

Summarizing posteriors through tables in RMarkdown

Johannes Karreth

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Simulate data and fit a model

```
n.sim <- 100; set.seed(123)
x1 <- rnorm(n.sim, mean = 5, sd = 2)
x2 <- rbinom(n.sim, size = 1, prob = 0.3)
e <- rnorm(n.sim, mean = 0, sd = 1)
b1 <- 1.2
b2 <- -3.1
a <- 1.5
y <- a + b1 * x1 + b2 * x2 + e

sim.dat <- data.frame(y, x1, x2)

bayes.mod <- function() {

  for(i in 1:N){
    y[i] ~ dnorm(mu[i], tau)
    mu[i] <- alpha + beta1 * x1[i] + beta2 * x2[i]
  }

  alpha ~ dnorm(0, .01)
  beta1 ~ dunif(-100, 100)
  beta2 ~ dunif(-100, 100)
  tau ~ dgamma(.01, .01)

}

sim.dat.jags <- as.list(sim.dat)

sim.dat.jags$N <- nrow(sim.dat)

bayes.mod.params <- c("alpha", "beta1", "beta2")

inits1 <- list("alpha" = 0, "beta1" = 0, "beta2" = 0)
inits2 <- list("alpha" = 1, "beta1" = 1, "beta2" = 1)
inits3 <- list("alpha" = -1, "beta1" = -1, "beta2" = -1)
bayes.mod.inits <- list(inits1, inits2, inits3)

library(R2jags)

## Loading required package: rjags
## Loading required package: coda
## Linked to JAGS 3.4.0
## Loaded modules: basemod,bugs
```

```
##
## Attaching package: 'R2jags'
##
## The following object is masked from 'package:coda':
##
##      traceplot

set.seed(123)

bayes.mod.fit <- jags(data = sim.dat.jags, inits = bayes.mod.inits,
  parameters.to.save = bayes.mod.params, n.chains = 3, n.iter = 9000,
  n.burnin = 1000,
  model.file = bayes.mod)

## module glm loaded

## Compiling model graph
##   Resolving undeclared variables
##   Allocating nodes
##   Graph Size: 511
##
## Initializing model
```

Create a summary table

```
devtools::source_url("https://raw.githubusercontent.com/jkarreth/JKmisc/master/mcmctab.R")
```

```
## SHA-1 hash of file is 0ee96c99f90759387c6cb1a882a5f0d82a6c7dfb
```

```
regtable <- mcmctab(as.mcmc(bayes.mod.fit))[-c(4), ] # remove the fourth row
```

Option 1: the pander package

```
library(pander)
pander(regtable)
```

	Mean	SD	Lower	Upper
alpha	1.844	0.295	1.264	2.401
beta1	1.136	0.053	1.033	1.238
beta2	-3.087	0.207	-3.486	-2.687

Option 2: the knitr package

```
library(knitr)
kable(regtable)
```

	Mean	SD	Lower	Upper
alpha	1.844	0.295	1.264	2.401
beta1	1.136	0.053	1.033	1.238
beta2	-3.087	0.207	-3.486	-2.687