Lathyrus ms2: Selection on reaction norms - multivariate modeling for phenotypic selection on plasticity 3 (Arnold et al. 2019 Phil. Trans. R. Soc. B)

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Code based on Arnold et al. 2019 Phil. Trans. R. Soc. B.

Fitting bivariate models of fitness and FFD, with random regressions for individuals, using a Poisson distribution for fitness, instead of Gaussian (and absolute instead of relative fitness). Using either total fitness or mean fitness per flowering event. Trying different alternatives for temperature (mean April, mean May, min April, mean April-May). Trying also to include number of flowers.

Using ids with 4 years of data

Total fitness

Temperature: mean April

Without number of flowers

Stack data:

2 2 2 2007

The bivariate MCMCglmm models with covariances between fitness and FFD intercepts and slopes are run using a stacked data set. We therefore need to first stack our data and add in index columns of 'traits', 'variable', and 'family', which are all required for MCMCglmm to fit a multivariate model of traits with different distributions. The values of the two variables need to be stacked into a single column and identified as either 'fitness' or 'FFD' with an index column. Likewise, the 'family' index column tells MCMCglmm what distribution family that response variable follows (here, Poisson or Gaussian). For this data-set, we put the fitness data first, then the repeated-measures FFD data.

```
# Create a single data-set "data.stack_pois", with single column at start to index observations
data.stack_pois <- c()</pre>
data.stack pois $0bs <- 1: (243 + 1455)
data.stack_pois$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Add first_yr to total data +
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data_4yrs_total_wfirstyr<-data_4yrs_total%>%
  right_join(data_4yrs[c(3,10)]%>%
               group_by(id)%>%
               summarise(first_yr=mean(first_yr)),by="id")
data.stack_pois$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                      data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois$temp <- c(rep(0, 243), data_4yrs$cmean_4)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness), data_4yrs$FFD)
# Create 3 index columns needed for MCMCqlmm
data.stack_pois$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack_pois$variable <- data.stack_pois$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois$family <- c(rep("poisson", 243), rep("gaussian", 1455))
data.stack_pois <- data.frame(data.stack_pois)</pre>
data.stack_pois$id <- as.factor(data.stack_pois$id)</pre>
data.stack_pois$year <- as.factor(data.stack_pois$year)</pre>
head(data.stack_pois)
     Obs id year temp fitness.FFD.stack traits variable family
## 1
     1 1 2006
                    0
                                    14 fitness fitness poisson
```

4 fitness fitness poisson

```
3 3 2007
## 3
                                      2 fitness fitness poisson
## 4
       4 4 2007
                    0
                                      6 fitness fitness poisson
## 5
       5 5 2007
                    0
                                      4 fitness fitness poisson
       6 6 2007
## 6
                    Λ
                                      2 fitness fitness poisson
# Scaling factor for MCMCglmm iterations
sc <- 100#0 # Increase this parameter for longer runs
priorBiv_RR_pois <- list(G = list(G1 = list(V = diag(1), nu = 1)),</pre>
                    # ^ random effect for year (fitted for FFD only)
                    R = list(R1 = list(V = diag(3), nu = 3, covu = TRUE),
                             # ^ 3-way var-cov matrix of (id + temp:id) for FFD,
                             # residual for fitness
                             R2 = list(V = diag(1), nu = 1))) # residual for FFD
modelBV_RR_pois <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
kable(summary(modelBV_RR_pois)$solutions,digits=c(3,3,3,0,3),caption="Fixed effects")
```

1-95% CI u-95% CI pMCMC post.mean eff.samp variableFFD 0.001 58.129 55.994 60.098 1000 variablefitness 0.9740.8441277 0.0011.088 at.level(variable, "FFD"):temp -2.417-3.922-0.7991000 0.004

Table 1: Fixed effects

kable(summary(modelBV_RR_pois) \$Gcovariances, digits=c(3,3,3,0), caption="Random effects")

Table 2: Random effects

	post.mean	l-95% CI	u-95% CI	eff.samp
at.level(variable, "FFD"):at.level(variable, "FFD").year	26.45	12.739	45.378	1000

kable(summary(modelBV_RR_pois)\$Rcovariances,digits=c(3,3,3,0),caption="Random effects")

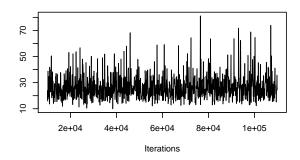
Table 3: Random effects

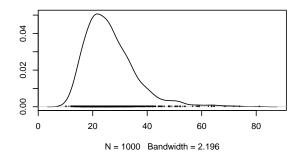
	post.mean	l-95% CI	u-95% CI	eff.samp
at.level(variable, "FFD").id:at.level(variable, "FFD").id	3.092	1.913	4.200	1102
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD").id	0.855	0.367	1.323	1000
at.level(variable, "fitness").id:at.level(variable, "FFD").id	-0.650	-0.953	-0.342	1000
at.level(variable, "FFD").id:at.level(variable, "FFD"):temp.id	0.855	0.367	1.323	1000
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD"):temp.id	0.717	0.369	1.112	1000
at.level(variable, "fitness").id:at.level(variable, "FFD"):temp.id	-0.128	-0.300	0.050	1197
at.level(variable, "FFD").id:at.level(variable, "fitness").id	-0.650	-0.953	-0.342	1000
at.level(variable, "FFD"):temp.id:at.level(variable, "fitness").id	-0.128	-0.300	0.050	1197
at.level(variable, "fitness").id:at.level(variable, "fitness").id	0.524	0.374	0.681	1066
at.level(variable, "FFD"):at.level(variable, "FFD").Obs	18.599	17.086	20.257	1262

Meaning of each row: "Among-id var in intercept for FFD", "Among-id covar between intercept and slope for FFD", "Among-individual covar between fitness and var in intercepts for FFD", "Among-id covar between intercept and slope for FFD", "Among-id var in slope for FFD", "Among-id covar between slope for FFD and intercept for fitness", "Among-individual covar between fitness and var in intercepts for FFD", "Among-id covar between slope for FFD and intercept for fitness", "Among-id var in intercept for fitness", "Within-id var in FFD between years".

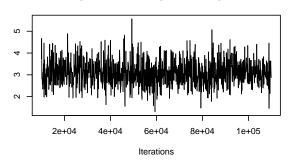
plot(modelBV_RR_pois\$VCV[,1:4])

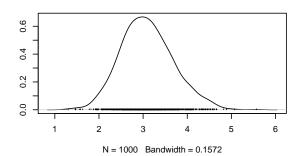
Trace of at.level(variable, "FFD"):at.level(variable, "FFD").ye Density of at.level(variable, "FFD"):at.level(variable, "FFD").ye



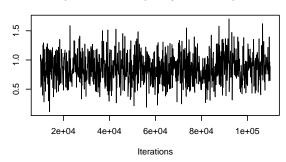


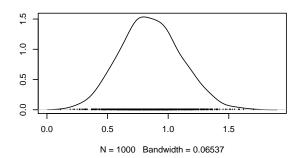
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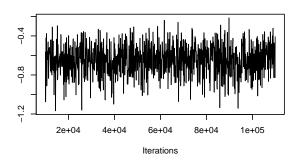


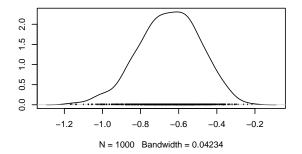
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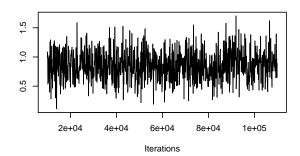
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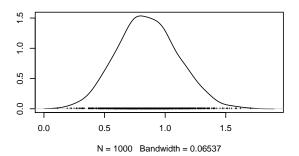




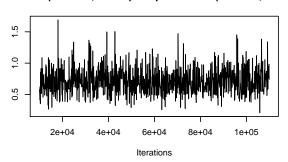
plot(modelBV_RR_pois\$VCV[,5:8])

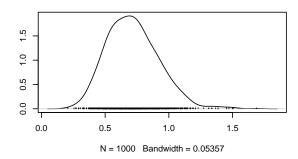
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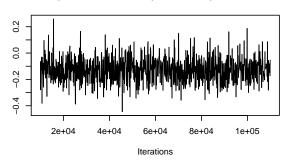


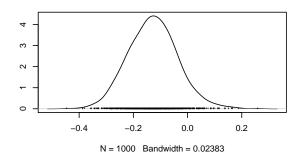
e of at.level(variable, "FFD"):temp.id:at.level(variable, "FFD")ity of at.level(variable, "FFD"):temp.id:at.level(variable, "FFD")



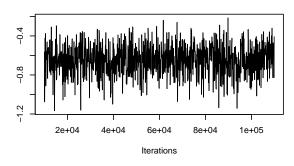


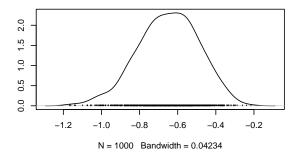
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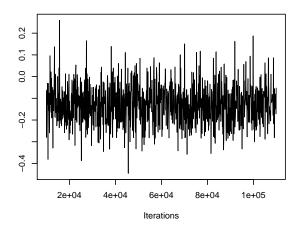
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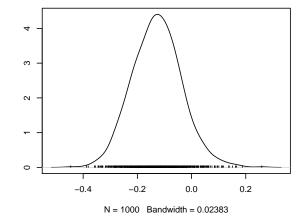




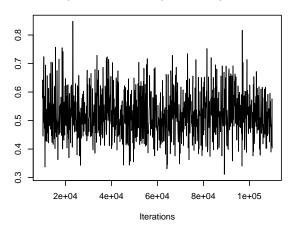
plot(modelBV_RR_pois\$VCV[,9:11])

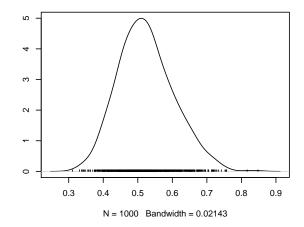
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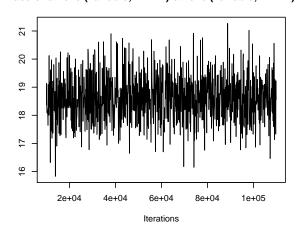


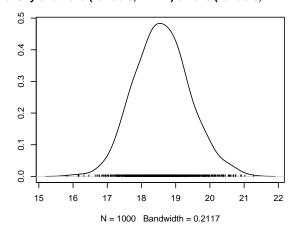
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Trace of at.level(variable, "FFD"):at.level(variable, "FFD").O Density of at.level(variable, "FFD"):at.level(variable, "FFD").C





Check for autocorrelation between successive stored iterations (suggested to be less than 0.1):

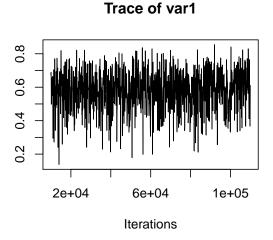
kable(diag(autocorr(modelBV_RR_pois\$VCV)[2, ,]),caption="Autocorrelation")

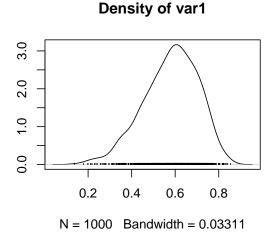
Table 4: Autocorrelation

	X
at.level(variable, "FFD"):at.level(variable, "FFD").year	-0.0091692
at.level(variable, "FFD").id:at.level(variable, "FFD").id	-0.0489505
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD").id	-0.0248751
at.level(variable, "fitness").id:at.level(variable, "FFD").id	-0.0258161
at.level(variable, "FFD").id:at.level(variable, "FFD"):temp.id	-0.0248751
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD"):temp.id	-0.0187903
at.level(variable, "fitness").id:at.level(variable, "FFD"):temp.id	-0.0331344
at.level(variable, "FFD").id:at.level(variable, "fitness").id	-0.0258161
at.level(variable, "FFD"):temp.id:at.level(variable, "fitness").id	-0.0331344
at.level(variable, "fitness").id:at.level(variable, "fitness").id	0.0337469
at.level(variable, "FFD"):at.level(variable, "FFD").Obs	0.0067911

Ensure that the among-individual correlation between intercepts and slopes for FFD is (approximately) the same as we estimated in our earlier univariate random regression model.

```
cor_BV_RR_pois_intslope <-
   modelBV_RR_pois$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\").id"]/
(sqrt(modelBV_RR_pois$VCV[,"at.level(variable, \"FFD\").id:at.level(variable, \"FFD\").id"])*
sqrt(modelBV_RR_pois$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.id"]))
plot(cor_BV_RR_pois_intslope)</pre>
```





```
posterior.mode(cor_BV_RR_pois_intslope)

## var1
## 0.594561

HPDinterval(cor_BV_RR_pois_intslope)

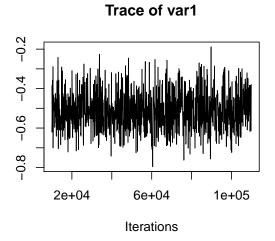
## lower upper
## var1 0.3298192 0.7825926
## attr(,"Probability")
## [1] 0.95
```

We find a strong positive correlation between among-individual variance in intercepts and slopes, at the

intercept (x = 0). Although it is a bit lower than in our earlier univariate random regression model (OK?).

Determining the among-individual correlation between FFD and fitness:

```
cor_BV_RR_pois_intfit <-
   modelBV_RR_pois$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\").id"]/
   (sqrt(modelBV_RR_pois$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id"])*
        sqrt(modelBV_RR_pois$VCV[,"at.level(variable, \"FFD\").id:at.level(variable, \"FFD\").id"]))
plot(cor_BV_RR_pois_intfit)</pre>
```



∞ - 0 -

Density of var1

```
posterior.mode(cor_BV_RR_pois_intfit)

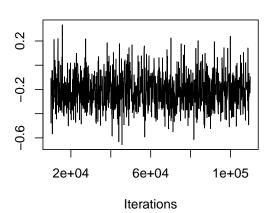
## var1
## -0.5706904

HPDinterval(cor_BV_RR_pois_intfit)

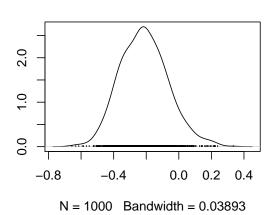
## lower upper
## var1 -0.6970216 -0.3091069
## attr(,"Probability")
## [1] 0.95
```

Negative correlation: Fitness increases when FFD decreases (i.e. is earlier).

```
cor_BV_RR_pois_slopefit <-
modelBV_RR_pois$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
   (sqrt(modelBV_RR_pois$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id"])*
        sqrt(modelBV_RR_pois$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.id
plot(cor_BV_RR_pois_slopefit)</pre>
```



Density of var1



```
posterior.mode(cor_BV_RR_pois_slopefit)

## var1
## -0.2251261

HPDinterval(cor_BV_RR_pois_slopefit)
```

```
## lower upper
## var1 -0.5012255 0.0667102
## attr(,"Probability")
## [1] 0.95
```

Negative correlation: Fitness increases when the slope for FFD decreases (i.e. is more negative, and therefore plasticity increases). Fitness is higher in more plastic individuals. However, this correlation is not significant because the CIs encompass zero!!!

Extract selection coefficients

[1,] 2.8949557 0.7508321 -0.6104302

Selection differentials or gradients should be calculated using relative fitness, and models are typically fitted assuming Gaussian errors. However, where the fitness measure follows a non-Gaussian distribution, as is typically the case with skewed distributions of fitness, a GLMM of absolute fitness will be preferable. The resulting covariances returned by the model will then be between the trait on the data scale and fitness on a 'latent' (link-function) scale. These estimates need to be transformed if data-scale estimates of selection are required. However, in the case of a GLMM with a log-link function (e.g. Poisson, over-dispersed Poisson, or negative binomial distribution), it is possible to exploit the fact that the latent-scale covariance with absolute fitness is equivalent to the data-scale covariance of relative fitness: consequently, and conveniently, the covariance components of Pind on the latent scale can simply be treated as selection differentials S. By extension, estimates of b as indicated above will also provide data-scale selection gradients.

```
# Extract 3x3 matrix of variance-covariance values for intercepts and slopes of X, and LBS
# These are in the 2nd-10th columns of model output
P.modelBV_RR_pois <- modelBV_RR_pois$VCV[,2:10]
P.modelBV_RR_pois.mode <- matrix(1:9, nrow = 3)
for (k in 1:9) P.modelBV_RR_pois.mode[k] <- posterior.mode(P.modelBV_RR_pois[,k])
P.modelBV_RR_pois.mode
### [,1] [,2] [,3]</pre>
```

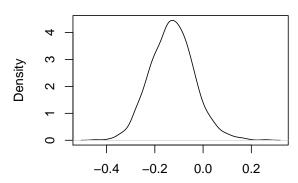
```
## [2,] 0.7508321 0.6280586 -0.1135805
## [3,] -0.6104302 -0.1135805 0.5308708
# Extract selection *differentials* (i.e. covariances) for intercept and slope:
# and calculate posterior mode and credible intervals for each
S.modelBV_RR_pois <- modelBV_RR_pois$VCV[, c(4,7)]</pre>
S.modelBV_RR_pois <- P.modelBV_RR_pois[, c(3,6)]</pre>
colnames(S.modelBV_RR_pois) <- c("S_intercepts", "S_slopes")</pre>
S.modelBV RR pois.mode <- P.modelBV RR pois.mode[1:2, 3]
S.modelBV_RR_pois.mode
## [1] -0.6104302 -0.1135805
posterior.mode(mcmc(S.modelBV_RR_pois))
## S_intercepts
                    S_slopes
     -0.6104302
                  -0.1135805
HPDinterval(mcmc(S.modelBV_RR_pois))
##
                     lower
## S intercepts -0.9529606 -0.34190093
## S slopes
                -0.3003430 0.05017838
## attr(,"Probability")
## [1] 0.95
# Plot posterior distribution of selection differentials
par(mfrow = c(1,2))
plot(density(S.modelBV_RR_pois[,1]), main = "S_intercepts")
plot(density(S.modelBV_RR_pois[,2]), main = "S_slopes")
```



Density -1.2 -0.8 -0.4

N = 1000 Bandwidth = 0.03595

S_slopes



N = 1000 Bandwidth = 0.02024

```
# Estimate selection gradients for intercept and slope (beta = S / P)
# on each sample of posterior and extract their mode
n <- length(modelBV_RR_pois$VCV[,2])  # sample size
beta_post_RR_pois <- matrix(NA, n ,2)

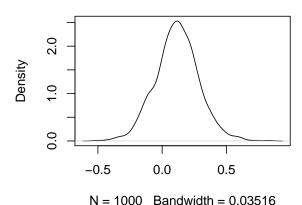
for (i in 1:n) {
    P3 <- matrix(rep(NA, 9), nrow = 3)  # 3x3 matrix of var-cov for individual X.int, X.slope and LBS
    for (k in 1:9) {P3[k] <- P.modelBV_RR_pois[i, k] }</pre>
```

```
P2 <- P3[1:2, 1:2] # 2x2 matrix of just trait intercept & slope var-cov
                   # selection differentials on traits (last column of P3)
  S \leftarrow P3[1:2, 3]
  beta_post_RR_pois[i,] <- solve(P2) %*% S # selection gradients beta = P^-1 * S
}
# Finally, extract and plot the selection gradients posterior modes
# and 95% credible intervals for both selection on intercepts (trait value)
# and slopes (trait plasticity).
# Note that credible intervals are not exactly confidence intervals. See here:
# https://statsdirect.com/help/basics/confidence_interval.htm and
# https://stats.stackexchange.com/questions/2272/
colnames(beta_post_RR_pois) <- c("beta_intercepts", "beta_slopes")</pre>
posterior.mode(mcmc(beta_post_RR_pois))
## beta_intercepts
                       beta_slopes
        -0.2569895
                         0.1445244
HPDinterval(mcmc(beta_post_RR_pois))
##
                        lower
                                   upper
## beta_intercepts -0.3733008 -0.1022348
## beta_slopes
                   -0.1892722 0.4764652
## attr(,"Probability")
## [1] 0.95
# Plot posterior distribution of selection gradients
par(mfrow = c(1,2))
plot(density(beta_post_RR_pois[,1]), main = "beta_intercepts")
plot(density(beta_post_RR_pois[,2]), main = "beta_slopes")
```



-0.5 -0.3 -0.1 0.0 N = 1000 Bandwidth = 0.01649

beta_slopes



NB selection differentials and gradients here are from covariances with latent-scale absolute fitness # These are equivalent to covariances with data-scale relative fitness: see main text of paper

With number of flowers

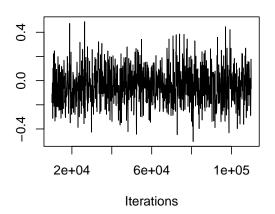
```
# Create a single data-set "data.stack_pois_nfl",
# with single column at start to index observations
data.stack pois nfl <- c()</pre>
data.stack_pois_nfl$0bs <- 1:(243 + 1455)</pre>
data.stack_pois_nfl$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois_nfl$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                     data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois_nfl$temp <- c(rep(0, 243), data_4yrs$cmean_4)</pre>
# Number of flowers column is only relevant for FFD, but is set to 0 for fitness values
data_4yrs$cn_fl<-scale(data_4yrs$n_fl,center=T,scale=F) # Standardize number of flowers
data.stack_pois_nfl$n_fl <- c(rep(0, 243), data_4yrs$cn_fl)</pre>
\# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois_nfl$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness), data_4yrs$FFD)
# Create 3 index columns needed for MCMCqlmm
data.stack_pois_nfl$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack_pois_nfl$variable <- data.stack_pois_nfl$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois_nfl$family <- c(rep("poisson", 243), rep("gaussian", 1455))</pre>
data.stack_pois_nfl <- data.frame(data.stack_pois_nfl)</pre>
data.stack_pois_nfl$id <- as.factor(data.stack_pois_nfl$id)</pre>
data.stack_pois_nfl$year <- as.factor(data.stack_pois_nfl$year)</pre>
head(data.stack_pois_nfl)
##
    Obs id year temp n_fl fitness.FFD.stack traits variable family
## 1 1 1 2006
                    0
                         0
                                          14 fitness fitness poisson
## 2 2 2 2007
                    0
                         0
                                           4 fitness fitness poisson
     3 3 2007
                                           2 fitness fitness poisson
## 3
                   0
                       0
                       0
## 4
      4 4 2007
                    0
                                            6 fitness fitness poisson
## 5 5 5 2007
                         0
                                            4 fitness fitness poisson
## 6 6 6 2007
                                            2 fitness fitness poisson
                    0
modelBV_RR_pois_nfl <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp + # single fixed effect of temp
                           at.level(variable, "FFD"):n_fl, # single fixed effect of n_fl
                         random = ~us(at.level(variable, "FFD")):year +
                           us(at.level(variable, "FFD") +
                                at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
```

```
# ^ residual variance within indivdiuals between years
# (labelled by 'Obs')
data = data.stack_pois_nfl,
prior = priorBiv_RR_pois,
family = NULL, # specified already in the data-set
nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

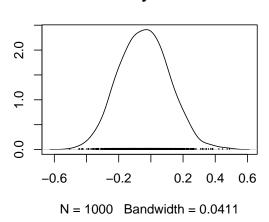
Determining the among-individual correlation between fitness and variation in slopes for FFD (we only focus in this result in this and the following models, to see if it reaches significance with any of the combinations):

```
cor_BV_RR_pois_nfl_slopefit <-
   modelBV_RR_pois_nfl$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
   (sqrt(modelBV_RR_pois_nfl$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id
        sqrt(modelBV_RR_pois_nfl$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.lot(cor_BV_RR_pois_nfl_slopefit)</pre>
```

Trace of var1



Density of var1



posterior.mode(cor_BV_RR_pois_nfl_slopefit)

```
## var1
## -0.04070236
HPDinterval(cor_BV_RR_pois_nfl_slopefit)
## lower upper
## var1 -0.3496906 0.2384892
## attr(,"Probability")
## [1] 0.95
```

Temperature: mean May

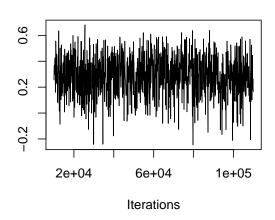
Mean-centred mean May temperature:

```
data_4yrs<-data_4yrs%>%
  mutate(cmean_5=scale(mean_5,center=T,scale=F))
```

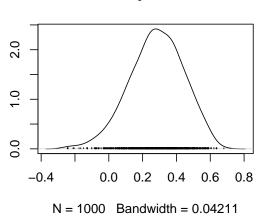
```
# Create a single data-set "data.stack_pois_may", with single column at start to index observations
data.stack_pois_may <- c()</pre>
data.stack_pois_may$0bs <- 1:(243 + 1455)</pre>
data.stack_pois_may$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois_may$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                     data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois_may$temp <- c(rep(0, 243), data_4yrs$cmean_5)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois_may$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness), data_4yrs$FFD)
# Create 3 index columns needed for MCMCqlmm
data.stack_pois_may$traits <- c(rep("fitness", 243), rep("FFD", 1455))
data.stack_pois_may$variable <- data.stack_pois_may$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois_may$family <- c(rep("poisson", 243), rep("gaussian", 1455))</pre>
data.stack_pois_may <- data.frame(data.stack_pois_may)</pre>
data.stack_pois_may$id <- as.factor(data.stack_pois_may$id)</pre>
data.stack_pois_may$year <- as.factor(data.stack_pois_may$year)</pre>
head(data.stack_pois_may)
     Obs id year temp fitness.FFD.stack traits variable family
## 1
       1 1 2006
                    0
                                      14 fitness fitness poisson
## 2
       2 2 2007
                                       4 fitness fitness poisson
                    0
## 3
       3 3 2007
                    0
                                       2 fitness fitness poisson
      4 4 2007
                    0
                                       6 fitness fitness poisson
       5 5 2007
## 5
                    0
                                       4 fitness fitness poisson
## 6
       6 6 2007
                                       2 fitness fitness poisson
modelBV_RR_pois_may <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                               at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois_may,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

```
cor_BV_RR_pois_may_slopefit <-
modelBV_RR_pois_may$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
   (sqrt(modelBV_RR_pois_may$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id
        sqrt(modelBV_RR_pois_may$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.
plot(cor_BV_RR_pois_may_slopefit)</pre>
```

Trace of var1



Density of var1



```
posterior.mode(cor_BV_RR_pois_may_slopefit)
```

var1 ## 0.279725

HPDinterval(cor_BV_RR_pois_may_slopefit)

```
## lower upper
## var1 -0.009500759 0.5907167
## attr(,"Probability")
## [1] 0.95
```

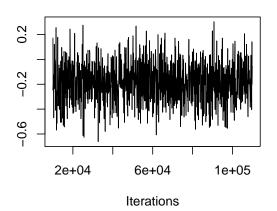
Temperature: min April

Mean-centred min April temperature:

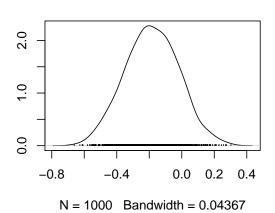
```
data_4yrs<-data_4yrs%>%
  mutate(cmin_4=scale(min_4,center=T,scale=F))
```

```
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois_min$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness), data_4yrs$FFD)
# Create 3 index columns needed for MCMCqlmm
data.stack_pois_min$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack_pois_min$variable <- data.stack_pois_min$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois_min$family <- c(rep("poisson", 243), rep("gaussian", 1455))</pre>
data.stack_pois_min <- data.frame(data.stack_pois_min)</pre>
data.stack_pois_min$id <- as.factor(data.stack_pois_min$id)</pre>
data.stack_pois_min$year <- as.factor(data.stack_pois_min$year)</pre>
head(data.stack_pois_may)
     Obs id year temp fitness.FFD.stack traits variable family
## 1
      1 1 2006
                    0
                                     14 fitness fitness poisson
## 2
      2 2 2007
                    Λ
                                      4 fitness fitness poisson
## 3 3 3 2007
                                      2 fitness fitness poisson
## 4 4 2007
                                      6 fitness fitness poisson
                    0
## 5
      5 5 2007
                                      4 fitness fitness poisson
                    0
## 6
       6 6 2007
                                      2 fitness fitness poisson
                    Ω
modelBV_RR_pois_min <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois_min,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

```
cor_BV_RR_pois_min_slopefit <-
    modelBV_RR_pois_min$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
    (sqrt(modelBV_RR_pois_min$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id
        sqrt(modelBV_RR_pois_min$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.
```



Density of var1



```
posterior.mode(cor_BV_RR_pois_min_slopefit)
```

```
## var1
## -0.1997217
```

HPDinterval(cor_BV_RR_pois_min_slopefit)

```
## lower upper
## var1 -0.5142569 0.1259144
## attr(,"Probability")
## [1] 0.95
```

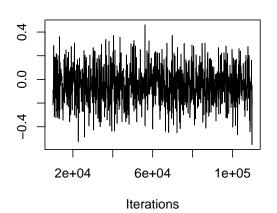
Temperature: mean April-May

Add April-May temperatures:

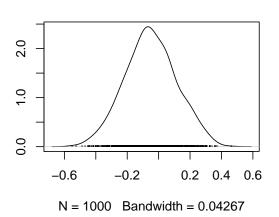
```
mean_weather_aprilmay$year<-as.integer(as.character(mean_weather_aprilmay$year))
data_4yrs<-data_4yrs%%
    right_join(unique(mean_weather_aprilmay),by="year")%>%
    mutate(cmean_45=scale(mean_45,center=T,scale=F))
```

```
# Create 3 index columns needed for MCMCqlmm
data.stack_pois_aprilmay$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack_pois_aprilmay$variable <- data.stack_pois_aprilmay$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois_aprilmay$family <- c(rep("poisson", 243), rep("gaussian", 1455))</pre>
data.stack_pois_aprilmay <- data.frame(data.stack_pois_aprilmay)</pre>
data.stack_pois_aprilmay$id <- as.factor(data.stack_pois_aprilmay$id)</pre>
data.stack_pois_aprilmay$year <- as.factor(data.stack_pois_aprilmay$year)</pre>
head(data.stack_pois_aprilmay)
     Obs id year temp fitness.FFD.stack traits variable family
## 1
      1 1 2006
                    0
                                     14 fitness fitness poisson
## 2
       2 2 2007
                    0
                                      4 fitness fitness poisson
       3 3 2007
## 3
                    0
                                      2 fitness fitness poisson
     4 4 2007
## 4
                    0
                                      6 fitness fitness poisson
## 5
      5 5 2007
                    0
                                       4 fitness fitness poisson
## 6
       6 6 2007
                    0
                                       2 fitness fitness poisson
modelBV_RR_pois_aprilmay <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois_aprilmay,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

```
cor_BV_RR_pois_aprilmay_slopefit <-
   modelBV_RR_pois_aprilmay$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id
   (sqrt(modelBV_RR_pois_aprilmay$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\
        sqrt(modelBV_RR_pois_aprilmay$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\")
plot(cor_BV_RR_pois_aprilmay_slopefit)</pre>
```



Density of var1



```
posterior.mode(cor_BV_RR_pois_aprilmay_slopefit)

## var1
## -0.07820992

HPDinterval(cor_BV_RR_pois_aprilmay_slopefit)

## lower upper
## var1 -0.3618492 0.2779801
## attr(,"Probability")
## [1] 0.95
```

Mean fitness per flowering event

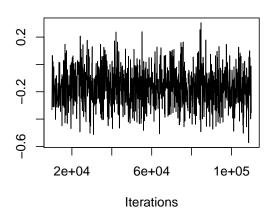
Using mean fitness per flowering event (sum of fitness divided by number of years when the plant flowered), instead of mean fitness over all years.

Temperature: mean April

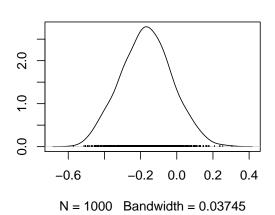
Without number of flowers

```
data.stack_pois2$temp <- c(rep(0, 243), data_4yrs$cmean_4)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack pois2$fitness.FFD.stack <- c(round(data 4yrs total$mean fitness fl), data 4yrs$FFD)
# Create 3 index columns needed for MCMCglmm
data.stack_pois2$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack pois2$variable <- data.stack pois2$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois2$family <- c(rep("poisson", 243), rep("gaussian", 1455))
data.stack_pois2 <- data.frame(data.stack_pois2)</pre>
data.stack_pois$id <- as.factor(data.stack_pois$id)</pre>
data.stack_pois$year <- as.factor(data.stack_pois$year)</pre>
head(data.stack_pois)
    Obs id year temp fitness.FFD.stack traits variable family
## 1
      1 1 2006
                                     14 fitness fitness poisson
       2 2 2007
## 2
                                      4 fitness fitness poisson
                    0
## 3
      3 3 2007
                    0
                                      2 fitness fitness poisson
## 4
      4 4 2007
                    0
                                       6 fitness fitness poisson
       5 5 2007
                                       4 fitness fitness poisson
## 5
                    0
                                       2 fitness fitness poisson
## 6
      6 6 2007
modelBV_RR_pois2 <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                          # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois2,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
Determining the among-individual correlation between fitness and variation in slopes for FFD:
cor_BV_RR_pois2_slopefit <-</pre>
  modelBV_RR_pois2$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
  (sqrt(modelBV_RR_pois2$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id"])
     sqrt(modelBV_RR_pois2$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.id
plot(cor_BV_RR_pois2_slopefit)
```

Temperature column is only relevant for FFD, but is set to 0 for fitness values



Density of var1



```
posterior.mode(cor_BV_RR_pois2_slopefit)
```

```
## var1
## -0.1766855
```

HPDinterval(cor_BV_RR_pois2_slopefit)

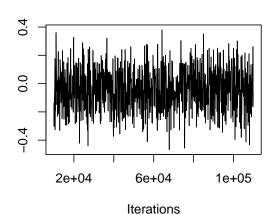
```
## lower upper
## var1 -0.4343329 0.1130045
## attr(,"Probability")
## [1] 0.95
```

With number of flowers

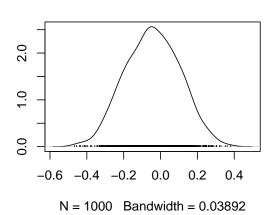
```
# Create a single data-set "data.stack pois2 nfl",
# with single column at start to index observations
data.stack pois2 nfl <- c()</pre>
data.stack_pois2_nfl$0bs <- 1:(243 + 1455)</pre>
data.stack_pois2_nfl$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois2_nfl$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                     data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois2_nfl$temp <- c(rep(0, 243), data_4yrs$cmean_4)</pre>
# Number of flowers column is only relevant for FFD, but is set to 0 for fitness values
data_4yrs$cn_fl<-scale(data_4yrs$n_fl,center=T,scale=F) # Standardize number of flowers
data.stack_pois2_nfl$n_fl <- c(rep(0, 243), data_4yrs$cn_fl)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack pois2 nfl$fitness.FFD.stack <- c(round(data 4yrs total$mean fitness fl), data 4yrs$FFD)
# Create 3 index columns needed for MCMCqlmm
data.stack_pois2_nfl$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
```

```
data.stack_pois2_nfl$variable <- data.stack_pois_nfl$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois2_nfl$family <- c(rep("poisson", 243), rep("gaussian", 1455))
data.stack_pois2_nfl <- data.frame(data.stack_pois2_nfl)</pre>
data.stack pois2 nfl$id <- as.factor(data.stack pois2 nfl$id)</pre>
data.stack_pois2_nfl$year <- as.factor(data.stack_pois2_nfl$year)</pre>
head(data.stack_pois2_nfl)
##
     Obs id year temp n_fl fitness.FFD.stack traits variable family
## 1
       1 1 2006
                    0
                         0
                                          16 fitness fitness poisson
       2 2 2007
## 2
                                           6 fitness fitness poisson
                    0
                         0
                       0
## 3
       3 3 2007
                    Ω
                                           3 fitness fitness poisson
       4 4 2007
                         0
                                           7 fitness fitness poisson
## 5
       5 5 2007
                    0
                         0
                                           4 fitness fitness poisson
## 6
       6 6 2007
                         0
                                           3 fitness fitness poisson
modelBV_RR_pois2_nfl <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         \# ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp + # single fixed effect of temp
                           at.level(variable, "FFD"):n_fl, # single fixed effect of n_fl
                         random = ~us(at.level(variable, "FFD")):year +
                           us(at.level(variable, "FFD") +
                                at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois2_nfl,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

```
cor_BV_RR_pois2_nfl_slopefit <-
modelBV_RR_pois2_nfl$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
  (sqrt(modelBV_RR_pois2_nfl$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").ic
        sqrt(modelBV_RR_pois2_nfl$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.id
```



Density of var1



posterior.mode(cor_BV_RR_pois2_nfl_slopefit)

```
## var1
## 0.03757234
```

HPDinterval(cor_BV_RR_pois2_nfl_slopefit)

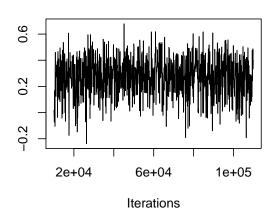
```
## lower upper
## var1 -0.3322421 0.2195263
## attr(,"Probability")
## [1] 0.95
```

Temperature: mean May

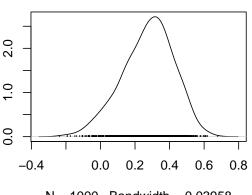
```
# Create a single data-set "data.stack_pois2_may", with single column at start to index observations
data.stack_pois2_may <- c()</pre>
data.stack_pois2_may$0bs <- 1:(243 + 1455)</pre>
data.stack_pois2_may$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois2_may$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                     data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois2_may$temp <- c(rep(0, 243), data_4yrs$cmean_5)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois2_may$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness_fl), data_4yrs$FFD)
# Create 3 index columns needed for MCMCglmm
data.stack_pois2_may$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack pois2 may$variable <- data.stack pois2 may$traits
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois2_may$family <- c(rep("poisson", 243), rep("gaussian", 1455))</pre>
```

```
data.stack_pois2_may <- data.frame(data.stack_pois2_may)</pre>
data.stack_pois2_may$id <- as.factor(data.stack_pois2_may$id)</pre>
data.stack_pois2_may$year <- as.factor(data.stack_pois2_may$year)</pre>
head(data.stack_pois2_may)
##
     Obs id year temp fitness.FFD.stack traits variable family
      1 1 2006
## 1
                                     16 fitness fitness poisson
## 2
       2 2 2007
                                      6 fitness fitness poisson
                    0
      3 3 2007
                                      3 fitness fitness poisson
## 3
                    0
## 4
     4 4 2007
                    Ω
                                      7 fitness fitness poisson
## 5
      5 5 2007
                                      4 fitness fitness poisson
      6 6 2007
                                      3 fitness fitness poisson
## 6
                    0
modelBV_RR_pois2_may <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack pois2 may,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

```
cor_BV_RR_pois2_may_slopefit <-
modelBV_RR_pois2_may_$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
  (sqrt(modelBV_RR_pois2_may_$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").i
        sqrt(modelBV_RR_pois2_may_$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.
```



Density of var1



N = 1000 Bandwidth = 0.03958

```
posterior.mode(cor_BV_RR_pois2_may_slopefit)

## var1
## 0.3352356

HPDinterval(cor_BV_RR_pois2_may_slopefit)

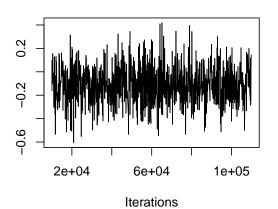
## lower upper
## var1 -0.0138436 0.555845
## attr(,"Probability")
## [1] 0.95
```

Temperature: min April

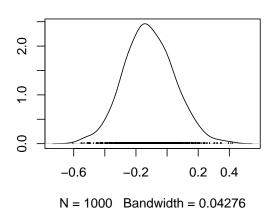
```
# Create a single data-set "data.stack_pois2_min", with single column at start to index observations
data.stack_pois2_min <- c()</pre>
data.stack_pois2_min$0bs <- 1:(243 + 1455)</pre>
data.stack_pois2_min$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois2_min$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                     data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois2_min$temp <- c(rep(0, 243), data_4yrs$cmin_4)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois2_min$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness_fl), data_4yrs$FFD)
# Create 3 index columns needed for MCMCglmm
data.stack_pois2_min$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack pois2 min$variable <- data.stack pois2 min$traits
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois2_min$family <- c(rep("poisson", 243), rep("gaussian", 1455))</pre>
```

```
data.stack_pois2_min <- data.frame(data.stack_pois2_min)</pre>
data.stack_pois2_min$id <- as.factor(data.stack_pois2_min$id)</pre>
data.stack_pois2_min$year <- as.factor(data.stack_pois2_min$year)</pre>
head(data.stack_pois2_min)
##
     Obs id year temp fitness.FFD.stack traits variable family
      1 1 2006
## 1
                    0
                                     16 fitness fitness poisson
       2 2 2007
## 2
                                      6 fitness fitness poisson
                    0
## 3
     3 3 2007
                                      3 fitness fitness poisson
                    0
## 4
     4 4 2007
                    Ω
                                      7 fitness fitness poisson
## 5
      5 5 2007
                                      4 fitness fitness poisson
      6 6 2007
                                      3 fitness fitness poisson
## 6
                    0
modelBV_RR_pois2_min <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack pois2 min,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

```
cor_BV_RR_pois2_min_slopefit <-
    modelBV_RR_pois2_min$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
    (sqrt(modelBV_RR_pois2_min$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id
        sqrt(modelBV_RR_pois2_min$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):t
```



Density of var1



```
posterior.mode(cor_BV_RR_pois2_min_slopefit)

## var1
## -0.09822202

HPDinterval(cor_BV_RR_pois2_min_slopefit)

## lower upper
## var1 -0.4151616 0.2124247
```

Temperature: mean April-May

attr(,"Probability")

Stack data:

[1] 0.95

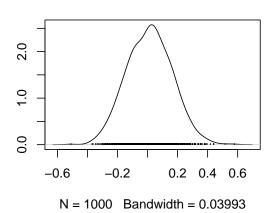
```
# Create a single data-set "data.stack_pois2_aprilmay",
# with single column at start to index observations
data.stack_pois2_aprilmay <- c()</pre>
data.stack_pois2_aprilmay$0bs <- 1:(243 + 1455)</pre>
data.stack_pois2_aprilmay$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois2_aprilmay$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                     data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois2_aprilmay$temp <- c(rep(0, 243), data_4yrs$cmean_45)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois2_aprilmay$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness_f1), data_4yrs$FFD)
# Create 3 index columns needed for MCMCglmm
data.stack pois2 aprilmay$traits <- c(rep("fitness", 243), rep("FFD", 1455))
data.stack_pois2_aprilmay$variable <- data.stack_pois2_aprilmay$traits
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
```

```
data.stack_pois2_aprilmay$family <- c(rep("poisson", 243), rep("gaussian", 1455))
data.stack_pois2_aprilmay <- data.frame(data.stack_pois2_aprilmay)</pre>
data.stack_pois2_aprilmay$id <- as.factor(data.stack_pois2_aprilmay$id)</pre>
data.stack_pois2_aprilmay$year <- as.factor(data.stack_pois2_aprilmay$year)
head(data.stack_pois2_aprilmay)
     Obs id year temp fitness.FFD.stack traits variable family
##
## 1
       1 1 2006
                                     16 fitness fitness poisson
       2 2 2007
                                      6 fitness fitness poisson
## 2
                    0
## 3
      3 3 2007
                    Ω
                                      3 fitness fitness poisson
## 4
      4 4 2007
                    0
                                      7 fitness fitness poisson
     5 5 2007
## 5
                    0
                                      4 fitness fitness poisson
       6 6 2007
                                      3 fitness fitness poisson
## 6
                    0
modelBV_RR_pois2_aprilmay <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois2_aprilmay,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```



2e+04 6e+04 1e+05 Iterations

Density of var1



```
posterior.mode(cor_BV_RR_pois2_aprilmay_slopefit)

## var1
## 0.02172168

HPDinterval(cor_BV_RR_pois2_aprilmay_slopefit)

## lower upper
## var1 -0.2724656 0.2983329
## attr(,"Probability")
```

Summary 4 years

[1] 0.95

Among-individual correlation between fitness and variation in slopes for FFD for each model:

```
cor_slopefit<-rbind(cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_slopefit),</pre>
                          rbind(HPDinterval(cor_BV_RR_pois_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_nfl_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois_nfl_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_may_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois_may_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_min_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois_min_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_aprilmay_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois_aprilmay_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois2_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois2_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois2_nfl_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois2_nfl_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois2_may_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois2_may_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois2_min_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois2_min_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois2_aprilmay_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois2_aprilmay_slopefit)[,1:2])))
row.names(cor_slopefit) <- c("Total fitness, mean April temperature, without n_fl",
```

```
"Total fitness, mean April temperature, with n_fl",

"Total fitness, mean May temperature",

"Total fitness, min April temperature",

"Total fitness, mean April-May temperature",

"Mean fitness per flowering event, mean April temperature, without n_fl",

"Mean fitness per flowering event, mean April temperature, with n_fl",

"Mean fitness per flowering event, mean May temperature",

"Mean fitness per flowering event, min April temperature",

"Mean fitness per flowering event, mean April-May temperature")

kable(cor_slopefit,digits=3)
```

	posterior.mode	lower	upper
Total fitness, mean April temperature, without n_fl	-0.225	-0.501	0.067
Total fitness, mean April temperature, with n_fl	-0.041	-0.350	0.238
Total fitness, mean May temperature	0.280	-0.010	0.591
Total fitness, min April temperature	-0.200	-0.514	0.126
Total fitness, mean April-May temperature	-0.078	-0.362	0.278
Mean fitness per flowering event, mean April temperature, without n_fl	-0.177	-0.434	0.113
Mean fitness per flowering event, mean April temperature, with n_fl	0.038	-0.332	0.220
Mean fitness per flowering event, mean May temperature	0.335	-0.014	0.556
Mean fitness per flowering event, min April temperature	-0.098	-0.415	0.212
Mean fitness per flowering event, mean April-May temperature	0.022	-0.272	0.298

Using ids with 5 years of data

All models without number of flowers.

Total fitness

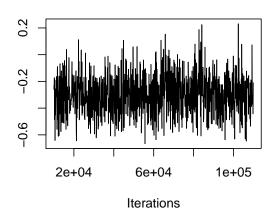
Temperature: mean April

Without number of flowers

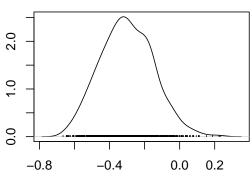
```
# Create 3 index columns needed for MCMCqlmm
data.stack_pois_5$traits <- c(rep("fitness", 156), rep("FFD", 1107))</pre>
data.stack_pois_5$variable <- data.stack_pois_5$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois_5$family <- c(rep("poisson", 156), rep("gaussian", 1107))</pre>
data.stack_pois_5 <- data.frame(data.stack_pois_5)</pre>
data.stack_pois_5$id <- as.factor(data.stack_pois_5$id)</pre>
data.stack_pois_5$year <- as.factor(data.stack_pois_5$year)</pre>
head(data.stack_pois_5)
     Obs id year temp fitness.FFD.stack traits variable family
       1 1 2006
## 1
                                     14 fitness fitness poisson
## 2
       2 2 2007
                                       4 fitness fitness poisson
                    0
       3 3 2007
## 3
                    0
                                       2 fitness fitness poisson
       4 4 2007
## 4
                    0
                                       6 fitness fitness poisson
## 5
       5 5 2007
                    0
                                       4 fitness fitness poisson
## 6
       6 6 2007
                                       2 fitness fitness poisson
modelBV_RR_pois_5 <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois_5,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
cor_BV_RR_pois_5_slopefit <-</pre>
  modelBV RR pois 5$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
  (sqrt(modelBV_RR_pois_5$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id"]
     sqrt(modelBV_RR_pois_5$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.
plot(cor_BV_RR_pois_5_slopefit)
```

Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:

data.stack_pois_5\fitness.FFD.stack <- c(round(data_5yrs_total\finesn_fitness), data_5yrs\fitness), data_5yrs\fitness



Density of var1

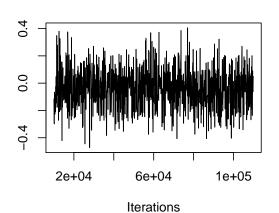


N = 1000 Bandwidth = 0.04008

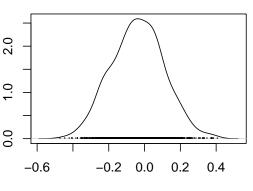
With number of flowers

```
# Create a single data-set "data.stack pois 5 nfl",
# with single column at start to index observations
data.stack_pois_5_nfl <- c()</pre>
data.stack_pois_5_nfl$0bs <- 1:(243 + 1455)</pre>
data.stack_pois_5_nfl$id <- c(data_4yrs_total$id, data_4yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois_5_nfl$year <- c(data_4yrs_total_wfirstyr$first_yr,</pre>
                     data_4yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois_5_nfl$temp <- c(rep(0, 243), data_4yrs$cmean_4)</pre>
# Number of flowers column is only relevant for FFD, but is set to 0 for fitness values
data_4yrs$cn_fl<-scale(data_4yrs$n_fl,center=T,scale=F) # Standardize number of flowers
data.stack_pois_5_nfl$n_fl <- c(rep(0, 243), data_4yrs$cn_fl)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois_5_nfl$fitness.FFD.stack <- c(round(data_4yrs_total$mean_fitness), data_4yrs$FFD)
# Create 3 index columns needed for MCMCglmm
```

```
data.stack_pois_5_nfl$traits <- c(rep("fitness", 243), rep("FFD", 1455))</pre>
data.stack_pois_5_nfl$variable <- data.stack_pois_5_nfl$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois_5_nfl$family <- c(rep("poisson", 243), rep("gaussian", 1455))
data.stack_pois_5_nfl <- data.frame(data.stack_pois_5_nfl)</pre>
data.stack_pois_5_nfl$id <- as.factor(data.stack_pois_5_nfl$id)</pre>
data.stack_pois_5_nfl$year <- as.factor(data.stack_pois_5_nfl$year)</pre>
head(data.stack_pois_5_nfl)
     Obs id year temp n_fl fitness.FFD.stack traits variable family
## 1
      1 1 2006
                    0
                          0
                                            14 fitness fitness poisson
       2 2 2007
                          0
## 2
                     0
                                             4 fitness fitness poisson
                                             2 fitness fitness poisson
## 3
       3 3 2007
                          0
## 4
       4 4 2007
                    Ω
                          0
                                             6 fitness fitness poisson
## 5
       5 5 2007
                          0
                                             4 fitness fitness poisson
       6 6 2007
## 6
                     0
                                             2 fitness fitness poisson
modelBV_RR_pois_5_nfl <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                          \# ^ means for each variable (and no overall mean (hence "-1"))
                          at.level(variable, "FFD"):temp + # single fixed effect of temp
at.level(variable, "FFD"):n_f1, # single fixed effect of n_fl
                          random = ~us(at.level(variable, "FFD")):year +
                            us(at.level(variable, "FFD") +
                                 at.level(variable, "FFD"):temp):id,
                        # ^ random intercepts for individual,
                        # and random slopes for temp/id
                        rcov = ~us(at.level(variable, "fitness")):id +
                          # ^ variance between indivdiuals in fitness
                          # (which is residual variance)
                          us(at.level(variable, "FFD")):Obs,
                          # ^ residual variance within indivdiuals between years
                        # (labelled by 'Obs')
                        data = data.stack_pois_5_nfl,
                        prior = priorBiv_RR_pois,
                        family = NULL, # specified already in the data-set
                        nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
cor_BV_RR_pois_5_nfl_slopefit <-</pre>
  modelBV_RR_pois_5_nfl$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
  (sqrt(modelBV_RR_pois_5_nfl$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").
     sqrt(modelBV_RR_pois_5_nfl$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):t
plot(cor_BV_RR_pois_5_nfl_slopefit)
```



Density of var1



N = 1000 Bandwidth = 0.03917

```
posterior.mode(cor_BV_RR_pois_5_nfl_slopefit)
```

```
## var1
## 0.009830252
```

HPDinterval(cor_BV_RR_pois_5_nfl_slopefit)

```
## lower upper
## var1 -0.3534902 0.2128344
## attr(,"Probability")
## [1] 0.95
```

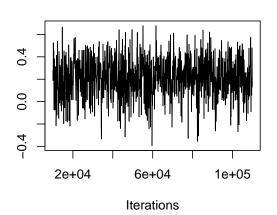
Temperature: mean May

Mean-centred mean May temperature:

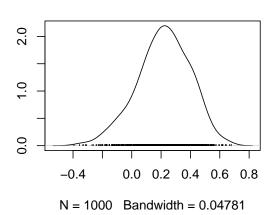
```
data_5yrs<-data_5yrs%>%
  mutate(cmean_5=scale(mean_5,center=T,scale=F))
```

```
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois_5_may$family <- c(rep("poisson", 156), rep("gaussian", 1107))
data.stack_pois_5_may <- data.frame(data.stack_pois_5_may)</pre>
data.stack_pois_5_may$id <- as.factor(data.stack_pois_5_may$id)</pre>
data.stack_pois_5_may$year <- as.factor(data.stack_pois_5_may$year)</pre>
head(data.stack_pois_5_may)
     Obs id year temp fitness.FFD.stack traits variable family
## 1
       1 1 2006
                                     14 fitness fitness poisson
## 2
       2 2 2007
                                      4 fitness fitness poisson
       3 3 2007
                                      2 fitness fitness poisson
## 3
                    0
       4 4 2007
## 4
                    0
                                      6 fitness fitness poisson
       5 5 2007
## 5
                    0
                                      4 fitness fitness poisson
                                      2 fitness fitness poisson
## 6 6 6 2007
                    0
modelBV_RR_pois_5_may <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois_5_may,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```

```
cor_BV_RR_pois_5_may_slopefit <-
modelBV_RR_pois_5_may$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
  (sqrt(modelBV_RR_pois_5_may$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").
        sqrt(modelBV_RR_pois_5_may$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):t
plot(cor_BV_RR_pois_5_may_slopefit)</pre>
```



Density of var1



```
posterior.mode(cor_BV_RR_pois_5_may_slopefit)

## var1
## 0.2204907

HPDinterval(cor_BV_RR_pois_5_may_slopefit)

## lower upper
## var1 -0.137329 0.5646195
## attr(,"Probability")
## [1] 0.95
```

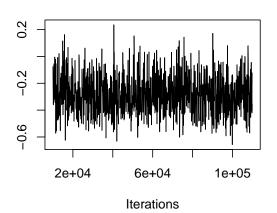
Mean fitness per flowering event

Temperature: mean April

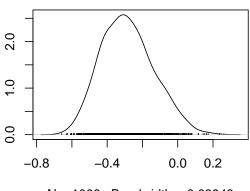
```
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack pois2_5$fitness.FFD.stack <- c(round(data_5yrs_total$mean_fitness_fl), data_5yrs$FFD)
# Create 3 index columns needed for MCMCqlmm
data.stack_pois2_5$traits <- c(rep("fitness", 156), rep("FFD", 1107))</pre>
data.stack_pois2_5$variable <- data.stack_pois2_5$traits</pre>
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois2_5$family <- c(rep("poisson", 156), rep("gaussian", 1107))
data.stack_pois2_5 <- data.frame(data.stack_pois2_5)</pre>
data.stack pois2 5$id <- as.factor(data.stack pois2 5$id)
data.stack_pois2_5$year <- as.factor(data.stack_pois2_5$year)</pre>
head(data.stack_pois2_5)
     Obs id year temp fitness.FFD.stack traits variable family
       1 1 2006
## 1
                                     16 fitness fitness poisson
## 2
       2 2 2007
                                      6 fitness fitness poisson
                    0
       3 3 2007
## 3
                    0
                                      3 fitness fitness poisson
       4 4 2007
## 4
                    0
                                      7 fitness fitness poisson
## 5
       5 5 2007
                    0
                                      4 fitness fitness poisson
## 6
       6 6 2007
                    0
                                      3 fitness fitness poisson
modelBV_RR_pois2_5 <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         \# ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack_pois2_5,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
cor_BV_RR_pois2_5_slopefit <-</pre>
  modelBV RR pois2 5$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
  (sqrt(modelBV_RR_pois2_5$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id"
```

plot(cor_BV_RR_pois2_5_slopefit)

sqrt(modelBV_RR_pois2_5\$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp



Density of var1



N = 1000 Bandwidth = 0.03949

```
posterior.mode(cor_BV_RR_pois2_5_slopefit)

## var1
## -0.3239812

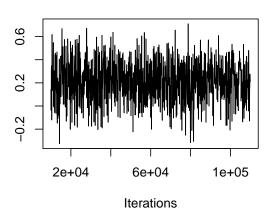
HPDinterval(cor_BV_RR_pois2_5_slopefit)

## lower upper
## var1 -0.5639433 -0.003528689
## attr(,"Probability")
## [1] 0.95
```

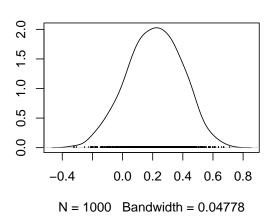
Temperature: mean May

```
# Create a single data-set "data.stack_pois2_5_may", with single column at start to index observations
data.stack_pois2_5_may <- c()</pre>
data.stack_pois2_5_may$0bs <- 1:(156 + 1107)</pre>
data.stack_pois2_5_may$id <- c(data_5yrs_total$id, data_5yrs$id)</pre>
# Year column is only relevant for FFD, but is set to first_yr for fitness values
data.stack_pois2_5_may$year <- c(data_5yrs_total_wfirstyr$first_yr,</pre>
                     data_5yrs$year)
# Temperature column is only relevant for FFD, but is set to 0 for fitness values
data.stack_pois2_5_may$temp <- c(rep(0, 156), data_5yrs$cmean_5)</pre>
# Create single column with first fitness values (ABSOLUTE VALUES), then FFD values:
data.stack_pois2_5_may$fitness.FFD.stack <- c(round(data_5yrs_total$mean_fitness_f1), data_5yrs$FFD)
# Create 3 index columns needed for MCMCglmm
data.stack_pois2_5_may$traits <- c(rep("fitness", 156), rep("FFD", 1107))
data.stack pois2 5 may variable <- data.stack pois2 5 may traits
# Fitness will be modelled with an overdispersed Poisson
# FFD will be modelled with a Gaussian distribution
# Specify this with the column 'family':
data.stack_pois2_5_may$family <- c(rep("poisson", 156), rep("gaussian", 1107))</pre>
```

```
data.stack_pois2_5_may <- data.frame(data.stack_pois2_5_may)</pre>
data.stack_pois2_5_may$id <- as.factor(data.stack_pois2_5_may$id)</pre>
data.stack_pois2_5_may$year <- as.factor(data.stack_pois2_5_may$year)</pre>
head(data.stack_pois2_5_may)
##
     Obs id year temp fitness.FFD.stack traits variable family
      1 1 2006
## 1
                    0
                                     16 fitness fitness poisson
## 2
       2 2 2007
                                      6 fitness fitness poisson
                    0
      3 3 2007
                                      3 fitness fitness poisson
## 3
                    0
## 4
      4 4 2007
                    Ω
                                      7 fitness fitness poisson
## 5
      5 5 2007
                    0
                                      4 fitness fitness poisson
      6 6 2007
## 6
                    0
                                      3 fitness fitness poisson
modelBV_RR_pois2_5_may <- MCMCglmm(fitness.FFD.stack ~ variable - 1 +</pre>
                         # ^ means for each variable (and no overall mean (hence "-1"))
                         at.level(variable, "FFD"):temp, # single fixed effect of temp
                       random = ~us(at.level(variable, "FFD")):year +
                         us(at.level(variable, "FFD") +
                              at.level(variable, "FFD"):temp):id,
                       # ^ random intercepts for individual,
                       # and random slopes for temp/id
                       rcov = ~us(at.level(variable, "fitness")):id +
                         # ^ variance between indivdiuals in fitness
                         # (which is residual variance)
                         us(at.level(variable, "FFD")):Obs,
                         # ^ residual variance within indivdiuals between years
                       # (labelled by 'Obs')
                       data = data.stack pois2 5 may,
                       prior = priorBiv_RR_pois,
                       family = NULL, # specified already in the data-set
                       nitt = 1100 * sc, thin = sc, burnin = 100 * sc, verbose = F)
```



Density of var1



```
posterior.mode(cor_BV_RR_pois2_5_may_slopefit)

## var1
## 0.2186931

HPDinterval(cor_BV_RR_pois2_5_may_slopefit)

## lower upper
## var1 -0.1415346 0.5413856
## attr(,"Probability")
## [1] 0.95
```

Summary 5 years

Among-individual correlation between fitness and variation in slopes for FFD for each model:

```
cor_slopefit_5<-rbind(cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_5_slopefit),</pre>
                          rbind(HPDinterval(cor_BV_RR_pois_5_slopefit)[,1:2])),
                      cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_5_nfl_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois_5_nfl_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois_5_may_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois_5_may_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois2_5_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois2_5_slopefit)[,1:2])),
                    cbind(posterior.mode=posterior.mode(cor_BV_RR_pois2_5_may_slopefit),
                          rbind(HPDinterval(cor_BV_RR_pois2_5_may_slopefit)[,1:2])))
row.names(cor_slopefit_5)<-c("Total fitness, mean April temperature, without n_fl",
                             "Total fitness, mean April temperature, with n_fl",
                           "Total fitness, mean May temperature",
                           "Mean fitness per flowering event, mean April temperature",
                           "Mean fitness per flowering event, mean May temperature")
kable(cor_slopefit_5,digits=3)
```

	posterior.mode	lower	upper
Total fitness, mean April temperature, without n_fl	-0.356	-0.589	-0.017
Total fitness, mean April temperature, with n_fl	0.010	-0.353	0.213

	posterior.mode	lower	upper
Total fitness, mean May temperature	0.220	-0.137	0.565
Mean fitness per flowering event, mean April temperature	-0.324	-0.564	-0.004
Mean fitness per flowering event, mean May temperature	0.219	-0.142	0.541

Only model with selection on RN slope: modelBV_RR_pois_5

kable(summary(modelBV_RR_pois_5)\$solutions,digits=c(3,3,3,0,3),caption="Fixed effects")

Table 7: Fixed effects

	post.mean	l-95% CI	u-95% CI	eff.samp	pMCMC
variableFFD variablefitness at.level(variable, "FFD"):temp	57.494	55.160	59.607	1000	0.001
	1.211	1.067	1.345	946	0.001
	-2.398	-3.816	-0.733	1147	0.002

kable(summary(modelBV_RR_pois_5)\$Gcovariances,digits=c(3,3,3,0),caption="Random effects")

Table 8: Random effects

	post.mean	l-95% CI	u-95% CI	eff.samp
at.level(variable, "FFD"):at.level(variable, "FFD").year	25.802	11.499	43.407	1127

kable(summary(modelBV_RR_pois_5)\$Rcovariances,digits=c(3,3,3,0),caption="Random effects")

Table 9: Random effects

	post.mean	l-95% CI	u-95% CI	eff.samp
at.level(variable, "FFD").id:at.level(variable, "FFD").id	3.405	2.111	4.676	1000
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD").id	1.041	0.541	1.723	1000
at.level(variable, "fitness").id:at.level(variable, "FFD").id	-0.620	-0.993	-0.305	1000
at.level(variable, "FFD").id:at.level(variable, "FFD"):temp.id	1.041	0.541	1.723	1000
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD"):temp.id	0.776	0.350	1.226	1000
at.level(variable, "fitness").id:at.level(variable, "FFD"):temp.id	-0.183	-0.387	-0.005	1000
at.level(variable, "FFD").id:at.level(variable, "fitness").id	-0.620	-0.993	-0.305	1000
at.level(variable, "FFD"):temp.id:at.level(variable, "fitness").id	-0.183	-0.387	-0.005	1000
at.level(variable, "fitness").id:at.level(variable, "fitness").id	0.482	0.327	0.653	959
at.level (variable, ``FFD") : at.level (variable, ``FFD"). Obs	18.740	16.838	20.345	1000

kable(diag(autocorr(modelBV_RR_pois_5\$VCV)[2, ,]),caption="Autocorrelation")

Table 10: Autocorrelation

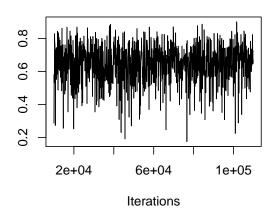
	x
at.level(variable, "FFD"):at.level(variable, "FFD").year	0.0310838
at.level(variable, "FFD").id:at.level(variable, "FFD").id	-0.0244161
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD").id	-0.0331450

	X
at.level(variable, "fitness").id:at.level(variable, "FFD").id	0.0224512
at.level(variable, "FFD").id:at.level(variable, "FFD"):temp.id	-0.0331450
at.level(variable, "FFD"):temp.id:at.level(variable, "FFD"):temp.id	-0.0264357
at.level(variable, "fitness").id:at.level(variable, "FFD"):temp.id	-0.0048342
at.level(variable, "FFD").id:at.level(variable, "fitness").id	0.0224512
at.level(variable, "FFD"):temp.id:at.level(variable, "fitness").id	-0.0048342
at.level(variable, "fitness").id:at.level(variable, "fitness").id	0.0433784
at.level(variable, "FFD"):at.level(variable, "FFD").Obs	0.0287729

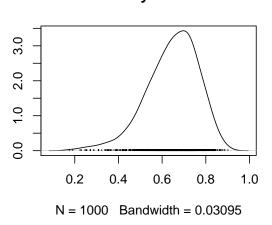
Among-individual correlation between intercepts and slopes for FFD:

```
cor_BV_RR_pois_5_intslope <-
   modelBV_RR_pois_5$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\").id"]/
(sqrt(modelBV_RR_pois_5$VCV[,"at.level(variable, \"FFD\").id:at.level(variable, \"FFD\").id"])*
sqrt(modelBV_RR_pois_5$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.id"])
plot(cor_BV_RR_pois_5_intslope)</pre>
```

Trace of var1



Density of var1



```
posterior.mode(cor_BV_RR_pois_5_intslope)
```

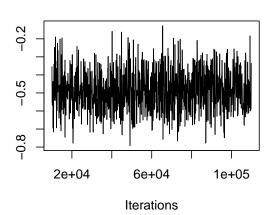
```
## var1
## 0.6885208
```

HPDinterval(cor_BV_RR_pois_5_intslope)

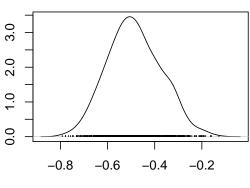
```
## lower upper
## var1 0.4102019 0.8531525
## attr(,"Probability")
## [1] 0.95
```

Among-individual correlation between FFD and fitness:

```
cor_BV_RR_pois_5_intfit <-
modelBV_RR_pois_5$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\").id"]/
   (sqrt(modelBV_RR_pois_5$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id"]
        sqrt(modelBV_RR_pois_5$VCV[,"at.level(variable, \"FFD\").id:at.level(variable, \"FFD\").id"]))
plot(cor_BV_RR_pois_5_intfit)</pre>
```



Density of var1



N = 1000 Bandwidth = 0.03019

posterior.mode(cor_BV_RR_pois_5_intfit)

var1 ## -0.4551948

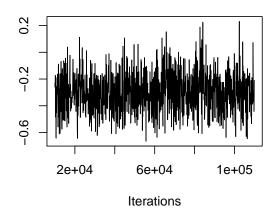
HPDinterval(cor_BV_RR_pois_5_intfit)

lower upper
var1 -0.7081592 -0.2775473
attr(,"Probability")
[1] 0.95

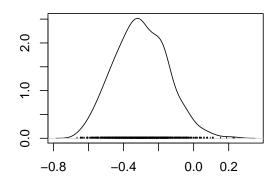
Among-individual correlation between fitness and variation in slopes for FFD:

cor_BV_RR_pois_5_slopefit <modelBV_RR_pois_5\$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"FFD\"):temp.id"]/
 (sqrt(modelBV_RR_pois_5\$VCV[,"at.level(variable, \"fitness\").id:at.level(variable, \"fitness\").id"]
 sqrt(modelBV_RR_pois_5\$VCV[,"at.level(variable, \"FFD\"):temp.id:at.level(variable, \"FFD\"):temp.
plot(cor_BV_RR_pois_5_slopefit)</pre>

Trace of var1



Density of var1



N = 1000 Bandwidth = 0.04008

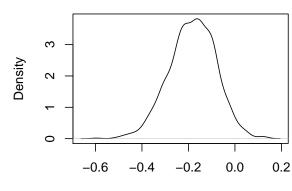
```
posterior.mode(cor_BV_RR_pois_5_slopefit)
         var1
## -0.3563453
HPDinterval(cor_BV_RR_pois_5_slopefit)
##
             lower
                          upper
## var1 -0.5893757 -0.01694865
## attr(,"Probability")
## [1] 0.95
Extract selection coefficients
# Extract 3x3 matrix of variance-covariance values for intercepts and slopes of X, and LBS
# These are in the 2nd-10th columns of model output
P.modelBV_RR_pois_5 <- modelBV_RR_pois_5$VCV[,2:10]</pre>
P.modelBV_RR_pois_5.mode <- matrix(1:9, nrow = 3)
for (k in 1:9) P.modelBV_RR_pois_5.mode[k] <- posterior.mode(P.modelBV_RR_pois_5[,k])
P.modelBV_RR_pois_5.mode
##
                          [,2]
                                      [,3]
               [,1]
## [1,] 3.5003523 1.0220040 -0.6122982
## [2,] 1.0220040 0.6412418 -0.1740813
## [3,] -0.6122982 -0.1740813 0.4853696
# Extract selection *differentials* (i.e. covariances) for intercept and slope:
# and calculate posterior mode and credible intervals for each
S.modelBV RR pois 5 \leftarrow \text{modelBV} RR pois 5\$\text{VCV}[, c(4,7)]
S.modelBV_RR_pois_5 <- P.modelBV_RR_pois_5[, c(3,6)]</pre>
colnames(S.modelBV_RR_pois_5) <- c("S_intercepts", "S_slopes")</pre>
S.modelBV_RR_pois_5.mode <- P.modelBV_RR_pois_5.mode[1:2, 3]</pre>
S.modelBV_RR_pois_5.mode
## [1] -0.6122982 -0.1740813
posterior.mode(mcmc(S.modelBV RR pois 5))
## S_intercepts
                     S_slopes
## -0.6122982
                  -0.1740813
HPDinterval(mcmc(S.modelBV_RR_pois_5))
##
                      lower
                                   upper
## S_intercepts -0.9929710 -0.305195789
## S_slopes
                -0.3867651 -0.005488248
## attr(,"Probability")
## [1] 0.95
# Plot posterior distribution of selection differentials
par(mfrow = c(1,2))
plot(density(S.modelBV_RR_pois_5[,1]), main = "S_intercepts")
plot(density(S.modelBV_RR_pois_5[,2]), main = "S_slopes")
```

S_intercepts

Oensity -1.4 -1.0 -0.6 -0.2

N = 1000 Bandwidth = 0.04132

S_slopes



N = 1000 Bandwidth = 0.02273

```
# Estimate selection gradients for intercept and slope (beta = S / P)
# on each sample of posterior and extract their mode
n <- length(modelBV_RR_pois_5$VCV[,2])</pre>
                                          # sample size
beta_post_RR_pois_5 <- matrix(NA, n ,2)</pre>
for (i in 1:n) {
 P3 <- matrix(rep(NA, 9), nrow = 3) # 3x3 matrix of var-cov for individual X.int, X.slope and LBS
  for (k in 1:9) {P3[k] <- P.modelBV_RR_pois_5[i, k] }</pre>
  P2 <- P3[1:2, 1:2]
                      # 2x2 matrix of just trait intercept & slope var-cov
  S \leftarrow P3[1:2, 3]
                  # selection differentials on traits (last column of P3)
  beta_post_RR_pois_5[i,] <- solve(P2) %*% S
                                               # selection gradients beta = P^-1 * S
}
# Finally, extract and plot the selection gradients posterior modes
# and 95% credible intervals for both selection on intercepts (trait value)
# and slopes (trait plasticity).
# Note that credible intervals are not exactly confidence intervals. See here:
# https://statsdirect.com/help/basics/confidence_interval.htm and
# https://stats.stackexchange.com/questions/2272/
colnames(beta_post_RR_pois_5) <- c("beta_intercepts", "beta_slopes")</pre>
posterior.mode(mcmc(beta post RR pois 5))
## beta_intercepts
                       beta_slopes
                        0.06080902
##
       -0.17559552
HPDinterval(mcmc(beta_post_RR_pois_5))
##
                        lower
                                     upper
## beta_intercepts -0.3499730 -0.03235167
                   -0.3508229 0.40774204
## beta_slopes
## attr(,"Probability")
## [1] 0.95
# Plot posterior distribution of selection gradients
par(mfrow = c(1,2))
plot(density(beta_post_RR_pois_5[,1]), main = "beta_intercepts")
plot(density(beta_post_RR_pois_5[,2]), main = "beta_slopes")
```

beta_intercepts beta_slopes 2 Density က 1.0 2 0.0 0 -0.5-0.3-0.10.1 -0.50.0 0.5 N = 1000 Bandwidth = 0.0183 N = 1000 Bandwidth = 0.04137

NB selection differentials and gradients here are from covariances with latent-scale absolute fitness # These are equivalent to covariances with data-scale relative fitness: see main text of paper