## Chapter 9

# Interpreting a Logistic Regression Model

This chapter is similar to Chapter 8 but focuses on logistic regression models. We first encountered these models as examples of classification algorithms in Chapter 2. Because of their popularity in the clinical domain, it's important to understand how these models are fit and how to interpret the summary output produced by software.

## 9.1 ER Readmissions Example from Chapter 2

In Chapter 2, we saw an example where information about two predictors – a disease severity score  $(x_1)$  and a social determinants score  $(x_2)$  – was used to predict a binary outcome: whether a patient would be readmitted to the ER within 30 days of discharge. We tried three different supervised learning algorithms, one of which was a **logistic regression** model (Section 2.3.1). The output from that model is repeated below.

```
m3 \leftarrow glm(y \sim x1 + x2, data = df, family = "binomial")
                                        summary(m3)
                                         Call:
                                         glm(formula = y \sim x1 + x2, family = "binomial", data = df)
                                         Deviance Residuals:
                                                          1Q
                                                                 Median
                                         -1.88232 -0.90614 -0.05965 0.86579
                                         Coefficients:
                                                     Estimate Std. Error z value Pr(>|z|)
                                         (Intercept) 0.9780 0.2945 3.321 0.000897 x1 0.1344 0.1372 0.980 0.327272
Social Determinants Score (x2)
                                         x2
                                                       -1.3981
                                                                 0.2316 -6.035 1.59e-09 ***
                                         Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
                                         (Dispersion parameter for binomial family taken to be 1)
                                             Null deviance: 277.26 on 199 degrees of freedom
                                         Residual deviance: 209.54 on 197 degrees of freedom
                                         AIC: 215.54
                                         Number of Fisher Scoring iterations: 4
           Disease Severity Score (x1)
```

## 9.2 Example: Low Birthweight Dataset

The goal of this study was to identify risk factors associated with giving birth to a low birth weight baby (a baby weighing less than 2500 grams). Infant mortality rates and birth defect rates are very high for low birth weight babies. A woman's behavior during pregnancy (including diet, smoking habits, and receiving prenatal care) can greatly alter the chances of carrying the baby to term and, consequently, of delivering a baby of normal birth weight.

Data were collected on 189 women, 59 of which had low birth weight babies and 130 of which had normal birth weight babies.

```
Low birth weight (0 = birth weight \geq 2500 g;
LOW
       1 = birth weight < 2500 g
      Age of mother in years
AGE
LWT
      Mother's weight in pounds at last menstrual period
      Race (1 = white, 2 = black, 3 = other)
RACE
SMOKE Smoking status during pregnancy (1 = yes, 0 = no)
      History of premature labor (0 = none, 1 = one, etc.)
PTL
ΗТ
      History of hypertension (0 = no, 1 = yes)
      Presence of uterine irritability (0 = no, 1 = yes)
UΙ
      Number of physician visits during the first trimester
FTV
BWT
       Birth weight in grams
```

SOURCE: Hosmer and Lemeshow (2000) *Applied Logistic Regression: Second Edition*. Data were collected at Baystate Medical Center, Springfield, Massachusetts during 1986.

We would like to predict LOW based on all of the other covariates except BWT. (Why not use BWT?) The GLM output of this model is:

```
Call:
qlm(formula = LOW ~ AGE + LWT + RACE + SMOKE + PTL + HT + UI +
   FTV, family = "binomial", data = d)
Deviance Residuals:
   Min
            1Q
                Median
                            3Q
-1.8946 -0.8212 -0.5316 0.9818
                                2.2125
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
(Intercept) 0.480623 1.196888
                               0.402
                                    0.68801
AGE
          -0.029549 0.037031 -0.798 0.42489
          -0.015424 0.006919 -2.229 0.02580 *
LWT
           1.272260 0.527357 2.413 0.01584 *
RACE2
           0.880496 0.440778 1.998 0.04576 *
RACE3
           0.938846 0.402147 2.335 0.01957 *
SMOKE
           PTL
ΗТ
           1.863303 0.697533 2.671 0.00756 **
           0.767648   0.459318   1.671   0.09467 .
UΙ
           0.065302 0.172394 0.379 0.70484
FTV
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

(Dispersion parameter for binomial family taken to be 1)

```
Null deviance: 234.67 on 188 degrees of freedom Residual deviance: 201.28 on 179 degrees of freedom AIC: 221.28
```

Number of Fisher Scoring iterations: 4

#### **Question 9.1**

In this model, is the effect of one predictor (say, AGE) impacted by the value(s) of any of the other predictor(s)? How does this differ from the other classification algorithms we've seen (KNN and decision trees)? What are the advantages and disadvantages of this choice?

#### **Question 9.2**

Comment on how the variable RACE enters into the model here. Does this make sense in light of what that variable means and how it potentially interacts with the other study variables?

### Question 9.3

Interpret the values of each of these coefficients. Based on the coefficient values and their standard errors, which predictor(s) do you think have the greatest impact on whether or not a woman has a low birthweight baby?