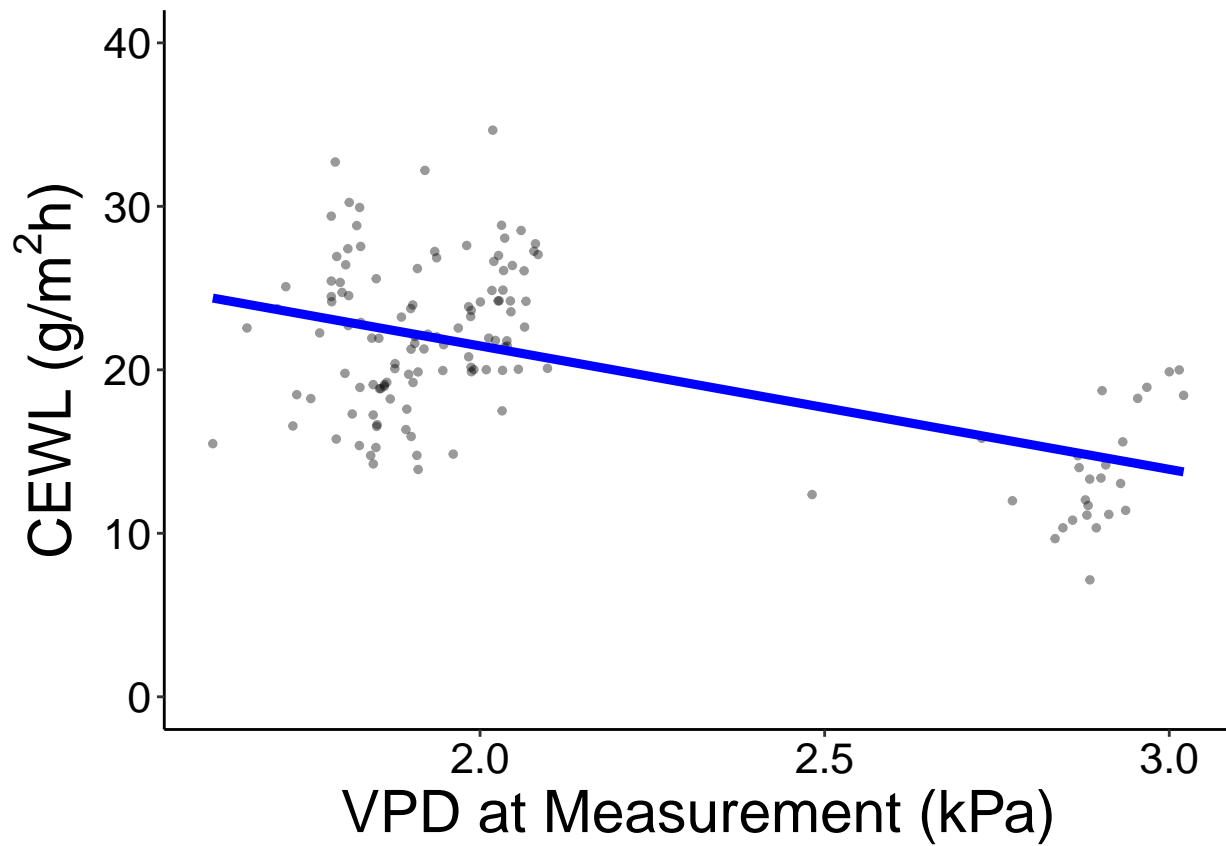
ggplot(dat) +
  aes(x = msmt_VPD_kPa,
       y = CEWL_g_m2h_mean) +
  geom_point(size = 1,

```

```

      alpha = 0.4) +
stat_smooth(formula = y ~ x,
            method = "lm",
            se = F,
            color = VPD_color,
            size = 1.6,
            alpha = 1 ) +
theme_classic() +
xlab("VPD at Measurement (kPa)") +
ylab(bquote('CEWL (g/'*m^2*'h)')) +
#ylab("") +
#xlim() +
ylim(0, 40) +
theme(text = element_text(color = "black",
                           family = "sans",
                           size = 22),
      axis.text = element_text(color = "black",
                                family = "sans",
                                size = 16),
      #axis.text.y = element_blank(),
      #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
    ) -> cap_CEWL_VPDm_fig
cap_CEWL_VPDm_fig

```

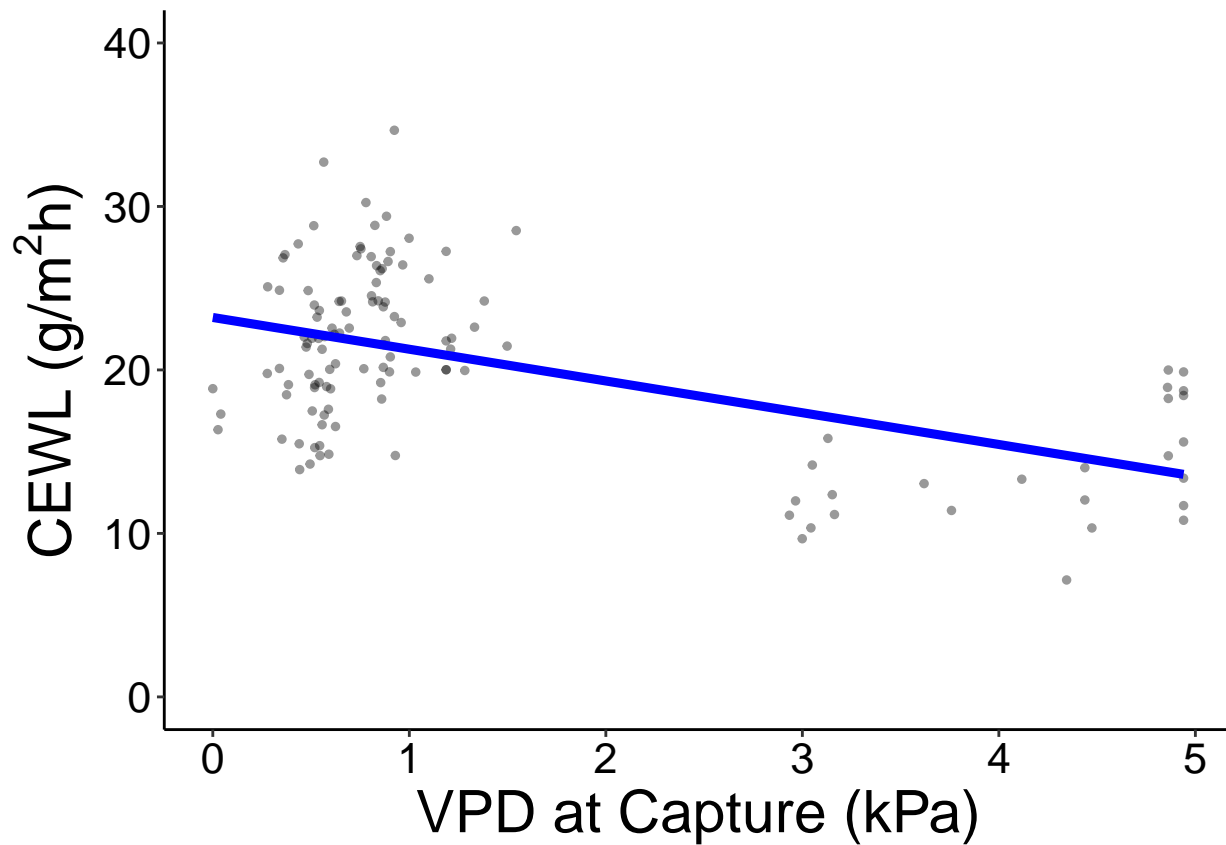


CEWL ~ VPD at Capture

```
ggplot(dat) +
  aes(x = VPD_kPa_int,
      y = CEWL_g_m2h_mean) +
  geom_point(size = 1,
            alpha = 0.4) +
  stat_smooth(formula = y ~ x,
             method = "lm",
             se = F,
             color = VPD_color,
             size = 1.6,
             alpha = 1) +
  theme_classic() +
  xlab("VPD at Capture (kPa)") +
  ylab(bquote('CEWL (g/'*m^2*'h)')) +
  #ylab("") +
  #xlim() +
  ylim(0, 40) +
  theme(text = element_text(color = "black",
                            family = "sans",
                            size = 22),
        axis.text = element_text(color = "black",
                                  family = "sans",
                                  size = 16),
        #axis.text.y = element_blank(),
        #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
  ) -> cap_CEWL_VPDc_fig
cap_CEWL_VPDc_fig
```

Warning: Removed 14 rows containing non-finite values (stat_smooth).

Warning: Removed 14 rows containing missing values (geom_point).



CEWL ~ Wind at Capture

```
ggplot(dat) +
  aes(x = wind_mph_interpol,
       y = CEWL_g_m2h_mean) +
  geom_point(size = 1,
             alpha = 0.4) +
  stat_smooth(formula = y ~ x,
              method = "lm",
              se = F,
              color = wind_color,
              size = 1.6,
              alpha = 1) +
  theme_classic() +
  xlab("Wind Speed at Capture (mph)") +
  ylab(bquote('CEWL (g/'*m^2*'h)')) +
  #ylab("") +
  #xlim() +
  ylim(0, 40) +
  theme(text = element_text(color = "black",
                             family = "sans",
                             size = 22),
        axis.text = element_text(color = "black",
                                  family = "sans",
                                  size = 16),
        #axis.text.y = element_blank(),
```

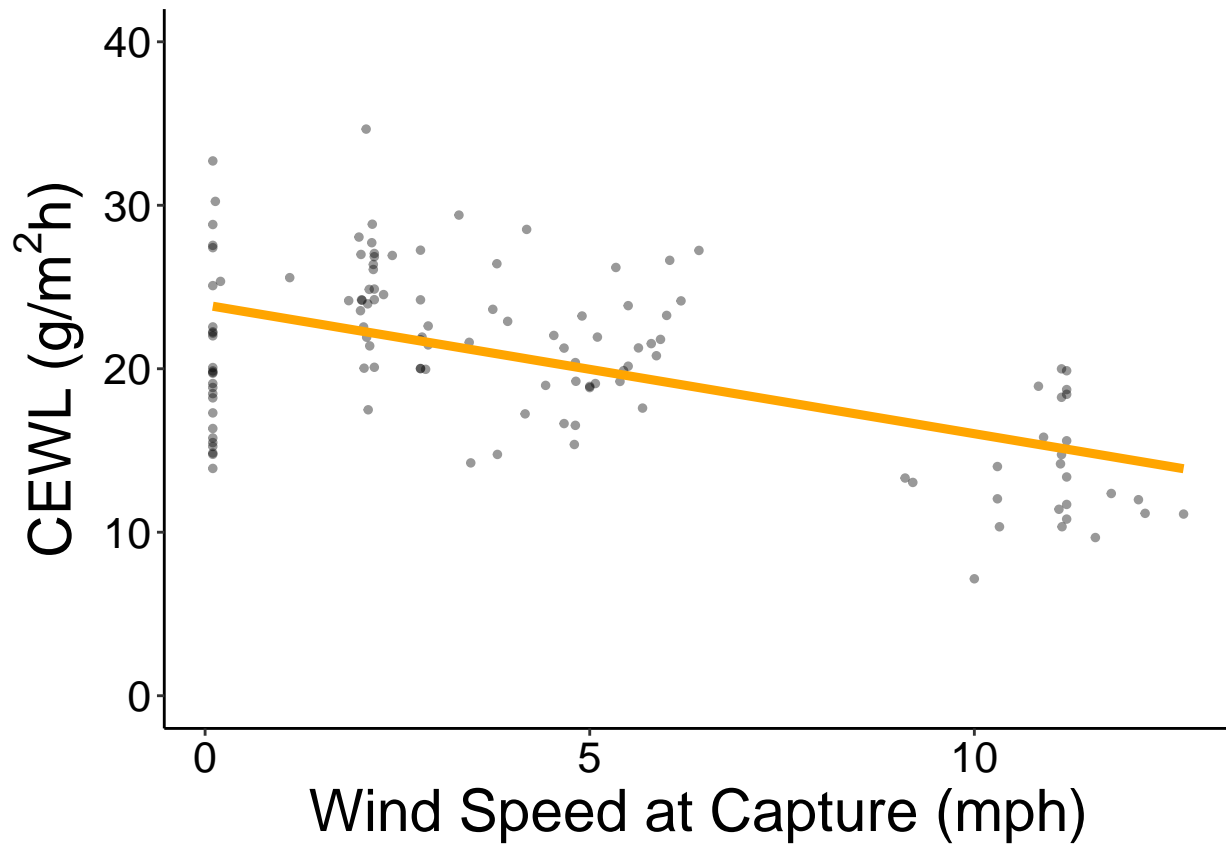
```

    #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
  ) -> cap_CEWL_wind_fig
cap_CEWL_wind_fig

```

```
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 14 rows containing missing values (geom_point).
```



CEWL ~ Date

```

ggplot(dat) +
  aes(x = as.factor(capture_date),
      y = CEWL_g_m2h_mean,
      group = as.factor(capture_date)) +
  geom_boxplot() +
  geom_jitter(size = 1,
              alpha = 0.4) +
  theme_classic() +
  xlab("Date") +
  ylab(bquote('CEWL (g/' * m^2 * 'h)')) +
  #ylab("") +
  #xlim() +
  ylim(0, 40) +
  #annotate("text", x = , y = ,
  #          label = "paste(italic(R) ^ 2, \" = 0.\")",
  #          parse = TRUE,
  #          size = 6) +

```



```

#annotate("text", x = , y = ,
#         label = "paste(italic(p), \" < 0.0001\")",
#         parse = TRUE,
#         size = 6) +
theme(text = element_text(color = "black",
                           family = "sans",
                           size = 22),
      axis.text = element_text(color = "black",
                                family = "sans",
                                size = 16),
      #axis.text.y = element_blank(),
      #plot.margin = unit(c(0.1,0,0.1,0.45), "cm")
    ) -> cap_CEWL_date_fig
cap_CEWL_date_fig

```

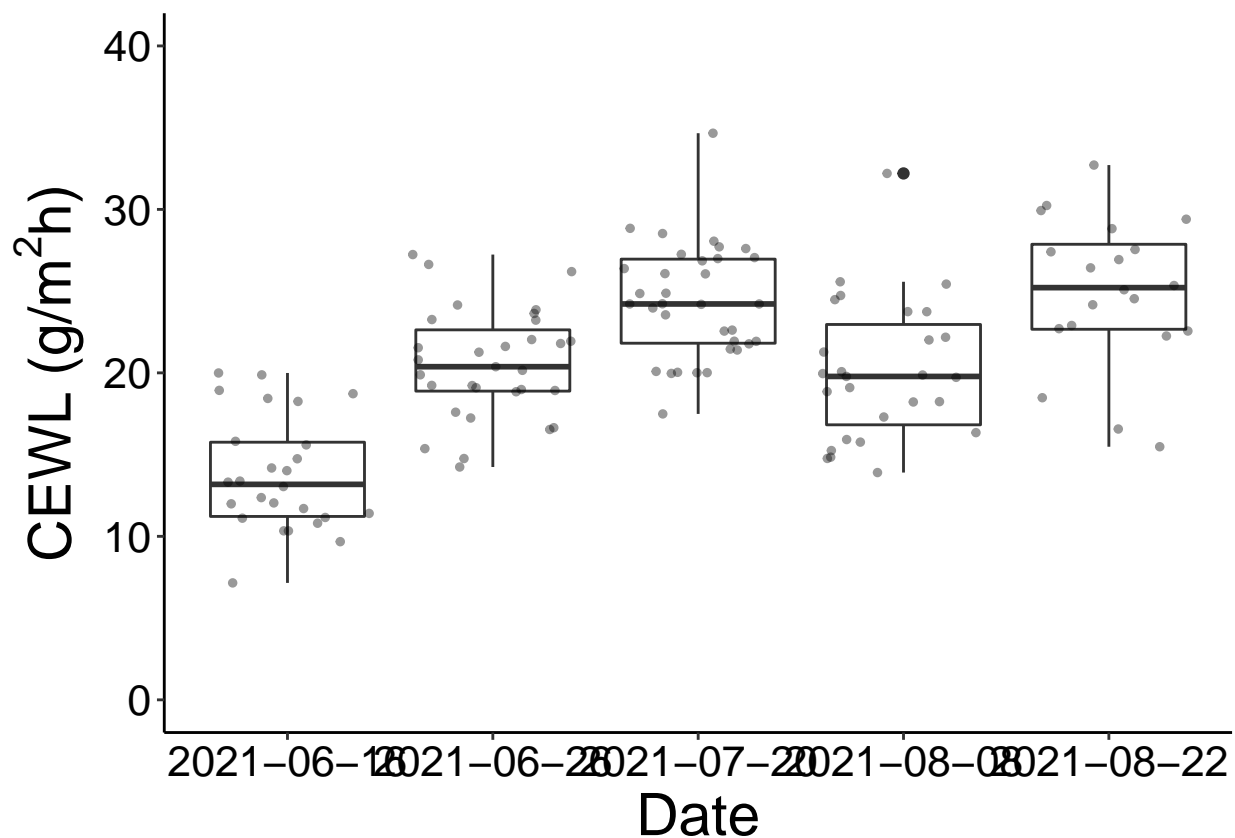


Figure Arrangements

```

# hematocrit
ggarrange(cap_hct_sorad_fig, cap_hct_VPD_fig,
          cap_hct_SMI_fig, cap_hct_wind_fig,
          ncol = 2, nrow = 2,
          legend = "none"
        ) -> cap_hct_multi_fig

```

```
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
```

```
## Warning: Removed 14 rows containing missing values (geom_point).
```

```
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
## Warning: Removed 14 rows containing missing values (geom_point).
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
## Warning: Removed 14 rows containing missing values (geom_point).
```

```
#cap_hct_multi_fig
# export figure
#ggsave(filename = "cap_hct_multi_fig.jpeg",
#       plot = cap_hct_multi_fig,
#       path = "./results_figures",
#       device = "jpeg",
#       dpi = 1200,
#       width = 12, height = 8)

# osmolality
ggarrange(cap_osml_sorad_fig, cap_osml_VPD_fig,
          cap_osml_SMI_fig, cap_osml_SVL_fig,
          ncol = 2, nrow = 2,
          legend = "none"
          ) -> cap_osml_multi_fig
```

```
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
## Warning: Removed 14 rows containing missing values (geom_point).
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
## Warning: Removed 14 rows containing missing values (geom_point).
```

```
#cap_osml_multi_fig
# export figure
#ggsave(filename = "cap_osml_multi_fig.jpeg",
#       plot = cap_osml_multi_fig,
#       path = "./results_figures",
#       device = "jpeg",
#       dpi = 1200,
#       width = 12, height = 8)

# CEWL
ggarrange(cap_CEWL_VPDm_fig, cap_CEWL_VPDc_fig,
          cap_CEWL_temp_fig, cap_CEWL_wind_fig,
          cap_CEWL_osml_fig, cap_CEWL_clotemp_fig,
          ncol = 2, nrow = 3,
          legend = "none"
          ) -> cap_CEWL_multi_fig
```

```
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
## Warning: Removed 14 rows containing missing values (geom_point).
## Warning: Removed 14 rows containing non-finite values (stat_smooth).
## Warning: Removed 14 rows containing missing values (geom_point).
```

```

#cap_CEWL_multi_fig
# export figure
#ggsave(filename = "cap_CEWL_multi_fig",
#       plot = cap_CEWL_multi_fig,
#       path = "./results_figures",
#       device = "jpeg",
#       dpi = 1200,
#       width = 12, height = 16)

# date differences
ggarrange(cap_osml_date_fig, cap_CEWL_date_fig,
          ncol = 1, nrow = 2,
          legend = "none"
          ) -> cap_date_diffs_multi_fig
##cap_date_diffs_multi_fig
# export figure
#ggsave(filename = "cap_date_diffs_multi_fig.jpeg",
#       plot = cap_date_diffs_multi_fig,
#       path = "./results_figures",
#       device = "jpeg",
#       dpi = 1200,
#       width = 6, height = 8)

```