

# Single-celled bottlenecks, germlines and the evolution of complex multi-cellularity

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## Pre-amble

### load packages

```
library(ape)
library(ggplot2)
library(tidyverse)

## -- Attaching packages ----- tidyverse 1.3.0 --

## v tibble   3.0.4      v dplyr    1.0.2
## v tidyr    1.1.2      v stringr  1.4.0
## v readr    1.4.0      vforcats  0.5.0
## v purrr   0.3.4

## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()   masks stats::lag()

library(knitr)
library(brms) #https://rdrr.io/cran;brms/f/vignettes;brms_phylogenetics.Rmd
```

```
## Loading required package: Rcpp
```

```

## Loading 'brms' package (version 2.15.0). Useful instructions
## can be found by typing help('brms'). A more detailed introduction
## to the package is available through vignette('brms_overview').

##
## Attaching package: 'brms'

## The following object is masked from 'package:stats':
##
##     ar

library(tidybayes)

##
## Attaching package: 'tidybayes'

## The following objects are masked from 'package:brms':
##
##     dstudent_t, pstudent_t, qstudent_t, rstudent_t

library(rotl) #see https://cran.r-project.org/web/packages/rotl/vignettes/rotl.html

```

## read in data

```
df<- read.csv('data/germline_data_1.1.csv')
```

## create phylogeny files

```

ResolvedNames <- tnr_s_match_names(df$species.updated.rotl, context_name = 'All life') #search for simil
ResolvedNames$IsInTree <- is_in_tree(ResolvedNames$ott_id) #T/F, did the above find a match that can be
ResolvedNamesInTree<- subset(ResolvedNames, IsInTree==T) #subset to only those where present in the phy
AllTree<- tol_induced_subtree(ResolvedNamesInTree$ott_id, label_format = 'id') #draw the phylogeny

## Warning in collapse_singles(tr, show_progress): Dropping singleton nodes
## with labels: mrcaott2ott142555, mrcaott2ott7623, ott916750, mrcaott2ott50189,
## mrcaott2ott108668, mrcaott2ott59852, mrcaott2ott8171, ott10210, ott99252,
## mrcaott2ott2645, mrcaott2ott35778, ott5298374, mrcaott2ott10930,
## mrcaott2ott2441, mrcaott2ott969, mrcaott2ott62529, mrcaott2ott8379,
## ott431495, ott853757, ott5316182, mrcaott248ott10053, mrcaott248ott20991,
## mrcaott248ott557, mrcaott557ott67236, ott216628, mrcaott557ott717698,
## mrcaott557ott864011, mrcaott557ott37775, mrcaott557ott904, mrcaott904ott121240,
## mrcaott904ott264912, mrcaott904ott8870, mrcaott904ott52717, ott33109,
## mrcaott904ott63159, mrcaott904ott96612, mrcaott904ott31366, mrcaott904ott9799,
## ott584111, mrcaott904ott5005, ott227063, ott801070, ott226780, ott1058517,
## ott5308424, ott225270, mrcaott252ott213153, ott921871, mrcaott252ott128594,
## mrcaott252ott1477, mrcaott1477ott591692, mrcaott1477ott2066, ott105574,

```

```

## ott423248, ott423246, ott215123, ott568878, mrcaott334ott335, ott852744,
## ott857751, mrcaott3355ott25934, mrcaott3355ott3511, mrcaott3355ott55292,
## mrcaott3355ott4623, mrcaott3355ott579768, mrcaott3355ott3515,
## mrcaott3355ott3520, mrcaott3520ott4662, mrcaott4662ott12148, ott361626,
## mrcaott4662ott285937, mrcaott4662ott913602, ott792012, mrcaott1439ott109938,
## mrcaott1439ott12986, ott695980, ott643238, mrcaott4474ott13510,
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## mrcaott338994ott431388, mrcaott66494ott407568, mrcaott407568ott432842,
## mrcaott407568ott1059896, mrcaott407568ott1059891, mrcaott1059891ott1059892,
## mrcaott1059891ott1059898, mrcaott1059895ott1059900, mrcaott5202ott159280,
## mrcaott5202ott30666, mrcaott5202ott40102, ott1058522, ott5296507,
## mrcaott237ott8444, ott4736806, ott994067, mrcaott8444ott9094,
## mrcaott8444ott62995, mrcaott8444ott8454, ott771683, mrcaott61771ott212475,
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## ott649199, mrcaott649193ott740531, ott657906, ott388859, ott910940,

```

```

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## mrcaott392ott3549, mrcaott3549ott17097, mrcaott3549ott7508, mrcaott3549ott5050,
## mrcaott3549ott6406, mrcaott3549ott143050, ott1032209, ott223661,
## mrcaott49ott6612, ott555379, ott257330, ott580673, ott1096612, ott3665427,
## ott1025977, ott335258, ott335259, ott910030, ott242963, ott576383, ott675306,
## ott1037807, ott987479, ott471706, ott5673589, ott251966, ott471705, ott492249,
## ott5673590, ott5677241, ott2942239, ott2942245, ott29723, ott750565, ott150273,
## ott176166, ott2844994, mrcaott

# tree with resolved polytomies:
ResolvedPolytomiesTree<- multi2di(AllTree)

# write to files
write.tree(AllTree, file='data/phylogeny_all.txt') #phylogeny
write.tree(AllTree, file='data/phylogeny_all_res_polytomy.txt') #phylogeny with resolved polytomies

```

```

write.csv(ResolvedNames, 'data/phylogeny_species_names.csv') #list of names
write.csv(ResolvedNamesInTree, 'data/phylogeny_species_names_in_tree.csv') #list of names that are pres

```

Preparing the phylogeny data

```

#read in phylogeny
phylo<- ape::read.tree('data/phylogeny_all.txt')
#subset df above with just those that are in the tree (note some missing)
names_in_tree<- read.csv('data/phylogeny_species_names_in_tree.csv')
names_in_tree$ott_id<- paste('ott',names_in_tree$ott_id, sep = '')
#phylo$tip.label %in% names_in_tree$ott_id #to check whether they're all present
names_in_tree<- subset(names_in_tree, names_in_tree$ott_id %in% phylo$tip.label)
#which species in the table are in the tree
df$species.updated.rotl<- tolower(df$species.updated.rotl)
df<- subset(df, tolower(df$species.updated.rotl) %in% tolower(names_in_tree$search_string))
#give them the ids in a column so that brms can match it up
df$species_id<- names_in_tree$ott_id

```

Covariance matrix produced using branch lengths of 1 (as in Fisher et al)

```

# set branch lengths to 1 for covariance matrix
phylo_1b <- compute.brlen(phylo, 1)
#create covariance matrix
CovarMatrix <- ape::vcv.phylo(phylo_1b)

```

```

#plot phylogeny to check
plot(AllTree, no.margin = TRUE, cex = 0.5, label.offset = 0.5)

```

## data tidying

Change the fission or budding observed column to ‘yes’ vs ‘no’ to make sure it is not doing some weird numeric thing

```

df$FissionOrBuddingObserved_Genus_nominal <- ifelse(df$FissionOrBuddingObserved_Genus == 1, 'yes','no')

early = c('1','1,2','2')

df$germline_timing_simple<- ifelse(df$germline_timing %in% early, 'early', df$germline_timing)
df$germline_timing_simple<- ifelse(df$germline_timing == '0', 'no_germline', df$germline_timing_simple)
df$germline_timing_simple<- ifelse(df$germline_timing == '3', 'adult', df$germline_timing_simple)

```

## Analyses

Conducting Bayesian analyses using BRMS, first without phylogeny included, then including phylogeny. Each analysis contains the code that defines the model, and briefly analyses them— producing summary stats and figs.

See below for analyses of:

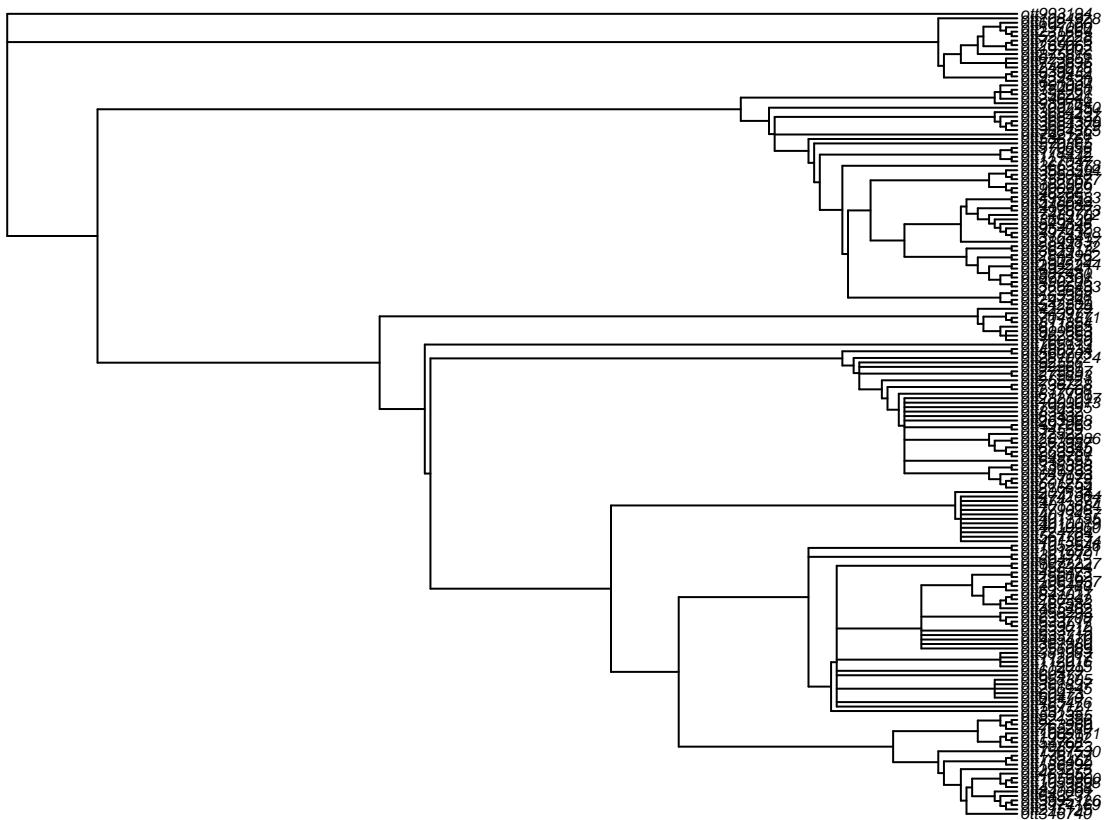


Figure 1: Simple phylogenetic tree constructed using the Open Tree of Life, needs manually checked for errors

Each analysis uses vague priors, 5 chains, 6,000,000 iterations, 100000 of which are discarded as warm-up, thinned by a factor of 1000.

(Currently use reproductive traits of genus rather than species)

- Phylogenetically naive:
  - Does a single-celled bottleneck correlate with increased cell number? (priors = normal dist, mean of 0, sd of 10)
  - Does a single-celled bottleneck correlate with increased cell types (per cell)? prior(normal(0, 10), "b")
  - Does germline timing correlate with increased cell number? prior = prior(normal(0, 10), "b")
  - Does germline timing correlate with increased cell types (per cell)?
- Phylogenetically informed:
  - Does a single-celled bottleneck correlate with increased cell number?
  - Does a single-celled bottleneck correlate with increased cell types (per cell)?
  - Does germline timing correlate with increased cell number?
  - Does germline timing correlate with increased cell types (per cell)?

To do: do any of these things need scaled?

## Phylogenetically naive

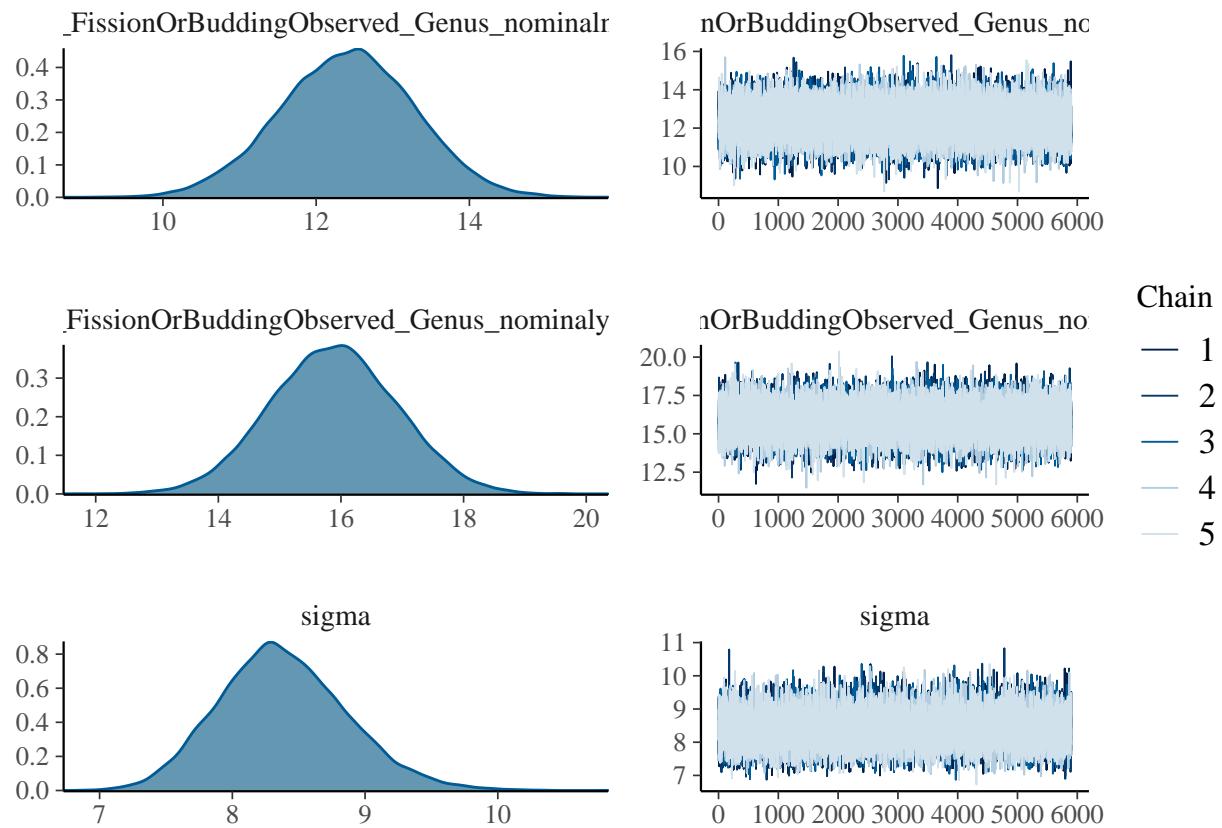
### Does a single-celled bottleneck correlate with increased cell number?

Using simple normally-distributed, relatively non-informative prior

One possibility is that the 0s and 1s are treated as numeric, when they should be categorical, so change to yes/no.

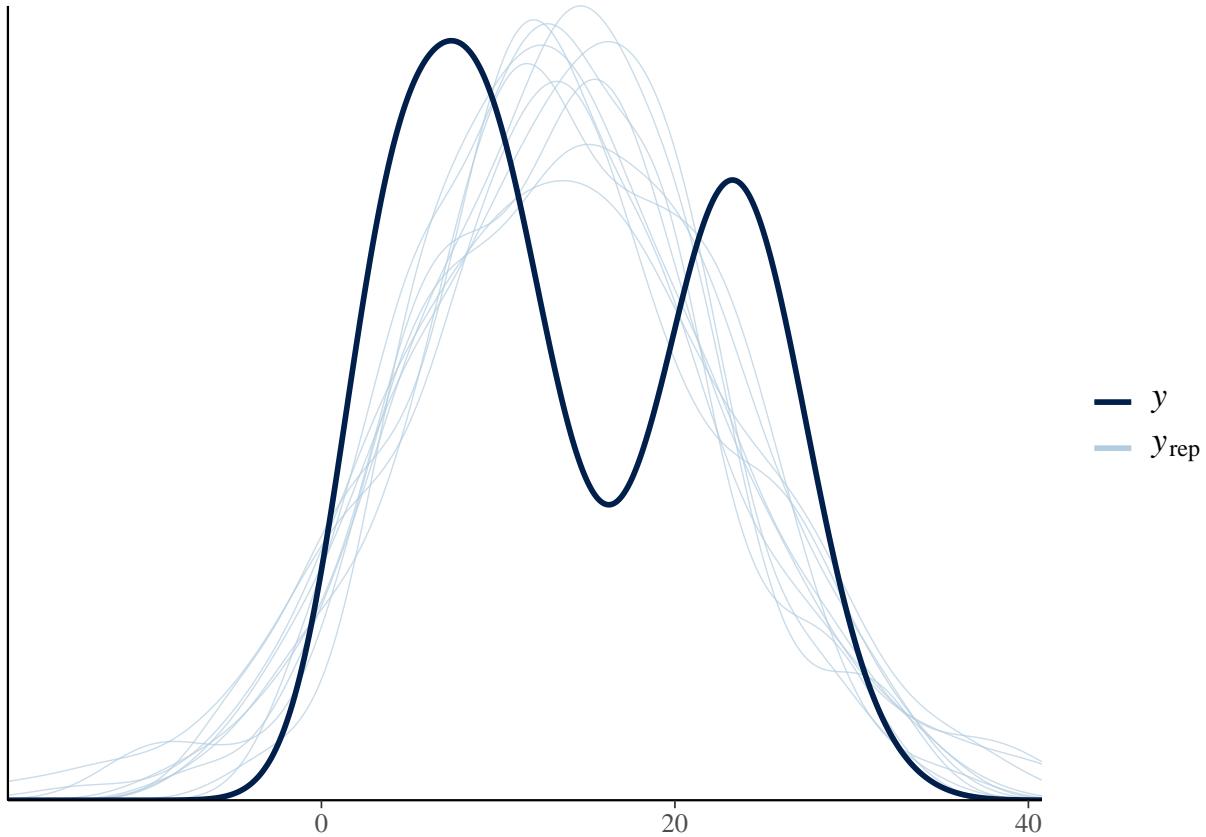
```
#fit the model
fit_fission_cell_num<-
  brm(data = df,
       family=gaussian(), # family of model
       formula = log(cell_number) ~ 0 + FissionOrBuddingObserved_Genus_nominal, #formula, the 0 means t
       iter = 6000000, warmup = 100000, chains = 5,thin = 1000, cores = 5, #chain settings
       prior = prior(normal(0, 10), "b"), #defining the priors- for things in 'class b' set this prior.
       file = 'fits/fit_FissionCellNumber' )

##### Assessing model:
plot(fit_fission_cell_num) #check that chains converged
```



```
pp_check(fit_fission_cell_num) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



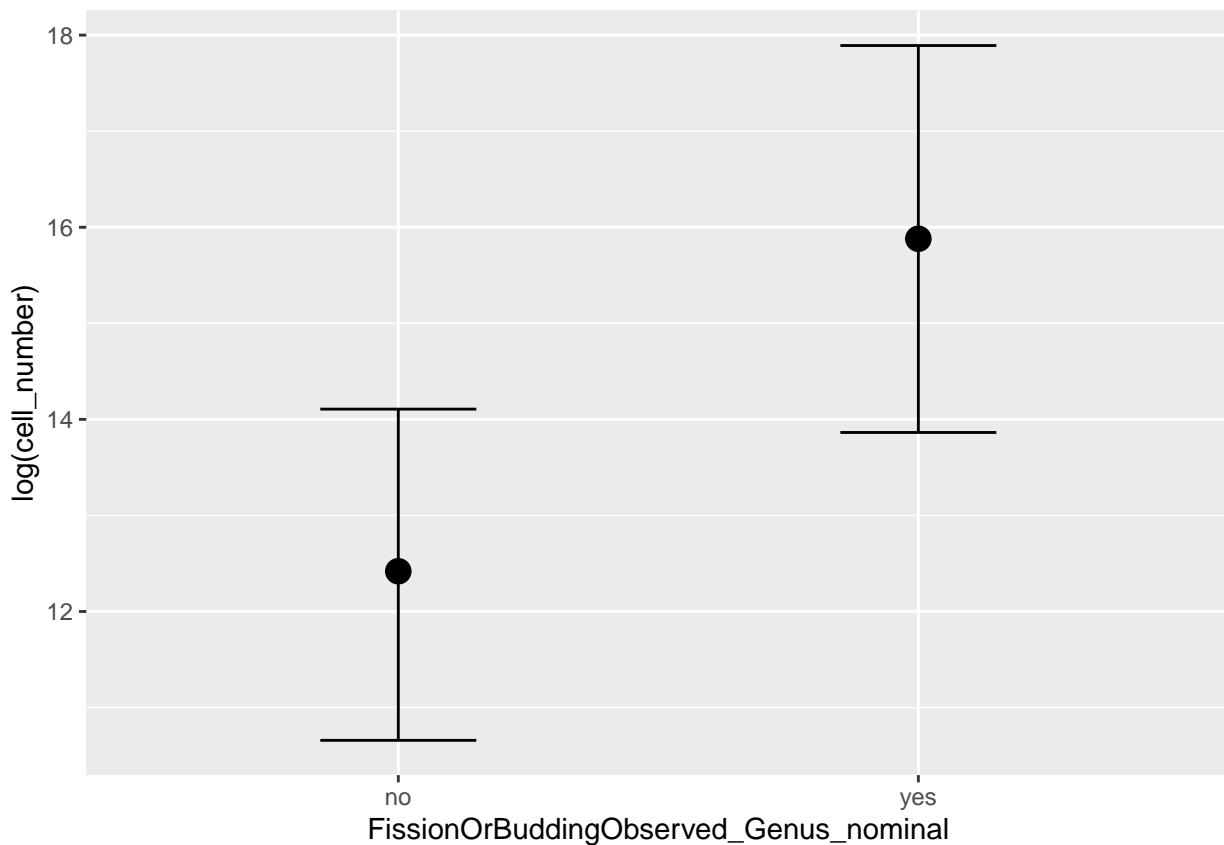
```
summary(fit_fission_cell_num) #summary of model
```

```
## Family: gaussian
## Links: mu = identity; sigma = identity
## Formula: log(cell_number) ~ 0 + FissionOrBuddingObserved_Genus_nominal
## Data: df (Number of observations: 155)
## Samples: 5 chains, each with iter = 6e+06; warmup = 1e+05; thin = 1000;
##          total post-warmup samples = 29500
##
## Population-Level Effects:
##                               Estimate Est.Error 1-95% CI u-95% CI
## FissionOrBuddingObserved_Genus_nominalno    12.41     0.88   10.66   14.11
## FissionOrBuddingObserved_Genus_nominalyes   15.88     1.03   13.86   17.89
##                                         Rhat Bulk_ESS Tail_ESS
## FissionOrBuddingObserved_Genus_nominalno  1.00   29496   29547
## FissionOrBuddingObserved_Genus_nominalyes 1.00   29056   28985
##
## Family Specific Parameters:
##           Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sigma      8.38     0.48    7.50    9.40 1.00    29919   29801
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

```
posterior_summary(fit_fission_cell_num, robust = T)
```

```
##                                     Estimate Est.Error      Q2.5
## b_FissionOrBudding0bserved_Genus_nominalno    12.418883 0.8870628 10.658512
## b_FissionOrBudding0bserved_Genus_nominalyes   15.879885 1.0370737 13.863384
## sigma                                         8.356744 0.4742410  7.501903
## lp__                                         -558.203857 1.0075698 -561.736062
##                                     Q97.5
## b_FissionOrBudding0bserved_Genus_nominalno    14.106579
## b_FissionOrBudding0bserved_Genus_nominalyes   17.891522
## sigma                                         9.403149
## lp__                                         -557.106225
```

```
plot(conditional_effects(fit_fission_cell_num, points = TRUE, ask = F))
```



```
hyp = hypothesis(fit_fission_cell_num, "FissionOrBudding0bserved_Genus_nominalno > FissionOrBudding0bserved_Genus_nominalyes")
```

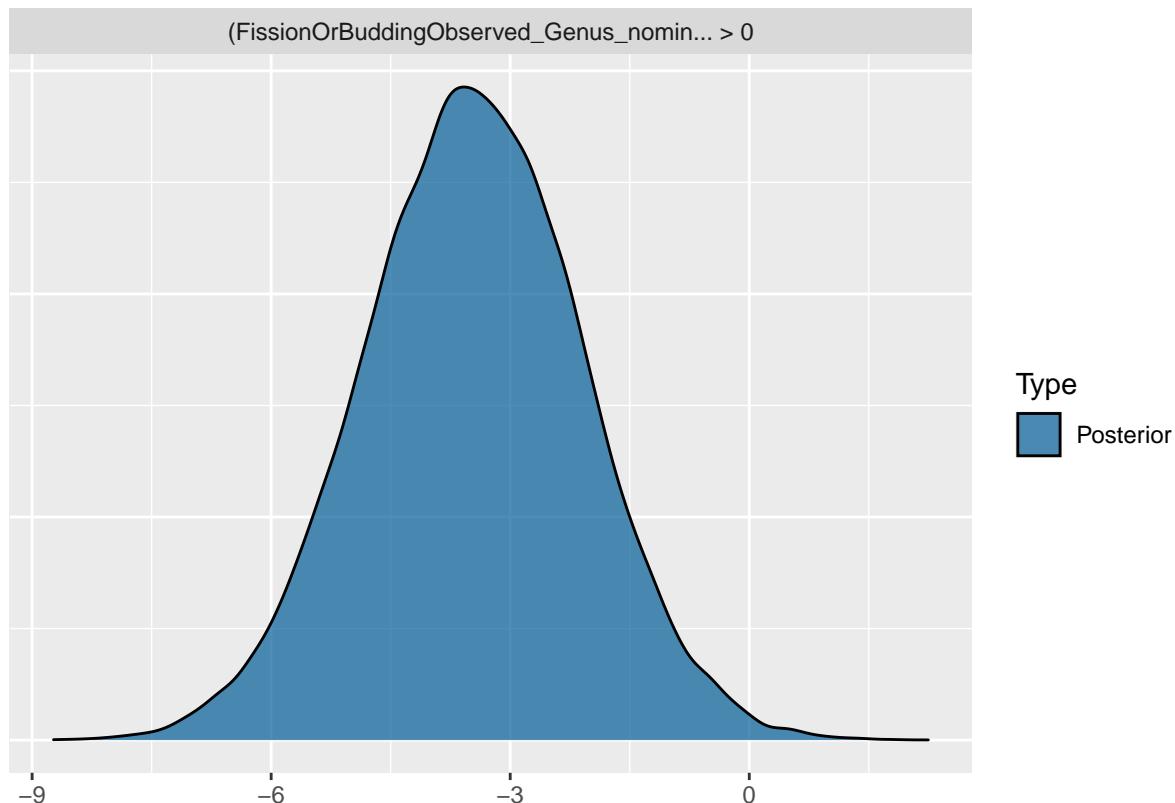
```
## Hypothesis Tests for class b:
##          Hypothesis Estimate Est.Error CI.Lower CI.Upper Evid.Ratio
## 1 (FissionOrBudding0bserved_Genus_nominalno > FissionOrBudding0bserved_Genus_nominalyes)  -3.47     1.36    -5.72    -1.24      0.01
## Post.Prob Star
## 1          0.01
## ---
```

```

## 'CI': 90%-CI for one-sided and 95%-CI for two-sided hypotheses.
## '*': For one-sided hypotheses, the posterior probability exceeds 95%;
## for two-sided hypotheses, the value tested against lies outside the 95%-CI.
## Posterior probabilities of point hypotheses assume equal prior probabilities.

```

```
plot(hyp)
```



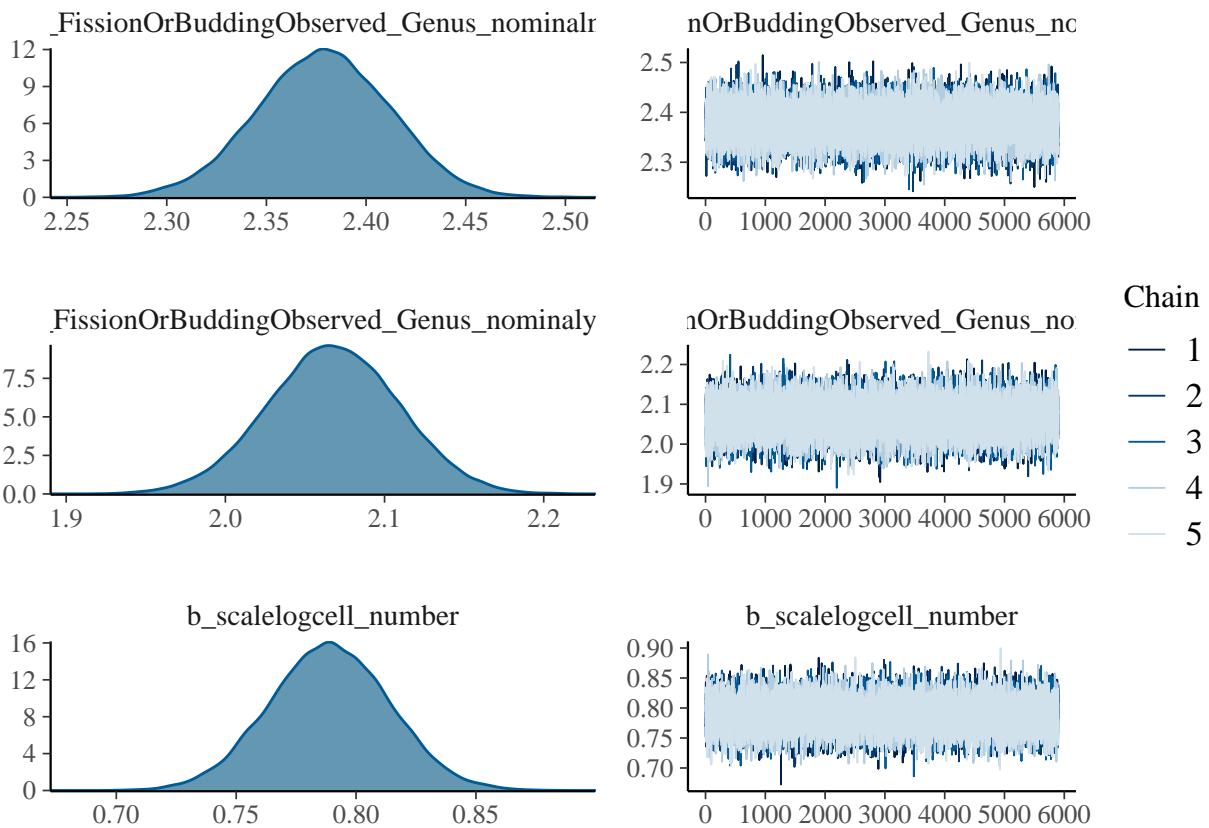
Does a single-celled bottleneck correlate with increased cell types (per cell)?

```

#fit the model
fit_fission_cell_type<-
  brm(data = df,
       family=poisson(),
       formula = cell_types ~ 0 + FissionOrBuddingObserved_Genus_nominal + scale(log(cell_number)), #for
       iter = 6000000, warmup = 100000, chains = 5, thin = 1000, cores = 5, #chain settings
       prior = prior(normal(0, 10), "b"), #defining the priors- for things in 'class b' set this prior.
       file = 'fits/fit_FissionCellType' )

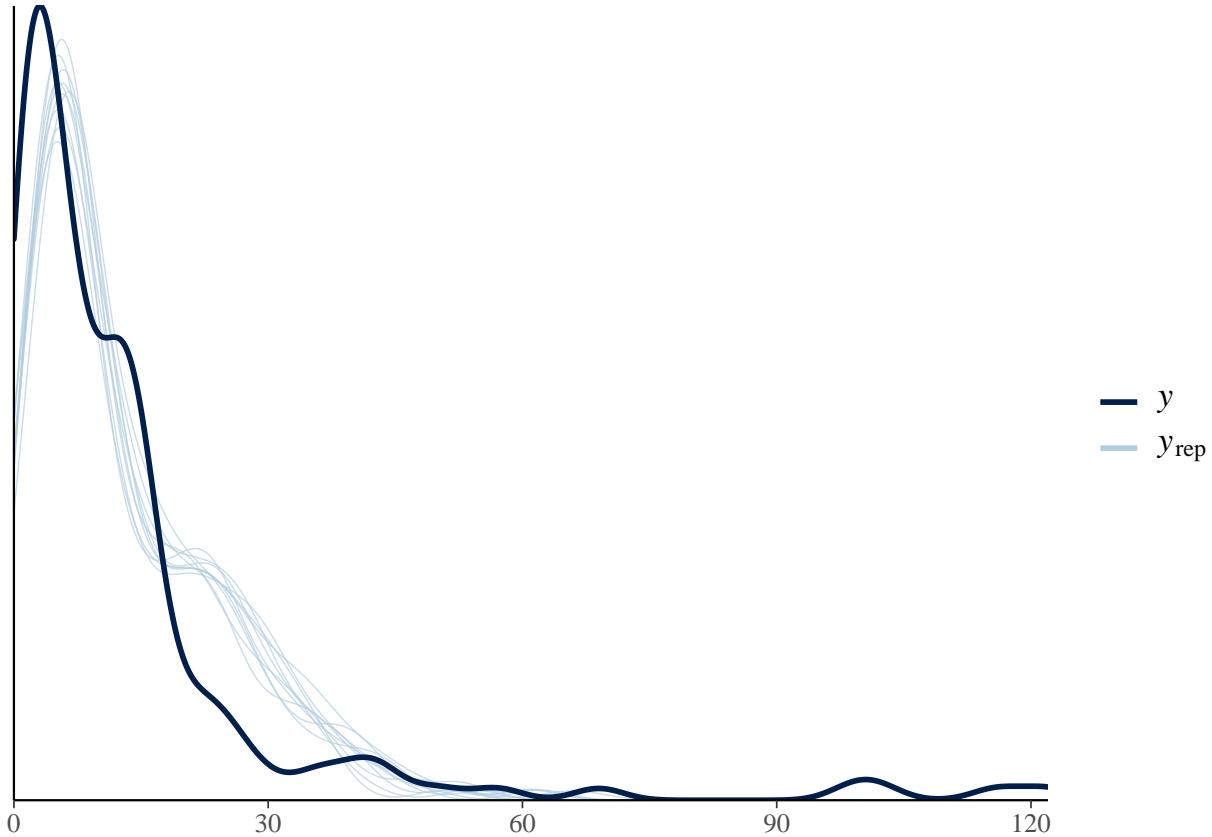
plot(fit_fission_cell_type) #check that chains converged

```



```
pp_check(fit_fission_cell_type) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



```
summary(fit_fission_cell_type) #summary of model
```

```
## Family: poisson
## Links: mu = log
## Formula: cell_types ~ 0 + FissionOrBudding0bserved_Genus_nominal + scale(log(cell_number))
## Data: df (Number of observations: 154)
## Samples: 5 chains, each with iter = 6e+06; warmup = 1e+05; thin = 1000;
##          total post-warmup samples = 29500
##
## Population-Level Effects:
##                               Estimate   Est.Error 1-95% CI u-95% CI
## FissionOrBudding0bserved_Genus_nominalno    2.38      0.03    2.31    2.44
## FissionOrBudding0bserved_Genus_nominalyes   2.07      0.04    1.99    2.15
## scalelogcell_number                         0.79      0.03    0.74    0.84
##                                         Rhat Bulk_ESS Tail_ESS
## FissionOrBudding0bserved_Genus_nominalno 1.00   29863    28122
## FissionOrBudding0bserved_Genus_nominalyes 1.00   29051    29217
## scalelogcell_number                      1.00   29251    29053
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

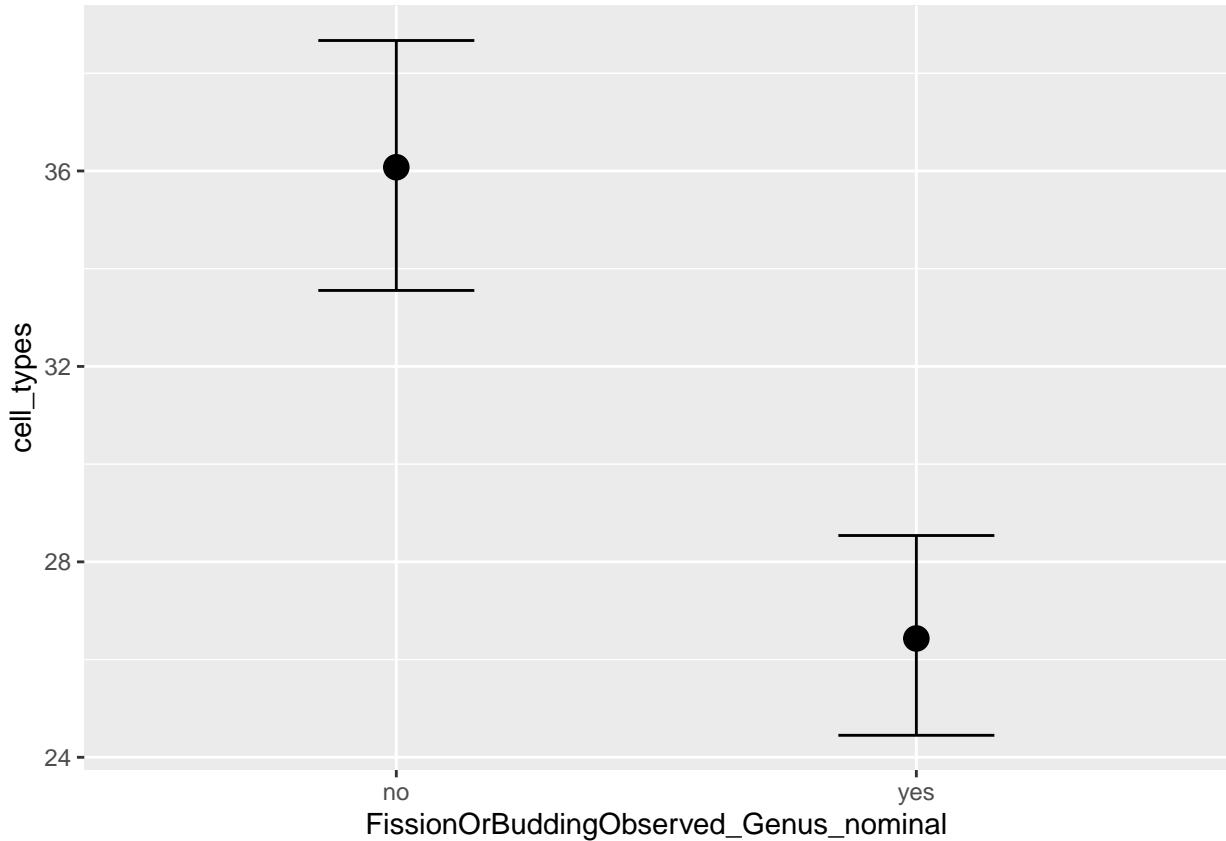
```
posterior_summary(fit_fission_cell_type, robust = T)
```

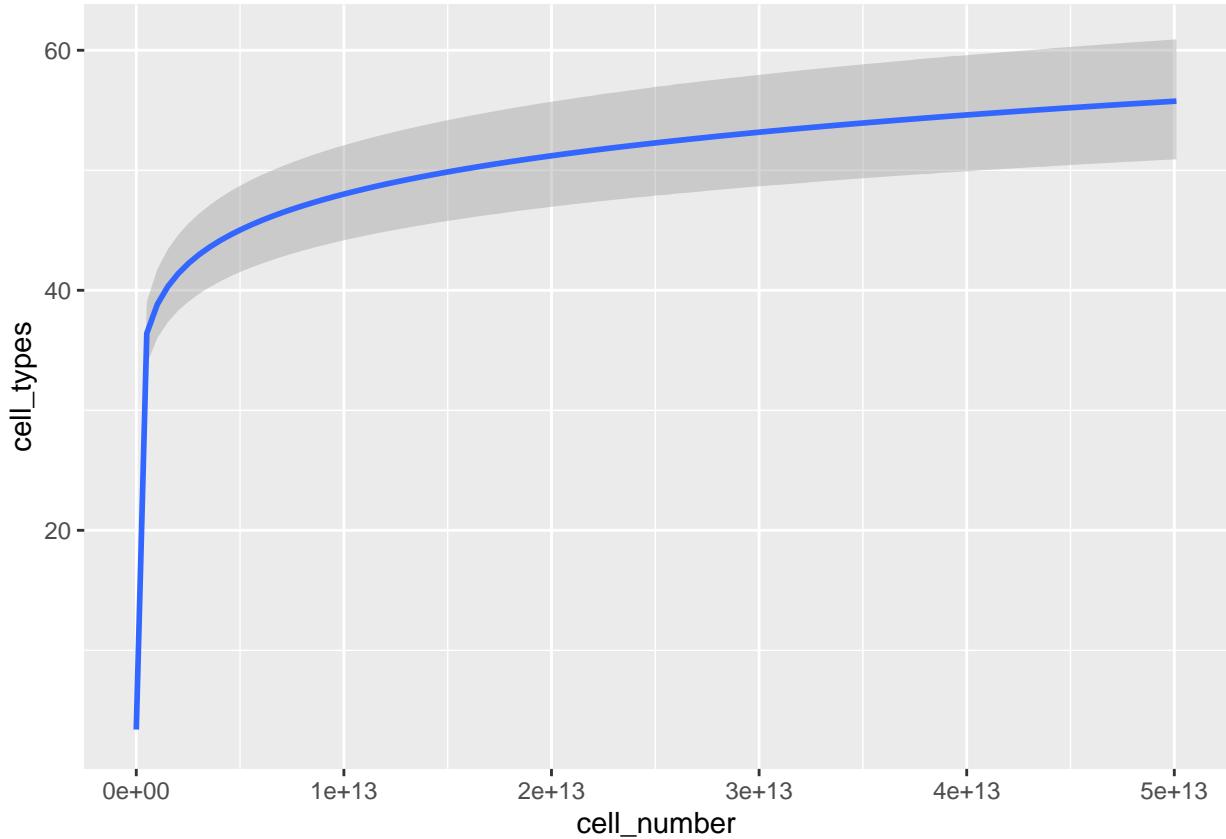
```

##                                     Estimate   Est.Error
## b_FissionOrBuddingObserved_Genus_nominalno    2.3787467  0.03347560
## b_FissionOrBuddingObserved_Genus_nominalyes   2.0677443  0.04129789
## b_scalelogcell_number                         0.7889378  0.02508357
## lp__                                         -1062.9584721 0.98458986
##                                     Q2.5      Q97.5
## b_FissionOrBuddingObserved_Genus_nominalno    2.3124781    2.4435956
## b_FissionOrBuddingObserved_Genus_nominalyes   1.9861762    2.1479040
## b_scalelogcell_number                         0.7397155    0.8380418
## lp__                                         -1066.4653841 -1061.8842775

```

```
plot(conditional_effects(fit_fission_cell_type, points = TRUE, ask = F))
```

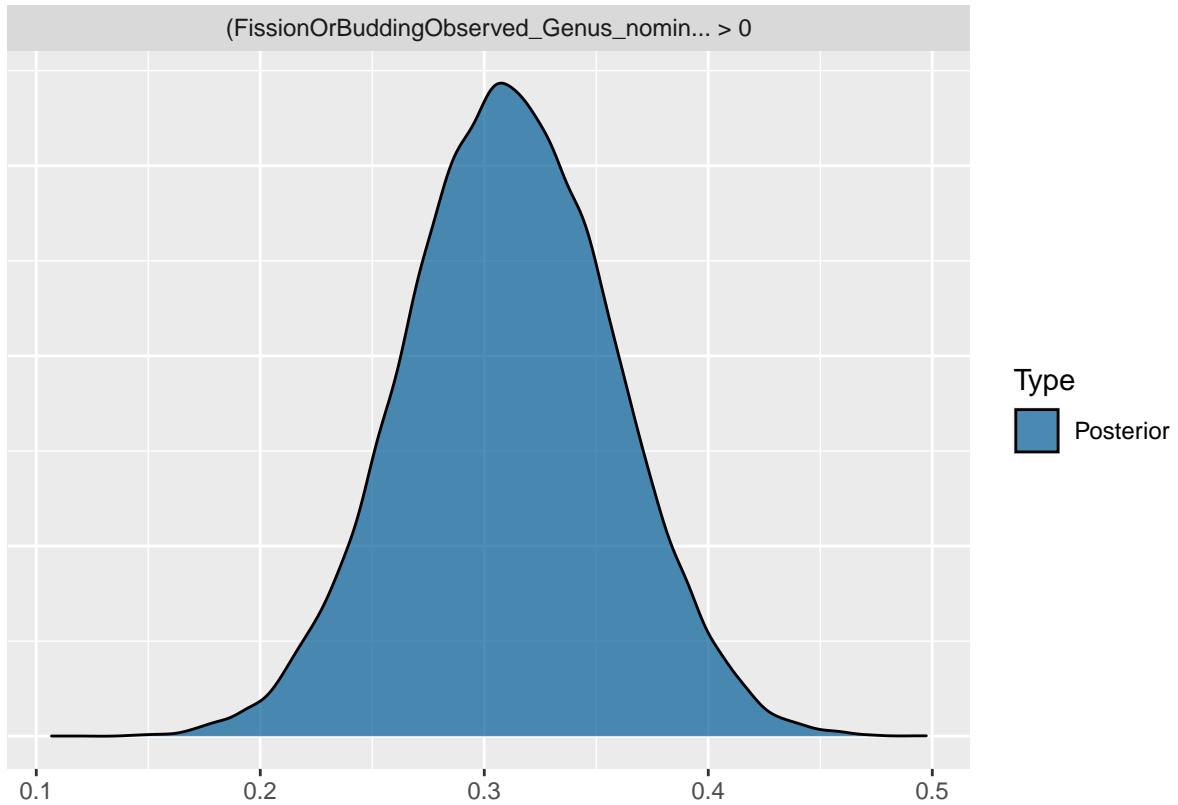




```
hyp = hypothesis(fit_fission_cell_type, "FissionOrBuddingObserved_Genus_nominalno > FissionOrBuddingObserved_Genus_nominalno")
hyp
```

```
## Hypothesis Tests for class b:
##          Hypothesis Estimate Est.Error CI.Lower CI.Upper Evid.Ratio
## 1 (FissionOrBudding... > 0      0.31      0.05     0.24     0.39      Inf
##   Post.Prob Star
## 1           1     *
## ---
## 'CI': 90%-CI for one-sided and 95%-CI for two-sided hypotheses.
## '*': For one-sided hypotheses, the posterior probability exceeds 95%;
## for two-sided hypotheses, the value tested against lies outside the 95%-CI.
## Posterior probabilities of point hypotheses assume equal prior probabilities.
```

```
plot(hyp)
```



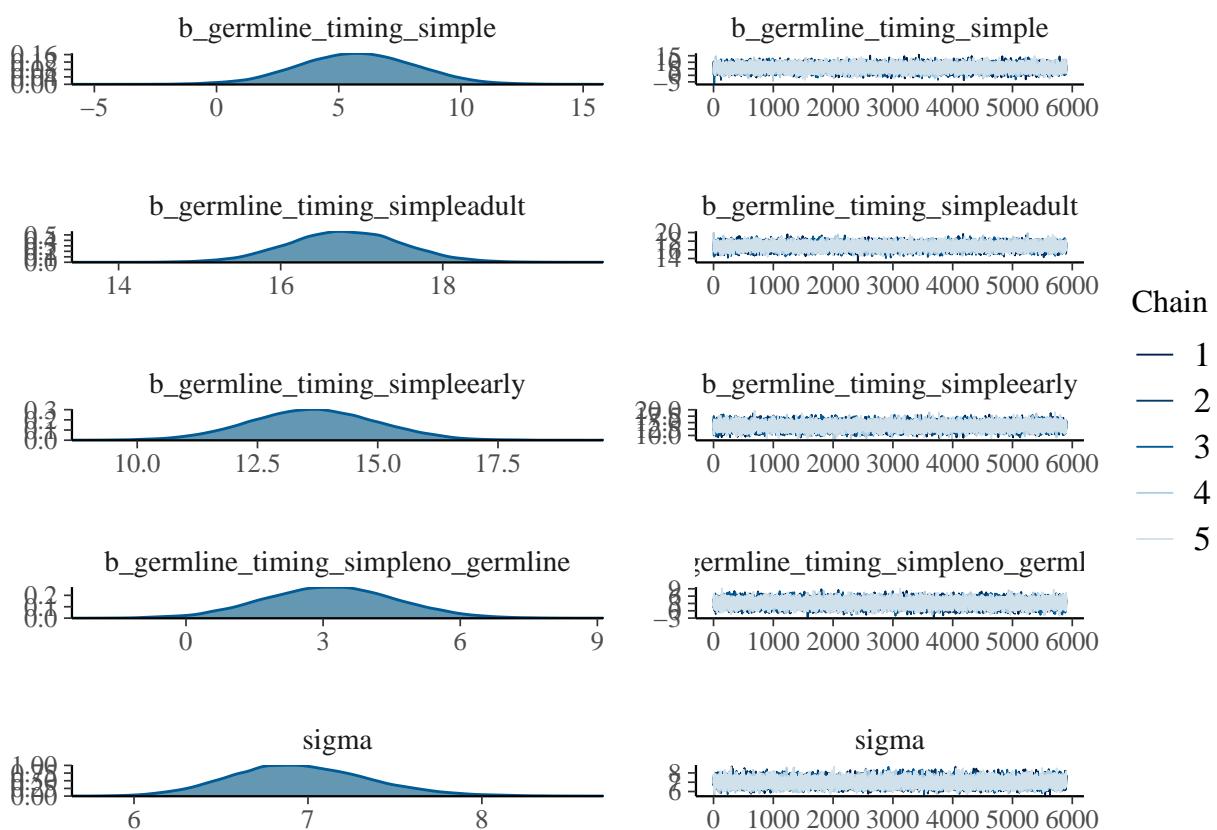
```
#### Does germline timing correlate with increased cell number?
```

```
prior <- get_prior(log(cell_number) ~ 0 + germline_timing_simple, family=gaussian(), data = df) #what p
```

```
## Warning: Rows containing NAs were excluded from the model.
```

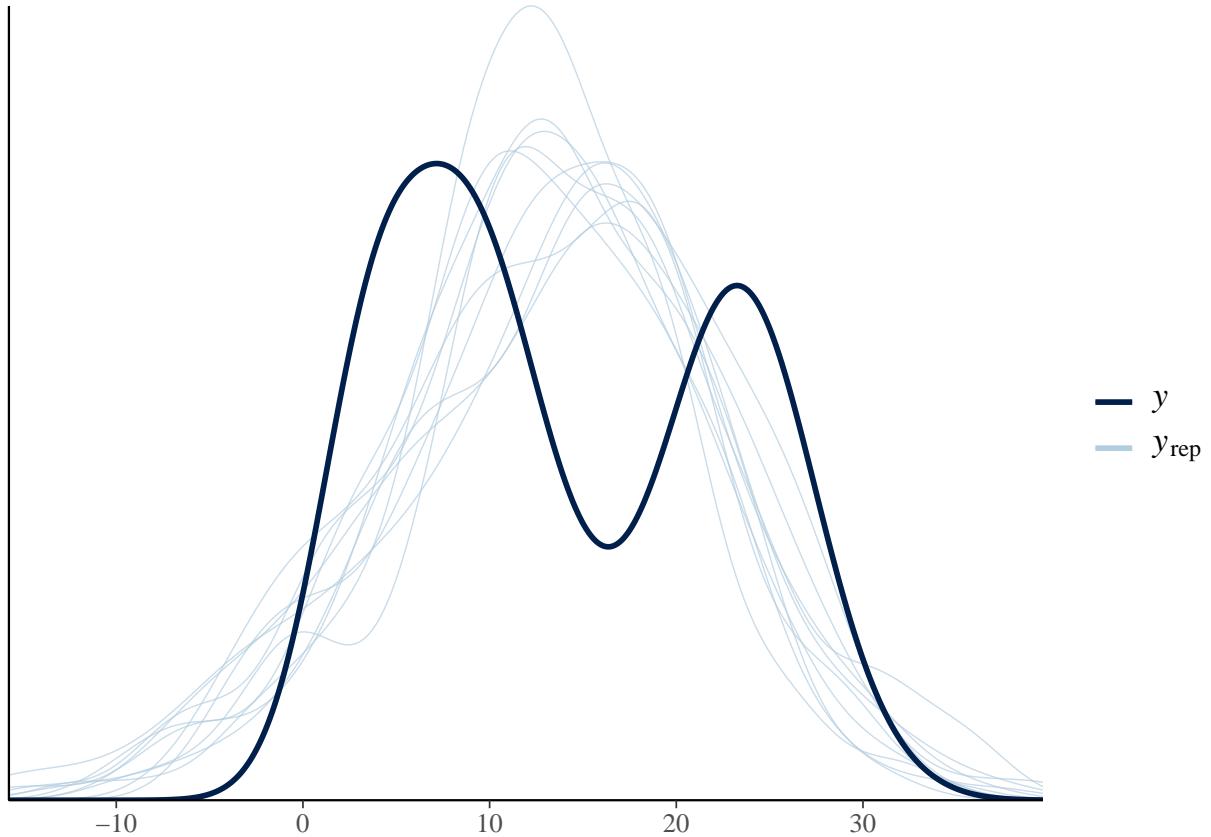
```
#fit the model
fit_germline_cell_num<-
  brm(data = df,
       family=gaussian(), # family of model
       formula = log(cell_number) ~ 0 + germline_timing_simple, #formula, the 0 means that there are es
       iter = 6000000, warmup = 100000, chains = 5,thin = 1000, cores = 5, #chain settings
       prior = prior(normal(0, 10), "b"), #defining the priors- for things in 'class b' set this prior.
       file = 'fits/fit_GermCellNum' )

plot(fit_germline_cell_num) #check that chains converged
```

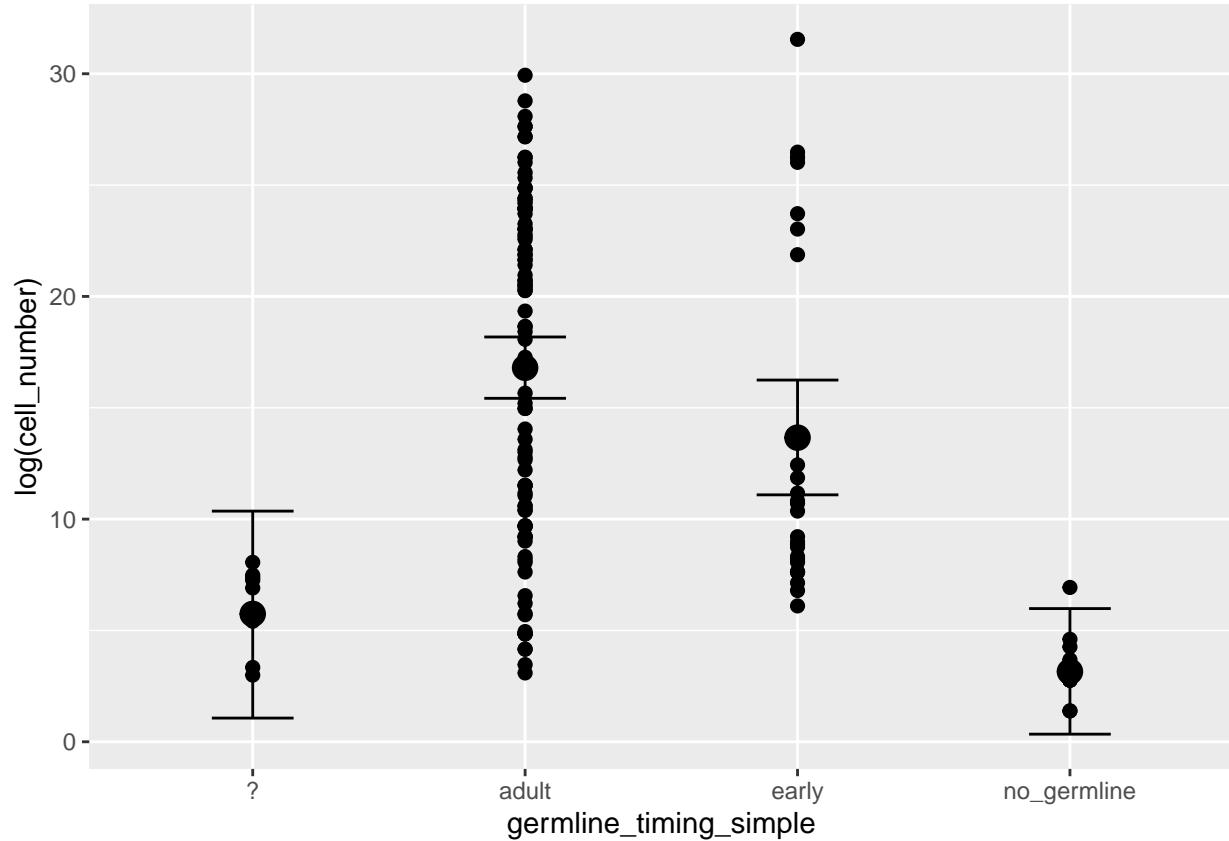


```
pp_check(fit_germline_cell_num) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



```
plot(conditional_effects(fit_germline_cell_num), points = TRUE)
```

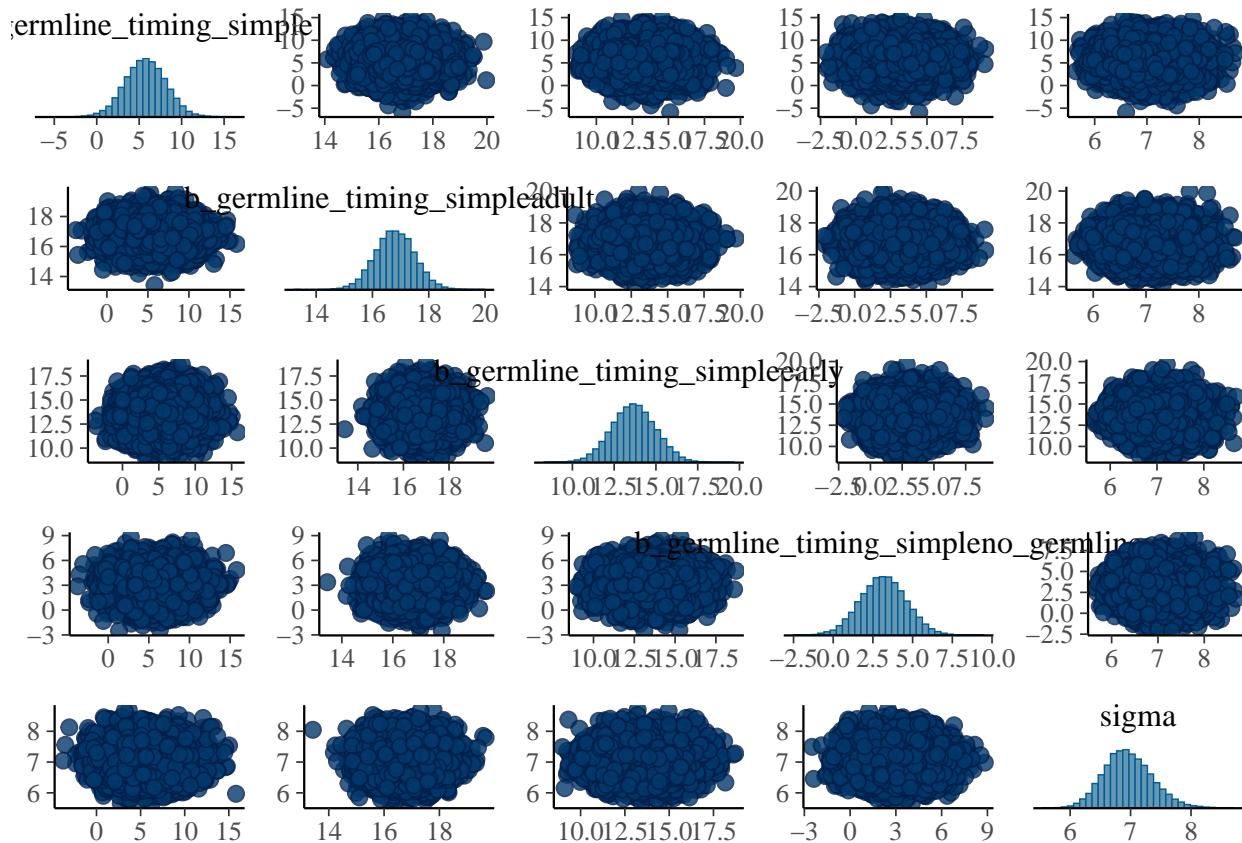


```
summary(fit_germline_cell_num)
```

```
## Family: gaussian
## Links: mu = identity; sigma = identity
## Formula: log(cell_number) ~ 0 + germline_timing_simple
## Data: df (Number of observations: 158)
## Samples: 5 chains, each with iter = 6e+06; warmup = 1e+05; thin = 1000;
##          total post-warmup samples = 29500
##
## Population-Level Effects:
##                               Estimate Est.Error 1-95% CI u-95% CI Rhat
## germline_timing_simple      5.75     2.38    1.06   10.36 1.00
## germline_timing_simpleadult 16.80     0.70   15.42   18.18 1.00
## germline_timing_simpleearly 13.67     1.32   11.09   16.24 1.00
## germline_timing_simpleno_germline 3.15     1.45    0.34    5.98 1.00
##                                         Bulk_ESS Tail_ESS
## germline_timing_simple        29223    28981
## germline_timing_simpleadult  28851    27031
## germline_timing_simpleearly  28632    28205
## germline_timing_simpleno_germline 29223    29179
##
## Family Specific Parameters:
##             Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sigma       6.96     0.40    6.23    7.80 1.00    29352   29552
##
```

```
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

```
pairs(fit_germline_cell_num)
```

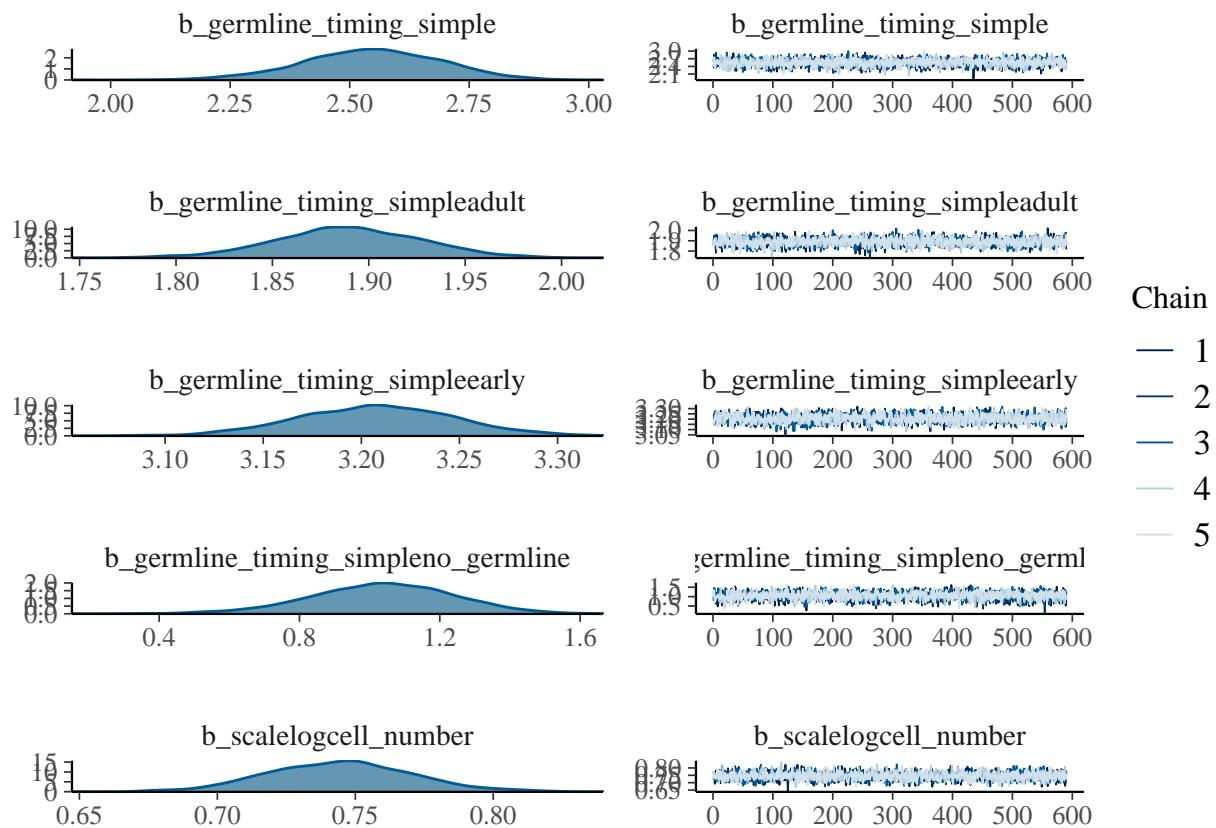


```
posterior_summary(fit_germline_cell_num, robust = T) #what are the medians for coefficients? if F, then
```

	Estimate	Est.Error	Q2.5
## b_germline_timing_simple	5.740959	2.3853286	1.0647546
## b_germline_timing_simpleadult	16.795643	0.6978221	15.4247193
## b_germline_timing_simpleearly	13.658220	1.3175933	11.0893763
## b_germline_timing_simpleno_germline	3.150338	1.4499312	0.3424271
## sigma	6.938587	0.3919637	6.2309749
## lp__	-546.467963	1.4160427	-550.7999339
			Q97.5
## b_germline_timing_simple	10.358601		
## b_germline_timing_simpleadult	18.178305		
## b_germline_timing_simpleearly	16.244379		
## b_germline_timing_simpleno_germline	5.982424		
## sigma	7.798130		
## lp__	-544.704614		

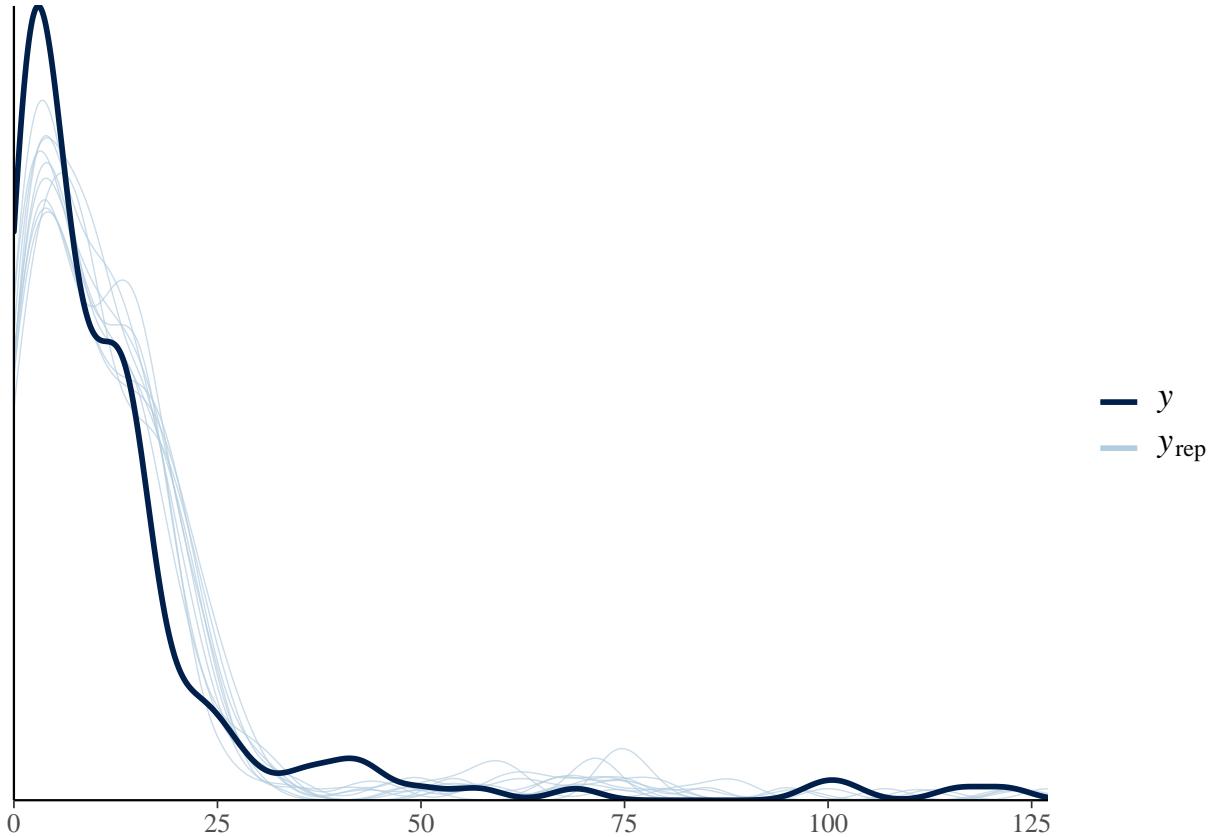
Does germline timing correlate with increased cell types (per cell)?

```
fit_type<-  
  brm(data = df,  
    family=poisson(),  
    formula = cell_types ~ 0 + germline_timing_simple + scale(log(cell_number)),  
    iter = 6000000, warmup = 100000, chains = 5, thin = 10000, cores = 5,  
    prior = prior(normal(0, 10), "b"), file = 'fits/fit_GermCellType')  
  
plot(fit_type) #check that chains converged
```



```
pp_check(fit_type) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



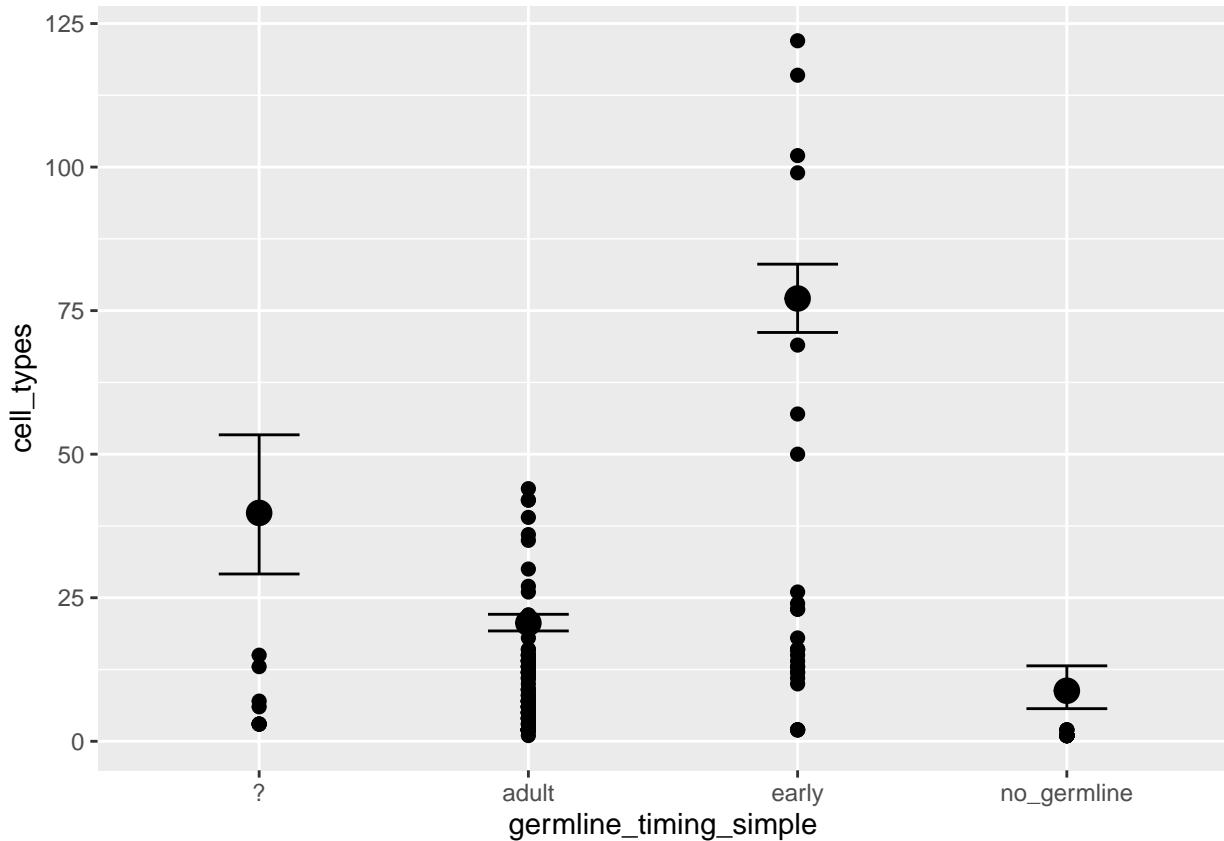
```
summary(fit_type) #summary of model
```

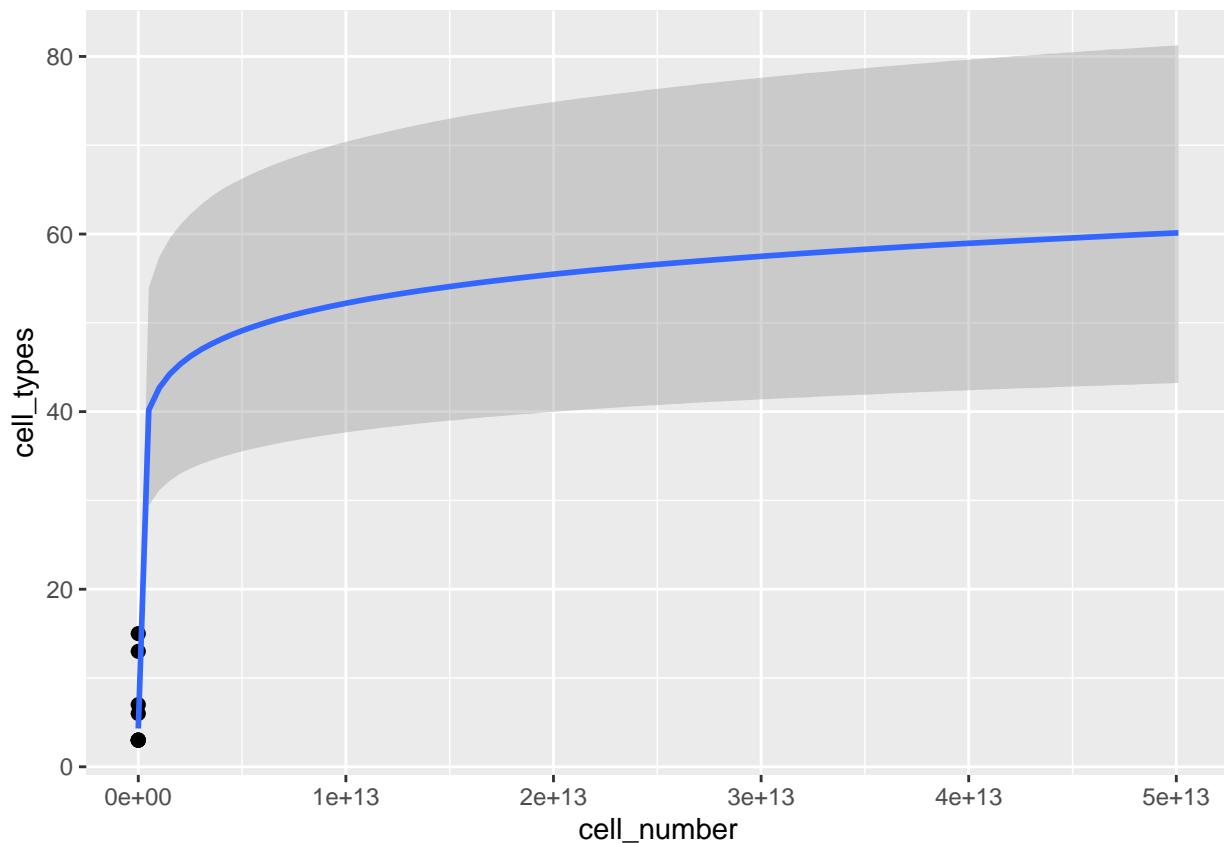
```
## Family: poisson
## Links: mu = log
## Formula: cell_types ~ 0 + germline_timing_simple + scale(log(cell_number))
## Data: df (Number of observations: 157)
## Samples: 5 chains, each with iter = 6e+06; warmup = 1e+05; thin = 10000;
##          total post-warmup samples = 2950
##
## Population-Level Effects:
##                               Estimate Est.Error 1-95% CI u-95% CI Rhat
## germline_timing_simple           2.55     0.14    2.26    2.82 1.00
## germline_timing_simpleadult     1.89     0.04    1.81    1.96 1.00
## germline_timing_simpleearly     3.21     0.04    3.13    3.28 1.00
## germline_timing_simpleno_germline 1.04     0.20    0.62    1.42 1.00
## scalelogcell_number            0.74     0.03    0.70    0.80 1.00
##                               Bulk_ESS Tail_ESS
## germline_timing_simple         2673      2593
## germline_timing_simpleadult   2830      2381
## germline_timing_simpleearly   2804      2903
## germline_timing_simpleno_germline 3223      2797
## scalelogcell_number           2872      2968
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

```
posterior_summary(fit_type, robust = T)
```

```
##                                     Estimate   Est.Error      Q2.5
## b_germline_timing_simple           2.5487216 0.14423226 2.2551736
## b_germline_timing_simpleadult    1.8894040 0.03775240 1.8144627
## b_germline_timing_simpleearly    3.2075050 0.03998024 3.1297714
## b_germline_timing_simpleno_germline 1.0396894 0.20202034 0.6211784
## b_scalelogcell_number            0.7445076 0.02645249 0.6956847
## lp_--                                -691.2963797 1.44798205 -695.5565536
##                                         Q97.5
## b_germline_timing_simple           2.8158235
## b_germline_timing_simpleadult    1.9622219
## b_germline_timing_simpleearly    3.2847058
## b_germline_timing_simpleno_germline 1.4226097
## b_scalelogcell_number            0.7974566
## lp_--                                -689.5067283
```

```
plot(conditional_effects(fit_type), points = TRUE, ask = F)
```





Phylogenetically informed:

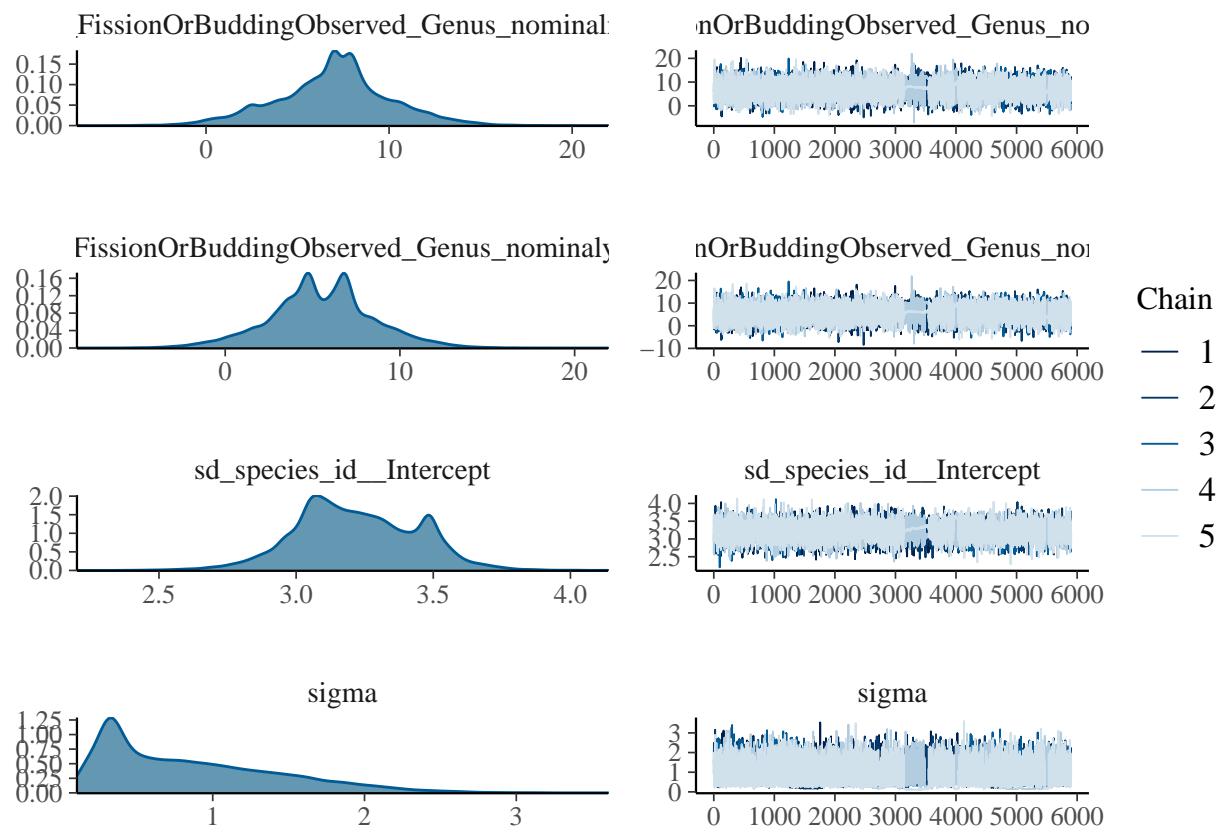
Does a single-celled bottleneck correlate with increased cell number?

```

fit_cellnumber_fission_phy<-
  brm(data = df,
       family=gaussian(), # family of model
       formula = log(cell_number) ~ 0 + FissionOrBuddingObserved_Genus_nominal + (1|gr(species_id, cov =
       iter = 6000000, warmup = 100000, chains = 5, thin = 1000, cores = 5,
       prior = prior(normal(0, 10), "b"), file = 'fits/fit_phy_FissionCellNumber', # same simple prior
       data2 = list(CovarMatrix = CovarMatrix))

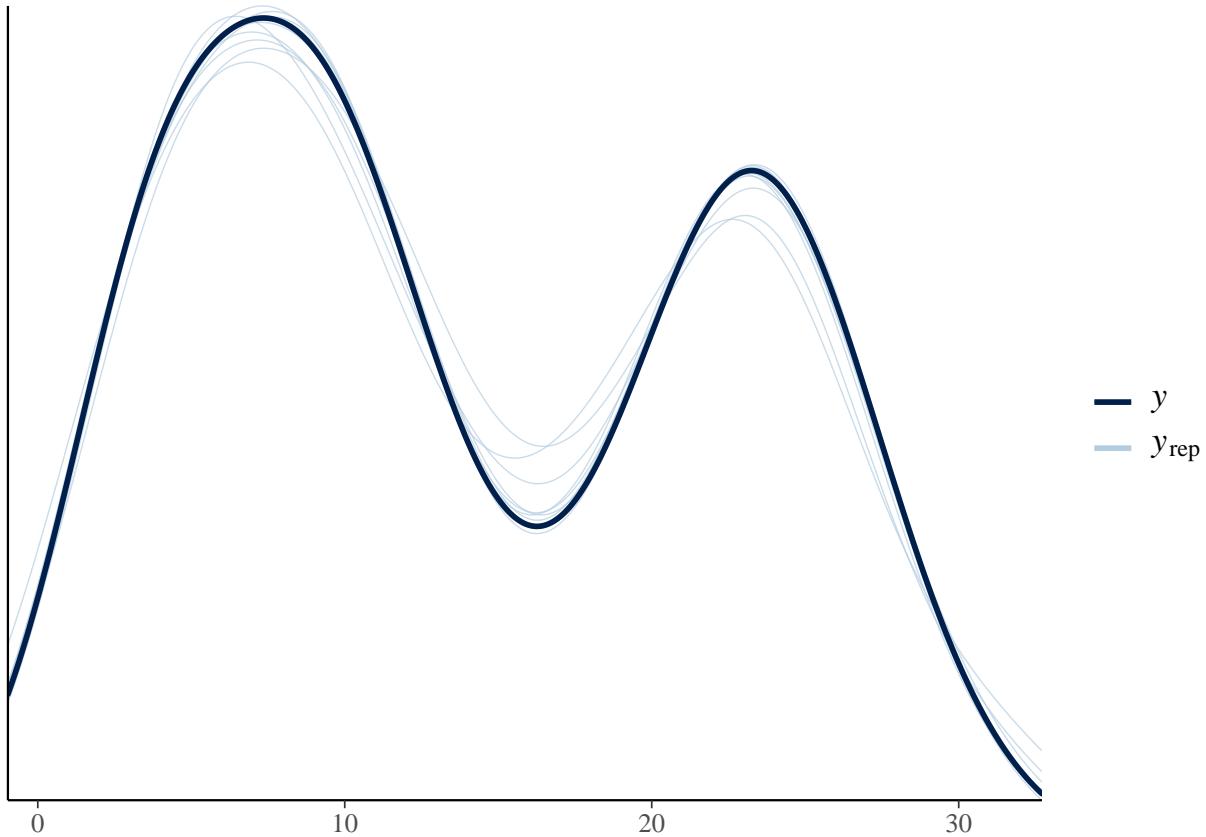
plot(fit_cellnumber_fission_phy) #check that chains converged

```



```
pp_check(fit_cellnumber_fission_phy) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



```

summary(fit_cellnumber_fission_phy) #summary of model

## Warning: Parts of the model have not converged (some Rhats are > 1.05). Be
## careful when analysing the results! We recommend running more iterations and/or
## setting stronger priors.

## Warning: There were 12874 divergent transitions after warmup.
## Increasing adapt_delta above 0.8 may help. See http://mc-stan.org/misc/
## warnings.html#divergent-transitions-after-warmup

## Family: gaussian
##   Links: mu = identity; sigma = identity
## Formula: log(cell_number) ~ 0 + FissionOrBuddingObserved_Genus_nominal + (1 | gr(species_id, cov =
## Data: df (Number of observations: 155)
## Samples: 5 chains, each with iter = 6e+06; warmup = 1e+05; thin = 1000;
##          total post-warmup samples = 29500
##
## Group-Level Effects:
## ~species_id (Number of levels: 155)
##             Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)    3.22     0.22     2.80     3.63 1.04      76     2143
## 
## Population-Level Effects:
##                               Estimate Est.Error l-95% CI u-95% CI
## FissionOrBuddingObserved_Genus_nominalno       7.00     3.07     0.62    13.20

```

```

## FissionOrBudding0bserved_Genus_nominalyes      5.51      2.99   -0.68    11.64
##                                         Rhat Bulk_ESS Tail_ESS
## FissionOrBudding0bserved_Genus_nominalno   1.06      526    1236
## FissionOrBudding0bserved_Genus_nominalyes  1.03      730    1621
##
## Family Specific Parameters:
##           Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sigma     0.86      0.57    0.15    2.18 1.03      95      64
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).

```

```
posterior_summary(fit_cellnumber_fission_phy, robust = T)
```

	Estimate	Est.Error
## b_FissionOrBudding0bserved_Genus_nominalno	7.06462590	2.5895889
## b_FissionOrBudding0bserved_Genus_nominalyes	5.45522328	2.6092922
## sd_species_id_Intercept	3.20384042	0.2264722
## sigma	0.72089274	0.5919395
## r_species_id[ott1002450,Intercept]	-1.85243086	2.8090915
## r_species_id[ott1017821,Intercept]	-2.80044951	2.7038441
## r_species_id[ott1052546,Intercept]	-2.73875225	2.7642125
## r_species_id[ott1059898,Intercept]	4.01304318	2.7785176
## r_species_id[ott1059900,Intercept]	5.82412830	2.7165660
## r_species_id[ott1061937,Intercept]	-0.14428369	2.6766553
## r_species_id[ott1069171,Intercept]	14.48010103	2.7532761
## r_species_id[ott1072227,Intercept]	-4.24083957	2.6928973
## r_species_id[ott108923,Intercept]	7.71513917	2.6102208
## r_species_id[ott1099013,Intercept]	16.80643889	2.6667915
## r_species_id[ott111442,Intercept]	5.52938040	2.6811066
## r_species_id[ott112015,Intercept]	-4.17709194	2.6744641
## r_species_id[ott112016,Intercept]	-4.20107568	2.7372812
## r_species_id[ott112017,Intercept]	-4.20765377	2.7074165
## r_species_id[ott127047,Intercept]	-0.35580876	2.7180326
## r_species_id[ott150272,Intercept]	1.86807718	2.6927313
## r_species_id[ott160850,Intercept]	4.39747634	2.7447592
## r_species_id[ott165368,Intercept]	18.85790714	2.7583775
## r_species_id[ott167121,Intercept]	-3.50129469	2.7580639
## r_species_id[ott178177,Intercept]	12.69432336	2.6601421
## r_species_id[ott178412,Intercept]	23.66802988	2.9719074
## r_species_id[ott181933,Intercept]	20.90184100	2.7906664
## r_species_id[ott182906,Intercept]	18.99877082	2.8370405
## r_species_id[ott186999,Intercept]	16.99113276	2.7565517
## r_species_id[ott187583,Intercept]	1.26693806	2.6848839
## r_species_id[ott199292,Intercept]	9.58302637	2.7507597
## r_species_id[ott207134,Intercept]	14.86303398	2.7465218
## r_species_id[ott215125,Intercept]	18.44067947	2.7236789
## r_species_id[ott216694,Intercept]	20.55006858	2.7228557
## r_species_id[ott223669,Intercept]	19.12623089	2.6900868
## r_species_id[ott225275,Intercept]	15.90837936	2.6453660
## r_species_id[ott237608,Intercept]	15.14576471	2.7744054
## r_species_id[ott246046,Intercept]	4.10050148	2.8871997
## r_species_id[ott247341,Intercept]	24.15825759	2.7530895

## r_species_id[ott256062,Intercept]	-2.17294819	2.8040605
## r_species_id[ott256089,Intercept]	-3.52491631	2.8130362
## r_species_id[ott256145,Intercept]	-2.08522229	2.6759763
## r_species_id[ott263960,Intercept]	14.89675567	2.7572527
## r_species_id[ott263980,Intercept]	14.88774825	2.7323777
## r_species_id[ott263987,Intercept]	11.39585914	2.6754832
## r_species_id[ott263988,Intercept]	17.21976655	2.6710946
## r_species_id[ott265121,Intercept]	20.27596373	2.7982612
## r_species_id[ott266342,Intercept]	0.32072246	2.6698828
## r_species_id[ott269063,Intercept]	-2.32046776	2.7214678
## r_species_id[ott275893,Intercept]	14.66677246	2.7158486
## r_species_id[ott275897,Intercept]	13.43791522	2.8018966
## r_species_id[ott2810724,Intercept]	-3.07024715	2.7981436
## r_species_id[ott2819986,Intercept]	17.07850384	2.7711695
## r_species_id[ott2821097,Intercept]	15.45500911	2.6395185
## r_species_id[ott2844172,Intercept]	2.83086212	2.8482366
## r_species_id[ott2844962,Intercept]	0.25547343	2.7371942
## r_species_id[ott2849837,Intercept]	-0.05825265	2.7480528
## r_species_id[ott2942244,Intercept]	4.70885092	2.7197860
## r_species_id[ott316441,Intercept]	15.59252344	2.7838582
## r_species_id[ott33153,Intercept]	-2.68853508	2.6941110
## r_species_id[ott336388,Intercept]	20.17661393	2.7708653
## r_species_id[ott34559,Intercept]	15.96944183	2.7292542
## r_species_id[ott346740,Intercept]	19.16350509	2.7894092
## r_species_id[ott3583594,Intercept]	2.11305880	2.7593074
## r_species_id[ott3587677,Intercept]	-0.68530482	2.7407637
## r_species_id[ott359012,Intercept]	-4.26750563	2.7914553
## r_species_id[ott361837,Intercept]	-2.64190158	2.6623175
## r_species_id[ott362913,Intercept]	5.03144730	2.7262401
## r_species_id[ott365439,Intercept]	3.27566767	2.6616060
## r_species_id[ott3663378,Intercept]	11.54967001	2.7989450
## r_species_id[ott3665433,Intercept]	5.69653135	2.7113222
## r_species_id[ott3684291,Intercept]	0.22115732	2.6739226
## r_species_id[ott3684365,Intercept]	0.30961312	2.6899387
## r_species_id[ott3684379,Intercept]	-3.92942596	2.8074330
## r_species_id[ott3684389,Intercept]	-1.64064479	2.6989463
## r_species_id[ott3684437,Intercept]	-3.57846033	2.8029167
## r_species_id[ott381979,Intercept]	-3.89687613	2.7058268
## r_species_id[ott381980,Intercept]	3.14747236	2.6965249
## r_species_id[ott381983,Intercept]	-4.48010146	2.7381493
## r_species_id[ott395048,Intercept]	-0.43657552	2.7761680
## r_species_id[ott3974169,Intercept]	17.80063262	2.7558329
## r_species_id[ott3995126,Intercept]	18.77244271	2.6970873
## r_species_id[ott4010019,Intercept]	16.82535782	2.7518213
## r_species_id[ott4010960,Intercept]	19.10451446	2.7397183
## r_species_id[ott4011155,Intercept]	10.62806401	2.9390731
## r_species_id[ott4013437,Intercept]	16.19825677	2.7242763
## r_species_id[ott4013674,Intercept]	20.35575993	2.8604399
## r_species_id[ott4013684,Intercept]	19.11799438	2.8062905
## r_species_id[ott422679,Intercept]	0.31217461	2.6767994
## r_species_id[ott431388,Intercept]	8.32818209	2.6443083
## r_species_id[ott446088,Intercept]	3.66570294	2.6914361
## r_species_id[ott4741377,Intercept]	15.38966560	2.6838422
## r_species_id[ott4742064,Intercept]	20.14242643	2.8238228

## r_species_id[ott481952,Intercept]	19.43833164	2.8227131
## r_species_id[ott48288,Intercept]	14.85349649	2.7291567
## r_species_id[ott485470,Intercept]	0.44638186	2.6846161
## r_species_id[ott485473,Intercept]	-2.06078688	2.6968523
## r_species_id[ott485476,Intercept]	-3.49008996	2.7191755
## r_species_id[ott485480,Intercept]	2.56903733	2.7001135
## r_species_id[ott485482,Intercept]	0.26986461	2.7084729
## r_species_id[ott486834,Intercept]	-0.74023063	2.7435296
## r_species_id[ott490206,Intercept]	6.01146782	2.7661244
## r_species_id[ott492241,Intercept]	0.93177169	2.7597267
## r_species_id[ott497063,Intercept]	15.07278049	2.7461882
## r_species_id[ott4974308,Intercept]	5.88885260	2.8797881
## r_species_id[ott4978773,Intercept]	1.16175026	2.7343386
## r_species_id[ott4979583,Intercept]	1.83138040	2.7005108
## r_species_id[ott518643,Intercept]	3.59204363	2.6888601
## r_species_id[ott542509,Intercept]	18.94816328	2.7020345
## r_species_id[ott54768,Intercept]	9.59079817	2.7571210
## r_species_id[ott549846,Intercept]	7.06127581	2.7288303
## r_species_id[ott560703,Intercept]	-1.97294064	2.7702370
## r_species_id[ott567703,Intercept]	18.06534720	2.7087620
## r_species_id[ott570365,Intercept]	0.60554246	2.7851363
## r_species_id[ott570656,Intercept]	12.09267016	2.7319978
## r_species_id[ott588761,Intercept]	3.69688440	2.7379335
## r_species_id[ott592355,Intercept]	14.97983887	2.7412680
## r_species_id[ott601255,Intercept]	23.16553691	2.7210835
## r_species_id[ott602180,Intercept]	-0.84721264	2.7104600
## r_species_id[ott60470,Intercept]	-2.67504695	2.7610017
## r_species_id[ott60473,Intercept]	-2.70017893	2.7153491
## r_species_id[ott60479,Intercept]	-3.48652780	2.7023470
## r_species_id[ott633708,Intercept]	1.12622173	2.6814608
## r_species_id[ott633710,Intercept]	3.15557170	2.7344595
## r_species_id[ott633711,Intercept]	1.76278892	2.7059625
## r_species_id[ott633717,Intercept]	-2.26409405	2.7441659
## r_species_id[ott633719,Intercept]	2.42368014	2.7063111
## r_species_id[ott643237,Intercept]	16.69984964	2.7296852
## r_species_id[ott645555,Intercept]	19.63684711	2.7968853
## r_species_id[ott649193,Intercept]	18.55450035	2.7795635
## r_species_id[ott675301,Intercept]	1.11082937	2.7405402
## r_species_id[ott724784,Intercept]	10.33238337	2.9094993
## r_species_id[ott72522,Intercept]	20.29659929	2.6985701
## r_species_id[ott727979,Intercept]	21.71416569	2.8690236
## r_species_id[ott733462,Intercept]	16.89958936	2.7111354
## r_species_id[ott736728,Intercept]	5.37033166	2.7363974
## r_species_id[ott742128,Intercept]	3.56133377	2.6983100
## r_species_id[ott7489702,Intercept]	3.80584590	2.7126916
## r_species_id[ott7567530,Intercept]	5.49684986	2.7330045
## r_species_id[ott765113,Intercept]	-0.52866604	2.7085757
## r_species_id[ott765280,Intercept]	13.86883792	2.6744774
## r_species_id[ott779028,Intercept]	16.42257505	2.7051891
## r_species_id[ott790395,Intercept]	15.99967293	2.6977480
## r_species_id[ott817791,Intercept]	14.61663453	2.8390008
## r_species_id[ott821356,Intercept]	15.18999811	2.8275455
## r_species_id[ott83430,Intercept]	11.33357986	2.6885018
## r_species_id[ott83432,Intercept]	5.11148068	2.8690091

## r_species_id[ott840001,Intercept]	18.43516581	2.7722402
## r_species_id[ott841027,Intercept]	0.29718964	2.7355523
## r_species_id[ott849781,Intercept]	13.98145426	2.7217247
## r_species_id[ott878345,Intercept]	5.76361294	2.7333749
## r_species_id[ott92556,Intercept]	13.10896349	2.7573013
## r_species_id[ott92561,Intercept]	14.58847774	2.6771453
## r_species_id[ott939432,Intercept]	4.38779929	2.6871800
## r_species_id[ott939454,Intercept]	4.42192810	2.6682235
## r_species_id[ott954042,Intercept]	16.28009470	2.8028405
## r_species_id[ott958293,Intercept]	-3.35840565	2.7726895
## r_species_id[ott958304,Intercept]	-4.24099051	2.7461189
## r_species_id[ott962359,Intercept]	2.68090529	2.7190010
## r_species_id[ott987480,Intercept]	0.19368309	2.7132913
## lp__	-399.71144620	137.6455875
##	Q2.5	Q97.5
## b_FissionOrBuddingObserved_Genus_nominalno	0.6153936	13.195213
## b_FissionOrBuddingObserved_Genus_nominalyes	-0.6794411	11.638574
## sd_species_id_Intercept	2.8047455	3.630059
## sigma	0.1534412	2.183785
## r_species_id[ott1002450,Intercept]	-8.2535254	4.507410
## r_species_id[ott1017821,Intercept]	-9.2129083	3.850850
## r_species_id[ott1052546,Intercept]	-9.2473592	3.854425
## r_species_id[ott1059898,Intercept]	-2.3087431	10.518682
## r_species_id[ott1059900,Intercept]	-0.5831382	12.294295
## r_species_id[ott1061937,Intercept]	-6.4365593	6.695234
## r_species_id[ott1069171,Intercept]	7.8528621	20.883961
## r_species_id[ott1072227,Intercept]	-10.6696703	2.511261
## r_species_id[ott108923,Intercept]	1.3654064	14.159741
## r_species_id[ott1099013,Intercept]	10.2768144	23.359566
## r_species_id[ott111442,Intercept]	-0.9207088	11.972166
## r_species_id[ott112015,Intercept]	-10.6606933	2.386327
## r_species_id[ott112016,Intercept]	-10.6022482	2.363083
## r_species_id[ott112017,Intercept]	-10.6540915	2.398650
## r_species_id[ott127047,Intercept]	-6.5244252	6.450267
## r_species_id[ott150272,Intercept]	-4.6891239	8.410393
## r_species_id[ott160850,Intercept]	-2.1814263	10.927224
## r_species_id[ott165368,Intercept]	12.6696491	25.828349
## r_species_id[ott167121,Intercept]	-9.8557888	3.145288
## r_species_id[ott178177,Intercept]	6.4304450	19.300072
## r_species_id[ott178412,Intercept]	16.4810238	30.084451
## r_species_id[ott181933,Intercept]	14.4299745	27.617541
## r_species_id[ott182906,Intercept]	12.1287514	25.699551
## r_species_id[ott186999,Intercept]	10.5078434	23.372559
## r_species_id[ott187583,Intercept]	-5.2000029	7.937452
## r_species_id[ott199292,Intercept]	3.1507855	16.023225
## r_species_id[ott207134,Intercept]	8.2895121	21.492542
## r_species_id[ott215125,Intercept]	12.0046116	24.849939
## r_species_id[ott216694,Intercept]	14.0932150	27.172168
## r_species_id[ott223669,Intercept]	12.5827142	25.795217
## r_species_id[ott225275,Intercept]	9.4446519	22.573019
## r_species_id[ott237608,Intercept]	8.4970227	21.482666
## r_species_id[ott246046,Intercept]	-2.3948808	10.454370
## r_species_id[ott247341,Intercept]	17.3672435	30.799265
## r_species_id[ott256062,Intercept]	-8.5688555	4.555120

## r_species_id[ott256089,Intercept]	-9.6511326	3.491819
## r_species_id[ott256145,Intercept]	-8.5432216	4.306427
## r_species_id[ott263960,Intercept]	8.3898498	21.332512
## r_species_id[ott263980,Intercept]	8.3715337	21.307315
## r_species_id[ott263987,Intercept]	4.9957440	17.854514
## r_species_id[ott263988,Intercept]	10.6184854	23.805139
## r_species_id[ott265121,Intercept]	13.5209187	26.770808
## r_species_id[ott266342,Intercept]	-5.8356101	7.291808
## r_species_id[ott269063,Intercept]	-8.7328513	4.116569
## r_species_id[ott275893,Intercept]	8.1221434	21.042547
## r_species_id[ott275897,Intercept]	6.9637121	20.058729
## r_species_id[ott2810724,Intercept]	-9.0700272	4.110758
## r_species_id[ott2819986,Intercept]	10.5575491	23.722539
## r_species_id[ott2821097,Intercept]	9.2464122	22.282312
## r_species_id[ott2844172,Intercept]	-3.5657695	9.224794
## r_species_id[ott2844962,Intercept]	-6.1645262	6.854245
## r_species_id[ott2849837,Intercept]	-6.2970802	6.858210
## r_species_id[ott2942244,Intercept]	-1.7905999	11.357933
## r_species_id[ott316441,Intercept]	8.7586729	22.280565
## r_species_id[ott33153,Intercept]	-9.1698201	3.965317
## r_species_id[ott336388,Intercept]	13.5516126	26.778124
## r_species_id[ott34559,Intercept]	9.3806520	22.584391
## r_species_id[ott346740,Intercept]	12.6480426	25.411975
## r_species_id[ott3583594,Intercept]	-4.2737786	8.854982
## r_species_id[ott3587677,Intercept]	-7.1759664	5.840956
## r_species_id[ott359012,Intercept]	-10.5232298	2.467659
## r_species_id[ott361837,Intercept]	-9.0906803	3.689092
## r_species_id[ott362913,Intercept]	-1.4602325	11.359886
## r_species_id[ott365439,Intercept]	-2.8674215	10.093817
## r_species_id[ott3663378,Intercept]	4.9880248	18.179411
## r_species_id[ott3665433,Intercept]	-0.8267493	12.380952
## r_species_id[ott3684291,Intercept]	-6.2346401	6.833449
## r_species_id[ott3684365,Intercept]	-6.2959529	6.872882
## r_species_id[ott3684379,Intercept]	-10.3242629	2.722197
## r_species_id[ott3684389,Intercept]	-8.1871576	4.936396
## r_species_id[ott3684437,Intercept]	-9.9204912	3.142227
## r_species_id[ott381979,Intercept]	-10.3520114	2.561869
## r_species_id[ott381980,Intercept]	-3.5255402	9.680731
## r_species_id[ott381983,Intercept]	-10.7514549	2.465423
## r_species_id[ott395048,Intercept]	-6.5306492	6.616935
## r_species_id[ott3974169,Intercept]	11.3670921	24.284927
## r_species_id[ott3995126,Intercept]	12.2511308	25.194752
## r_species_id[ott4010019,Intercept]	10.5800765	23.760267
## r_species_id[ott4010960,Intercept]	12.5779476	25.446885
## r_species_id[ott4011155,Intercept]	4.3694689	17.244428
## r_species_id[ott4013437,Intercept]	9.7964007	22.604851
## r_species_id[ott4013674,Intercept]	13.7536902	26.748760
## r_species_id[ott4013684,Intercept]	12.5436223	25.407325
## r_species_id[ott422679,Intercept]	-6.0433408	6.705244
## r_species_id[ott431388,Intercept]	1.8792033	14.800622
## r_species_id[ott446088,Intercept]	-2.7957409	10.121815
## r_species_id[ott4741377,Intercept]	8.9221781	21.722311
## r_species_id[ott4742064,Intercept]	13.5650631	26.515198
## r_species_id[ott481952,Intercept]	12.7069011	25.874878

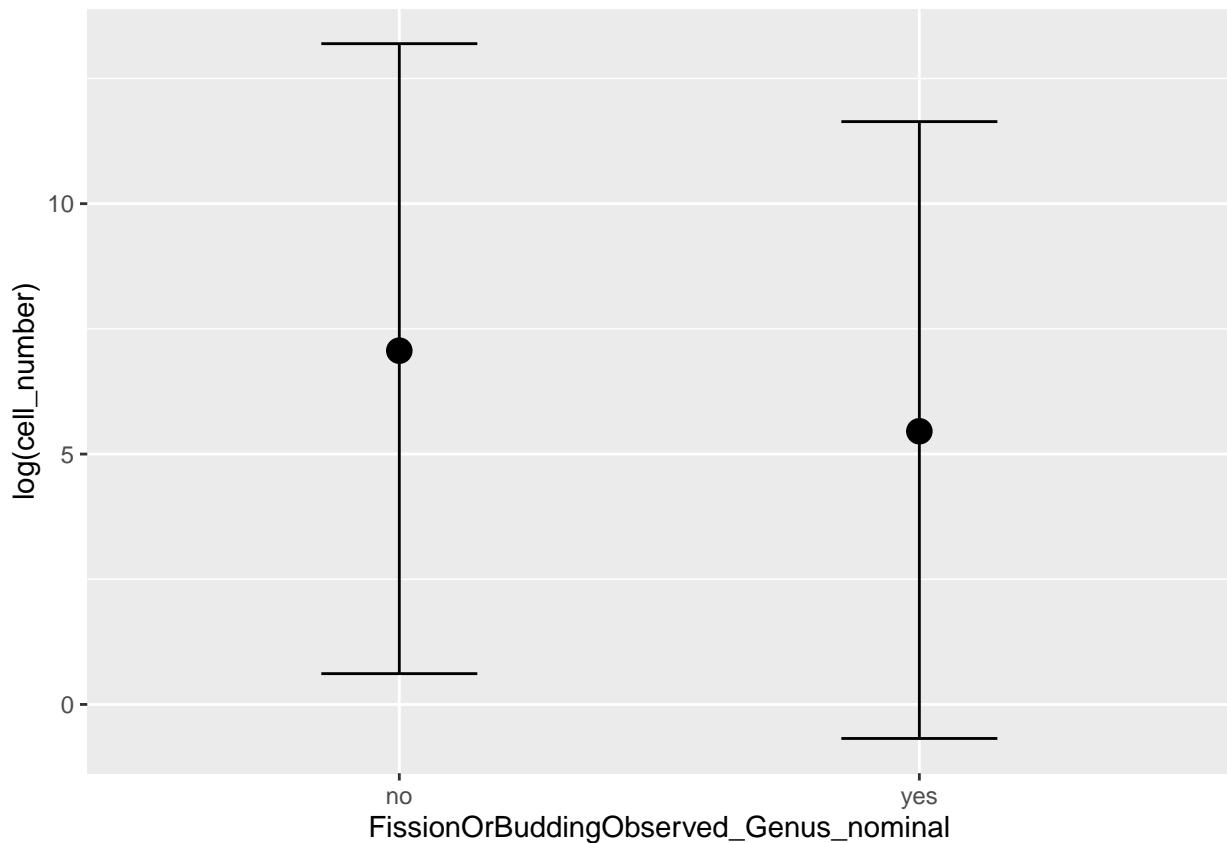
## r_species_id[ott48288,Intercept]	8.2484201	21.429629
## r_species_id[ott485470,Intercept]	-5.9854746	7.246224
## r_species_id[ott485473,Intercept]	-8.5832512	4.606154
## r_species_id[ott485476,Intercept]	-9.8571933	3.199908
## r_species_id[ott485480,Intercept]	-3.9950507	9.210695
## r_species_id[ott485482,Intercept]	-5.8149346	7.240549
## r_species_id[ott486834,Intercept]	-7.1633583	5.949320
## r_species_id[ott490206,Intercept]	-0.5466732	12.553794
## r_species_id[ott492241,Intercept]	-5.2772837	7.837014
## r_species_id[ott497063,Intercept]	8.6915411	21.693613
## r_species_id[ott4974308,Intercept]	-0.4828801	12.412550
## r_species_id[ott4978773,Intercept]	-5.1480894	7.891810
## r_species_id[ott4979583,Intercept]	-4.5609448	8.490329
## r_species_id[ott518643,Intercept]	-2.8578090	10.251227
## r_species_id[ott542509,Intercept]	12.6055087	25.650458
## r_species_id[ott54768,Intercept]	3.0906864	15.982108
## r_species_id[ott549846,Intercept]	0.6143222	13.447608
## r_species_id[ott560703,Intercept]	-8.3359703	4.813498
## r_species_id[ott567703,Intercept]	11.5668594	24.353932
## r_species_id[ott570365,Intercept]	-5.7910680	7.145330
## r_species_id[ott570656,Intercept]	5.6397451	18.552881
## r_species_id[ott588761,Intercept]	-2.7524379	10.092405
## r_species_id[ott592355,Intercept]	8.4001517	21.246589
## r_species_id[ott601255,Intercept]	16.6179137	29.604265
## r_species_id[ott602180,Intercept]	-7.2758515	5.577403
## r_species_id[ott60470,Intercept]	-9.1525329	3.786283
## r_species_id[ott60473,Intercept]	-9.1425320	3.715511
## r_species_id[ott60479,Intercept]	-9.8946869	3.060752
## r_species_id[ott633708,Intercept]	-5.4465843	7.770535
## r_species_id[ott633710,Intercept]	-3.5359003	9.750626
## r_species_id[ott633711,Intercept]	-4.6078295	8.527661
## r_species_id[ott633717,Intercept]	-8.6960403	4.483246
## r_species_id[ott633719,Intercept]	-4.0054521	9.106923
## r_species_id[ott643237,Intercept]	10.5442387	23.569382
## r_species_id[ott645555,Intercept]	13.0318503	26.048557
## r_species_id[ott649193,Intercept]	12.0564825	24.905895
## r_species_id[ott675301,Intercept]	-5.3667912	7.687026
## r_species_id[ott724784,Intercept]	3.9919737	16.818289
## r_species_id[ott72522,Intercept]	13.5598568	26.889265
## r_species_id[ott727979,Intercept]	15.2729611	28.061368
## r_species_id[ott733462,Intercept]	10.3682289	23.292054
## r_species_id[ott736728,Intercept]	-0.6671030	12.526205
## r_species_id[ott742128,Intercept]	-2.9410691	9.921909
## r_species_id[ott7489702,Intercept]	-2.3130235	10.817222
## r_species_id[ott7567530,Intercept]	-0.9381590	11.964005
## r_species_id[ott765113,Intercept]	-6.6718029	6.330556
## r_species_id[ott765280,Intercept]	7.3496208	20.316973
## r_species_id[ott779028,Intercept]	9.9072820	22.780252
## r_species_id[ott790395,Intercept]	9.5008236	22.492745
## r_species_id[ott817791,Intercept]	8.1044548	21.230039
## r_species_id[ott821356,Intercept]	8.6940649	21.516381
## r_species_id[ott83430,Intercept]	5.3467771	18.370723
## r_species_id[ott83432,Intercept]	-1.2166550	12.070073
## r_species_id[ott840001,Intercept]	11.8687001	24.716835

```

## r_species_id[ott841027,Intercept]      -5.9138623   7.292688
## r_species_id[ott849781,Intercept]      7.7855769  20.916326
## r_species_id[ott878345,Intercept]     -0.2970872  12.864811
## r_species_id[ott92556,Intercept]       6.5514766  19.478697
## r_species_id[ott92561,Intercept]       8.0554953  21.276904
## r_species_id[ott939432,Intercept]     -2.0418734  10.982796
## r_species_id[ott939454,Intercept]     -2.0257961  10.953848
## r_species_id[ott954042,Intercept]      9.4738346  22.910687
## r_species_id[ott958293,Intercept]     -9.7746836  3.248121
## r_species_id[ott958304,Intercept]     -10.6755199 2.547868
## r_species_id[ott962359,Intercept]     -3.8244462  9.103583
## r_species_id[ott987480,Intercept]     -5.9046678  7.219286
## lp__                                -571.1418682 -175.711027

```

```
plot(conditional_effects(fit_cellnumber_fission_phy, points = TRUE, ask = F))
```



```

hyp = hypothesis(fit_cellnumber_fission_phy, "FissionOrBuddingObserved_Genus_nominalno > FissionOrBuddingObserved_Genus_nominalyes")
hyp

```

```

## Hypothesis Tests for class b:
##                               Hypothesis Estimate Est.Error CI.Lower CI.Upper Evid.Ratio
## 1 (FissionOrBuddingObserved_Genus_nominalno > FissionOrBuddingObserved_Genus_nominalyes)
##   Post.Prob Star
## 1      0.94
## ---

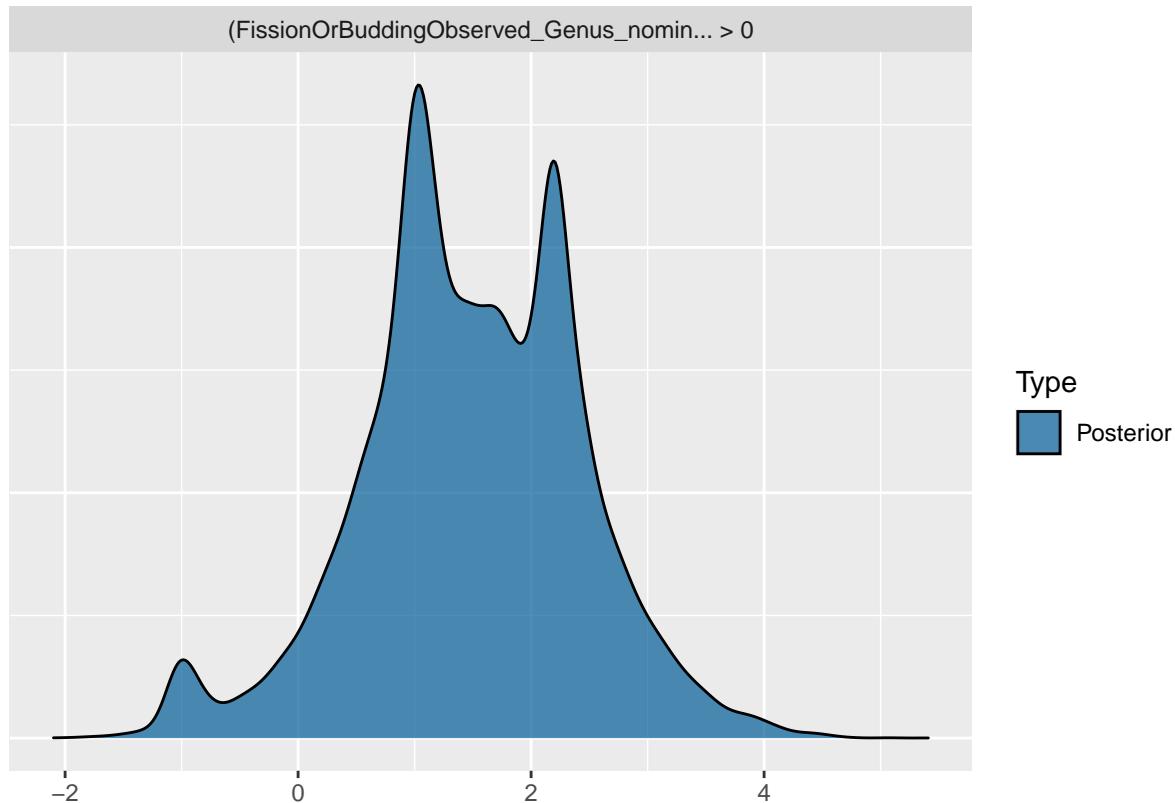
```

```

## 'CI': 90%-CI for one-sided and 95%-CI for two-sided hypotheses.
## '*': For one-sided hypotheses, the posterior probability exceeds 95%;
## for two-sided hypotheses, the value tested against lies outside the 95%-CI.
## Posterior probabilities of point hypotheses assume equal prior probabilities.

```

```
plot(hyp)
```



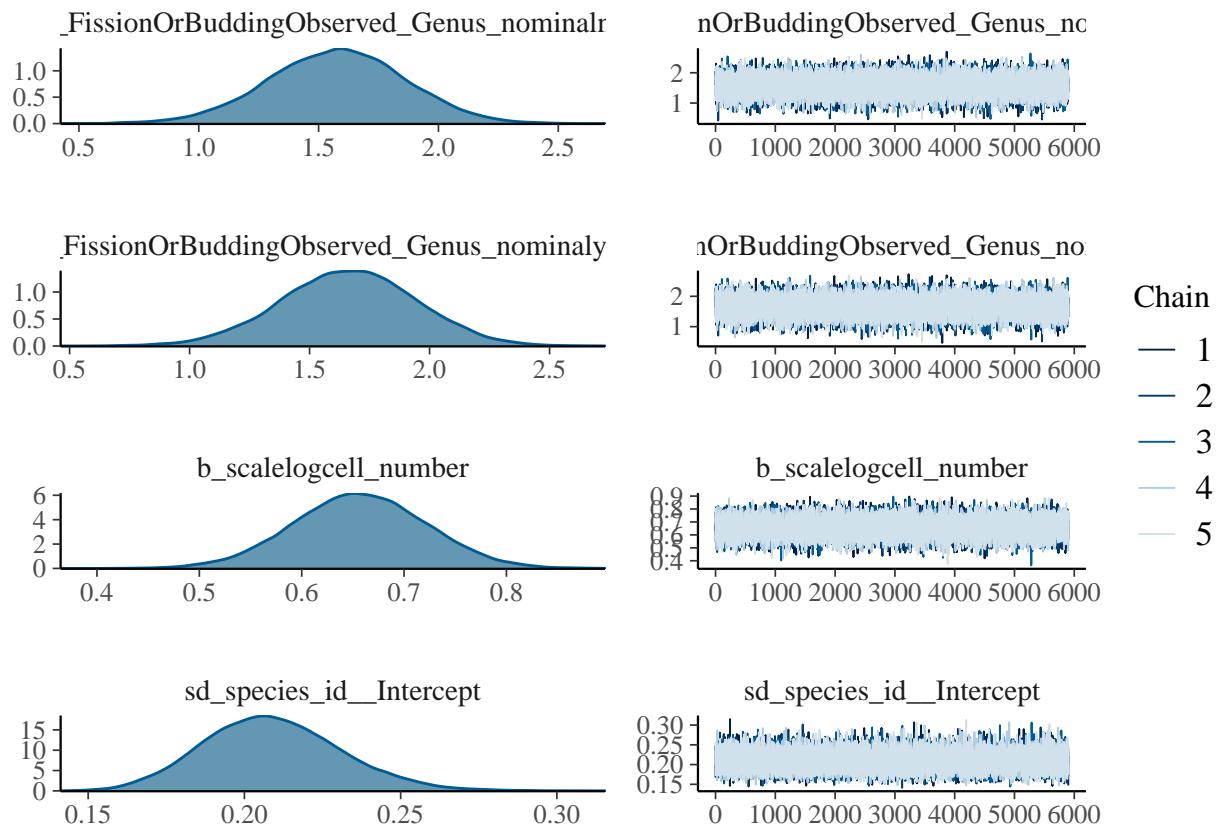
```
#### Does a single-celled bottleneck correlate with increased cell types (per cell)?
```

```

#fit the model
phylo_fit_fission_cell_type<-
  brm(data = df,
       family=poisson(),
       formula = cell_types ~ 0 + FissionOrBuddingObserved_Genus_nominal + scale(log(cell_number)) + (1|b),
       iter = 600000, warmup = 10000, chains = 5, thin = 100, cores = 5, #chain settings
       prior = prior(normal(0, 10), "b"), #defining the priors- for things in 'class b' set this prior.
       data2 = list(CovarMatrix = CovarMatrix),
       file = 'fits/fit_phy_FissionCellType')

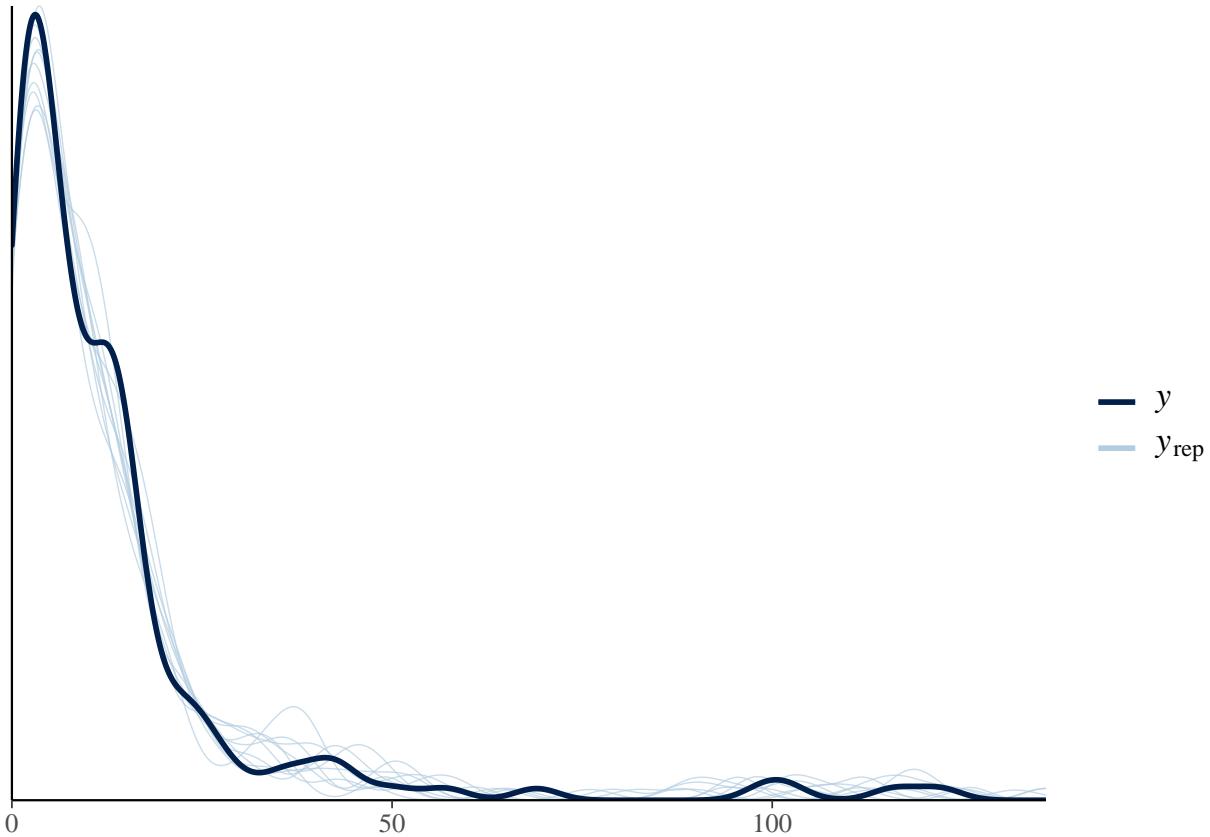
plot(phylo_fit_fission_cell_type) #check that chains converged

```



```
pp_check(phylo_fit_fission_cell_type) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



```
summary(phylo_fit_fission_cell_type) #summary of model
```

```
## Family: poisson
## Links: mu = log
## Formula: cell_types ~ 0 + FissionOrBuddingObserved_Genus_nominal + scale(log(cell_number)) + (1 | gr
## Data: df (Number of observations: 154)
## Samples: 5 chains, each with iter = 6e+05; warmup = 10000; thin = 100;
##          total post-warmup samples = 29500
##
## Group-Level Effects:
## ~species_id (Number of levels: 154)
##             Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)    0.21     0.02     0.17     0.25 1.00    28860    28825
##
## Population-Level Effects:
##                                         Estimate Est.Error l-95% CI u-95% CI
## FissionOrBuddingObserved_Genus_nominalno      1.58     0.29     1.01     2.13
## FissionOrBuddingObserved_Genus_nominalyes      1.66     0.28     1.10     2.21
## scalelogcell_number                          0.66     0.07     0.53     0.78
##                                         Rhat Bulk_ESS Tail_ESS
## FissionOrBuddingObserved_Genus_nominalno  1.00    29830   28977
## FissionOrBuddingObserved_Genus_nominalyes  1.00    29526   29464
## scalelogcell_number                      1.00    29588   28713
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
```

```
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

```
posterior_summary(phylo_fit_fission_cell_type, robust = T)
```

	Estimate	Est.Error
##		
## b_FissionOrBuddingObserved_Genus_nominalno	1.57975901	0.28431764
## b_FissionOrBuddingObserved_Genus_nominalyes	1.66406774	0.28292158
## b_scalelogcell_number	0.65641396	0.06464607
## sd_species_id_Intercept	0.20796192	0.02185879
## r_species_id[ott1017821,Intercept]	-0.13872134	0.41541290
## r_species_id[ott1052546,Intercept]	-0.14202203	0.41569741
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## r_species_id[ott160850,Intercept]	-0.23383634	0.41276043
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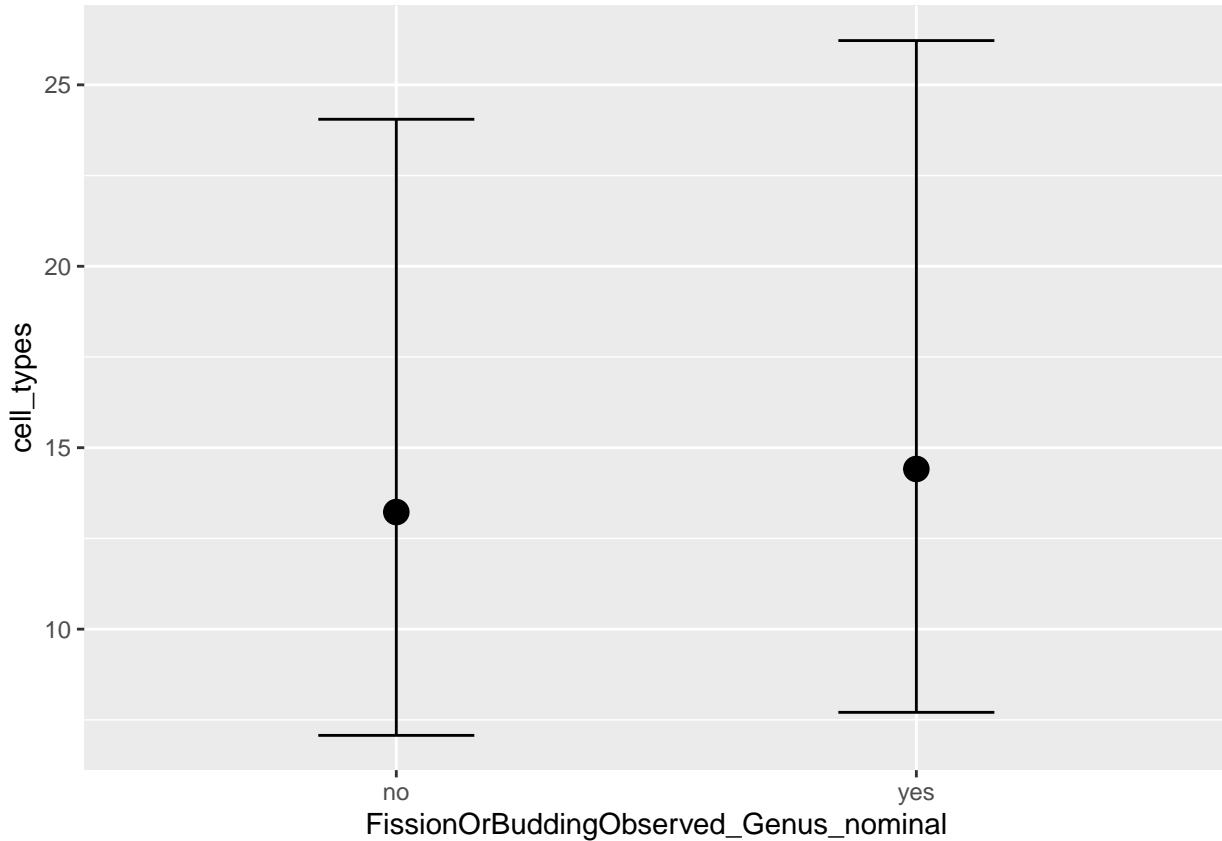
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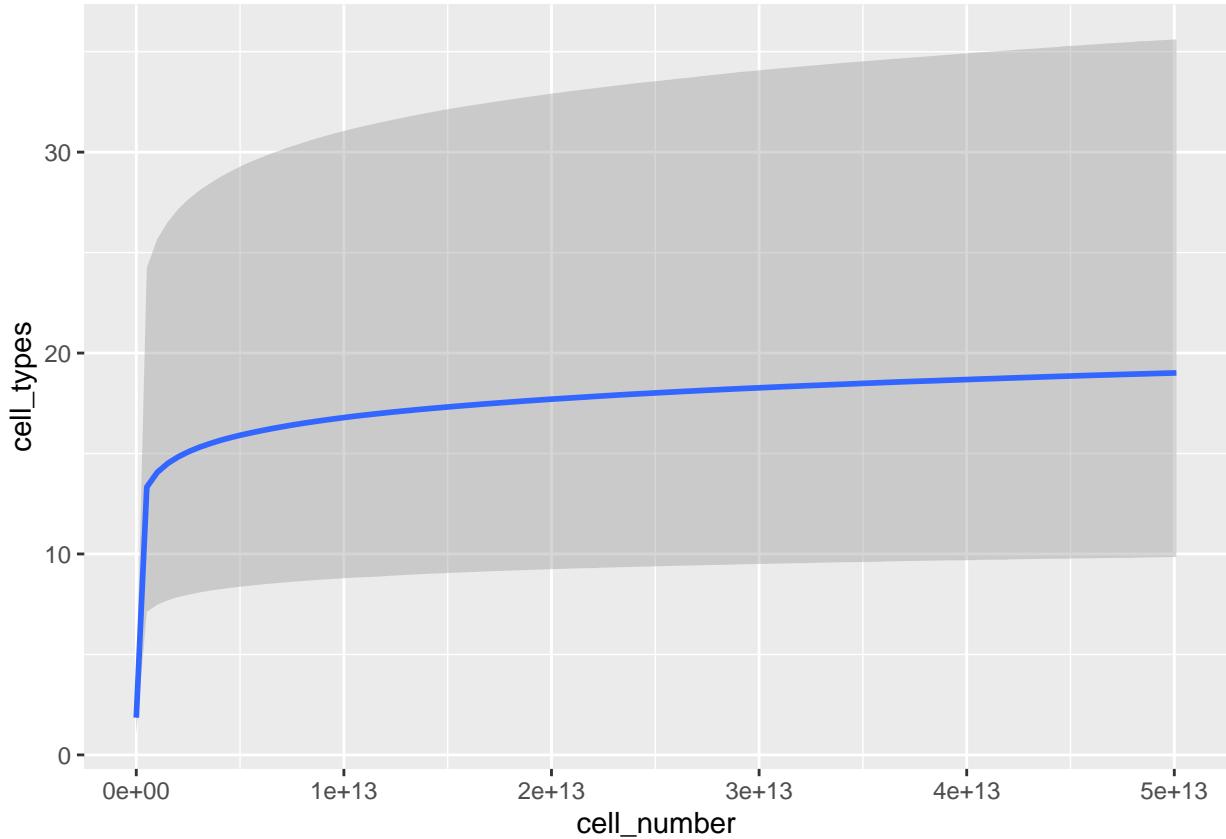
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## r_species_id[ott567703,Intercept]	-0.63138404	0.73581209
## r_species_id[ott570365,Intercept]	0.03025473	1.45846591
## r_species_id[ott570656,Intercept]	0.24602508	1.60250875
## r_species_id[ott588761,Intercept]	0.41574981	1.77535901
## r_species_id[ott592355,Intercept]	-0.23632126	1.14243191
## r_species_id[ott601255,Intercept]	-1.59412895	-0.04043146
## r_species_id[ott602180,Intercept]	-1.09704273	0.60984004
## r_species_id[ott60470,Intercept]	-1.38834310	0.23207451
## r_species_id[ott60473,Intercept]	-1.37317344	0.23147113
## r_species_id[ott60479,Intercept]	-1.26760724	0.25514981
## r_species_id[ott633708,Intercept]	-1.30118089	0.38900256
## r_species_id[ott633710,Intercept]	-1.23677470	0.26369651
## r_species_id[ott633711,Intercept]	-1.33297310	0.35071271
## r_species_id[ott633717,Intercept]	-1.29904514	0.49538443
## r_species_id[ott633719,Intercept]	-1.23374461	0.27522232
## r_species_id[ott643237,Intercept]	0.62644994	1.93118883
## r_species_id[ott645555,Intercept]	-0.90802714	0.49156172
## r_species_id[ott649193,Intercept]	-1.49149031	0.04658647
## r_species_id[ott675301,Intercept]	0.72806158	2.09556900
## r_species_id[ott724784,Intercept]	-0.65147670	0.74264223
## r_species_id[ott72522,Intercept]	-0.92380656	0.50722384
## r_species_id[ott727979,Intercept]	-1.60748581	-0.03082879
## r_species_id[ott733462,Intercept]	-0.41782073	0.98231458
## r_species_id[ott736728,Intercept]	-0.84225179	0.71460551
## r_species_id[ott742128,Intercept]	0.21578808	1.54221763
## r_species_id[ott7489702,Intercept]	0.51028568	1.86821372
## r_species_id[ott7567530,Intercept]	0.03026486	1.44355950
## r_species_id[ott765113,Intercept]	-0.76656775	0.61722355
## r_species_id[ott765280,Intercept]	0.14758048	1.49805535
## r_species_id[ott779028,Intercept]	-0.74751544	0.63643802
## r_species_id[ott790395,Intercept]	-1.13620773	0.29980708
## r_species_id[ott817791,Intercept]	-1.02015806	0.44059896
## r_species_id[ott821356,Intercept]	0.29795671	1.62815496
## r_species_id[ott83430,Intercept]	-1.07038325	0.36886410
## r_species_id[ott83432,Intercept]	-1.06417982	0.37928588
## r_species_id[ott840001,Intercept]	0.59976537	1.90773366
## r_species_id[ott841027,Intercept]	-1.37060482	0.42003493
## r_species_id[ott849781,Intercept]	-1.29546499	0.28037609
## r_species_id[ott878345,Intercept]	-1.17033057	0.42114902
## r_species_id[ott92556,Intercept]	-0.85036469	0.56160472
## r_species_id[ott92561,Intercept]	-0.92251290	0.48844157
## r_species_id[ott939432,Intercept]	-1.12186313	0.33012440
## r_species_id[ott939454,Intercept]	-1.12197393	0.33259937
## r_species_id[ott954042,Intercept]	0.37468730	1.67690522
## r_species_id[ott958293,Intercept]	-1.29845250	0.33833056
## r_species_id[ott958304,Intercept]	-1.36665888	0.31634816
## r_species_id[ott962359,Intercept]	-1.08682718	0.55493372
## r_species_id[ott987480,Intercept]	0.79648474	2.17846157

```
## lp_--
```

```
-598.80436333 -556.01021506
```

```
plot(conditional_effects(phylo_fit_fission_cell_type, points = TRUE, ask = F))
```

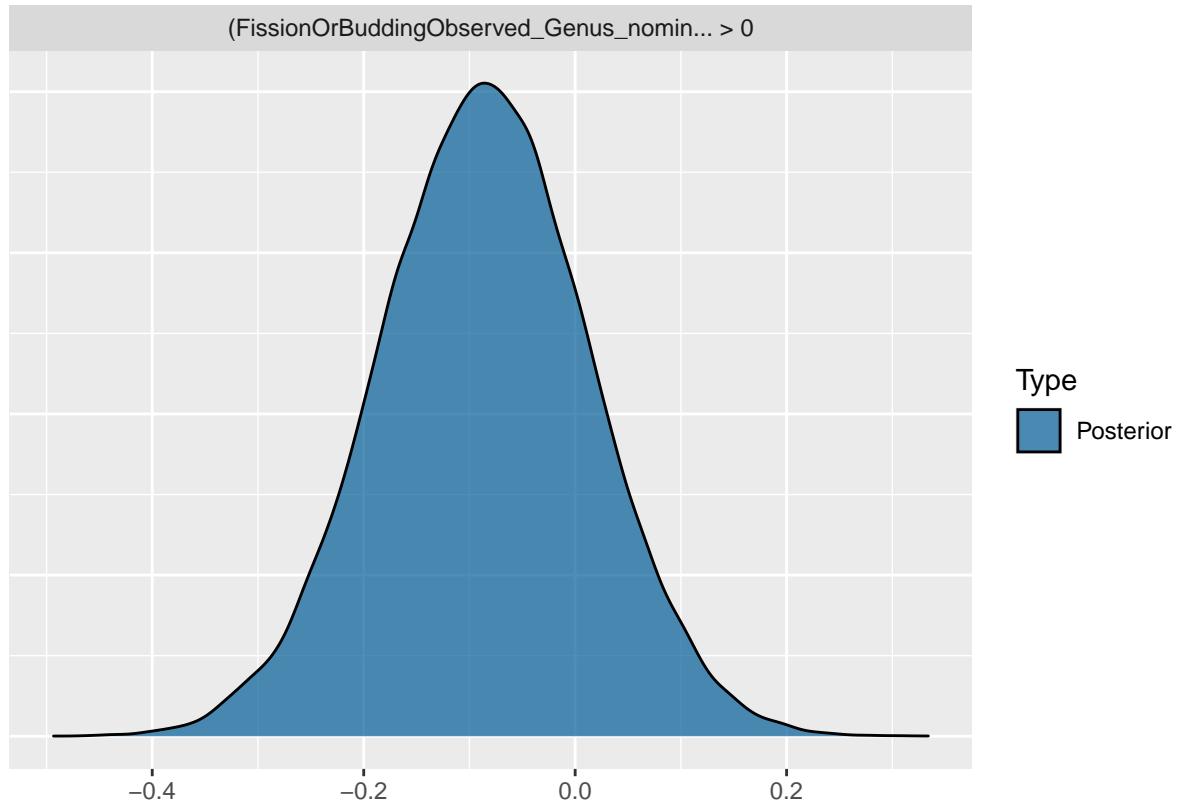




```
hyp = hypothesis(phylo_fit_fission_cell_type, "FissionOrBuddingObserved_Genus_nominalno > FissionOrBud
hyp
```

```
## Hypothesis Tests for class b:
##          Hypothesis Estimate Est.Error CI.Lower CI.Upper Evid.Ratio
## 1 (FissionOrBudding... > 0     -0.09      0.1    -0.25     0.08     0.23
##   Post.Prob Star
## 1      0.19
## ---
## 'CI': 90%-CI for one-sided and 95%-CI for two-sided hypotheses.
## '*': For one-sided hypotheses, the posterior probability exceeds 95%;
## for two-sided hypotheses, the value tested against lies outside the 95%-CI.
## Posterior probabilities of point hypotheses assume equal prior probabilities.
```

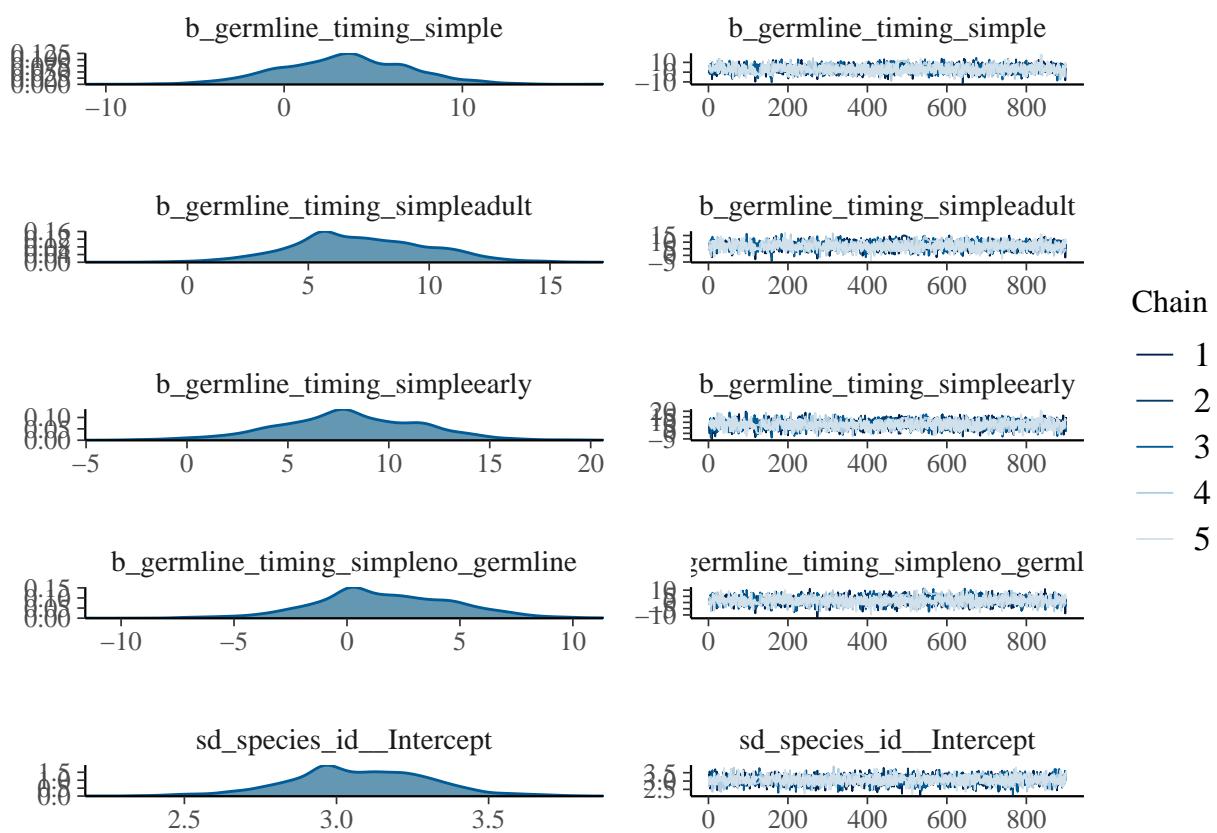
```
plot(hyp)
```

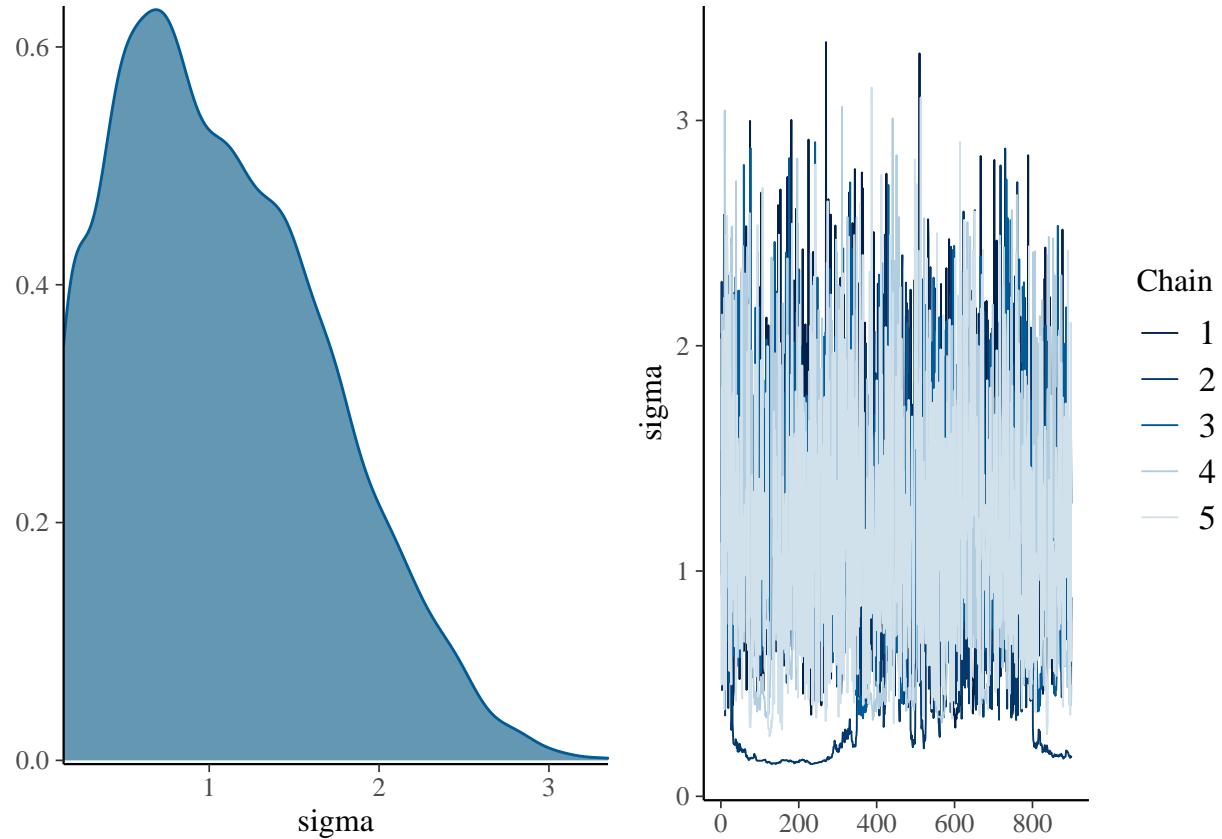


Does germline timing correlate with increased cell number?

```
#fit the model
phylofit_germline_cell_num<-
  brm(data = df,
       family=gaussian(), # family of model
       formula = log(cell_number) ~ 0 + germline_timing_simple + (1|gr(species_id, cov = CovarMatrix)),
       data2 = list(CovarMatrix = CovarMatrix) ,
       iter = 1000000, warmup = 100000, chains = 5,thin = 1000, cores = 5, #chain settings
       prior = prior(normal(0, 10), "b"), #defining the priors- for things in 'class b' set this prior.
       file = 'fits/fit_phy_GermCellNum' )

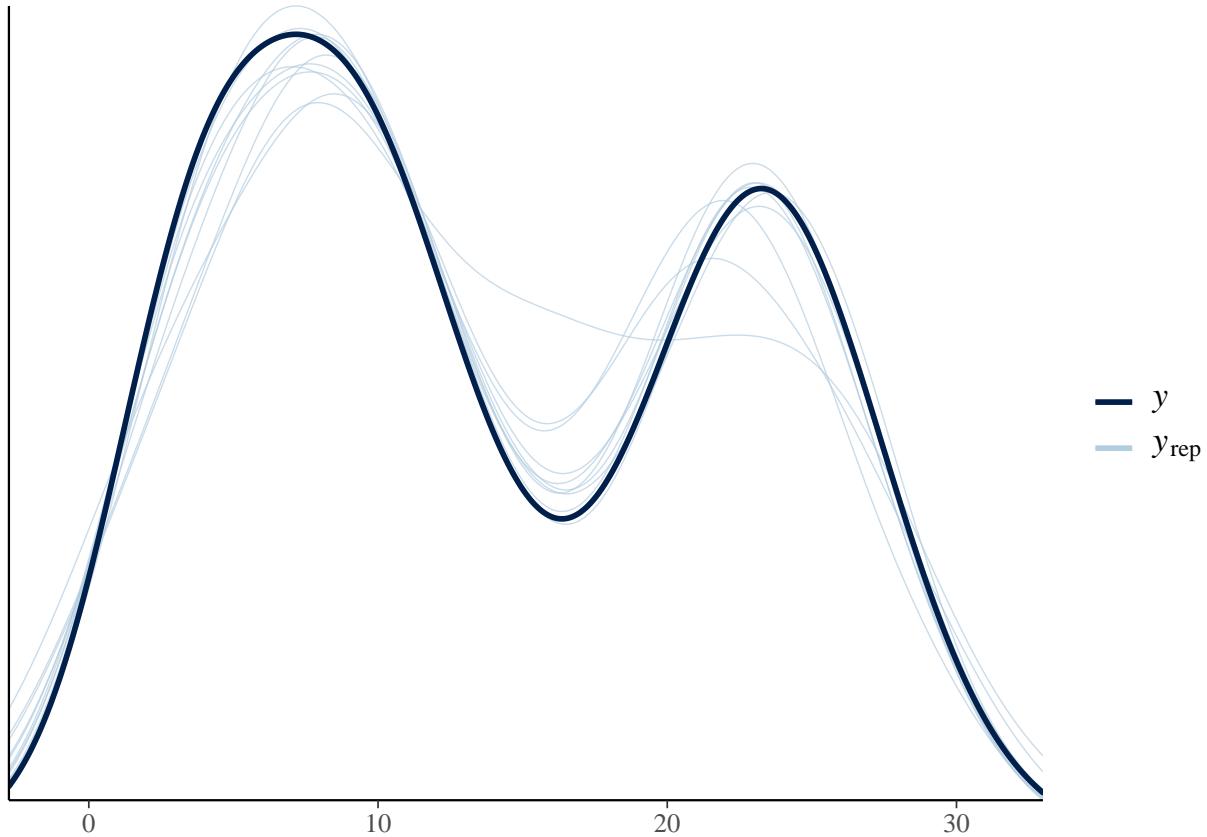
plot(phylofit_germline_cell_num) #check that chains converged
```



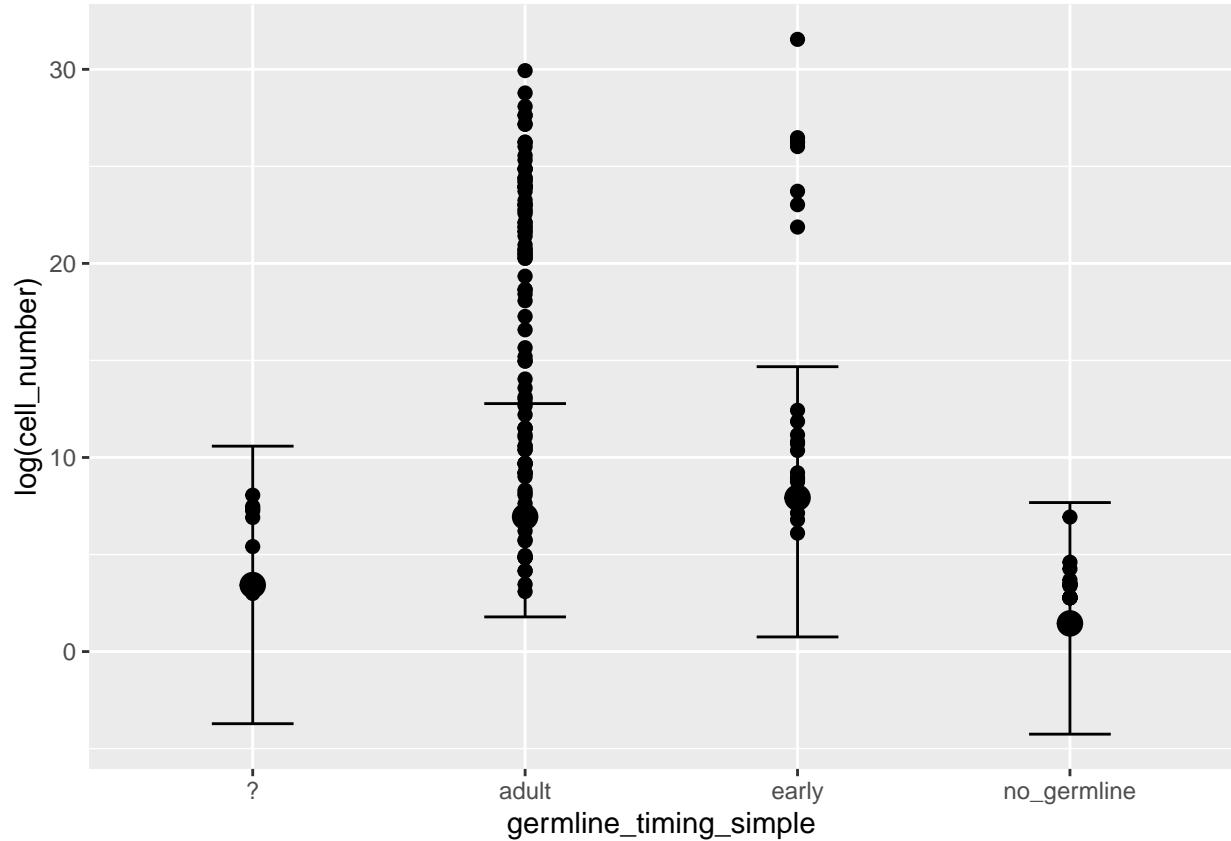


```
pp_check(phylofit_germline_cell_num) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



```
plot(conditional_effects(phylofit_germline_cell_num), points = TRUE)
```



```
summary(phylofit_germline_cell_num)
```

```
## Warning: Parts of the model have not converged (some Rhats are > 1.05). Be
## careful when analysing the results! We recommend running more iterations and/or
## setting stronger priors.

## Warning: There were 1219 divergent transitions after warmup.
## Increasing adapt_delta above 0.8 may help. See http://mc-stan.org/misc/
## warnings.html#divergent-transitions-after-warmup

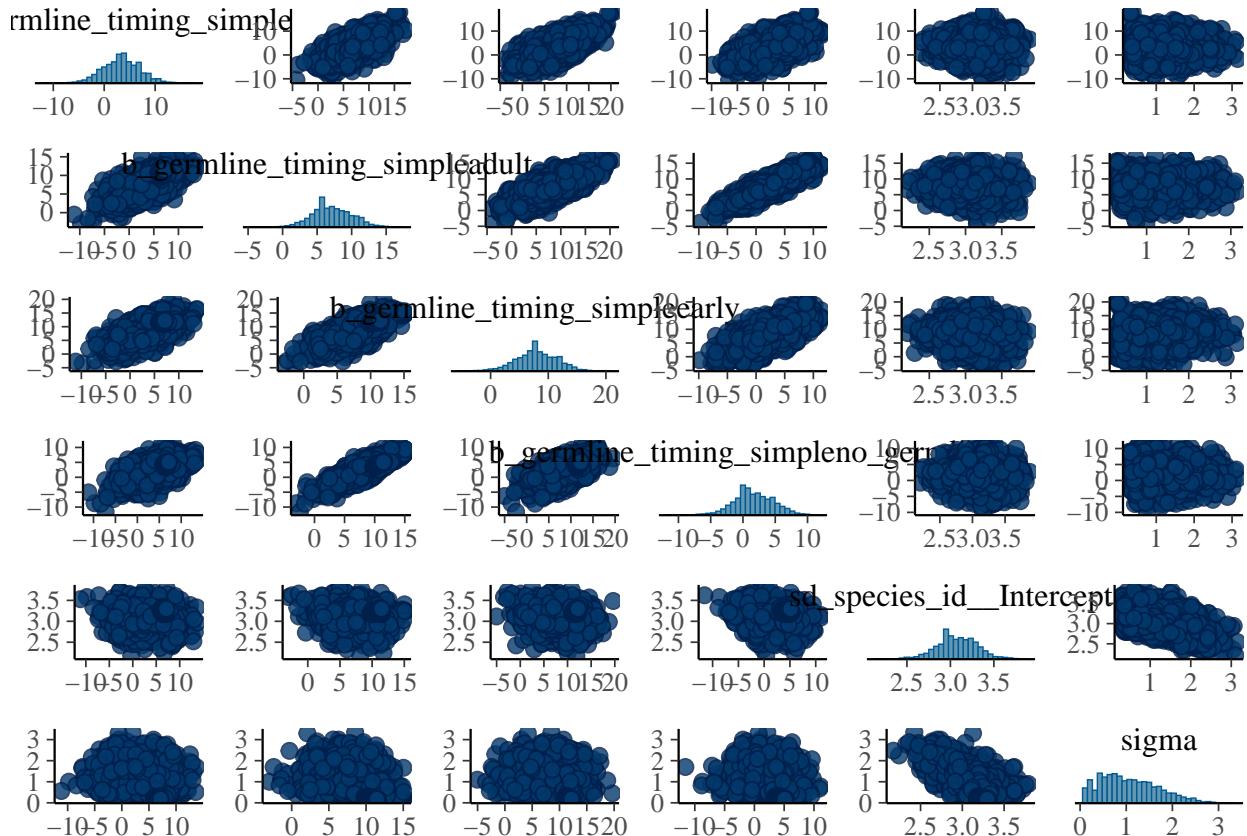
## Family: gaussian
##   Links: mu = identity; sigma = identity
## Formula: log(cell_number) ~ 0 + germline_timing_simple + (1 | gr(species_id, cov = CovarMatrix))
##   Data: df (Number of observations: 158)
## Samples: 5 chains, each with iter = 1e+06; warmup = 1e+05; thin = 1000;
##           total post-warmup samples = 4500
##
## Group-Level Effects:
## ~species_id (Number of levels: 158)
##             Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)    3.07     0.23    2.60    3.54 1.01      979     1539
##
## Population-Level Effects:
##             Estimate Est.Error 1-95% CI u-95% CI Rhat
## germline_timing_simple          3.39     3.62    -3.72    10.58 1.05
```

```

## germline_timing_simpleadult      7.16      2.85      1.79      12.78 1.01
## germline_timing_simpleearly     8.03      3.54      0.75      14.68 1.07
## germline_timing_simpleno_germline 1.67      3.06     -4.26      7.68 1.01
##                                         Bulk_ESS Tail_ESS
## germline_timing_simple          733       756
## germline_timing_simpleadult    454      1575
## germline_timing_simpleearly    908      1652
## germline_timing_simpleno_germline 516       604
##
## Family Specific Parameters:
##             Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sigma      1.07      0.62   0.16     2.42 1.13        26       15
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).

```

```
pairs(phylofit_germline_cell_num)
```



```
posterior_summary(phylofit_germline_cell_num, robust = T) #what are the medians for coefficients? if F,
```

	Estimate	Est.Error	Q2.5
## b_germline_timing_simple	3.42609702	3.6456708	-3.7159027
## b_germline_timing_simpleadult	6.94899423	2.7580361	1.7860413
## b_germline_timing_simpleearly	7.92744534	3.4688554	0.7547506

## b_germline_timing_simpleno_germline	1.45206392	2.9379109	-4.2551620
## sd_species_id__Intercept	3.06991955	0.2213336	2.6024103
## sigma	0.99502613	0.6726738	0.1572766
## r_species_id[ott1002450,Intercept]	1.99248457	3.1427736	-4.6126736
## r_species_id[ott1017821,Intercept]	-2.44198237	2.8902686	-8.7713890
## r_species_id[ott1052546,Intercept]	-2.49751459	2.9512065	-8.6516385
## r_species_id[ott1059898,Intercept]	2.69648232	2.9565248	-3.5877554
## r_species_id[ott1059900,Intercept]	4.42144197	2.8602242	-1.7698357
## r_species_id[ott1061937,Intercept]	5.21777302	3.2056694	-1.2970632
## r_species_id[ott1069171,Intercept]	12.89776745	3.0907144	6.3493268
## r_species_id[ott1072227,Intercept]	1.36479575	3.1496192	-5.2944196
## r_species_id[ott108923,Intercept]	6.42116650	2.8962205	0.0585727
## r_species_id[ott1099013,Intercept]	16.71123899	2.9976380	10.3016515
## r_species_id[ott111442,Intercept]	4.14997692	2.9174191	-2.3898715
## r_species_id[ott112015,Intercept]	1.38380173	3.1223632	-5.3487446
## r_species_id[ott112016,Intercept]	1.32546634	3.1646132	-5.2844625
## r_species_id[ott112017,Intercept]	1.39743015	3.1020625	-5.2775496
## r_species_id[ott127047,Intercept]	-1.32046531	2.9488586	-7.6543353
## r_species_id[ott150272,Intercept]	0.91356133	3.6896357	-6.1488996
## r_species_id[ott160850,Intercept]	4.28155395	3.0200759	-2.0567609
## r_species_id[ott165368,Intercept]	18.26608355	3.6564003	11.0940846
## r_species_id[ott167121,Intercept]	1.97279873	3.1423902	-4.5767761
## r_species_id[ott178177,Intercept]	11.49194801	2.9418538	5.3123222
## r_species_id[ott178412,Intercept]	21.65296687	3.6160306	14.5234964
## r_species_id[ott181933,Intercept]	20.88945002	3.0706221	14.4472348
## r_species_id[ott182906,Intercept]	17.85575975	3.7717719	10.2898994
## r_species_id[ott186999,Intercept]	15.48258608	3.0682561	8.9880251
## r_species_id[ott187583,Intercept]	0.48420714	3.7079441	-6.7719007
## r_species_id[ott199292,Intercept]	8.15618309	2.9052406	1.7394949
## r_species_id[ott207134,Intercept]	14.82046344	3.0890899	8.2342038
## r_species_id[ott215125,Intercept]	17.04780517	3.0296852	10.6943253
## r_species_id[ott216694,Intercept]	20.57633699	2.9991104	14.2350655
## r_species_id[ott223669,Intercept]	18.32127823	3.5780760	11.1501061
## r_species_id[ott225275,Intercept]	15.95346285	3.0387438	9.6030852
## r_species_id[ott237608,Intercept]	13.61451538	3.1159507	7.2030857
## r_species_id[ott246046,Intercept]	2.86706788	2.9690903	-3.3822520
## r_species_id[ott247341,Intercept]	23.13567656	3.7908746	15.5790049
## r_species_id[ott256062,Intercept]	-1.85228487	3.0312856	-8.1407918
## r_species_id[ott256089,Intercept]	-2.73974852	2.9268494	-9.0407435
## r_species_id[ott256145,Intercept]	1.94302256	3.1780957	-4.6403117
## r_species_id[ott263960,Intercept]	13.47673814	2.9594704	7.1377876
## r_species_id[ott263980,Intercept]	13.44234583	2.8967691	7.0527766
## r_species_id[ott263987,Intercept]	10.20848233	2.9407667	3.8640036
## r_species_id[ott263988,Intercept]	17.08183159	2.9831368	10.8030290
## r_species_id[ott265121,Intercept]	18.61695575	3.2678123	12.0950826
## r_species_id[ott266342,Intercept]	-0.29619258	3.6732210	-7.3026446
## r_species_id[ott269063,Intercept]	-3.56660684	2.9687525	-9.9243034
## r_species_id[ott275893,Intercept]	13.30450586	2.9866750	6.9807543
## r_species_id[ott275897,Intercept]	13.50833109	3.0039604	7.0727087
## r_species_id[ott2810724,Intercept]	2.76668807	3.0895863	-3.6323004
## r_species_id[ott2819986,Intercept]	16.96108671	3.1398783	10.4588710
## r_species_id[ott2821097,Intercept]	15.69010390	2.9406497	9.3759467
## r_species_id[ott2844172,Intercept]	1.50774439	2.8509191	-4.6514498
## r_species_id[ott2844962,Intercept]	-0.57565840	3.7060614	-7.6647618

## r_species_id[ott2849837,Intercept]	3.46389802	3.6532630	-3.7215632
## r_species_id[ott2942244,Intercept]	3.90556403	3.5896535	-3.5817513
## r_species_id[ott316441,Intercept]	14.46891891	3.8060252	6.5829094
## r_species_id[ott33153,Intercept]	2.76654615	3.1500854	-3.8409630
## r_species_id[ott336388,Intercept]	20.19959878	2.9828704	13.8594616
## r_species_id[ott34559,Intercept]	15.93241800	3.0438044	9.6030731
## r_species_id[ott346740,Intercept]	17.62853833	3.0595821	11.1997227
## r_species_id[ott3583594,Intercept]	1.25776392	3.6156530	-5.7921409
## r_species_id[ott3587677,Intercept]	-1.60144594	3.6725880	-8.7130917
## r_species_id[ott359012,Intercept]	1.15370950	3.1317373	-5.5065089
## r_species_id[ott361837,Intercept]	1.38683611	3.1577040	-5.1174253
## r_species_id[ott362913,Intercept]	3.64090147	2.9667473	-2.5892624
## r_species_id[ott365439,Intercept]	2.64362445	3.6546246	-4.4362197
## r_species_id[ott3663378,Intercept]	11.47421845	3.0462018	5.1049713
## r_species_id[ott3665433,Intercept]	5.81339435	2.9501901	-0.6665857
## r_species_id[ott3684291,Intercept]	3.68059006	3.6682166	-3.6402663
## r_species_id[ott3684365,Intercept]	3.68524142	3.8216736	-3.7348955
## r_species_id[ott3684379,Intercept]	-0.26437453	3.7188512	-7.7094875
## r_species_id[ott3684389,Intercept]	1.87115411	3.7640400	-5.5216986
## r_species_id[ott3684437,Intercept]	0.10531841	3.7667654	-7.3671871
## r_species_id[ott381979,Intercept]	0.07545832	3.1714416	-6.4692737
## r_species_id[ott381980,Intercept]	3.15061568	3.0181040	-3.0659893
## r_species_id[ott381983,Intercept]	1.42510730	3.1601900	-5.2907148
## r_species_id[ott395048,Intercept]	-0.91550393	3.6129023	-8.1141451
## r_species_id[ott3974169,Intercept]	16.42041740	2.9891857	10.1048277
## r_species_id[ott3995126,Intercept]	17.25348914	2.9784545	10.9391138
## r_species_id[ott4010019,Intercept]	16.89671441	3.0527541	10.5078719
## r_species_id[ott4010960,Intercept]	17.51016712	3.0617215	11.2412477
## r_species_id[ott4011155,Intercept]	9.62347559	2.9721812	3.3378038
## r_species_id[ott4013437,Intercept]	14.71955896	2.9335587	8.3782738
## r_species_id[ott4013674,Intercept]	18.70181814	3.2871587	12.1143216
## r_species_id[ott4013684,Intercept]	17.51054434	3.0075544	11.1886814
## r_species_id[ott422679,Intercept]	-0.97423142	2.9103515	-7.2017202
## r_species_id[ott431388,Intercept]	6.97739851	2.8877422	0.6814831
## r_species_id[ott446088,Intercept]	1.32319190	3.5773095	-5.7996969
## r_species_id[ott4741377,Intercept]	13.99963649	2.8507502	7.6265131
## r_species_id[ott4742064,Intercept]	18.51964072	3.1167009	12.1215465
## r_species_id[ott481952,Intercept]	17.76931948	3.1216327	11.3128112
## r_species_id[ott48288,Intercept]	13.97031156	3.6629563	6.6845715
## r_species_id[ott485470,Intercept]	0.80390618	2.9314680	-5.4854365
## r_species_id[ott485473,Intercept]	-1.94982316	2.9416871	-8.3141669
## r_species_id[ott485476,Intercept]	2.03448142	3.1948426	-4.5187330
## r_species_id[ott485480,Intercept]	2.85573507	2.8905770	-3.4734327
## r_species_id[ott485482,Intercept]	-0.19741340	3.6021613	-7.2245065
## r_species_id[ott486834,Intercept]	-0.48002539	2.9151835	-6.6600791
## r_species_id[ott490206,Intercept]	6.07762948	2.9605807	-0.2296249
## r_species_id[ott492241,Intercept]	0.41730912	3.5978065	-6.8361696
## r_species_id[ott497063,Intercept]	15.25999836	2.9443174	8.8406783
## r_species_id[ott4974308,Intercept]	3.53660223	3.7307520	-3.4657145
## r_species_id[ott4978773,Intercept]	0.37881144	3.5804178	-6.9353975
## r_species_id[ott4979583,Intercept]	1.00861474	3.6287590	-6.2045681
## r_species_id[ott518643,Intercept]	2.75854245	3.6555856	-4.4680771
## r_species_id[ott542509,Intercept]	18.18213933	3.6517812	11.1272922
## r_species_id[ott54768,Intercept]	8.18420835	2.8923099	1.9507345

## r_species_id[ott549846,Intercept]	4.74083427	3.6619935	-2.4375890
## r_species_id[ott560703,Intercept]	-1.63007544	2.9447662	-7.8502168
## r_species_id[ott567703,Intercept]	16.55500608	2.9938550	10.2395458
## r_species_id[ott570365,Intercept]	-0.59361246	2.9089921	-6.8047860
## r_species_id[ott570656,Intercept]	10.74616928	2.8642878	4.3919411
## r_species_id[ott588761,Intercept]	2.33120224	2.8842811	-3.8938261
## r_species_id[ott592355,Intercept]	13.40483816	3.0234342	6.9420724
## r_species_id[ott601255,Intercept]	21.69473354	3.0268391	15.5007789
## r_species_id[ott602180,Intercept]	2.93457208	3.1365028	-3.5999943
## r_species_id[ott60470,Intercept]	1.39618962	3.1203269	-5.2535733
## r_species_id[ott60471,Intercept]	0.16468220	3.1121325	-6.5743876
## r_species_id[ott60473,Intercept]	1.35678423	3.0928190	-5.2429519
## r_species_id[ott60477,Intercept]	1.97370612	3.1279173	-4.5848178
## r_species_id[ott60479,Intercept]	1.96210543	3.1961951	-4.6032556
## r_species_id[ott633708,Intercept]	1.27321610	2.9592504	-5.0549390
## r_species_id[ott633710,Intercept]	3.10589836	3.0613512	-3.2264370
## r_species_id[ott633711,Intercept]	2.19313503	3.0130682	-4.1751793
## r_species_id[ott633717,Intercept]	-1.75346194	2.9318813	-7.8962363
## r_species_id[ott633719,Intercept]	2.51462638	2.9838484	-3.4990162
## r_species_id[ott643237,Intercept]	16.96590634	2.9280293	10.6062376
## r_species_id[ott645555,Intercept]	18.04690989	3.1404421	11.5311445
## r_species_id[ott649193,Intercept]	17.15458996	2.9779022	10.9212025
## r_species_id[ott675301,Intercept]	4.57942842	3.7989146	-2.9202518
## r_species_id[ott724784,Intercept]	9.19116642	2.9593729	2.7569315
## r_species_id[ott72522,Intercept]	20.04617326	3.1756108	13.6771267
## r_species_id[ott727979,Intercept]	20.20716137	2.9584707	14.0357246
## r_species_id[ott733462,Intercept]	15.32888217	2.9895088	8.9487199
## r_species_id[ott736728,Intercept]	6.04643791	2.8926198	-0.2394106
## r_species_id[ott742128,Intercept]	2.37489489	2.9415381	-3.8099022
## r_species_id[ott7489702,Intercept]	3.31325834	3.6790343	-3.9640158
## r_species_id[ott7567530,Intercept]	4.33282817	2.9681354	-1.9628493
## r_species_id[ott765113,Intercept]	-0.02461901	2.8583603	-6.4118349
## r_species_id[ott765280,Intercept]	12.41985895	2.8923847	6.1641610
## r_species_id[ott779028,Intercept]	14.97761511	2.9209996	8.7158426
## r_species_id[ott790395,Intercept]	15.89123643	2.9848584	9.6201912
## r_species_id[ott817791,Intercept]	14.63765678	2.9837927	8.4801986
## r_species_id[ott821356,Intercept]	13.71334146	2.9260621	7.5753622
## r_species_id[ott83430,Intercept]	11.93109994	2.8644073	5.5938699
## r_species_id[ott83432,Intercept]	4.08223448	3.1574516	-2.1933020
## r_species_id[ott840001,Intercept]	16.97426602	2.9690336	10.5690754
## r_species_id[ott841027,Intercept]	-0.21681980	3.6640707	-7.3510218
## r_species_id[ott849781,Intercept]	14.31689554	3.0814300	7.8551707
## r_species_id[ott878345,Intercept]	6.47210781	2.9256936	0.1275615
## r_species_id[ott92556,Intercept]	11.79522305	2.9340454	5.4919601
## r_species_id[ott92561,Intercept]	14.53863187	3.0413480	8.1480597
## r_species_id[ott939432,Intercept]	4.43490975	2.9992834	-1.6122673
## r_species_id[ott939454,Intercept]	4.42285415	2.9029234	-1.8886801
## r_species_id[ott954042,Intercept]	15.17407791	3.8380761	7.4936067
## r_species_id[ott958293,Intercept]	1.92558216	3.1350384	-4.6770866
## r_species_id[ott958304,Intercept]	1.33031325	3.1731860	-5.2235961
## r_species_id[ott962359,Intercept]	1.34967534	2.9298475	-5.0495190
## r_species_id[ott987480,Intercept]	4.09015188	3.7093487	-3.3000607
## r_species_id[ott989764,Intercept]	6.74657781	2.9847310	0.3861690
## lp_	-464.69202078	107.0026800	-603.9722980

##	Q97.5
## b_germline_timing_simple	10.584152
## b_germline_timing_simpleadult	12.777312
## b_germline_timing_simpleearly	14.678657
## b_germline_timing_simpleno_germline	7.678403
## sd_species_id_Intercept	3.540341
## sigma	2.421592
## r_species_id[ott1002450,Intercept]	8.080142
## r_species_id[ott1017821,Intercept]	3.096677
## r_species_id[ott1052546,Intercept]	3.092212
## r_species_id[ott1059898,Intercept]	8.360771
## r_species_id[ott1059900,Intercept]	10.018602
## r_species_id[ott1061937,Intercept]	11.290418
## r_species_id[ott1069171,Intercept]	18.360529
## r_species_id[ott1072227,Intercept]	7.485817
## r_species_id[ott108923,Intercept]	12.115871
## r_species_id[ott1099013,Intercept]	22.360271
## r_species_id[ott111442,Intercept]	9.765671
## r_species_id[ott112015,Intercept]	7.414733
## r_species_id[ott112016,Intercept]	7.569822
## r_species_id[ott112017,Intercept]	7.568846
## r_species_id[ott127047,Intercept]	4.486614
## r_species_id[ott150272,Intercept]	8.339389
## r_species_id[ott160850,Intercept]	10.036152
## r_species_id[ott165368,Intercept]	25.491013
## r_species_id[ott167121,Intercept]	8.106051
## r_species_id[ott178177,Intercept]	16.824146
## r_species_id[ott178412,Intercept]	27.739002
## r_species_id[ott181933,Intercept]	26.518745
## r_species_id[ott182906,Intercept]	25.421414
## r_species_id[ott186999,Intercept]	20.964725
## r_species_id[ott187583,Intercept]	7.796460
## r_species_id[ott199292,Intercept]	13.881953
## r_species_id[ott207134,Intercept]	20.395623
## r_species_id[ott215125,Intercept]	22.608542
## r_species_id[ott216694,Intercept]	26.236077
## r_species_id[ott223669,Intercept]	25.674663
## r_species_id[ott225275,Intercept]	21.583097
## r_species_id[ott237608,Intercept]	19.229417
## r_species_id[ott246046,Intercept]	8.558250
## r_species_id[ott247341,Intercept]	30.538422
## r_species_id[ott256062,Intercept]	3.632911
## r_species_id[ott256089,Intercept]	2.963400
## r_species_id[ott256145,Intercept]	8.124576
## r_species_id[ott263960,Intercept]	18.978043
## r_species_id[ott263980,Intercept]	19.045404
## r_species_id[ott263987,Intercept]	15.909342
## r_species_id[ott263988,Intercept]	22.690679
## r_species_id[ott265121,Intercept]	24.221400
## r_species_id[ott266342,Intercept]	7.269712
## r_species_id[ott269063,Intercept]	2.086037
## r_species_id[ott275893,Intercept]	18.882209
## r_species_id[ott275897,Intercept]	19.213665
## r_species_id[ott2810724,Intercept]	9.034601

```

## r_species_id[ott2819986,Intercept]      22.726545
## r_species_id[ott2821097,Intercept]      21.355180
## r_species_id[ott2844172,Intercept]      7.200757
## r_species_id[ott2844962,Intercept]      6.831467
## r_species_id[ott2849837,Intercept]      10.873544
## r_species_id[ott2942244,Intercept]      11.086764
## r_species_id[ott316441,Intercept]       21.974262
## r_species_id[ott33153,Intercept]        8.871727
## r_species_id[ott336388,Intercept]       25.708741
## r_species_id[ott34559,Intercept]        21.650336
## r_species_id[ott346740,Intercept]       23.188794
## r_species_id[ott3583594,Intercept]      8.921274
## r_species_id[ott3587677,Intercept]      5.917915
## r_species_id[ott359012,Intercept]       7.242999
## r_species_id[ott361837,Intercept]       7.531543
## r_species_id[ott362913,Intercept]       9.283738
## r_species_id[ott365439,Intercept]       9.964813
## r_species_id[ott3663378,Intercept]      17.027642
## r_species_id[ott3665433,Intercept]      11.271034
## r_species_id[ott3684291,Intercept]      10.977956
## r_species_id[ott3684365,Intercept]      11.288833
## r_species_id[ott3684379,Intercept]      7.490038
## r_species_id[ott3684389,Intercept]      9.599077
## r_species_id[ott3684437,Intercept]      7.544370
## r_species_id[ott381979,Intercept]       6.104060
## r_species_id[ott381980,Intercept]       8.687702
## r_species_id[ott381983,Intercept]       7.421631
## r_species_id[ott395048,Intercept]       6.611994
## r_species_id[ott3974169,Intercept]      22.042064
## r_species_id[ott3995126,Intercept]      22.965367
## r_species_id[ott4010019,Intercept]      22.548945
## r_species_id[ott4010960,Intercept]      23.248067
## r_species_id[ott4011155,Intercept]      15.354980
## r_species_id[ott4013437,Intercept]      20.304470
## r_species_id[ott4013674,Intercept]      24.359384
## r_species_id[ott4013684,Intercept]      23.145865
## r_species_id[ott422679,Intercept]       4.738644
## r_species_id[ott431388,Intercept]       12.553543
## r_species_id[ott446088,Intercept]       8.858681
## r_species_id[ott4741377,Intercept]      19.513781
## r_species_id[ott4742064,Intercept]      24.255354
## r_species_id[ott481952,Intercept]       23.459936
## r_species_id[ott48288,Intercept]        21.313354
## r_species_id[ott485470,Intercept]       6.318466
## r_species_id[ott485473,Intercept]       3.663854
## r_species_id[ott485476,Intercept]       8.266321
## r_species_id[ott485480,Intercept]       8.371440
## r_species_id[ott485482,Intercept]       7.142573
## r_species_id[ott486834,Intercept]       5.051734
## r_species_id[ott490206,Intercept]       11.590345
## r_species_id[ott492241,Intercept]       7.724865
## r_species_id[ott497063,Intercept]       20.789252
## r_species_id[ott4974308,Intercept]      11.307968
## r_species_id[ott4978773,Intercept]      7.895545

```

```

## r_species_id[ott4979583,Intercept]      8.408380
## r_species_id[ott518643,Intercept]      10.140600
## r_species_id[ott542509,Intercept]      25.670022
## r_species_id[ott54768,Intercept]       13.626398
## r_species_id[ott549846,Intercept]      12.427059
## r_species_id[ott560703,Intercept]      4.096789
## r_species_id[ott567703,Intercept]      22.038235
## r_species_id[ott570365,Intercept]      5.349493
## r_species_id[ott570656,Intercept]      16.363404
## r_species_id[ott588761,Intercept]      8.044289
## r_species_id[ott592355,Intercept]      18.994390
## r_species_id[ott601255,Intercept]      27.258008
## r_species_id[ott602180,Intercept]      9.010865
## r_species_id[ott60470,Intercept]       7.548191
## r_species_id[ott60471,Intercept]       6.270965
## r_species_id[ott60473,Intercept]       7.631676
## r_species_id[ott60477,Intercept]       8.200983
## r_species_id[ott60479,Intercept]       8.055455
## r_species_id[ott633708,Intercept]      7.047696
## r_species_id[ott633710,Intercept]      8.671270
## r_species_id[ott633711,Intercept]      7.744483
## r_species_id[ott633717,Intercept]      3.987609
## r_species_id[ott633719,Intercept]      8.202503
## r_species_id[ott643237,Intercept]      22.423775
## r_species_id[ott645555,Intercept]      23.765726
## r_species_id[ott649193,Intercept]      22.966079
## r_species_id[ott675301,Intercept]      11.922598
## r_species_id[ott724784,Intercept]      15.069359
## r_species_id[ott72522,Intercept]       25.684691
## r_species_id[ott727979,Intercept]      25.941179
## r_species_id[ott733462,Intercept]      21.054448
## r_species_id[ott736728,Intercept]      11.946389
## r_species_id[ott742128,Intercept]      7.965908
## r_species_id[ott7489702,Intercept]     10.748996
## r_species_id[ott7567530,Intercept]    10.175107
## r_species_id[ott765113,Intercept]      5.536461
## r_species_id[ott765280,Intercept]      18.051418
## r_species_id[ott779028,Intercept]      20.673285
## r_species_id[ott790395,Intercept]      21.519641
## r_species_id[ott817791,Intercept]      20.208496
## r_species_id[ott821356,Intercept]      19.353658
## r_species_id[ott83430,Intercept]       17.731100
## r_species_id[ott83432,Intercept]       11.018759
## r_species_id[ott840001,Intercept]      22.610799
## r_species_id[ott841027,Intercept]      7.302418
## r_species_id[ott849781,Intercept]      19.963530
## r_species_id[ott878345,Intercept]      12.482267
## r_species_id[ott92556,Intercept]       17.358841
## r_species_id[ott92561,Intercept]       20.181519
## r_species_id[ott939432,Intercept]      10.125099
## r_species_id[ott939454,Intercept]      9.987044
## r_species_id[ott954042,Intercept]      22.653094
## r_species_id[ott958293,Intercept]      7.839058
## r_species_id[ott958304,Intercept]      7.348908

```

```

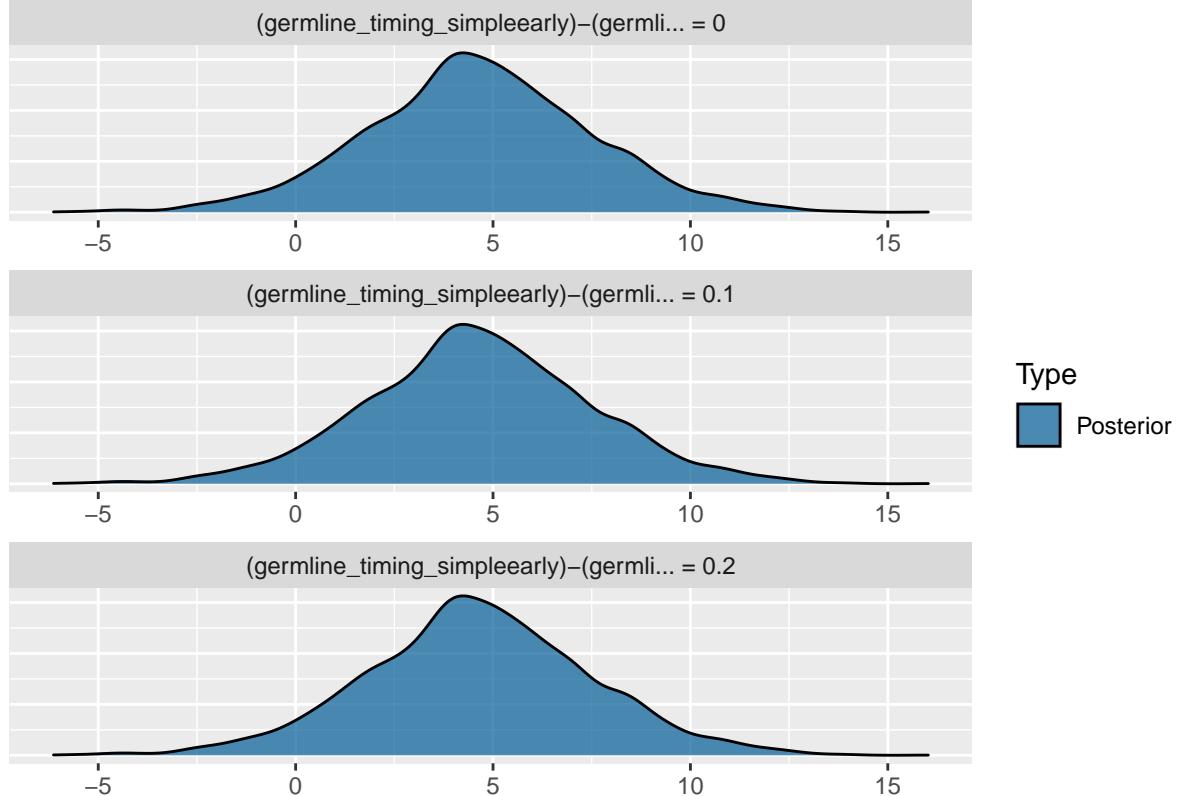
## r_species_id[ott962359,Intercept]      7.056204
## r_species_id[ott987480,Intercept]      11.605380
## r_species_id[ott989764,Intercept]      12.362309
## lp__                                -185.063333

hyp = hypothesis(phylofit_germline_cell_num, c("germline_timing_simpleearly = germline_timing_simple",
hyp

## Hypothesis Tests for class b:
##          Hypothesis Estimate Est.Error CI.Lower CI.Upper Evid.Ratio
## 1 (germline_timing_simpleearly = 0     4.64      2.86    -1.12    10.49      NA
## 2 (germline_timing_simpleearly = 0     0.87      2.39    -4.02     5.51      NA
## 3 (germline_timing_simpleearly = 0     6.36      2.71     0.74    11.56      NA
##   Post.Prob Star
## 1        NA
## 2        NA
## 3        NA      *
## ---
## 'CI': 90%-CI for one-sided and 95%-CI for two-sided hypotheses.
## '*': For one-sided hypotheses, the posterior probability exceeds 95%;
## for two-sided hypotheses, the value tested against lies outside the 95%-CI.
## Posterior probabilities of point hypotheses assume equal prior probabilities.

plot(hyp)

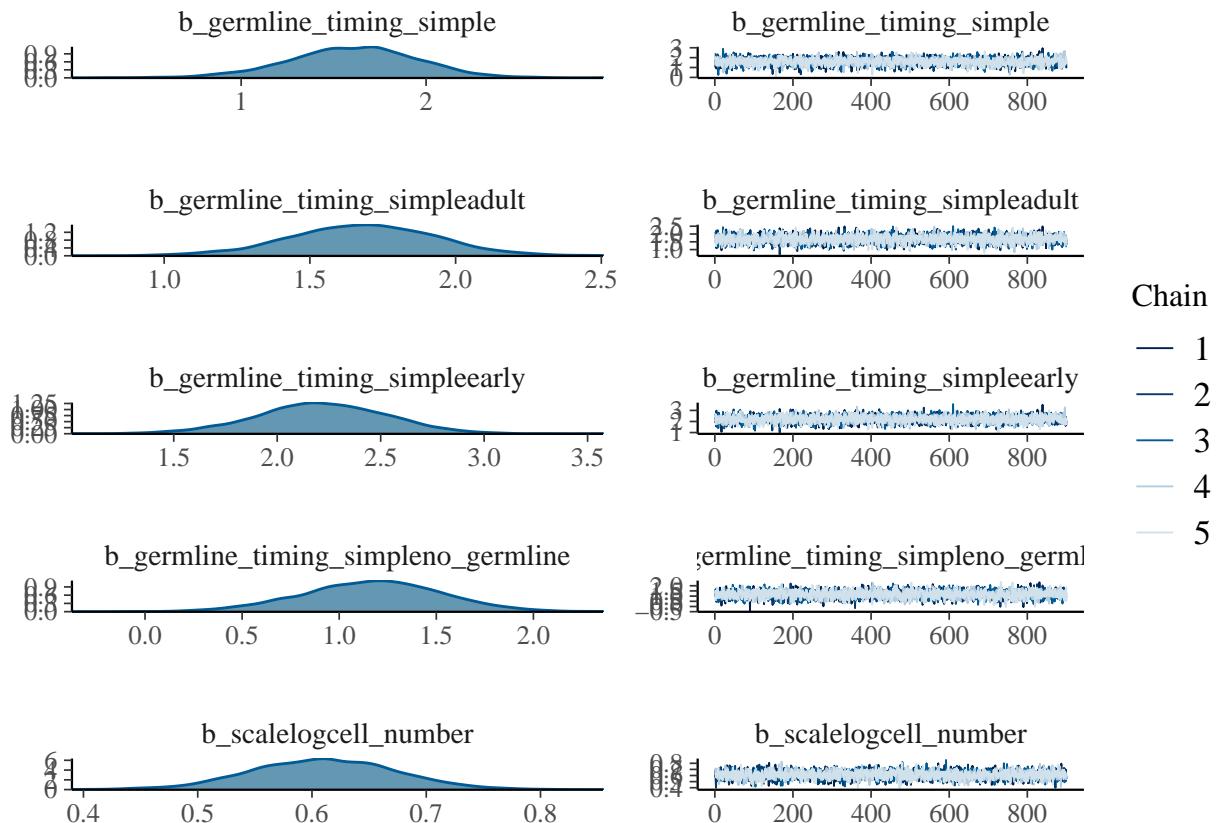
```

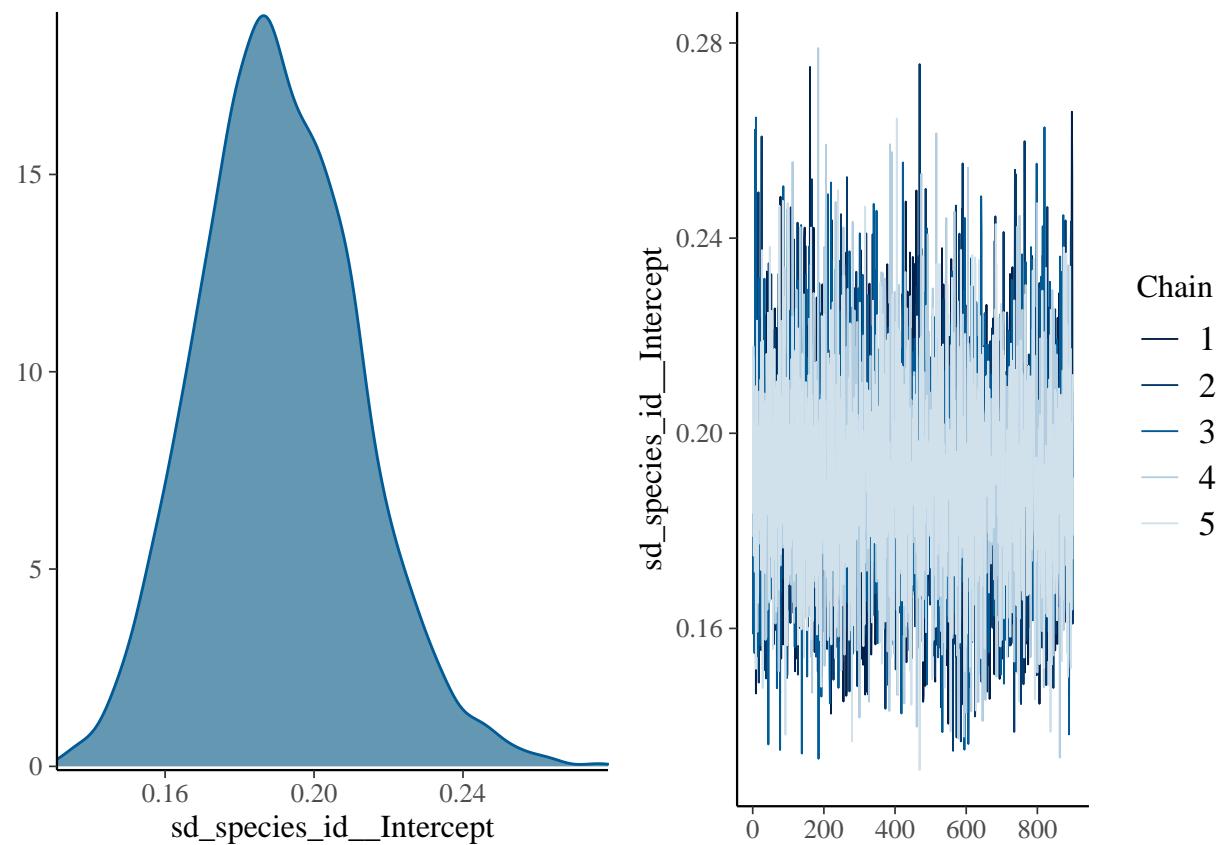


Does germline timing correlate with increased cell types (per cell)?

```
fit_type_phy<-
  brm(data = df,
      family=poisson(),
      formula = cell_types ~ 0 + germline_timing_simple + scale(log(cell_number)) + (1|gr(species_id, co
      iter = 1000000, warmup = 100000, chains = 5, thin = 1000, cores = 5,
      prior = prior(normal(0, 10), "b"), file = 'fits/fit_phy_GermCellType', # same simple prior
      data2 = list(CovarMatrix = CovarMatrix))

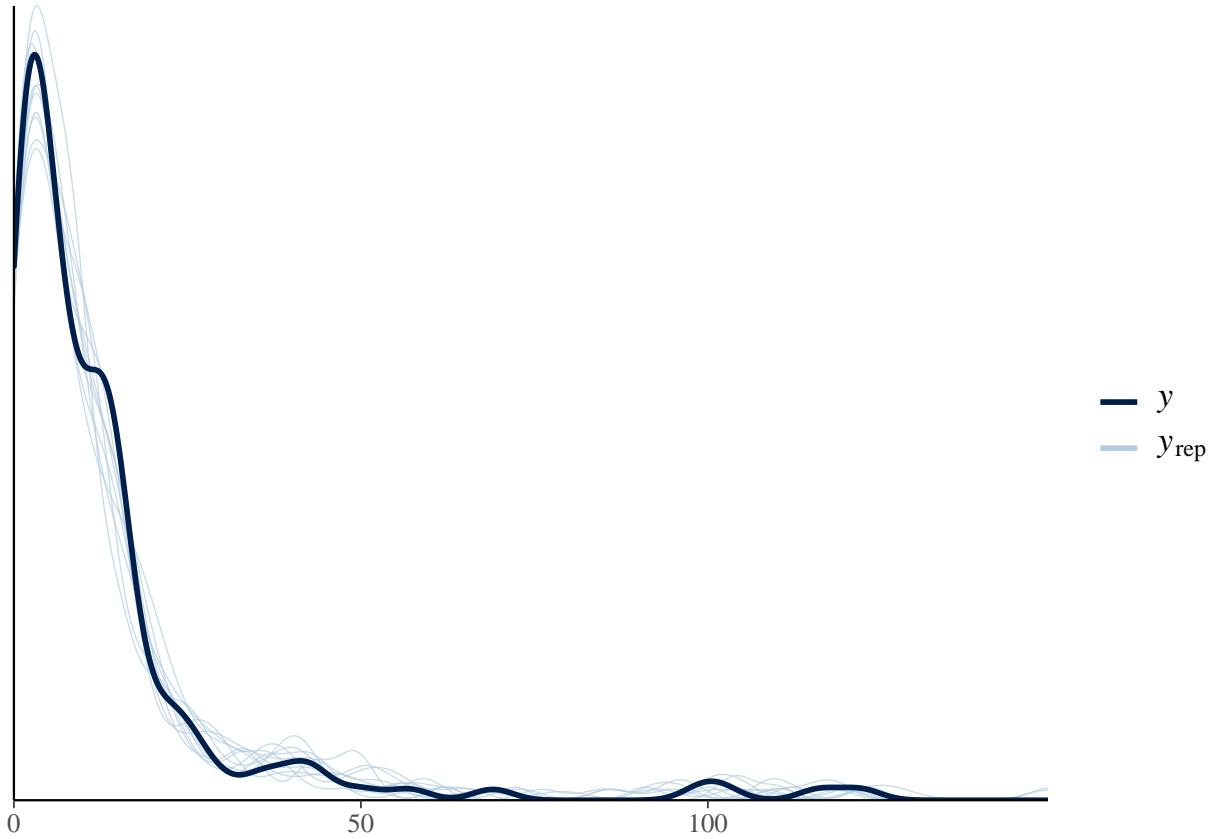
plot(fit_type_phy) #check that chains converged
```





```
pp_check(fit_type_phy) #check the predictions
```

```
## Using 10 posterior samples for ppc type 'dens_overlay' by default.
```



```
summary(fit_type_phy) #summary of model
```

```
## Family: poisson
## Links: mu = log
## Formula: cell_types ~ 0 + germline_timing_simple + scale(log(cell_number)) + (1 | gr(species_id, cov
## Data: df (Number of observations: 157)
## Samples: 5 chains, each with iter = 1e+06; warmup = 1e+05; thin = 1000;
##          total post-warmup samples = 4500
##
## Group-Level Effects:
## ~species_id (Number of levels: 157)
##             Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)    0.19     0.02     0.15     0.24 1.00    4597    4296
##
## Population-Level Effects:
##                               Estimate Est.Error l-95% CI u-95% CI Rhat
## ## germline_timing_simple           1.60     0.35     0.89     2.26 1.00
## ## germline_timing_simpleadult     1.68     0.25     1.17     2.17 1.00
## ## germline_timing_simpleearly     2.22     0.32     1.58     2.82 1.00
## ## germline_timing_simpleno_germline 1.17     0.36     0.45     1.86 1.00
## ## scalelogcell_number            0.61     0.06     0.49     0.73 1.00
## ##                                         Bulk_ESS Tail_ESS
## ## germline_timing_simple          4328     4042
## ## germline_timing_simpleadult    4395     4376
## ## germline_timing_simpleearly    4580     4207
```

```

## germline_timing_simpleno_germline      4673      4361
## scalelogcell_number                   4593      4486
##
## Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS
## and Tail_ESS are effective sample size measures, and Rhat is the potential
## scale reduction factor on split chains (at convergence, Rhat = 1).

```

```
posterior_summary(fit_type_phy, robust = T)
```

	Estimate	Est.Error	Q2.5
## b_germline_timing_simple	1.613862e+00	0.34229785	8.900840e-01
## b_germline_timing_simpleadult	1.683416e+00	0.24941465	1.174107e+00
## b_germline_timing_simpleearly	2.212725e+00	0.31195555	1.583920e+00
## b_germline_timing_simpleno_germline	1.175380e+00	0.35251608	4.527177e-01
## b_scalelogcell_number	6.101713e-01	0.06365627	4.865558e-01
## sd_species_id_Intercept	1.895796e-01	0.02117920	1.512694e-01
## r_species_id[ott1017821,Intercept]	-1.320057e-01	0.37621521	-8.752720e-01
## r_species_id[ott1052546,Intercept]	-1.327601e-01	0.38675465	-8.700756e-01
## r_species_id[ott1059898,Intercept]	6.956797e-01	0.33521919	2.696798e-02
## r_species_id[ott1059900,Intercept]	7.127724e-01	0.32469653	6.029043e-02
## r_species_id[ott1061937,Intercept]	-5.790327e-01	0.41726509	-1.415903e+00
## r_species_id[ott1069171,Intercept]	6.282759e-01	0.31541029	4.312902e-02
## r_species_id[ott1072227,Intercept]	-3.835338e-01	0.40294286	-1.202793e+00
## r_species_id[ott108923,Intercept]	7.422910e-01	0.31742357	1.219379e-01
## r_species_id[ott1099013,Intercept]	-4.105269e-01	0.32951555	-1.050182e+00
## r_species_id[ott111442,Intercept]	9.204121e-01	0.33209181	2.836621e-01
## r_species_id[ott112015,Intercept]	-3.784360e-01	0.40328873	-1.209839e+00
## r_species_id[ott112016,Intercept]	-3.755042e-01	0.40809570	-1.179680e+00
## r_species_id[ott112017,Intercept]	-3.859473e-01	0.40661784	-1.183408e+00
## r_species_id[ott127047,Intercept]	6.900156e-01	0.33181285	2.906827e-02
## r_species_id[ott150272,Intercept]	1.000047e+00	0.35452628	3.291106e-01
## r_species_id[ott160850,Intercept]	-2.773533e-01	0.37531692	-1.003980e+00
## r_species_id[ott165368,Intercept]	1.621118e+00	0.33568207	9.709559e-01
## r_species_id[ott167121,Intercept]	-3.755459e-01	0.36739297	-1.118421e+00
## r_species_id[ott178177,Intercept]	1.721172e-01	0.33809655	-5.016634e-01
## r_species_id[ott178412,Intercept]	4.955721e-01	0.30554917	-1.132571e-01
## r_species_id[ott181933,Intercept]	-3.338145e-01	0.34459883	-1.019382e+00
## r_species_id[ott182906,Intercept]	1.103289e+00	0.33851275	4.433805e-01
## r_species_id[ott186999,Intercept]	4.264693e-01	0.31764556	-2.236822e-01
## r_species_id[ott187583,Intercept]	-8.278403e-01	0.39791756	-1.621770e+00
## r_species_id[ott199292,Intercept]	8.110085e-01	0.32797561	1.745638e-01
## r_species_id[ott207134,Intercept]	-8.029681e-02	0.31328730	-6.952835e-01
## r_species_id[ott215125,Intercept]	1.319569e+00	0.29764254	7.397137e-01
## r_species_id[ott216694,Intercept]	-7.410767e-01	0.35552224	-1.448605e+00
## r_species_id[ott223669,Intercept]	1.660082e+00	0.33954264	9.850681e-01
## r_species_id[ott225275,Intercept]	9.320068e-01	0.30422771	3.422907e-01
## r_species_id[ott237608,Intercept]	-1.431173e-01	0.35030221	-8.315568e-01
## r_species_id[ott246046,Intercept]	2.789052e-01	0.33135693	-3.688767e-01
## r_species_id[ott247341,Intercept]	1.149895e+00	0.35331209	4.614924e-01
## r_species_id[ott256062,Intercept]	-5.335395e-01	0.40136615	-1.309569e+00
## r_species_id[ott256089,Intercept]	-4.657957e-01	0.35547547	-1.180120e+00
## r_species_id[ott256145,Intercept]	-3.812313e-01	0.41271320	-1.168378e+00
## r_species_id[ott263960,Intercept]	8.690192e-01	0.29711291	2.839766e-01
## r_species_id[ott263980,Intercept]	-5.503092e-01	0.36352350	-1.275152e+00

```

## r_species_id[ott263987,Intercept] -4.317320e-01 0.36556780 -1.152812e+00
## r_species_id[ott263988,Intercept] -4.413921e-01 0.33313591 -1.079220e+00
## r_species_id[ott265121,Intercept] -4.803899e-01 0.33271585 -1.152188e+00
## r_species_id[ott266342,Intercept] -8.651876e-01 0.41760990 -1.682602e+00
## r_species_id[ott269063,Intercept] -1.795505e-01 0.41737849 -1.028845e+00
## r_species_id[ott275893,Intercept] -1.398137e-01 0.34028910 -8.046466e-01
## r_species_id[ott275897,Intercept] -1.219377e-01 0.33452841 -7.651139e-01
## r_species_id[ott2810724,Intercept] -1.861746e-01 0.36157209 -8.784147e-01
## r_species_id[ott2819986,Intercept] -4.886254e-01 0.34975464 -1.190270e+00
## r_species_id[ott2821097,Intercept] -3.649568e-01 0.32391265 -1.008585e+00
## r_species_id[ott2844172,Intercept] 1.217628e+00 0.32187949 5.919885e-01
## r_species_id[ott2844962,Intercept] 9.763933e-01 0.34538065 2.695796e-01
## r_species_id[ott2849837,Intercept] 8.672245e-01 0.36532180 1.847016e-01
## r_species_id[ott2942244,Intercept] 8.027351e-01 0.34314852 9.865480e-02
## r_species_id[ott316441,Intercept] 1.018003e+00 0.34516340 3.460523e-01
## r_species_id[ott33153,Intercept] -2.441729e-01 0.37159114 -9.815870e-01
## r_species_id[ott336388,Intercept] -5.035005e-01 0.35713778 -1.200381e+00
## r_species_id[ott34559,Intercept] -4.569033e-01 0.34296789 -1.127589e+00
## r_species_id[ott346740,Intercept] 8.581557e-01 0.30518910 2.694688e-01
## r_species_id[ott3583594,Intercept] 1.055770e+00 0.35978976 3.491975e-01
## r_species_id[ott3587677,Intercept] 1.360876e+00 0.35913420 6.611012e-01
## r_species_id[ott359012,Intercept] -3.615247e-01 0.43186647 -1.237853e+00
## r_species_id[ott361837,Intercept] -3.788062e-01 0.40723288 -1.186241e+00
## r_species_id[ott362913,Intercept] 1.002142e+00 0.31929447 3.802148e-01
## r_species_id[ott365439,Intercept] 4.892054e-01 0.37033641 -2.386455e-01
## r_species_id[ott3663378,Intercept] 6.216634e-01 0.31762264 -6.556566e-03
## r_species_id[ott3665433,Intercept] 9.649679e-01 0.32205313 3.367407e-01
## r_species_id[ott3684291,Intercept] 4.352979e-01 0.35561451 -2.601264e-01
## r_species_id[ott3684365,Intercept] 3.614566e-01 0.38175033 -4.359073e-01
## r_species_id[ott3684379,Intercept] 5.190988e-01 0.39660627 -2.902331e-01
## r_species_id[ott3684389,Intercept] 3.998355e-01 0.41107950 -4.162766e-01
## r_species_id[ott3684437,Intercept] 4.208434e-01 0.37629767 -2.983627e-01
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## r_species_id[ott381980,Intercept] -4.979959e-01 0.34588837 -1.198349e+00
## r_species_id[ott381983,Intercept] -3.755452e-01 0.39921426 -1.200578e+00
## r_species_id[ott395048,Intercept] 1.361184e+00 0.36434368 6.727905e-01
## r_species_id[ott3974169,Intercept] 1.199039e+00 0.29873822 6.109624e-01
## r_species_id[ott3995126,Intercept] 1.166316e+00 0.30252630 5.934373e-01
## r_species_id[ott4010019,Intercept] 6.262726e-02 0.30879574 -5.419050e-01
## r_species_id[ott4010960,Intercept] 7.085555e-02 0.31338406 -5.468632e-01
## r_species_id[ott4011155,Intercept] 2.251828e-01 0.32349423 -3.869468e-01
## r_species_id[ott4013437,Intercept] 1.374208e-01 0.30652330 -4.715778e-01
## r_species_id[ott4013674,Intercept] -6.224201e-03 0.30889343 -6.187875e-01
## r_species_id[ott4013684,Intercept] 4.780737e-02 0.30436749 -5.549839e-01
## r_species_id[ott422679,Intercept] -1.152730e-01 0.32592483 -7.726485e-01
## r_species_id[ott431388,Intercept] 1.049885e+00 0.30867806 4.612936e-01
## r_species_id[ott446088,Intercept] 1.043963e+00 0.36168583 3.396957e-01
## r_species_id[ott4741377,Intercept] 1.052793e-01 0.30892501 -5.146019e-01
## r_species_id[ott4742064,Intercept] 6.431004e-02 0.30412416 -5.348935e-01
## r_species_id[ott481952,Intercept] 1.146571e+00 0.29415384 5.721267e-01
## r_species_id[ott48288,Intercept] 1.107873e+00 0.32968562 4.309378e-01
## r_species_id[ott485470,Intercept] -4.882041e-01 0.34356081 -1.189752e+00
## r_species_id[ott485473,Intercept] -5.165923e-01 0.38599016 -1.305146e+00
## r_species_id[ott485476,Intercept] -3.763729e-01 0.36626065 -1.098182e+00

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## r_species_id[ott485480,Intercept] -6.107805e-01 0.40102465 -1.445477e+00
## r_species_id[ott485482,Intercept] -7.653628e-01 0.38189203 -1.517469e+00
## r_species_id[ott486834,Intercept] -1.921825e-01 0.36292181 -9.047032e-01
## r_species_id[ott490206,Intercept] 9.988855e-01 0.31539995 3.760750e-01
## r_species_id[ott492241,Intercept] 8.215615e-01 0.35338947 1.085340e-01
## r_species_id[ott497063,Intercept] -4.437932e-01 0.33515440 -1.132447e+00
## r_species_id[ott4974308,Intercept] 6.617621e-01 0.36636565 -5.309627e-02
## r_species_id[ott4978773,Intercept] 8.676252e-01 0.35530820 1.651940e-01
## r_species_id[ott4979583,Intercept] 8.036261e-01 0.35878839 9.233255e-02
## r_species_id[ott518643,Intercept] 6.421357e-01 0.36493056 -6.747939e-02
## r_species_id[ott542509,Intercept] 1.479780e+00 0.34603074 8.173515e-01
## r_species_id[ott54768,Intercept] 7.593986e-01 0.31807427 1.402636e-01
## r_species_id[ott549846,Intercept] 5.178710e-01 0.36090529 -1.833122e-01
## r_species_id[ott560703,Intercept] -7.080214e-02 0.36688528 -7.996504e-01
## r_species_id[ott567703,Intercept] 7.581902e-02 0.31222969 -5.298788e-01
## r_species_id[ott570365,Intercept] 6.752173e-01 0.32013065 3.572399e-03
## r_species_id[ott570656,Intercept] 8.938374e-01 0.32398013 2.689032e-01
## r_species_id[ott588761,Intercept] 9.807428e-01 0.30945151 3.719624e-01
## r_species_id[ott592355,Intercept] 4.764503e-01 0.32153858 -1.374226e-01
## r_species_id[ott601255,Intercept] -7.612516e-01 0.35601852 -1.497443e+00
## r_species_id[ott602180,Intercept] -1.301254e-01 0.41326496 -9.575781e-01
## r_species_id[ott60470,Intercept] -3.855166e-01 0.40897077 -1.175136e+00
## r_species_id[ott60471,Intercept] -1.072685e-01 0.39265371 -9.074495e-01
## r_species_id[ott60473,Intercept] -3.838808e-01 0.40881099 -1.182631e+00
## r_species_id[ott60477,Intercept] -3.821318e-01 0.37251048 -1.101541e+00
## r_species_id[ott60479,Intercept] -3.721285e-01 0.37392424 -1.096923e+00
## r_species_id[ott633708,Intercept] -4.456668e-01 0.38843554 -1.238417e+00
## r_species_id[ott633710,Intercept] -5.010891e-01 0.34752457 -1.186866e+00
## r_species_id[ott633711,Intercept] -6.241796e-01 0.37813917 -1.384452e+00
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## r_species_id[ott633719,Intercept] -4.924346e-01 0.35136948 -1.185172e+00
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## r_species_id[ott649193,Intercept] -6.856349e-01 0.35606896 -1.392336e+00
## r_species_id[ott675301,Intercept] 1.249319e+00 0.36250297 5.436957e-01
## r_species_id[ott724784,Intercept] 4.950574e-02 0.31419472 -5.572837e-01
## r_species_id[ott72522,Intercept] -2.647577e-01 0.33374875 -9.039555e-01
## r_species_id[ott727979,Intercept] -7.710108e-01 0.37889118 -1.508077e+00
## r_species_id[ott733462,Intercept] 3.201453e-01 0.31965036 -3.379665e-01
## r_species_id[ott736728,Intercept] -1.176102e-01 0.34785485 -8.385756e-01
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## r_species_id[ott7489702,Intercept] 5.737862e-01 0.35405220 -1.476073e-01
## r_species_id[ott7567530,Intercept] 7.010155e-01 0.32621781 4.275577e-02
## r_species_id[ott765113,Intercept] -8.737064e-02 0.31537223 -7.281576e-01
## r_species_id[ott765280,Intercept] 8.118104e-01 0.31400451 2.441184e-01
## r_species_id[ott779028,Intercept] -2.589175e-02 0.30707605 -6.320431e-01
## r_species_id[ott790395,Intercept] -4.518194e-01 0.33493644 -1.093026e+00
## r_species_id[ott817791,Intercept] -3.347904e-01 0.34170094 -9.949187e-01
## r_species_id[ott821356,Intercept] 9.635533e-01 0.30503384 3.720782e-01
## r_species_id[ott83430,Intercept] -3.983458e-01 0.32609437 -1.025249e+00
## r_species_id[ott83432,Intercept] -3.816801e-01 0.33931622 -1.022614e+00
## r_species_id[ott840001,Intercept] 1.263230e+00 0.30727429 6.878205e-01
## r_species_id[ott841027,Intercept] -8.636857e-01 0.40687239 -1.687187e+00
## r_species_id[ott849781,Intercept] -5.352084e-01 0.35590229 -1.269349e+00

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## r_species_id[ott878345,Intercept] -4.286977e-01 0.35524953 -1.157656e+00
## r_species_id[ott92556,Intercept] -1.583363e-01 0.32098298 -8.005879e-01
## r_species_id[ott92561,Intercept] -2.352381e-01 0.32852658 -8.862393e-01
## r_species_id[ott939432,Intercept] -3.824572e-01 0.34521905 -1.062975e+00
## r_species_id[ott939454,Intercept] -3.718348e-01 0.33974750 -1.080271e+00
## r_species_id[ott954042,Intercept] 4.585395e-01 0.34190517 -2.342827e-01
## r_species_id[ott958293,Intercept] -4.720568e-01 0.37227548 -1.202426e+00
## r_species_id[ott958304,Intercept] -3.821677e-01 0.40086793 -1.182091e+00
## r_species_id[ott962359,Intercept] -2.769978e-01 0.37294406 -1.051810e+00
## r_species_id[ott987480,Intercept] 1.302857e+00 0.36066175 6.190324e-01
## r_species_id[ott989764,Intercept] 3.422009e-01 0.29939534 -2.403670e-01
## lp_ -5.919607e+02 10.93706128 -6.146191e+02
##
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## b_germline_timing_simple 2.260670e+00
## b_germline_timing_simpleadult 2.167713e+00
## b_germline_timing_simpleearly 2.819800e+00
## b_germline_timing_simpleno_germline 1.862193e+00
## b_scelogcell_number 7.325088e-01
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## r_species_id[ott1017821,Intercept] 6.066463e-01
## r_species_id[ott1052546,Intercept] 5.961988e-01
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## r_species_id[ott108923,Intercept] 1.376201e+00
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## r_species_id[ott112015,Intercept] 4.109330e-01
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## r_species_id[ott127047,Intercept] 1.343197e+00
## r_species_id[ott150272,Intercept] 1.697914e+00
## r_species_id[ott160850,Intercept] 4.689000e-01
## r_species_id[ott165368,Intercept] 2.293505e+00
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## r_species_id[ott178177,Intercept] 8.167691e-01
## r_species_id[ott178412,Intercept] 1.130074e+00
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## r_species_id[ott182906,Intercept] 1.798972e+00
## r_species_id[ott186999,Intercept] 1.066171e+00
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## r_species_id[ott207134,Intercept] 5.608879e-01
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## r_species_id[ott216694,Intercept] -3.004971e-02
## r_species_id[ott223669,Intercept] 2.354876e+00
## r_species_id[ott225275,Intercept] 1.560557e+00
## r_species_id[ott237608,Intercept] 5.430362e-01
## r_species_id[ott246046,Intercept] 9.440899e-01
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## r_species_id[ott256089,Intercept] 2.134858e-01

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## r_species_id[ott256145,Intercept]      4.156694e-01
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## r_species_id[ott263988,Intercept]      1.966490e-01
## r_species_id[ott265121,Intercept]      1.987558e-01
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## r_species_id[ott275897,Intercept]      5.611716e-01
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## r_species_id[ott3974169,Intercept]     1.808084e+00
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## r_species_id[ott48288,Intercept]      1.806502e+00

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## r_species_id[ott485470,Intercept]      1.843978e-01
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## r_species_id[ott490206,Intercept]      1.625513e+00
## r_species_id[ott492241,Intercept]      1.519779e+00
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## r_species_id[ott790395,Intercept]     2.080392e-01
## r_species_id[ott817791,Intercept]     3.403504e-01
## r_species_id[ott821356,Intercept]     1.588549e+00
## r_species_id[ott83430,Intercept]      2.647235e-01
## r_species_id[ott83432,Intercept]      2.839053e-01

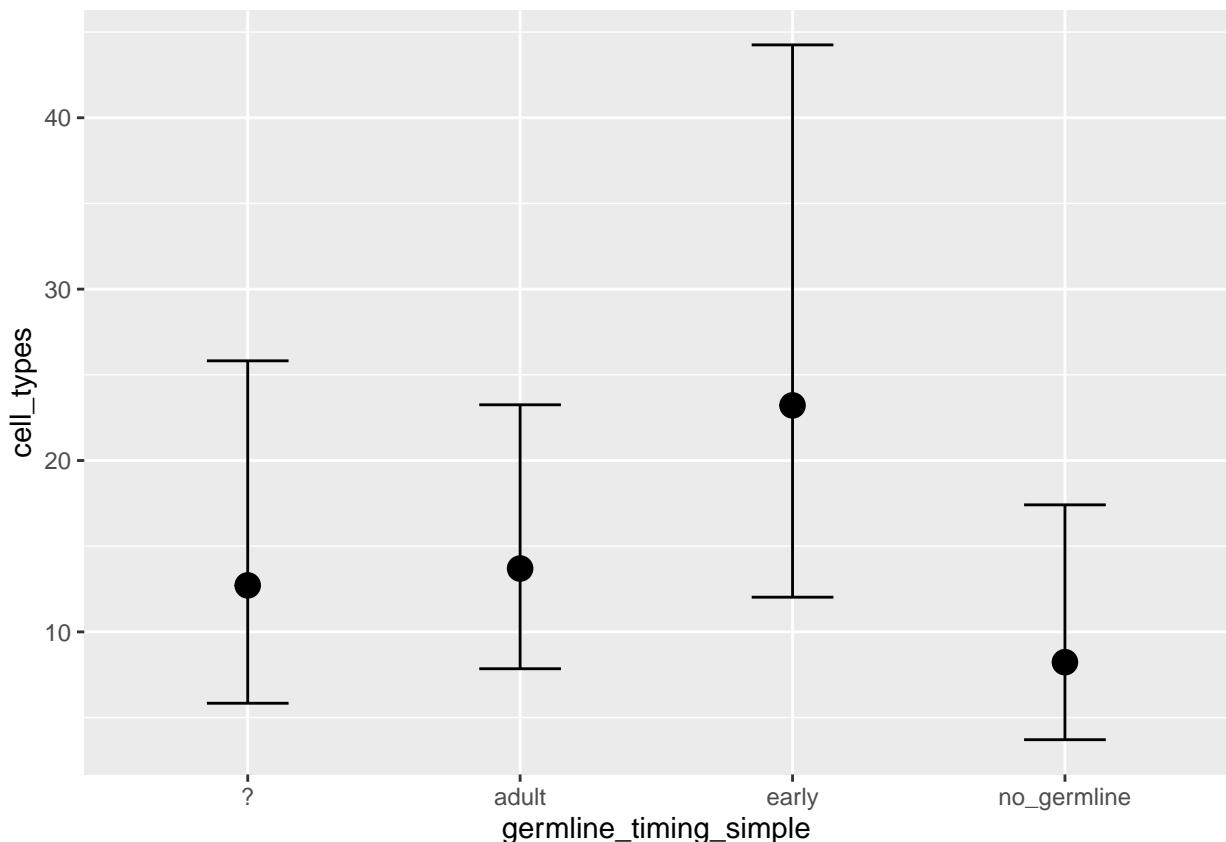
```

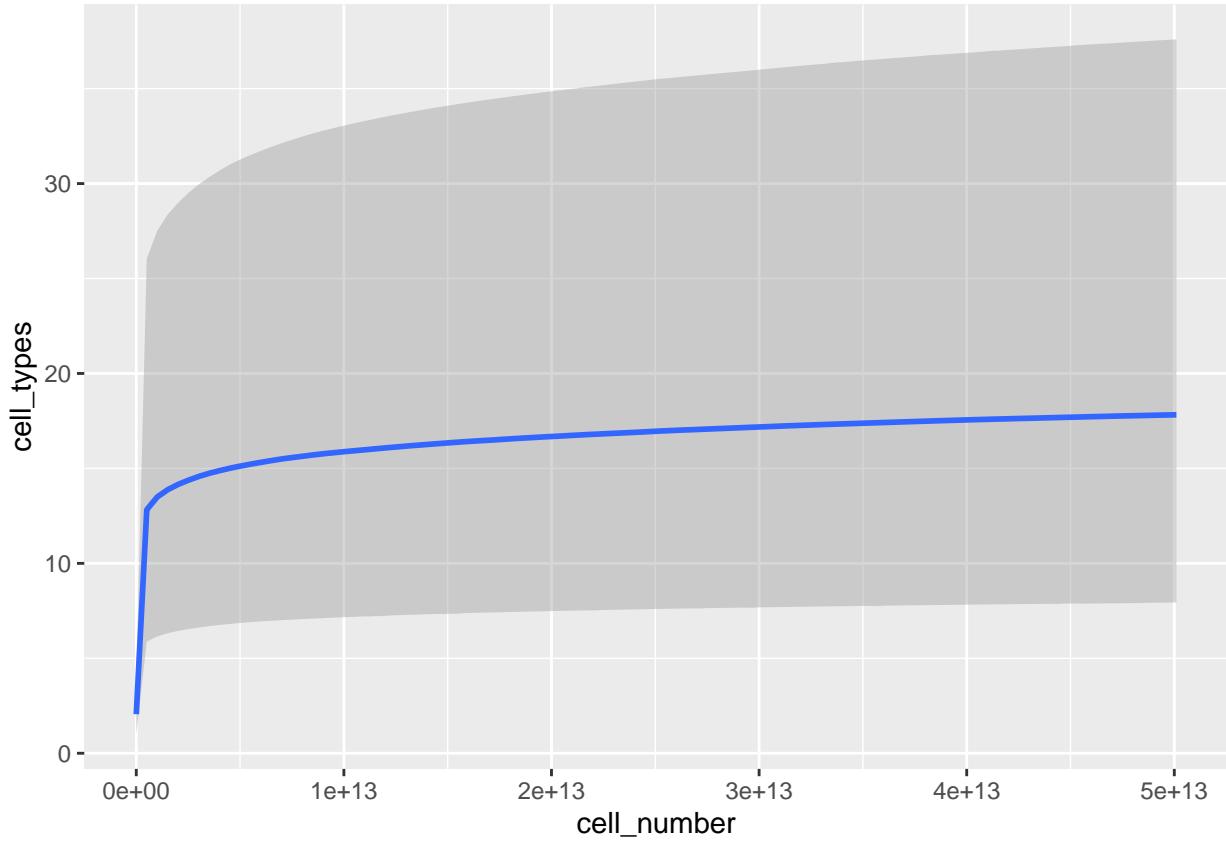
```

## r_species_id[ott840001,Intercept]    1.854120e+00
## r_species_id[ott841027,Intercept]    -6.355871e-02
## r_species_id[ott849781,Intercept]    1.955769e-01
## r_species_id[ott878345,Intercept]    2.879141e-01
## r_species_id[ott92556,Intercept]     4.974022e-01
## r_species_id[ott92561,Intercept]     4.099571e-01
## r_species_id[ott939432,Intercept]    2.598973e-01
## r_species_id[ott939454,Intercept]    2.639871e-01
## r_species_id[ott954042,Intercept]    1.167595e+00
## r_species_id[ott958293,Intercept]    2.822249e-01
## r_species_id[ott958304,Intercept]    4.169826e-01
## r_species_id[ott962359,Intercept]    4.596690e-01
## r_species_id[ott987480,Intercept]    2.040528e+00
## r_species_id[ott989764,Intercept]    9.509299e-01
## lp__
-5.716001e+02

```

```
plot(conditional_effects(fit_type_phy, points = TRUE, ask = F))
```





```
hyp = hypothesis(fit_type_phy, c("germline_timing_simpleearly = germline_timing_simple", "germline_timing_simplelate = germline_timing_simple", "germline_timing_simplemid = germline_timing_simple", "germline_timing_simpleearly < germline_timing_simplelate", "germline_timing_simplemid < germline_timing_simplelate", "germline_timing_simplemid < germline_timing_simpleearly"))
hyp
```

```
## Hypothesis Tests for class b:
##          Hypothesis Estimate Est.Error CI.Lower CI.Upper Evid.Ratio
## 1 (germline_timing_simpleearly = 0      0.61      0.27     0.11     1.15      NA
## 2 (germline_timing_simplelate = 0      0.53      0.21     0.13     0.93      NA
## 3 (germline_timing_simplemid = 0      1.05      0.33     0.41     1.72      NA
##   Post.Prob Star
## 1        NA   *
## 2        NA   *
## 3        NA   *
## ---
## 'CI': 90%-CI for one-sided and 95%-CI for two-sided hypotheses.
## '*': For one-sided hypotheses, the posterior probability exceeds 95%;
## for two-sided hypotheses, the value tested against lies outside the 95%-CI.
## Posterior probabilities of point hypotheses assume equal prior probabilities.
```

```
plot(hyp)
```

