SOC542 Statistical Methods in Sociology II Multiple Regression

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Plan

- Recap
- ► Multiple regression: An overview
- ► Lab: Multiple regression in R

Recap

What we have learned so far

- 1. Fundamentals of frequentist inference
- 2. Simple linear regression
- 3. Probability and Bayesian inference

OLS assumptions review

- \triangleright x and y are independently and identically distributed (IID).
 - The sample x must contain some variability. Specifically, var(x) > 0.
- ightharpoonup The conditional distribution of u given x has a mean of zero.
 - ▶ Errors are independent $E[u_i|x_i] = E[u_i] = 0$.
 - Errors have constant variance $var(u_i) = \sigma^2$.
 - Errors are uncorrelated.
- ▶ If these assumptions are met, then OLS is **BLUE**
 - ► The Best Linear conditionally Unbiased Estimator

Simple linear regression

Let's say we estimate a simple linear regression of the form:

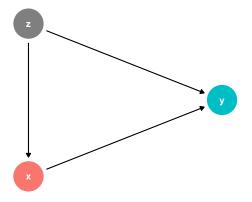
$$\hat{y} = \hat{\beta_0} + \hat{\beta_1} x + \hat{u}$$

- ▶ In this case, we assume that the outcome *y* is a linear function of a single predictor *x*.
- ▶ But what if we think have reason to believe that y is also a function of other predictors?

Omitted variable bias

- Omitted variable bias occurs when we leave out (or omit) a predictor that should be in our model.
- It exists when
 - x is correlated with the omitted variable z.
 - The omitted variable is a predictor of the dependent variable *y*.

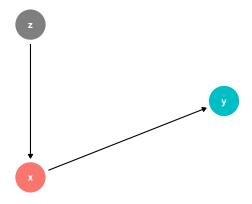
Omitted variable bias



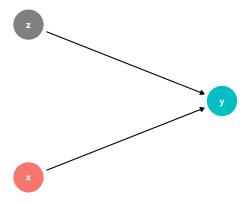
Consequences omitted variable bias

- ▶ The assumption that $E(u_i|x_i) = 0$ is violated.
 - If z is correlated with x but not included, then the error term u captures the unmeasured effect of z and thus u is correlated with x.
- ▶ The slope coefficient β_1 will be *biased*.
 - ► The mean of the sampling distribution of the OLS estimator may not equal the true effect of x.
 - $\hat{\beta}_1 = \beta_1 + bias$
- ▶ The OLS estimator is *inconsistent* as $\hat{\beta}_1$ does not converge in probability to β_1 .
 - ▶ Bias remains even with large samples.
- ► The greater the correlation between *x* and *u*, the greater the bias.

(Not) Omitted variable bias



(Not) Omitted variable bias

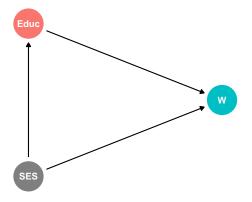


Example

Wealth_i =
$$\beta_0 + \beta_1 Educ + u$$

- Let's say we have a model of wealth as a function of education:
- We estimate the model and see a strong, positive relationship between education and wealth (i.e. $\hat{\beta}_1$ is positive)
- What other factors might be correlated with education and predict wealth?
- Education is correlated with parental socioeconomic status (SES) and predicts wealth. Our estimate of the effect of education is biased without taking SES into account.

Drawing the DAG



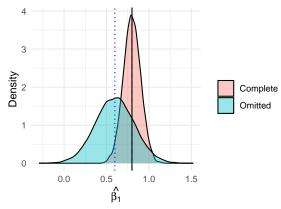
```
N <- 100

x <- rnorm(N,2,1)

z <- 0.1*x + rnorm(N,5,2)

y <- 0.8*x + -2*z + rnorm(N, 0, 1)
```

```
coefs.omitted <- c()
coefs.complete <- c()
sims <- 1E4
for (i in 1:1E4) {
    x <- rnorm(N,2,1)
    z <- 0.1*x + rnorm(N,5,1)
    y <- 0.8*x + -2*z + rnorm(N, 0, 1)
    m.omit <- lm(y ~ x)
    m.both <- lm(y ~ x + z)
    coefs.omitted[i] <- m.omit$coefficients[2]
    coefs.complete[i] <- m.both$coefficients[2]
}</pre>
```



Distribution of $\hat{\beta}_1$ over 10000 simulations.

The multiple regression model

▶ In a multiple regression model, we specify a linear relationship between an outcome and a set of *k* predictors.

$$E[y|x_1, x_2, ..., x_k] = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + ... + \beta_k x_k + u$$

Independent variables and controls

- ► The predictors added to the model can be considered as additional **independent variables** or as **controls**.
- ▶ In general, we use the former term when we have a theoretical reason to be interested in a effect of a variable and the latter when we expect it to matter but are not interested in analyzing the relationship directly.
- We typically add control variables to address potential omitted variable bias.

Interpreting coefficients

► Consider the following population model:

$$y_i = \beta_0 + \beta_1 x + \beta_2 z + u$$

 \triangleright β_1 is the effect of a unit change in x when z is held constant.

$$\beta_1 = \frac{\Delta y}{\Delta x}$$
, holding z constant

Interpreting the intercept

Consider the same model:

$$y_i = \beta_0 + \beta_1 x + \beta_2 z + u$$

 \triangleright β_0 is the expected value of y_i when x=0 and z=0.

The OLS estimator

- ▶ The OLS estimator minimizes the sum of the squared residuals.
- Over n observations and k predictors, we minimize the following quantity:

$$\sum_{i=1}^{n} (y_i - \beta_0 - \beta_1 x_{1i} - \beta_1 x_{2i} - \dots - \beta_k x_{ki})^2$$

► Thus, the predicted values are

$$\hat{y}_i = \hat{\beta}_0 + \hat{\beta}_1 x_{1i} + \hat{\beta}_2 x_{2i} + \dots + \hat{\beta}_k x_{ki}$$

▶ And the residuals are defined as $\hat{u}_i = y_i - \hat{y}_i$ for i = 1, ..., n.

OLS in matrix form

▶ We can write the equation for multiple regression more compactly using matrix notation. Here X is an n by k matrix of predictors and B is vector of coefficients with length k.

$$y = \beta_0 + \beta X + u$$

▶ Ordinary least squares estimates can be computed directly using matrix multiplication, where X is a matrix of predictors and the first column is a vector of 1s (for the intercept) and y is the outcome.

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

OLS in matrix form

```
Intercept <- rep(1,N)</pre>
X <- cbind(Intercept,x,z)</pre>
Betas <- solve(t(X) %*% X) %*% (t(X) %*% y)
print(t(Betas))
##
        Intercept
                            x
## [1,] 0.1205118 0.6961067 -2.035357
m \leftarrow lm(y \sim x + x + z)
print(m$coefficients)
## (Intercept)
                           x
     ##
t() is the transpose operation, solve() finds the inverse of a matrix, and %*% is the matrix multiplication operator.
```

OLS in matrix form

- ➤ OLS estimates are derived directly from algebraic manipulation of the data.
- ► OLS is a special case. Other approaches we will encounter in a few weeks require more complicated maximum likelihood estimation.
- Bayesian regression with uniform priors will converge to the least squares solution, despite a radically different estimation procedure.

Model fit and the Standard Error of the Regression

- ▶ The Standard Error of the Regression (SER) is an estimate of the standard deviation of the error term u_i . It captures the spread of y around the regression line.
- ► For a single regressor,

$$SER = \sqrt{\sigma_{\hat{u}}} = \sqrt{\frac{1}{n-2} \sum_{i=1}^{n} \hat{u}_{i}^{2}} = \sqrt{\frac{SSR}{n-2}}$$

- n − 2 accounts for degrees of freedom used by slope and intercept.
- ► A smaller SER indicates better fit.

Model fit and the Standard Error of the Regression

If we have multiple predictors we need to include an degrees of freedom adjustment k, where k is the number of predictors. The −1 accounts for the intercept.

$$SER = \sqrt{\sigma_{\hat{u}}} = \sqrt{\frac{1}{n-k-1}\sum_{i=1}^{n}\hat{u_i}^2} = \sqrt{\frac{SSR}{n-k-1}}$$

ightharpoonup The adjustment has a small effect when n is large.

Model fit and R^2

 \triangleright We define R^2 in the sample way as a simple regression:

$$R^{2} = \frac{\sum_{i=1}^{n} (\hat{y}_{i} - \bar{y})^{2}}{\sum_{i=1}^{n} (y_{i} - \bar{y})^{2}} = \frac{ESS}{TSS}$$
$$R^{2} = 1 - \frac{SSR}{TSS}$$

Adjusted R^2

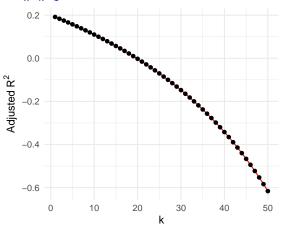
- ▶ R^2 increases mechanistically as we add predictors because the SSR declines as long as $\hat{\beta}_k \neq 0$, inflating model fit.
- ▶ A degrees of freedom correction is used to adjust for this:

Adjusted
$$R^2 = 1 - \frac{n-1}{n-k-1} \frac{SSR}{TSS}$$

Properties of adjusted R^2

- ▶ Adjusted R^2 is always less than R^2 .
- Adding a predictor can increase Adjusted R^2 , but it can decline if the change to the SSR is weaker than the offset n-1/n-k-1.
- Adjusted R^2 can be negative if the reduction in SSR does not offset n 1/n k 1.

The penalty $\frac{n-1}{n-k-1}$ increases as we add predictors



This example shows the effect of the degree of freedom adjustment, assuming $\beta_k = 0$ for all k > 1.

Bayesian R^2

- ▶ There is no direct analogue for R^2 in Bayesian statistics
 - ► Recall that frequentist models assume *fixed* parameters, whereas Bayesian parameters have *distributions*.
- If we treat the Bayesian estimates as fixed, for example by taking the median of the posterior distribution $\hat{\beta}_k$, we could calculate something using the formula above, but it would not account for the *uncertainty* contained in the posterior distribution.

Bayesian R^2

- Instead, we use posterior simulations to repeat the calculation across all samples from the posterior.
- ▶ Bayesian R^2 therefore has a posterior distribution. We can summarize this into a single metric using the same approach as the regression coefficients, e.g. using the median of the posterior distribution.

 $^{^{1}}$ "Everything that depends upon parameters has a posterior distribution" - McElreath 98.

²See GHV 170-171

Significance tests: t-tests

- ► Like simple linear regression, we can interpret the statistical significance of regression coefficients using the t-statistics.
- ▶ Typically, we are interested in testing the null hypothesis that $\beta_k = 0$. We get the t-statistic by dividing a coefficient by its standard error:

$$t = \frac{\hat{\beta}_k - 0}{SE(\hat{\beta}_k)} = \frac{\hat{\beta}_k}{SE(\hat{\beta}_k)}$$

▶ We can the use the t-statistic to look up the relevant *p-value*.

Confidence interval

Most regression software provides a 95% confidence interval around each estimate. For β_k this would take the following form:

$$[\hat{\beta}_j - 1.96SE(\hat{\beta}_j), \hat{\beta}_j + 1.96SE(\hat{\beta}_j)]$$

▶ Recall that only 5% of the probability density of a t-distribution is greater than |1.96|.

Joint tests

- The F-statistic is used to test a joint hypothesis.
- ▶ If we consider a two variable example, we might test the following *null hypothesis*:

$$H_N: \beta_1 = 0, \beta_2 = 0$$

- A joint test has q restrictions. In this case, q = 2.
- ▶ The alternative hypothesis H_A is that one or more of the q restrictions does not hold.

Joint tests and the F-statistic

- ➤ Since we expect the predictors to have a *joint sampling distribution*, we cannot conduct a joint test by summarizing a series of paired tests (e.g. a t-test for every predictor) because the t-statistics are not independent.
- ▶ Instead, we must calculate the F-statistic. Where q = 2 it is defined as:

$$F = \frac{1}{2} \left(\frac{t_1^2 + t_2^2 - 2\hat{\rho}_{t_1, t_2} t_1 t_2}{1 - \hat{\rho}_{t_1, t_2}^2} \right)$$

Joint tests and the F-statistic

If $\hat{\rho}_{t_1,t_2} = 0$ the equation simplifies to the average of the squared t-statistics:

$$F = \frac{1}{2}(t_1^2 + t_2^2)$$

- ▶ The p-value can then be derived from the relevant F-distribution, where $F \sim F_{a,\infty}$
- Typically, we use an F-test to test the restriction that $\beta_1 = 0, \beta_2 = 0, ..., \beta_k = 0.$

Joint tests and the F-statistic

If we assume the residuals are *homoskedastic*, we can test the restriction $\beta_1=0,\beta_2=0,...,\beta_k=0$ using the following formula:

$$F_0 = \frac{(SSR_r - SSR_u)/q}{SSR_u/(n-k+1)}$$

- ▶ The SSR_r is obtained from *restricted* model where we calculate the SSR assuming the null hypothesis is true. The SSR from the fitted model, SSR_u , is known as the *unrestricted* SSR.
- ▶ The test statistic is assessed using an *F*-distribution with q degrees of freedom and n k + 1 observations.
- ▶ In most cases the homoskedasticity assumption is likely violated, so we use the more complicated formula from the previous slide, known as the heteroskedasticity robust F-statistic.

Interpreting regression output

```
Call:
lm(formula = y \sim x + z)
Residuals:
    Min
          10 Median 30
                                      Max
-2.17211 -0.49561 0.00997 0.54232 3.09640
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 0.12051 0.53775 0.224
                                        0.823
          0.69611 0.09055 7.688 1.23e-11 ***
         -2.03536 0.10526 -19.336 < 2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 0.9607 on 97 degrees of freedom
Multiple R-squared: 0.7973, Adjusted R-squared: 0.7932
F-statistic: 190.8 on 2 and 97 DF, p-value: < 2.2e-16
```

Bayesian approaches

- "We have essentially no interest in using hypothesis tests for regression because we almost never encounter problems where it would make sense to think of the coefficients as being exactly zero" - GHV 147
- Bayesian regression is assessed by analyzing the posterior distribution of parameters to understand uncertainty.
- ► Nonetheless, Bayesian equivalents to t-tests and F-tests can be used if desired ³

³See Kruschke and Liddell 2018.

Multicollinearity

- ▶ Multicollinearity occurs when a predictor *x* is highly correlated one or more other predictors *z*.
 - **Perfect multicollinearity** arises when cor(x, z) = 1 or \$ -1\$.
 - Usually due to some type of misspecification. e.g. accidentally including the same variable twice.
 - Imperfect multicollinearity means that two or more regressors are highly correlated.

Multicollinearity and its implications

▶ Assume the following model and that x and z are highly correlated:

$$\hat{y}_i = \hat{\beta}_0 + \hat{\beta}_1 x_i + \hat{\beta}_2 z_i + u_i$$

- ► The variance of $\hat{\beta}_1^4$ is inversely proportional to $1 \rho_{x,z}^2$, where $\rho_{x,z}$ is the correlation between x and z.
 - If $\rho_{x,z}$ is large, then this term is small and thus the variance is large.
- Multicollinearity increases variance and reduces precision, potentially making β_1 non-identifiable.

The same issue also applies to $\hat{\beta}_2$

```
N <- 100

x <- rnorm(N,2,1)

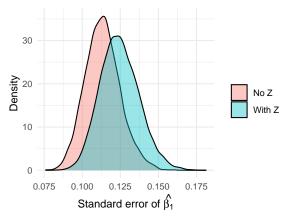
x2 <- rnorm(N,0,1)

z <- 0.7*x + rnorm(N,0,1)

y <- 0.5*x + -0.5*x2 + 0.5*z + rnorm(N, 1, 1)
```

```
m1 \leftarrow summary(lm(y \sim x + x2))
m2 \leftarrow summary(lm(y \sim x + x2 + z))
round(m1$coefficients,2) # omitted variable bias
##
            Estimate Std. Error t value Pr(>|t|)
               1.66 0.25 6.66
## (Intercept)
              0.59 0.11 5.22
## x
## x2
            -0.59 0.11 -5.42
round(m2$coefficients,2) # multicolinearity
##
            Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                1.48 0.23 6.31 0.00
              0.29 0.13 2.28 0.02
## x
             -0.58 0.10 -5.79 0.00
## x2
## 2.
              0.49 0.12 4.24 0.00
```

```
se.omitted <- c()
se.complete <- c()
sims < -1E4
for (i in 1:1E4) {
    x \leftarrow rnorm(N, 2, 1)
    x2 \leftarrow rnorm(N,0,1)
    z \leftarrow 0.7*x + rnorm(N.0.1)
    y \leftarrow 0.5*x + -0.5*x2 + 0.5*z + rnorm(N, 1, 1)
    m.omit \leftarrow summary(lm(y \sim x + x2))
    m.complete \leftarrow summary(lm(y \sim x + x2 + z))
     se.omitted[i] <- m.omit$coefficients[2,2]</pre>
    se.complete[i] <- m.complete$coefficients[2,2]</pre>
```



Distribution of standard error over 10000 simulations.

Fixing multicollinearity

- ▶ In general, multicollinearity is less severe than omitted variable bias.
 - ► The inflated variance will lead to more Type II errors than Type I errors
 - Omitted variable bias can produce Type I errors, sign errors, and magnitude errors.

Fixing multicollinearity

- ► Solution 1: Use more data. If we have a larger sample then we might be able to learn from additional variation in x and z.
- Solution 2: If we are only concerned about x then we could exclude z. But this risks omitted variable bias if z is also a predictor of y.
- ➤ Solution 3: Transform or combine predictors (e.g. factor analysis).

Revisiting our assumptions

- \triangleright $E(u_i|x_{1i},x_{2i},...,x_{ki})=0$
- ► All $y_i, x_{1i}, x_{2i}, ..., x_{ki}$ are IID.
- Large outliers are unlikely.
- No perfect multicollinearity.

Spurious relationships and confounding

- Sometimes we observe spurious relationships in regression models where a correlation between two variables exists, despite the absence of any causal relationship.
 - e.g. Finding that hurricanes with female names caused more deaths than male named hurricanes.
- Sometimes this occur purely due to chance, but it can be due to confounding: a confounding variable z influences both x and y.
- Adding more predictors can often help to reduce the risk of spurious associations.
 - ▶ If we control for the confounder *z*, the spurious relationship between *y* and *x* disappears.

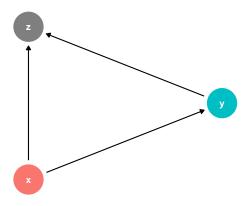
Masked relationships

- Assume a true relationship between y and x.
- We estimate a model $y = \beta_0 + \beta_1 x$.
- The results do not show evidence of an association (i.e. $\hat{\beta}_1 \approx 0$).
- \triangleright We estimate a second model including a new predictor z.
- Controlling for z allows us to observe a relationship between y and x.

Colliders

- \triangleright z is caused by y and x.
- In this case, z is a **collider** and controlling for z can introduce bias.
- Using DAGs can help to formalize relationships between variables and identify potential colliders.

Colliders⁵

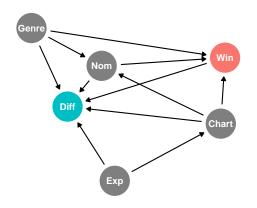


⁵See Elwert, Felix, and Christopher Winship. 2014. "Endogenous Selection Bias: The Problem of Conditioning on a Collider Variable." *Annual Review of Sociology* 40(1):31–53. doi: 10.1146/annurev-soc-071913-043455.

Variable selection

- It is often conventional practice to include a wide array of potential confounders in a regression model ("kitchen sink" or "garbage can" regressions), but this approach can cause problems!
- We must carefully consider omitted variable bias, multicollinearity, and collider bias when specifying models.
- We must use domain knowledge and theory to guide model specifications, we cannot identify these issues from the data alone.
- ▶ DAGs are a useful tool for representing our assumptions, guiding variable selection, and identifying problematic specifications.

DAGs in the wild⁶



⁶ Stylized DAG based on analysis from Negro, Giacomo, Balázs Kovács, and Glenn R. Carroll. 2022. "What's Next? Artists' Music after Grammy Awards." *American Sociological Review* 87(4):644–74. doi: 10.1177/00031224221103257.

Next week

Non-linear predictors

- ► Dummy variables
- ► Categorical variables
- Non-linear transformations

Lab

► Estimating and interpreting multiple regression models