

Multiple Linear Regression: Collinearity and Categories

a **statsTeachR** resource

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Recap: Least squares for MLR

As in simple linear regression, we want to find the β that minimizes the residual sum of squares.

$$RSS(\beta) = \sum_i \epsilon_i^2 = \epsilon^T \epsilon$$

After taking the derivative, setting equal to zero, we obtain:

$$\hat{\beta} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

Hat matrix

$$\mathbf{H} = \mathbf{X}(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T$$

Some properties of the hat matrix:

- It is a projection matrix: $\mathbf{H}\mathbf{H} = \mathbf{H}$
- It is symmetric: $\mathbf{H}^T = \mathbf{H}$
- The residuals are $\hat{\epsilon} = (\mathbf{I} - \mathbf{H})\mathbf{y}$
- The inner product of $(\mathbf{I} - \mathbf{H})\mathbf{y}$ and $\mathbf{H}\mathbf{y}$ is zero (predicted values and residuals are uncorrelated).

Projection space interpretation

The hat matrix projects \mathbf{y} onto the column space of \mathbf{X} .

Alternatively, minimizing the $RSS(\beta)$ is equivalent to minimizing the Euclidean distance between \mathbf{y} and the column space of \mathbf{X} .

Lung Data Example (con't from previous clas)

```
mlr2 <- lm(disease ~ crowding + education + airqual,  
           data=dat, x=TRUE, y=TRUE)  
coef(mlr2)  
  
## (Intercept)      crowding      education      airqual  
##      -7.7505         1.3128         1.4377         0.2881  
  
X = mlr2$x  
y = mlr2$y  
(betaHat = solve( t(X) %*% X) %*% t(X) %*% y )  
  
##                [,1]  
## (Intercept) -7.7505  
## crowding      1.3128  
## education     1.4377  
## airqual       0.2881
```

Key points so far

- Our model is $\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\epsilon}$ with $\boldsymbol{\epsilon} \sim (0, \sigma^2 \mathbf{I})$
- The design matrix \mathbf{X} contains the terms included in the model
- We have least squares solutions under some conditions

Least squares estimates

$$\hat{\beta} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

A condition on $(\mathbf{X}^T \mathbf{X})$

- If $(\mathbf{X}^T \mathbf{X})$ is singular, there are infinitely many least squares solutions, making $\hat{\beta}$ non-identifiable (can't choose between different solutions)

Non-identifiability

- Can happen if \mathbf{X} is not of full rank, i.e. the columns of \mathbf{X} are linearly dependent (for example, including weight in Kg and lb as predictors)
- Can happen if there are fewer data points than terms in \mathbf{X} : $n < p$ (having 100 predictors and only 50 observations)
- Generally, the $p \times p$ matrix $(\mathbf{X}^T \mathbf{X})$ is invertible if and only if it has rank p .

Infinite solutions

Suppose I fit a model $y_i = \beta_0 + \beta_1 x_{i1} + \epsilon_i$.

- I have estimates $\hat{\beta}_0 = 1, \hat{\beta}_1 = 2$
- I put in a new variable $x_2 = x_1$
- My new model is $y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \epsilon_i$
- Possible least squares estimates that are equivalent to my first model:
 - ▶ $\hat{\beta}_0 = 1, \hat{\beta}_1 = 2, \hat{\beta}_2 = 0$
 - ▶ $\hat{\beta}_0 = 1, \hat{\beta}_1 = 0, \hat{\beta}_2 = 2$
 - ▶ $\hat{\beta}_0 = 1, \hat{\beta}_1 = 1002, \hat{\beta}_2 = -1000$
 - ▶ ...

Non-identifiability

- Often due to data coding errors (variable duplication, scale changes)
- Pretty easy to detect and resolve
- Can be addressed using *penalties* (might come up much later)
- A bigger problem is near-unidentifiability (collinearity)

Causes of collinearity

- Arises when variables are highly correlated, but not exact duplicates
- Commonly arises in data (perfect correlation is usually there by mistake)
- Might exist between several variables, i.e. a linear combination of several variables exists in the data
- A variety of tools exist (correlation analyses, multiple R^2 , eigen decompositions)

Effects of collinearity

Suppose I fit a model $y_i = \beta_0 + \beta_1 x_{i1} + \epsilon_i$.

- I have estimates $\hat{\beta}_0 = 1, \hat{\beta}_1 = 2$
- I put in a new variable $x_2 = x_1 + \text{error}$, where *error* is pretty small
- My new model is $y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \epsilon_i$
- Possible least squares estimates that are nearly equivalent to my first model:
 - ▶ $\hat{\beta}_0 = 1, \hat{\beta}_1 = 2, \hat{\beta}_2 = 0$
 - ▶ $\hat{\beta}_0 = 1, \hat{\beta}_1 = 0, \hat{\beta}_2 = 2$
 - ▶ $\hat{\beta}_0 = 1, \hat{\beta}_1 = 1002, \hat{\beta}_2 = -1000$
 - ▶ ...
- A unique solution exists, but it is hard to find

Effects of collinearity

- Collinearity results in a “flat” RSS
- Makes identifying a unique solution difficult
- Dramatically inflates the variance of LSEs

Non-identifiability example: lung data

```
mlr3 <- lm(disease ~ airqual, data=dat)
coef(mlr3)
```

```
## (Intercept)      airqual
##      35.4445      0.3537
```

```
dat$x2 <- dat$airqual/100
mlr4 <- lm(disease ~ airqual + x2, data=dat, x=TRUE)
coef(mlr4)
```

```
## (Intercept)      airqual      x2
##      35.4445      0.3537      NA
```

```
X = mlr4$x
solve( t(X) %*% X)
```

```
## Error: system is computationally singular:
reciprocal condition number = 3.57906e-20
```

Collinearity example: lung data

```
dat$crowd2 <- dat$crowding + rnorm(nrow(dat), sd=.1)
mlr5 <- lm(disease ~ crowding, data=dat)
summary(mlr5)$coef
```

##	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	12.992	3.4750	3.739	3.130e-04
## crowding	1.509	0.1394	10.826	2.232e-18

```
mlr6 <- lm(disease ~ crowding + crowd2, data=dat)
summary(mlr6)$coef
```

##	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	12.978	3.489	3.7201	0.0003354
## crowding	5.067	7.106	0.7131	0.4775184
## crowd2	-3.558	7.103	-0.5009	0.6176152

Some take away messages

- Collinearity can (and does) happen, so be careful
- Often contributes to the problem of variable selection, which we'll touch on later

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

- Choose one group as the baseline
- Create 0/1 terms to include in the model x_1, x_2, \dots, x_{k-1}
- Pose the model

$$y_i = \beta_0 + \beta_1 x_{i1} + \dots + \beta_{k-1} x_{i,k-1} + \epsilon_i$$

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?

$$\mathbf{x}_1 = \begin{bmatrix} 1 & 1 & 0 & 0 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & 1 & 0 & 0 \\ 1 & 0 & 1 & 0 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 1 & 0 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 0 & 1 \\ \vdots & \vdots & \vdots & \vdots \\ 1 & 0 & 0 & 1 \end{bmatrix} \quad \text{or} \quad \mathbf{x}_2 = \begin{bmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 1 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 1 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 1 \end{bmatrix} \quad \text{or} \quad \mathbf{x}_3 = \begin{bmatrix} 1 & 0 & 0 \\ \vdots & \vdots & \vdots \\ 1 & 0 & 0 \\ 0 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 0 & 1 & 0 \\ \vdots & \vdots & \vdots \\ 0 & 0 & 1 \\ \vdots & \vdots & \vdots \\ 0 & 0 & 1 \end{bmatrix}$$

ANOVA model interpretation

Using the model $y_i = \beta_0 + \beta_1 x_{i1} + \dots + \beta_{k-1} x_{i,k-1} + \epsilon_i$, interpret $\beta_0 =$

$\beta_1 =$

Equivalent model

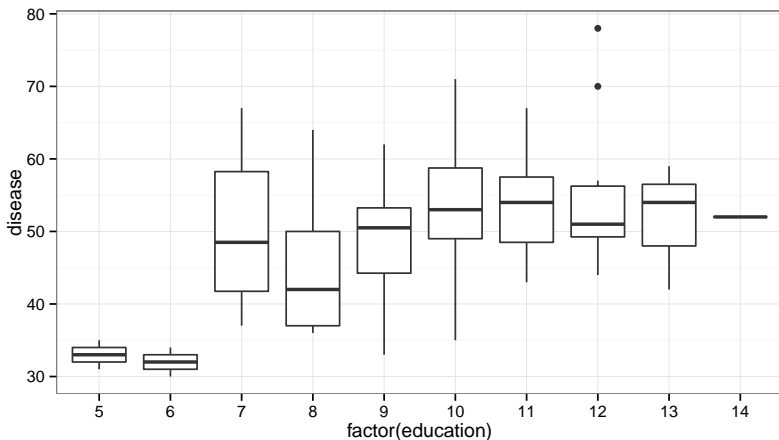
Define the model $y_i = \beta_1 x_{i1} + \dots + \beta_k x_{i,k} + \epsilon_i$ where there are indicators for each possible group

$$\beta_1 =$$

$$\beta_2 =$$

Categorical predictor example: lung data

```
require(ggplot2)  
qplot(factor(education), disease, geom="boxplot", data=dat)
```



Categorical predictor example: lung data

```
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

Categorical predictor example: lung data

```
mlr8 <- lm(disease ~ factor(education) - 1, data=dat)
summary(mlr8)$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

Today's big ideas

- Multiple linear regression models, projections, collinearity, categorical variables