

# Analysis of fit

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

```
### 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.

First, we create the null model with no fixed effects and a market random effect. This preliminary step allows us to partition the variance into hospital and market levels.

```
#### This is the code for a null model with one random effect (hospital referral region)
m1 = lmer(episode ~ 1 + (1 | hrr) , data = k12ReducedRG)

summary(m1)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: episode ~ 1 + (1 | hrr)
## Data: k12ReducedRG
##
## REML criterion at convergence: 59231.3
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -4.4593 -0.5832  0.0381  0.5955  6.8181
##
## Random effects:
## Groups Name Variance Std.Dev.
## hrr (Intercept) 1564856 1251
## Residual 8795309 2966
## Number of obs: 3132, groups: hrr, 304
##
## Fixed effects:
## Estimate Std. Error t value
## (Intercept) 18912.76 96.48 196
```

To check if this null model approximates our data, we can use STAN to draw samples. We use the parameters found above to specify our STAN code.

```
####prepare data for STAN

m1_stan_data = list(episode = k12ReducedRG$episode,
                    hrr = as.integer(as.factor(k12ReducedRG$hrr)),
                    K = length(unique(k12ReducedRG$hrr)))

#### Specify prior parameters
```

```

N = nrow(k12ReducedRG)

### Draw samples with stan
stan_null = stan(file = "m1.stan", data = m1_stan_data, iter = 2000, chains = 4)

## Warning: There were 1958 transitions after warmup that exceeded the maximum treedepth. Increase max_
## http://mc-stan.org/misc/warnings.html#maximum-treedepth-exceeded

## Warning: There were 4 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

print(stan_null, pars = c("beta", "sigma_e", "sigma_w"),
      probs=c(0.025, 0.5, 0.975))

## Inference for Stan model: m1.
## 4 chains, each with iter=2000; warmup=1000; thin=1;
## post-warmup draws per chain=1000, total post-warmup draws=4000.
##
##               mean se_mean      sd    2.5%    50%    97.5% n_eff Rhat
## beta[1]  19008.61   37.39 128.72 18730.04 19037.47 19199.73   12 1.53
## sigma_e   3135.59   57.31 148.57  2913.07  3205.28  3334.60    7 2.45
## sigma_w    545.35  231.12 586.96    2.48   50.68  1352.75    6 2.83
##
## Samples were drawn using NUTS(diag_e) at Mon Dec 18 13:40:23 2017.
## For each parameter, n_eff is a crude measure of effective sample size,
## and Rhat is the potential scale reduction factor on split chains (at
## convergence, Rhat=1).

```

The mean intercept in the stan model (18987) is similar to the observed data model (18912).

---

Next, we add the hospital level predictors.

```

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

m2 = lmer(episode ~ qstarrating +
  qieffort +
  jchaoaccredited +
  urbanlocation +
  mdaffiliation +
  ownershipstatus +
  hospitalbedsize +
  cmirank +
  dshpctrank +
  mdadjadmitrank +
  (1 | hrr),
  data = k12ReducedRG)

summary(m2)

## Linear mixed model fit by REML ['lmerMod']

```

```

## Formula:
## episode ~ qstarrating + qieffort + jchaoaccredited + urbanlocation +
##      mdaffiliation + ownershipstatus + hospitalbedsize + cmirank +
##      dshpctrank + mdadjadmitrank + (1 | hrr)
## Data: k12ReducedRG
##
## REML criterion at convergence: 52030.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -5.4273 -0.5418 -0.0134  0.5431 10.7281
##
## Random effects:
## Groups Name Variance Std.Dev.
## hrr (Intercept) 689961 830.6
## Residual 3006563 1733.9
## Number of obs: 2920, groups: hrr, 304
##
## Fixed effects:
## Estimate Std. Error t value
## (Intercept) 14119.06 256.32 55.08
## qstarrating -50.10 45.13 -1.11
## qieffort 108.84 82.86 1.31
## jchaoaccredited 373.81 95.31 3.92
## urbanlocation 656.51 95.18 6.90
## mdaffiliation 109.08 54.75 1.99
## ownershipstatus 335.50 58.98 5.69
## hospitalbedsize 565.36 71.54 7.90
## cmirank 1579.75 42.28 37.36
## dshpctrank -27.77 34.93 -0.80
## mdadjadmitrank -72.11 32.30 -2.23
##
## Correlation of Fixed Effects:
## (Intr) qstrrt qifftrt jchccr urbnlc mdfflt ownrsh hsptlb cmirnk
## qstarrating -0.660
## qieffort -0.076 -0.059
## jchaoccrdtd -0.179 0.001 -0.064
## urbanlocatn -0.086 0.020 -0.103 -0.022
## mdaffiliatn -0.181 0.099 -0.066 0.016 -0.024
## ownrshpstts -0.495 0.024 0.190 -0.108 -0.100 0.057
## hospitlbdsz -0.288 0.134 -0.157 -0.109 -0.156 -0.210 -0.002
## cmirank 0.210 -0.117 -0.138 -0.050 -0.262 -0.176 -0.056 -0.414
## dshpctrank -0.277 0.216 0.082 0.005 0.088 -0.118 0.062 -0.194 0.095
## mdadjdmtrnk -0.260 0.017 -0.006 0.037 0.108 -0.115 0.117 0.001 -0.010
## dshpct
## qstarrating
## qieffort
## jchaoccrdtd
## urbanlocatn
## mdaffiliatn
## ownrshpstts
## hospitlbdsz
## cmirank
## dshpctrank

```

```
## mdadjdmtrnk 0.007
```

Next, we check the model using STAN.

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

###change N
N = nrow(k12ReducedRG)

### prepare data for STAN
m2_stan_data = 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,
  K = length(unique(k12ReducedRG$hrr)),
  id = as.integer(as.factor(k12ReducedRG$Provider)))
```

```
### run STAN simulation
```

```
m2_stan = stan(file = "m2.stan", data = m2_stan_data, iter = 1000, chains = 2)
```

```
## Warning: There were 1000 transitions after warmup that exceeded the maximum treedepth. Increase max_
```

```
## http://mc-stan.org/misc/warnings.html#maximum-treedepth-exceeded
```

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

```
print(m2_stan, pars = c("beta", "sigma_e", "sigma_w"),
  probs=c(0.025, 0.5, 0.975))
```

```
## Inference for Stan model: m2.
```

```
## 2 chains, each with iter=1000; warmup=500; thin=1;
```

```
## post-warmup draws per chain=500, total post-warmup draws=1000.
```

```
##
```

	mean	se_mean	sd	2.5%	50%	97.5%	n_eff	Rhat
## beta[1]	19251.48	24.70	85.60	19059.61	19255.31	19403.53	12	1.10
## beta[2]	35.66	21.40	37.60	-10.57	21.71	121.77	3	2.53
## beta[3]	-141.96	70.90	101.48	-293.36	-158.04	110.68	2	1.60
## beta[4]	245.37	197.88	203.42	-32.88	239.57	556.85	1	4.13
## beta[5]	-114.83	66.81	137.65	-390.93	-71.71	97.42	4	2.36
## beta[6]	-112.49	183.18	194.72	-443.24	-100.73	162.15	1	3.87
## beta[7]	137.30	237.74	289.08	-143.15	15.92	750.13	1	4.51
## beta[8]	-69.54	55.30	66.31	-142.98	-95.21	85.12	1	2.13
## beta[9]	81.90	60.48	118.96	-220.59	36.66	289.46	4	1.57
## beta[10]	-61.41	24.65	62.60	-159.93	-71.01	89.94	6	1.67
## sigma_e	3014.89	32.67	58.78	2931.32	3007.04	3140.62	3	1.25
## sigma_w	1.05	0.54	0.54	0.48	0.98	1.81	1	9.94

```

##
## Samples were drawn using NUTS(diag_e) at Mon Dec 18 16:15:55 2017.
## For each parameter, n_eff is a crude measure of effective sample size,
## and Rhat is the potential scale reduction factor on split chains (at
## convergence, Rhat=1).
summary(m2)

## Linear mixed model fit by REML ['lmerMod']
## Formula:
## episode ~ qstarrating + qieffort + jchaoaccredited + urbanlocation +
##      mdaffiliation + ownershipstatus + hospitalbedsize + cmirank +
##      dshpctrank + mdadjadmitrank + (1 | hrr)
##      Data: k12ReducedRG
##
## REML criterion at convergence: 52030.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -5.4273 -0.5418 -0.0134  0.5431 10.7281
##
## Random effects:
##   Groups   Name              Variance Std.Dev.
##   hrr      (Intercept) 689961    830.6
##   Residual              3006563 1733.9
## Number of obs: 2920, groups: hrr, 304
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)   14119.06    256.32   55.08
## qstarrating    -50.10     45.13   -1.11
## qieffort       108.84     82.86    1.31
## jchaoaccredited 373.81     95.31    3.92
## urbanlocation  656.51     95.18    6.90
## mdaffiliation  109.08     54.75    1.99
## ownershipstatus 335.50     58.98    5.69
## hospitalbedsize 565.36     71.54    7.90
## cmirank        1579.75     42.28   37.36
## dshpctrank     -27.77     34.93   -0.80
## mdadjadmitrank -72.11     32.30   -2.23
##
## Correlation of Fixed Effects:
##              (Intr) qstrrt qiffrt jchccr urbnlc mdfflt ownrsh hsptlb cmirnk
## qstarrating -0.660
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## ownrshpstts -0.495  0.024  0.190 -0.108 -0.100  0.057
## hospitlbdsz -0.288  0.134 -0.157 -0.109 -0.156 -0.210 -0.002
## cmirank      0.210 -0.117 -0.138 -0.050 -0.262 -0.176 -0.056 -0.414
## dshpctrank   -0.277  0.216  0.082  0.005  0.088 -0.118  0.062 -0.194  0.095
## mdadjdmtrnk -0.260  0.017 -0.006  0.037  0.108 -0.115  0.117  0.001 -0.010
##
##              dshpct
## qstarrating

```

```
## qieffort
## jchaoccrdtd
## urbanlocatn
## mdaffiliatn
## ownrshpstts
## hospitlbdsz
## cmirank
## dshpctrank
## mdadjdmtrnk 0.007
```