

# IDS 702: MODULE 4.3

## MULTILEVEL/HIERARCHICAL LINEAR MODELS (ILLUSTRATION I)

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# THE RADON ANALYSIS

There are 919 total observations in the data. The data is in the file `Radon.txt` on Sakai.

Variable	Description
radon	radon levels for each house
log_radon	log(radon)
state	state
floor	lowest living area of each house: 0 for basement, 1 for first floor
countyname	county names
countyID	ID for the county names (1-85)
fips	state + county fips code
uranium	county-level soil uranium
log_uranium	log(uranium)

# THE RADON ANALYSIS

```
Radon <- read.csv("data/Radon.txt", header = T, sep = ",")
Radon$floor <- factor(Radon$floor, levels = c(0, 1), labels = c("Basement", "First Floor"))
str(Radon)
```

```
## 'data.frame':    919 obs. of  9 variables:
## $ radon      : num  2.2 2.2 2.9 1 3.1 2.5 1.5 1 0.7 1.2 ...
## $ state      : Factor w/ 1 level "MN": 1 1 1 1 1 1 1 1 1 1 ...
## $ log_radon  : num  0.788 0.788 1.065 0 1.131 ...
## $ floor      : Factor w/ 2 levels "Basement","First Floor": 2 1 1 1 1 1 1 1 1 1 ...
## $ countyname : Factor w/ 85 levels "AITKIN","ANOKA",...: 1 1 1 1 2 2 2 2 2 2 ...
## $ countyID   : int   1 1 1 1 2 2 2 2 2 2 ...
## $ fips       : int  27001 27001 27001 27001 27003 27003 27003 27003 27003 27003 ...
## $ uranium    : num  0.502 0.502 0.502 0.502 0.429 ...
## $ log_uranium: num  -0.689 -0.689 -0.689 -0.689 -0.847 ...
```

```
head(Radon)
```

```
##   radon state log_radon      floor countyname countyID  fips  uranium
## 1   2.2   MN 0.7884574 First Floor   AITKIN         1 27001 0.502054
## 2   2.2   MN 0.7884574   Basement   AITKIN         1 27001 0.502054
## 3   2.9   MN 1.0647107   Basement   AITKIN         1 27001 0.502054
## 4   1.0   MN 0.0000000   Basement   AITKIN         1 27001 0.502054
## 5   3.1   MN 1.1314021   Basement   ANOKA          2 27003 0.428565
## 6   2.5   MN 0.9162907   Basement   ANOKA          2 27003 0.428565
##   log_uranium
## 1 -0.6890476
## 2 -0.6890476
## 3 -0.6890476
## 4 -0.6890476
## 5 -0.8473129
## 6 -0.8473129
```

# THE RADON ANALYSIS

`table(Radon$countyname)` *#we don't have enough data in some counties, so we should look to borrow information across counties.*

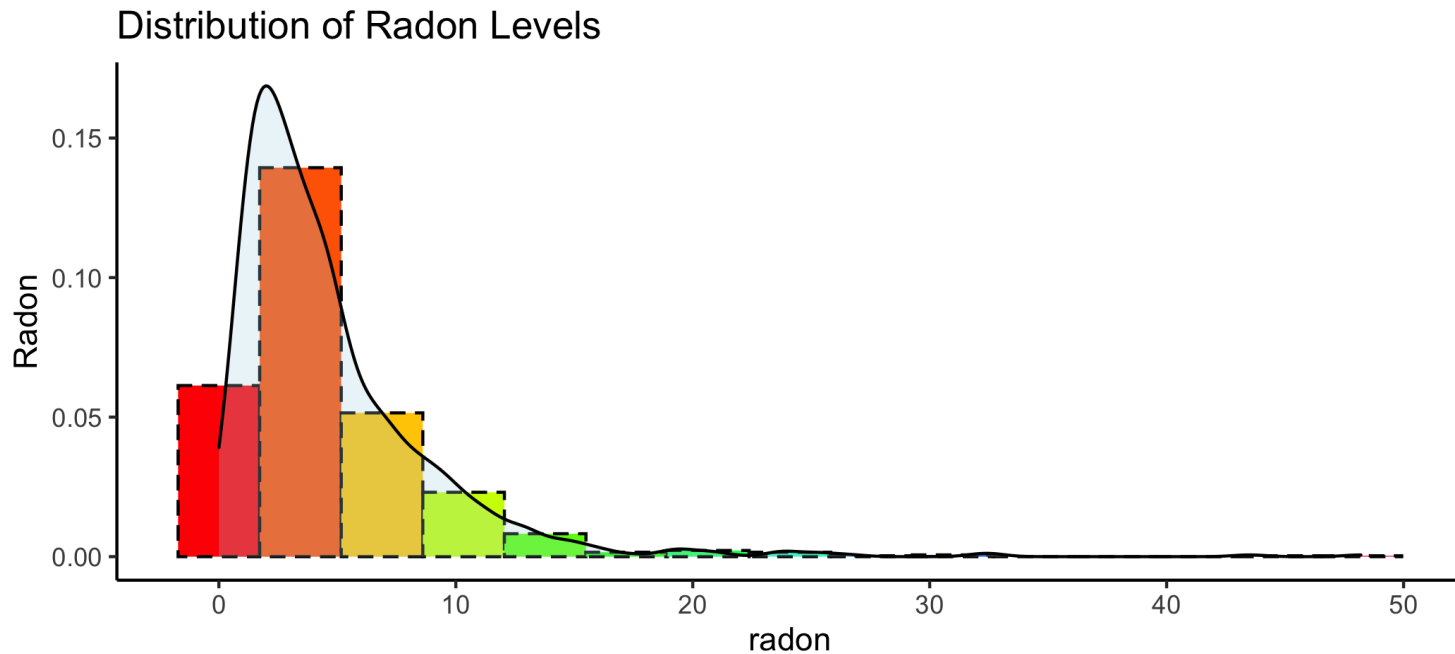
```
##
##      AITKIN          ANOKA          BECKER          BELTRAMI
##      4              52             3              7
##      BENTON        BIG STONE       BLUE EARTH      BROWN
##      4              3              14             4
##      CARLTON       CARVER          CASS           CHIPPEWA
##      10            6              5              4
##      CHISAGO       CLAY           CLEARWATER      COOK
##      6              14            4              2
##      COTTONWOOD    CROW WING      DAKOTA         DODGE
##      4              12            63             3
##      DOUGLAS       FARIBAULT      FILLMORE       FREEBORN
##      9              6              2              9
##      GOODHUE       HENNEPIN      HOUSTON        HUBBARD
##      14            105           6              5
##      ISANTI        ITASCA        JACKSON        KANABEC
##      3              11            5              4
##      KANDIYOHI     KITTSON      KOOCHICHING    LAC QUI PARLE
##      4              3              7              2
##      LAKE LAKE OF THE WOODS      LE SUEUR       LINCOLN
##      9              4              5              4
##      LYON          MAHNOMEN      MARSHALL       MARTIN
##      8              1              9              7
##      MCLEOD        MEEKER        MILLE LACS     MORRISON
##      13            5              2              9
##      MOWER         MURRAY        NICOLLET       NOBLES
##      13            1              4              3
##      NORMAN        OLMSTED      OTTER TAIL     PENNINGTON
##      3              23            8              3
##      PINE          PIPESTONE     POLK           POPE
##      6              4              4              2
##      RAMSEY        REDWOOD      RENVILLE       RICE
##      32            5              3              11
##      ROCK          ROSEAU       SCOTT          SHERBURNE
##      2              14            13             8
##      SIBLEY        ST LOUIS     STEARNS        STEELE
##      4              116           25             10
##      STEVENS       SWIFT        TODD           TRAVERSE
##      2              4              3              4
##      WABASHA       WADENA      WASECA         WASHINGTON
##      7              5              4              46
##      WATONWAN     WILKIN      WINONA         WRIGHT
##      3              1              13             13
##      YELLOW MEDICINE
```



# THE RADON ANALYSIS

The raw radon levels can only take on positive values.

```
ggplot(Radon,aes(radon)) +  
  geom_histogram(aes(y=..density..),color="black",linetype="dashed",  
                fill=rainbow(15),bins=15) + theme(legend.position="none") +  
  geom_density(alpha=.25, fill="lightblue") + scale_fill_brewer(palette="Blues") +  
  labs(title="Distribution of Radon Levels",y="Radon") + theme_classic()
```

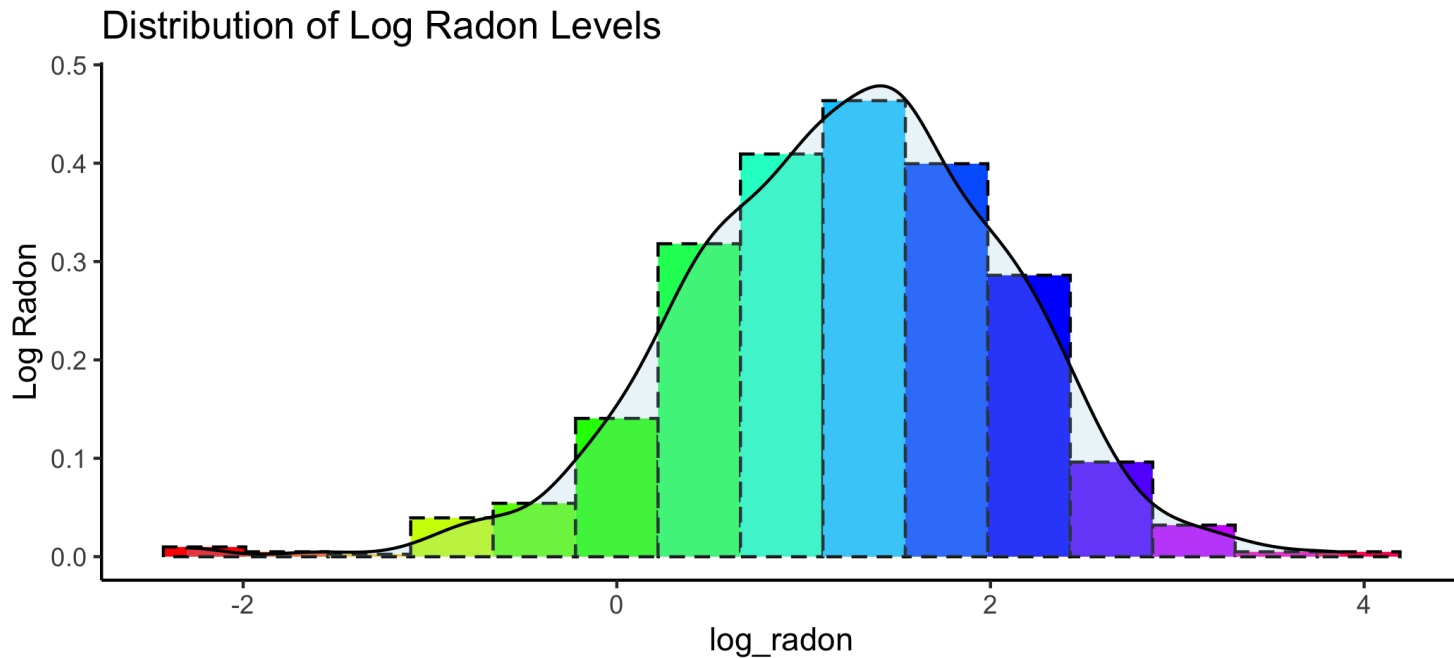


Obviously very skewed.

# THE RADON ANALYSIS

Let's look at `log_radon` instead.

```
ggplot(Radon, aes(log_radon)) +  
  geom_histogram(aes(y=..density..), color="black", linetype="dashed",  
                fill=rainbow(15), bins=15) + theme(legend.position="none") +  
  geom_density(alpha=.25, fill="lightblue") + scale_fill_brewer(palette="Blues") +  
  labs(title="Distribution of Log Radon Levels", y="Log Radon") + theme_classic()
```

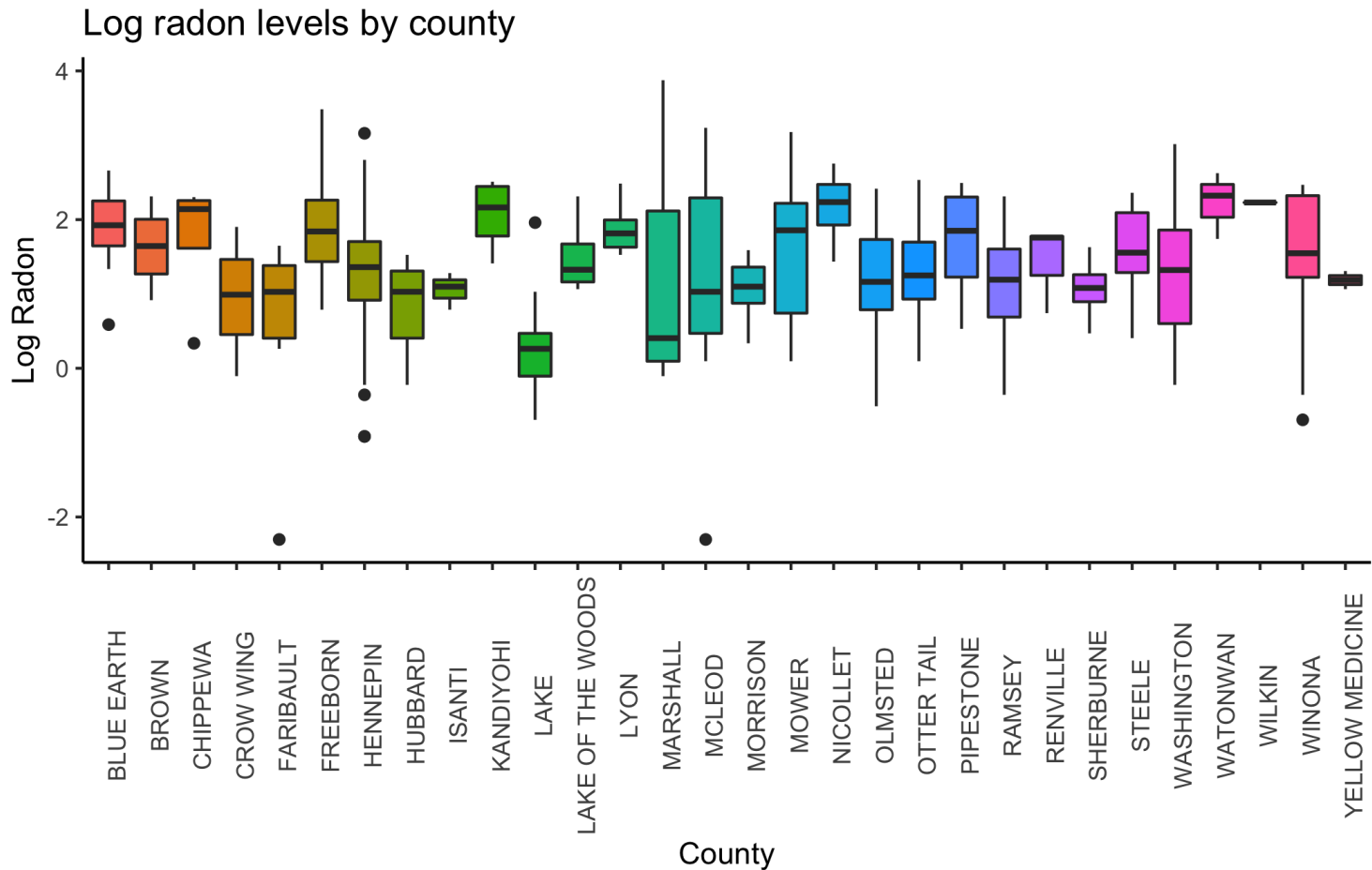


# THE RADON ANALYSIS

Are there any variations of radon levels by county? There are too many counties, so, let's do it for a random sample of counties.

```
set.seed(1000)
sample_county <- sample(unique(Radon$countyname),25,replace=F)
ggplot(Radon[is.element(Radon$countyname,sample_county),],
       aes(x=countyname, y=log_radon, fill=countyname)) +
  geom_boxplot() +
  labs(title="Log radon levels by county",
       x="County",y="Log Radon") + theme_classic() +
  theme(legend.position="none",axis.text.x = element_text(angle = 90))
```

# THE RADON ANALYSIS



Looks like the levels vary by county. However, there are many counties with very little data.

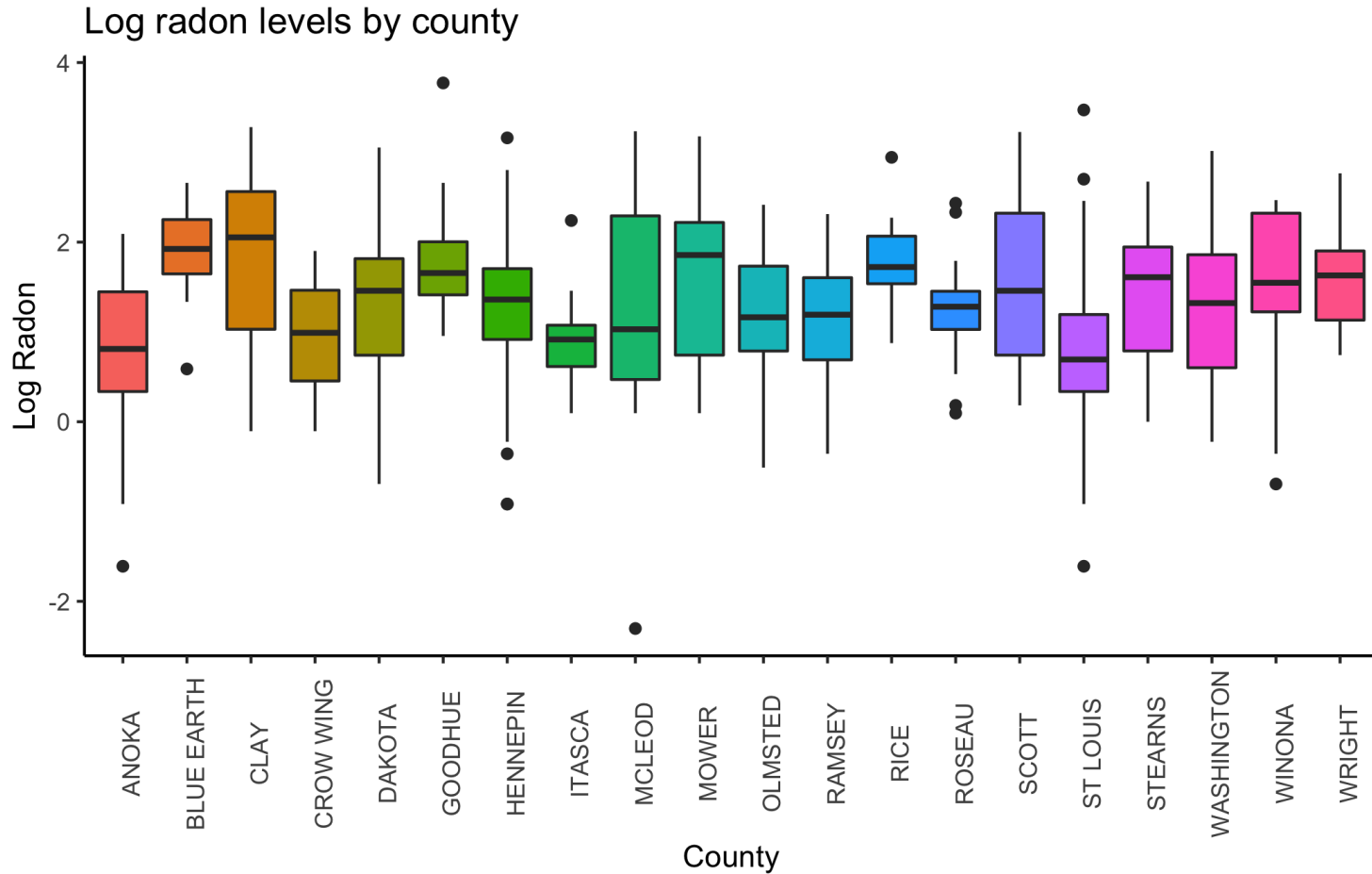


# THE RADON ANALYSIS

Let's focus on counties with at least 11 houses.

```
sample_county <- which(table(Radon$countyID) > 10)
ggplot(Radon[is.element(Radon$countyID,sample_county),],
       aes(x=countyname, y=log_radon, fill=countyname)) +
  geom_boxplot() +
  labs(title="Log radon levels by county",
       x="County",y="Log Radon") + theme_classic() +
  theme(legend.position="none",axis.text.x = element_text(angle = 90))
```

# THE RADON ANALYSIS

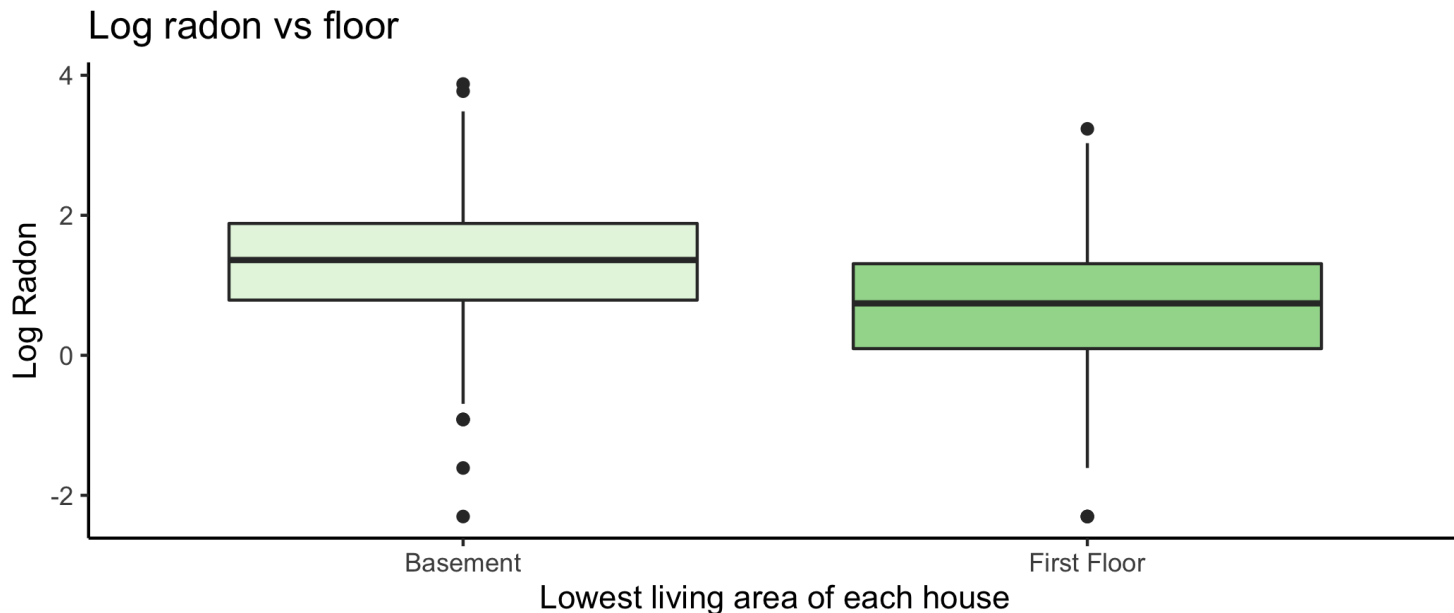


What can you conclude from this plot?

# THE RADON ANALYSIS

Next, the relationship with `floor`, the only individual-level (different observation for each house) variable we have.

```
ggplot(Radon, aes(x=floor, y=log_radon, fill=floor)) +  
  geom_boxplot() + scale_fill_brewer(palette="Greens") +  
  labs(title="Log radon vs floor", x="Lowest living area of each house", y="Log Radon") +  
  theme_classic() + theme(legend.position="none")
```

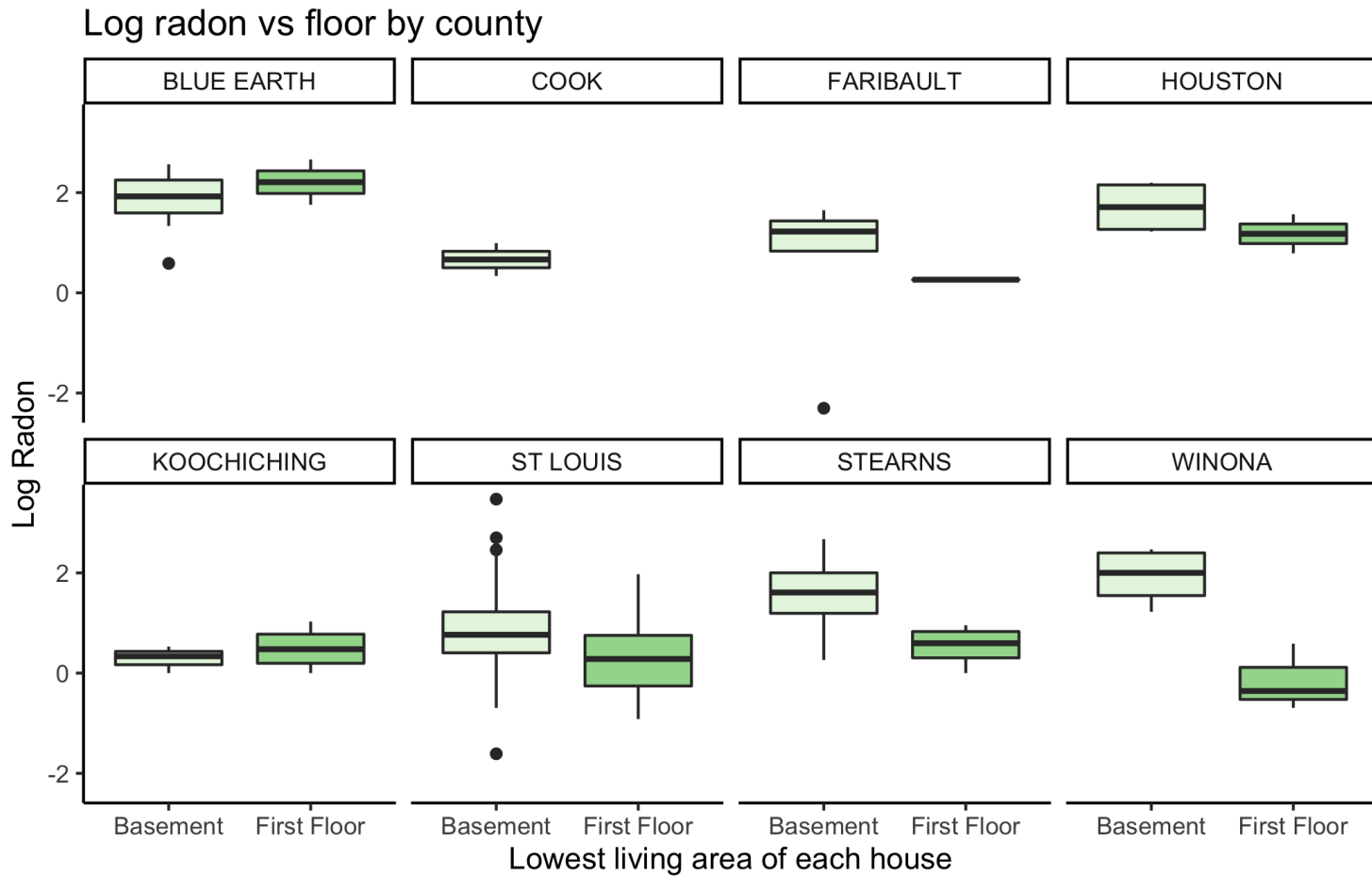


# THE RADON ANALYSIS

Let's look at the same relationship for a random sample of counties.

```
sample_county <- sample(unique(Radon$countyname),8,replace=F)
ggplot(Radon[is.element(Radon$countyname,sample_county),],
       aes(x=floor, y=log_radon, fill=floor)) +
  geom_boxplot() +
  scale_fill_brewer(palette="Greens") +
  labs(title="Log radon vs floor by county",
       x="Lowest living area of each house",y="Log Radon") +
  theme_classic() + theme(legend.position="none") +
  facet_wrap( ~ countyname,ncol=4)
```

# THE RADON ANALYSIS



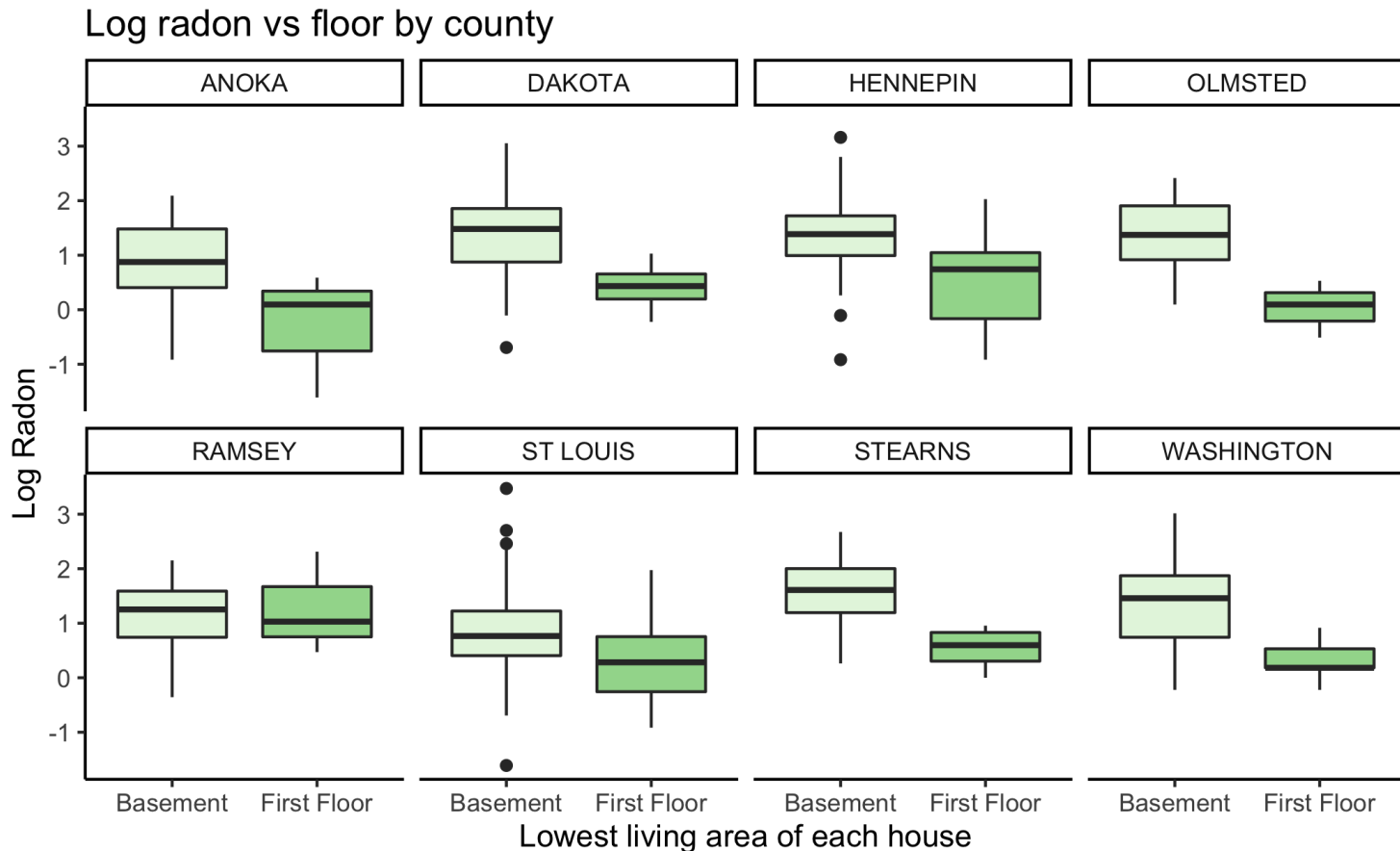
Again, not enough data for some counties.

# THE RADON ANALYSIS

Let's focus on counties with at least 16 houses.

```
sample_county <- which(table(Radon$countyID) > 15)
ggplot(Radon[is.element(Radon$countyID,sample_county),],
       aes(x=floor, y=log_radon, fill=floor)) +
  geom_boxplot() +
  scale_fill_brewer(palette="Greens") +
  labs(title="Log radon vs floor by county",
       x="Lowest living area of each house",y="Log Radon") +
  theme_classic() + theme(legend.position="none") +
  facet_wrap( ~ countyname,ncol=4)
```

# THE RADON ANALYSIS



Even though the overall direction is the same, it looks like the actual differences between floor = 0 and floor = 1 differs for some counties.

# THE RADON ANALYSIS

- Let's start by only focusing on `floor`.
- We will try a varying-slope, varying-intercept linear model.
- Let  $y_{ij}$  and  $x_{1ij}$  be the log radon level and indicator variable `floor` respectively for house  $i$  in county  $j$ .
- Mathematically, we have

$$\begin{aligned}y_{ij} &= (\beta_0 + \gamma_{0j}) + (\beta_1 + \gamma_{1j})x_{1ij} + \epsilon_{ij}; \quad i = 1, \dots, n_j; \quad j = 1, \dots, 85 \\ \epsilon_{ij} &\sim N(0, \sigma^2) \\ (\gamma_{0j}, \gamma_{1j}) &\sim N_2(\mathbf{0}, \Sigma).\end{aligned}$$

- Alternative representation:

$$\begin{aligned}\log(\text{radon}_{ij}) &= (\beta_0 + \gamma_{0j}) + (\beta_1 + \gamma_{1j}) \text{floor}_{ij} + \epsilon_{ij}; \quad i = 1, \dots, n_j; \quad j = 1, \dots, 85 \\ \epsilon_{ij} &\sim N(0, \sigma^2) \\ (\gamma_{0j}, \gamma_{1j}) &\sim N_2(\mathbf{0}, \Sigma).\end{aligned}$$



# THE RADON ANALYSIS

- We skipped this before but  $\Sigma$  actually takes the form

$$\Sigma = \begin{bmatrix} \tau_0^2 & \rho\tau_0\tau_1 \\ \rho\tau_0\tau_1 & \tau_1^2 \end{bmatrix}$$

where

- $\tau_0^2$  describes the across county variation attributed to the random/varying intercept,
- $\tau_1^2$  describes the across county variation attributed to the random/varying slope (that is, floor), and
- $\rho$  describes the correlation between  $\gamma_{0j}$  and  $\gamma_{1j}$ .

# THE RADON ANALYSIS

In R, we have

```
Model1 <- lmer(log_radon ~ floor + (floor | countyname), data = Radon)
summary(Model1)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: log_radon ~ floor + (floor | countyname)
## Data: Radon
##
## REML criterion at convergence: 2168.3
##
## Scaled residuals:
##   Min       1Q   Median       3Q      Max
## -4.4044 -0.6224  0.0138  0.6123  3.5682
##
## Random effects:
##   Groups      Name                Variance Std.Dev. Corr
##   countyname (Intercept)          0.1216   0.3487
##             floorFirst Floor  0.1181   0.3436  -0.34
## Residual                        0.5567   0.7462
## Number of obs: 919, groups:  countyname, 85
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)      1.46277   0.05387  27.155
## floorFirst Floor -0.68110   0.08758  -7.777
##
## Correlation of Fixed Effects:
##              (Intr)
## florFrstFlr -0.381
```

# INTERPRETATION OF FIXED EFFECTS

- Intuitively, we have an overall "average" regression line for all houses across all counties in Minnesota which has slope -0.68 and intercept 1.46.
- That is, the general estimated line for any of the houses in Minnesota is:

$$\log(\widehat{\text{radon}}_i) = 1.46 - 0.68 \times \text{floor}_i$$

- For any house in Minnesota with a basement as the lowest living area, the baseline radon level is  $e^{1.46} = 4.31$ .
- Then, for any house in Minnesota, having a first floor as the lowest living area, instead of a basement, reduces the radon level by a multiplicative effect of  $e^{-0.68} = 0.51$ , that is, about a 49% reduction.
- However, if the house is in Dakota county for example, we also need to add on the random intercepts and slopes for that county.

# INTERPRETATION OF FIXED EFFECTS

- For Dakota county, we have

```
(ranef(Model1)$countyname)["DAKOTA",]
```

```
##           (Intercept) floorFirst Floor  
## DAKOTA   -0.1099052      -0.08786805
```

so that the estimated regression line for Dakota county is actually

$$\log(\widehat{\text{radon}}_i) = (1.46 - 0.11) + (-0.68 - 0.09) \times \text{floor}_i = 1.35 - 0.77 \times \text{floor}_i$$

- Thus, for any house in Dakota county in Minnesota with a basement as the lowest living area, the baseline radon level is actually  $e^{1.35} = 3.86$ , which is **lower than the overall state wide average**.
- And for any house in Dakota county in Minnesota, having the first floor be the lowest living area then reduces the radon level by a multiplicative effect of  $e^{-0.77} = 0.46$ , that is about a 54% reduction, **more than the overall state wide effect**.

# THE RADON ANALYSIS

Again,

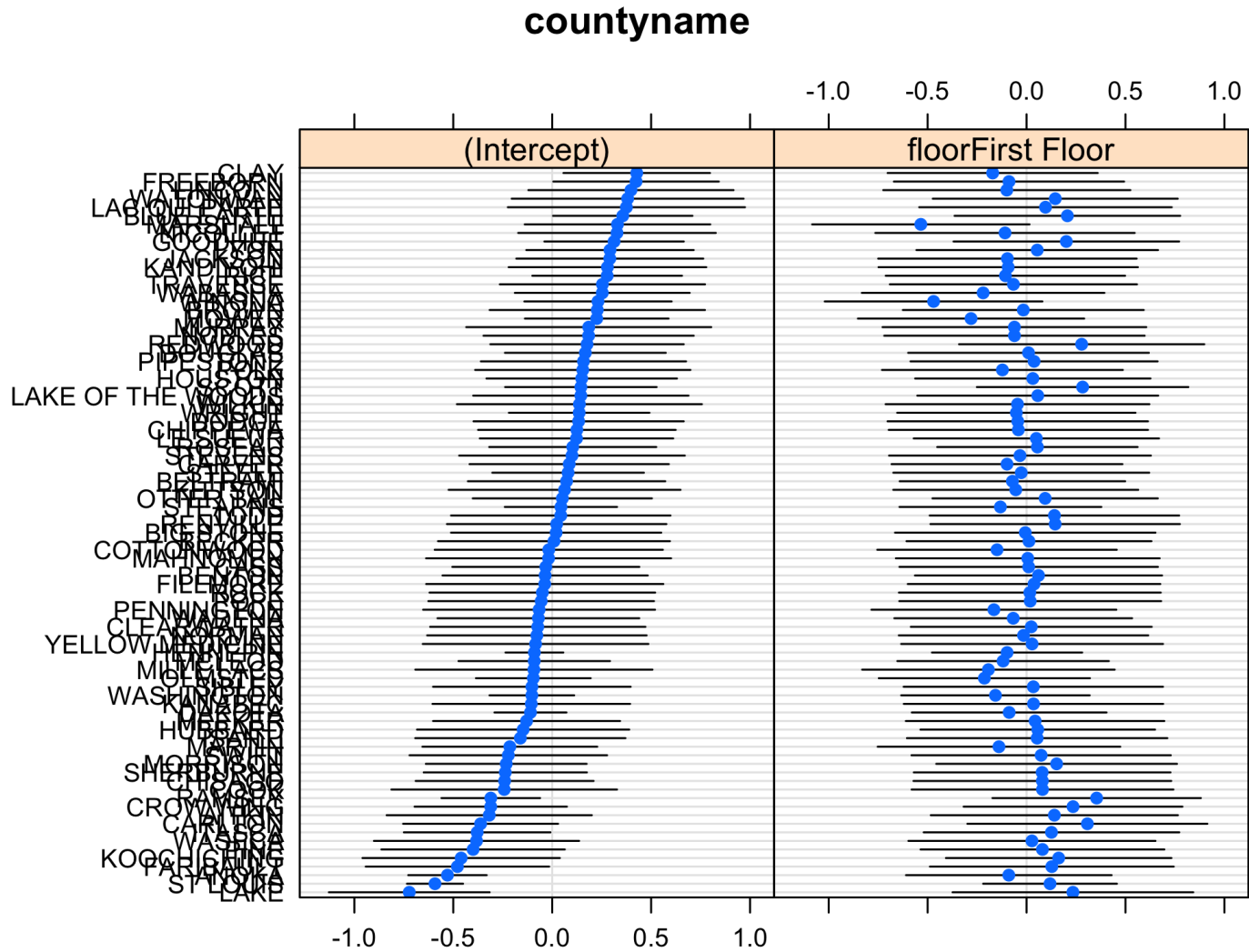
```
summary(Model1)
```

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: log_radon ~ floor + (floor | countyname)
##   Data: Radon
##
## REML criterion at convergence: 2168.3
##
## Scaled residuals:
##   Min       1Q   Median       3Q      Max
## -4.4044 -0.6224  0.0138  0.6123  3.5682
##
## Random effects:
##   Groups      Name                Variance Std.Dev. Corr
##   countyname (Intercept)          0.1216   0.3487
##             floorFirst Floor    0.1181   0.3436  -0.34
##   Residual                        0.5567   0.7462
## Number of obs: 919, groups:  countyname, 85
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)      1.46277   0.05387  27.155
## floorFirst Floor -0.68110   0.08758  -7.777
##
## Correlation of Fixed Effects:
##              (Intr)
## florFrstFlr -0.381
```

# INTERPRETATION OF RANDOM EFFECTS

- The estimated standard error  $\hat{\sigma} = 0.75$  describes the within-county or remaining unexplained variation.
- The estimated  $\hat{\tau}_0 = 0.35$  describes the across-county variation attributed to the random intercept.
- The estimated  $\hat{\tau}_1 = 0.34$  describes the across-county variation attributed to the random slope (the predictor, floor).
- Those two sources of county variation are actually quite similar.
- The estimated correlation between  $\gamma_{0j}$  and  $\gamma_{1j}$  is  $\hat{\rho} = -0.34$ .
- You can visualize the random effects by typing `dotplot(ranef(Model1, condVar=TRUE))$countname` in R.
- So many counties! So, you will need to zoom out on your computer.

# INTERPRETATION OF RANDOM EFFECTS



# WHAT'S NEXT?

MOVE ON TO THE READINGS FOR THE NEXT MODULE!