Regression: dummy variables

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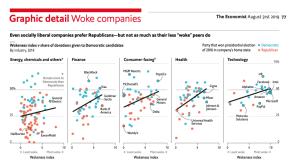
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Outline

Dummy variables for categorical covariates

Categorical predictors

- Assume X is a categorical / nominal / factor variable with k levels: e.g. 'Industry'.
- If you use a single predictor with continuous values of 1,2,..., K this assumes that a "one unit increase" has a clear meaning.
- You need to create indicator or dummy variables so that each level stands on its own and can be estimated separately.



Categorical predictors

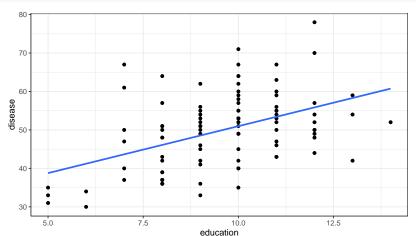
Important to distinguish between...

- "nominal" categorical variables: e.g. ones with no natural ordering, such as Industry, country, etc...
- "ordinal" categorical variables: e.g. ones with a natural ordering, such as education level, or age grouping.

Categorical predictor example: lung data

Education could plausibly be continuous (e.g. you could interpret a one-unit increase), but likely a linear assumption is not great. Thinking of education as a "factor" may be more practical.

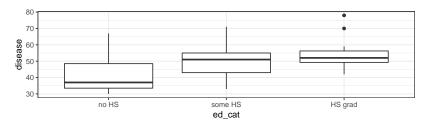
```
qplot(education, disease, data=dat) + geom_point() +
  geom_smooth(method="lm", se=FALSE)
```



Defining a categorical variable

We could define educational level relative to high-school (HS) achievement.

$$\mathsf{ed_cat}_i = egin{cases} \mathsf{no}\ \mathsf{HS}, & \mathsf{if}\ \mathsf{education}_i < 5 \\ \mathsf{some}\ \mathsf{HS}, & \mathsf{if}\ 5 \leq \mathsf{education}_i < 8 \\ \mathsf{HS}\ \mathsf{grad}, & \mathsf{if}\ 8 \leq \mathsf{education}_i \end{cases}$$



Indicator variables

- Let x be a categorical variable with k levels .
- Choose one group as the baseline (e.g. "no HS").
- Create (k-1) binary variables to encode the information about which group each observation belongs to.

```
dat$someHS <- as.numeric(dat$ed_cat=="some HS")</pre>
dat$HSgrad <- as.numeric(dat$ed_cat=="HS grad")</pre>
dat[8:13, c("disease", "education", "ed_cat", "someHS", "HSgrad")]
##
     disease education ed_cat someHS HSgrad
## 8
         58
                  10 some HS
        52
                  14 HS grad
## 9
## 10 57
                  12 HS grad
      4.3
                  11 some HS 1
## 11
## 12 48 8 some HS 1
## 13
        34
               6 no HS
```

Standard model interpretation

```
## note that R doesn't need the two dummy variables we created
## it creates them for us.
mod1 <- lm(disease ~ crowding + ed_cat, data=dat)</pre>
```

Interpret:

$$dis_i = \beta_0 + \beta_1 \cdot crowding_i + \beta_2 \cdot someHS_i + \beta_3 \cdot HSgrad_i + \epsilon_i$$
.

$$\beta_0 =$$

$$\beta_1 =$$

$$\beta_2 =$$

Categorical predictor example: lung data

```
coefs <- coef(mod1)</pre>
ggplot(dat, aes(x=crowding, y=disease, color=ed_cat, shape=ed_cat)) +
  geom_point() + scale_color_manual(values=c("#1b9e77", "#d95f02", "#7570b3"))+
  geom_abline(intercept = coefs[1], slope = coefs[2], color="#1b9e77")+
  geom_abline(intercept = coefs[1]+coefs[3], slope = coefs[2], color="#d95f02")
  geom_abline(intercept = coefs[1]+coefs[4], slope = coefs[2], color="#7570b3")
  70
  60
                                                                   ed cat
disease
                                                                      no HS
                                                                      some HS

    HS grad

  40
  30
                  20
```

Categorical predictor example: lung data

$$\mathsf{dis}_i = \beta_0 + \beta_1 \cdot \mathsf{crowding}_i + \beta_2 \cdot \mathsf{someHS}_i + \beta_3 \cdot \mathsf{HSgrad}_i + \epsilon_i$$

Categorical predictor example: interaction

$$\widehat{\mathsf{dis}}_i = \beta_0 + \beta_1 \cdot c_i + \beta_2 \cdot \mathsf{someHS}_i + \beta_3 \cdot \mathsf{HSgrad}_i + \beta_4 \cdot c_i \cdot \mathsf{someHS}_i + \beta_5 \cdot c_i \cdot \mathsf{HSgrad}_i$$

In terms of the betas, what are the equations of the regression lines for the 'no HS', 'some HS' and 'HS grad' categories?