# Autocorrelation stats

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# Load and Preprocess Datasets

We load two datasets: one with a preparation technique (repeated frond selection) and one without.

```
# This dataset contains replicates for which a preparation technique was performed (repeated first born
original_dataset_2 <- read.csv("https://raw.githubusercontent.com/Cuddington-Lab/thermal-experiments/ma
original_dataset_2$prep <- rep("yes",times=length(original_dataset_2$Experiment_Number))

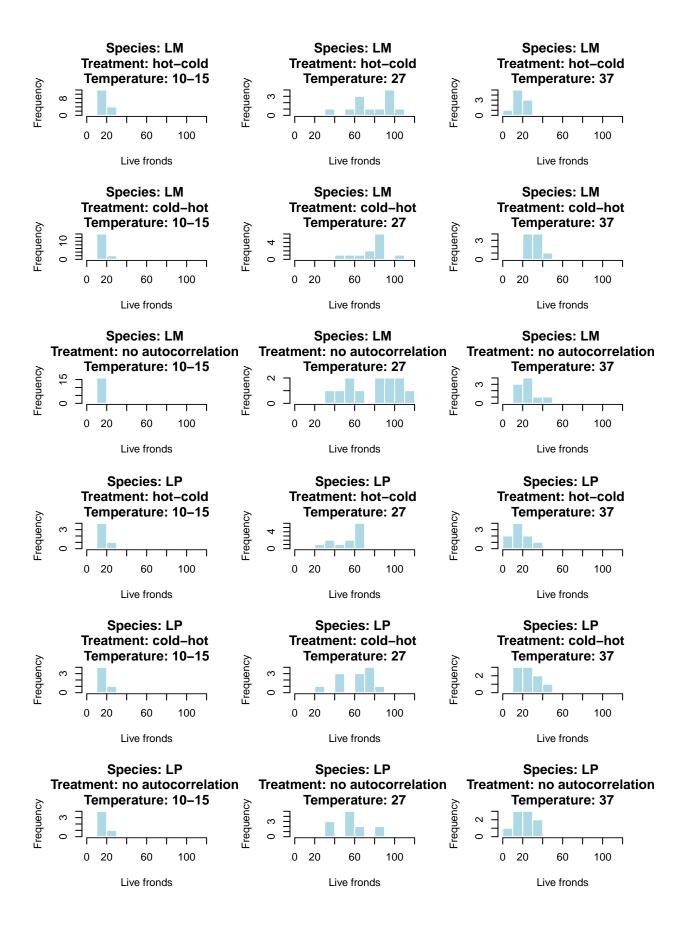
# This dataset contains replicates of experiments performed without a preparation technique
original_dataset_1 <- read.csv("https://raw.githubusercontent.com/Cuddington-Lab/thermal-experiments/ma
original_dataset_1$prep <- rep("no",times=length(original_dataset_1$Experiment_Number))</pre>
```

#### **Combine Datasets**

We combine both datasets and filter out rows based on specific conditions for standard deviation (Obs\_sd) and autocorrelation (Obs ac) to clean the data.

# View dataset and response variable

	Species	Exp_run	Treatment	total_living_fronds
395	LM	2733	cold-hot	90
396	$_{ m LP}$	2733	cold-hot	74
397	LM	2733	hot-cold	91
398	$_{ m LP}$	2733	hot-cold	63
399	LM	2733	no autocorrelation	101
400	$_{ m LP}$	2733	no autocorrelation	88



# View number of replicates

```
LM
##
##
           no autocorrelation cold-hot hot-cold
##
     10-15
                             16
                                       16
##
     27
                             12
                                       12
                                                 12
##
     37
                              9
                                        9
                                                  9
LP
##
##
           no autocorrelation cold-hot hot-cold
##
     10-15
                              5
                                        5
##
     27
                             12
                                       12
                                                 12
                                        9
##
     37
                                                  9
```

## Mixed-Effects Model Fitting

We fit candidate simple or mixed-effects models to the data, including random effects for Experiment run (Exp\_run).

```
## [1] "Comparing simple x mixed model:"
## p_value: 0
## [1] "Comparing simple x mixed model:"
## p_value: 0
```

# Model Comparison and Results

We compare the performance of the models and extract estimates, confidence intervals, and p values.

```
## [1] "Model comparison for species: LM"
##
##
##
                        Name
                                                 AIC
##
## no random effects
                       no random effects
                                              892.90
## experiment number
                        experiment number
                                              772.12
   [1] "Model comparison for species: LP"
##
##
##
                        Name
                                                 AIC
                                              793.81
## no random effects
                        no random effects
                                              680.89
## experiment number
                        experiment number
```

### Model significance testing

```
## Treatment:Mean_temperature
                               35.9543 4
                                                  0.0000002957 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## [1] "Residual degrees of freedom - species : LM : 101"
## Wald test type 3 for significance of predictor: LP
## Analysis of Deviance Table (Type III Wald chisquare tests)
## Response: total_living_fronds
##
                                Chisq Df
                                                  Pr(>Chisq)
## (Intercept)
                             319.8839 1 < 0.0000000000000000 ***
## Treatment
                               0.8332 2
                                                      0.6593
                                       2 <0.000000000000000 ***
                              85.9939
## Mean_temperature
## Treatment:Mean_temperature
                              4.4267 4
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## [1] "Residual degrees of freedom - species : LP : 68"
```

#### R2 for common duckweeds

```
## Marginal R2: NA
## Conditional R2: NA
```

#### Visualization of Results for common duckweeds

```
## Generalized linear mixed model fit by maximum likelihood (Laplace
     Approximation) [glmerMod]
  Family: poisson (log)
## Formula: total_living_fronds ~ Treatment * Mean_temperature + (1 | Exp_run)
      Data: dataset_species
## Control: glmerControl(optimizer = "nloptwrap")
##
##
       AIC
                 BIC
                       logLik deviance df.resid
##
      772.1
              799.2
                       -376.1
                                 752.1
                                            101
##
## Scaled residuals:
##
      Min
               1Q Median
                                3Q
                                       Max
## -3.3375 -0.3678 -0.0017 0.3794 3.6435
##
## Random effects:
## Groups Name
                        Variance Std.Dev.
## Exp_run (Intercept) 0.03766 0.1941
## Number of obs: 111, groups: Exp_run, 37
## Fixed effects:
##
                                        Estimate Std. Error z value
## (Intercept)
                                         2.82097
                                                    0.07771 36.303
## Treatmentcold-hot
                                                    0.08459
                                                              0.638
                                         0.05397
## Treatmenthot-cold
                                         0.06070
                                                    0.08446
                                                              0.719
## Mean_temperature27
                                         1.51541
                                                    0.10126
                                                             14.965
## Mean temperature37
                                         0.46435
                                                    0.11965
                                                              3.881
## Treatmentcold-hot:Mean_temperature27 -0.05601
                                                    0.09635 -0.581
## Treatmenthot-cold:Mean_temperature27 -0.05632
                                                    0.09619
                                                             -0.585
## Treatmentcold-hot:Mean_temperature37  0.07823
                                                    0.12134
                                                             0.645
## Treatmenthot-cold:Mean_temperature37 -0.55293
                                                    0.13329 - 4.148
```

```
Pr(>|z|)
##
## (Intercept)
                                       < 0.0000000000000002 ***
## Treatmentcold-hot
                                                  0.523492
## Treatmenthot-cold
                                                  0.472348
## Mean temperature27
                                       < 0.000000000000000 ***
## Mean temperature37
                                                  0.000104 ***
## Treatmentcold-hot:Mean temperature27
                                                  0.560991
## Treatmenthot-cold:Mean_temperature27
                                                  0.558218
## Treatmentcold-hot:Mean_temperature37
                                                  0.519091
## Treatmenthot-cold:Mean_temperature37
                                                 0.0000335 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation of Fixed Effects:
##
                (Intr) Trtmntc- Trtmnth- Mn_t27 Mn_t37 Trtmntc-:M_27
## Trtmntcld-h
                -0.559
## Trtmntht-cl
                -0.560 0.514
                               0.430
## Mn tmprtr27
                -0.767 0.429
## Mn_tmprtr37
                -0.649 0.363
                               0.364
                                          0.498
## Trtmntc-:M 27 0.491 -0.878
                                -0.452
                                         -0.486 - 0.319
## Trtmnth-:M_27  0.492  -0.452  -0.878  -0.486  -0.319  0.511
## Trtmntc-:M 37 0.390 -0.697
                                -0.359 -0.299 -0.531 0.612
## Trtmnth-:M_37 0.355 -0.326
                                -0.634 -0.272 -0.483 0.286
                Trtmnth-:M_27 Trtmntc-:M_37
## Trtmntcld-h
## Trtmntht-cl
## Mn_tmprtr27
## Mn_tmprtr37
## Trtmntc-:M_27
## Trtmnth-:M 27
## Trtmntc-:M_37 0.315
## Trtmnth-:M_37 0.556
                               0.477
## optimizer (nloptwrap) convergence code: 0 (OK)
## Model failed to converge with max|grad| = 0.00674263 (tol = 0.002, component 1)
Model validation for common duckweeds
## [1] "Model dispersion ratio : 0.780382048699719"
## [1] "Shapiro test for normality of random factors 0.051"
Post-hoc test for common duckweeds
## [1] "Pairwise interactions"
## Mean_temperature = 10-15:
## Treatment
                          rate
                                     SE df asymp.LCL asymp.UCL
## no autocorrelation 16.79316 1.304921 Inf
                                            14.42080
                                                     19.55580
                    17.72437 1.355084 Inf
                                            15.25786 20.58959
## cold-hot
                      17.84400 1.361503 Inf 15.36545 20.72236
## hot-cold
##
## Mean_temperature = 27:
## Treatment
                          rate
                                     SE df asymp.LCL asymp.UCL
## no autocorrelation 76.43063 4.964703 Inf 67.29395 86.80784
## cold-hot
                     76.27457 4.955841 Inf 67.15434 86.63343
```

```
76.76596 4.983742 Inf 67.59391 87.18258
## hot-cold
##
## Mean temperature = 37:
                                     SE df asymp.LCL asymp.UCL
  Treatment
                          rate
## no autocorrelation 26.71753 2.431967 Inf 22.35195 31.93574
## cold-hot
                     30.49379 2.690734 Inf 25.65089 36.25103
## hot-cold
                      16.33133 1.701186 Inf 13.31540 20.03035
##
## Confidence level used: 0.95
## Intervals are back-transformed from the log scale
## Mean_temperature = 10-15:
## contrast
                                   ratio
                                            SE df null z.ratio p.value
## no autocorrelation / (cold-hot) 0.947 0.0801 Inf
                                                      1 -0.638 0.7992
## no autocorrelation / (hot-cold) 0.941 0.0795 Inf
                                                      1 -0.719 0.7524
   (cold-hot) / (hot-cold)
##
                                   0.993 0.0827 Inf
                                                      1 -0.081 0.9964
##
## Mean_temperature = 27:
## contrast
                                   ratio
                                            SE df null z.ratio p.value
## no autocorrelation / (cold-hot) 1.002 0.0462 Inf
                                                      1
                                                          0.044 0.9989
## no autocorrelation / (hot-cold) 0.996 0.0458 Inf
                                                      1 -0.095 0.9950
## (cold-hot) / (hot-cold)
                                   0.994 0.0458 Inf
                                                      1 -0.139 0.9893
##
## Mean temperature = 37:
## contrast
                                            SE df null z.ratio p.value
                                   ratio
## no autocorrelation / (cold-hot) 0.876 0.0762 Inf
                                                      1 -1.520 0.2816
## no autocorrelation / (hot-cold) 1.636 0.1687 Inf
                                                      1
                                                          4.774 <.0001
   (cold-hot) / (hot-cold)
                                   1.867 0.1880 Inf
##
                                                      1
                                                          6.203 <.0001
##
## P value adjustment: tukey method for comparing a family of 3 estimates
## Tests are performed on the log scale
```

