BIOS7345 Lab 2

Least-squares regression

Sarah Lotspeich

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Prologue: Pipes

Before we begin, there are two basic pipes functions I will be using.

Single pipe: %>%Double pipe: %<>%

[1] 5.5

The %>% operator "pipes" an argument into a function. For example, the following two lines of code are equivalent.

```
# without pipes
mean(seq(1, 10))
## [1] 5.5
# with pipes
seq(1, 10) %>% mean()
```

The single pipe is super useful when you have lots of nested arguments, and it saves you a lot of confusing parentheses.

The %<>% operator "pipes" the object on the left-hand side into a function, and then reassigns the new value to the original name. Some simple code:

```
# create a vector
x <- seq(1, 10)

# say I want to squareroot transform the whole vector without pipes
x <- sqrt(x)
x

## [1] 1.000000 1.414214 1.732051 2.0000000 2.236068 2.449490 2.645751
## [8] 2.828427 3.000000 3.162278

# create a vector
x <- seq(1, 10)
# with pipes
x %<>% sqrt()
x

## [1] 1.000000 1.414214 1.732051 2.0000000 2.236068 2.449490 2.645751
## [8] 2.828427 3.000000 3.162278
```

Data: Entertainer Career Arcs

These data come from a super cool social media site for data people called data.world. The link to the documentation can be found here, but the TLDR is:

- 1. we have a sample of <u>63 notable entertainers</u> (actors, actresses, musicians, etc.) from the last hundred years, and
- 2. for each we know gender (gender_traditional), the age when they first "broke through" (breakthrough_age), and the age when they won their first award (first_award_age).

Spoiler alert: you won't know all of the names. Darn millenials.

```
stars <- read.csv(file = "https://query.data.world/s/xsweyho62srvyorzoly3w2uctn76rz",
header = TRUE, stringsAsFactors = FALSE)</pre>
```

Objective

We are interested in the following research questions:

- 1. What is the association between the age at which an entertainer broke through and the age at which they won their first award?
- 2. Do men and women have different expected ages at which they would win their first award?
- 3. Is the association between ages of breakthrough and first award different for men and women?

We can answer them using three separate, pre-specified regression models.

Models

Simple linear regression with one continuous predictor

The first model we will fit is for first_award_age on breakthrough_age. Specify the model, using proper LaTex, below:

 $E(\text{first award age}|\text{breakthrough age}) = \beta_0 + \beta_1(\text{breakthrough age}).$

Now, begin fitting the model by restructuring these data into your response vector (y) and design matrix (X). I recommend using data.matrix() for this.

```
# create response vector

# create design matrix using data.matrix()
```

Begin by estimating the model coefficients, $\hat{\boldsymbol{\beta}} = \begin{bmatrix} \hat{\beta}_0 & \hat{\beta}_1 \end{bmatrix}^T$ according to the Rencher Theorem 7.3a:

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

```
# use matrix operations to calculate least-squares estimates beta0, beta1
# for later, go ahead and round these to 3 digits
```

Now we need an unbiased estimate for the variance of the error terms (σ^2), which we obtain using Rencher Equation (7.23):

$$s^2 = \frac{1}{n-k-1} (\boldsymbol{y} - \boldsymbol{X}\boldsymbol{\hat{\beta}})^T (\boldsymbol{y} - \boldsymbol{X}\boldsymbol{\hat{\beta}}),$$

where k = the number of predictors in your model and n = sample size.

```
# sample size
# number of predictors (excl. intercept)
# use (7.23) to estimate the variance of the error terms
```

The last thing this model needs are measures of uncertainty (e.g. standard errors) for the coefficient estimates. Using our estimate s^2 above, we can calculate these directly according to Rencher (7.27):

$$\hat{Cov}(\hat{\boldsymbol{\beta}}) = s^2(\boldsymbol{X}^T\boldsymbol{X})^{-1}.$$

```
# estimate the SEs for the estimated coefficients
# round these off
```

To wrap up this model let's make a nice table and print it using xtable() (which we totally remember from last week's lab). Recall: what do we need to add in the chunk header to make this print nicely?

```
# create a table for the model
```

Try using the dplyr::mutate() function to append columns for the Wald Z test statistic and p-value for significance tests for $H_0: \beta_0 = 0$ and $H_0: \beta_1 = 1$. Consider using ifelse() to clean up the p-values if they're really small.

```
# append Z test statistics and p-values
# print model table
```

Do future you a favor and put this into a function called my_ls() with parameters for the response vectors, y, the design matrix, X, and a caption for the model table to be printed.

```
# write a function to take in y, X and print the model table
my_ls <- function(y, X, cap) {
}</pre>
```

For comparison, you may now fit this model using rms::ols().

```
# fit the same model using ols()
```

Simple linear regression with one categorical predictor

The next model we are going to fit is for first_award_age on traditional_gender. Once again, Specify the model, using proper LaTex, below:

```
E(\text{first award age}|\text{gender}) = \beta_0 + \beta_1(\text{gender}).
```

Our response vector (y) is the same as above, but we code categorical design matrices differently. Treating males as referrent, create the design matrix for this model now.

```
# create design matrix using data.matrix()
```

Use my_ls() to calculate the same three values for this model, and create a table of the same format.

```
# use my_ls() to fit this model
```

Check yourself. Since we want males to be our reference group, we need to recode gender_traditional as a factor to make sure that ols() will know.

```
# make gender_traditional a factor
# fit the same model using ols()
```

Multiple linear regression with interacting continuous and categorical predictors

Our last model seeks to incorporate separate effects of breakthrough_age for male and female entertainers, which we accomplish via an interaction between predictors breakthrough_age and traditional_gender. Thus, we specify the following model:

 $E(\text{first award age}|\text{breakthrough age, gender}) = \beta_0 + \beta_1(\text{breakthrough age}) + \beta_2(\text{gender}) + \beta_3(\text{breakthrough age}) \times (\text{gender}).$

Once more, we can recycle y from the first model, but our design matrix X now needs:

- 1. an intercept,
- 2. the continuous breakthrough_age values,
- 3. the indicator for female (as in model 2), and
- 4. the interaction between breakthrough_age and the female indicator.

```
# create design matrix using data.matrix() for model: first_award_age ~
# breakthrough_age*gender_traditional
```

Use my_ls() again to calculate the same three values for this model, and create a table of the same format.

```
# fit this model using my_ls()
```

And check yourself against rms::ols().

```
# fit the same model using ols()
```

Predicting based on models

Manually calculating predictions

Begin by plotting breakthrough_age against first_award_age.

```
# make a scatterplot of breakthrough_age vs. first_award_age
```

Manually calculate the predicted values for first_award_age based on your beta_hat1 estimates for breakthrough_age from 10 to 55 years old.

```
# create design matrix to predict age at first award for breakthrough ages # 10-55 # use matrix operations to predict y
```

Add a line to your mod1_plot with the predicted values from the model.

```
# create a dataframe with columns for x and predicted y

# add geom_line() layer to mod1_plot with the model predictions for
# first_award_age
```

Lastly, let's add the model equation to the plot (because we're fancy). Before we put it into the plot, let's make sure we're comfortable with paste().

```
# test your paste() statement to make sure you like how it looks
# now add an annotate() layer to mod1_plot with the model equation
# and print your plot
```

Invariance of predicted values to linear transformation

Often, we like to transform predictors so that the coefficients can be interpreted more meaningfully. Say we're more interested in 5 year changes in breakthrough_age than 1 year changes. Let's transform breakthrough_age, refit model 1 (recall: first_award_age ~ breakthrough_age), and compare predicted values between the original and linearly transformed models.

```
# add a column for breakthrough_age_centered by subtracting the median
# create the design matrix again
# use matrix operations to calculate least-squares estimates beta0, beta1
# create design matrix to predict age at first award for breakthrough ages
# 10-55
# and predict first_award_age for ages 10-55 again using matrix operations
# create a dataframe with columns for x and predicted y
```

Create the same plot as before with breakthrough_age_5yrs against first_award_age, overlaid with your transformed model predictions. Use ggpubr:ggarrange() to print this plot side-by-side with the original.

```
# test your paste() statement to make sure you like how it looks
# create the same plot using the transformed model
# print this plot next to the original
```

This is follows from Rencher Theorem 7.3e.

Full rank transformation on predictor

Take any full-rank 2×2 matrix K_1 (e.g. sampling the elements independently from a Normal distribution).

```
\# generate full-rank 2x2 transformation matrix \mathit{K}
```

Just for fun, let's make sure that this matrix is full-rank.

You: "But Sarah, didn't we just design it to be?"

Sarah: "Yes, but wouldn't you like to know how to calculate the rank of a matrix in R anyway?"

```
# check Rank(K)
```

There are some other helpful matrix operations in R that I would recommend for checking homework, spending a Friday night at home, etc. that can be found here.

Now, consider a model adjusting for breakthrough_age and tranditional_gender without the interaction. Construct this design matrix,

```
# create the design matrix for model: first_award_age ~ breakthrough_age +
# traditional_gender
```

and fit this model.

```
# fit the model for y ~ X5
```

Next, let's do a full-rank linear transformation of our design matrix X5.

```
# full-rank linear transformation of X5 be careful: should you be # transforming the intercept column?
```

What do we know about $Rank(\mathbf{Z})$? First, check that X5 is full-rank.

```
# check rank(X5)
```

Now, since X5 is full-rank and Z is a full-rank, square matrix, we have that multiplying X5 by Z does not change the rank (by Rencher <u>Theorem 2.4.</u>). Check that the rank of Z equals the rank of X5.

```
\# check that Rank(Z) = Rank(X5)
```

Fit the model for first_award_age on the transformed design matrix Z now.

```
# fit the model for y \sim Z
```

Last, we compare the fitted values for the 63 entertainers based on the model before and after full-rank linear transformation.

```
# check that predicted values for the models are the same
```

This is Rencher Theorem 7.3e Corollary 1.