

R Codes for Preliminary Checks (Bayesian Method)

SSHPA / AISDL
2018

```
## column names (variables): scid, bmattitude, bmuse, freqstillgood, capab,
#bmdiff,
## theodiff, moddiff, resdiff, codediff, plan, trysoft, learntime
## the following codes run well with the hypothetical data set btestdata.csv
## bmdat <- read.table("c://dr.vuong/BMSurvey/btestdata.csv", sep=",",
header=T)

m1 <- map(
  alist(
    bmuse ~ dbinom(1,p) ,
    logit(p) <- a + bfreq*freqstillgood,
    a ~ dnorm(0,10),
    bfreq ~ dnorm(0,10)
  ),
  data=bmdat)

m3 <- map(
  alist(
    bmuse ~ dbinom(1,p) ,
    logit(p) <- a + bfreq*freqstillgood + bcap*capab,
    a ~ dnorm(0,10),
    bfreq ~ dnorm(0,10),
    bcap ~ dnorm(0,10)
  ),
  data=bmdat)

## Note for m1: a model with the suspicion that when one considers the old
## way of
## analysis is still good, one is hesitant to adopt the Bayesian method

m2 <- map(
  alist(
    bmattitude ~ dbinom(1,p) ,
    logit(p) <- a + bfreq*freqstillgood + bcap*capab,
    a ~ dnorm(0,10),
    bfreq ~ dnorm(0,10),
    bcap ~ dnorm(0,10)
  ),
  data=bmdat)

## Note for m2: a model with the suspicion that when one considers the old
## way
## of analysis is inadequate and one's capability is good enough,
## one will tend to learn and use Bayesian method

> compare(m1,m2,m3)
      WAIC pWAIC dWAIC weight     SE     dSE
m3 28.6    8.6    0.0    0.90 13.79     NA
m1 33.1    2.4    4.5    0.09  9.99 10.31
m2 42.3    9.2   13.7    0.00 14.75   8.33

> precis(m3)
```

```

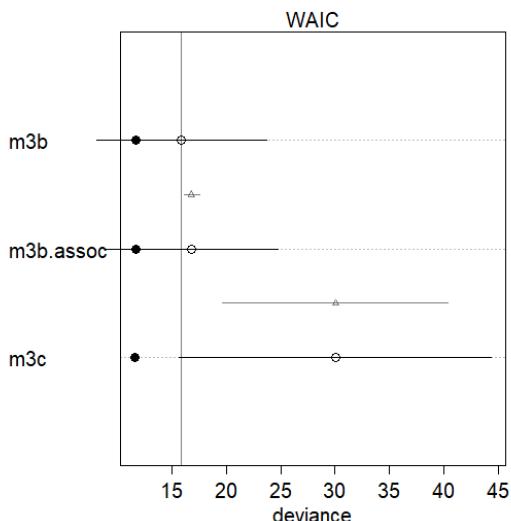
      Mean StdDev 5.5% 94.5%
a     -1.71   6.60 -12.26  8.84
bfreq -2.01   6.59 -12.54  8.53
bcap    7.43   3.58   1.71 13.16

> logistic(-1.71)
[1] 0.1531637
> logistic(-1.71+7.43)
[1] 0.996731

> compare(m3b,m3b.assoc,m3c)
      WAIC pWAIC dWAIC weight    SE    dSE
m3b      15.3    1.9    0.0    0.61  7.86    NA
m3b.assoc 16.2    2.3    0.9    0.39  7.80    0.45
m3c      30.5    9.4   15.1    0.00 14.44 10.58

> plot(compare(m3b,m3b.assoc,m3c))

```

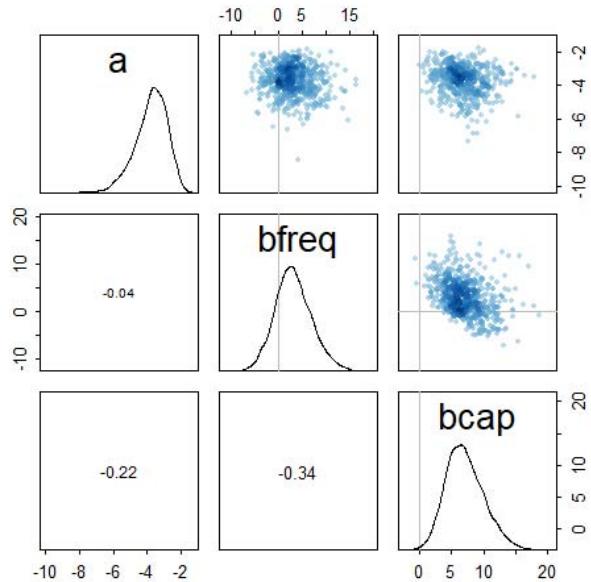


```

> precis(m3b.assoc)
      Mean StdDev 5.5% 94.5%
a     -3.50   0.85 -4.85 -2.14
bfreq  1.85   3.34 -3.49  7.19
bcap    5.32   2.42  1.46  9.18

# HMC Stan
m3b.assoc.stan <- map2stan(m3b.assoc, data=bmdat, iter=10000, warmup=1000)
# check Gaussian
pairs(m3b.assoc.stan)

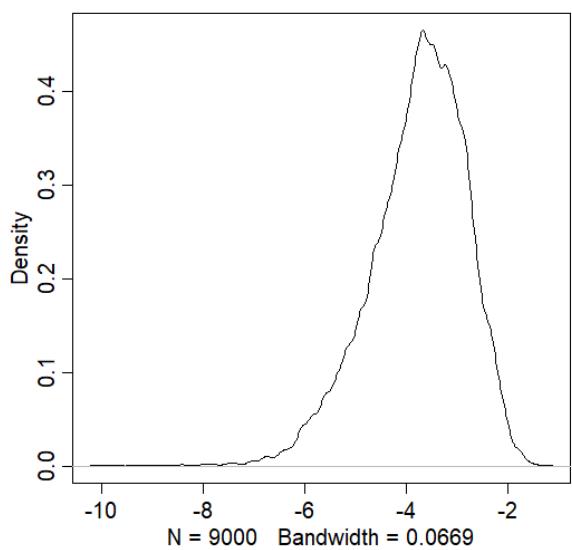
```



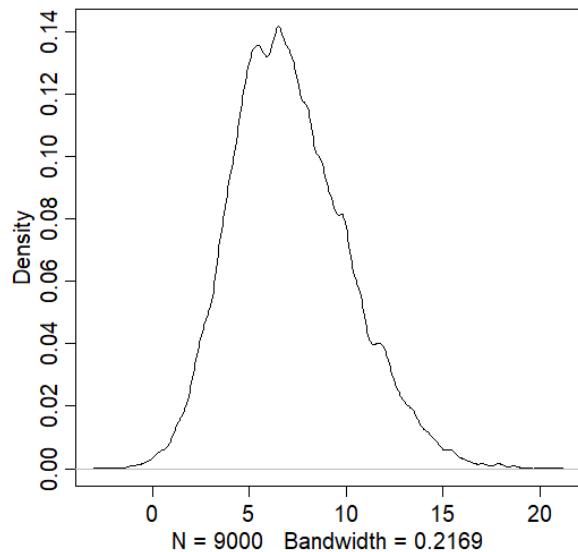
```

> precis(m3b.assoc.stan)
      Mean StdDev lower 0.89 upper 0.89 n_eff Rhat
a     -3.78    0.94    -5.24    -2.31    4728     1
bfreq  3.02    3.89    -2.91     9.54    4114     1
bcap   7.11    2.98     2.36    11.77    3652     1
> post <- extract.samples(m3b.assoc.stan)
> str(post)
List of 3
 $ a     : num [1:9000(1d)] -4.42 -4.6 -2.95 -3.94 -2.11 ...
 $ bfreq: num [1:9000(1d)] 0.298 4.228 4.957 5.496 3.666 ...
 $ bcap  : num [1:9000(1d)] 6.41 5.34 3.91 3.98 3.93 ...
> dens(post$a)

```

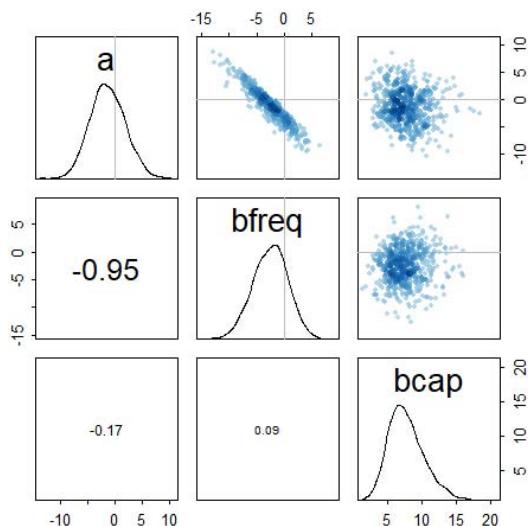


```
> dens(post$bcap)
```

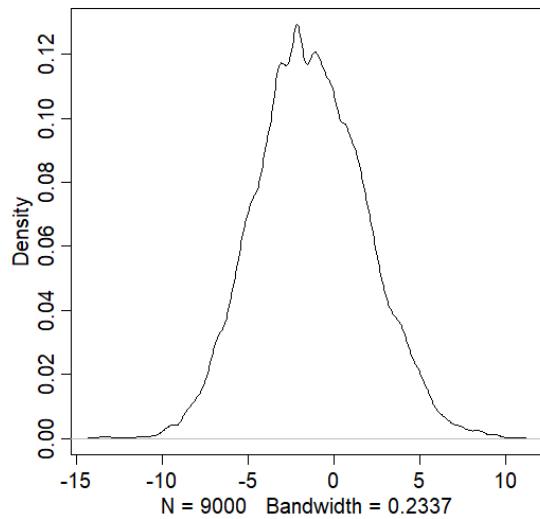


```
# MCMC Stan for m3b model
> m3b.stan <- map2stan(m3b, data=bmdat, iter=10000, warmup=1000)
> precis(m3b.stan)
  Mean StdDev lower 0.89 upper 0.89 n_eff Rhat
a     -1.33    3.21      -6.24      4.01  2489    1
bfreq -2.55    3.20      -7.88      2.29  2480    1
bcap    7.78    2.61      3.55     11.61  2511    1

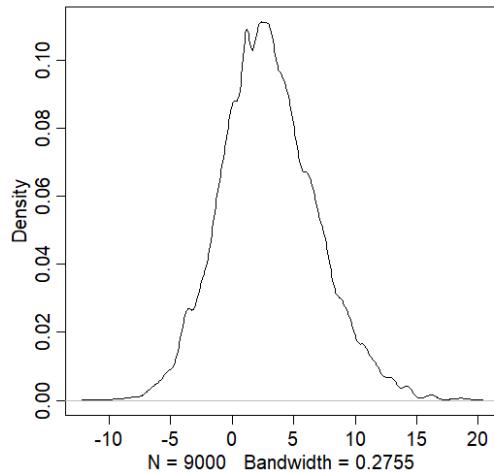
> pairs(m3b.stan)
```



```
> post2 <- extract.samples(m3b.stan)
> dens(post2$a)
```



```
> dens(post$bfreq)
```



Stan MCMC performs quite well.

References

- McElreath R. (2018). Statistical Rethinking: A Bayesian Course with Examples in R and Stan. New York: Chapman and Hall/CRC.
- Vuong QH. (2018). The (ir)rational consideration of the cost of science in transition economies. *Nature Human Behaviour*, 2(1), 5.
- Vuong QH, La VP, Vuong TT,... & Ho MT. (2018). Cultural additivity: Behavioural insights from the interaction of Confucianism, Buddhism, and Taoism in folktales. *Palgrave Communications*, 4(1), 143.