Indexing

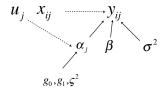
ESS 575 Models for Ecological Data

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An essential coding trick: Indexing groups



$$\begin{split} & \left[\boldsymbol{\alpha}, \boldsymbol{\beta},, \boldsymbol{\sigma}^2, \mathbf{g}, \boldsymbol{\varsigma}^2, | \mathbf{y} \right] \propto \prod_{i=1}^{n_j} \prod_{j=1}^{J} \operatorname{normal} \left(y_{ij} | \boldsymbol{\alpha}_j + \boldsymbol{\beta} x_{ij}, \boldsymbol{\sigma}^2 \right) \\ & \times \operatorname{normal} \left(\boldsymbol{\alpha}_j | g_0 + g_1 u_j, \boldsymbol{\varsigma}^2 \right) \\ & \times \operatorname{normal} \left(\boldsymbol{\beta} | 0,.001 \right) \operatorname{normal} \left(g_0 | 0,1000 \right) \times \operatorname{normal} \left(g_1 | 0,1000 \right) \\ & \times \operatorname{inverse gamma} \left(\boldsymbol{\sigma}^2 |.001,.001 \right) \operatorname{uniform} \left(\boldsymbol{\varsigma} | 0,200 \right) \end{split}$$

Indexing groups

> u

8 108 4.543959 38.93163

[109,] 8 109 1.287844 34.65796 [110,] 8 110 6.642313 40.62259 [111,] 8 111 7.404183 40.46518 [112,] 8 112 8.252571 41.47995 [113,] 8 113 9.558780 46.14771

Γ108,

Indexing groups

```
model{
beta \sim dnorm(0,.0001)
sigma \sim dunif(0.50)
tau.p <- 1/sigma^2
q0 \sim dnorm(0,.0001)
g1 \sim dnorm(0,.0001)
varsigma \sim dunif(0,50)
tau.g <- 1/varsigma^2
for (i in 1:length(y)){
 mu[i] <- alpha[group[i]]+ beta*x[i]</pre>
  y[i] ~ dnorm(mu[i],tau.p)
  for(j in 1:n.group){
  mu.g[j] <- g0 + g1*u[j]
  alpha[j]~dnorm(mu.g[j],tau.g)
```

Creating an index for a group

Assume you have a data frame (or tibble) called y with a column for sites containing a character code describing each site, i.e. y\$site. There are multiple observations within each site. How would you create a sequential index such that each site has its own, unique integer value corresponding to the site code, something like y\$site.index?¹

```
library(dplyr) # or library(tidyverse)
y = y %>% mutate(site.index = as.integer(as.factor(site)))
or, using base R
    y$site.index = as.integer(as.factor(site))
```

 $^{^1}$ The tidyverse packages, particularly dplyr are spectacular. $2 + \sqrt{2} + \sqrt{2} = \sqrt{2}$

Creating nested indices (2)

```
##Make index for site (each unique)
dat = dat %>% mu-
tate(Site.index = as.numeric(as.factor(Site)))
##Make index for strata
dat = dat %>% mu-
tate(Stratum.index = as.numeric(as.factor(Stratum)))
##Make index for sites within strata
dat = dat %>% group_by(Stratum) %>%
mutate(Site.within.strat.index = 1:n()) %>%
ungroup
```

| > | aat | | | | | |
|---|-------------|-------------|-------------|-------------|---------------|-----------------|
| # | A tibble: 6 | x 6 | | | | |
| | Stratum | Site | Count | Site.index | Stratum.index | Site.within.str |
| | <fct></fct> | <fct></fct> | <int></int> | <dbl></dbl> | <dbl></dbl> | <int></int> |
| 1 | Poudre | Beave | 44 | 1 | 2 | 1 |
| 2 | Poudre | LoneP | 23 | 4 | 2 | 2 |
| 3 | Poudre | Butes | 67 | 2 | 2 | 3 |
| 4 | BigThompson | HighB | 204 | 3 | 1 | 1 |
| 5 | BigThompson | Willo | 23 | 6 | 1 | 2 |
| 6 | BigThompson | Sheep | 123 | 5 | 1 | 3 |
| | | | | | | |

²see code MakeIndex.R

In House