Summarizing posteriors through tables in RMarkdown

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Simulate data and fit a model

```
n.sim <- 100; set.seed(123)
x1 \leftarrow rnorm(n.sim, mean = 5, sd = 2)
x2 \leftarrow rbinom(n.sim, size = 1, prob = 0.3)
e \leftarrow rnorm(n.sim, mean = 0, sd = 1)
b1 <- 1.2
b2 <- -3.1
a < -1.5
y \leftarrow a + b1 * x1 + b2 * x2 + e
sim.dat <- data.frame(y, x1, x2)</pre>
bayes.mod <- function() {</pre>
for(i in 1:N){
y[i] ~ dnorm(mu[i], tau)
mu[i] \leftarrow 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)</pre>
sim.dat.jags$N <- nrow(sim.dat)</pre>
bayes.mod.params <- c("alpha", "beta1", "beta2")</pre>
inits1 <- list("alpha" = 0, "beta1" = 0, "beta2" = 0)
inits2 <- list("alpha" = 1, "beta1" = 1, "beta2" = 1)</pre>
inits3 <- list("alpha" = -1, "beta1" = -1, "beta2" = -1)
bayes.mod.inits <- list(inits1, inits2, inits3)</pre>
library(R2jags)
```

```
## Loading required package: rjags
## Loading required package: coda
## Linked to JAGS 4.2.0
```

```
## 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,</pre>
 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 information:
##
      Observed stochastic nodes: 100
      Unobserved stochastic nodes: 4
##
##
      Total graph size: 516
## Initializing model
```

Create a summary table

```
devtools::source_url("https://raw.githubusercontent.com/jkarreth/JKmisc/master/mcmctab.R")
## SHA-1 hash of file is aaf011f39a5b02aa830972cda46ebd378fc45d9f
regtable <- mcmctab(as.mcmc(bayes.mod.fit))[-c(4), ] # remove the fourth row</pre>
```

Option 1: the pander package

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

	Mean	SD	Lower	Upper	Pr
alpha	1.849	0.294	1.259	2.427	1
beta1	1.134	0.052	1.032	1.237	1
$\mathbf{beta2}$	-3.086	0.212	-3.494	-2.667	1

Option 2: the knitr package

library(knitr)
kable(regtable)

	Mean	SD	Lower	Upper	Pr
alpha	1.849	0.294	1.259	2.427	1
beta1	1.134	0.052	1.032	1.237	1
beta2	-3.086	0.212	-3.494	-2.667	1