

Bayesian Final Project

2026-01-23

```
library(readxl)
library(dplyr)
library(tidyr)
library(lubridate)
library(stringr)

load("beatspy.RData")
```

Revised Frequentist Models

The mixed-effects models are likely overfitting (boundary (singular) fit issue)

```
#logistic regression models fitted within each sector
library(purrr)
library(modelsummary)

sector_models = m3_df |>
  group_split(gics_sector_name) |>
  setNames(unique(m3_df$gics_sector_name)) |>
  map(~ glm(
    beat_spy ~ log_pe + div_yield,
    data = .x,
    family = binomial(link = "logit")
  ))

modelsummary(sector_models)
```

```
#mixed-effects w/ firm-level random intercepts
library(lme4)
```

```
## Loading required package: Matrix

##
## Attaching package: 'Matrix'

## The following objects are masked from 'package:tidyverse':
##      expand, pack, unpack
```

	Health Care	Information Technology	Consumer Staples	Industrials	Utilities	Financials	Materials
(Intercept)	-0.846 (1.388)	1.524 (0.741)	0.684 (1.268)	-0.559 (0.571)	0.512 (0.566)	-2.651 (1.001)	0.231 (0.726)
log_pe	0.215 (0.403)	-0.304 (0.212)	-0.262 (0.373)	0.049 (0.153)	0.022 (0.168)	0.791 (0.293)	0.000 (0.214)
div_yield	-0.044 (0.097)	-0.279 (0.089)	-0.181 (0.090)	-0.012 (0.052)	-0.184 (0.060)	-0.038 (0.096)	-0.074 (0.081)
Num.Obs.	94	283	306	189	603	321	577
AIC	131.3	387.2	395.5	257.9	826.0	427.6	803.8
BIC	138.9	398.2	406.7	267.7	839.2	438.9	816.9
Log.Lik.	-62.641	-190.622	-194.752	-125.971	-410.016	-210.811	-398.899
RMSE	0.49	0.49	0.47	0.49	0.49	0.48	0.50

```

mixed_intercept = glmer(
  beat.spy ~ log_pe + div_yield +
    (1 | Ticker),
  data = m3_df,
  family = binomial(link = "logit")
)

## boundary (singular) fit: see help('isSingular')

mixed_intercept |> summary()

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: beat.spy ~ log_pe + div_yield + (1 | Ticker)
## Data: m3_df
##
##          AIC      BIC      logLik -2*log(L)  df.resid
##        4626.4   4650.9   -2309.2     4618.4      3396
##
## Scaled residuals:
##       Min      1Q  Median      3Q     Max
## -1.1735 -0.9373 -0.7145  1.0129  4.7846
##
## Random effects:
##   Groups Name        Variance Std.Dev.
##   Ticker (Intercept) 0         0
##   Number of obs: 3400, groups: Ticker, 406
##
## Fixed effects:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.17522   0.19108   0.917   0.359
## log_pe      0.03057   0.05524   0.553   0.580
## div_yield   -0.15169   0.01974  -7.684 1.55e-14 ***
## ---

```

```

## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) log_pe
## log_pe   -0.946
## div_yield -0.539  0.299
## optimizer (Nelder_Mead) convergence code: 0 (OK)
## boundary (singular) fit: see help('isSingular')

#mixed effects w/ firm-level random slopes
mixed_random_slopes = glmer(
  beat.spy ~ log_pe + div_yield +
    (1 + log_pe + div_yield | Ticker),
  data = m3_df,
  family = binomial(link = "logit"),
  control = glmerControl(optimizer = "bobyqa")
)

## boundary (singular) fit: see help('isSingular')

mixed_random_slopes |> summary()

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula: beat.spy ~ log_pe + div_yield + (1 + log_pe + div_yield | Ticker)
## Data: m3_df
## Control: glmerControl(optimizer = "bobyqa")
##
##      AIC      BIC      logLik -2*log(L)  df.resid
## 4632.9  4688.0  -2307.4   4614.9      3391
##
## Scaled residuals:
##      Min      1Q Median      3Q      Max
## -1.1874 -0.9363 -0.6846  1.0067  2.9505
##
## Random effects:
## Groups Name      Variance Std.Dev. Corr
## Ticker (Intercept) 0.289481 0.53803
##           log_pe     0.012842 0.11332 -1.00
##           div_yield  0.007972 0.08928 -1.00  1.00
## Number of obs: 3400, groups: Ticker, 406
##
## Fixed effects:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept)  0.24602   0.20147   1.221   0.222
## log_pe       0.02221   0.05797   0.383   0.702
## div_yield   -0.17073   0.02119  -8.057 7.84e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) log_pe
## log_pe   -0.946
## div_yield -0.539  0.299
## optimizer (Nelder_Mead) convergence code: 0 (OK)
## boundary (singular) fit: see help('isSingular')

```

```

## log_pe      -0.949
## div_yield -0.565  0.335
## optimizer (bobyqa) convergence code: 0 (OK)
## boundary (singular) fit: see help('isSingular')

#mixed-effects model w/ sector-level random slopes
mixed_sector_slopes = glmer(
  beat.spy ~ log_pe + div_yield +
    (1 + log_pe + div_yield | gics_sector_name),
  data = m3_df,
  family = binomial(link = "logit")
)

## boundary (singular) fit: see help('isSingular')

mixed_sector_slopes |> summary()

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial  ( logit )
## Formula:
## beat.spy ~ log_pe + div_yield + (1 + log_pe + div_yield | gics_sector_name)
##   Data: m3_df
##
##       AIC     BIC   logLik -2*log(L)  df.resid
##   4597.2   4652.4   -2289.6     4579.2      3391
##
## Scaled residuals:
##   Min     1Q Median     3Q    Max
## -1.4695 -0.8955 -0.6795  0.9989  3.4987
##
## Random effects:
##   Groups           Name        Variance Std.Dev. Corr
##   gics_sector_name (Intercept) 0.400691 0.63300
##                   log_pe       0.004766 0.06904 -0.99
##                   div_yield   0.002169 0.04658 -0.99  0.96
##   Number of obs: 3400, groups: gics_sector_name, 11
##
## Fixed effects:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.32205   0.29108 -1.106   0.2686
## log_pe       0.12805   0.06343  2.019   0.0435 *
## div_yield   -0.11209   0.02587 -4.333 1.47e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##   (Intr) log_pe
## log_pe   -0.885
## div_yield -0.743  0.472
## optimizer (Nelder_Mead) convergence code: 0 (OK)
## boundary (singular) fit: see help('isSingular')

```

```

#mixed-effects model with sector-level variation + firm-level variation within sector
mixed_nested = glmer(
  beat_spy ~ log_pe + div_yield +
    (1 | gics_sector_name/Ticker),
  data = m3_df,
  family = binomial(link = "logit")
)

## boundary (singular) fit: see help('isSingular')

mixed_nested |> summary()

## Generalized linear mixed model fit by maximum likelihood (Laplace
## Approximation) [glmerMod]
## Family: binomial  ( logit )
## Formula: beat_spy ~ log_pe + div_yield + (1 | gics_sector_name/Ticker)
## Data: m3_df
##
##      AIC      BIC      logLik -2*log(L)  df.resid
##  4593.4   4624.0   -2291.7    4583.4      3395
##
## Scaled residuals:
##      Min      1Q  Median      3Q      Max
## -1.4931 -0.9133 -0.6750  1.0000  3.8468
##
## Random effects:
## Groups            Name        Variance Std.Dev.
## Ticker:gics_sector_name (Intercept) 2.908e-10 1.705e-05
## gics_sector_name     (Intercept) 7.238e-02 2.690e-01
## Number of obs: 3400, groups:
## Ticker:gics_sector_name, 406; gics_sector_name, 11
##
## Fixed effects:
##             Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.29217   0.24022 -1.216   0.2239
## log_pe       0.13164   0.06247  2.107   0.0351 *
## div_yield   -0.11290   0.02243 -5.034 4.81e-07 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ',' 1
##
## Correlation of Fixed Effects:
##          (Intr) log_pe
## log_pe   -0.894
## div_yield -0.595  0.414
## optimizer (Nelder_Mead) convergence code: 0 (OK)
## boundary (singular) fit: see help('isSingular')

```

Bayesian Analysis

```

library(brms)
library(tidybayes)

```

```
library(bayesplot)
library(posterior)
```

Model 1 (Logistic)

$$Y_{i,t} \sim Bernoulli(p_{i,t})$$

$$\text{logit}(p_{i,t}) = \beta_0 + \beta_1 + \log(PE_{i,t}) + \beta_2 \text{DivYield}_{i,t}$$

```
bayes_model1 <- brm(
  beat_spy ~ log_pe + div_yield,
  data = m3_df,
  family = bernoulli(link = "logit"),
  seed = 123
)
```

```
## Compiling Stan program...
```

```
## Trying to compile a simple C file
```

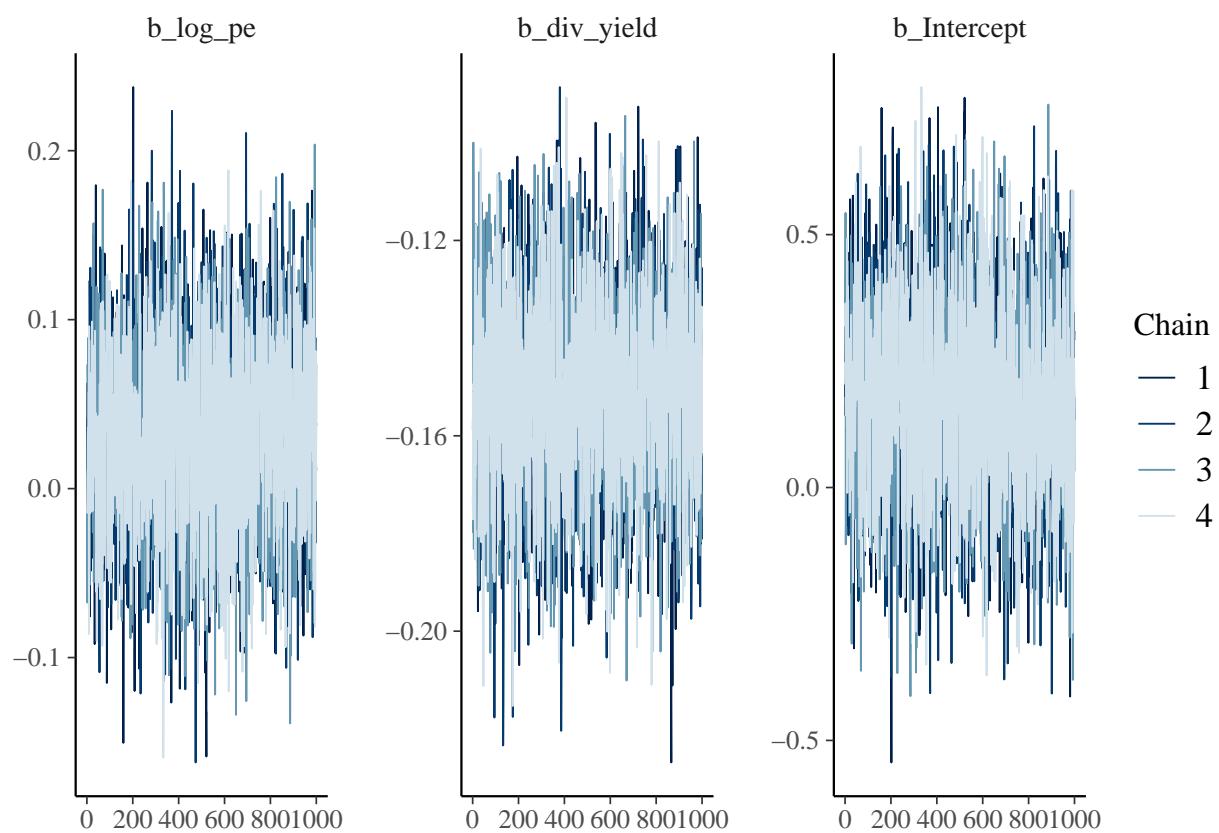
```
## Start sampling
```

```
summary(bayes_model1)$fixed
```

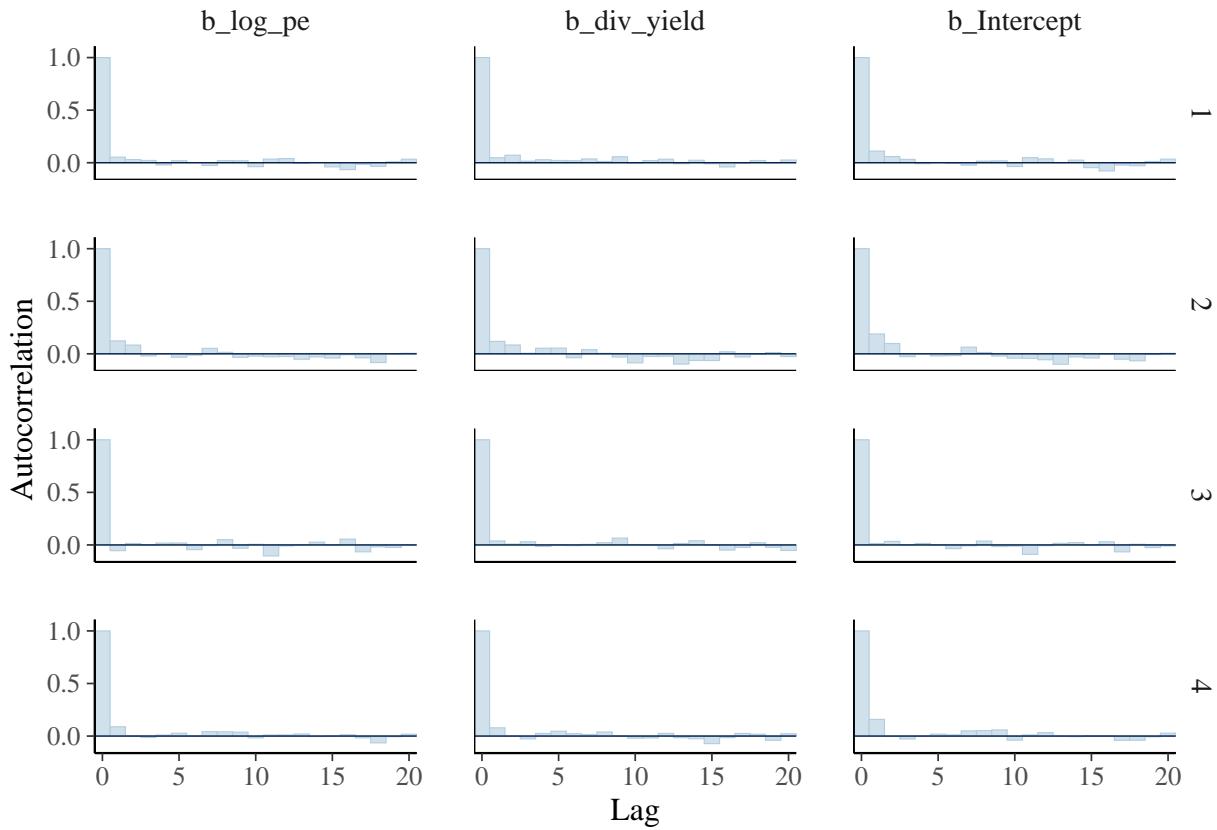
	Estimate	Est.Error	1-95% CI	u-95% CI	Rhat	Bulk_ESS
## Intercept	0.17273278	0.18791917	-0.1830719	0.5376506	1.001122	3007.931
## log_pe	0.03149262	0.05432937	-0.0760189	0.1343352	1.001685	3368.702
## div_yield	-0.15189077	0.01951898	-0.1906835	-0.1138507	1.000259	2819.099
## Tail_ESS						
## Intercept	2906.451					
## log_pe	3373.531					
## div_yield	2796.484					

```
draws_model1 = bayes_model1 |>
  as_draws_array()

mcmc_trace(draws_model1,
           pars = c("b_log_pe", "b_div_yield", "b_Intercept"))
```



```
mcmc_acf_bar(
  draws_model1,
  pars = c("b_log_pe", "b_div_yield", "b_Intercept")
)
```



Model 2 (Hierarchical)

```

bayes_model2 <- brm(
  beat.spy ~ log_pe + div_yield + (1 | gics_sector_name/Ticker),
  data = m3_df,
  family = bernoulli(link = "logit"),
  seed = 123
)

## Compiling Stan program...

## Trying to compile a simple C file

## Start sampling

summary(bayes_model2)

## Family: bernoulli
## Links: mu = logit
## Formula: beat.spy ~ log_pe + div_yield + (1 | gics_sector_name/Ticker)
## Data: m3_df (Number of observations: 3400)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;

```

```

##          total post-warmup draws = 4000
##
## Multilevel Hyperparameters:
## ~gics_sector_name (Number of levels: 11)
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)     0.34      0.10     0.19     0.59 1.00     1348     2025
##
## ~gics_sector_name:Ticker (Number of levels: 406)
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept)     0.06      0.05     0.00     0.17 1.00     1607     1590
##
## Regression Coefficients:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept     -0.32      0.26    -0.82     0.17 1.00     2110     2593
## log_pe        0.14      0.06     0.02     0.26 1.00     3616     2987
## div_yield    -0.11      0.02    -0.15    -0.07 1.00     3638     2948
##
## Draws were sampled 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).

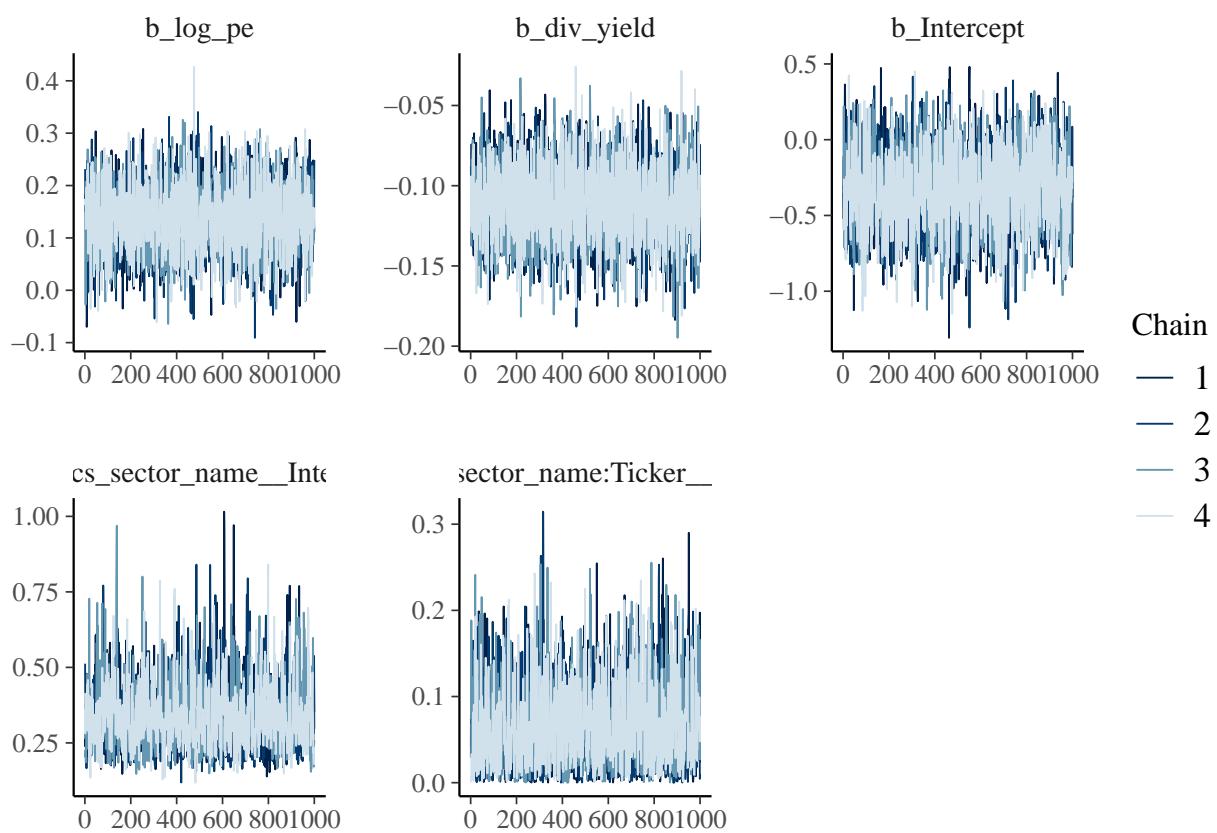
```

```

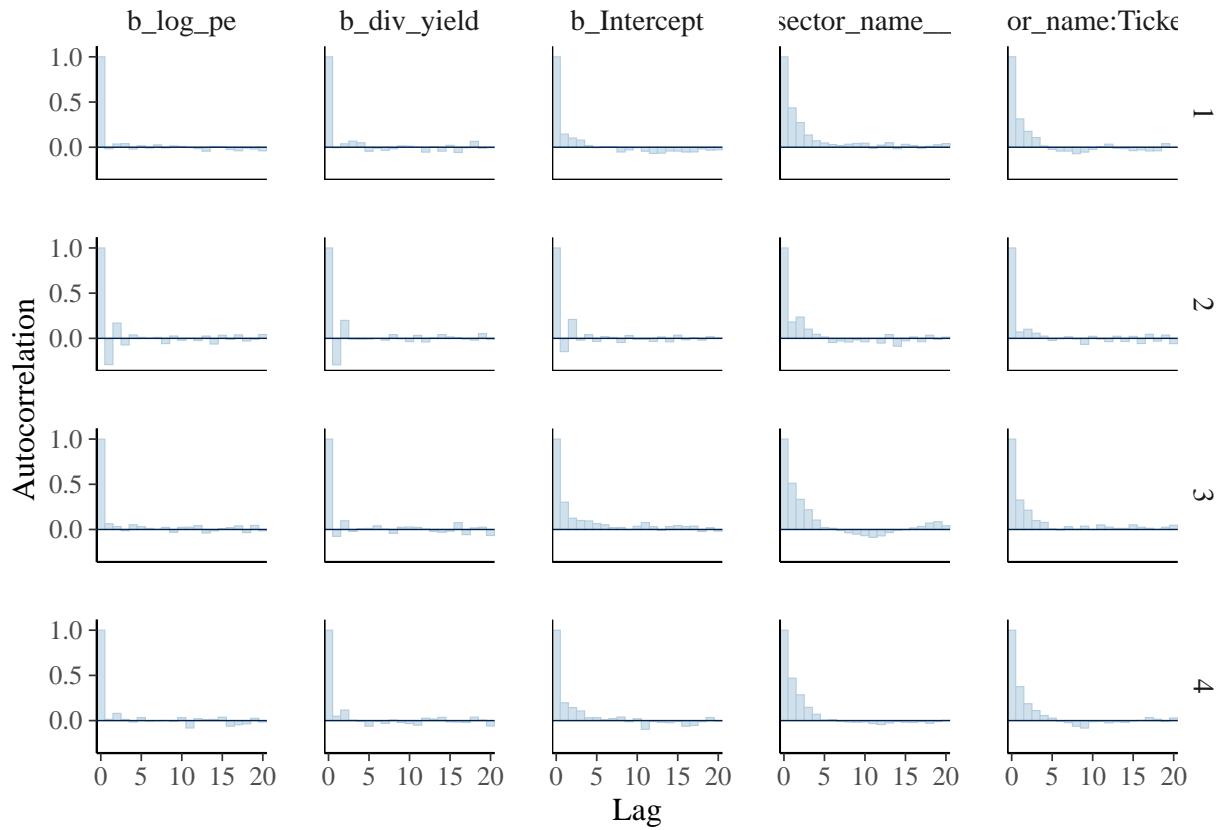
draws_model2 = bayes_model2 |>
  as_draws_array()

mcmc_trace(
  draws_model2,
  pars = c("b_log_pe",
           "b_div_yield",
           "b_Intercept",
           "sd_gics_sector_name__Intercept",
           "sd_gics_sector_name:Ticker__Intercept")
)

```



```
mcmc_acf_bar(
  draws_model2,
  pars = c("b_log_pe", "b_div_yield", "b_Intercept", "sd_gics_sector_name__Intercept", "sd_gics_sector_name__Intercept"))
```



Model Comparison

```
loo_compare(loo(bayes_model1),
            loo(bayes_model2))
```

```
##          elpd_diff se_diff
## bayes_model2    0.0      0.0
## bayes_model1 -22.2     7.0
```

The difference in expected log predictive density is 22.2 (SE = 7.0). LOOCV provides strong evidence that the multilevel specification improves out-of-sample predictive performance.