MA678_homework_08

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Getting to know stan

Read through the tutorial on Stan https://github.com/stan-dev/rstan/wiki/RStan-Getting-Started

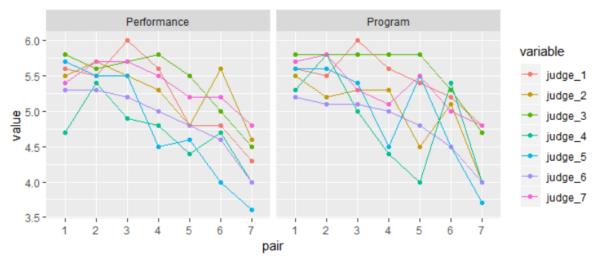
• Explore Stan website and Stan reference manual and try to connect them with Gelman and Hill 16 - 17.

Data analysis

Using stan:

The folder olympics has seven judges' ratings of seven figure skaters (on two criteria: "technical merit" and "artistic impression") from the 1932 Winter Olympics. Take a look at

http://www.stat.columbia.edu/~gelman/arm/examples/olympics/olympics1932.t xt



```
Program Performance pair
                                   Judge
##
## 1:
          5.6
                       5.6
                               1 judge 1
## 2:
          5.5
                       5.5
                               1 judge_2
                               1 judge_3
          5.8
                       5.8
## 3:
## 4:
          5.3
                       4.7
                               1 judge_4
          5.6
                       5.7
## 5:
                               1 judge_5
## 6:
          5.2
                       5.3
                               1 judge_6
```

use stan to fit a non-nested multilevel model (varying across skaters and judges) for the technical merit ratings.

```
https://github.com/stan-dev/example-models/blob/master/ARM/Ch.17/17.3_flight_simulator.stan https://github.com/stan-dev/example-models/blob/master/ARM/Ch.17/17.3_non-nested_models.R
```

```
fit_program<-lmer(Program~1+(1|pair) + (1|Judge),olympics_long)</pre>
dataList.1 <- list(N=49, n judges=7, n pairs=7, judge=as.integer(olymp</pre>
ics_long$Judge), pair=as.integer(olympics_long$pair), y=olympics_long$P
rogram)
skating_stan<-"
data {
 int<lower=0> N;
  int<lower=0> n_judges;
  int<lower=0> n pairs;
  int<lower=0,upper=n judges> judge[N];
  int<lower=0,upper=n_pairs> pair[N];
 vector[N] y;
}
parameters {
  real<lower=0> sigma;
  real<lower=0> sigma gamma;
  real<lower=0> sigma_delta;
 vector[n judges] gamma;
 vector[n pairs] delta;
 real mu;
}
model {
 vector[N] y_hat;
  sigma ~ uniform(0, 100);
  sigma_gamma ~ uniform(0, 100);
  sigma delta ~ uniform(0, 100);
  mu \sim normal(0, 100);
  gamma ~ normal(0, sigma_gamma);
  delta ~ normal(0, sigma delta);
 for (i in 1:N)
    y_hat[i] = mu + gamma[judge[i]] + delta[pair[i]];
 y ~ normal(y hat, sigma);
}
"
```

```
pilots <- read.table
("http://www.stat.columbia.edu/~gelman/arm/examples/pilots/pilots.dat",
header=TRUE)</pre>
```

flight_simulator.sf1 <- stan(model_code=skating_stan, data=dataList.1, iter=2000, chains=4)

Multilevel logistic regression

The folder speed.dating contains data from an experiment on a few hundred students that randomly assigned each participant to 10 short dates with participants of the opposite sex (Fisman et al., 2006). For each date, each person recorded several subjective numerical ratings of the other person (attractiveness, compatibility, and some other characteristics) and also wrote down whether he or she would like to meet the other person again. Label $y_{ij}=1$ if person i is interested in seeing person j again 0 otherwise. And r_{ij1}, \ldots, r_{ij6} as person i's numerical ratings of person j on the dimensions of attractiveness, compatibility, and so forth. Please look at

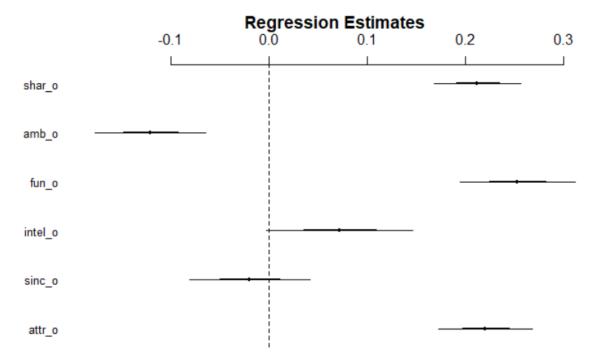
http://www.stat.columbia.edu/~gelman/arm/examples/speed.dating/Speed%20D ating%20Data%20Key.doc for details.

```
dating<-fread("http://www.stat.columbia.edu/~gelman/arm/examples/speed.
dating/Speed%20Dating%20Data.csv")</pre>
```

1. Fit a classical logistic regression predicting $Pr(y_{ij} = 1)$ given person i's 6 ratings of person j. Discuss the importance of attractiveness, compatibility, and so forth in this predictive model.

```
#classical logistic model without mixed effcects
#with attr o: attractiveness, sinc o: sincerity, intel o: intelligence,
fun_o: fun, amb_o: ambition, shar_o: shared interest/hobbies
log 1 <- glm(match~attr o +sinc o +intel o +fun o +amb o +shar o,data=d
ating,family=binomial)
summary(log_1)
##
## Call:
## glm(formula = match ~ attr_o + sinc_o + intel_o + fun_o + amb_o +
      shar_o, family = binomial, data = dating)
##
##
## Deviance Residuals:
##
      Min
                10
                     Median
                                 3Q
                                         Max
## -1.5300 -0.6362 -0.4420 -0.2381
                                      3.1808
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
##
                         0.21859 -25.714 < 2e-16 ***
## (Intercept) -5.62091
## attr_o 0.22047
                         0.02388
                                   9.233 < 2e-16 ***
## sinc o
                         0.03067 -0.651
                                           0.5152
              -0.01996
## intel_o 0.07176 0.03716 1.931
                                           0.0535 .
```

```
## fun o
             0.25315
                          0.02922
                                   8.665 < 2e-16 ***
## amb o
              -0.12099
                          0.02838 -4.264 2.01e-05 ***
                          0.02209
## shar_o
               0.21225
                                   9.608 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 6466.6 on 7030
                                     degrees of freedom
## Residual deviance: 5611.0 on 7024 degrees of freedom
    (1347 observations deleted due to missingness)
## AIC: 5625
##
## Number of Fisher Scoring iterations: 5
coefplot(log_1)
```



The result of this model can be written like:

$$P(match = 1) \\ = logit^{-1}(-5.6 + 0.22attr - 0.02sinc + 0.07intel + 0.25fun - 0.12amb \\ + 0.21shar)$$

(If we apply the "divided by 4" method to explain coefficients)

Attractiveness: 0.22/4 = 0.055, one point higher in attractiveness will lead to 5.5% higher willingness of another date.

Sincerity: -0.02/4 = -0.005, one point higher in sincerity will lead to 0.5% lower willingness of another date, which is contrary to our expectation.

Intelligence: 0.07/4 = 0.0175, one point higher in intelligence will lead to 1.75% higher willingness of another date.

Fun: 0.25/4 = 0.0625, one point higher in humor will lead to 6.25% higher willingness of another date.

Ambition: -0.12/4 = -0.03, one point higher in ambition will lead to 3% lower willingness of another date.

Shared interest: 0.21/4 = 0.0525, one point higher in shared interest will lead to 5.25% higher willingness of another date.

According to these results, the most important to determine a person's willingness to date again is humor and attractiveness.

2. Expand this model to allow varying intercepts for the persons making the evaluation; that is, some people are more likely than others to want to meet someone again. Discuss the fitted model.

```
log mix <- lmer(match~gender+scale(attr o) +scale(sinc o) +scale(intel</pre>
o) +scale(fun o) +scale(amb o) +scale(shar o)+(1|iid),data=dating,famil
y=binomial(link="logit"))
summary(log_mix)
## Generalized linear mixed model fit by maximum likelihood (Laplace
     Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula:
## match ~ gender + scale(attr o) + scale(sinc o) + scale(intel o) +
       scale(fun_o) + scale(amb_o) + scale(shar_o) + (1 | iid)
##
      Data: dating
##
## Control:
## structure(list(optimizer = c("bobyqa", "Nelder_Mead"), calc.derivs =
TRUE,
       use.last.params = FALSE, restart edge = FALSE, boundary.tol = 1e
##
-05,
      tolPwrss = 1e-07, compDev = TRUE, nAGQ@initStep = TRUE, checkCon
##
trol = list(
##
           check.nobs.vs.rankZ = "ignore", check.nobs.vs.nlev = "stop",
           check.nlev.gtreq.5 = "ignore", check.nlev.gtr.1 = "stop",
##
           check.nobs.vs.nRE = "stop", check.rankX = "message+drop.cols
##
           check.scaleX = "warning", check.formula.LHS = "stop",
##
           check.response.not.const = "stop"), checkConv = list(
##
           check.conv.grad = list(action = "warning", tol = 0.001,
##
               relTol = NULL), check.conv.singular = list(action = "ign
##
ore",
```

```
##
               tol = 1e-04), check.conv.hess = list(action = "warning",
               tol = 1e-06)), optCtrl = list()), class = c("glmerContro
##
1",
## "merControl"))
##
##
        AIC
                 BIC
                       logLik deviance df.resid
##
     5543.2
              5605.0 -2762.6
                                5525.2
                                           7022
##
## Scaled residuals:
       Min
                10 Median
                                3Q
## -1.7458 -0.4453 -0.2877 -0.1454 10.3764
##
## Random effects:
## Groups Name
                       Variance Std.Dev.
## iid
           (Intercept) 0.4294
                                0.6553
## Number of obs: 7031, groups: iid, 551
##
## Fixed effects:
##
                  Estimate Std. Error z value Pr(>|z|)
                              0.07079 -30.122 < 2e-16 ***
## (Intercept)
                  -2.13226
                              0.09322
## gender
                   0.15452
                                        1.658
                                                0.0974 .
## scale(attr_o)
                   0.46048
                              0.05203
                                        8.850 < 2e-16 ***
                                                0.6658
## scale(sinc_o) -0.02474
                              0.05728 -0.432
## scale(intel o) 0.10874
                              0.06203
                                        1.753
                                                0.0796 .
## scale(fun o)
                   0.51341
                              0.06192
                                        8.291 < 2e-16 ***
## scale(amb_o)
                              0.05468 -4.311 1.63e-05 ***
                  -0.23570
## scale(shar_o)
                  0.48473
                              0.05045
                                        9.609 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##
               (Intr) gender scl(t_) scl(sn_) scl(n_) scl(f_) scl(m_)
## gender
               -0.672
## scale(ttr_) -0.202 0.109
## scale(snc ) -0.022 0.048 -0.123
## scale(ntl_) 0.026 -0.055 -0.039 -0.466
## scale(fun_) -0.156 0.015 -0.246 -0.150
                                              -0.132
## scale(amb_) 0.143 -0.092 -0.062 -0.014
                                              -0.370
                                                      -0.187
## scale(shr_) -0.135 0.009 -0.100 -0.054
                                              -0.005
                                                      -0.268
                                                              -0.203
## convergence code: 0
## unable to evaluate scaled gradient
## Model failed to converge: degenerate Hessian with 6 negative eigenv
alues
#ranef(log_mix)
P(match = 1)
= logit^{-1}(-2.13 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc))
```

+0.11scale(intel) +0.51scale(fun) -0.23scale(amb) +0.48scale(shar) $+iid_i$)

Among fixed effects: Gender: a male dating partner will receive approximately $0.16/4 = 0.0375 \, 4\%$ higher score from the person who give rates.

Attractiveness: 0.46/4 = 0.115, one point higher than average attractiveness rate will lead to 11.5% higher willingness of another date.

Sincerity: -0.02/4 = -0.005, one point higher than average sincerity rate will lead to 0.5% lower willingness of another date, which is contrary to our expectation.

Intelligence: 0.11/4 = 0.0275, one point higher than average intelligence rate will lead to 2.75% higher willingness of another date.

Fun: 0.51/4 = 0.1275, one point higher than average humor rate will lead to 12.75% higher willingness of another date.

Ambition: -0.23/4 = -0.0575, one point higher than average ambition rate will lead to 5.75% lower willingness of another date.

Shared interest: 0.48/4 = 0.12, one point higher than average shared interest rate will lead to 12% higher willingness of another date.

For random effects: If we take the first 5 people for example: 1 0.493948779

- 2 -0.181380981
- 3 -0.467790677
- 4 -0.107449058
- 5 0.124412503

The model with random effects for these five people can be written as follow:

```
P(match = 1) \\ = logit^{-1}(-1.64 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.31 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.59 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.24 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.01 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.01 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.01 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.01 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.01 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.11scale(intel) + 0.51scale(fun) - 0.23scale(amb) + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.01 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.48scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.01 + 0.15gender + 0.46scale(attr) - 0.02scale(sinc) \\ + 0.48scale(shar)) \\ P(match = 1) \\ P(match = 1
```

3. Expand further to allow varying intercepts for the persons being rated. Discuss the fitted model.

```
log_mix2 <- glmer(match~gender+scale(attr_o) +scale(sinc_o) +scale(inte</pre>
1_o) +scale(fun_o) +scale(amb_o) +scale(shar_o)+(1|iid)+(1|pid),data=da
ting, family=binomial)
## Warning in checkConv(attr(opt, "derivs"), opt$par, ctrl =
## control$checkConv, : Model failed to converge with max|grad| = 0.217
492
## (tol = 0.001, component 1)
summary(log_mix2)
## Generalized linear mixed model fit by maximum likelihood (Laplace
     Approximation) [glmerMod]
## Family: binomial ( logit )
## Formula:
## match ~ gender + scale(attr o) + scale(sinc o) + scale(intel o) +
       scale(fun_o) + scale(amb_o) + scale(shar_o) + (1 | iid) +
##
       (1 | pid)
      Data: dating
##
##
##
                 BIC
                       logLik deviance df.resid
        AIC
##
     5257.6
              5326.1
                      -2618.8
                                5237.6
                                           7021
##
## Scaled residuals:
       Min
                10 Median
                                30
                                       Max
## -3.7839 -0.3829 -0.2197 -0.0918 9.1741
##
## Random effects:
## Groups Name
                       Variance Std.Dev.
           (Intercept) 0.594
                                0.7707
## iid
           (Intercept) 1.254
## pid
                                1.1199
## Number of obs: 7031, groups: iid, 551; pid, 537
##
## Fixed effects:
##
                  Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                  -2.53701
                              0.11724 -21.639 < 2e-16 ***
                                        1.180
## gender
                              0.14929
                   0.17624
                                                0.2378
## scale(attr o)
                              0.06370 10.000 < 2e-16 ***
                   0.63703
## scale(sinc o)
                   0.03504
                              0.06782
                                        0.517
                                                0.6054
## scale(intel o) 0.17216
                              0.07357
                                        2.340
                                                0.0193 *
## scale(fun o)
                   0.57822
                              0.07097
                                        8.147 3.73e-16 ***
                              0.06463 -2.608
## scale(amb o)
                  -0.16857
                                                0.0091 **
## scale(shar o)
                              0.06155
                                        9.575 < 2e-16 ***
                   0.58936
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
               (Intr) gender scl(t_) scl(sn_) scl(n_) scl(f_) scl(m_)
```

```
## gender -0.647
## scale(ttr ) -0.221 0.094
## scale(snc_) -0.049 0.037 -0.065
## scale(ntl ) -0.009 -0.044 -0.024
                                    -0.438
## scale(fun_) -0.140 0.009 -0.220 -0.123
                                              -0.098
## scale(amb_) 0.072 -0.070 -0.051
                                      0.011
                                              -0.334 -0.168
## scale(shr ) -0.139  0.005 -0.072 -0.057
                                              -0.020 -0.234 -0.159
## convergence code: 0
## Model failed to converge with max|grad| = 0.217492 (tol = 0.001, com
ponent 1)
a iid <- data.frame(ranef(log mix2))[1:5,4]</pre>
a_pid <- data.frame(ranef(log_mix2))[552:556,4]</pre>
a_{iid}
## [1] 0.452970133 -0.463750802 -0.795488077 -0.323614171 -0.004300494
a_pid
## [1] 0.97137439 0.11231247 -1.78993378 -0.63479896 0.04949994
```

This model has added another varying intercept, if we still take the first five people as example, now their models will become (taking the changed fixed effects into consideration):

```
P(match = 1) \\ = logit^{-1}(-1.11 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.88 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-5.12 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-3.48 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.49 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.49 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.49 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.49 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.49 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.49 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar)) \\ P(match = 1) \\ = logit^{-1}(-2.49 + 0.16gender + 0.64scale(attr)0.035scale(sinc) \\ + 0.17scale(intel) + 0.58scale(fun) - 0.16scale(amb) + 0.59scale(shar) \\ P(match = 1) \\ P(m
```

4. You will now fit some models that allow the coefficients for attractiveness, compatibility, and the other attributes to vary by person. Fit a no-pooling model: for each person i, fit a logistic regression to the data y_{ij} for the 10 persons j whom he or she rated, using as predictors the 6 ratings r_{ij1} , ..., r_{ij6} .

(Hint: with 10 data points and 6 predictors, this model is difficult to fit. You will need to simplify it in some way to get reasonable fits.)

```
#No pooling model for each person i
no_pooling <- glm(match~attr_o + sinc_o + intel_o + fun_o + amb_o + sha
r_o + factor(iid)-1,data=dating)</pre>
```

5. Fit a multilevel model, allowing the intercept and the coefficients for the 6 ratings to vary by the rater i.

```
#Vary characteristics by person
vary_slope <- glmer(match~(1+attr_o+sinc_o+intel_o+fun_o+amb_o+shar_o|i</pre>
id) + attr o + sinc o + intel o + fun o + amb o + shar o,data=dating,fa
mily=binomial)
## Warning in optwrap(optimizer, devfun, start, rho$lower, control =
## control, : convergence code 1 from bobyga: bobyga -- maximum number
of
## function evaluations exceeded
## Warning in (function (fn, par, lower = rep.int(-Inf, n), upper =
## rep.int(Inf, : failure to converge in 10000 evaluations
## Warning in checkConv(attr(opt, "derivs"), opt$par, ctrl =
## control$checkConv, : unable to evaluate scaled gradient
## Warning in checkConv(attr(opt, "derivs"), opt$par, ctrl =
## control$checkConv, : Model failed to converge: degenerate Hessian wi
th 3
## negative eigenvalues
```

6. Compare the inferences from the multilevel model in (5) to the no-pooling model in (4) and the complete-pooling model from part (1) of the previous exercise.

```
anova(vary slope, log 1, no pooling)
## Data: dating
## Models:
## log 1: match ~ attr o + sinc o + intel o + fun o + amb o + shar o
## vary_slope: match ~ (1 + attr_o + sinc_o + intel o + fun o + amb o +
shar o
## vary slope:
                  iid) + attr o + sinc o + intel o + fun o + amb o + s
## no pooling: match ~ attr o + sinc o + intel o + fun o + amb o + shar
0 +
## no_pooling:
                  factor(iid) - 1
                           BIC logLik deviance Chisq Chi Df Pr(>Chis
##
              Df
                    AIC
q)
## log_1
              7 5625.0 5673.0 -2805.5
                                         5611.0
## vary_slope 35 5576.8 5816.8 -2753.4
                                         5506.8 104.2
                                                           28 1.047e-
10 ***
```

```
## no_pooling 558 5607.8 9434.6 -2245.9 4491.8 1015.0 523 < 2.2e-
16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1</pre>
```

From the anova test, we can see that the deviance and the AIC of no pooling model are the lowest, which is weird.