

Multiple Linear Regression: Categorical Predictors

Author: Nicholas G Reich, Jeff Goldsmith

*This material is part of the **statsTeachR** project*

*Made available under the Creative Commons Attribution-ShareAlike 3.0 Unported
License: http://creativecommons.org/licenses/by-sa/3.0/deed.en_US*

Multiple Linear Regression: recapping model definition

In matrix notation...

$$\vec{\mathbf{y}} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon}$$

where $E(\boldsymbol{\epsilon}) = 0$ and $\text{Cov}(\boldsymbol{\epsilon}) = \sigma^2 I$

In individual observation notation...

$$y_i = \beta_0 + \beta_1 x_{1,i} + \dots + \beta_p x_{p,i} + \epsilon_i$$

where $\epsilon_i \stackrel{iid}{\sim} (0, \sigma^2)$

Categorical predictors

- Assume X is a categorical / nominal / factor variable with k levels
- With only one categorical X , we have classic one-way ANOVA design
- Can't use a single predictor with levels $1, 2, \dots, K$ – this has the wrong interpretation
- Need to create *indicator* or *dummy* variables

Indicator variables

$$y_i = \beta_0 + \beta_1 x_i$$
$$\hookrightarrow y_i = \beta_0 + \beta_1 x_{1,i} + \beta_2 x_{2,i}$$

- Let x be a categorical variable with k levels (e.g. with $k = 3$ “low”, “med”, “hi”).
- Choose one group as the baseline (e.g. “low”).
- Create $(k - 1)$ binary terms to include in the model:

$$x_{1,i} = \mathbb{1}(x_i = \text{“med”})$$
$$x_{2,i} = \mathbb{1}(x_i = \text{“hi”})$$

$$\rightarrow \mathbb{1}(x_i = \text{“hi”}) = \begin{cases} 1 & \text{if } x_i = \text{“hi”} \\ 0 & \text{o.w.} \end{cases}$$

- For a model with no additional predictors, pose the model

$$y_i = \beta_0 + \beta_1 x_{1,i} + \dots + \beta_{k-1} x_{k-1,i} + \epsilon_i$$

and estimate parameters using least squares

- Note distinction between *predictors* and *terms*

Categorical predictor design matrix

Which of the following is a “correct” design matrix for a categorical predictor with 3 levels?

$$X_1 = \begin{matrix} & x_1 & x_2 & x_3 & x_4 \\ \begin{bmatrix} 1 & 1 & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & 1 & 0 & 0 \\ 1 & 0 & 1 & 0 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 1 & 0 \\ 1 & 0 & 0 & 1 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 0 & 1 \end{bmatrix} \end{matrix}$$

$$x_4 = 2 - x_3 - x_2 - x_1$$

$$\text{or } X_2 = \begin{bmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 1 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 1 & 0 \\ 1 & 0 & 1 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 1 \end{bmatrix}$$

$$\text{or } X_3 = \begin{bmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 0 & 1 & 0 \\ 0 & 0 & 1 \\ \vdots & \vdots & \vdots \\ 0 & 0 & 1 \end{bmatrix}$$

x_{1i}
 x_{2i}
 x_{3i}

ANOVA model interpretation

Using the model $y_i = \beta_0 + \beta_1 x_{1,i} + \dots + \beta_{k-1} x_{k-1,i} + \epsilon_i$, interpret

$$\beta_0 = E(y | \text{all } x\text{'s} = 0)$$

$$= E(y | x_1 = 0, x_2 = 0)$$

= expected/average y for baseline group

β_1 = difference in $E(y)$ between base line group and group 1.

$$\beta_0 + \beta_1 = E(y | x_1 = 1)$$

Equivalent model

x_1 x_2 x_3

Define the model $y_i = \beta_1 x_{i1} + \dots + \beta_k x_{i,k} + \epsilon_i$ where there are indicators for each possible group

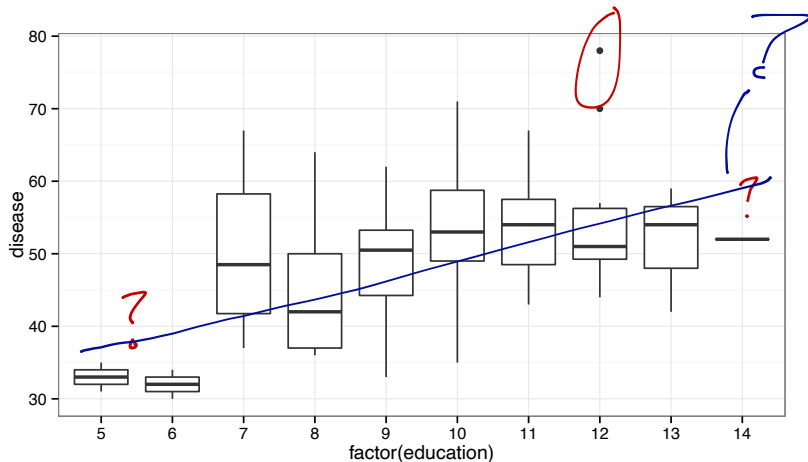
$$\beta_1 = E(y | x_1 = 1)$$

= expected value of y in group 1

$$\beta_2 = E(y | x_2 = 1)$$

Categorical predictor example: lung data

```
qplot(factor(education), disease, geom="boxplot", data=dat)
```



Categorical predictor example: lung data

$$\underline{dis_i} = \beta_0 + \beta_1 educ_{6,i} + \beta_2 educ_{7,i} + \cdots + \beta_{14} educ_{14,i}$$

```
mlr7 <- lm(disease ~ factor(education), data=dat)
summary(mlr7)$coef
```

##	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	33.00	4.913	6.7173	1.689e-09
## factor(education)6	-1.00	7.768	-0.1287	8.979e-01
## factor(education)7	17.33	6.017	2.8808	4.969e-03
## factor(education)8	11.18	5.329	2.0975	3.879e-02
## factor(education)9	15.50	5.353	2.8953	4.765e-03
## factor(education)10	20.38	5.188	3.9289	1.683e-04
## factor(education)11	20.53	5.382	3.8155	2.505e-04
## factor(education)12	22.20	5.601	3.9633	1.489e-04
## factor(education)13	18.67	6.948	2.6868	8.609e-03
## factor(education)14	19.00	9.825	1.9338	5.632e-02

$\beta_0 + \beta_4$

Categorical predictor releveling

$$\begin{aligned} X_1 &\leftarrow \text{cont} + \text{class}(X_2) \\ X_2 &\leftarrow \text{categ.} \\ \ln(y \sim X_1 + X_2) \end{aligned}$$

$$dis_i = \beta_0 + \beta_1 educ_{5,i} + \beta_2 educ_{6,i} + \beta_1 educ_{7,i} + \beta_2 educ_{9,i} + \dots + \beta_{14} educ_{14,i}$$

```
dat$educ_new <- relevel(factor(dat$education), ref="8")  
mlr8 <- lm(disease ~ educ_new, data=dat)  
summary(mlr8)$coef
```

	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	44.176	2.064	21.4059	7.303e-37
## educ_new5	-11.176	5.329	-2.0975	3.879e-02
## educ_new6	-12.176	6.361	-1.9143	5.880e-02
## educ_new7	6.157	4.041	1.5238	1.311e-01
## educ_new9	4.324	2.964	1.4588	1.482e-01
## educ_new10	9.208	2.654	3.4695	8.059e-04
## educ_new11	9.357	3.014	3.1042	2.559e-03
## educ_new12	11.024	3.391	3.2507	1.626e-03
## educ_new13	7.490	5.329	1.4057	1.633e-01
## educ_new14	7.824	8.756	0.8935	3.740e-01

Categorical predictor: no baseline group

$$dis_i = \beta_1 educ_{5,i} + \beta_2 educ_{6,i} + \cdots + \beta_{14} educ_{14,i}$$

+ 0

```
mlr9 <- lm(disease ~ factor(education) - 1, data=dat)
summary(mlr9)$coef
```

##	Estimate	Std. Error	t value	Pr(> t)
## factor(education)5	33.00	4.913	6.717	1.689e-09
## factor(education)6	32.00	6.017	5.318	7.716e-07
## factor(education)7	50.33	3.474	14.489	3.846e-25
## factor(education)8	44.18	2.064	21.406	7.303e-37
## factor(education)9	48.50	2.127	22.799	6.282e-39
## factor(education)10	53.38	1.669	31.991	1.359e-50
## factor(education)11	53.53	2.197	24.366	3.801e-41
## factor(education)12	55.20	2.691	20.514	1.713e-35
## factor(education)13	51.67	4.913	10.517	2.758e-17
## factor(education)14	52.00	8.509	6.111	2.561e-08

Creating categories using cut()

$$dis_i = \beta_1 educ_{low,i} + \beta_2 educ_{med,i} + \beta_3 educ_{hi,i}$$

continuous

3

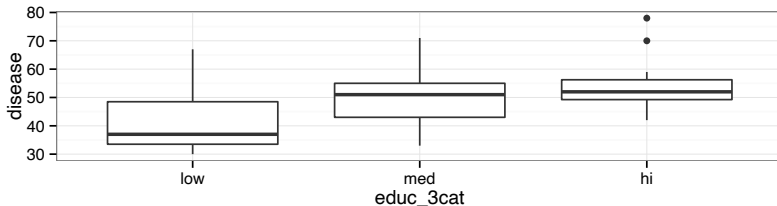
? cut

```
dat$educ_3cat <- cut(dat$education, breaks=3,  
                     labels=c("low", "med", "hi"))
```

```
mlr10 <- lm(disease ~ educ_3cat - 1, data=dat)  
coef(mlr10)
```

```
## educ_3catlow educ_3catmed educ_3cat  
##          42.27          50.24          54.21
```

```
qplot(educ_3cat, disease, geom="boxplot", data=dat)
```



Today's big ideas

- Multiple linear regression: categorical variables