KU Leuven Summer School Segment 2B More Missing Data

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Stylized context (very similar to Segment 2A)

 X_1 : Binary indicator of high blood pressure (1=Yes)

 X_2 : Binary indicator of regular exercise (1=No)

Y: Binary indicator of heart disease (1=Yes)

Statistical model:

$$\mathsf{logit}\{\mathit{Pr}(\mathit{Y}=1|\mathit{X}_{1},\mathit{X}_{2})\} = \beta_{0} + \beta_{1}\mathit{X}_{1} + \beta_{2}\mathit{X}_{2}$$

But there is a "disturbance in the force:"

- Obtain (X₁, Y) for all n study subjects from electronic health records.
- ▶ Obtain X_2 for **only some** study subjects from survey, turns out to have a low response rate.

First (mystery) dataset

summary(dat1)

```
x1
                      x2
##
##
   Min.
         :0.000
                 Min.
                        :0
                             Min. :0.000
##
   1st Qu.:0.000 1st Qu.:0 1st Qu.:0.000
##
   Median: 0.000 Median: 0 Median: 0.000
   Mean :0.406 Mean
                             Mean :0.353
##
                       :0
   3rd Qu.:1.000 3rd Qu.:1
                             3rd Qu.:1.000
##
##
   Max. :1.000
                 Max. :1
                             Max. :1.000
                       :603
##
                 NA's
```

First dataset, continued

```
table(dat1, exclude=NULL)
## , , y = 0
##
##
   x2
## x1
      0
          1 <NA>
    0 141 27 244
##
##
    1 32 65 138
##
## , , y = 1
##
##
     x2
## x1
        0
          1 <NA>
##
    0 53
           15 114
##
    1
       13
           51 107
```

Second (mystery) dataset

```
table(dat2, exclude=NULL)
## , , y = 0
##
##
     x2
## x1
        0
          1 <NA>
    0 188 28 196
##
   1 43 84 108
##
##
## , , y = 1
##
##
     x2
## x1
        0
          1 <NA>
```

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0 75 29

5 32 134

1

##

##

Third (mystery) dataset

```
table(dat3, exclude=NULL)
## , , y = 0
##
##
   x2
## x1
      0
          1 <NA>
    0 351 61
##
## 1 81 154 0
##
## , , y = 1
##
##
     x2
## x1
        0
          1 <NA>
##
    0 136 13 33
    1 30 32 109
##
```

Answer by package - JAGS

```
genmod.string <- "model{</pre>
  ### prior distribution
  alpha0 \sim dnorm(0, 0.1)
  alpha1 \sim dnorm(0, 0.1)
   beta0 ~ dnorm(0, 0.1)
   beta1 ~ dnorm(0, 0.1)
   beta2 ~ dnorm(0, 0.1)
  ### statistical model
  for (i in 1:n) {
    x2[i] ~ dbern(pr.x2[i])
    logit(pr.x2[i]) <- alpha0 + alpha1*x1[i]</pre>
    y[i] ~ dbern(pr.y[i])
    logit(pr.y[i]) <- beta0 + beta1*x1[i] + beta2*x2[i]</pre>
711
```

Pause to comment on this prior and stat model specification

JAGS, continued

```
### generative model, data go in
mod <- jags.model(textConnection(genmod.string),</pre>
         data=list(x1=dat1$x1, x2=dat1$x2, y=dat1$y,
         n.chains=4)
update(mod, 2000) # burn-in
### MC output comes out
opt1.JAGS <- coda.samples(mod, n.iter=10000,
  variable.names=c("alpha0", "alpha1", "beta0", "beta1",
    "beta2", "x2[7]", "x2[8]"))
```

JAGS, continued

##

summary(opt1.JAGS)

```
##
## Iterations = 3001:13000
## Thinning interval = 1
## Number of chains = 4
## Sample size per chain = 10000
##
## 1. Empirical mean and standard deviation for each variable,
     plus standard error of the mean:
##
##
##
                  SD Naive SE Time-series SE
           Mean
## alpha0 -1.519 0.170 0.000851 0.00300
## alpha1 2.477 0.244 0.001222
                                   0.00434
## beta0 -0.918 0.104 0.000518 0.00130
## beta1 0.227 0.195 0.000976
                                 0.00322
## beta2 0.508 0.256 0.001279 0.00509
## x2[7] 0.231 0.421 0.002106 0.00218
## x2[8] 0.000 0.000 0.000000
                                    0.00000
##
## 2. Quantiles for each variable:
```

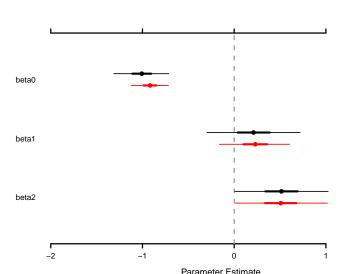
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And for comparison: complete-case analysis

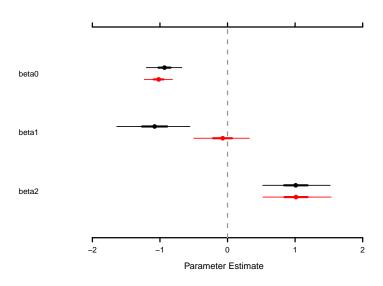
```
cmplt <- !is.na(dat1$x2)</pre>
cmplt[1:8]
## [1]
       TRUE FALSE FALSE FALSE FALSE FALSE
                                                 TRUE
mod <- jags.model(textConnection(genmod.string),</pre>
  data=list(x1=dat1$x1[cmplt],
             x2=dat1$x2[cmplt],
             y=dat1$y[cmplt],
opt1.cc.JAGS <- coda.samples(mod,</pre>
                  variable.names=c("beta0","beta1","beta2"),
                  n.iter=10000)
```

Comparison: Dataset 1, complete-case versus latent

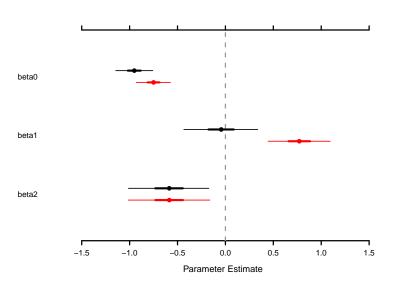
```
MCMCplot(opt1.cc.JAGS, opt1.JAGS,
    params=c("beta0","beta1","beta2"))
```



Same comparison, but for Dataset 2



Same comparison, but for Dataset 3



So what have we actually done here? (thinking space 1)

Let's think harder about how missing values became that way

Let R by a binary indicator, taking the value 1 if X_2 is observed, 0 if it's missing.

Need to think about the distribution of (X_1, X_2, Y, R) .

And concede that in fact for a given patient we will observe an event which has one of these two forms:

- $(X_1 = x_1, X_2 = x_2, Y = y, R = 1)$
- $(X_1 = x_1, Y = y, R = 0)$

Aside to think about: Sometimes this would be written as (X_1, Y, R, X_2R) are the observable variables.

In generality, think of this generative model

$$f(\alpha, \beta, x_1, x_2, y, r) = f(\alpha, \beta) f(x_1) f(x_2 | x_1, \alpha) \times f(y | x_1, x_2, \beta) f(r | x_1, x_2, y).$$

- ▶ Have made *conditional independence* assumptions here.
- ▶ With terms in red, think if our answer depends on them, then we will have to know their forms.

Applying Bayes theorem to this generative model gets us to

$$f\left(\alpha,\beta,x_2^{(mis)}|x_1,x_2^{(obs)},y,r\right) \propto f(\alpha,\beta)f(x_2|x_1)f(y|x_1,x_2) \times f(r|x_1,x_2,y)$$

(and again, think hard about the meaning of \propto here) So we have a hall-pass to stick with the analysis above so long as. . .

Ignorable missingess

In words, chance of missingess

on the underlying value that may/may not be obscured. Two related things to ponder. In situations where we *aren't* comfortable making this assumption:

- Could we include a further unknown parameter (say λ) in the generative model and keep/augment the $f(r|x_1, x_2, y, \lambda)$ term?
- Can the data empirically provide evidence for/against the assumption?

Now for a grand reveal concerning the three mystery datasets

$$logit{Pr(Y = 1|X_1, X_2)} = + X_1$$

Dataset 1

$$Pr(R=1|X_1,X_2,Y) =$$

Dataset 2

$$Pr(R=1|X_1,X_2,Y) =$$

Dataset 3

$$Pr(R = 1|X_1, X_2, Y) =$$

And then a final thought to come back to

If we don't feel comfortable assuming ignorable missingness, why not just work with

$$f(\alpha,\beta,\lambda)f(x_1)f(x_2|x_1,\alpha)f(y|x_1,x_2,\beta)f(r|x_1,x_2,y,\lambda)$$

Thought, continued