MCMC examples

```
library(MCMCglmm)
library(brms)
library(rstanarm)
library(lme4)
options(brms.backend = "cmdstanr")
library(broom.mixed)
library(purrr)
library(dplyr)
library(tidybayes)
library(bayesplot)
library(bayestestR)
library(ggplot2); theme_set(theme_bw())
library(shinystan)
library(tidyverse)
```

• a little more on priors:

```
– parameter-expanded priors: y_j|\mu,\xi_j\sim N(\mu+\alpha\sigma_j,\sigma_j^2),\ \sigma_j\sim N(0,\sigma_\xi^2); \alpha\sim N(\alpha_0,\sigma_\alpha),\ \sigma_\alpha\sim \text{inverse-Gamma}(\nu)
```

```
df(v/alpha.V, df1 = 1, df2 = nu, ncp = (alpha.mu^2)/alpha.V)
2 * dt(sqrt(v)/sqrt(alpha.V), df = nu, ncp = alpha.mu/sqrt(alpha.V))
```

... always set alpha.mu=0, can set V = 1 (or diag() in more complex cases) wlog; sqrt(alpha.V) (scale) and nu are the only relevant parameters

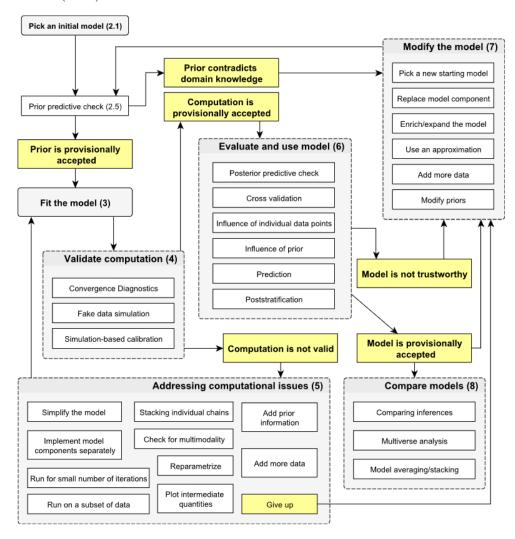
effective sample size

- number of samples, corrected for autocorrelation
- ESS may be > sample size! (e.g. antithetic sampling)

- efficiency of a sampler is not (samples/time), but (effective samples/time)
- effective sample size >1000 for both tail and bulk quantities (Vehtari et al. 2021)

Bayesian workflow

Gelman et al. (2020)



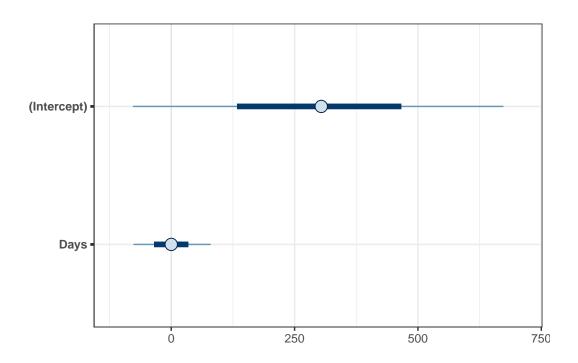
simulation-based calibration

Talts et al. (2020)

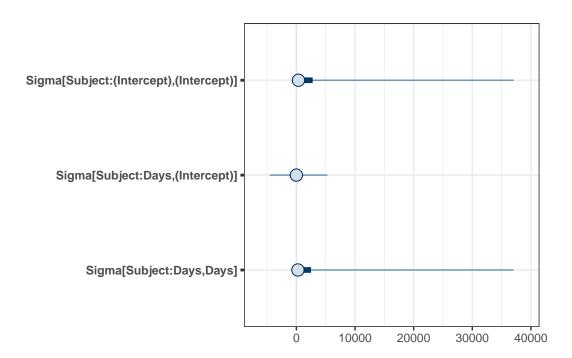
default priors/prior predictive simulations:

• rstanarm default priors: https://cran.r-project.org/web/packages/rstanarm/vignettes/priors.html
Using the good old sleepstudy example:

```
priorpred <- stan_lmer(Reaction ~ Days + (Days|Subject),</pre>
                          prior_PD = TRUE, data = sleepstudy, chains = 1,
                          refresh = 0)
  prior_summary(priorpred)
Priors for model 'priorpred'
Intercept (after predictors centered)
  Specified prior:
    ~ normal(location = 299, scale = 2.5)
  Adjusted prior:
    ~ normal(location = 299, scale = 141)
Coefficients
  Specified prior:
    ~ normal(location = 0, scale = 2.5)
  Adjusted prior:
    ~ normal(location = 0, scale = 49)
Auxiliary (sigma)
  Specified prior:
    ~ exponential(rate = 1)
  Adjusted prior:
    ~ exponential(rate = 0.018)
Covariance
 ~ decov(reg. = 1, conc. = 1, shape = 1, scale = 1)
See help('prior_summary.stanreg') for more details
  plot(priorpred, pars = c("(Intercept)", "Days"))
```

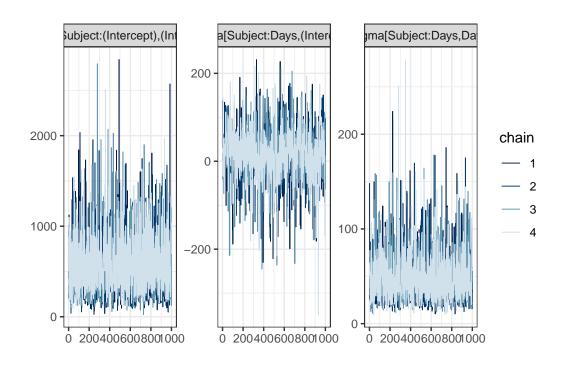


plot(priorpred, regex_pars = "Sigma")

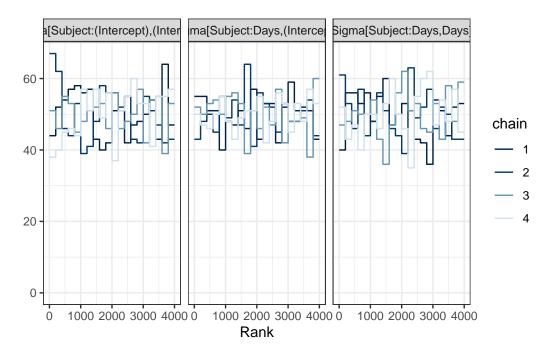


launch_shinystan(stanfit)

mcmc_trace(stanfit, regex_pars= "Sigma")



mcmc_rank_overlay(stanfit, regex_pars= "Sigma")



- MCMC diagnostics
 - trace plots, improved trace plots
 - R-hat Vehtari et al. (2021)
 - divergences (HMC only)

See http://bbolker.github.io/bbmisc/bayes/examples.html

doing stuff with the results

```
tidy(stanfit, effects=c("fixed", "ran_pars"), conf.int = TRUE)
```

A tibble: 6 x 6 term estimate std.error conf.low conf.high group <chr> <dbl> <dbl> <dbl> <dbl> <chr> 1 (Intercept) 251. 6.34 240. 262. <NA> 2 Days 10.5 1.65 7.52 13.2 <NA> 3 sd_(Intercept).Subject 23.8 NA Subject NANA4 sd_Days.Subject 6.94 NANANASubject 5 cor_(Intercept).Days.Subject 0.0806 NASubject NANA6 sd_Observation.Residual 26.0 Residual NANANA

¿¿ why don't we get confidence intervals?? Do it by hand ...

```
(as_draws(stanfit)
       |> tidyr::pivot_longer(everything())
       |> group_by(name)
       |> summarise(estimate = median(value),
                    lwr = quantile(value, 0.025),
                    upr = quantile(value, 0.975))
       |> filter(!stringr::str_detect(name, "^b\\["))
  )
# A tibble: 9 x 4
 name
                                           estimate
                                                         lwr
                                                                upr
  <chr>
                                              <dbl>
                                                       <dbl>
                                                              <dbl>
1 (Intercept)
                                              251.
                                                      238.
                                                              265.
2 .chain
                                                2.5
                                                        1
                                                                4
3 .draw
                                             2000.
                                                      101.
                                                             3900.
4 .iteration
                                              500.
                                                       26.0
                                                              975.
5 Days
                                               10.5
                                                        6.86
                                                               13.8
                                                             1295.
6 Sigma[Subject:(Intercept),(Intercept)]
                                              510.
                                                      156.
7 Sigma[Subject:Days,(Intercept)]
                                               16.0 -106.
                                                              112.
8 Sigma[Subject:Days,Days]
                                               42.9
                                                       18.1
                                                              108.
9 sigma
                                               26.0
                                                       23.2
                                                               29.4
  form1 <- Reaction ~ Days + (Days|Subject)</pre>
  get_prior(form1, sleepstudy)
                                class
                                                    group resp dpar nlpar lb ub
                      prior
                                            coef
                     (flat)
                                     b
                     (flat)
                                     b
                                            Days
                     lkj(1)
                                   cor
                     lkj(1)
                                                  Subject
                                   cor
 student_t(3, 288.7, 59.3) Intercept
     student_t(3, 0, 59.3)
                                                                            0
     student_t(3, 0, 59.3)
                                    sd
                                                 Subject
                                                                            0
     student_t(3, 0, 59.3)
                                    sd
                                            Days Subject
                                                                            0
     student_t(3, 0, 59.3)
                                    sd Intercept Subject
                                                                            0
     student_t(3, 0, 59.3)
                                                                            0
                                sigma
       source
      default
```

```
(vectorized)
      default
 (vectorized)
      default
      default
 (vectorized)
 (vectorized)
 (vectorized)
      default
  b_prior <- c(set_prior("normal(200, 50)", "Intercept"),</pre>
                set prior("normal(0, 10)", "b"),
                set_prior("normal(0, 1)", "sigma")
  b <- brm(form1, sleepstudy,</pre>
            prior = b_prior,
                                    ## reproducibility
            seed = 101,
            sample_prior = 'only', ## for prior predictive sim
            chains = 1, iter = 500, ## very short sample for convenience
            silent = 2, refresh = 0 ## be vewy vewy quiet ...
  p_df <- sleepstudy |> add_predicted_draws(b)
'spaghetti plot' of prior preds
  gg0 <- ggplot(p_df,aes(Days, .prediction, group=interaction(Subject,.draw))) +
           geom_line(alpha = 0.1)
  b_prior4 <- c(set_prior("normal(200, 5)", "Intercept"),</pre>
                 set_prior("normal(0, 2)", "b"),
                 set_prior("normal(0, 1)", "sd"),
                 set_prior("normal(0, 1)", "sigma")
                 )
  cc <-capture.output(</pre>
      b_reg <- brm(form1, sleepstudy,</pre>
                prior = b_prior4,
                seed = 101,
                init = 0,
                control = list(adapt_delta = 0.95)
```

)

Start sampling

Chain 1 Informational Message: The current Metropolis proposal is about to be rejected because the control of t

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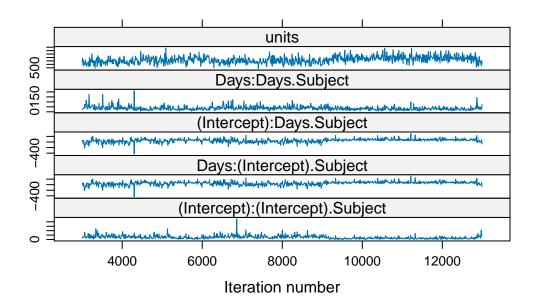
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Parameter Rhat ESS MCSE
b_Days 1.0002 3744 0.01764
b_Intercept 0.9997 3297 0.05154

```
## debug(MCMCglmm:::priorformat)
  m <- MCMCglmm(Reaction ~ Days, random = ~us(1+Days):Subject,</pre>
                data = sleepstudy,
                verbose=FALSE,
                prior = list(G=list(G1=list(V=diag(2), nu = 0.1))))
  broom.mixed::tidy(m)
# A tibble: 6 x 5
 effect group
                   term
                                       estimate std.error
  <chr> <chr> <chr>
                                            <dbl>
                                                    <dbl>
1 fixed <NA> (Intercept)
                                            252.
                                                      6.65
2 fixed <NA>
                   Days
                                            10.5
                                                     1.61
3 ran_pars Subject var__(Intercept)
                                            508.
                                                   357.
4 ran_pars Subject cov__(Intercept).Days
                                            49.1
                                                    63.5
5 ran_pars Subject var__Days
                                                    19.9
                                            33.3
6 ran_pars Residual var__Observation
                                          700.
                                                    98.6
  try(MCMCglmm(Reaction ~ Days, random = ~us(1+Days):Subject,
                data = sleepstudy,
                verbose=FALSE,
                prior = list(G=list(G1=list(V=diag(2), nu = 0.1,
                                           alpha.mu = 0, alpha.V = diag(2))))))
Error in priorformat(if (NOpriorG) { :
  alpha.mu is the wrong length for some prior$G/prior$R elements
  m2 <- MCMCglmm(Reaction ~ Days, random = ~us(1+Days):Subject,</pre>
                data = sleepstudy,
                verbose=FALSE,
                prior = list(G=list(G1=list(V=diag(2), nu = 0.1,
                                           alpha.mu = rep(0,2),
                                           alpha.V = diag(2))))
  lattice::xyplot(m2$VCV)
```



Run longer (and thin)? Strengthen prior?

Gelman, Andrew, Aki Vehtari, Daniel Simpson, Charles C. Margossian, Bob Carpenter, Yuling Yao, Lauren Kennedy, Jonah Gabry, Paul-Christian Bürkner, and Martin Modrák. 2020. "Bayesian Workflow." arXiv:2011.01808 [Stat], November. http://arxiv.org/abs/2011.01808.

Talts, Sean, Michael Betancourt, Daniel Simpson, Aki Vehtari, and Andrew Gelman. 2020. "Validating Bayesian Inference Algorithms with Simulation-Based Calibration." arXiv:1804.06788 [Stat], October. http://arxiv.org/abs/1804.06788.

Vehtari, Aki, Andrew Gelman, Daniel Simpson, Bob Carpenter, and Paul-Christian Bürkner. 2021. "Rank-Normalization, Folding, and Localization: An Improved R-hat for Assessing Convergence of MCMC (with Discussion)." *Bayesian Analysis* 16 (2): 667–718. https://doi.org/10.1214/20-BA1221.