

Fit Test Try 2

Blain Morin

December 21, 2017

$$Y_i = \alpha_{j[i]} + \sum \beta_p X_{pi} + \epsilon_i, \text{for hospitals}(i = 1 \text{ to } N)$$
$$\alpha_j = a + \sum b_k W_{kj} + u_j, \text{for markets}(j = 1 \text{ to } J)$$

```
### load required packages
library(lme4)

## Loading required package: Matrix
library(tidyverse)

## Loading tidyverse: ggplot2
## Loading tidyverse: tibble
## Loading tidyverse: tidyr
## Loading tidyverse: readr
## Loading tidyverse: purrr
## Loading tidyverse: dplyr

## Conflicts with tidy packages -----

## expand(): tidyr, Matrix
## filter(): dplyr, stats
## lag():    dplyr, stats
### rstan requires having rtools installed
library(rstan)

## Warning: package 'rstan' was built under R version 3.4.3
## Loading required package: StanHeaders
## Warning: package 'StanHeaders' was built under R version 3.4.3
## rstan (Version 2.16.2, packaged: 2017-07-03 09:24:58 UTC, GitRev: 2e1f913d3ca3)
## For execution on a local, multicore CPU with excess RAM we recommend calling
## rstan_options(auto_write = TRUE)
## options(mc.cores = parallel::detectCores())

##
## Attaching package: 'rstan'

## The following object is masked from 'package:tidyr':
##
##      extract

### read in data
k12ReducedRG = read_csv("k12ReducedRG.csv")

## Parsed with column specification:
## cols(
##   .default = col_double(),
##   episode = col_integer(),
```

```
## Provider = col_character(),
## hrr = col_character(),
## avgagehrr = col_integer(),
## `Rank for Variable dshpct` = col_integer(),
## `Rank for Variable cmi` = col_integer(),
## `Rank for Variable mdadjadmit` = col_integer(),
## qstarrating = col_integer(),
## urbanlocation = col_integer(),
## joinnetwork = col_integer(),
## jchaoaccredited = col_integer(),
## qieffort = col_integer(),
## reform = col_integer(),
## mdaffiliation = col_integer(),
## ownershipstatus = col_integer(),
## hospitalbedsize = col_integer()
## )

## See spec(...) for full column specifications.
```

Remove missing data for Stan:

```
###remove missing data colums
k12ReducedRG = k12ReducedRG %>%
  select(-reform, -joinnetwork)

###change data to only complete cases
k12ReducedRG = k12ReducedRG[complete.cases(k12ReducedRG),]

dim(k12ReducedRG)

## [1] 2920 28
```

Get m1 data ready for stan:

```
modelldata = list(episode = k12ReducedRG$episode,
  hrr = as.integer(as.factor(k12ReducedRG$hrr)),
  K = length(unique(k12ReducedRG$hrr)),
  N = nrow(k12ReducedRG))

modelistan = stan("model1.stan", data = modelldata, chains = 4, iter = 2000)
```

```
## Warning: There were 13 transitions after warmup that exceeded the maximum treedepth. Increase max_tr
## http://mc-stan.org/misc/warnings.html#maximum-treedepth-exceeded

## Warning: There were 2 chains where the estimated Bayesian Fraction of Missing Information was low. S
## http://mc-stan.org/misc/warnings.html#bfmi-low

## Warning: Examine the pairs() plot to diagnose sampling problems
```

Compare max and min from stan simulations to observed data:

```
###Extract Maxes
modellmaxes = extract(model1stan, pars = c("maximum"))

###Extract Mins
modellmins = extract(model1stan, pars = c("minimum"))

###Mean Max
mean(modellmaxes$maximum)
```

```
## [1] 29936.65
```

```
summary(modellmaxes$maximum)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  27605   29215   29817   29937   30510   35555
```

```
###Mean Min
```

```
mean(modellmins$minimum)
```

```
## [1] 8322.247
```

```
summary(modellmins$minimum)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   1355    7763    8460    8322    9042   10711
```

```
###Compare observed
```

```
summary(k12ReducedRG$episode)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   7119   17126   19269   19076   21028   41469
```

Get m2 data ready for stan:

```
###rename columns
k12ReducedRG = k12ReducedRG %>%
  rename(cmirank = "Rank for Variable cmi",
         dshpctrank = "Rank for Variable dshpct",
         mdadjadmitrank = "Rank for Variable mdadjadmit")
```

```
### prepare data for STAN
```

```
model2data = list(episode = k12ReducedRG$episode,
                  hrr = as.integer(as.factor(k12ReducedRG$hrr)),
                  qstar = k12ReducedRG$qstarrating,
                  qieffort = k12ReducedRG$qieffort,
                  accredited = k12ReducedRG$jchaoaccredited,
                  urban = k12ReducedRG$urbanlocation,
                  mdaffiliation = k12ReducedRG$mdaffiliation,
                  ownership = k12ReducedRG$ownershipstatus,
                  bedsize = k12ReducedRG$hospitalbedsize,
                  cmi = k12ReducedRG$cmirank,
                  dsh = k12ReducedRG$dshpctrank,
                  mdadjadmitrank = k12ReducedRG$mdadjadmitrank,
```



```

##
##
## SAMPLING FOR MODEL 'model2' NOW (CHAIN 1).
##
## Gradient evaluation took 0.002 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 20 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration: 1 / 600 [ 0%] (Warmup)
## Iteration: 60 / 600 [ 10%] (Warmup)
## Iteration: 120 / 600 [ 20%] (Warmup)
## Iteration: 180 / 600 [ 30%] (Warmup)
## Iteration: 240 / 600 [ 40%] (Warmup)
## Iteration: 300 / 600 [ 50%] (Warmup)
## Iteration: 301 / 600 [ 50%] (Sampling)
## Iteration: 360 / 600 [ 60%] (Sampling)
## Iteration: 420 / 600 [ 70%] (Sampling)
## Iteration: 480 / 600 [ 80%] (Sampling)
## Iteration: 540 / 600 [ 90%] (Sampling)
## Iteration: 600 / 600 [100%] (Sampling)
##
## Elapsed Time: 160.592 seconds (Warm-up)
##                275.245 seconds (Sampling)
##                435.837 seconds (Total)
##
##
## SAMPLING FOR MODEL 'model2' NOW (CHAIN 2).
##
## Gradient evaluation took 0 seconds
## 1000 transitions using 10 leapfrog steps per transition would take 0 seconds.
## Adjust your expectations accordingly!
##
##
## Iteration: 1 / 600 [ 0%] (Warmup)
## Iteration: 60 / 600 [ 10%] (Warmup)
## Iteration: 120 / 600 [ 20%] (Warmup)
## Iteration: 180 / 600 [ 30%] (Warmup)
## Iteration: 240 / 600 [ 40%] (Warmup)
## Iteration: 300 / 600 [ 50%] (Warmup)
## Iteration: 301 / 600 [ 50%] (Sampling)
## Iteration: 360 / 600 [ 60%] (Sampling)
## Iteration: 420 / 600 [ 70%] (Sampling)
## Iteration: 480 / 600 [ 80%] (Sampling)
## Iteration: 540 / 600 [ 90%] (Sampling)
## Iteration: 600 / 600 [100%] (Sampling)
##
## Elapsed Time: 164.989 seconds (Warm-up)
##                272.795 seconds (Sampling)
##                437.784 seconds (Total)
##
## Warning: There were 600 transitions after warmup that exceeded the maximum treedepth. Increase max_t
## http://mc-stan.org/misc/warnings.html#maximum-treedepth-exceeded
## Warning: Examine the pairs() plot to diagnose sampling problems

```

Compare simulation max and mean to observed

```
###Extract maxes
model2maxes = extract(model2stan, pars = c("maximum"))
```

```
###Extract mins
model2mins = extract(model2stan, pars = c("minimum"))
```

```
###Mean max
mean(model2maxes$maximum)
```

```
## [1] 17549.68
```

```
summary(model2maxes$maximum)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##    11120  14255   16725   17550   21065   26605
```

```
###Mean min
mean(model2mins$minimum)
```

```
## [1] -4097.518
```

```
summary(model2mins$minimum)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## -11744.9 -7222.8 -5036.7 -4097.5   -772.8   4218.3
```

```
###Compare observed
summary(k12ReducedRG$episode)
```

```
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##      7119   17126   19269   19076   21028   41469
```