Lecture 7: Hierarchical Modeling

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2020-11-29

Announcements

- Fill out ESCI evaluations online!
 - What did you like? Would you like to see more courses like this?
 - What can be improved?

Comparing Multiple Related Groups

- Hierarchy of nested populations
- Models which account for this are called *hiearchical* or *multi-level* models

Some examples:

- Patient outcomes within several different hospitals
- People within counties in the United States (e.g. Asthma mortality example)
- Athlete performance in sports
- Genes within a group of animals

- A study was performed for the Educational Testing Service (ETS) to evaluate the effects of coaching programs on SAT preparation
- Each of eight different schools used a short-term SAT prep coaching program
- Compute the average SAT score in those who did take the program minus those that did not participate in the program
- We observe the average difference varies by school. What accounts for these differences?

- Interested in "real" differences due to training
- Want to reduce effect of chance variability
- How do we estimate the effect of the program in each of the schools?
- Two extremes:
 - Estimate the effect of the program in every school independently
 - A separate prior distribution for each school effect
 - Or assume the effect is the same in every school
 - Combine all the data
 - A compromise between the above 2 options?

$$y_j \sim N(heta_j, \sigma_j^2)$$

- y_j is the observed effects of the program in school j
 - Based on a sample of test scores from those in the program and those not in the program
- θ_j are the true *unknown* effects of the program in school j
- Variances, σ_j^2 , are *known*
 - Determined by the number of students in the sample

```
J <- 8

y = c(28, 8, -3, 7, -1, 1, 18, 12)

sigma <- c(15, 10, 16, 11, 9, 11, 10, 18)
```

- Assuming the effect of the program on each school is identical.
- What are the chances of seeing a value as large as 28?
- As small as -3?

If effects are actual equal, what is it?

```
## Compute the precision frome each school
prec <- 1/sigma^2

## global estimate is a weighted vareage
mu_global <- sum(prec * y / sum(prec))
mu_global</pre>
```

```
## [1] 7.685617
```

- Assume the effect of the program on each school is identical, i.e. $\theta_i = \mu$
- What are the chances of seeing a value as large as 28?
- As small as -3?

```
# 1000 datasets with mean mu_global but different sigmas

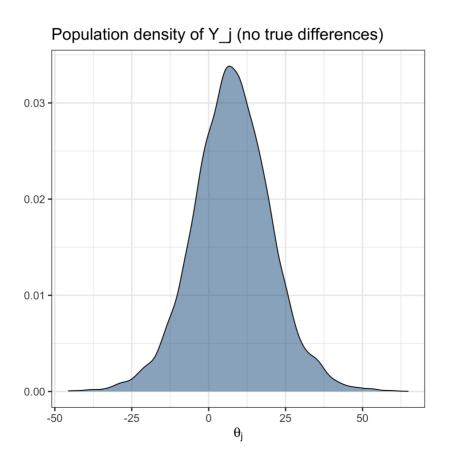
## Chance of seeing a value greater than 28
mean(sapply(1:1000, function(x)
    max(rnorm(J, mean=mu_global, sd=sigma))) >= 28)
```

[1] 0.378

```
## Chance of seeing a value less than -3
mean(sapply(1:1000, function(x)
  min(rnorm(J, mean=mu_global, sd=sigma))) <= -3)</pre>
```

[1] 0.81

Density of $Y_j \sim N(\theta_{\text{pooled}}, \sigma_j^2)$



$$y_j \sim N(heta_j, \sigma_j^2)$$

- θ_j are the true unknown effects of the program in school j
- y_i is the observed effects of the program in school j
 - Based on a sample of test scores from those in the program and those not in the program
 - \circ Number of people in the sample determine the magnitude of σ_j^2

- How do we estimate θ_i ?
 - \circ Independent: $\hat{ heta}_{j}^{(MLE)} = y_{j}$ is the MLE
 - $\circ \text{ Identical effects: } \hat{\theta}_j^{(pool)} = \frac{\sum_i \frac{1}{\sigma_i^2} y_i}{\sum_i \frac{1}{\sigma_i^2}}$
 - Same effect for all schools: estimate using a weighted average of the observed effects
- Compromise: $\hat{ heta}_j^{(shrink)} = w heta_j^{(MLE)} + (1-w) heta_j^{(pool)}$

Eight Schools

theta j mle <- y

```
theta_j_mle

## [1] 28 8 -3 7 -1 1 18 12

theta_j_pooled <- rep(sum(1/sigma^2 * y) / sum(1/sigma^2), J)
theta_j_pooled

## [1] 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.685617 7.6856
```

Is there a middle ground between two extremes?

A hierarchical model:

$$egin{aligned} heta_i &\sim N(\mu, au^2) \ y_j &\sim N(heta_j, \sigma_j^2) \end{aligned}$$

- Add a *shared* normal prior distribution to θ_j
- Assume the global mean, μ is also unknown
- How do we choose prior for μ ?

$$\mu \sim N(\mu_0, au_0^2)$$
? or $p(\mu) \propto 1$?

- Need to estimate all of $(\mu, \theta_1, \dots, \theta_8)$ with MCMC
- au^2 determines how much weight weight we put on the independent estimate vs the pooled estimate

Intuition behind shrinkage

- $Y_j = \theta_j + \epsilon_j$ and for simplicity assume that the variance of ϵ , σ^2 for all j
 - \circ θ_i represents true differences between schools (signal)
 - $\circ \epsilon_j$ is sampling variability (noise, chance variation)
- $ullet \ Var(Y_j) = Var(heta_j) + Var(\epsilon_j) = au^2 + \sigma^2$
 - The variance of the observed outcomes is the sum of signal variance, τ^2 , and the sampling variance σ_j^2
- Consequence: the observed outcomes always have higher variance across groups than the signal
 - $\circ \ \operatorname{Var}(Y_j) > \operatorname{Var}(heta_j)$
- Solution: reduce the variance by shrinking them!
 - \circ Want the variance of the shrunken estimates to be close to au^2

Comments:

- The global average, μ , is a parameter so also has uncertainty
- How dow we determine how much to shrink, e.g. how do we determine τ^2 ?
- Is the training program effective in school j?
 - What is $P(\theta_i > 0 \mid y)$?
- On avearge (over all schools) is the training program effective?
 - \circ What is $P(\mu > 0 \mid y)$?

- If τ^2 is large, the prior for θ_j is not very strong
 - \circ If $\tau^2 \to \infty$ equivalent to the no pooling model
- If τ^2 is small, we assume a priori that θ_j are very close
 - \circ if $au^2 o 0$ equivalent to the complete pooling model, $heta_j = \mu$

Estimating parameters

- MH: Need to generate a proposal from a 9-dimensional posterior distribution
 - \circ Eight parameters for θ_i and one for μ
- Gibbs: sample each of the 9 parameters from the complete conditionals
 - \circ Sample $p(\theta_j \mid \theta_{-j}, \mu)$
 - \circ Sample $p(\mu \mid \theta_1, \dots \theta_8, \mu)$
- Stan

Eight Schools Estimation

```
J <- 8

y = c(28, 8, -3, 7, -1, 1, 18, 12)

sigma <- c(15, 10, 16, 11, 9, 11, 10, 18)

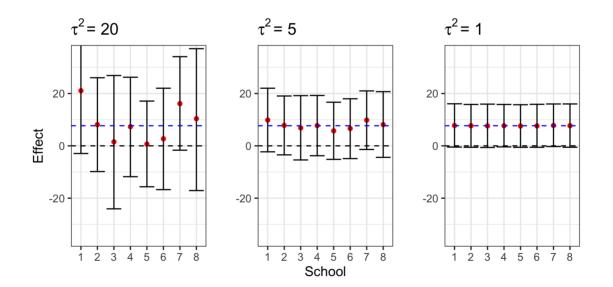
## fix the variance of the prior to a number

tau <- 20
```

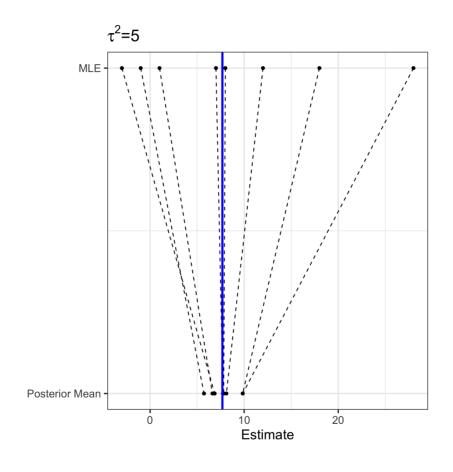
Eight Schools in Stan

```
// saved as 8schools.stan
data {
 // estimated treatment effects
 real v[J];
 real<lower=0> sigma[J]; // standard error of effect estimates
 real<lower=0> tau; // shrinkage standard deviation
parameters {
 real mu; // population treatment effect
 vector[J] eta; // unscaled deviation from mu by school
transformed parameters {
 vector[J] theta = mu + tau * eta; // school treatment effects
model {
 target += normal lpdf(y | theta, sigma); // log-likelihood
```

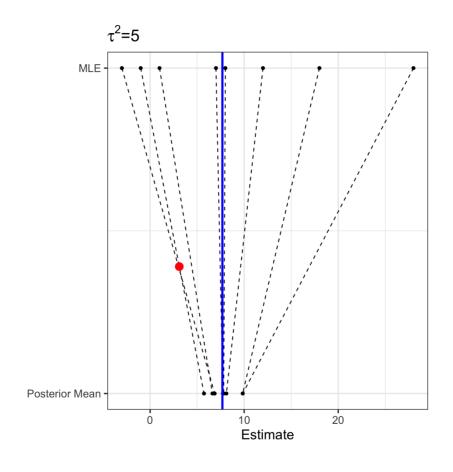
```
## `summarise()` ungrouping output (override with `.groups` argument)
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```



MLE vs Posterior Mean



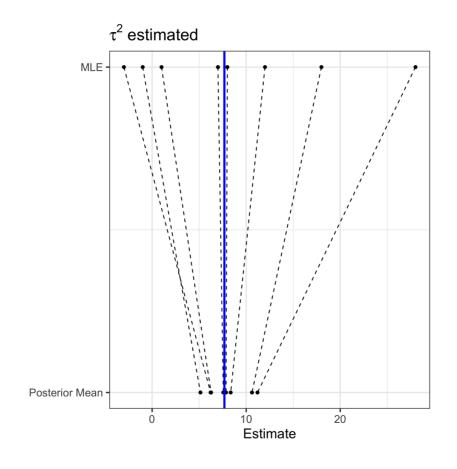
MLE vs Posterior Mean



Eight Schools in Stan

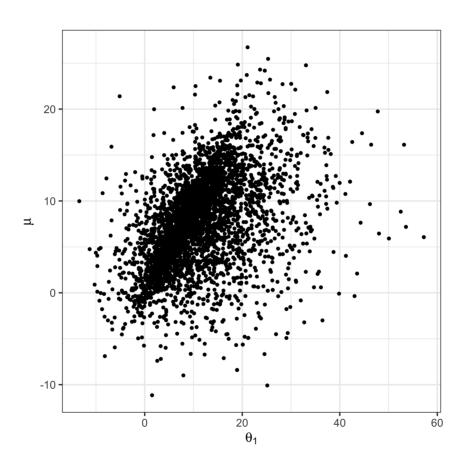
```
// saved as 8schools stan
data {
 real y[J]; // estimated treatment effects
 real<lower=0> sigma[J]; // standard error of effect estimates
parameters {
 real mu; // population treatment effect
 real<lower=0> tau; // standard deviation in treatment effects
 vector[J] eta;  // unscaled deviation from mu by school
transformed parameters {
 vector[J] theta = mu + tau * eta; // school treatment effects
model {
 target += -2log(tau); // prior for tau
 target += normal lpdf(eta | 0, 1); // normal model for the
 target += normal lpdf(y | theta, sigma); // log-likelihood
```

MLE vs Posterior Mean

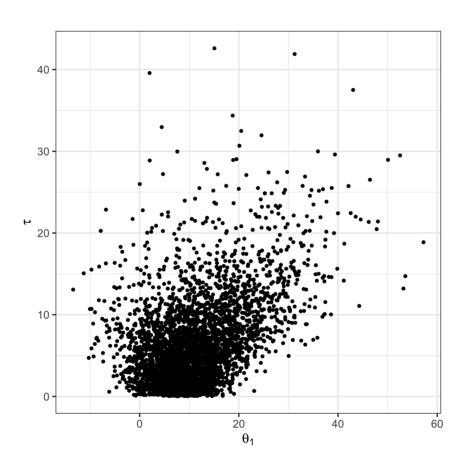


Posterior mean of $\tau^2 = 6.4633808$

Eight Schools Scatter Plot: θ_1 vs μ



Eight Schools Scatter Plot: θ_1 vs τ

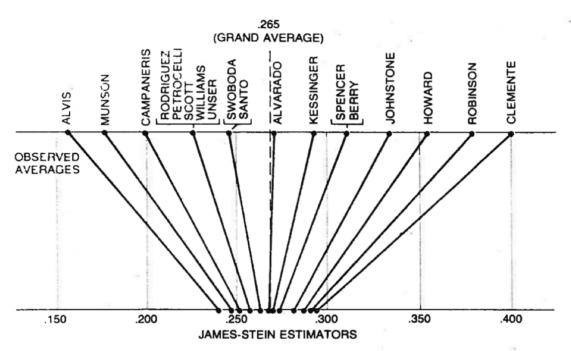


Hierarchical modeling in sports

1. 1970 Batting Averages for 18 Major League Players and Transformed Values $X_i, \; \theta_i$

i	Player	Y, = batting average for first 45 at bats (1)	p; = batting average for remainder of season (2)	At bats for remainder of season (3)	X; (4)	θ _i (5)
2	F. Robinson (Balt, AL)	.378	.298	426	-1.66	-2.79
3	F. Howard (Wash, AL)	.356	.276	521	-1.97	-3.11
4	Johnstone (Cal, AL)	.333	.222	275	-2.28	-3.96
5 6	Berry (Chi, AL)	.311	.273	418	-2.60	−3.17
6	Spencer (Cal, AL)	.311	.270	466	-2.60	-3.20
7	Kessinger (Chi, NL)	.289	.263	586	-2.92	-3.32
8	L. Alvarado (Bos, AL)	.267	.210	138	-3.26	-4.15
9	Santo (Chi, NL)	.244	.269	510	-3.60	-3.23
10	Swoboda (NY, NL)	.244	.230	200	-3.60	-3.83
11	Unser (Wash, AL)	.222	.264	277	-3.95	-3.30
12	Williams (Chi, AL)	.222	.256	270	-3.95	-3.43
13	Scott (Bos, AL)	.222	.303	435	-3.95	-2.71
14	Petrocelli (Bos, AL)	.222	.264	538	-3.95	-3.30
15	E. Rodriguez (KC, AL)	.222	.226	186	-3.95	-3.89
16	Campaneris (Oak, AL)	.200	.285	558	-4.32	-2.98
17	Munson (NY, AL)	.178	.316	408	-4.70	-2.53
18	Alvis (Mil, NL)	.156	.200	70	-5.10	-4.32

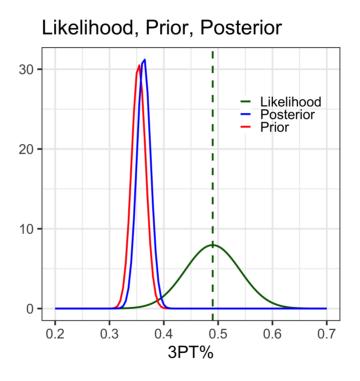
Hierarchical modeling in sports



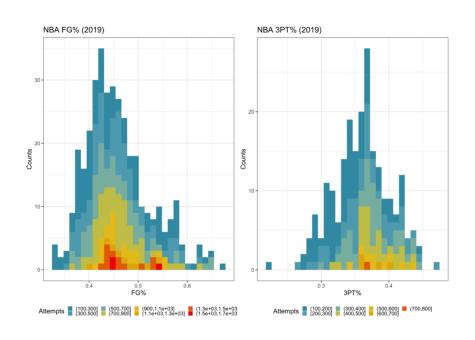
JAMES-STEIN ESTIMATORS for the 18 baseball players were calculated by "shrinking" the individual batting averages toward the overall "average of the averages." In this case the grand average is .265 and each of the averages is shrunk about 80 percent of the distance to this value. Thus the theorem on which Stein's method is based asserts that the true batting abilities are more tightly clustered than the preliminary batting averages would seem to suggest they are.

Basketball Example

After 100 shots Robert Covington's 3PT% was 0.49



Basketball Example



Eight Schools and Independence

- The effects θ_j in each school are *not* independent
- E.g. if we know θ_1 is really large, it means θ_j is also more likely to be large
- However, before we see data the distributions of θ_j 's, are indistinguishable
 - Don't know which θ 's will be large and which small
 - This kind of dependence is known as exchangeability

An exchangeable sequence of random variables is a finite (or infinite) sequence $X_1, X_2, ... X_n$ of random variables such that for any finite permutation π of the indices 1, 2, 3, ..., the joint probability distribution of the permuted sequence

$$X_{\pi(1)}, X_{\pi(2)}, X_{\pi(3)}, \dots$$

is the same as the joint probability distribution of the original sequence.

- Consider a set of J experiments in which experiment j as data y_j and parameter θ_j
- The J experiments are related, but no information in the indices that distinguish θ_j
- Assume an exchangeable prior distribution: $p(\theta_1, \dots \theta_J) = p(\theta_{\pi_1}, \dots \theta_{\pi_J})$ where π is any permutation of the indices $1 \dots J$
- Equivalent to a prior assumption about symmetry among the parameters $(\theta_1, \dots \theta_J)$

- Example: All i.i.d random variables are exchangeable
- Example: Non-independent random variables can also be exchangeable
- I flip a coin 5 times and observe 2 heads. What is the distribution over the order in which they were flipped?
 - \circ Let X_i be a 1 if the *i*th outcome of the flip was heads and zero otherwise
 - \circ X_i are not independent because the number of observed heads is fixed
 - $\circ p(X_1, X_2, \dots X_5)$ is exchangeable

• Example: multivariate normal with common mean μ and equicorrelation

$$Cov(Y) = egin{pmatrix} 1 &
ho &
ho \
ho & 1 &
ho \
ho &
ho & 1 \end{pmatrix}$$

Non-exchangeable random variables

Examples of *not* exchangeable variables include

- Time series data (closer in time = more correlated)
- Spatial data (closer in space = more correlated)
- Typically, ignorance (about indices) implies exchangeability
- When exchangeability doesn't hold, can often assume conditional exchangeability

Mixture of i.i.d random variables

• i.i.d random variables are exchangeable:

$$p\left(heta_{1},\ldots, heta_{n}
ight)=\prod_{j=1}^{J}p(heta_{j}\mid\phi)$$

• Mixtures of i.i.d random variables:

$$p\left(heta_{1},\ldots, heta_{n}
ight)=\int\left\{\prod_{j=1}^{n}p\left(heta_{j}|\phi
ight)
ight\}p(\phi)d\phi$$

- If ϕ were known, θ_i would be i.i.d.
- \circ Since ϕ is not known, integrate over uncertainty
- $\circ \theta_j$ are a mixture of i.i.d random variables
- Dependent but exchangeable

de Finetti's theorem

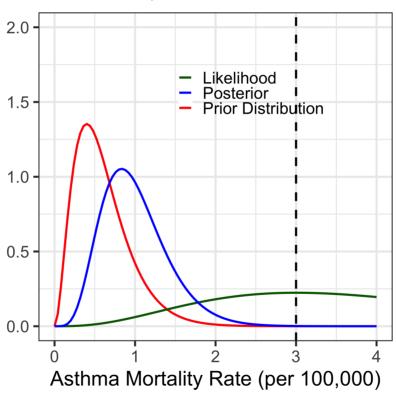
- Theorem: As $J \to \infty$ all exchangeable distributions can be expressed as a mixture of independent and identical distributions
- For a finite J, no guarantee that the variables can be represented as a mixture of i.i.d random variables
- But, if variables are exchangeable usually can be modeled as *approximately* a mixture of i.i.d
- Mixture's of i.i.d random variables \rightarrow a hierarchical model!
 - For this reason de Finetti's theorem is often cited
- Good discussion in 2.7-2.8 of Hoff book

Poisson Example, 1 county

- In a particular county 3 people out of a population of 100,000 died of asthma
- Assume a Poisson sampling model with rate λ (units are rate of deaths per 100,000 people)
- Want to know the true mortality rate, λ
- We discussed how to specify a prior distribution for λ

Asthma Mortality

Likelihood, Prior and Posterior



Poisson Example, many counties

- Assume we measure asthma mortality, y_i in n counties
- $Y_j \sim \operatorname{Pois}(\lambda_j)$
- Does it make sense to model λ_i exchangeable?
- If we don't know how j relates to geographic location

$$\circ \ \lambda_j \sim \operatorname{Gam}(a,b)$$
 and $p(a,b)$

• If we do know which geographic information relates to each j shouldn't model θ_j as exchangeabale

Poisson Example, many counties, many states

- Assume we measure asthma mortality, y_{ij} in county j in state i
- $Y_{ij} \sim \operatorname{Pois}(\lambda_{ij})$
- Does it make sense to model λ_{ij} as exchangeable?
- Model: $\lambda_{ij} \sim \operatorname{Gam}(a_i, b_i)$
- Can model county rates within state as exchangeable
- Can model parameters for state averages, (a_i, b_i) , as exchangeable