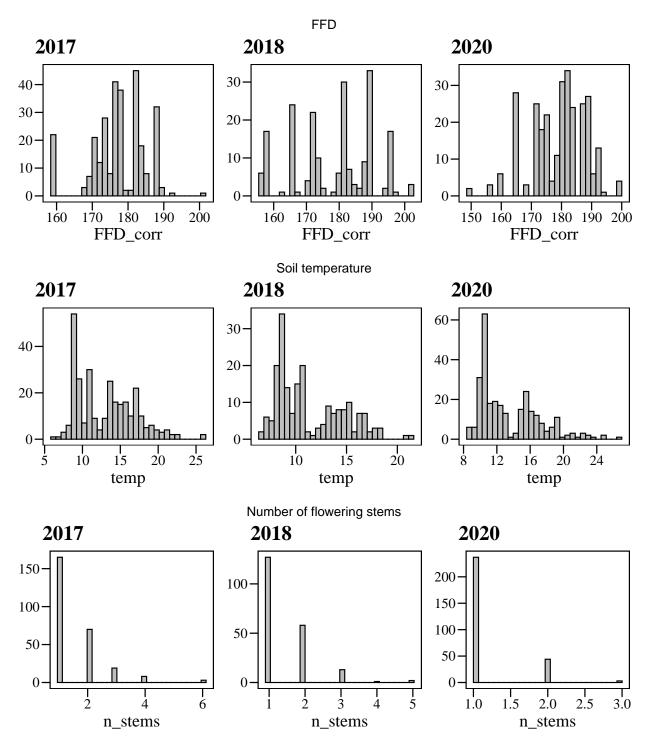
Response of Pinguicula vulgaris to geothermal heating Analyses

Alicia Valdés

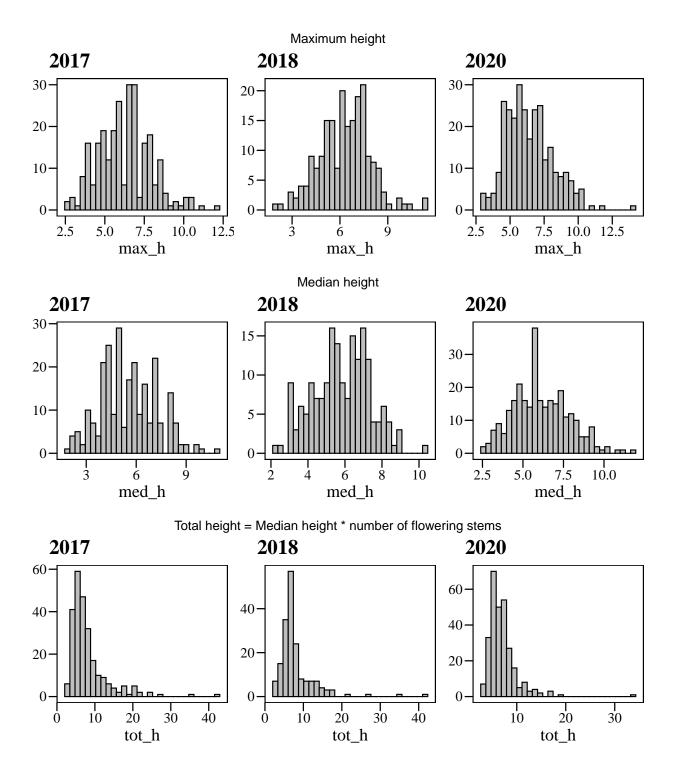
Contents

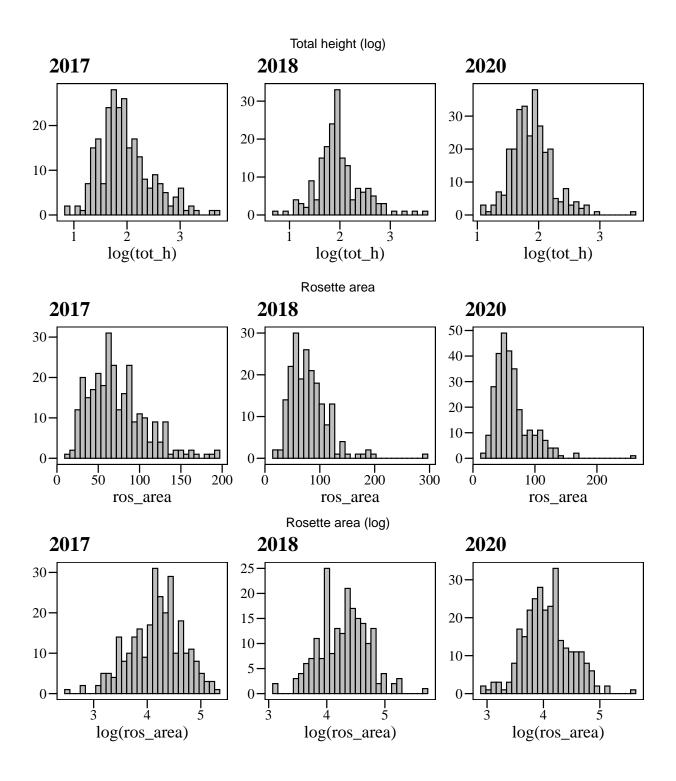
Va	ariable distributions	2
Relationships bewteen fitness and condition variables		
	Maximum height	5
	Median height	6
	Rosette area (log)	8
	Total height (log)	9
1.	Models for FFD	11
	1.1. Effect of temperature on FFD	11
2.	Models for fitness	15
	2.1 Effect of temperature on fitness	15
	Fitness as number of seeds	15
	Fitness as $0/1$	17
	2.2. Effect of treatment on fitness (2020)	19
	Fitness as number of seeds	19
	Fitness as $0/1$	20
	2.3. Effect of temperature AND treatment on fitness (2020)	21
	Fitness as number of seeds	21
	Fitness as $0/1$	23
3.	Selection models	27
	3.1. Effect of temperature on selection on FFD $\ \ldots \ \ldots \ \ldots \ \ldots \ \ldots \ \ldots$	27
	3.2. Effect of treatment on selection on FFD (2020) $\ \ldots \ \ldots \ \ldots \ \ldots \ \ldots \ \ldots$	30
	3.3. Effects of temperature AND treatment on selection on FFD (2020)	31
	3.4. Effects of all 2-way interactions between temperature, treatment and FFD (2020)	33
	3.5. Effects of all 2-way interactions between temperature, treatment and FFD $+$ 3-way interaction (2020)	35

Variable distributions



Most of them have 1 flowering stem, and the variation is not so large. Probably use another condition variable.





Relationships bewteen fitness and condition variables

Chose between maximum height, median height, rosette area (log) and total height (log) as condition variables.

Used first Poisson models, but they showed significant overdispersion and zero-inflation. Best fitting models were zero-inflated negative binomial GLMs, so I used those

Maximum height

(Intercept) 4.50062

Zero-inflation model:

0.11903

max h

```
cond1_17<-glmmTMB(n_seeds~max_h,ziformula=~1,family="nbinom2",ping_17_data)
cond1_18<-glmmTMB(n_seeds~max_h,ziformula=~1,family="nbinom2",ping_18_data)</pre>
cond1_20<-glmmTMB(n_seeds~max_h,ziformula=~1,family="nbinom2",ping_20_data)
summary(cond1_17)
## Family: nbinom2 ( log )
## Formula:
                     n_seeds ~ max_h
## Zero inflation:
                             ~1
## Data: ping_17_data
##
##
        AIC
                 BIC logLik deviance df.resid
##
     2081.7
              2095.9 -1036.8
                                2073.7
                                            255
##
##
## Overdispersion parameter for nbinom2 family (): 2.7
##
## Conditional model:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 5.15492
                           0.22372 23.041
                                             <2e-16 ***
               0.05572
                           0.03405
                                              0.102
## max_h
                                     1.636
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.1005
                          0.1244 -0.807
summary(cond1_18)
## Family: nbinom2 ( log )
## Formula:
                     n_seeds ~ max_h
## Zero inflation:
                            ~1
## Data: ping_18_data
##
##
        AIC
                BIC
                       logLik deviance df.resid
##
     1202.4
             1215.5
                       -597.2
                               1194.4
                                            194
##
##
## Overdispersion parameter for nbinom2 family (): 2.91
##
## Conditional model:
##
              Estimate Std. Error z value Pr(>|z|)
```

2.906 0.00366 **

0.27313 16.478 < 2e-16 ***

0.04096

Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1

```
Estimate Std. Error z value Pr(>|z|)
                0.4520
                           0.1458
                                     3.1 0.00193 **
## (Intercept)
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
summary(cond1_20)
## Family: nbinom2 (log)
                    n_seeds ~ max_h
## Formula:
## Zero inflation:
                        ~1
## Data: ping_20_data
##
       AIC
                BIC logLik deviance df.resid
##
    1923.2
            1937.7 -957.6
                              1915.2
##
## Overdispersion parameter for nbinom2 family (): 6.29
## Conditional model:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 5.02433
                         0.14697 34.19 < 2e-16 ***
## max_h
               0.06723
                          0.02252
                                   2.99 0.00283 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
              Estimate Std. Error z value Pr(>|z|)
                0.1023
                          0.1210 0.845
## (Intercept)
                                            0.398
Median height
cond2_17<-glmmTMB(n_seeds~med_h,ziformula=~1,family="nbinom2",ping_17_data)
cond2_18<-glmmTMB(n_seeds~med_h,ziformula=~1,family="nbinom2",ping_18_data)</pre>
cond2_20<-glmmTMB(n_seeds~med_h,ziformula=~1,family="nbinom2",ping_20_data)
summary(cond2_17)
## Family: nbinom2 (log)
## Formula:
                    n_seeds ~ med_h
## Zero inflation:
                           ~1
## Data: ping_17_data
##
##
       AIC
                BIC
                      logLik deviance df.resid
##
    1930.1
                      -961.1
                              1922.1
                                           241
             1944.1
##
##
## Overdispersion parameter for nbinom2 family (): 2.74
## Conditional model:
              Estimate Std. Error z value Pr(>|z|)
##
```

```
## (Intercept) 5.64057
                         0.17938
                                  31.44
                                          <2e-16 ***
## med h
           -0.02697
                         0.03066 -0.88
                                           0.379
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Zero-inflation model:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.05717 0.12783 -0.447
summary(cond2_18)
## Family: nbinom2 ( log )
## Formula:
                   n_seeds ~ med_h
## Zero inflation:
## Data: ping_18_data
##
       AIC
             BIC logLik deviance df.resid
##
    1085.5 1098.3 -538.8 1077.5
##
## Overdispersion parameter for nbinom2 family (): 2.63
## Conditional model:
             Estimate Std. Error z value Pr(>|z|)
                       0.28597 19.046
## (Intercept) 5.44661
                                          <2e-16 ***
## med_h
          -0.02995
                         0.04849 -0.618
                                           0.537
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Zero-inflation model:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.4664
                         0.1536
                                 3.037 0.00239 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
summary(cond2_20)
## Family: nbinom2 ( log )
## Formula:
                   n_seeds ~ med_h
## Zero inflation:
                       ~1
## Data: ping_20_data
##
       AIC
##
               BIC logLik deviance df.resid
##
   1930.7 1945.2 -961.4 1922.7
                                         270
##
## Overdispersion parameter for nbinom2 family (): 5.94
## Conditional model:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 5.60256 0.13281 42.18
                                          <2e-16 ***
## med h
             -0.02478
                         0.02141
                                  -1.16
                                           0.247
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
## Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.1023 0.1210 0.845 0.398
```

Rosette area (log)

summary(cond3_17)

```
## Family: nbinom2 (log)
## Formula:
                  n_seeds ~ log(ros_area)
## Zero inflation:
                       ~1
## Data: ping_17_data
##
##
      AIC
              BIC logLik deviance df.resid
##
    2076.4 2090.7 -1034.2 2068.4
##
## Overdispersion parameter for nbinom2 family ():
## Conditional model:
              Estimate Std. Error z value Pr(>|z|)
                3.4967
                         0.4501 7.768 7.97e-15 ***
## (Intercept)
                          0.1072 4.458 8.28e-06 ***
## log(ros_area)
                0.4780
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
            Estimate Std. Error z value Pr(>|z|)
```

summary(cond3_18)

```
## Overdispersion parameter for nbinom2 family (): 3.16
##
## Conditional model:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                 2.7849
                           0.6127 4.545 5.49e-06 ***
## log(ros_area) 0.5757
                            0.1420 4.055 5.01e-05 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.4520
                         0.1458
                                    3.1 0.00193 **
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
summary(cond3_20)
## Family: nbinom2 ( log )
## Formula:
           n_seeds ~ log(ros_area)
## Zero inflation:
                       ~1
## Data: ping_20_data
##
              BIC logLik deviance df.resid
       AIC
##
    1914.7 1929.1 -953.3 1906.7
                                         270
##
##
## Overdispersion parameter for nbinom2 family (): 6.71
## Conditional model:
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) 3.86037 0.37036 10.423 < 2e-16 ***
## log(ros_area) 0.39389
                         0.09164 4.298 1.72e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Zero-inflation model:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.1023 0.1210 0.845 0.398
Total height (log)
cond4_17<-glmmTMB(n_seeds~log(tot_h),ziformula=~1,family="nbinom2",</pre>
                 ping_17_data)
cond4_18<-glmmTMB(n_seeds~log(tot_h),ziformula=~1,family="nbinom2",</pre>
                 ping 18 data)
cond4_20<-glmmTMB(n_seeds~log(tot_h),ziformula=~1,family="nbinom2",
                 ping_20_data)
summary(cond4_17)
```

Family: nbinom2 (log)

```
n_seeds ~ log(tot_h)
## Zero inflation:
                     ~1
## Data: ping_17_data
##
##
       AIC
              BIC logLik deviance df.resid
##
    1917.9 1931.9 -955.0 1909.9
##
##
## Overdispersion parameter for nbinom2 family ():
##
## Conditional model:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 4.6430
                       0.2337 19.863 < 2e-16 ***
## log(tot_h)
               0.4100
                         0.1125 3.643 0.000269 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Zero-inflation model:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.05716   0.12783   -0.447   0.655
summary(cond4_18)
## Family: nbinom2 ( log )
## Formula:
                   n_seeds ~ log(tot_h)
## Zero inflation:
## Data: ping_18_data
##
              BIC logLik deviance df.resid
##
       AIC
##
    1066.2 1079.0 -529.1 1058.2
                                         175
##
##
## Overdispersion parameter for nbinom2 family (): 3.42
##
## Conditional model:
             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 3.8697 0.3106 12.458 < 2e-16 ***
## log(tot_h)
                         0.1476 4.481 7.44e-06 ***
               0.6615
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
             Estimate Std. Error z value Pr(>|z|)
                        0.1536 3.037 0.00239 **
## (Intercept) 0.4664
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
summary(cond4_20)
## Family: nbinom2 ( log )
                   n_seeds ~ log(tot_h)
## Formula:
## Zero inflation:
                       ~1
## Data: ping_20_data
```

```
##
##
        AIC
                 BIC
                       logLik deviance df.resid
              1907.8
##
     1893.4
                       -942.7
                                1885.4
##
##
## Overdispersion parameter for nbinom2 family (): 7.9
##
## Conditional model:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
               4.19879
                           0.18918 22.194 < 2e-16 ***
## log(tot_h)
                0.65456
                           0.09879
                                     6.626 3.45e-11 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
##
               Estimate Std. Error z value Pr(>|z|)
                 0.1023
                            0.1210
                                     0.845
                                              0.398
## (Intercept)
```

Rosette area and total height have significant effects on fitness on the 3 years. Total height has larger effect sizes than rosette area in 2 out of 3 years (checked with models with scaled predictors, not shown). I decided to use total height as condition variable.

1. Models for FFD

1.1. Effect of temperature on FFD

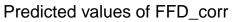
Using linear models, including quadratic effects of temp (included first but removed in 2017 because they were not significant).

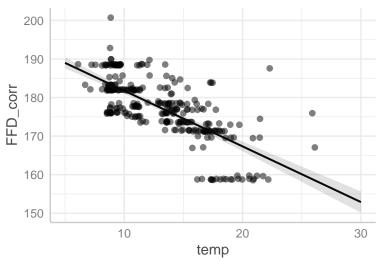
```
FFD_2017_1<-lm(FFD_corr~temp,ping_17_data)
summary(FFD_2017_1)</pre>
```

```
##
## lm(formula = FFD corr ~ temp, data = ping 17 data)
##
## Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
                                            Max
  -14.2025 -2.6261 -0.2459
                               2.9045
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 196.20860
                           1.10402 177.72
                                              <2e-16 ***
                                    -17.81
               -1.44386
                           0.08105
                                              <2e-16 ***
## temp
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 5.342 on 290 degrees of freedom
## Multiple R-squared: 0.5225, Adjusted R-squared: 0.5209
## F-statistic: 317.4 on 1 and 290 DF, p-value: < 2.2e-16
```

```
FFD_2018_1<-lm(FFD_corr~temp+I(temp^2),ping_18_data)
summary(FFD_2018_1)
##
## Call:
## lm(formula = FFD_corr ~ temp + I(temp^2), data = ping_18_data)
## Residuals:
##
       Min
                 1Q
                      Median
                                   30
                      0.0669
## -12.4932 -4.3128
                               3.0298 16.1994
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 235.5409
                           5.7042 41.293 < 2e-16 ***
## temp
               -6.7120
                           0.9608 -6.986 4.21e-11 ***
## I(temp^2)
                0.1369
                           0.0380
                                    3.604 0.000397 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 5.604 on 198 degrees of freedom
## Multiple R-squared: 0.7895, Adjusted R-squared: 0.7874
## F-statistic: 371.3 on 2 and 198 DF, p-value: < 2.2e-16
FFD_2020_1<-lm(FFD_corr~temp+I(temp^2),ping_20_data)
summary(FFD_2020_1)
##
## Call:
## lm(formula = FFD_corr ~ temp + I(temp^2), data = ping_20_data)
##
## Residuals:
       Min
                 1Q
                     Median
                                   3Q
                                           Max
## -12.3407 -2.5589
                     0.2415
                               2.8193 14.0155
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 216.81764
                           3.78519 57.281 < 2e-16 ***
## temp
               -3.37575
                           0.51862 -6.509 3.41e-10 ***
                0.03967
                           0.01681
                                     2.360 0.0189 *
## I(temp^2)
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 4.435 on 284 degrees of freedom
## Multiple R-squared: 0.7598, Adjusted R-squared: 0.7581
## F-statistic: 449.2 on 2 and 284 DF, p-value: < 2.2e-16
Model diagnostics (not shown) indicated that linear models are OK.
2017:
plot(ggpredict(FFD_2017_1),add.data=T)
```

```
## $temp
```



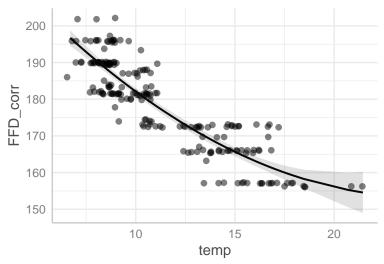


2018:

plot(ggpredict(FFD_2018_1),add.data=T)

\$temp



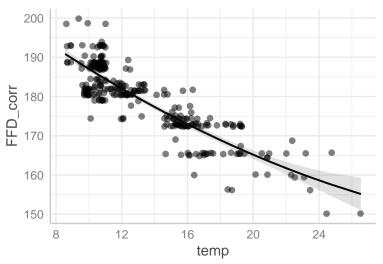


2020:

plot(ggpredict(FFD_2020_1),add.data=T)

\$temp

Predicted values of FFD_corr



The effects look almost linear in 2018 and 2020.

Predictions of FFD_corr for minimum and maximum temperatures:

```
ggpredict(FFD_2017_1,terms="temp[minmax]")
```

```
## # Predicted values of FFD_corr
## # x = temp
##

## x | Predicted | 95% CI
## ------
## 6.20 | 187.26 | [186.01, 188.51]
## 26.00 | 158.67 | [156.52, 160.81]
```

```
# 187.26-158.67=28.59 days earlier on warmer soils ggpredict(FFD_2018_1,terms="temp[minmax]")
```

```
## # Predicted values of FFD_corr
## # x = temp
##

## x | Predicted | 95% CI
## ------
## 6.70 | 196.72 | [194.59, 198.85]
## 21.45 | 154.57 | [148.95, 160.18]
```

```
# 196.72-154.57=42.15 days earlier on warmer soils ggpredict(FFD_2020_1,terms="temp[minmax]")
```

```
## # Predicted values of FFD_corr
## # x = temp
##

## x | Predicted | 95% CI
## ------
## 8.55 | 190.85 | [189.57, 192.14]
## 26.55 | 155.16 | [151.03, 159.28]
```

2. Models for fitness

2.1 Effect of temperature on fitness

Fitness as number of seeds

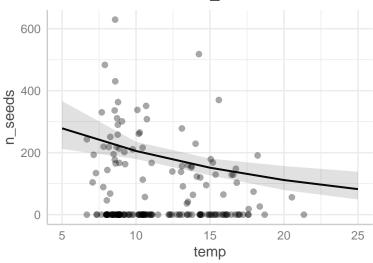
Including total height (log) as condition variable. I first included quadratic effects of temperature, but then removed them because they were never significant. I used only plants from control treatment in 2020. Using negative binomial GLMs with zero inflation (best fitting models).

```
summary(fitness_2017_1)
```

```
## Family: nbinom2 (log)
                    n_seeds ~ temp + log(tot_h)
## Formula:
## Zero inflation:
                            ~1
## Data: ping_17_data
##
##
       AIC
                BIC
                      logLik deviance df.resid
##
    1918.9
             1936.4
                      -954.5
                               1908.9
##
##
## Overdispersion parameter for nbinom2 family (): 3.02
##
## Conditional model:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 4.41331
                          0.32506 13.577 < 2e-16 ***
## temp
               0.01469
                          0.01468
                                   1.001 0.31706
## log(tot_h)
              0.43193
                          0.11440
                                   3.776 0.00016 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.05716
                          0.12783 -0.447
                                             0.655
summary(fitness_2018_1)
```

```
##
##
       ATC
                BIC logLik deviance df.resid
             1074.8 -524.4
##
    1058.8
                              1048.8
##
## Overdispersion parameter for nbinom2 family (): 3.88
## Conditional model:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 4.74135
                          0.40672 11.658 < 2e-16 ***
## temp
              -0.06094
                          0.01901 -3.205 0.00135 **
## log(tot_h)
              0.57202
                          0.14216
                                  4.024 5.73e-05 ***
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                0.4664
                          0.1536 3.037 0.00239 **
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
summary(fitness_2020_1)
## Family: nbinom2 (log)
                    n_seeds ~ temp + log(tot_h)
## Formula:
## Zero inflation:
                           ~1
## Data: subset(ping_20_data, treatment == "C")
##
##
       AIC
                BIC
                      logLik deviance df.resid
##
     551.0
              562.5
                     -270.5
                                541.0
                                            68
##
##
## Overdispersion parameter for nbinom2 family (): 6.76
##
## Conditional model:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 4.03855
                          0.57778
                                   6.990 2.75e-12 ***
              -0.00284
                          0.02233 -0.127 0.89879
## temp
## log(tot_h)
              0.72693
                          0.24914
                                    2.918 0.00353 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.08224
                          0.23428 -0.351
                                             0.726
Effect of temperature only significant in 2018.
2018:
plot(ggeffect(fitness_2018_1,type="zero_inflated",terms="temp"),
    add.data=T) # * negative effect of temp
```

Predicted counts of n_seeds



Fitness as 0/1

Proportion of plants in each year that produced 0 seeds:

```
nrow(subset(ping_17_data,n_seeds==0))/nrow(ping_17_data)
```

[1] 0.4623288

```
nrow(subset(ping_18_data,n_seeds==0))/nrow(ping_18_data)
```

[1] 0.6069652

```
nrow(subset(ping_20_data,n_seeds==0))/nrow(ping_20_data)
```

[1] 0.5052265

Around half of the plants in each year produced no seeds.

Therefore, I tried also fitting logistic regressions (binomial GLMs) with fitness as a 0/1 variable (0 = no seeds produced, 1 = at least 1 seed produced).

Create 0/1 variable for fitness:

Including total height (log) as condition variable. I first included quadratic effects of temperature, but then removed them because they were never significant. I used only plants from control treatment in 2020.

```
fitness_01_2017_1<-glm(fitness_01~temp+log(tot_h),ping_17_data,
                       family="binomial")
fitness_01_2018_1<-glm(fitness_01~temp+log(tot_h),ping_18_data,</pre>
                       family="binomial")
fitness_01_2020_1<-glm(fitness_01~temp+log(tot_h),
                   subset(ping_20_data,treatment=="C"),family="binomial")
summary(fitness 01 2017 1)
##
## Call:
## glm(formula = fitness_01 ~ temp + log(tot_h), family = "binomial",
      data = ping_17_data)
##
## Deviance Residuals:
##
                1Q Median
      \mathtt{Min}
                                   3Q
                                           Max
## -1.5217 -1.1834 0.9672
                               1.1606
                                        1.3193
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.714507
                           0.778864 -0.917
                                               0.359
              -0.007203
                           0.035244 -0.204
                                               0.838
## log(tot_h)
              0.437526
                           0.276854
                                    1.580
                                               0.114
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 339.44 on 244 degrees of freedom
## Residual deviance: 336.61 on 242 degrees of freedom
     (47 observations deleted due to missingness)
## AIC: 342.61
##
## Number of Fisher Scoring iterations: 4
summary(fitness_01_2018_1)
##
## Call:
## glm(formula = fitness_01 ~ temp + log(tot_h), family = "binomial",
      data = ping_18_data)
##
## Deviance Residuals:
##
      Min
                 1Q
                    Median
                                   3Q
                                           Max
## -1.2840 -0.9867 -0.8985
                             1.3376
                                        1.5835
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -2.01683
                           0.97709 - 2.064
                                             0.0390 *
## temp
               0.01566
                           0.04732
                                     0.331
                                             0.7407
## log(tot_h)
               0.68903
                           0.35970
                                     1.916
                                             0.0554 .
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

##

```
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 238.67 on 178 degrees of freedom
## Residual deviance: 234.86 on 176 degrees of freedom
     (22 observations deleted due to missingness)
## AIC: 240.86
## Number of Fisher Scoring iterations: 4
summary(fitness_01_2020_1)
##
## Call:
## glm(formula = fitness_01 ~ temp + log(tot_h), family = "binomial",
       data = subset(ping_20_data, treatment == "C"))
##
##
## Deviance Residuals:
     Min
              1Q Median
                               3Q
                                      Max
## -1.448 -1.199
                  0.997
                            1.151
                                    1.289
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 1.48968
                           1.92751
                                     0.773
                                              0.440
## temp
               -0.06519
                           0.06593 -0.989
                                              0.323
## log(tot_h) -0.27212
                           0.76150 - 0.357
                                              0.721
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 101.08 on 72 degrees of freedom
## Residual deviance: 100.07 on 70 degrees of freedom
     (6 observations deleted due to missingness)
## AIC: 106.07
##
## Number of Fisher Scoring iterations: 4
```

No significant effects.

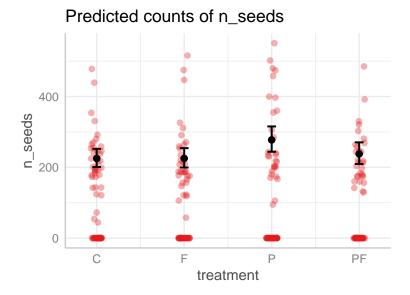
2.2. Effect of treatment on fitness (2020)

Fitness as number of seeds

Using a negative binomial GLM with zero inflation. Including total height (log) as condition variable.

```
##
##
        AIC
                 BIC
                       logLik deviance df.resid
                       -939.0
##
     1892.1
              1917.4
                                1878.1
##
##
## Overdispersion parameter for nbinom2 family (): 8.36
##
## Conditional model:
##
               Estimate Std. Error z value Pr(>|z|)
                                     21.962 < 2e-16 ***
## (Intercept) 4.168918
                          0.189827
## treatmentF
               0.000115
                          0.083925
                                      0.001
                                              0.9989
                                      2.406
                                              0.0161 *
## treatmentP
               0.209158
                          0.086936
## treatmentPF 0.055546
                          0.086118
                                      0.645
                                              0.5189
## log(tot_h)
               0.636962
                          0.096591
                                      6.594 4.27e-11 ***
##
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Zero-inflation model:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                 0.1023
                            0.1210
                                      0.845
```

```
plot(ggeffect(fitness_2020_treat,term="treatment"),add.data=T)
```



Fitness significantly larger in pollination treatment than in control.

Fitness as 0/1

Logistic regression (binomial GLM) with fitness as a 0/1 variable (0 = no seeds produced, 1 = at least 1 seed produced).

```
##
## Call:
  glm(formula = fitness_01 ~ treatment + log(tot_h), family = "binomial",
       data = ping_20_data)
## Deviance Residuals:
              10 Median
                               30
                                      Max
## -1.272 -1.123 -1.041
                            1.196
                                    1.348
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
                0.4323
                            0.7252
                                     0.596
                                              0.551
## (Intercept)
## treatmentF
                -0.1080
                            0.3388 -0.319
                                              0.750
## treatmentP
                -0.2802
                            0.3442 - 0.814
                                              0.416
## treatmentPF
               -0.3448
                            0.3376
                                    -1.021
                                               0.307
## log(tot_h)
                -0.1873
                            0.3672 -0.510
                                               0.610
##
  (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 379.13 on 273 degrees of freedom
## Residual deviance: 377.51 on 269 degrees of freedom
     (13 observations deleted due to missingness)
## AIC: 387.51
## Number of Fisher Scoring iterations: 3
```

No significant effects.

2.3. Effect of temperature AND treatment on fitness (2020)

Fitness as number of seeds

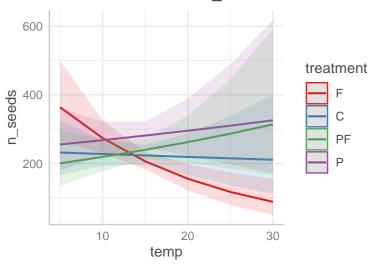
Conditional model:

Using a negative binomial GLM with zero inflation. Including total height (log) as condition variable. Including temperature and treatment as predictors both in the "count" part of the model and in the "zero-inflation" part of the model.

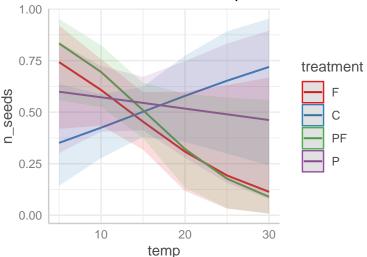
```
fitness_2020_temp_treat_2<-glmmTMB(n_seeds~temp*treatment+log(tot_h),
                                   ziformula=~.,ping_20_data,family="nbinom2")
summary(fitness_2020_temp_treat_2)
## Family: nbinom2 ( log )
## Formula:
                     n_seeds ~ temp * treatment + log(tot_h)
## Zero inflation:
## Data: ping_20_data
##
##
        AIC
                 BIC
                       logLik deviance df.resid
##
     1895.6
              1964.2
                       -928.8
                                1857.6
                                             255
##
## Overdispersion parameter for nbinom2 family (): 9.11
```

```
##
                    Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                    4.276828
                               0.322948 13.243 < 2e-16 ***
                   -0.003804
                               0.019334
                                        -0.197
                                                  0.8440
## temp
                    0.714743 0.359863
                                          1.986
                                                  0.0470 *
## treatmentF
## treatmentP
                    0.027426
                               0.380013
                                          0.072
                                                  0.9425
## treatmentPF
                               0.398618 -0.640
                                                  0.5222
                   -0.255127
## log(tot h)
                    0.606530
                               0.093641
                                          6.477 9.34e-11 ***
                               0.025858 -2.044
## temp:treatmentF
                   -0.052864
                                                  0.0409 *
## temp:treatmentP
                     0.013564
                               0.027583
                                          0.492
                                                  0.6229
## temp:treatmentPF 0.021741
                               0.028032
                                          0.776
                                                  0.4380
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Zero-inflation model:
##
                   Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                   -1.23940
                               1.23191 -1.006
                                                 0.3144
                    0.06220
                               0.06345
                                         0.980
                                                 0.3269
## temp
## treatmentF
                    2.60611
                               1.41471
                                         1.842
                                                 0.0655 .
## treatmentP
                    1.43741
                               1.31724
                                         1.091
                                                 0.2752
## treatmentPF
                    3.31885
                               1.39915
                                         2.372
                                                 0.0177 *
## log(tot_h)
                    0.16037
                               0.37874
                                         0.423
                                                 0.6720
## temp:treatmentF -0.18668
                               0.10267 -1.818
                                                 0.0690 .
                                                 0.3691
## temp:treatmentP -0.08437
                               0.09393 -0.898
## temp:treatmentPF -0.21935
                               0.09975 -2.199
                                                 0.0279 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
plot(ggpredict(fitness_2020_temp_treat_2,terms=c("temp","treatment"),
               type="fixed"))
```

Predicted counts of n_seeds



Predicted zero-inflation probabilities of n_s



First plot: fitness decreases with temperature only for plants in F treatment. This seems contrary what I would expect - see how can we explain this! Second plot: the probability of zero inflation is represented; values of 1 in the Y-axis indicate a 100% probability of zero inflation (i.e. of producing 0 seeds), so it is like an "inversed" logistic model. The probability of zero inflation decreases with temperature for plants in the PF treatment (and almost in the F treatment, p=0.07). In other words, the probability of producing seeds increases with temperature in those treatments. This agrees with my expectations, as pollination and prey supplementation might partly compensate for the maladaptive nature of plasticity on heated soils, "restoring" interactions that might be lost in these areas because of different sensitivities to warming between plants and interacting animals.

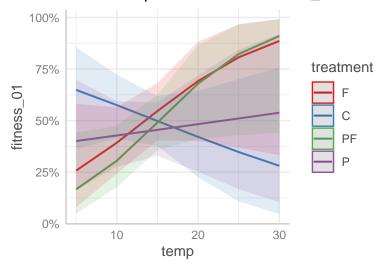
Fitness as 0/1

Logistic regression (binomial GLM) with fitness as a 0/1 variable (0 = no seeds produced, 1 = at least 1 seed produced).

```
##
## Call:
   glm(formula = fitness_01 ~ temp * treatment + log(tot_h), family = "binomial",
       data = ping_20_data)
##
##
## Deviance Residuals:
##
       Min
                  1Q
                       Median
                                     3Q
                                             Max
                     -0.8799
##
   -1.6791
           -1.0851
                                1.1915
                                          1.6359
##
##
  Coefficients:
                     Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                      1.23943
                                 1.23190
                                            1.006
                                                    0.3144
## temp
                     -0.06220
                                 0.06345
                                           -0.980
                                                    0.3269
## treatmentF
                     -2.60614
                                 1.41470
                                           -1.842
                                                    0.0654 .
## treatmentP
                     -1.43744
                                 1.31721
                                           -1.091
                                                    0.2752
```

```
## treatmentPF
                    -3.31888
                                 1.39913
                                         -2.372
                                                   0.0177 *
## log(tot_h)
                    -0.16038
                                 0.37874
                                          -0.423
                                                   0.6720
                     0.18668
## temp:treatmentF
                                 0.10267
                                           1.818
                                                   0.0690 .
                                                   0.3690
## temp:treatmentP
                     0.08438
                                 0.09393
                                           0.898
##
  temp:treatmentPF
                     0.21936
                                 0.09975
                                           2.199
                                                   0.0279 *
##
                     '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
##
##
   (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 379.13
                              on 273
                                       degrees of freedom
## Residual deviance: 369.56 on 265
                                       degrees of freedom
##
     (13 observations deleted due to missingness)
## AIC: 387.56
##
## Number of Fisher Scoring iterations: 4
plot(ggpredict(fitness_01_2020_temp_treat,terms=c("temp","treatment")))
```

Predicted probabilities of fitness_01



The probability of producing seeds increases with temperature for plants in the PF treatment (and almost in the F treatment, p=0.07). This agrees with the results of the previous zero-inflated model and with my expectations.

Relationship among fitness_01 and temp for the different treatments:

```
##
## Call:
## glm(formula = fitness_01 ~ temp + log(tot_h), family = "binomial",
## data = subset(ping_20_data, treatment == "C"))
##
## Deviance Residuals:
## Min 1Q Median 3Q Max
```

```
## -1.448 -1.199
                   0.997 1.151
                                    1.289
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 1.48968
                           1.92751
                                     0.773
              -0.06519
                           0.06593 -0.989
                                              0.323
## temp
## log(tot_h) -0.27212
                           0.76150 -0.357
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 101.08 on 72 degrees of freedom
## Residual deviance: 100.07 on 70 degrees of freedom
     (6 observations deleted due to missingness)
## AIC: 106.07
##
## Number of Fisher Scoring iterations: 4
summary(glm(fitness_01~temp+log(tot_h),subset(ping_20_data,treatment=="P"),
            family="binomial"))
##
## Call:
## glm(formula = fitness_01 ~ temp + log(tot_h), family = "binomial",
##
       data = subset(ping_20_data, treatment == "P"))
##
## Deviance Residuals:
      Min
                 1Q
                     Median
                                   3Q
                                           Max
## -1.2693 -1.1016 -0.9815
                                        1.5178
                               1.2339
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.62914
                           1.65290
                                     0.381
                                              0.703
                0.02390
                           0.06964
                                     0.343
                                              0.731
## temp
                           0.71274 -0.834
                                              0.404
## log(tot_h) -0.59443
##
## (Dispersion parameter for binomial family taken to be 1)
      Null deviance: 89.354 on 64 degrees of freedom
##
## Residual deviance: 88.525 on 62 degrees of freedom
     (3 observations deleted due to missingness)
## AIC: 94.525
##
## Number of Fisher Scoring iterations: 4
summary(glm(fitness_01~temp+log(tot_h),subset(ping_20_data,treatment=="F"),
            family="binomial"))
##
## Call:
## glm(formula = fitness_01 ~ temp + log(tot_h), family = "binomial",
##
       data = subset(ping_20_data, treatment == "F"))
##
## Deviance Residuals:
```

```
Median
                                   3Q
                                           Max
                 1Q
                               1.1547
## -1.4604 -1.0807 -0.9269
                                        1.4481
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -2.50267
                                              0.237
                           2.11630 -1.183
                                     1.610
## temp
                0.13558
                           0.08419
                                              0.107
## log(tot_h)
                0.36324
                           0.82340
                                     0.441
                                              0.659
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 92.867 on 66 degrees of freedom
## Residual deviance: 90.086 on 64 degrees of freedom
     (2 observations deleted due to missingness)
## AIC: 96.086
##
## Number of Fisher Scoring iterations: 4
summary(glm(fitness_01~temp+log(tot_h),subset(ping_20_data,treatment=="PF"),
            family="binomial")) # +*
##
## Call:
## glm(formula = fitness_01 ~ temp + log(tot_h), family = "binomial",
       data = subset(ping_20_data, treatment == "PF"))
##
## Deviance Residuals:
                      Median
##
      Min
                                   3Q
                                           Max
                 1Q
## -1.6580 -0.9408 -0.8586
                               1.1553
                                        1.6238
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -2.47263
                                   -1.289
                           1.91842
                                             0.1974
                           0.07778
                0.15909
                                     2.045
                                             0.0408 *
## temp
                0.03583
                           0.79012
                                     0.045
                                             0.9638
## log(tot h)
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 94.477 on 68 degrees of freedom
##
## Residual deviance: 90.001 on 66 degrees of freedom
     (2 observations deleted due to missingness)
## AIC: 96.001
##
## Number of Fisher Scoring iterations: 4
```

Effect of temperature only significant in the PF treatment.

3. Selection models

3.1. Effect of temperature on selection on FFD

Standardize traits and relativize fitness:

```
# Standardize FFD
ping_17_data$FFD_corr_std<-as.numeric(scale(ping_17_data$FFD_corr))</pre>
ping_18_data$FFD_corr_std<-as.numeric(scale(ping_18_data$FFD_corr))</pre>
ping_20_data$FFD_corr_std<-as.numeric(scale(ping_20_data$FFD_corr))</pre>
# Standardize total height (first log, then scale)
ping 17 data$tot h std<-as.numeric(scale(log(ping 17 data$tot h)))</pre>
ping_18_data$tot_h_std<-as.numeric(scale(log(ping_18_data$tot_h)))</pre>
ping_20_data$tot_h_std<-as.numeric(scale(log(ping_20_data$tot_h)))</pre>
# Relativize fitness
ping_17_data$n_seeds_rel<-with(ping_17_data,n_seeds/mean(n_seeds,na.rm=T))
ping_18_data$n_seeds_rel<-with(ping_18_data,n_seeds/mean(n_seeds,na.rm=T))
ping 20 data$n seeds rel<-with(ping 20 data,n seeds/mean(n seeds,na.rm=T))
# pinq_20_data, treament C
ping_20_data_C<-subset(ping_20_data,treatment=="C")%>%
  mutate(FFD_corr_std=as.numeric(scale(FFD_corr)),
         tot_h_std=as.numeric(scale(log(tot_h))),
         n_seeds_rel=n_seeds/mean(n_seeds,na.rm=T))
```

Using linear models. I first included quadratic effects of temperature, but then removed them because they were never significant. I used only plants from control treatment in 2020.

Model diagnostics (not shown) looked bad, model residuals did not meet the assumption of normality, so using 95% bias-corrected and accelerated (BCa) bootstrap intervals for the model estimates to determine significance. Same for all selection models below.

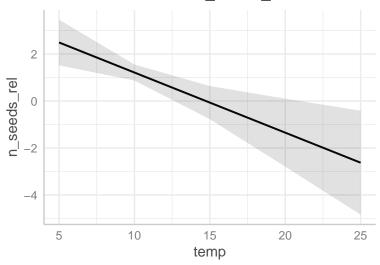
```
summary(selection_2017_1)
```

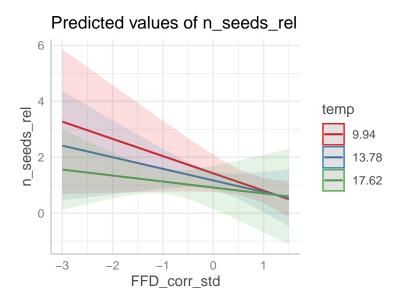
```
##
## Call:
## lm(formula = n_seeds_rel ~ FFD_corr_std * temp + tot_h_std, data = subset(ping_17_data,
## !is.na(n_seeds_rel) & !is.na(tot_h_std)))
##
## Residuals:
## Min 1Q Median 3Q Max
## -2.0571 -0.9139 -0.4028 0.8556 5.0597
##
```

```
## Coefficients:
##
                   Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                   0.948656 0.424566
                                       2.234 0.026377 *
                   ## FFD_corr_std
## temp
                    0.008093
                             0.032155
                                        0.252 0.801494
                    0.294441 0.080229
                                       3.670 0.000299 ***
## tot h std
                              0.021003
                                       1.208 0.228209
## FFD_corr_std:temp 0.025373
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.216 on 240 degrees of freedom
## Multiple R-squared: 0.0601, Adjusted R-squared: 0.04444
## F-statistic: 3.837 on 4 and 240 DF, p-value: 0.004832
BCIs_selection_2017_1
##
                         lower
                                   upper
## FFD_corr_std
                   -1.01037006 0.21112755
## temp
                   -0.05393528 0.06850920
                    0.11478074 0.48333447
## tot h std
## FFD corr std temp -0.01372701 0.06461923
summary(selection_2018_1)
##
## Call:
## lm(formula = n_seeds_rel ~ FFD_corr_std * temp + tot_h_std, data = subset(ping_18_data,
##
      !is.na(n_seeds_rel) & !is.na(tot_h_std)))
##
## Residuals:
      Min
              10 Median
                             3Q
                                    Max
## -2.1999 -1.0248 -0.4156 0.8851 6.0338
## Coefficients:
                   Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                   ## FFD_corr_std
                   0.02831
                              0.48547
                                      0.058 0.95356
## temp
                   -0.26006
                              0.08072 -3.222 0.00152 **
                    0.49726
                              0.11100 4.480 1.35e-05 ***
## tot_h_std
## FFD_corr_std:temp -0.05927
                              0.04170 - 1.421 0.15708
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.463 on 174 degrees of freedom
## Multiple R-squared: 0.1614, Adjusted R-squared: 0.1421
## F-statistic: 8.373 on 4 and 174 DF, p-value: 3.356e-06
BCIs_selection_2018_1
##
                        lower
                                    upper
## FFD_corr_std
                   -0.7529034 0.755218875
                   -0.4111267 -0.115655859
## temp
```

```
## tot h std
                     0.2542839 0.763273652
## FFD_corr_std_temp -0.1214831 0.001855899
summary(selection_2020_C_1)
##
## Call:
## lm(formula = n_seeds_rel ~ FFD_corr_std * temp + tot_h_std, data = subset(ping_20_data_C,
       !is.na(n_seeds_rel) & !is.na(tot_h_std)))
##
## Residuals:
##
       Min
                10 Median
                                  3Q
## -1.43960 -0.94334 -0.04187 0.75455 3.07677
## Coefficients:
##
                   Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                  2.07408 1.17071 1.772 0.0809 .
## FFD corr std
                   -1.13764 0.49832 -2.283 0.0256 *
## temp
                   -0.06586
                               0.08624 -0.764 0.4477
                               0.13841 0.575
## tot_h_std
                    0.07960
                                                0.5671
## FFD_corr_std:temp 0.05242
                               0.02698 1.943 0.0562 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.099 on 68 degrees of freedom
## Multiple R-squared: 0.09254, Adjusted R-squared: 0.03916
## F-statistic: 1.734 on 4 and 68 DF, p-value: 0.1527
BCIs_selection_2020_C_1
##
                          lower
                                     upper
## FFD_corr_std
                   -2.098458916 -0.1775734
## temp
                   -0.220064647 0.1088901
## tot_h_std
                  -0.171761449 0.4159084
## FFD_corr_std_temp  0.008688879  0.1014103
plot(ggpredict(selection_2018_1,terms=c("temp")))
```

Predicted values of n_seeds_rel





2018: higher fitness in colder soils. 2020: early flowering relatively more favoured in colder soils (similar to result for Cerastium but it seems that there is no selection on FFD in warmer soils here - if we look only at plants in the C treatment).

3.2. Effect of treatment on selection on FFD (2020)

Using a linear model.

```
## Call:
## lm(formula = n_seeds_rel ~ FFD_corr_std * treatment + tot_h_std,
      data = subset(ping_20_data, !is.na(n_seeds_rel) & !is.na(tot_h_std)))
##
## Residuals:
      Min
               10 Median
                               30
                                      Max
## -1.8694 -1.0191 -0.6208 0.8837 3.7217
##
## Coefficients:
                            Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                                       0.144355
                                                7.115 1.04e-11 ***
                            1.027134
## FFD_corr_std
                           -0.003572
                                       0.142071 -0.025
                                                           0.980
                           -0.025346
                                       0.209006 -0.121
## treatmentF
                                                           0.904
## treatmentP
                                                0.223
                           0.047172
                                       0.211190
                                                           0.823
## treatmentPF
                           -0.117259
                                       0.206653 -0.567
                                                           0.571
## tot_h_std
                            0.183138
                                       0.076229
                                                  2.402
                                                           0.017 *
## FFD_corr_std:treatmentF -0.163085
                                       0.227570 -0.717
                                                           0.474
                                       0.204835 -0.414
## FFD_corr_std:treatmentP -0.084798
                                                           0.679
## FFD_corr_std:treatmentPF -0.194303
                                       0.206308 -0.942
                                                           0.347
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.228 on 265 degrees of freedom
## Multiple R-squared: 0.03315,
                                   Adjusted R-squared:
## F-statistic: 1.136 on 8 and 265 DF, p-value: 0.3394
BCIs_selection_2020_2
```

```
##
                                  lower
                                            upper
## FFD_corr_std
                            -0.27971064 0.2129141
## treatmentF
                            -0.39671000 0.3629940
## treatmentP
                            -0.38922188 0.5011998
## treatmentPF
                            -0.49368173 0.2447608
## tot h
                             0.01817649 0.3719280
## FFD_corr_std:treatmentF -0.48579939 0.2174295
## FFD_corr_std:treatmentP -0.47802734 0.3019227
## FFD_corr_std:treatmentPF -0.52234768 0.1770212
```

No significant effects (only total height).

3.3. Effects of temperature AND treatment on selection on FFD (2020)

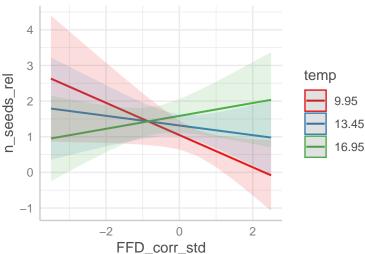
Using a linear model.

Call:

##

```
## lm(formula = n_seeds_rel ~ temp * FFD_corr_std + treatment *
##
      FFD_corr_std + tot_h_std, data = subset(ping_20_data, !is.na(n_seeds_rel) &
##
      !is.na(tot h std)))
##
## Residuals:
##
      Min
               1Q Median
                              3Q
                                    Max
## -1.6893 -1.0016 -0.3455 0.8113 3.4259
##
## Coefficients:
##
                          Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                           0.28060
                                     0.61951 0.453 0.650965
                           0.07672
                                     0.04641
                                              1.653 0.099499 .
## temp
## FFD_corr_std
                          ## treatmentF
                         -0.11716 0.20377 -0.575 0.565815
## treatmentP
                          0.07054 0.20533
                                             0.344 0.731466
## treatmentPF
                          -0.14646
                                   0.20050 -0.730 0.465745
## tot_h_std
                                              2.076 0.038863 *
                          0.15415 0.07425
## temp:FFD corr std
                           0.09042 0.02083
                                             4.342 2.02e-05 ***
## FFD_corr_std:treatmentF
                           0.21368
                                     0.23747
                                               0.900 0.369040
## FFD corr std:treatmentP
                           0.09114
                                     0.20317
                                              0.449 0.654104
## FFD_corr_std:treatmentPF 0.02324
                                     0.20717 0.112 0.910779
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.191 on 263 degrees of freedom
## Multiple R-squared: 0.09781,
                                 Adjusted R-squared: 0.06351
## F-statistic: 2.851 on 10 and 263 DF, p-value: 0.002187
BCIs selection 2020 4
##
                                 lower
                                           upper
                          -0.023617946 0.1738861
## temp
## FFD_corr_std
                          -2.040020629 -0.6110404
## treatmentF
                          -0.496925530 0.2730249
## treatmentP
                          -0.354602116 0.5047448
## treatmentPF
                          -0.512460343 0.2202682
## tot h
                           0.003394836 0.3326810
## temp:FFD_corr_std
                          0.050378402 0.1379978
## FFD_corr_std:treatmentF -0.159222227 0.6057428
## FFD_corr_std:treatmentP -0.284608626 0.4577974
## FFD_corr_std:treatmentPF -0.339589420 0.3896713
plot(ggpredict(selection_2020_4,terms=c("FFD_corr_std","temp")))
```

Predicted values of n_seeds_rel



3.4. Effects of all 2-way interactions between temperature, treatment and FFD (2020)

Using a linear model.

```
##
## Call:
## lm(formula = n_seeds_rel ~ temp * FFD_corr_std + treatment *
       FFD_corr_std + temp * treatment + tot_h_std, data = subset(ping_20_data,
##
##
       !is.na(n_seeds_rel) & !is.na(tot_h_std)))
##
## Residuals:
                1Q Median
##
       Min
                                3Q
                                       Max
## -1.6351 -0.9620 -0.2950 0.8474 3.4640
## Coefficients:
##
                            Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                             1.51592
                                        1.22386
                                                  1.239 0.216598
                                        0.09136
                                                 -0.172 0.863484
## temp
                            -0.01572
## FFD corr std
                            -1.60380
                                        0.43737
                                                 -3.667 0.000298 ***
## treatmentF
                             0.26473
                                        1.62297
                                                  0.163 0.870555
## treatmentP
                            -2.07773
                                        1.65538
                                                 -1.255 0.210555
## treatmentPF
                            -3.42343
                                        1.69283
                                                 -2.022 0.044168 *
## tot_h_std
                             0.13784
                                        0.07423
                                                  1.857 0.064441
                                        0.02078
## temp:FFD_corr_std
                                                  4.198 3.71e-05 ***
                             0.08721
## FFD corr std:treatmentF
                             0.14024
                                        0.45052
                                                   0.311 0.755833
## FFD_corr_std:treatmentP
                             0.60736
                                        0.44590
                                                  1.362 0.174345
## FFD_corr_std:treatmentPF
                             0.77045
                                        0.44195
                                                  1.743 0.082464 .
```

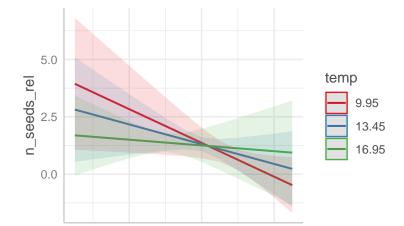
```
## temp:treatmentF
                           -0.02779
                                      0.11943 -0.233 0.816178
                            0.16131
                                      0.12339
                                                1.307 0.192249
## temp:treatmentP
## temp:treatmentPF
                                      0.12495
                            0.24364
                                                1.950 0.052263 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.182 on 260 degrees of freedom
## Multiple R-squared: 0.1211, Adjusted R-squared: 0.0772
## F-statistic: 2.757 on 13 and 260 DF, p-value: 0.001094
```

BCIs_selection_2020_5

```
##
                                 lower
                                            upper
                           -0.18447092 0.1607550
## temp
## FFD_corr_std
                           -2.47534382 -0.7150114
## treatmentF
                           -3.02921143 3.3993082
## treatmentP
                           -5.34210848 1.8442826
## treatmentPF
                           -6.36350324 -0.3376139
                           -0.01351256 0.3140180
## tot h
## temp:FFD_corr_std
                            0.04547299 0.1360521
## FFD_corr_std:treatmentF -0.67222847 0.9639395
## FFD_corr_std:treatmentP -0.36608287 1.4637105
## FFD_corr_std:treatmentPF -0.03973203 1.5656117
## temp:treatmentF
                           -0.25832508 0.2037471
## temp:treatmentP
                           -0.11298434 0.4178062
## temp:treatmentPF
                            0.01438222 0.4559907
```

Predicted values of n seeds rel

plot(ggpredict(selection_2020_5,terms=c("FFD_corr_std","temp")))



0

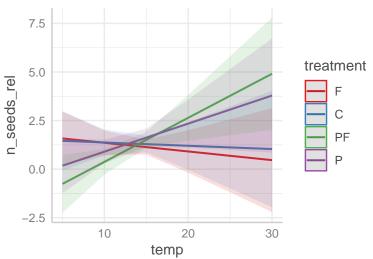
FFD_corr_std

-2

```
plot(ggpredict(selection_2020_5,terms=c("temp","treatment")))
```

2

Predicted values of n_seeds_rel



Similar result as in the model for fitness (section 2.3): Fitness increases with temperature in the PF treatment.

3.5. Effects of all 2-way interactions between temperature, treatment and FFD + 3-way interaction (2020)

Using a linear model.

```
##
## Call:
## lm(formula = n_seeds_rel ~ temp * FFD_corr_std * treatment +
       tot_h_std, data = subset(ping_20_data, !is.na(n_seeds_rel) &
##
##
       !is.na(tot h std)))
##
## Residuals:
##
       Min
                1Q Median
                                 3Q
                                        Max
  -2.1380 -0.9110 -0.2512 0.8369
##
##
## Coefficients:
                                   Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                                   2.005304
                                              1.230379
                                                          1.630
                                                                  0.1044
                                                         -0.663
## temp
                                  -0.061382
                                              0.092571
                                                                  0.5079
## FFD_corr_std
                                  -1.053468
                                              0.491097
                                                         -2.145
                                                                  0.0329 *
## treatmentF
                                   0.030751
                                              1.784860
                                                          0.017
                                                                  0.9863
## treatmentP
                                  -3.430192
                                              1.719298
                                                         -1.995
                                                                  0.0471 *
## treatmentPF
                                  -4.126835
                                              1.718899
                                                         -2.401
                                                                  0.0171 *
                                                         1.960
## tot_h_std
                                   0.144532
                                              0.073758
                                                                  0.0511 .
## temp:FFD_corr_std
                                   0.047660
                                              0.026434
                                                          1.803
                                                                  0.0726 .
## temp:treatmentF
                                  -0.005886
                                              0.138464
                                                        -0.043
                                                                  0.9661
## temp:treatmentP
                                   0.296603
                                              0.132757
                                                          2.234
                                                                  0.0263 *
                                              0.130234
## temp:treatmentPF
                                   0.315421
                                                          2.422
                                                                  0.0161 *
```

```
-0.083891
## FFD_corr_std:treatmentF
                                           1.061109 -0.079
                                                             0.9370
## FFD_corr_std:treatmentP
                               -1.273561
                                           0.852098 -1.495
                                                             0.1362
                                                             0.6382
## FFD corr std:treatmentPF
                               -0.427416
                                           0.908010 -0.471
## temp:FFD_corr_std:treatmentF
                                                             0.9099
                                0.009697
                                           0.085613
                                                     0.113
## temp:FFD_corr_std:treatmentP
                                0.142455
                                           0.055345
                                                     2.574
                                                             0.0106 *
## temp:FFD_corr_std:treatmentPF 0.087626
                                                     1.490
                                           0.058807
                                                             0.1374
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.172 on 257 degrees of freedom
## Multiple R-squared: 0.1467, Adjusted R-squared: 0.09359
## F-statistic: 2.762 on 16 and 257 DF, p-value: 0.0004132
```

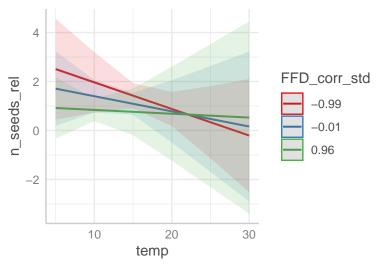
BCIs_selection_2020_6

```
##
                                       lower
## temp
                                -0.226726224 0.10929902
## FFD_corr_std
                                -1.939769426 -0.10589412
## treatmentF
                                -3.446587290 3.39754964
## treatmentP
                                -6.917407767 0.52682140
## treatmentPF
                                -7.125951010 -0.81714668
## tot h
                                -0.002727111 0.32661562
## temp:FFD_corr_std
                                0.005237069 0.09109947
                                -0.265373374 0.25384419
## temp:treatmentF
## temp:treatmentP
                                -0.010610699 0.56375640
## temp:treatmentPF
                                 0.058883052 0.55315635
## FFD_corr_std:treatmentF
                                -2.096140173 1.73915921
## FFD corr std:treatmentP
                                -3.169129726 0.39599662
## FFD_corr_std:treatmentPF
                                -2.142337872 1.60857445
## temp:FFD_corr_std:treatmentF
                                -0.137023324 0.16672187
## temp:FFD_corr_std:treatmentP
                                 0.025066586 0.27064164
## temp:FFD_corr_std:treatmentPF -0.051802621 0.21571639
```

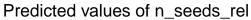
2-way significant interactions:

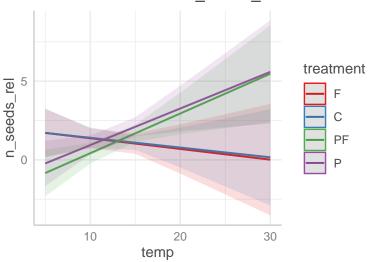
```
plot(ggpredict(selection_2020_6,terms=c("temp","FFD_corr_std")))
```

Predicted values of n_seeds_rel



plot(ggpredict(selection_2020_6,terms=c("temp","treatment")))



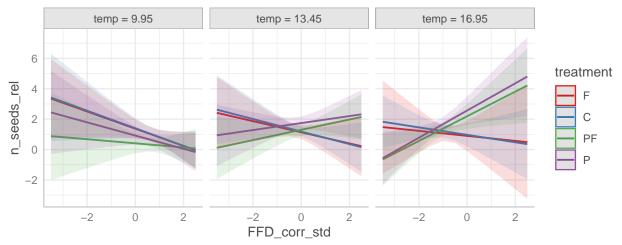


Similar results as before.

3-way interaction (two different representations):

plot(ggpredict(selection_2020_6,terms=c("FFD_corr_std","treatment","temp")))

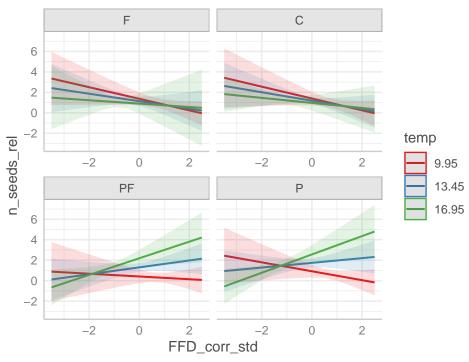
Predicted values of n_seeds_rel



At low temperatures (left panel above), it is always better to flower early. At medium and high temperatures (center and right panel above), it is better to flower early for plants in the C and F treatments, but better to flower late for plants in the P and PF treatments (only temp * FFD * treatmentP is significantly different from control).

plot(ggpredict(selection_2020_6,terms=c("FFD_corr_std","temp","treatment")))





In other words, for plants in C and F treatments, it is always better to flower early, irrespective of soil temperature. For plants in P and PF treatments (only in P if we look at significance), it is better to flower early at cold soil temperatures, but late at warm soil temperatures.

Do we want to include this 3-way interaction, or stay just with 2-way interactions?