

Meta-analysis Manuscript Wrangling

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```
## Warning: 1 components of '...' were not used.  
##  
## We detected these problematic arguments:  
## * 'method'  
##  
## Did you misspecify an argument?
```

Motivation and Hypotheses

Variability has been identified as important to organismal success and ecosystem dynamics (Vasseur et al 2014). To further understand the impacts of variability of performance, we conducted to analyses on data that explicitly accounted for acclimation (Acclimation model) and studies focuses solely on non-linear averaging (Non-linear averaging model).

H-Acclimation: If reared in fluctuating environments, when exposed to different thermal environments, organisms with larger fluctuation ranges will perform worse than those reared in constant environments. Additional covariates, such as age, size, and expsoure temperature will be correlated negative responses.

H-Non-linear averaging: Organisms will perform better in constant environments than fluctuating environments. Additional covariates, such as age, size, and large fluctuation range will be correlated negative responses.

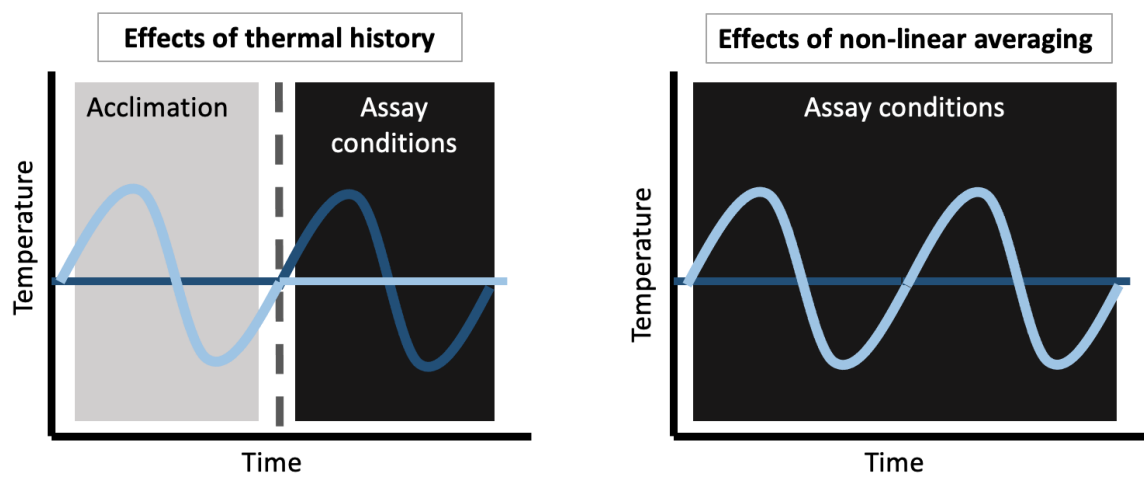


Figure 1: Conceptual figure demonstrating how acclimation and non-linear averaging account for different aspects of performance.

Acclimation Model

This data and analysis accounts for the temperatures at which organisms are reared and how their performance compares when exposed to different temperature. The results from this model has very similar results from the model I originally ran for my thesis...

Acclimation with temperature modifiers

```
##
## Multivariate Meta-Analysis Model (k = 332; method: REML)
##
## Variance Components:
##
##          estim      sqrt  nlvls  fixed          factor
## sigma^2.1 0.0000 0.0002    11    no          study_id
## sigma^2.2 0.4359 0.6602    33    no      study_id/experiment_id
## sigma^2.3 0.2693 0.5190    61    no  study_id/experiment_id/response_id
##
## Test for Residual Heterogeneity:
## QE(df = 328) = 3625.8727, p-val < .0001
##
## Test of Moderators (coefficients 2:4):
## QM(df = 3) = 122.6186, p-val < .0001
##
## Model Results:
##
##              estimate      se      zval      pval      ci.lb
## intrcpt              4.9972 0.9940   5.0274 <.0001   3.0490
## flux_range          -0.4634 0.0989  -4.6872 <.0001  -0.6572
## mean_temp_reared    -0.2538 0.0439  -5.7870 <.0001  -0.3398
## flux_range:mean_temp_reared  0.0220 0.0044   5.0383 <.0001   0.0135
##
##              ci.ub
## intrcpt          6.9454 ***
## flux_range       -0.2696 ***
## mean_temp_reared -0.1678 ***
## flux_range:mean_temp_reared  0.0306 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Acclimation with trait modifiers

```
## Warning: Rows with NAs omitted from model fitting.
```

```
## Warning: Redundant predictors dropped from the model.
```

```
##
## Multivariate Meta-Analysis Model (k = 331; method: REML)
##
## Variance Components:
##
##          estim      sqrt  nlvls  fixed          factor
```

```
## sigma^2.1 0.0000 0.0001 11 no study_id
## sigma^2.2 0.4994 0.7067 33 no study_id/experiment_id
## sigma^2.3 0.2272 0.4767 60 no study_id/experiment_id/response_id
##
## Test for Residual Heterogeneity:
## QE(df = 327) = 3704.6953, p-val < .0001
##
## Test of Moderators (coefficients 2:4):
## QM(df = 3) = 3.6051, p-val = 0.3074
##
## Model Results:
##
## estimate se zval pval ci.lb ci.ub
## intrcpt 0.9898 0.9078 1.0904 0.2755 -0.7894 2.7690
## exp_age -0.1968 0.3207 -0.6137 0.5394 -0.8254 0.4317
## size -0.7382 0.4090 -1.8047 0.0711 -1.5399 0.0635 .
## exposure_temp 0.0010 0.0031 0.3254 0.7449 -0.0050 0.0070
##
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Acclimation with all modifiers

```
## Warning: Rows with NAs omitted from model fitting.
```

```
## Warning: Redundant predictors dropped from the model.
```

```
##
## Multivariate Meta-Analysis Model (k = 331; method: REML)
##
## Variance Components:
##
## estim sqrt nlvls fixed factor
## sigma^2.1 0.0000 0.0001 11 no study_id
## sigma^2.2 0.4698 0.6854 33 no study_id/experiment_id
## sigma^2.3 0.2474 0.4974 60 no study_id/experiment_id/response_id
##
## Test for Residual Heterogeneity:
## QE(df = 324) = 3580.2892, p-val < .0001
##
## Test of Moderators (coefficients 2:7):
## QM(df = 6) = 127.0957, p-val < .0001
##
## Model Results:
##
## estimate se zval pval ci.lb
## intrcpt 6.0093 1.3387 4.4889 <.0001 3.3855
## flux_range -0.4757 0.0991 -4.7983 <.0001 -0.6700
## mean_temp_reared -0.2595 0.0440 -5.9001 <.0001 -0.3457
## exp_age -0.0836 0.3184 -0.2626 0.7929 -0.7078
## size -0.6691 0.4072 -1.6432 0.1003 -1.4671
## exposure_temp 0.0032 0.0031 1.0098 0.3126 -0.0030
## flux_range:mean_temp_reared 0.0226 0.0044 5.1550 <.0001 0.0140
```

```

##                                ci.ub
## intrcpt                      8.6331 ***
## flux_range                   -0.2814 ***
## mean_temp_reared             -0.1733 ***
## exp_age                      0.5405
## size                         0.1290
## exposure_temp                0.0093
## flux_range:mean_temp_reared  0.0312 ***
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Effects of non-linear averaging

This uses the full dataset from my thesis (with more data from the additional search). Interesting that this time, mean temp is the only significant predictor?

Non-linear averaging model with temperature modifiers

```

##
## Multivariate Meta-Analysis Model (k = 366; method: REML)
##
## Variance Components:
##
##      estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0000  0.0008    28    no      study_id
## sigma^2.2  0.2561  0.5061    45    no      study_id/experiment_id
## sigma^2.3  0.6116  0.7821   100    no      study_id/experiment_id/response_id
##
## Test for Residual Heterogeneity:
## QE(df = 362) = 6776.0661, p-val < .0001
##
## Test of Moderators (coefficients 2:4):
## QM(df = 3) = 95.1905, p-val < .0001
##
## Model Results:
##
##      estimate      se      zval      pval      ci.lb
## intrcpt          1.1548  0.3148   3.6686  0.0002   0.5378
## flux_range       -0.0207  0.0221  -0.9397  0.3474  -0.0640
## mean_temp_constant -0.0465  0.0127  -3.6559  0.0003  -0.0714
## flux_range:mean_temp_constant  0.0012  0.0010   1.2075  0.2272  -0.0007
##
##      ci.ub
## intrcpt          1.7718 ***
## flux_range          0.0225
## mean_temp_constant -0.0216 ***
## flux_range:mean_temp_constant  0.0031
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

Non-linear averaging model with trait modifiers

```
## Warning: Rows with NAs omitted from model fitting.

##
## Multivariate Meta-Analysis Model (k = 359; method: REML)
##
## Variance Components:
##
##      estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0000  0.0004    27    no      study_id
## sigma^2.2  0.2846  0.5335    44    no      study_id/experiment_id
## sigma^2.3  0.4882  0.6987    98    no      study_id/experiment_id/response_id
##
## Test for Residual Heterogeneity:
## QE(df = 355) = 6522.6407, p-val < .0001
##
## Test of Moderators (coefficients 2:4):
## QM(df = 3) = 2.8138, p-val = 0.4212
##
## Model Results:
##
##      estimate      se      zval      pval      ci.lb      ci.ub
## intrcpt      0.4893  0.3499   1.3983  0.1620  -0.1965   1.1751
## exp_age      0.0793  0.0730   1.0857  0.2776  -0.0638   0.2224
## size      -0.2398  0.2014  -1.1908  0.2337  -0.6345   0.1549
## org_level   -0.2252  0.2773  -0.8122  0.4167  -0.7688   0.3183
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Non-linear averaging model with all modifiers

```
## Warning: Rows with NAs omitted from model fitting.

##
## Multivariate Meta-Analysis Model (k = 359; method: REML)
##
## Variance Components:
##
##      estim      sqrt  nlvls  fixed      factor
## sigma^2.1  0.0464  0.2154    27    no      study_id
## sigma^2.2  0.3284  0.5731    44    no      study_id/experiment_id
## sigma^2.3  0.4578  0.6766    98    no      study_id/experiment_id/response_id
##
## Test for Residual Heterogeneity:
## QE(df = 352) = 6377.2072, p-val < .0001
##
## Test of Moderators (coefficients 2:7):
## QM(df = 6) = 97.4285, p-val < .0001
##
## Model Results:
##
```

```
##               estimate      se      zval      pval      ci.lb
## intrcpt          1.3702  0.4645   2.9495  0.0032   0.4597
## flux_range       -0.0209  0.0223  -0.9399  0.3473  -0.0646
## mean_temp_constant -0.0466  0.0129  -3.6196  0.0003  -0.0719
## exp_age           0.0706  0.0737   0.9580  0.3381  -0.0738
## size             -0.1461  0.2205  -0.6626  0.5076  -0.5783
## org_level        -0.2797  0.2824  -0.9905  0.3220  -0.8331
## flux_range:mean_temp_constant  0.0012  0.0010   1.2001  0.2301  -0.0008
##               ci.ub
## intrcpt          2.2806  **
## flux_range         0.0227
## mean_temp_constant -0.0214  ***
## exp_age            0.2150
## size               0.2861
## org_level          0.2738
## flux_range:mean_temp_constant  0.0031
##
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

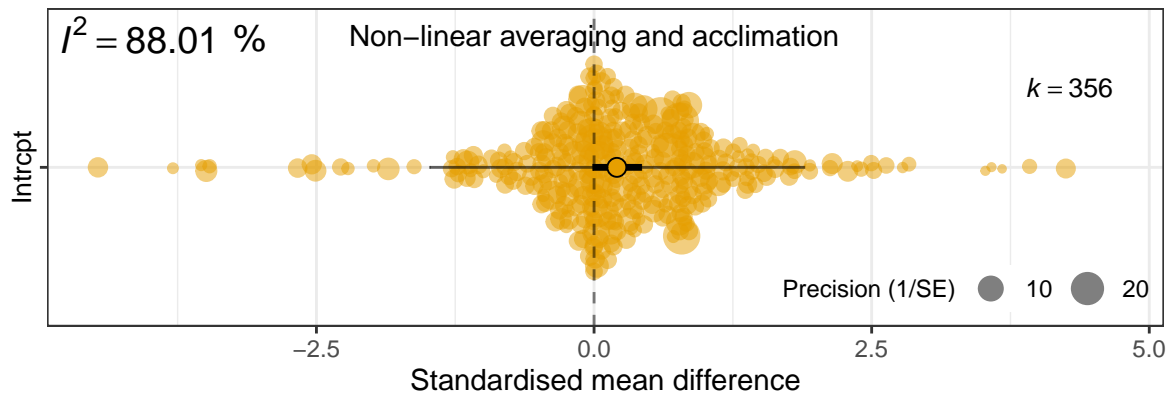
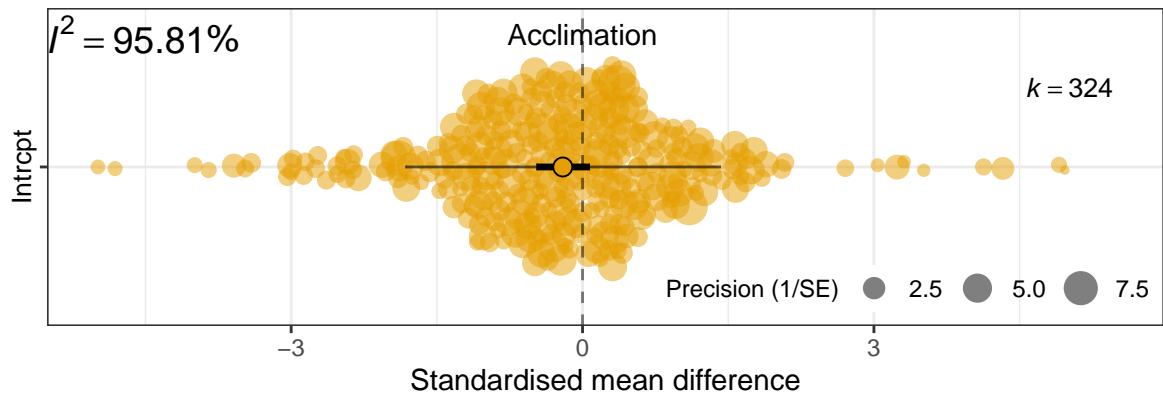
#Model selection

```
AIC(temp_acclimation_model, trait_acclimation_model, acclimation_model, temp_full_var_model, trait_full_var_model)
```

```
## Warning: Models not all fitted to the same data.
```

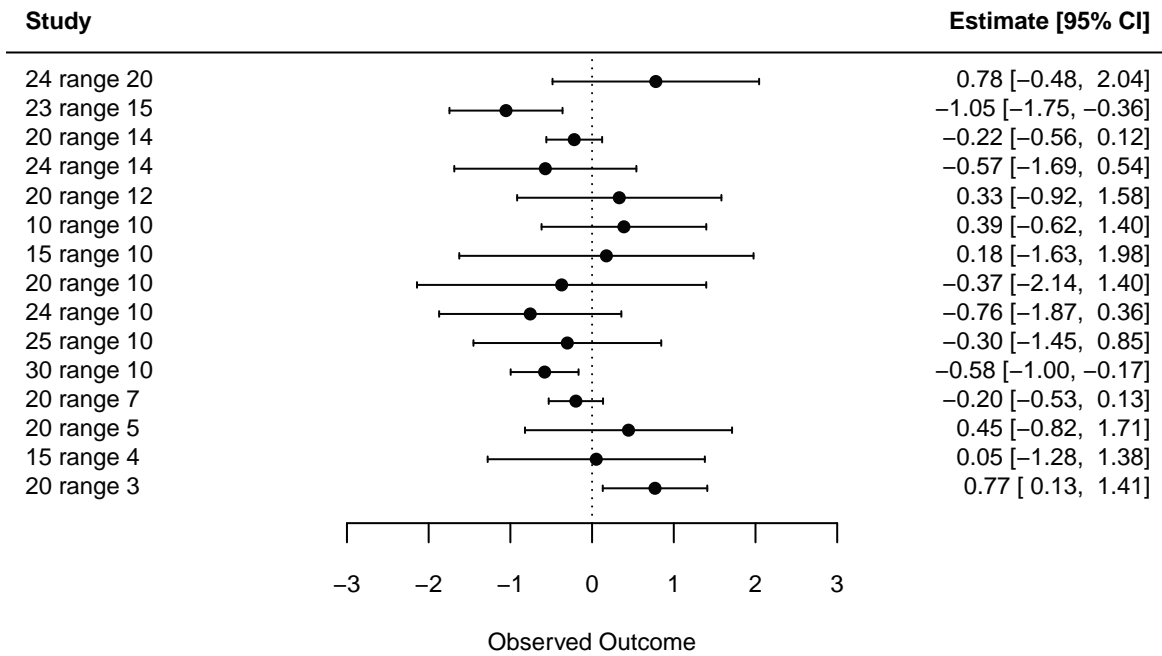
```
##           df      AIC
## temp_acclimation_model    7 2513.012
## trait_acclimation_model    7 2622.262
## acclimation_model         10 2504.172
## temp_full_var_model        7 4235.311
## trait_full_var_model        7 4187.426
## full_var_model            10 4100.259
```

Figures

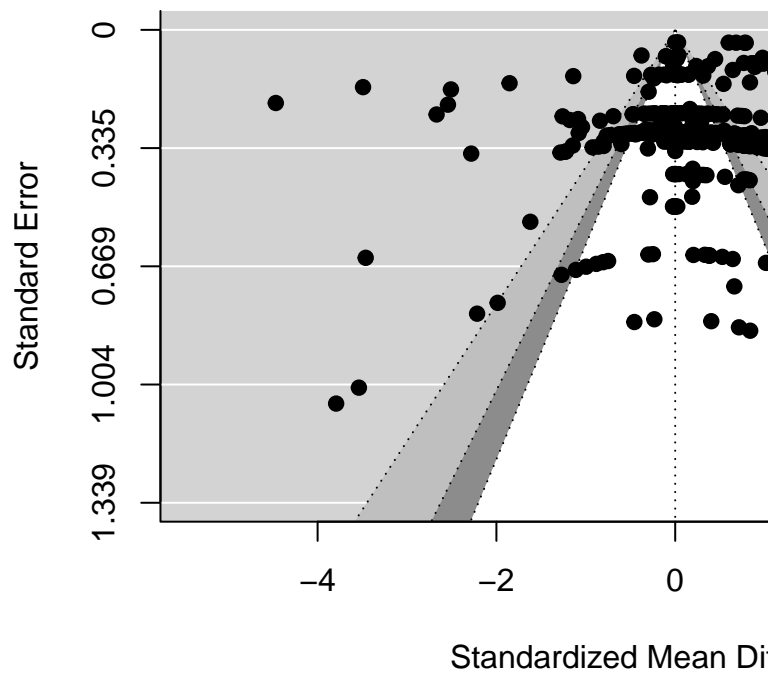
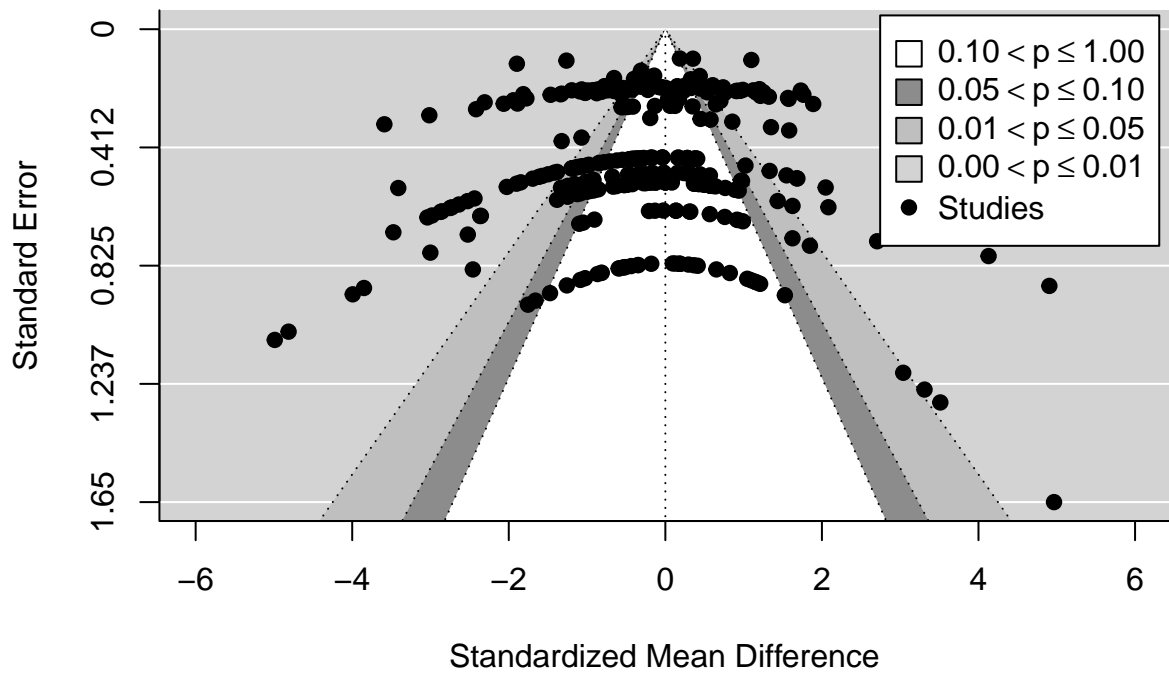


'summarise()' has grouped output by 'mean_temp_reared'. You can override using the '.groups' argument

'summarise()' has grouped output by 'mean_temp_constant'. You can override using the '.groups' argument



Funnel plot of acclimation data

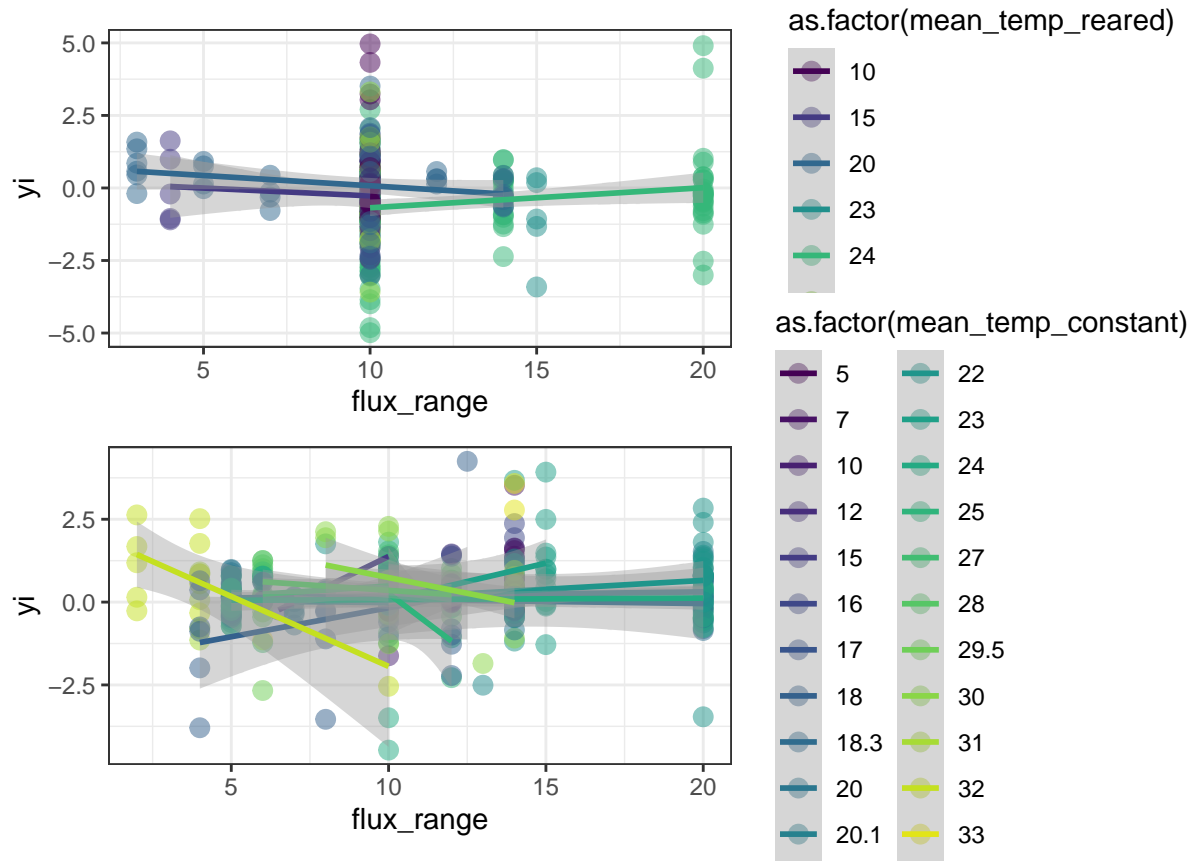


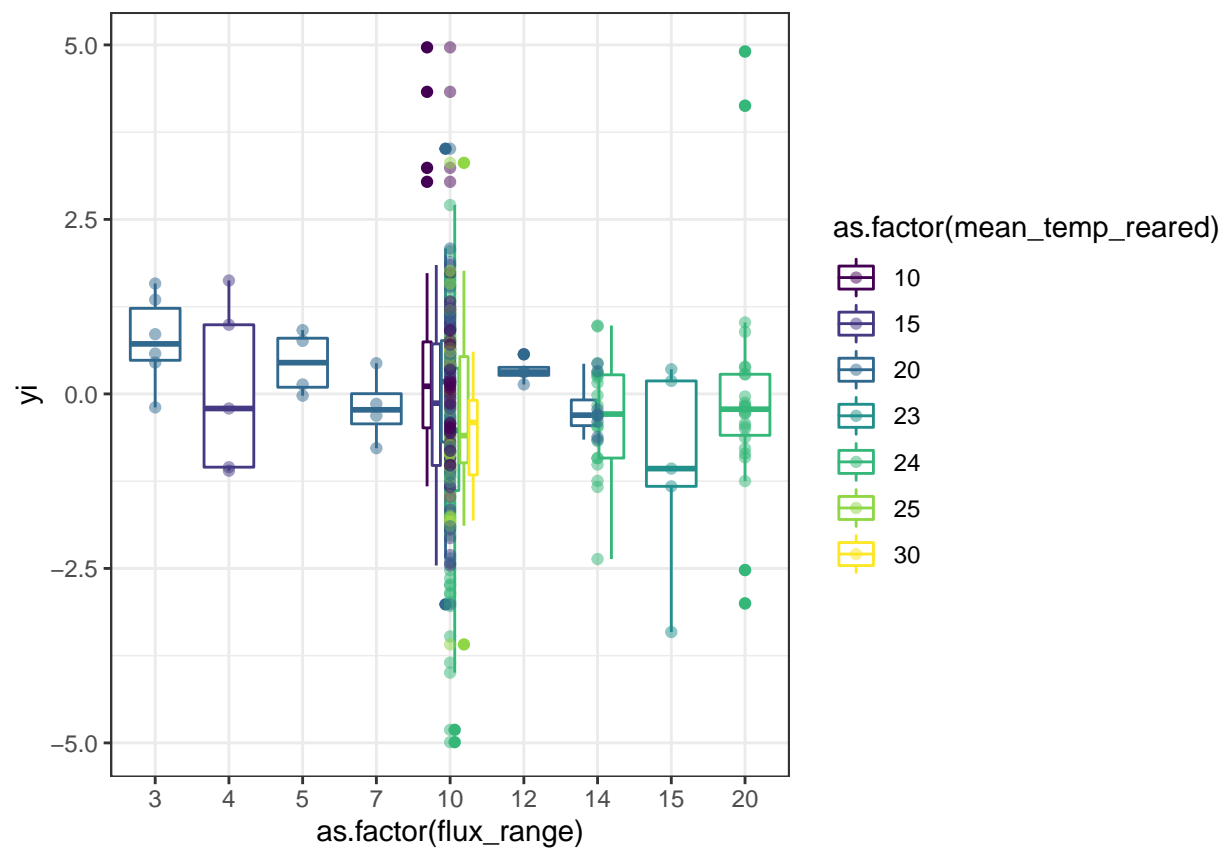
Funnel plot of non-linear averaging + acclimation data

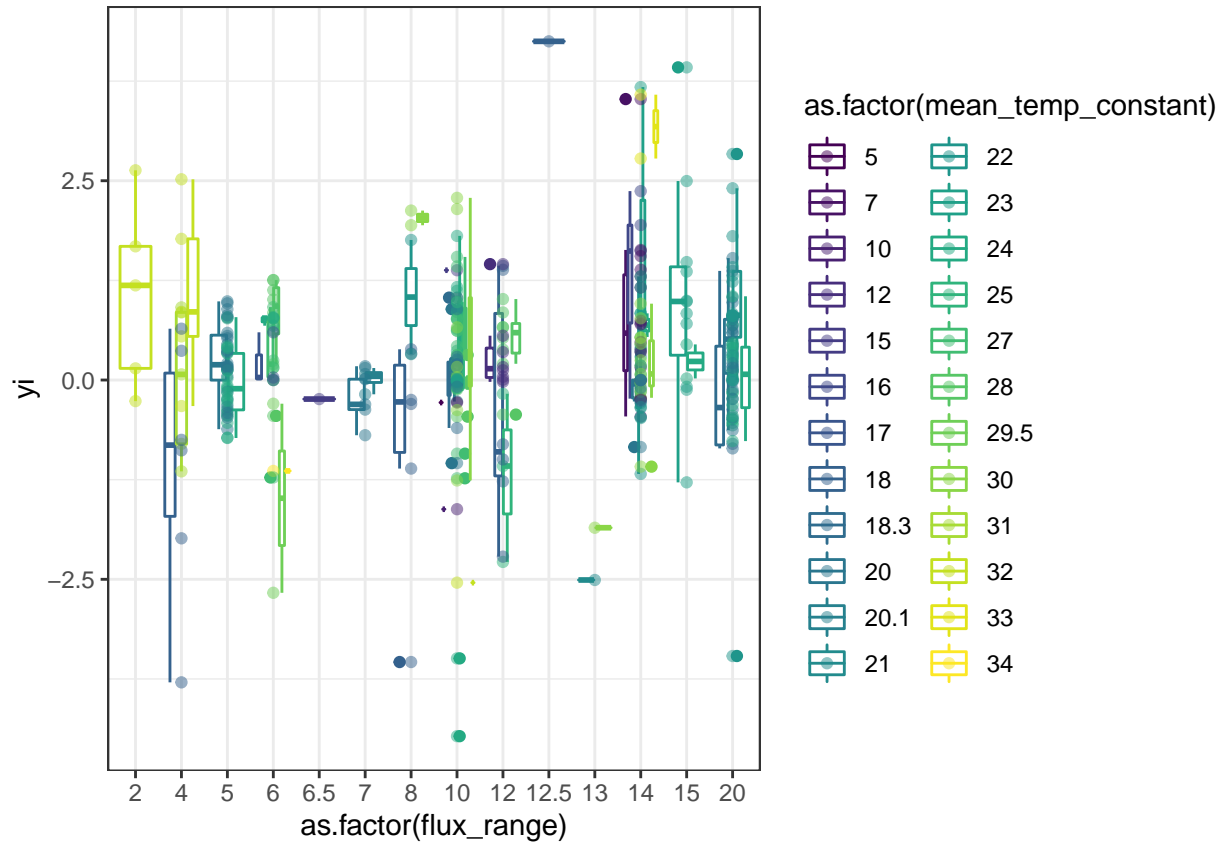
```
## 'geom_smooth()' using formula 'y ~ x'
## 'geom_smooth()' using formula 'y ~ x'
```

```
## Warning in qt((1 - level)/2, df): NaNs produced
```

```
## Warning in max(ids, na.rm = TRUE): no non-missing arguments to max; returning
## -Inf
```







Supplementary code and models

Acclimation data split out by CT data and other thermal performance metrics

CT aggregated model

```
## Warning: Redundant predictors dropped from the model.
```

Remaining acclimation metrics model

```
## Warning: Rows with NAs omitted from model fitting.
```

```
## Warning: Redundant predictors dropped from the model.
```