

BDA - Project

Anonymous

12/6/2021

Introduction

The motivation

Neonatal mortality is related to the number of deaths in the first 28 days of a child's life. This is a high-risk period and according to UNICEF data, the average number of deaths is estimated to be 17 per 1000 live births in 2019 [1]. Neonatal mortality is classified into an early neonatal mortality which is between the first 7 days of birth and a late neonatal mortality which occurs after 7 days of birth. On the other hand, perinatal mortality refers to the death of the child during the fetal period in conjunction to early neonatal mortality. It is clear that there are huge differences in perinatal and neonatal mortality rates between continents and between developed and emerging countries. This project will attempt to model the probability of perinatal and late neonatal mortality in 2020 in different regions in Colombia (a developing country in Latin America) and compare it with the US, which is among the richest countries in the world.

The main modeling idea

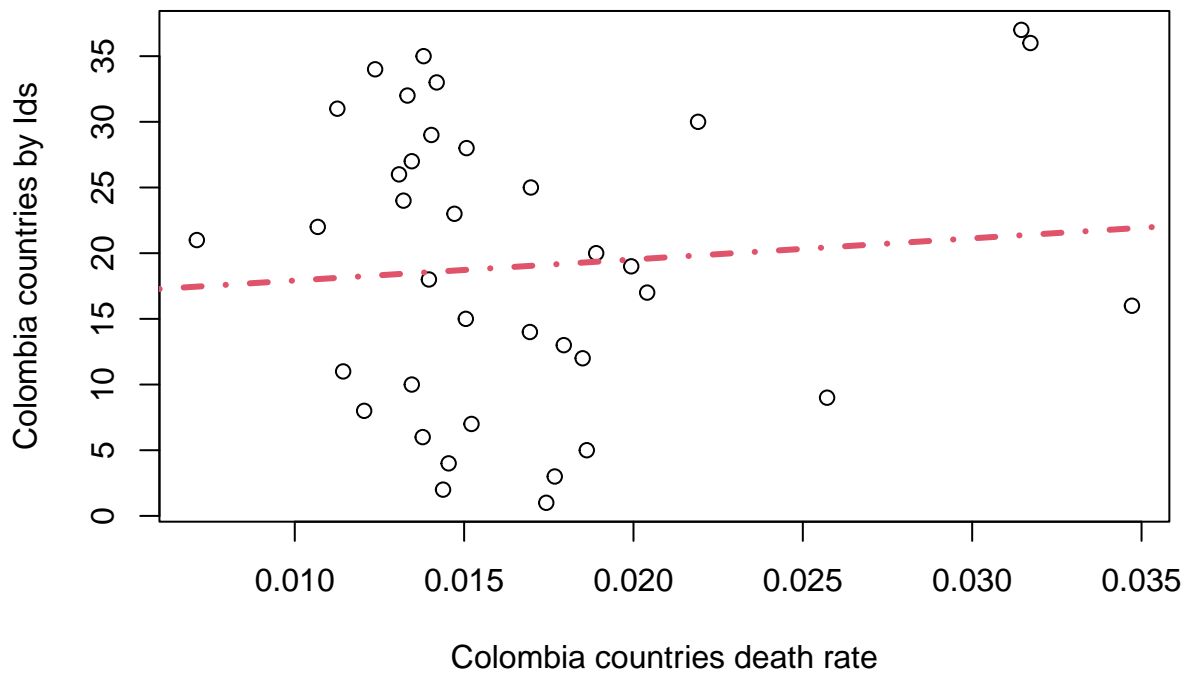
The general idea for modeling the probability of perinatal and late neonatal mortality is to treat the cases of death from this condition as “successes” among the total number of births in the same time period. With this in mind, we came up with a beta-binomial model that is natural in these cases.

Some illustrative figure

```
# Read Excel file and convert it to DataFrame
library("readxl")
library("rstan")
exceldata = read_excel("Colombia2020.xlsx")
dfData= data.frame(exceldata)
deaths = dfData$Deaths
births = dfData$Lbirths+ dfData$Deaths
dfData$rate = (deaths/births)

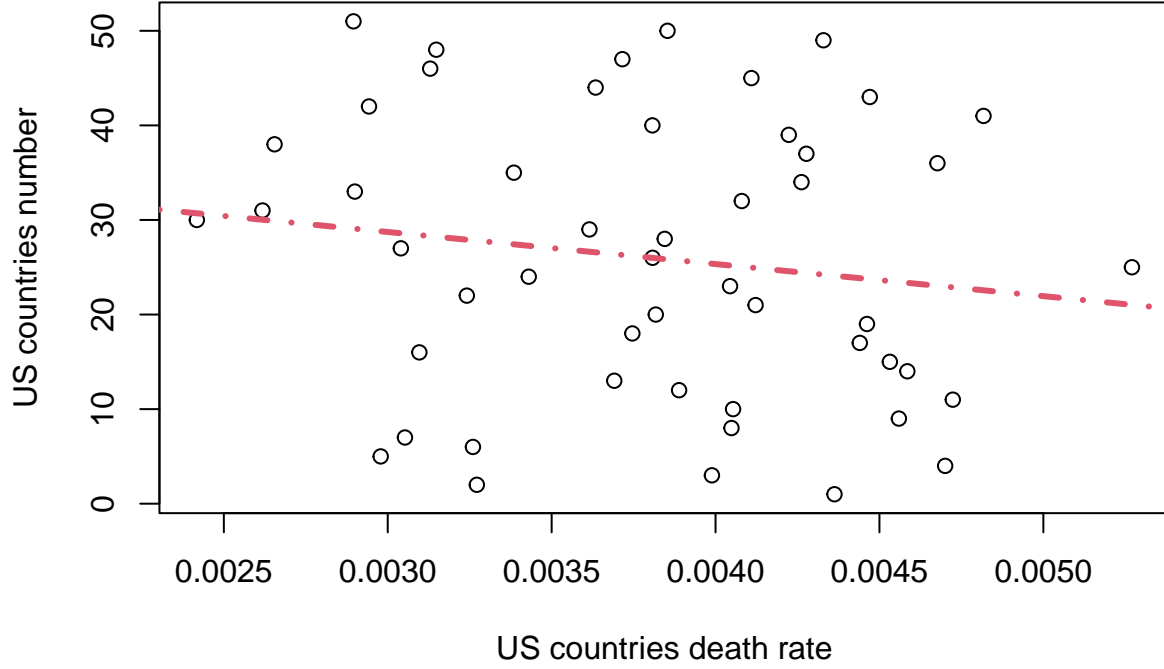
dfData$id <- seq.int(nrow(dfData))
plot(dfData$id ~ dfData$rate, title="Perinatal and neonatal mortality for Colombia",
     ylab="Colombia countries by Ids",
     xlab="Colombia countries death rate")
exceldata = read_excel("a.xlsx")
dfA= data.frame(exceldata)
exceldata = read_excel("b.xlsx")
```

```
dfB= data.frame(exceldata)
lm1 <- lm(id ~ rate, data = dfData)
abline(lm1, col = 2, lty = 4, lw = 3)
```



And here about the US data:

```
exceldata = read_excel("US2020.xlsx")
dfData= data.frame(exceldata)
deaths = dfData$Deaths
births = dfData$Births
dfData$rate = (deaths/births)
dfData$id <- seq.int(nrow(dfData))
plot(dfData$id ~ dfData$rate, title="The perinatal and neonatal mortality for US",
      xlab="US countries death rate", ylab="US countries number")
exceldata = read_excel("aUS.xlsx")
dfA= data.frame(exceldata)
exceldata = read_excel("bUS.xlsx")
dfB= data.frame(exceldata)
lm1 <- lm(id ~ rate, data = dfData)
abline(lm1, col = 2, lty = 4, lw = 3)
```



Data

For Colombia, data were found on the website of the National Institute of Health of the country, data were collected manually in an excel file from epidemiological reports from the year 2017 to 2020. The data collected were the number of births and the number of deaths in the perinatal and late neonatal period by year and in 38 regions. Data for the United States was collected from the CDC wonder website in the Infant Death Records. In the case of Colombia, there are no published models for this type of infant mortality and only predictive models for the whole country were found in terms of US data.

Model description

The following section describe the two models used in this problem, hierarchical and separate.

Separate model

The parameter of interest is θ which corresponds to the probability late neonatal and perinatal death among regions. The likelihood is computed as:

$$p(y_i|\theta_i) = \binom{n_i}{y_i} \theta_i^{y_i} (1 - \theta_i)^{n_i - y_i}$$

Being y_i the number of deaths, n_i the number of births for each region.

Prior

Informative priors were chosen in both models since we also have data from previous years. The mean of a and b values for the previous years were used in the hierarchical model and separate models.

The prior in this case was obtained using beta distribution since parameter a resemble number of successes and $a + b$ the total number of trials.

So, the prior in this case was computed as:

$$\theta_i \sim \text{Beta}(\bar{a}_i, \bar{b}_i)$$

Being \bar{a}_i, \bar{b}_i the mean of of deaths and live births between each region and computed from previous years.

Hierarchical model

The likelihood remains the same as in the previous model. But in this case we use a transformation to compute hyper-prior.

Parameter μ is equal to the probability of death, that is $a/(a+b)$ and η corresponds to the total population ($a+b$).

Hyper-priors

$$\eta \sim \text{exponential}(x), \mu_i \sim \text{Beta}(a_i, b_i)$$

Prior

$$\theta_i \sim \text{Beta}(\alpha, \beta)$$

given that $\alpha = \mu\eta$ and $\beta = (1 - \mu)\eta$.

Stan code

Separate model

```
write("// Stan Separate model
//Beta-binomial Separate model

data {
  int<lower=0> N; // Number of states
  int<lower=0> y[N]; //Number of neonatal deaths
  int<lower=0> n[N]; //Number of births
  real aMean[N]; //Minimun a value
  real bMean[N]; //Minimun b Value
}

// The parameters accepted by the model
```

```

parameters {
  vector<lower=0, upper=1>[N] theta;
}

// The model to be estimated. We model the output
// 'y' to be normally distributed with mean 'mu'
// and standard deviation 'sigma'.
model {

  //Priors

  for (j in 1:N) {
    theta[j] ~ beta(aMean[j], bMean[j]);
  }

  //Likelihood
  for (k in 1:N){
    y[k] ~ binomial(n[k], theta[k]);
  }

}

generated quantities {

  //Log Likelihood ratios
  vector[N] log_lik;
  real ypred[N];

  for(j in 1:N){
    log_lik[j] = binomial_lpmf(y[j] | n[j], theta[j]);
  }

  for(j in 1:N){
    ypred[j]= binomial_rng(n[j],theta[j]);
  }

}
// The posterior predictive distribution",
"separate_model.stan")

```

Hierarchical model

```

write("// Stan Beta-binomial Hierarchical model
data {

  int<lower=0> N; // Number of states
  int<lower=0> y[N]; //Number of neonatal deaths
  int<lower=0> n[N];
  real a[N]; //Mean a value
  real b[N]; //Mean b Value

```

```

    real<lower=0> e;
}

// The parameters accepted by the model
parameters {
    real<lower=0,upper=1> mu;
    real<lower=0> eta;
    real<lower=0,upper=1> theta[N];
}

transformed parameters {
    real<lower=0> alpha;
    real<lower=0> beta;
    alpha = eta* mu ;
    beta = eta*(1-mu);
}

model {
    //Hyper-priors
    eta ~ exponential(e);

    //Prior
    for (k in 1:N){

        mu ~ beta(a[k],b[k]);
        theta[k] ~ beta(alpha,beta);

    //Likelihood
        y[k] ~ binomial(n[k], theta[k]);
    }

}

generated quantities {

    vector[N] log_lik;
    real ypred[N];

    for(i in 1:N){
        log_lik[i] = binomial_lpmf(y[i]|n[i], theta[i]);
    }

    for(j in 1:N){
        ypred[j]= binomial_rng(n[j],theta[j]);
    }
}

// The posterior predictive distribution",
"hierachichal_model.stan")

```

Running the model

US data

Separate model

First we need to read the data and transform it into a data frame.

```
library("readxl")
library("rstan")
exceldata = read_excel("US2020.xlsx")
dfData= data.frame(exceldata)
```

Then we read the files that have the mean and of the number of live births and the number of deaths in previous years. In the case of US average was obtained from 2007-2018

```
exceldata = read_excel("aUS.xlsx")
dfA= data.frame(exceldata)
exceldata = read_excel("bUS.xlsx")
dfB= data.frame(exceldata)
```

Then we compute the model with the data

```
library(loo)

deaths = dfData$Deaths
births = dfData$Births

stan_data <- list(
  N = 51,
  y = deaths,
  n = births,
  aMean = dfA$Mean,
  bMean = dfB$Mean
)

sm <- rstan::stan_model(file = "separate_model.stan")
separate_modelUS <- rstan::sampling(sm, data = stan_data, refresh= 0)
```

Hierarchical model

```
sm <- rstan::stan_model(file = "hierarchical_model.stan")
stan_data_hierachichal <- list(
  N=51,
  y = deaths,
  n = births,
  a = dfA$Mean,
  b = dfB$Mean,
  e=1/60000
)
hierarchical_modelUS <- rstan::sampling(sm, data = stan_data_hierachichal,refresh=0)
```

Colombia data

First we need to read the data and transform it into a data frame.

```
exceldata = read_excel("Colombia2020.xlsx")
dfData= data.frame(exceldata)
```

Then we read the files that have the mean and of the number of live births and the number of deaths in previous years. In the case of Colombia these values were obtained using the mean from 2017 to 2019.

```
exceldata = read_excel("a.xlsx")
dfA= data.frame(exceldata)
exceldata = read_excel("b.xlsx")
dfB= data.frame(exceldata)
```

Separate model

```
library(bayesplot)
library(loo)

deaths = dfData$Deaths
births = dfData$Lbirths+ dfData$Deaths

stan_data <- list(
  N = 37,
  y = deaths,
  n = births,
  aMean = dfA$Mean,
  bMean = dfB$Mean
)

sm <- rstan::stan_model(file = "separate_model.stan")
separate_modelCol <- rstan::sampling(sm, data = stan_data, refresh= 0)
```

Hierarchical model

```
sm <- rstan::stan_model(file = "hierarchical_model.stan")

stan_data_hierachichal <- list(
  N=37,
  y = deaths,
  n = births,
  a = dfA$Mean,
  b = dfB$Mean,
  e=1/20000
)

hierarchical_modelCol <- rstan::sampling(sm, data = stan_data_hierachichal,refresh=0)
```


Convergence diagnostics (\hat{R} , ESS, divergences) and what was done if the convergence was not good with the first try.

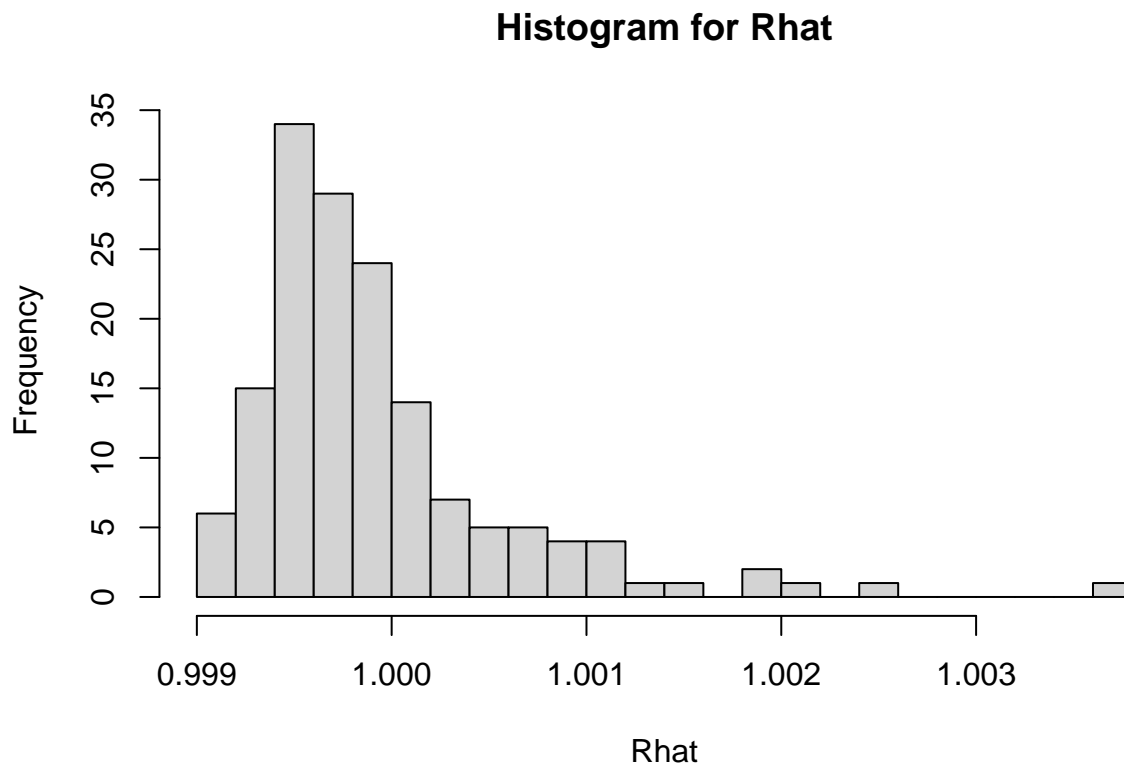
For convergence diagnostics Rhat and Bulk_ESS values were used. This statistic compares between chain and within chain draws for each of the model parameters. If chains have not converged well, Rhat will be larger than 1.

US Data

Separate model

Below there are the histograms of Rhat for each one of the models:

```
s <- summary(separate_modelUS)$summary
Rhat <- s[,10]
hist(Rhat,
     main="Histogram for Rhat",
     xlab="Rhat",
     breaks=20)
```



```
check_divergences(separate_modelUS)
```

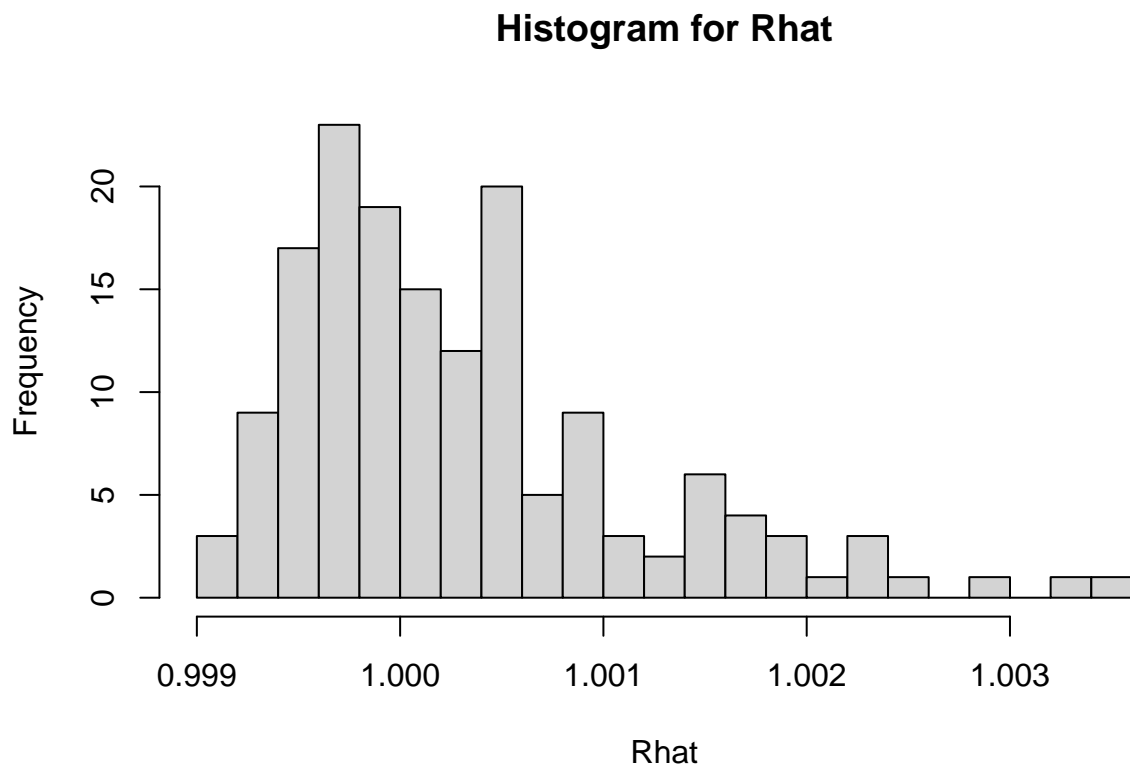
```
## 0 of 4000 iterations ended with a divergence.
```

```
check_treedepth(separate_modelUS)
```

0 of 4000 iterations saturated the maximum tree depth of 10.

Hierarchical model

```
s <- summary(hierarchical_modelUS)$summary  
Rhat <- s[,10]  
hist(Rhat,  
     main="Histogram for Rhat",  
     xlab="Rhat",  
     breaks=20)
```



```
check_divergences(hierarchical_modelUS)
```

0 of 4000 iterations ended with a divergence.

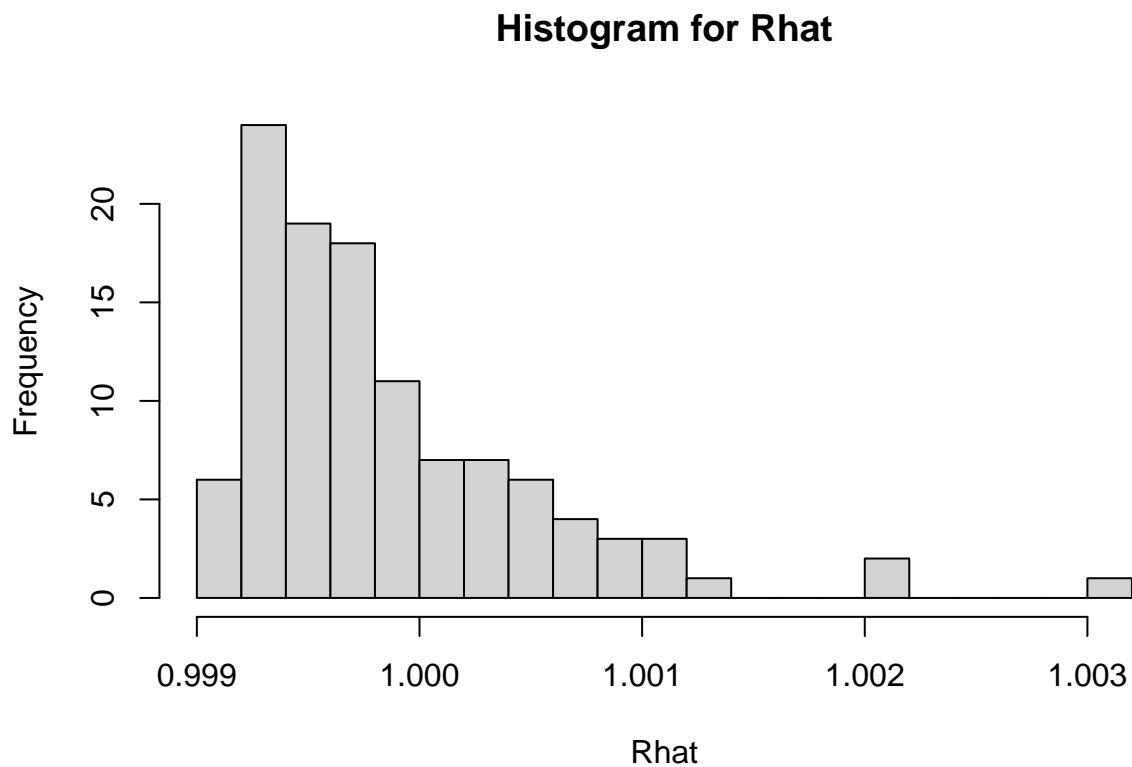
```
check_treedepth(hierarchical_modelUS)
```

0 of 4000 iterations saturated the maximum tree depth of 10.

Colombia data

Separate model

```
s <- summary(separate_modelCol)$summary
Rhat <- s[,10]
hist(Rhat,
     main="Histogram for Rhat",
     xlab="Rhat",
     breaks=20)
```



```
check_divergences(separate_modelCol)
```

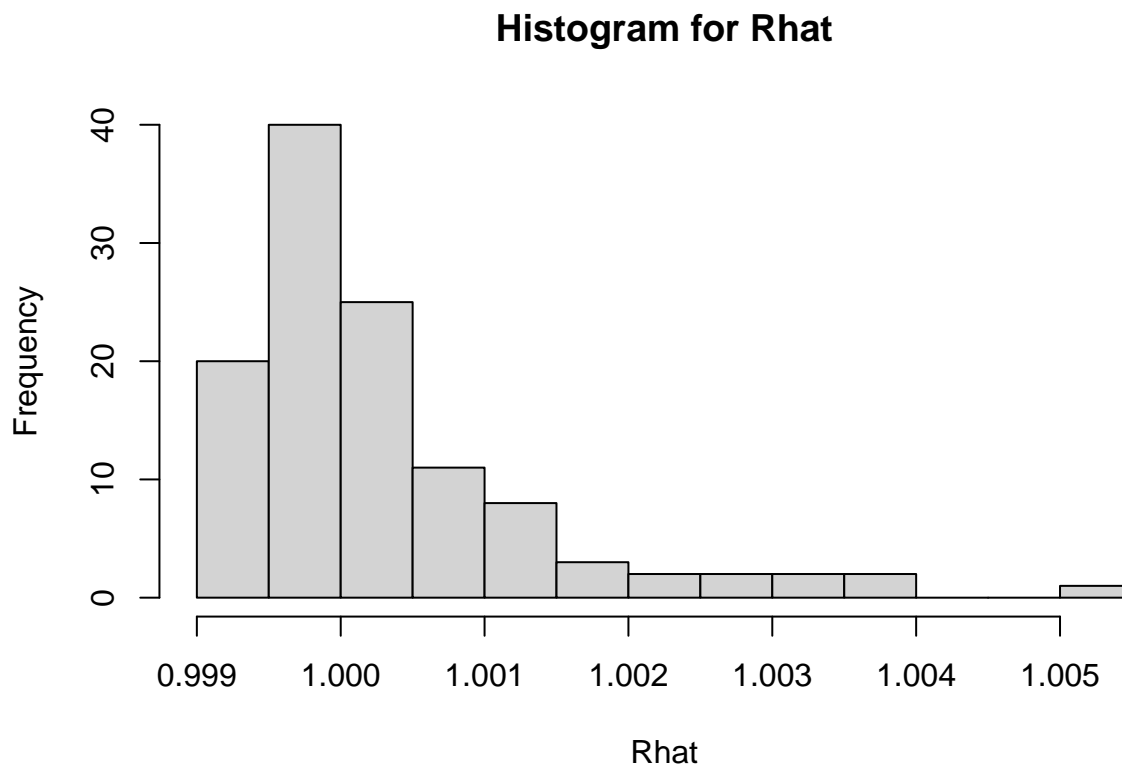
```
## 0 of 4000 iterations ended with a divergence.
```

```
check_treedepth(separate_modelCol)
```

```
## 0 of 4000 iterations saturated the maximum tree depth of 10.
```

Hierarchical model

```
s <- summary(hierarchical_modelCol)$summary
Rhat <- s[,10]
hist(Rhat,
     main="Histogram for Rhat",
     xlab="Rhat",
     breaks=20)
```



```
check_divergences(hierarchical_modelCol)
```

```
## 0 of 4000 iterations ended with a divergence.
```

```
check_treedepth(hierarchical_modelCol)
```

```
## 0 of 4000 iterations saturated the maximum tree depth of 10.
```

In all of these histograms we see that Rhat values are really close to 1 and none of Rhat values are below 1.05 which is recommended in Stan official website.

Additionally if you check the **Appendix** section and **Convergence monitoring** all values of Bulk Effective sample size are above 100 so the models are considered good.

Finally, there were not any warnings regarding iterations ending in divergence or saturating the maximum tree depth of 10 for any of the models.

Model improvements

Separate model

For the separate model, the prior was first tried to be computed using another hyper-prior information assuming that parameters a (number of deaths) and b (number of living births) will follow a normal distribution based on past data. The following lines contain the previous model priors:

```
model {

  //Hyper priors
  for (i in 1:N) {
    a[i] ~ normal(aMean[i], aStd[i]) T[L,]; //Number of successes parameter
    b[i] ~ normal(bMean[i], bStd[i]) T[L,]; //Number of no Success
  }

  //Priors

  for (j in 1:N) {
    theta[j] ~ beta(a[j], b[j]);
  }

  //Likelihood
  for (k in 1:N){
    y[k] ~ binomial(n[k], theta[k]);
  }
}
```

However, doing the k-pareto analysis, there were many values between 0.7 and 1 and even higher than 1, so this choice was discarded.

Hierarchical model

Many values were chosen for hyper-priors of parameters μ and η . We start using gamma function for parameter η and one single distribution for μ . Like this:

```
model {

model {
  //Hyper-priors
  eta ~ gamma(s,t);

  mu ~ beta(a,b);

  //Prior
  for (k in 1:N){
    theta[k] ~ beta(alpha,beta);

  //Likelihood
```

```

      y[k] ~ binomial(n[k], theta[k]);
    }
  }
}

```

However this common parameter η distribution was changed to exponential since it is a weakly hyper-prior and μ different distributions were obtained depending on the region. This changes improved k-pareto diagnostics a little bit.

Model comparison

US Data

Separate model

```

separate_extract_log_lik <- extract_log_lik(separate_modelUS,
                                           parameter_name = "log_lik",
                                           merge_chains = FALSE);
r_eff <- relative_eff(exp(separate_extract_log_lik), cores = 2)
separate_model_loo <- loo(separate_extract_log_lik, r_eff = r_eff, cores = 2)
print(separate_model_loo)

```

```

##
## Computed from 4000 by 51 log-likelihood matrix
##
##           Estimate   SE
## elpd_loo    -232.7  9.4
## p_loo        30.9  4.6
## looic        465.5 18.9
## -----
## Monte Carlo SE of elpd_loo is NA.
##
## Pareto k diagnostic values:
##           Count Pct.    Min. n_eff
## (-Inf, 0.5] (good)   18   35.3%   451
## (0.5, 0.7] (ok)     24   47.1%    98
## (0.7, 1] (bad)      9   17.6%    23
## (1, Inf) (very bad) 0    0.0%   <NA>
## See help('pareto-k-diagnostic') for details.

```

Hierarchical model

```

hierarchical_extract_log_lik <- extract_log_lik(hierarchical_modelUS,
                                                parameter_name = "log_lik",
                                                merge_chains = FALSE);
r_eff <- relative_eff(exp(hierarchical_extract_log_lik), cores = 2)
hierarchical_model_loo <- loo(hierarchical_extract_log_lik, r_eff = r_eff, cores = 2)
print(hierarchical_model_loo)

```

```
##
## Computed from 4000 by 51 log-likelihood matrix
##
##           Estimate   SE
## elpd_loo   -229.3  5.7
## p_loo       34.0  2.7
## looic       458.5 11.5
## -----
## Monte Carlo SE of elpd_loo is NA.
##
## Pareto k diagnostic values:
##           Count Pct.   Min. n_eff
## (-Inf, 0.5] (good)    3    5.9%   864
## (0.5, 0.7] (ok)      17   33.3%   123
## (0.7, 1] (bad)       26   51.0%    20
## (1, Inf) (very bad)  5    9.8%    11
## See help('pareto-k-diagnostic') for details.
```

Colombia Data

Separate model

```
separate_extract_log_lik <- extract_log_lik(separate_modelCol,
                                             parameter_name = "log_lik",
                                             merge_chains = FALSE);
r_eff <- relative_eff(exp(separate_extract_log_lik), cores = 2)
separate_model_loo <- loo(separate_extract_log_lik, r_eff = r_eff, cores = 2)
print(separate_model_loo)
```

```
##
## Computed from 4000 by 37 log-likelihood matrix
##
##           Estimate   SE
## elpd_loo   -165.0  6.7
## p_loo       21.8  3.9
## looic       330.1 13.5
## -----
## Monte Carlo SE of elpd_loo is NA.
##
## Pareto k diagnostic values:
##           Count Pct.   Min. n_eff
## (-Inf, 0.5] (good)   11   29.7%   739
## (0.5, 0.7] (ok)      18   48.6%   413
## (0.7, 1] (bad)       7    18.9%    81
## (1, Inf) (very bad)  1    2.7%    12
## See help('pareto-k-diagnostic') for details.
```

Hierarchical model

```
hierarchical_extract_log_lik <- extract_log_lik(hierarchical_modelCol,
                                              parameter_name = "log_lik",
                                              merge_chains = FALSE);
r_eff <- relative_eff(exp(hierarchical_extract_log_lik), cores = 2)
hierarchical_model_loo <- loo(hierarchical_extract_log_lik, r_eff = r_eff, cores = 2)
print(hierarchical_model_loo)
```

```
##
## Computed from 4000 by 37 log-likelihood matrix
##
##           Estimate   SE
## elpd_loo   -174.1  5.8
## p_loo       32.6  4.2
## looic       348.1 11.7
## -----
## Monte Carlo SE of elpd_loo is NA.
##
## Pareto k diagnostic values:
##           Count Pct.    Min. n_eff
## (-Inf, 0.5] (good)     0    0.0%    <NA>
## (0.5, 0.7] (ok)       8   21.6%    118
## (0.7, 1] (bad)      23   62.2%     30
## (1, Inf) (very bad)  6   16.2%      3
## See help('pareto-k-diagnostic') for details.
```

Looking at k-pareto analysis since the majority of k values are below 0.7 in the separate model, this model was chosen as the best for both data sets. In colombian data the elpd_loo is greater in the separate model, that is another reason why this model was chosen.

Posterior checking and results

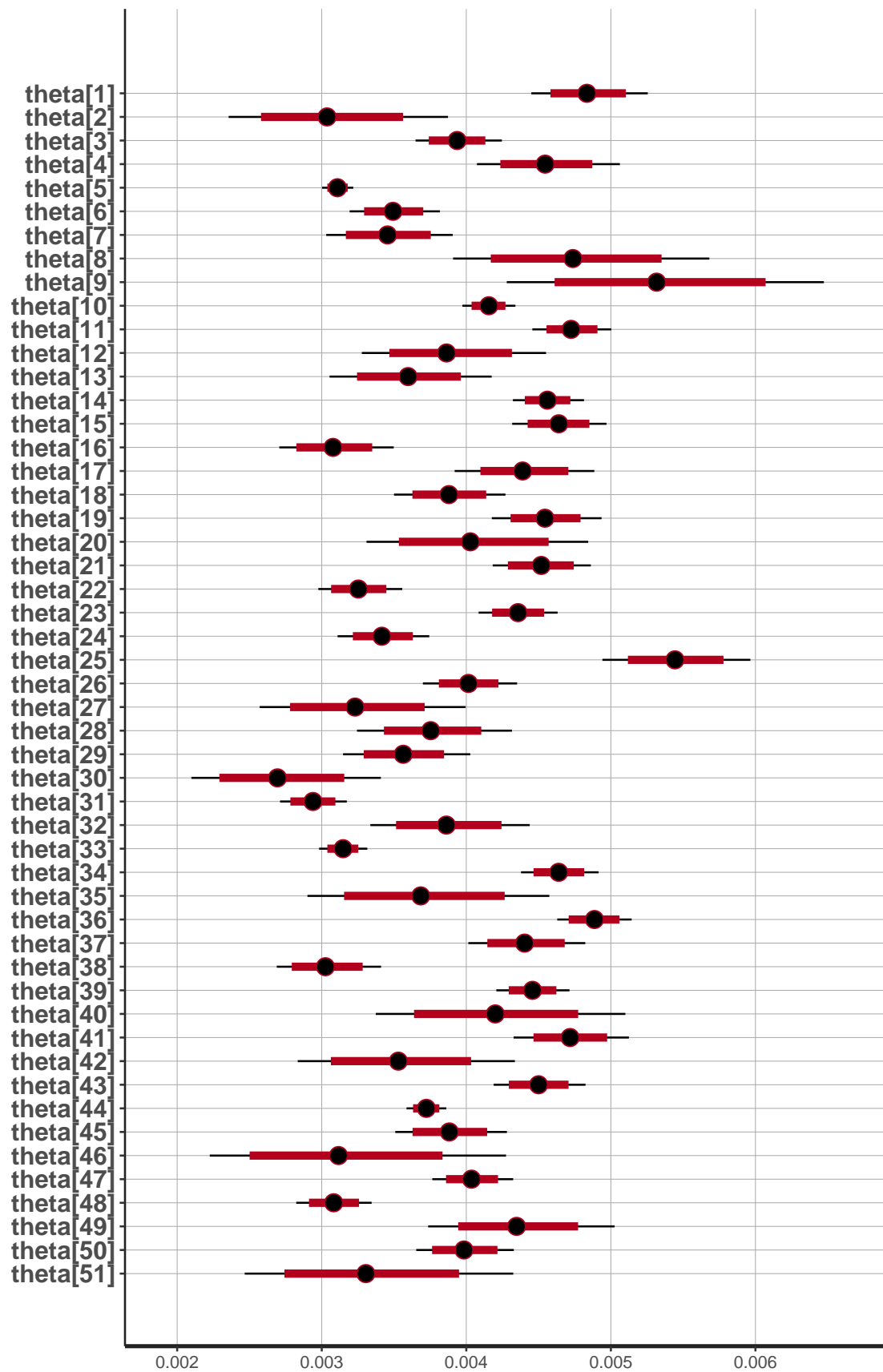
As the best model was the separate, we will use the posterior draws of this model to compare the theta parameter for the two countries.

The US:

```
plot(separate_modelUS, pars=c('theta'))
```

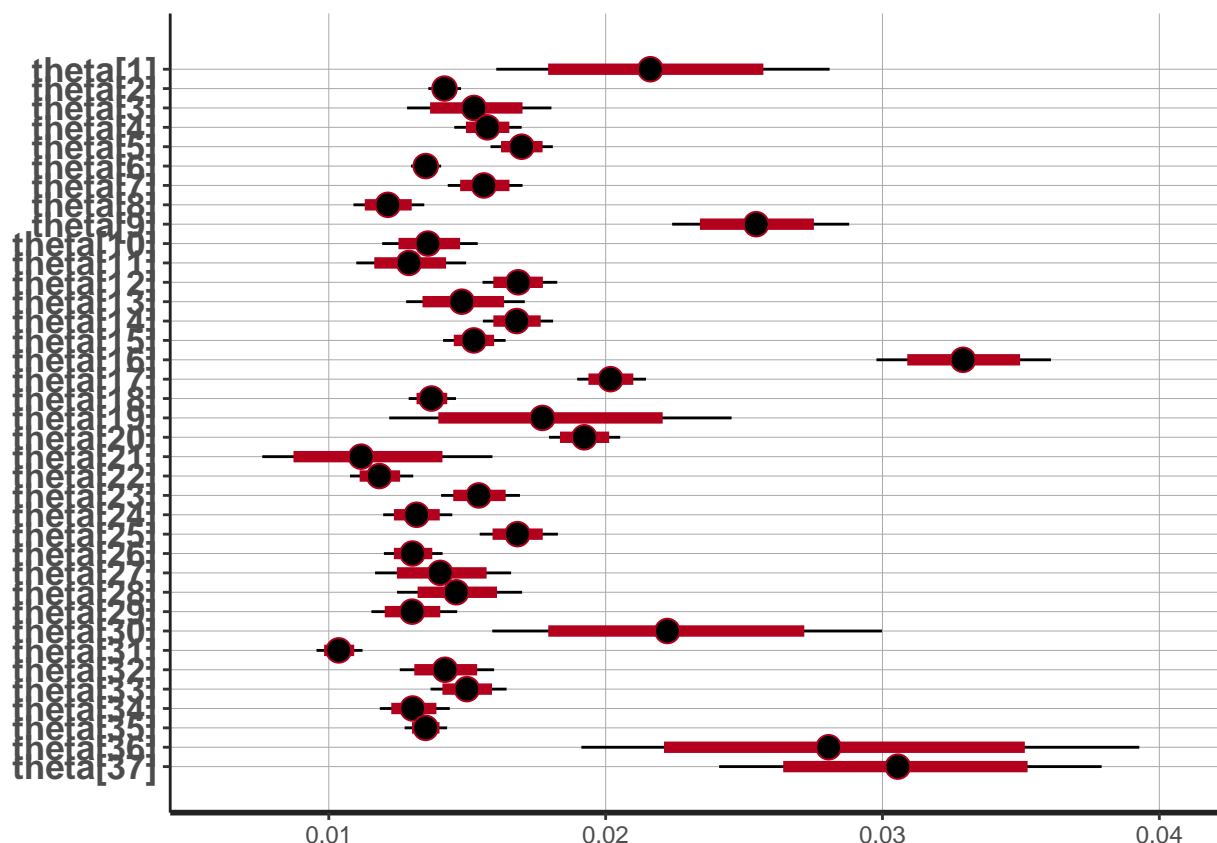
```
## ci_level: 0.8 (80% intervals)
```

```
## outer_level: 0.95 (95% intervals)
```

Colombia:

```
plot(separate_modelCol, pars=c('theta'))
```



As you can see from the figures, which represent the confidence interval and the mean value of the probability of death in the perinatal and late neonatal period, the probability parameter is drastically larger among all Colombian states compared to the ones in the US.

Regarding within-country comparison and using these posterior draws summary graph we can rank the top 5 regions of both countries with worst perinatal and late neonatal mortality rate.

In Colombia the top 5 worst states are: 5. Cordoba ($\bar{\theta}$) of 0.0202, 4. Buenaventura ($\bar{\theta}$) of 0.0284 3. Vaupes ($\bar{\theta}$) of 0.0284, 2. Vichada ($\bar{\theta}$) of 0.0306 and 1. Choco with a ($\bar{\theta}$) of 0.0330.

In the US, the top 5 worst states are 5. South Carolina ($\bar{\theta}$) of 0.00472, 4. Alabama ($\bar{\theta}$) of 0.00484 3. Ohio ($\bar{\theta}$) of 0.00488, 2. Columbia ($\bar{\theta}$) of 0.00534 and 1. Mississippi with a ($\bar{\theta}$) of 0.00545.

It is interesting that Choco, Vichada and Vaupes are among the top 5 poorest states in Colombia [2] while South Carolina, Columbia, Mississippi and Alabama are between the top 10 poorest states in the US [3].

With this in mind, there is reason to suspect that the state wealth variable is related to the rate of perinatal and late neonatal mortality.

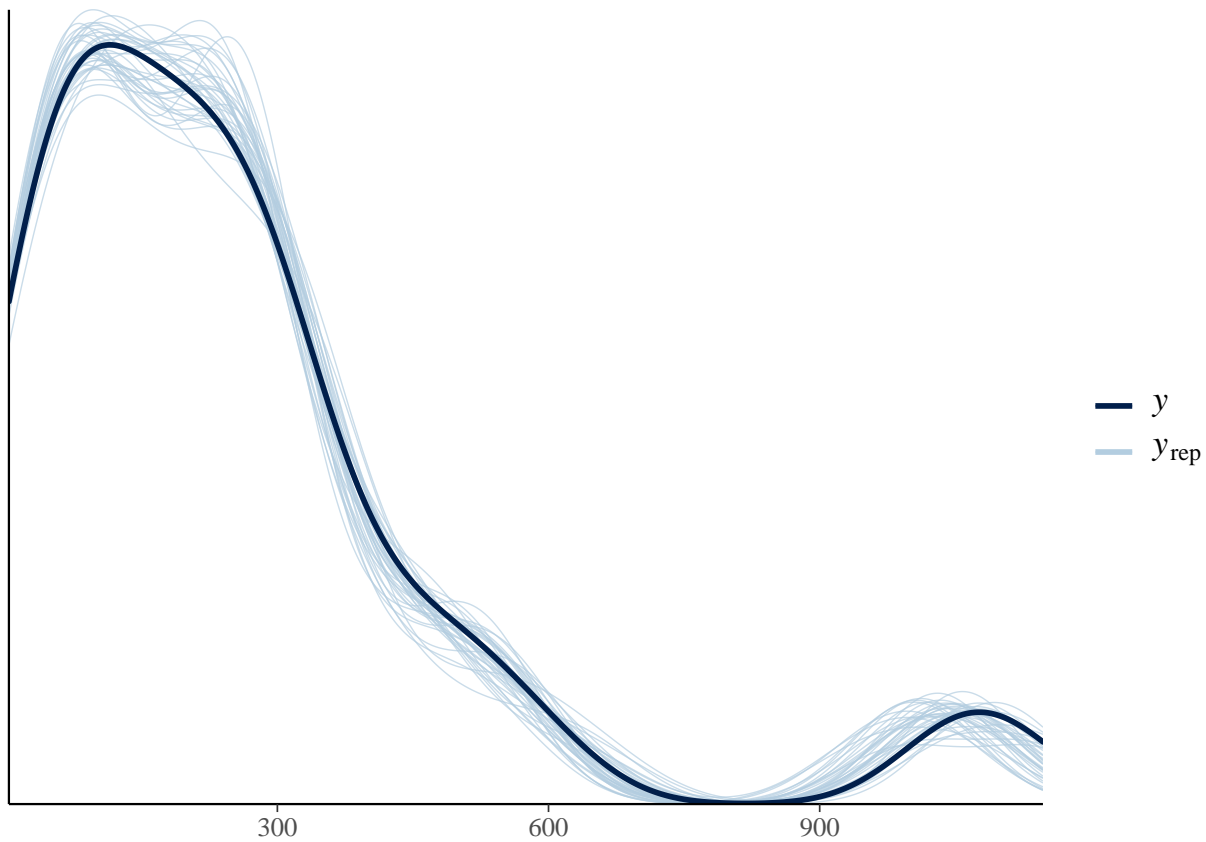
Posterior predictive checking

The posterior predictive checking was done using the best model which was the separate one in Colombian data.

```
y_pred <- as.matrix(separate_modelCol, pars = "ypred")
dim(y_pred)
```

```
## [1] 4000 37
```

```
ppc_dens_overlay(deaths, y_pred[1:37, ])
```



As you can see from the figure, the predictions are quite similar to the real value of y (number of deaths in neonatal and perinatal period). The predictive performance is quite good.

Sensitivity analysis with respect to prior choices (i.e. checking whether the result changes a lot if prior is changed)

Separate model

As mentioned in model improvements, some other priors were chosen before, specifically hyper-priors

```
//Hyper priors
for (i in 1:N) {
  a[i] ~ normal(aMean[i], aStd[i]) T[L,]; //Number of successes parameter
  b[i] ~ normal(bMean[i], bStd[i]) T[L,]; //Number of no Success
}
```

The results of the posterior draws were almost the same but when performing k-pareto diagnostics, results were worse (more k values above 0.7).

Hierarchical

As mentioned in model improvements, some other priors were chosen before, specifically hyper-priors

```
//Hyper-priors  
eta ~ gamma(s,t);
```

Many values were manually selected from s, t to resemble the distribution of the total number of births, but the k-pareto diagnostics gave worse results (more k values above 0.7).

Discussion of issues and potential improvements.

The hierarchical model did not perform well in k-pareto diagnostics, many k values were above 0.7. Maybe to improve our model we can work with the data points of the previous years, first to improve k-values and second to make a predictive model by state so that the probability of perinatal and neonatal mortality by state can be forecasted in future years. Probably using models that fit data series such as ARIMA or ARIMAX.

Conclusion what was learned from the data analysis.

From our point of view, we learned and reinforced how to model using hierarchical and non-hierarchical models in case of binomial trials. Also we learn how to work on the model (choosing priors is not easy!) and also how to do all kinds of calculations to get information about convergence and how well the model fits the data. We also learned about model comparison and basically we have the tools to do Bayesian modeling in any area.

Self-reflection of what the group learned while making the project.

We as a group learned how we can work together first in finding the data then doing some research to find the appropriate model for the problem we want to solve. We learned how we can improve models by sharing our knowledge and experience and finally doing our best together to improve the proposed model and solution to the related problem

References

- [1] “Neonatal Mortality.” UNICEF DATA, 20 July 2021, <https://data.unicef.org/topic/child-survival/neonatal-mortality/>.
- [2] Semana Magazine. “Pobreza En Colombia: Estos Son Los Departamentos Más Afectados.” Semana.com Últimas Noticias De Colombia y El Mundo, 21 Dec. 2020, <https://www.semana.com/economia/articulo/pobreza-en-colombia-estos-son-los-departamentos-mas-afectados/202026/>.
- [3] November 10, 2021. “Top 10 Poorest States in the U.S.” Friends Committee On National Legislation, <https://www.fcnl.org/updates/2021-11/top-10-poorest-states-us>. Vehtari, Aki, and Markus Paasiniemi. “BDA3 Demos Comparison of 2 Groups with Binomial.” Aalto University. Niemi, Jarad. “Hierarchical Models.” PPT file. Iowa State University, 2019. Hu, Jim Albert and

Jingchen. “Probability and Bayesian Modeling.” Chapter 10 Bayesian Hierarchical Modeling, 30 July 2020, <https://bayesball.github.io/BOOK/bayesian-hierarchical-modeling.html>. Brody-Moore, Peter, “Bayesian Hierarchical Meta-Analysis of Asymptomatic Ebola Seroprevalence” (2019). CMC Senior Theses. 2228.https://scholarship.claremont.edu/cmc_theses/2228

Appendix

Convergence monitoring

US Data

Separate model

```
sUS<- monitor(separate_modelUS)$Bulk_ESS
```

```
## Inference for the input samples (4 chains: each with iter = 2000; warmup = 0):
##
##           Q5      Q50      Q95      Mean   SD   Rhat Bulk_ESS
## theta[1]      0.0      0.0      0.0      0.0  0.0    1     7149
## theta[2]      0.0      0.0      0.0      0.0  0.0    1     6492
## theta[3]      0.0      0.0      0.0      0.0  0.0    1     7188
## theta[4]      0.0      0.0      0.0      0.0  0.0    1     6583
## theta[5]      0.0      0.0      0.0      0.0  0.0    1     5984
## theta[6]      0.0      0.0      0.0      0.0  0.0    1     6756
## theta[7]      0.0      0.0      0.0      0.0  0.0    1     6728
## theta[8]      0.0      0.0      0.0      0.0  0.0    1     5760
## theta[9]      0.0      0.0      0.0      0.0  0.0    1     6410
## theta[10]     0.0      0.0      0.0      0.0  0.0    1     6668
## theta[11]     0.0      0.0      0.0      0.0  0.0    1     7088
## theta[12]     0.0      0.0      0.0      0.0  0.0    1     6024
## theta[13]     0.0      0.0      0.0      0.0  0.0    1     6442
## theta[14]     0.0      0.0      0.0      0.0  0.0    1     6359
## theta[15]     0.0      0.0      0.0      0.0  0.0    1     5993
## theta[16]     0.0      0.0      0.0      0.0  0.0    1     6336
## theta[17]     0.0      0.0      0.0      0.0  0.0    1     6823
## theta[18]     0.0      0.0      0.0      0.0  0.0    1     5161
## theta[19]     0.0      0.0      0.0      0.0  0.0    1     5801
## theta[20]     0.0      0.0      0.0      0.0  0.0    1     6581
## theta[21]     0.0      0.0      0.0      0.0  0.0    1     6998
## theta[22]     0.0      0.0      0.0      0.0  0.0    1     6372
## theta[23]     0.0      0.0      0.0      0.0  0.0    1     6945
## theta[24]     0.0      0.0      0.0      0.0  0.0    1     6642
## theta[25]     0.0      0.0      0.0      0.0  0.0    1     5609
## theta[26]     0.0      0.0      0.0      0.0  0.0    1     6632
## theta[27]     0.0      0.0      0.0      0.0  0.0    1     7208
## theta[28]     0.0      0.0      0.0      0.0  0.0    1     6532
## theta[29]     0.0      0.0      0.0      0.0  0.0    1     6744
## theta[30]     0.0      0.0      0.0      0.0  0.0    1     5865
## theta[31]     0.0      0.0      0.0      0.0  0.0    1     7939
## theta[32]     0.0      0.0      0.0      0.0  0.0    1     7022
```

## theta[33]	0.0	0.0	0.0	0.0	0.0	1	6424
## theta[34]	0.0	0.0	0.0	0.0	0.0	1	5808
## theta[35]	0.0	0.0	0.0	0.0	0.0	1	6944
## theta[36]	0.0	0.0	0.0	0.0	0.0	1	6922
## theta[37]	0.0	0.0	0.0	0.0	0.0	1	6922
## theta[38]	0.0	0.0	0.0	0.0	0.0	1	5154
## theta[39]	0.0	0.0	0.0	0.0	0.0	1	7185
## theta[40]	0.0	0.0	0.0	0.0	0.0	1	5864
## theta[41]	0.0	0.0	0.0	0.0	0.0	1	6218
## theta[42]	0.0	0.0	0.0	0.0	0.0	1	6573
## theta[43]	0.0	0.0	0.0	0.0	0.0	1	7306
## theta[44]	0.0	0.0	0.0	0.0	0.0	1	7162
## theta[45]	0.0	0.0	0.0	0.0	0.0	1	7523
## theta[46]	0.0	0.0	0.0	0.0	0.0	1	6585
## theta[47]	0.0	0.0	0.0	0.0	0.0	1	7408
## theta[48]	0.0	0.0	0.0	0.0	0.0	1	6210
## theta[49]	0.0	0.0	0.0	0.0	0.0	1	6987
## theta[50]	0.0	0.0	0.0	0.0	0.0	1	5848
## theta[51]	0.0	0.0	0.0	0.0	0.0	1	6334
## log_lik[1]	-7.6	-5.1	-3.8	-5.3	1.2	1	7083
## log_lik[2]	-3.9	-2.8	-2.7	-3.0	0.5	1	2601
## log_lik[3]	-4.8	-3.9	-3.8	-4.1	0.4	1	1821
## log_lik[4]	-4.8	-3.7	-3.5	-3.8	0.5	1	2746
## log_lik[5]	-8.1	-5.8	-4.7	-6.0	1.1	1	6023
## log_lik[6]	-5.8	-4.1	-3.6	-4.3	0.7	1	5872
## log_lik[7]	-6.3	-4.1	-3.3	-4.4	1.0	1	6635
## log_lik[8]	-5.1	-3.4	-2.8	-3.6	0.8	1	5596
## log_lik[9]	-5.2	-3.3	-2.8	-3.6	0.8	1	5341
## log_lik[10]	-6.0	-4.6	-4.3	-4.8	0.6	1	4891
## log_lik[11]	-5.1	-4.2	-4.1	-4.4	0.4	1	1694
## log_lik[12]	-3.9	-3.1	-3.0	-3.3	0.3	1	1912
## log_lik[13]	-4.2	-3.2	-3.1	-3.4	0.4	1	2236
## log_lik[14]	-5.2	-4.3	-4.2	-4.4	0.4	1	2258
## log_lik[15]	-5.2	-4.0	-3.9	-4.2	0.5	1	2988
## log_lik[16]	-4.3	-3.4	-3.3	-3.6	0.4	1	1717
## log_lik[17]	-4.5	-3.6	-3.5	-3.7	0.4	1	1798
## log_lik[18]	-5.0	-3.8	-3.6	-4.0	0.5	1	3194
## log_lik[19]	-4.8	-3.8	-3.7	-4.0	0.4	1	2348
## log_lik[20]	-4.0	-3.0	-2.8	-3.1	0.4	1	2646
## log_lik[21]	-7.4	-5.0	-3.9	-5.2	1.1	1	7130
## log_lik[22]	-4.5	-3.7	-3.6	-3.9	0.3	1	1728
## log_lik[23]	-7.7	-5.2	-4.1	-5.5	1.1	1	6652
## log_lik[24]	-4.6	-3.8	-3.6	-3.9	0.4	1	1883
## log_lik[25]	-4.8	-3.7	-3.6	-3.9	0.5	1	3069
## log_lik[26]	-5.8	-4.1	-3.7	-4.4	0.7	1	5671
## log_lik[27]	-3.8	-2.8	-2.7	-3.0	0.4	1	2930
## log_lik[28]	-4.3	-3.3	-3.2	-3.5	0.4	1	2300
## log_lik[29]	-4.4	-3.5	-3.3	-3.6	0.4	1	2195
## log_lik[30]	-4.1	-2.8	-2.6	-3.0	0.5	1	4003
## log_lik[31]	-8.3	-5.6	-4.0	-5.8	1.4	1	7872
## log_lik[32]	-4.6	-3.4	-3.2	-3.6	0.5	1	3006
## log_lik[33]	-9.6	-6.4	-4.6	-6.7	1.5	1	6411
## log_lik[34]	-8.7	-5.9	-4.3	-6.1	1.4	1	5835
## log_lik[35]	-4.2	-2.9	-2.7	-3.1	0.5	1	3517

## log_lik[36]	-6.7	-4.8	-4.2	-5.0	0.8	1	6293
## log_lik[37]	-4.9	-3.8	-3.6	-3.9	0.4	1	3088
## log_lik[38]	-6.6	-4.3	-3.3	-4.5	1.0	1	5217
## log_lik[39]	-7.1	-4.9	-4.1	-5.2	1.0	1	7330
## log_lik[40]	-4.3	-3.0	-2.8	-3.2	0.5	1	4493
## log_lik[41]	-4.9	-3.9	-3.7	-4.0	0.4	1	2700
## log_lik[42]	-5.2	-3.3	-2.7	-3.5	0.8	1	6586
## log_lik[43]	-4.8	-4.0	-3.9	-4.1	0.4	1	1775
## log_lik[44]	-6.6	-4.9	-4.5	-5.2	0.7	1	5395
## log_lik[45]	-5.4	-3.9	-3.6	-4.1	0.6	1	4809
## log_lik[46]	-3.2	-2.4	-2.3	-2.6	0.3	1	1943
## log_lik[47]	-7.7	-5.2	-4.0	-5.4	1.2	1	7300
## log_lik[48]	-4.9	-3.9	-3.7	-4.0	0.4	1	2545
## log_lik[49]	-4.0	-3.2	-3.1	-3.3	0.3	1	2001
## log_lik[50]	-5.1	-3.9	-3.7	-4.1	0.5	1	3583
## log_lik[51]	-3.8	-2.6	-2.4	-2.8	0.5	1	4518
## ypred[1]	246.0	279.0	315.0	279.9	20.8	1	4617
## ypred[2]	20.0	31.0	43.0	31.0	6.9	1	4542
## ypred[3]	282.0	317.0	354.0	317.3	21.6	1	4910
## ypred[4]	143.0	168.0	196.0	168.4	16.2	1	4515
## ypred[5]	1341.0	1415.5	1490.0	1415.0	45.0	1	4437
## ypred[6]	191.0	220.0	249.0	219.8	17.8	1	4689
## ypred[7]	99.0	120.0	142.0	120.3	13.3	1	4903
## ypred[8]	37.0	50.0	65.0	50.6	8.5	1	4339
## ypred[9]	35.0	49.0	64.0	49.1	8.7	1	4648
## ypred[10]	860.0	920.0	982.0	920.2	37.2	1	3957
## ypred[11]	547.0	596.0	645.0	596.5	30.0	1	4477
## ypred[12]	50.0	66.0	83.0	65.9	9.9	1	4256
## ypred[13]	60.0	77.0	95.0	77.2	10.7	1	4854
## ypred[14]	610.0	660.0	713.0	660.8	31.5	1	4478
## ypred[15]	342.0	379.0	419.0	379.0	23.7	1	4152
## ypred[16]	96.0	116.0	139.0	116.8	13.3	1	4235
## ypred[17]	134.0	159.0	186.0	159.2	15.6	1	4408
## ypred[18]	180.0	209.0	239.0	209.1	18.1	1	4659
## ypred[19]	239.0	271.0	304.0	271.0	19.8	1	4486
## ypred[20]	36.0	50.0	64.0	49.7	8.6	1	4357
## ypred[21]	286.0	320.0	357.0	320.7	21.5	1	4720
## ypred[22]	197.0	225.0	256.0	225.1	18.0	1	4319
## ypred[23]	436.0	480.0	524.0	479.9	26.4	1	4691
## ypred[24]	200.0	230.0	262.0	230.4	18.6	1	4888
## ypred[25]	174.0	201.0	230.0	201.3	17.3	1	4551
## ypred[26]	262.0	294.0	329.0	294.6	20.5	1	5292
## ypred[27]	26.0	37.0	50.0	37.5	7.4	1	4676
## ypred[28]	77.0	96.0	116.0	95.9	11.8	1	5013
## ypred[29]	105.0	127.0	150.0	127.3	13.7	1	4621
## ypred[30]	22.0	32.0	45.0	32.5	6.9	1	4052
## ypred[31]	264.0	297.0	333.0	297.7	21.0	1	4909
## ypred[32]	71.0	89.0	109.0	89.3	11.7	1	4655
## ypred[33]	660.0	712.0	764.0	712.0	32.0	1	4629
## ypred[34]	507.0	552.0	601.0	552.3	28.5	1	4619
## ypred[35]	28.0	39.0	53.0	39.5	7.7	1	4802
## ypred[36]	609.0	660.0	710.0	660.3	31.2	1	4438
## ypred[37]	191.0	219.0	250.0	219.6	18.2	1	4495
## ypred[38]	106.0	128.0	151.0	128.2	13.8	1	4083

## ypred[39]	557.0	605.0	655.0	605.3	29.9	1	4442
## ypred[40]	32.0	44.0	58.0	44.5	8.1	1	4739
## ypred[41]	235.0	267.0	300.0	267.6	19.8	1	4313
## ypred[42]	30.0	42.0	56.0	42.0	7.9	1	4592
## ypred[43]	325.0	362.0	401.0	362.8	23.0	1	4474
## ypred[44]	1335.0	1410.0	1487.0	1410.5	46.0	1	5001
## ypred[45]	158.0	183.0	212.0	183.8	16.5	1	4835
## ypred[46]	9.0	17.0	26.0	17.1	5.0	1	4499
## ypred[47]	364.0	403.0	445.0	403.0	24.6	1	4812
## ypred[48]	233.0	265.0	298.0	265.1	19.9	1	4921
## ypred[49]	63.0	79.0	98.0	79.6	10.8	1	5057
## ypred[50]	224.0	255.0	289.0	255.6	19.8	1	4104
## ypred[51]	13.0	22.0	31.0	21.8	5.6	1	4806
## lp_--	-199370.7	-199361.3	-199354.0	-199361.7	5.0	1	1589
##	Tail_ESS						
## theta[1]	2951						
## theta[2]	2838						
## theta[3]	2755						
## theta[4]	3085						
## theta[5]	3017						
## theta[6]	2522						
## theta[7]	3133						
## theta[8]	3016						
## theta[9]	2921						
## theta[10]	3333						
## theta[11]	2852						
## theta[12]	3210						
## theta[13]	3012						
## theta[14]	2975						
## theta[15]	2748						
## theta[16]	2479						
## theta[17]	2311						
## theta[18]	2779						
## theta[19]	2823						
## theta[20]	2854						
## theta[21]	3204						
## theta[22]	3213						
## theta[23]	2967						
## theta[24]	3029						
## theta[25]	2859						
## theta[26]	2659						
## theta[27]	2970						
## theta[28]	2766						
## theta[29]	2852						
## theta[30]	2996						
## theta[31]	2857						
## theta[32]	2637						
## theta[33]	2824						
## theta[34]	2962						
## theta[35]	2875						
## theta[36]	2982						
## theta[37]	3035						
## theta[38]	3243						
## theta[39]	2916						

## theta[40]	2762
## theta[41]	2828
## theta[42]	2696
## theta[43]	2909
## theta[44]	2991
## theta[45]	2923
## theta[46]	2601
## theta[47]	3251
## theta[48]	3139
## theta[49]	2725
## theta[50]	2922
## theta[51]	2774
## log_lik[1]	2951
## log_lik[2]	2666
## log_lik[3]	2034
## log_lik[4]	2836
## log_lik[5]	3068
## log_lik[6]	2522
## log_lik[7]	3271
## log_lik[8]	3016
## log_lik[9]	3410
## log_lik[10]	3651
## log_lik[11]	2048
## log_lik[12]	2133
## log_lik[13]	2657
## log_lik[14]	2411
## log_lik[15]	3050
## log_lik[16]	1725
## log_lik[17]	1671
## log_lik[18]	2992
## log_lik[19]	2540
## log_lik[20]	2668
## log_lik[21]	3210
## log_lik[22]	2753
## log_lik[23]	2967
## log_lik[24]	1975
## log_lik[25]	2678
## log_lik[26]	3129
## log_lik[27]	2426
## log_lik[28]	2615
## log_lik[29]	2414
## log_lik[30]	3197
## log_lik[31]	2857
## log_lik[32]	2707
## log_lik[33]	2824
## log_lik[34]	2962
## log_lik[35]	3211
## log_lik[36]	3039
## log_lik[37]	3001
## log_lik[38]	3251
## log_lik[39]	3453
## log_lik[40]	3354
## log_lik[41]	2562
## log_lik[42]	2988

## log_lik[43]	2217
## log_lik[44]	2991
## log_lik[45]	2923
## log_lik[46]	2380
## log_lik[47]	3251
## log_lik[48]	2999
## log_lik[49]	1842
## log_lik[50]	2874
## log_lik[51]	3100
## ypred[1]	3941
## ypred[2]	3915
## ypred[3]	3941
## ypred[4]	3923
## ypred[5]	3871
## ypred[6]	3770
## ypred[7]	3565
## ypred[8]	3779
## ypred[9]	4146
## ypred[10]	3819
## ypred[11]	3671
## ypred[12]	3600
## ypred[13]	3972
## ypred[14]	4016
## ypred[15]	3899
## ypred[16]	3956
## ypred[17]	3032
## ypred[18]	3715
## ypred[19]	3864
## ypred[20]	3703
## ypred[21]	3984
## ypred[22]	3931
## ypred[23]	3868
## ypred[24]	3969
## ypred[25]	4078
## ypred[26]	3899
## ypred[27]	3974
## ypred[28]	3866
## ypred[29]	3839
## ypred[30]	3706
## ypred[31]	4221
## ypred[32]	3856
## ypred[33]	4092
## ypred[34]	3944
## ypred[35]	3748
## ypred[36]	3738
## ypred[37]	3846
## ypred[38]	3633
## ypred[39]	4062
## ypred[40]	3785
## ypred[41]	3966
## ypred[42]	3417
## ypred[43]	3926
## ypred[44]	4050
## ypred[45]	3845

```
## ypred[46]      3562
## ypred[47]      3777
## ypred[48]      4071
## ypred[49]      3959
## ypred[50]      3495
## ypred[51]      3727
## lp__           2097
##
## For each parameter, Bulk_ESS and Tail_ESS are crude measures of
## effective sample size for bulk and tail quantities respectively (an ESS > 100
## per chain is considered good), and Rhat is the potential scale reduction
## factor on rank normalized split chains (at convergence, Rhat <= 1.05).
```

Hierarchical model

```
sUS<- monitor(hierarchical_modelUS)$Bulk_ESS
```

```
## Inference for the input samples (4 chains: each with iter = 2000; warmup = 0):
##
##           Q5      Q50      Q95      Mean      SD      Rhat Bulk_ESS
## mu           0.0      0.0      0.0      0.0      0.0      1      7396
## eta        6811.1  10357.3  15427.2  10635.3  2650.7      1      2768
## theta[1]      0.0      0.0      0.0      0.0      0.0      1      5594
## theta[2]      0.0      0.0      0.0      0.0      0.0      1      5917
## theta[3]      0.0      0.0      0.0      0.0      0.0      1      6486
## theta[4]      0.0      0.0      0.0      0.0      0.0      1      5672
## theta[5]      0.0      0.0      0.0      0.0      0.0      1      6006
## theta[6]      0.0      0.0      0.0      0.0      0.0      1      5547
## theta[7]      0.0      0.0      0.0      0.0      0.0      1      5063
## theta[8]      0.0      0.0      0.0      0.0      0.0      1      6676
## theta[9]      0.0      0.0      0.0      0.0      0.0      1      5765
## theta[10]     0.0      0.0      0.0      0.0      0.0      1      5398
## theta[11]     0.0      0.0      0.0      0.0      0.0      1      7061
## theta[12]     0.0      0.0      0.0      0.0      0.0      1      5546
## theta[13]     0.0      0.0      0.0      0.0      0.0      1      5313
## theta[14]     0.0      0.0      0.0      0.0      0.0      1      5234
## theta[15]     0.0      0.0      0.0      0.0      0.0      1      4905
## theta[16]     0.0      0.0      0.0      0.0      0.0      1      5177
## theta[17]     0.0      0.0      0.0      0.0      0.0      1      5311
## theta[18]     0.0      0.0      0.0      0.0      0.0      1      5252
## theta[19]     0.0      0.0      0.0      0.0      0.0      1      5580
## theta[20]     0.0      0.0      0.0      0.0      0.0      1      5319
## theta[21]     0.0      0.0      0.0      0.0      0.0      1      6688
## theta[22]     0.0      0.0      0.0      0.0      0.0      1      5460
## theta[23]     0.0      0.0      0.0      0.0      0.0      1      7293
## theta[24]     0.0      0.0      0.0      0.0      0.0      1      5837
## theta[25]     0.0      0.0      0.0      0.0      0.0      1      5047
## theta[26]     0.0      0.0      0.0      0.0      0.0      1      5020
## theta[27]     0.0      0.0      0.0      0.0      0.0      1      4602
## theta[28]     0.0      0.0      0.0      0.0      0.0      1      6364
## theta[29]     0.0      0.0      0.0      0.0      0.0      1      5792
## theta[30]     0.0      0.0      0.0      0.0      0.0      1      4550
```

## theta[31]	0.0	0.0	0.0	0.0	0.0	1	5375
## theta[32]	0.0	0.0	0.0	0.0	0.0	1	5449
## theta[33]	0.0	0.0	0.0	0.0	0.0	1	6056
## theta[34]	0.0	0.0	0.0	0.0	0.0	1	7405
## theta[35]	0.0	0.0	0.0	0.0	0.0	1	6171
## theta[36]	0.0	0.0	0.0	0.0	0.0	1	5622
## theta[37]	0.0	0.0	0.0	0.0	0.0	1	6129
## theta[38]	0.0	0.0	0.0	0.0	0.0	1	5001
## theta[39]	0.0	0.0	0.0	0.0	0.0	1	5531
## theta[40]	0.0	0.0	0.0	0.0	0.0	1	6514
## theta[41]	0.0	0.0	0.0	0.0	0.0	1	6037
## theta[42]	0.0	0.0	0.0	0.0	0.0	1	4420
## theta[43]	0.0	0.0	0.0	0.0	0.0	1	5902
## theta[44]	0.0	0.0	0.0	0.0	0.0	1	4845
## theta[45]	0.0	0.0	0.0	0.0	0.0	1	6515
## theta[46]	0.0	0.0	0.0	0.0	0.0	1	5331
## theta[47]	0.0	0.0	0.0	0.0	0.0	1	6597
## theta[48]	0.0	0.0	0.0	0.0	0.0	1	6028
## theta[49]	0.0	0.0	0.0	0.0	0.0	1	5608
## theta[50]	0.0	0.0	0.0	0.0	0.0	1	5471
## theta[51]	0.0	0.0	0.0	0.0	0.0	1	6244
## alpha	27.3	41.7	61.9	42.7	10.6	1	2770
## beta	6784.0	10315.7	15365.0	10592.6	2640.1	1	2769
## log_lik[1]	-5.4	-3.9	-3.7	-4.1	0.6	1	2260
## log_lik[2]	-4.3	-2.9	-2.7	-3.1	0.6	1	4172
## log_lik[3]	-5.5	-4.0	-3.8	-4.2	0.6	1	1837
## log_lik[4]	-5.4	-3.7	-3.5	-4.0	0.7	1	2494
## log_lik[5]	-6.6	-4.8	-4.5	-5.1	0.8	1	2177
## log_lik[6]	-5.6	-3.8	-3.6	-4.1	0.7	1	2884
## log_lik[7]	-5.6	-3.6	-3.3	-3.9	0.8	1	3537
## log_lik[8]	-3.8	-2.9	-2.8	-3.1	0.4	1	1834
## log_lik[9]	-4.0	-3.0	-2.8	-3.1	0.5	1	3489
## log_lik[10]	-6.2	-4.5	-4.3	-4.8	0.7	1	1703
## log_lik[11]	-6.2	-4.4	-4.1	-4.6	0.8	1	1795
## log_lik[12]	-4.2	-3.2	-3.0	-3.3	0.4	1	1973
## log_lik[13]	-4.5	-3.3	-3.1	-3.5	0.5	1	2278
## log_lik[14]	-6.1	-4.4	-4.2	-4.7	0.7	1	2252
## log_lik[15]	-5.8	-4.1	-3.9	-4.3	0.7	1	2097
## log_lik[16]	-5.5	-3.6	-3.3	-3.9	0.8	1	3955
## log_lik[17]	-5.1	-3.7	-3.5	-3.9	0.6	1	2390
## log_lik[18]	-5.2	-3.8	-3.6	-4.0	0.6	1	2265
## log_lik[19]	-5.5	-3.9	-3.7	-4.2	0.6	1	2132
## log_lik[20]	-4.0	-3.0	-2.8	-3.1	0.4	1	2010
## log_lik[21]	-5.5	-4.0	-3.8	-4.2	0.6	1	1478
## log_lik[22]	-5.7	-3.9	-3.6	-4.2	0.7	1	2534
## log_lik[23]	-5.8	-4.2	-4.0	-4.4	0.7	1	1855
## log_lik[24]	-5.5	-3.9	-3.6	-4.1	0.7	1	2660
## log_lik[25]	-6.2	-4.0	-3.6	-4.3	0.9	1	3501
## log_lik[26]	-5.4	-3.9	-3.7	-4.2	0.6	1	2210
## log_lik[27]	-4.7	-3.1	-2.7	-3.3	0.7	1	5183
## log_lik[28]	-4.6	-3.4	-3.2	-3.6	0.5	1	2236
## log_lik[29]	-5.0	-3.5	-3.3	-3.8	0.6	1	2449
## log_lik[30]	-6.1	-3.7	-2.7	-4.0	1.1	1	4651
## log_lik[31]	-6.5	-4.1	-3.7	-4.5	1.0	1	3846

## log_lik[32]	-4.5	-3.3	-3.2	-3.5	0.5	1	1999
## log_lik[33]	-6.3	-4.4	-4.2	-4.7	0.8	1	2639
## log_lik[34]	-5.9	-4.2	-4.0	-4.5	0.7	1	1667
## log_lik[35]	-4.0	-2.9	-2.7	-3.1	0.5	1	4020
## log_lik[36]	-6.0	-4.4	-4.1	-4.6	0.7	1	2255
## log_lik[37]	-5.2	-3.8	-3.6	-4.0	0.6	1	2098
## log_lik[38]	-6.5	-3.9	-3.3	-4.2	1.1	1	4687
## log_lik[39]	-5.8	-4.3	-4.1	-4.5	0.6	1	2036
## log_lik[40]	-3.8	-2.9	-2.8	-3.0	0.4	1	2367
## log_lik[41]	-5.7	-4.0	-3.7	-4.2	0.7	1	2675
## log_lik[42]	-5.0	-3.1	-2.7	-3.4	0.8	1	4975
## log_lik[43]	-5.7	-4.1	-3.9	-4.3	0.7	1	2291
## log_lik[44]	-6.5	-4.8	-4.5	-5.0	0.7	1	1833
## log_lik[45]	-5.1	-3.8	-3.6	-4.0	0.6	1	2169
## log_lik[46]	-3.7	-2.6	-2.3	-2.7	0.5	1	4983
## log_lik[47]	-5.6	-4.1	-3.9	-4.3	0.6	1	2097
## log_lik[48]	-5.9	-4.0	-3.7	-4.3	0.8	1	2635
## log_lik[49]	-4.4	-3.3	-3.1	-3.5	0.5	1	2284
## log_lik[50]	-5.3	-3.9	-3.7	-4.1	0.6	1	1991
## log_lik[51]	-4.3	-2.8	-2.4	-3.0	0.6	1	6377
## ypred[1]	215.0	249.0	285.0	248.8	21.5	1	4631
## ypred[2]	26.0	37.0	50.0	37.0	7.4	1	4505
## ypred[3]	282.0	322.0	363.0	322.5	24.8	1	4636
## ypred[4]	141.0	168.0	198.0	168.5	17.3	1	4929
## ypred[5]	1281.0	1363.0	1451.0	1364.6	51.7	1	4645
## ypred[6]	181.0	212.0	244.0	212.1	19.3	1	4557
## ypred[7]	91.0	113.0	139.0	113.8	14.3	1	4197
## ypred[8]	30.0	43.0	57.0	42.9	8.1	1	4557
## ypred[9]	28.0	39.0	52.0	39.3	7.5	1	4735
## ypred[10]	827.0	897.0	968.0	896.9	42.8	1	4653
## ypred[11]	535.0	589.0	645.0	589.3	33.7	1	4612
## ypred[12]	51.0	67.0	85.0	67.0	10.4	1	4476
## ypred[13]	63.0	81.0	101.0	81.1	11.7	1	4022
## ypred[14]	602.0	658.0	719.0	658.6	35.9	1	4570
## ypred[15]	323.0	365.0	408.0	365.9	26.0	1	4275
## ypred[16]	101.0	124.0	150.0	124.7	14.8	1	4468
## ypred[17]	131.0	157.0	185.0	157.5	16.6	1	4620
## ypred[18]	173.0	203.0	236.0	204.2	19.2	1	4837
## ypred[19]	226.0	261.0	299.0	261.5	22.0	1	4805
## ypred[20]	35.0	48.0	63.0	48.0	8.6	1	4404
## ypred[21]	253.0	291.0	331.0	291.2	23.5	1	5032
## ypred[22]	197.0	231.0	267.0	231.0	21.1	1	4971
## ypred[23]	397.0	443.0	493.0	443.5	29.2	1	4937
## ypred[24]	203.0	236.0	272.0	236.3	21.2	1	4794
## ypred[25]	156.0	185.0	215.0	184.9	18.3	1	4383
## ypred[26]	243.0	281.0	318.0	280.7	22.6	1	4520
## ypred[27]	28.0	40.0	53.0	40.2	7.7	1	4526
## ypred[28]	79.0	99.0	121.0	99.2	12.8	1	4826
## ypred[29]	107.0	132.0	158.0	131.9	15.3	1	4502
## ypred[30]	26.0	37.0	51.0	37.7	7.7	1	4061
## ypred[31]	241.0	277.0	318.0	278.2	23.4	1	4766
## ypred[32]	74.0	93.0	114.0	93.4	12.2	1	4696
## ypred[33]	608.0	666.0	725.0	666.6	36.0	1	5181
## ypred[34]	454.0	506.0	557.0	505.7	30.9	1	4737

## ypred[35]	27.0	39.0	52.0	39.2	7.6	1	4294
## ypred[36]	571.0	626.0	684.0	626.2	34.3	1	4742
## ypred[37]	180.0	211.0	244.0	211.0	19.4	1	4988
## ypred[38]	99.0	123.0	149.0	123.6	15.2	1	4458
## ypred[39]	518.0	571.0	627.0	571.1	33.2	1	4576
## ypred[40]	29.0	41.0	54.0	41.0	7.9	1	4766
## ypred[41]	229.0	266.0	302.0	265.6	21.9	1	4685
## ypred[42]	28.0	41.0	55.0	41.1	8.1	1	4027
## ypred[43]	317.0	357.0	401.0	357.0	25.9	1	4763
## ypred[44]	1295.0	1379.0	1468.0	1380.2	52.0	1	4804
## ypred[45]	163.0	193.0	225.0	193.1	18.8	1	4303
## ypred[46]	12.0	20.0	29.0	20.2	5.2	1	4315
## ypred[47]	332.0	373.0	419.0	374.2	26.0	1	4639
## ypred[48]	241.0	278.0	316.0	278.5	23.0	1	5136
## ypred[49]	59.0	76.0	96.0	76.8	11.1	1	4608
## ypred[50]	214.0	248.0	283.0	248.1	21.2	1	4599
## ypred[51]	15.0	23.0	33.0	23.4	5.7	1	4663
## lp_--	-199242.3	-199232.2	-199224.1	-199232.6	5.5	1	1469
##	Tail_ESS						
## mu	3023						
## eta	2795						
## theta[1]	2923						
## theta[2]	3189						
## theta[3]	2888						
## theta[4]	2797						
## theta[5]	3039						
## theta[6]	3061						
## theta[7]	2962						
## theta[8]	2943						
## theta[9]	3002						
## theta[10]	2717						
## theta[11]	2586						
## theta[12]	2604						
## theta[13]	3099						
## theta[14]	3123						
## theta[15]	2872						
## theta[16]	3082						
## theta[17]	3058						
## theta[18]	3074						
## theta[19]	2911						
## theta[20]	3009						
## theta[21]	3001						
## theta[22]	2565						
## theta[23]	2776						
## theta[24]	3242						
## theta[25]	2805						
## theta[26]	2705						
## theta[27]	2475						
## theta[28]	3042						
## theta[29]	2735						
## theta[30]	2915						
## theta[31]	3233						
## theta[32]	2740						
## theta[33]	3014						

## theta[34]	3007
## theta[35]	2624
## theta[36]	3035
## theta[37]	2973
## theta[38]	3162
## theta[39]	2674
## theta[40]	3155
## theta[41]	2824
## theta[42]	2730
## theta[43]	2943
## theta[44]	2918
## theta[45]	3111
## theta[46]	3041
## theta[47]	3135
## theta[48]	3250
## theta[49]	3240
## theta[50]	3082
## theta[51]	2825
## alpha	2850
## beta	2795
## log_lik[1]	2673
## log_lik[2]	3244
## log_lik[3]	2369
## log_lik[4]	2596
## log_lik[5]	2620
## log_lik[6]	3192
## log_lik[7]	2962
## log_lik[8]	1996
## log_lik[9]	2981
## log_lik[10]	1901
## log_lik[11]	1982
## log_lik[12]	2526
## log_lik[13]	2505
## log_lik[14]	2631
## log_lik[15]	2499
## log_lik[16]	3284
## log_lik[17]	2621
## log_lik[18]	2506
## log_lik[19]	2743
## log_lik[20]	2115
## log_lik[21]	1921
## log_lik[22]	2876
## log_lik[23]	1722
## log_lik[24]	2850
## log_lik[25]	2724
## log_lik[26]	2416
## log_lik[27]	3152
## log_lik[28]	2329
## log_lik[29]	2562
## log_lik[30]	3339
## log_lik[31]	3165
## log_lik[32]	2514
## log_lik[33]	2878
## log_lik[34]	1763

## log_lik[35]	3438
## log_lik[36]	2911
## log_lik[37]	2284
## log_lik[38]	3573
## log_lik[39]	1779
## log_lik[40]	2439
## log_lik[41]	2627
## log_lik[42]	3637
## log_lik[43]	2443
## log_lik[44]	2332
## log_lik[45]	2355
## log_lik[46]	3041
## log_lik[47]	2055
## log_lik[48]	3112
## log_lik[49]	2898
## log_lik[50]	2397
## log_lik[51]	3160
## ypred[1]	3783
## ypred[2]	3790
## ypred[3]	3505
## ypred[4]	3870
## ypred[5]	3167
## ypred[6]	3641
## ypred[7]	3552
## ypred[8]	3892
## ypred[9]	3843
## ypred[10]	3273
## ypred[11]	3551
## ypred[12]	4055
## ypred[13]	3451
## ypred[14]	3706
## ypred[15]	3071
## ypred[16]	3777
## ypred[17]	3792
## ypred[18]	3552
## ypred[19]	3645
## ypred[20]	3512
## ypred[21]	3448
## ypred[22]	3671
## ypred[23]	3676
## ypred[24]	4064
## ypred[25]	3301
## ypred[26]	3884
## ypred[27]	4121
## ypred[28]	3666
## ypred[29]	3762
## ypred[30]	3671
## ypred[31]	3656
## ypred[32]	3667
## ypred[33]	3359
## ypred[34]	3760
## ypred[35]	3758
## ypred[36]	3516
## ypred[37]	3698


```
## ypred[38]      3890
## ypred[39]      3682
## ypred[40]      3889
## ypred[41]      3709
## ypred[42]      3834
## ypred[43]      3790
## ypred[44]      3513
## ypred[45]      3877
## ypred[46]      3588
## ypred[47]      3374
## ypred[48]      3818
## ypred[49]      3831
## ypred[50]      3472
## ypred[51]      4095
## lp__           2100
##
## For each parameter, Bulk_ESS and Tail_ESS are crude measures of
## effective sample size for bulk and tail quantities respectively (an ESS > 100
## per chain is considered good), and Rhat is the potential scale reduction
## factor on rank normalized split chains (at convergence, Rhat <= 1.05).
```

Colombia Data

Separate model

```
sUS<- monitor(separate_modelCol)$Bulk_ESS
```

```
## Inference for the input samples (4 chains: each with iter = 2000; warmup = 0):
##
##           Q5      Q50      Q95      Mean      SD      Rhat Bulk_ESS Tail_ESS
## theta[1]    0.0    0.0    0.0    0.0  0.0    1      8690    2599
## theta[2]    0.0    0.0    0.0    0.0  0.0    1      8780    2971
## theta[3]    0.0    0.0    0.0    0.0  0.0    1      8299    3168
## theta[4]    0.0    0.0    0.0    0.0  0.0    1      9520    2681
## theta[5]    0.0    0.0    0.0    0.0  0.0    1      8388    3466
## theta[6]    0.0    0.0    0.0    0.0  0.0    1     10700    2842
## theta[7]    0.0    0.0    0.0    0.0  0.0    1      9434    2489
## theta[8]    0.0    0.0    0.0    0.0  0.0    1      9531    2475
## theta[9]    0.0    0.0    0.0    0.0  0.0    1      7660    2946
## theta[10]   0.0    0.0    0.0    0.0  0.0    1      8582    3033
## theta[11]   0.0    0.0    0.0    0.0  0.0    1      9065    2749
## theta[12]   0.0    0.0    0.0    0.0  0.0    1      8937    3133
## theta[13]   0.0    0.0    0.0    0.0  0.0    1      7721    2783
## theta[14]   0.0    0.0    0.0    0.0  0.0    1      7402    3238
## theta[15]   0.0    0.0    0.0    0.0  0.0    1      8973    2833
## theta[16]   0.0    0.0    0.0    0.0  0.0    1      7727    2860
## theta[17]   0.0    0.0    0.0    0.0  0.0    1      8363    2601
## theta[18]   0.0    0.0    0.0    0.0  0.0    1      7806    2798
## theta[19]   0.0    0.0    0.0    0.0  0.0    1      7714    2678
## theta[20]   0.0    0.0    0.0    0.0  0.0    1      8363    2955
## theta[21]   0.0    0.0    0.0    0.0  0.0    1      8345    3505
## theta[22]   0.0    0.0    0.0    0.0  0.0    1      9161    2837
```

## theta[23]	0.0	0.0	0.0	0.0	0.0	1	8808	2983
## theta[24]	0.0	0.0	0.0	0.0	0.0	1	8080	2800
## theta[25]	0.0	0.0	0.0	0.0	0.0	1	9208	2825
## theta[26]	0.0	0.0	0.0	0.0	0.0	1	9539	2931
## theta[27]	0.0	0.0	0.0	0.0	0.0	1	8166	2699
## theta[28]	0.0	0.0	0.0	0.0	0.0	1	9051	2950
## theta[29]	0.0	0.0	0.0	0.0	0.0	1	8388	2971
## theta[30]	0.0	0.0	0.0	0.0	0.0	1	9231	2829
## theta[31]	0.0	0.0	0.0	0.0	0.0	1	8577	3178
## theta[32]	0.0	0.0	0.0	0.0	0.0	1	7655	2766
## theta[33]	0.0	0.0	0.0	0.0	0.0	1	7799	2648
## theta[34]	0.0	0.0	0.0	0.0	0.0	1	8068	3195
## theta[35]	0.0	0.0	0.0	0.0	0.0	1	8977	2888
## theta[36]	0.0	0.0	0.0	0.0	0.0	1	9355	3208
## theta[37]	0.0	0.0	0.0	0.0	0.0	1	7103	2823
## log_lik[1]	-4.4	-2.8	-2.4	-3.0	0.7	1	7248	3194
## log_lik[2]	-5.6	-4.5	-4.4	-4.7	0.5	1	3554	2936
## log_lik[3]	-5.9	-3.8	-3.1	-4.1	0.9	1	7899	3168
## log_lik[4]	-6.6	-4.6	-3.8	-4.8	0.9	1	7980	2824
## log_lik[5]	-8.4	-5.6	-4.2	-5.9	1.3	1	8469	3466
## log_lik[6]	-6.1	-4.7	-4.4	-4.9	0.6	1	5253	2948
## log_lik[7]	-5.0	-3.8	-3.7	-4.0	0.5	1	2704	2381
## log_lik[8]	-4.4	-3.6	-3.5	-3.7	0.3	1	1431	1936
## log_lik[9]	-4.2	-3.4	-3.3	-3.5	0.3	1	1628	2219
## log_lik[10]	-4.2	-3.4	-3.3	-3.5	0.3	1	1838	2107
## log_lik[11]	-5.4	-3.6	-3.0	-3.8	0.8	1	7232	2749
## log_lik[12]	-7.8	-5.2	-3.9	-5.4	1.2	1	8586	3182
## log_lik[13]	-8.3	-5.2	-3.5	-5.4	1.5	1	7730	2783
## log_lik[14]	-4.7	-3.9	-3.8	-4.0	0.3	1	2134	2462
## log_lik[15]	-4.9	-4.0	-3.8	-4.1	0.4	1	2017	2218
## log_lik[16]	-5.6	-3.9	-3.6	-4.2	0.7	1	5023	2860
## log_lik[17]	-5.1	-4.2	-4.0	-4.3	0.4	1	2161	2211
## log_lik[18]	-5.2	-4.2	-4.0	-4.3	0.4	1	2848	2717
## log_lik[19]	-3.9	-2.5	-2.4	-2.8	0.5	1	2986	2613
## log_lik[20]	-5.1	-4.1	-3.9	-4.2	0.4	1	2881	2737
## log_lik[21]	-5.3	-3.1	-2.1	-3.3	1.0	1	7917	3505
## log_lik[22]	-6.7	-4.5	-3.6	-4.7	1.0	1	8694	2837
## log_lik[23]	-5.4	-3.9	-3.6	-4.1	0.6	1	4897	2959
## log_lik[24]	-4.5	-3.7	-3.6	-3.8	0.4	1	1852	1882
## log_lik[25]	-4.7	-3.8	-3.7	-4.0	0.4	1	1775	2070
## log_lik[26]	-4.7	-3.9	-3.7	-4.0	0.4	1	1704	1723
## log_lik[27]	-4.2	-3.1	-3.0	-3.3	0.4	1	2433	2349
## log_lik[28]	-4.2	-3.3	-3.1	-3.4	0.4	1	2357	2512
## log_lik[29]	-5.5	-3.8	-3.4	-4.0	0.7	1	6728	2971
## log_lik[30]	-3.2	-2.4	-2.3	-2.6	0.3	1	1728	2006
## log_lik[31]	-7.0	-4.8	-3.8	-5.0	1.0	1	8258	3178
## log_lik[32]	-5.1	-3.6	-3.3	-3.8	0.6	1	5215	2797
## log_lik[33]	-5.4	-3.9	-3.6	-4.1	0.6	1	5904	3104
## log_lik[34]	-5.4	-3.8	-3.6	-4.1	0.7	1	5398	3195
## log_lik[35]	-5.6	-4.3	-4.1	-4.5	0.5	1	3753	3086
## log_lik[36]	-4.0	-2.6	-2.4	-2.8	0.6	1	3470	3224
## log_lik[37]	-4.0	-2.9	-2.8	-3.1	0.4	1	1961	2062
## ypred[1]	14.0	22.0	32.0	22.4	5.7	1	4684	3983
## ypred[2]	967.0	1030.0	1094.0	1029.9	38.5	1	5109	4110

```
## ypred[3]      48.0      63.0      79.0      62.9  9.5      1      4991      3905
## ypred[4]     263.0     296.0     330.0     296.2 20.7      1      4720      3724
## ypred[5]     333.0     369.0     406.0     369.3 22.6      1      4846      3949
## ypred[6]    1021.0    1086.0    1152.0    1086.4 40.2      1      5087      4029
## ypred[7]     230.0     261.0     295.0     261.4 19.5      1      5410      4116
## ypred[8]     140.0     164.0     190.0     164.3 15.5      1      4683      4077
## ypred[9]      92.0     113.0     134.0     113.0 12.8      1      4943      3809
## ypred[10]     89.0     109.0     131.0     109.3 12.8      1      4787      4020
## ypred[11]     62.0      79.0      98.0      79.5 10.8      1      4775      4031
## ypred[12]    255.0     287.0     321.0     287.7 20.3      1      4977      4168
## ypred[13]     71.0      90.0     110.0      90.1 11.7      1      4955      3978
## ypred[14]    268.0     301.0     335.0     301.1 20.4      1      4299      3864
## ypred[15]    309.0     347.0     385.0     346.7 23.1      1      4611      4025
## ypred[16]    176.0     203.0     233.0     203.7 17.0      1      4829      3640
## ypred[17]    459.0     503.0     549.0     503.4 27.2      1      5249      3430
## ypred[18]    442.0     484.0     529.0     484.2 26.5      1      4616      3242
## ypred[19]      9.0      16.0      25.0      16.1  4.9      1      5209      3689
## ypred[20]    380.0     419.0     463.0     420.1 25.3      1      4491      3691
## ypred[21]      7.0      14.0      23.0      14.4  4.7      1      4795      3866
## ypred[22]    173.0     200.0     229.0     200.7 17.2      1      4914      3879
## ypred[23]    199.0     228.0     259.0     228.3 18.5      1      5675      3720
## ypred[24]    174.0     202.0     231.0     201.9 17.1      1      5241      3877
## ypred[25]    243.0     275.0     309.0     275.6 19.8      1      4735      3866
## ypred[26]    256.0     287.0     323.0     288.1 20.8      1      5248      4093
## ypred[27]     49.0      64.0      82.0      64.8  9.9      1      5234      3768
## ypred[28]     63.0      80.0      98.0      80.6 10.9      1      4760      4048
## ypred[29]    105.0     127.0     150.0     127.2 13.7      1      4824      3864
## ypred[30]     10.0      17.0      26.0      17.3  5.0      1      4914      3572
## ypred[31]    242.0     273.0     308.0     273.5 20.0      1      4440      3853
## ypred[32]    110.0     133.0     157.0     133.3 14.4      1      4591      3800
## ypred[33]    186.0     213.0     242.0     213.1 17.2      1      5173      4213
## ypred[34]    181.0     209.0     238.0     209.3 17.3      1      5081      3732
## ypred[35]    532.0     578.0     629.0     579.1 29.4      1      5113      3628
## ypred[36]      9.0      17.0      26.0      17.0  5.1      1      4833      3506
## ypred[37]     30.0      43.0      57.0      42.9  8.0      1      5206      3866
## lp__         -97359.8 -97351.8 -97345.8 -97352.2 4.3      1      1520      2263
##
## For each parameter, Bulk_ESS and Tail_ESS are crude measures of
## effective sample size for bulk and tail quantities respectively (an ESS > 100
## per chain is considered good), and Rhat is the potential scale reduction
## factor on rank normalized split chains (at convergence, Rhat <= 1.05).
```

Hierarchical model

```
sUS<- monitor(hierarchical_modelCol)$Bulk_ESS
```

```
## Inference for the input samples (4 chains: each with iter = 2000; warmup = 0):
##
##           Q5      Q50      Q95      Mean      SD      Rhat Bulk_ESS Tail_ESS
## mu          0.0       0.0       0.0       0.0     0.0        1      6991      3157
## eta        593.5     943.4    1497.7     979.0  281.3        1      3377      3313
## theta[1]     0.0       0.0       0.0       0.0     0.0        1      6301      3132
```

## theta[2]	0.0	0.0	0.0	0.0	0.0	1	6684	2862
## theta[3]	0.0	0.0	0.0	0.0	0.0	1	6279	3027
## theta[4]	0.0	0.0	0.0	0.0	0.0	1	7548	3142
## theta[5]	0.0	0.0	0.0	0.0	0.0	1	5908	2880
## theta[6]	0.0	0.0	0.0	0.0	0.0	1	6869	3285
## theta[7]	0.0	0.0	0.0	0.0	0.0	1	6106	2897
## theta[8]	0.0	0.0	0.0	0.0	0.0	1	6687	3149
## theta[9]	0.0	0.0	0.0	0.0	0.0	1	5875	3212
## theta[10]	0.0	0.0	0.0	0.0	0.0	1	6571	2933
## theta[11]	0.0	0.0	0.0	0.0	0.0	1	5589	3255
## theta[12]	0.0	0.0	0.0	0.0	0.0	1	5247	2641
## theta[13]	0.0	0.0	0.0	0.0	0.0	1	5922	2861
## theta[14]	0.0	0.0	0.0	0.0	0.0	1	5646	2845
## theta[15]	0.0	0.0	0.0	0.0	0.0	1	6196	2736
## theta[16]	0.0	0.0	0.0	0.0	0.0	1	4677	3071
## theta[17]	0.0	0.0	0.0	0.0	0.0	1	6131	2796
## theta[18]	0.0	0.0	0.0	0.0	0.0	1	6678	3032
## theta[19]	0.0	0.0	0.0	0.0	0.0	1	5904	2815
## theta[20]	0.0	0.0	0.0	0.0	0.0	1	6451	3116
## theta[21]	0.0	0.0	0.0	0.0	0.0	1	5200	2926
## theta[22]	0.0	0.0	0.0	0.0	0.0	1	6139	3231
## theta[23]	0.0	0.0	0.0	0.0	0.0	1	5719	2860
## theta[24]	0.0	0.0	0.0	0.0	0.0	1	6802	3279
## theta[25]	0.0	0.0	0.0	0.0	0.0	1	5989	2986
## theta[26]	0.0	0.0	0.0	0.0	0.0	1	6371	3326
## theta[27]	0.0	0.0	0.0	0.0	0.0	1	6444	3174
## theta[28]	0.0	0.0	0.0	0.0	0.0	1	6364	2919
## theta[29]	0.0	0.0	0.0	0.0	0.0	1	5376	2528
## theta[30]	0.0	0.0	0.0	0.0	0.0	1	5889	2674
## theta[31]	0.0	0.0	0.0	0.0	0.0	1	5735	3354
## theta[32]	0.0	0.0	0.0	0.0	0.0	1	5465	2648
## theta[33]	0.0	0.0	0.0	0.0	0.0	1	6982	3060
## theta[34]	0.0	0.0	0.0	0.0	0.0	1	6443	3212
## theta[35]	0.0	0.0	0.0	0.0	0.0	1	5556	2792
## theta[36]	0.0	0.0	0.0	0.0	0.0	1	4787	2961
## theta[37]	0.0	0.0	0.0	0.0	0.0	1	4838	2725
## alpha	8.6	13.9	22.0	14.4	4.1	1	3382	3284
## beta	584.7	929.5	1475.8	964.6	277.1	1	3377	3313
## log_lik[1]	-3.6	-2.5	-2.4	-2.7	0.5	1	2806	2537
## log_lik[2]	-6.3	-4.6	-4.4	-4.9	0.7	1	1815	2200
## log_lik[3]	-4.9	-3.3	-3.1	-3.5	0.6	1	2301	2338
## log_lik[4]	-5.6	-3.9	-3.7	-4.2	0.7	1	1504	2101
## log_lik[5]	-5.7	-4.1	-3.9	-4.4	0.6	1	2291	2659
## log_lik[6]	-6.2	-4.6	-4.4	-4.9	0.7	1	2014	2263
## log_lik[7]	-5.5	-3.9	-3.7	-4.2	0.7	1	1756	2159
## log_lik[8]	-5.2	-3.7	-3.5	-3.9	0.6	1	2138	2460
## log_lik[9]	-6.2	-3.7	-3.3	-4.1	1.0	1	4163	3563
## log_lik[10]	-5.0	-3.5	-3.3	-3.7	0.6	1	1778	2034
## log_lik[11]	-4.9	-3.2	-3.0	-3.5	0.7	1	2476	2770
## log_lik[12]	-5.9	-4.0	-3.8	-4.3	0.8	1	1862	1942
## log_lik[13]	-5.1	-3.5	-3.3	-3.7	0.7	1	2208	2762
## log_lik[14]	-5.5	-4.0	-3.8	-4.2	0.7	1	2128	2482
## log_lik[15]	-5.8	-4.0	-3.8	-4.3	0.7	1	1911	1770
## log_lik[16]	-7.6	-4.4	-3.6	-4.8	1.4	1	4210	3071

## log_lik[17]	-6.1	-4.3	-4.0	-4.6	0.8	1	2145	2210
## log_lik[18]	-5.9	-4.2	-4.0	-4.5	0.7	1	1835	2187
## log_lik[19]	-4.0	-2.6	-2.4	-2.8	0.6	1	3661	2784
## log_lik[20]	-5.9	-4.2	-3.9	-4.5	0.7	1	2154	2328
## log_lik[21]	-4.9	-2.7	-2.0	-3.0	0.9	1	5308	3195
## log_lik[22]	-5.3	-3.7	-3.5	-4.0	0.7	1	2227	2536
## log_lik[23]	-5.3	-3.8	-3.6	-4.1	0.6	1	1670	2384
## log_lik[24]	-5.4	-3.8	-3.6	-4.0	0.7	1	1713	2246
## log_lik[25]	-5.7	-4.0	-3.7	-4.2	0.7	1	1725	2235
## log_lik[26]	-5.7	-4.0	-3.7	-4.2	0.7	1	1989	2533
## log_lik[27]	-4.5	-3.2	-3.0	-3.4	0.6	1	2083	2617
## log_lik[28]	-4.7	-3.3	-3.1	-3.6	0.6	1	2124	2532
## log_lik[29]	-5.1	-3.6	-3.4	-3.8	0.6	1	1827	2284
## log_lik[30]	-4.3	-2.7	-2.3	-2.9	0.7	1	4274	2674
## log_lik[31]	-5.7	-4.0	-3.8	-4.3	0.7	1	2182	2563
## log_lik[32]	-5.0	-3.5	-3.3	-3.8	0.6	1	2077	2128
## log_lik[33]	-5.3	-3.8	-3.6	-4.0	0.6	1	2029	2316
## log_lik[34]	-5.4	-3.8	-3.6	-4.0	0.7	1	2067	2359
## log_lik[35]	-5.9	-4.3	-4.1	-4.6	0.7	1	1900	2465
## log_lik[36]	-6.3	-3.8	-2.5	-4.1	1.2	1	4866	2961
## log_lik[37]	-7.0	-4.1	-2.8	-4.4	1.4	1	4472	2359
## ypred[1]	9.0	16.0	25.0	16.6	5.0	1	4322	3810
## ypred[2]	972.0	1045.0	1120.0	1045.7	45.5	1	5102	3760
## ypred[3]	53.0	70.0	90.0	70.6	11.3	1	5055	3723
## ypred[4]	237.0	274.0	313.0	274.4	23.0	1	5346	3760
## ypred[5]	358.0	401.0	448.0	401.7	27.2	1	4951	3924
## ypred[6]	1035.0	1109.0	1188.0	1110.2	46.5	1	5314	4037
## ypred[7]	218.0	254.0	290.0	254.2	21.8	1	4937	3524
## ypred[8]	138.0	165.0	196.0	165.7	17.9	1	5772	3666
## ypred[9]	83.0	104.0	129.0	105.0	13.8	1	4248	3295
## ypred[10]	86.0	109.0	134.0	109.3	14.4	1	4953	3937
## ypred[11]	55.0	72.0	92.0	72.5	11.5	1	4289	3736
## ypred[12]	273.0	312.0	354.0	311.9	24.3	1	4178	3431
## ypred[13]	85.0	106.0	130.0	106.3	13.8	1	4716	3798
## ypred[14]	264.0	301.0	342.0	301.6	23.9	1	4621	3295
## ypred[15]	299.0	341.0	385.0	341.7	26.1	1	4863	3810
## ypred[16]	167.0	198.0	230.0	198.1	19.3	1	4810	3725
## ypred[17]	453.0	502.0	556.0	502.5	31.6	1	5004	3756
## ypred[18]	444.0	493.0	546.0	493.4	30.9	1	5079	3719
## ypred[19]	9.0	15.0	24.0	15.7	4.8	1	4900	4047
## ypred[20]	363.0	409.0	457.0	409.0	28.7	1	5625	3534
## ypred[21]	6.0	13.0	21.0	13.1	4.6	1	4094	3644
## ypred[22]	154.0	185.0	216.0	184.9	19.0	1	4801	3224
## ypred[23]	186.0	218.0	252.0	218.6	20.1	1	4688	3714
## ypred[24]	172.0	203.0	236.0	203.2	19.7	1	4894	3721
## ypred[25]	239.0	275.0	315.0	275.7	23.2	1	5048	3784
## ypred[26]	252.0	290.0	331.0	290.3	23.9	1	5298	3762
## ypred[27]	46.0	62.0	82.0	63.0	10.7	1	4674	3860
## ypred[28]	63.0	82.0	103.0	82.7	12.3	1	4598	3536
## ypred[29]	112.0	137.0	164.0	137.7	15.8	1	4804	3607
## ypred[30]	7.0	14.0	22.0	14.0	4.4	1	4638	3904
## ypred[31]	261.0	299.0	340.0	299.9	24.0	1	5124	4026
## ypred[32]	102.0	126.0	152.0	126.3	15.4	1	4501	3389
## ypred[33]	172.0	203.0	235.0	203.0	19.4	1	5005	3666

```

## ypred[34]      169.0    199.0    233.0    200.0   19.4     1    4791    3236
## ypred[35]      535.0    592.0    649.0    592.3   34.3     1    4289    3788
## ypred[36]        7.0     13.0     20.0     12.8    4.2     1    3903    3682
## ypred[37]       23.0     34.0     48.0     34.6    7.6     1    4409    4001
## lp__          -97360.6 -97352.2 -97345.9 -97352.6   4.5     1    1476    2319
##
## For each parameter, Bulk_ESS and Tail_ESS are crude measures of
## effective sample size for bulk and tail quantities respectively (an ESS > 100
## per chain is considered good), and Rhat is the potential scale reduction
## factor on rank normalized split chains (at convergence, Rhat <= 1.05).

```