

Unit 3 Lecture 1: Logistic Regression

October 4, 2022

```
library(tidyverse)
library(splines)
library(cowplot)
library(stat471)
```

Linear regression

In the context of splines, we worked with `age` and `income` from the income data. In fact, these data have more columns as well, which we might want to use for predicting income:

```
income_data <- read_tsv("income_data.tsv")
income_data
```

```
## # A tibble: 3,000 x 8
##   income year  age maritl      race  education  jobclass  health
##   <dbl> <dbl> <dbl> <chr>    <chr>    <chr>    <chr>    <chr>
## 1  75.0  2006   18 1. Never Married 1. White 1. < HS Grad 1. Indus~ 1. <=~
## 2  70.5  2004   24 1. Never Married 1. White 4. College Grad 2. Infor~ 2. >=~
## 3 131.   2003   45 2. Married      1. White 3. Some College 1. Indus~ 1. <=~
## 4 155.   2003   43 2. Married      3. Asian 4. College Grad 2. Infor~ 2. >=~
## 5  75.0  2005   50 4. Divorced     1. White 2. HS Grad    2. Infor~ 1. <=~
## 6 127.   2008   54 2. Married      1. White 4. College Grad 2. Infor~ 2. >=~
## 7 170.   2009   44 2. Married      4. Other 3. Some College 1. Indus~ 2. >=~
## 8 112.   2008   30 1. Never Married 3. Asian 3. Some College 2. Infor~ 1. <=~
## 9 119.   2006   41 1. Never Married 2. Black 3. Some College 2. Infor~ 2. >=~
## 10 129.  2004   52 2. Married      1. White 2. HS Grad    2. Infor~ 2. >=~
## # ... with 2,990 more rows
## # i Use `print(n = ...)` to see more rows
```

Let's split our data into train and test:

```
set.seed(4710)
train_samples <- sample(1:nrow(income_data), 0.8 * nrow(income_data))
income_train <- income_data %>% filter(row_number() %in% train_samples)
income_test  <- income_data %>% filter(!(row_number() %in% train_samples))
```

We can run a linear regression using `lm()`, which we have already used for spline fits:

```
lm(income ~ age, data = income_train)

##
## Call:
## lm(formula = income ~ age, data = income_train)
##
## Coefficients:
## (Intercept)      age
##    81.9531    0.7225
```

We can specify multiple predictors using the + syntax:

```
lm(income ~ age + education, data = income_train)

##
## Call:
## lm(formula = income ~ age + education, data = income_train)
##
## Coefficients:
##              (Intercept)                age
##              60.8669                0.5683
##      education2. HS Grad      education3. Some College
##              11.2193                24.5819
##      education4. College Grad  education5. Advanced Degree
##              40.3823                64.8561
```

We can include all predictors except for the response using the . syntax:

```
lm(income ~ ., data = income_train)

##
## Call:
## lm(formula = income ~ ., data = income_train)
##
## Coefficients:
##              (Intercept)                year
##             -2550.4114                1.2973
##                age          maritl2. Married
##                0.3631                17.7661
##      maritl3. Widowed          maritl4. Divorced
##                4.1442                4.7896
##      maritl5. Separated          race2. Black
##               13.0303               -5.5724
##                race3. Asian          race4. Other
##               -3.4709               -3.8735
##      education2. HS Grad      education3. Some College
##               10.7993                23.3761
##      education4. College Grad  education5. Advanced Degree
##               37.0926                59.2820
##      jobclass2. Information          health2. >=Very Good
##               5.0515                7.3695
```

How do we interpret the coefficient for age? What about for education2. HS Grad?

We can make predictions for a given linear regression model using predict():

```
lm_fit <- lm(income ~ ., data = income_train) # first save the model fit
predictions <- predict(lm_fit, newdata = income_test)
```

We can then assess test error metrics like RMSE based on these predictions:

```
sqrt(mean((income_test$income - predictions)^2))

## [1] 33.12452
```

In Unit 2, we found that, if modeling income as a spline function of age, the best choice of degrees of freedom is around 3. How does the RMSE of this model compare to that of the model above using all of the variables?

Logistic regression

Now, let's move on to logistic regression. Recall the default data:

```
default_data <- read_tsv("default_data.tsv")
default_data
```

```
## # A tibble: 10,000 x 4
##   default student balance income
##   <chr>    <chr>      <dbl>  <dbl>
## 1 No      No          730.  44362.
## 2 No      Yes          817.  12106.
## 3 No      No         1074.  31767.
## 4 No      No          529.  35704.
## 5 No      No          786.  38463.
## 6 No      Yes          920.   7492.
## 7 No      No          826.  24905.
## 8 No      Yes          809.  17600.
## 9 No      No         1161.  37469.
## 10 No     No           0   29275.
## # ... with 9,990 more rows
## # i Use `print(n = ...)` to see more rows
```

The rest of the activity will be easier if we code default as 0-1:

```
default_data <- default_data %>% mutate(default = as.numeric(default == "Yes"))
default_data
```

```
## # A tibble: 10,000 x 4
##   default student balance income
##   <dbl> <chr>      <dbl>  <dbl>
## 1      0 No          730.  44362.
## 2      0 Yes          817.  12106.
## 3      0 No         1074.  31767.
## 4      0 No          529.  35704.
## 5      0 No          786.  38463.
## 6      0 Yes          920.   7492.
## 7      0 No          826.  24905.
## 8      0 Yes          809.  17600.
## 9      0 No         1161.  37469.
## 10     0 No           0   29275.
## # ... with 9,990 more rows
## # i Use `print(n = ...)` to see more rows
```

Let's split the default data into training and test sets:

```
set.seed(4710)
train_samples <- sample(1:nrow(default_data), 0.8 * nrow(default_data))
default_train <- default_data %>% filter(row_number() %in% train_samples)
default_test <- default_data %>% filter(!(row_number() %in% train_samples))
```

Running a logistic regression

The way to run a logistic regression is through the `glm` function:

```
glm_fit <- glm(default ~ student + balance + income,
  family = "binomial",
```

```
data = default_train
)
coef(glm_fit)
```

```
## (Intercept) studentYes balance income
## -1.088496e+01 -5.701843e-01 5.635043e-03 7.560014e-06
```

- What is the coefficient estimate for **student**?
- Does this suggest that being a student increases or decreases the probability of default, other things being equal?
- According to this estimate, how does being a student impact the log-odds of default? How does it impact the odds of default?

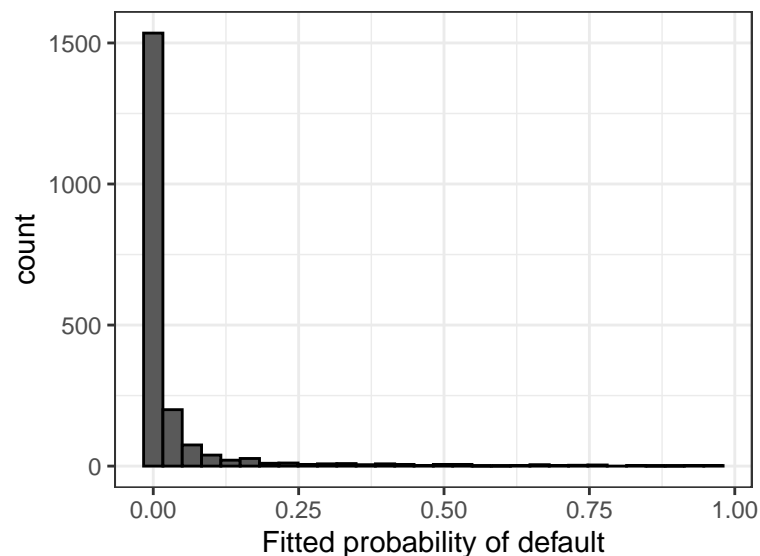
Fitted probabilities and making predictions

We can extract the fitted probabilities of default for a test set using the `predict` function:

```
fitted_probabilities <- predict(glm_fit,
  newdata = default_test,
  type = "response" # to get output on probability scale
)
head(fitted_probabilities)
```

```
##          1          2          3          4          5          6
## 0.0099981865 0.0019920721 0.0112420255 0.0117355375 0.0001321415 0.0051567773
```

```
tibble(fitted_probabilities) %>%
  ggplot(aes(x = fitted_probabilities)) +
  geom_histogram(color = "black") +
  labs(x = "Fitted probability of default")
```



We can now make predictions based on the fitted probabilities using the standard 0.5 threshold:

```
predictions <- as.numeric(fitted_probabilities > 0.5)
head(predictions)
```

```
## [1] 0 0 0 0 0 0
```

Evaluating the classifier

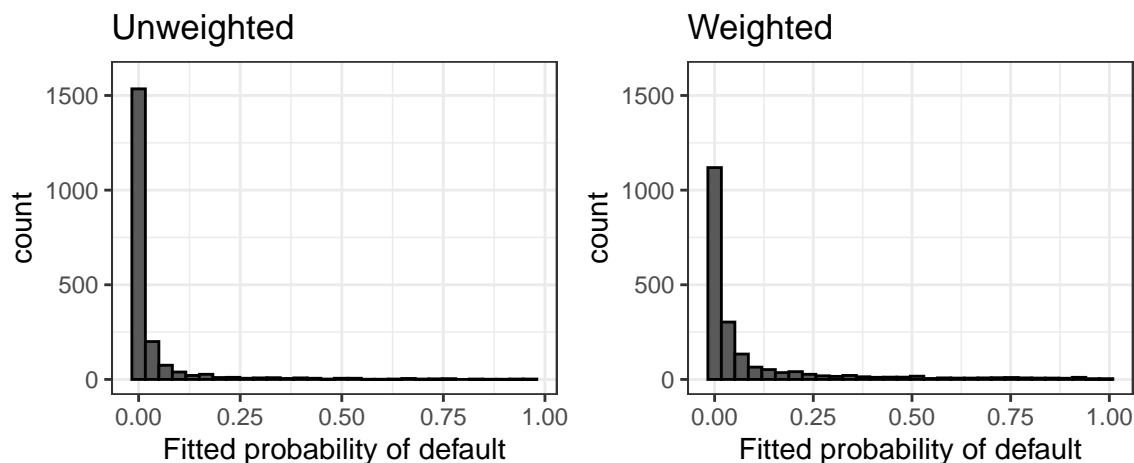
Let's evaluate the performance of the above logistic regression classifier on the test set rate. We can use the `classification_metrics()` function from the `stat471` package.

```
classification_metrics(  
  test_responses = default_test$default,  
  test_predictions = predictions  
)
```

```
## # A tibble: 1 x 5  
##   misclass_err w_misclass_err   TPR   TNR     F  
##         <dbl> <lgl>         <dbl> <dbl> <dbl>  
## 1         0.028 NA         0.370 0.995 0.539
```

If we want the classifier to pay more attention to the positive class, we'll need to upweight it. Fortunately, logistic regression accommodates observation weights, via the `weights` argument to `glm()`. Let's upweight the positive class by a factor of 5:

```
# fit the weighted GLM  
glm_fit_weighted <- glm(default ~ student + balance + income,  
  family = "binomial",  
  weights = 5 * (default_train$default == 1) + 1 * (default_train$default == 0),  
  data = default_train  
)  
# extract the fitted probabilities  
fitted_probabilities_weighted <- predict(glm_fit_weighted,  
  newdata = default_test,  
  type = "response" # to get output on probability scale  
)  
# plot the predicted probabilities based on the unweighted and weighted logistic  
# regression fits  
plot_grid(  
  tibble(fitted_probabilities) %>%  
    ggplot(aes(x = fitted_probabilities)) +  
    geom_histogram(color = "black") +  
    scale_y_continuous(limits = c(0, 1600)) +  
    labs(  
      x = "Fitted probability of default",  
      title = "Unweighted"  
    ),  
  tibble(fitted_probabilities_weighted) %>%  
    ggplot(aes(x = fitted_probabilities_weighted)) +  
    geom_histogram(color = "black") +  
    scale_y_continuous(limits = c(0, 1600)) +  
    labs(  
      x = "Fitted probability of default",  
      title = "Weighted"  
    )  
)
```



How did the predicted probabilities change? Is this what we expected?

Let's make predictions by thresholding at 0.5 and evaluate the resulting classifier:

```
predictions_weighted <- as.numeric(fitted_probabilities_weighted > 0.5)
bind_rows(
  classification_metrics(
    test_responses = default_test$default,
    test_predictions = predictions
  ) %>%
  mutate(weighting = FALSE, .before = 1),
  classification_metrics(
    test_responses = default_test$default,
    test_predictions = predictions_weighted
  ) %>%
  mutate(weighting = TRUE, .before = 1)
)
```

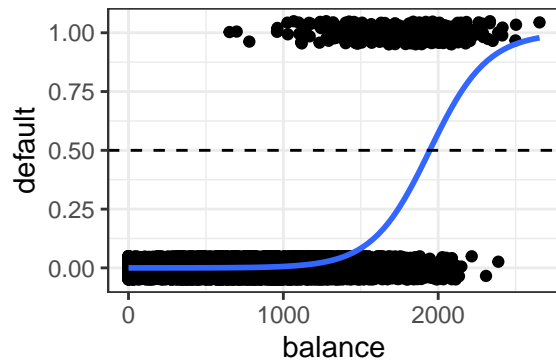
```
## # A tibble: 2 x 6
##   weighting misclass_err w_misclass_err   TPR   TNR     F
##   <lgl>         <dbl> <lgl>         <dbl> <dbl> <dbl>
## 1 FALSE         0.028 NA             0.370 0.995 0.539
## 2 TRUE          0.041 NA             0.726 0.968 0.830
```

How did these test error metrics change, compared to the unweighted case? Is this what we expected?

Plotting a univariate logistic regression fit

Univariate logistic regression fits can be plotted using `geom_smooth()`:

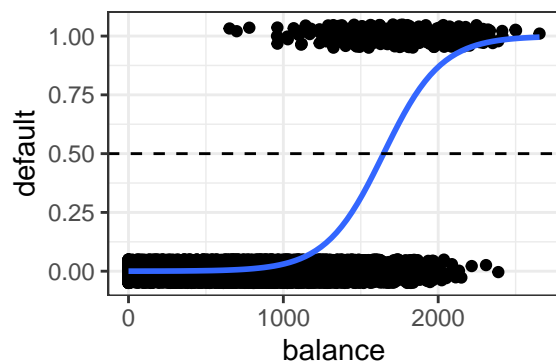
```
default_train %>%
  ggplot(aes(x = balance, y = default)) +
  geom_jitter(height = .05) +
  geom_smooth(
    method = "glm",
    formula = "y~x",
    method.args = list(family = "binomial"), # extra argument
    se = FALSE
  ) +
  geom_hline(yintercept = 0.5, linetype = "dashed")
```



Roughly at what value of balance do we switch from predicting no default to predicting default?

`geom_smooth()` accommodates weighted logistic regression as well:

```
default_train %>%
  ggplot(aes(x = balance, y = default)) +
  geom_jitter(height = .05) +
  geom_smooth(
    method = "glm",
    formula = "y~x",
    method.args = list(family = "binomial"),
    aes(weight = ifelse(default == 1, 5, 1)), # specify weights inside aes()
    se = FALSE
  ) +
  geom_hline(yintercept = 0.5, linetype = "dashed")
```



Now, for roughly what value of balance do we switch from predicting no default to predicting default? How does this compare to the above? Is this what we expected?

Exercise

For logistic regression, if we use `type = "response"` within `predict()` we get the fitted probabilities, whereas `type = "link"` gives us the probabilities on the log-odds scale (also called the “score” in the slides). For `glm_fit` defined above, use `predict()` with `type = "response"` to define a vector called `probabilities` and with `type = "link"` to define a vector called `scores`. Then, create a scatter plot of `probabilities` versus `scores`. What is the relationship between the two?