

Logistic regression and prediction

Agenda

- + Exam 1
 - + Wednesday September 21, in class
 - + Covers material up through today (inclusive)
 - + Closed notes
 - + Bring a calculator (cannot use phone or laptop)
 - + I won't ask you to write R code, but you may need to interpret R output
 - + Questions similar to assignments and class activities
- + Today: more logistic regression

Data

Data on 5720 Vietnamese children, admitted to hospital with possible dengue fever. Variables include:

- + Dengue: whether the patient actually has dengue fever, based on a lab test (0 = no, 1 = yes)
- + Temperature: patient's body temperature (in Celsius)
- + Abdominal: whether the patient has abdominal pain (0 = no, 1 = yes)
- + HCT: patient's hematocrit (proportion of red blood cells)
- + Age: patient's age (in years)
- + Sex: patient's sex
- + + several others

Last time

$$Y_i \sim \text{Bernoulli}(\pi_i)$$

$$\log\left(\frac{\pi_i}{1 - \pi_i}\right) = \beta_0 + \beta_1 \text{Temperature}_i + \beta_2 \text{Abdominal}_i \\ + \beta_3 \text{Temperature}_i \cdot \text{Abdominal}_i$$

Does the model improve when we add hematocrit (the proportion of red blood cells)?

Model

$$Y_i \sim \text{Bernoulli}(\pi_i)$$

$$\begin{aligned} \log\left(\frac{\pi_i}{1 - \pi_i}\right) &= \beta_0 + \beta_1 \text{Temperature}_i + \beta_2 \text{Abdominal}_i \\ &\quad + \beta_3 \text{Temperature}_i \cdot \text{Abdominal}_i \\ &\quad + \beta_4 \text{HCT}_i \end{aligned}$$

Class activity, Part I

https://sta214-f22.github.io/class_activities/ca_lecture_11.html

Class activity

What is the estimated change in odds associated with a 1 point increase in hematocrit, holding temperature and abdominal pain constant?

Class activity

How does the deviance change when we add hematocrit to the model?

Class activity

Researchers want to test whether there is a relationship between hematocrit and the probability a patient has dengue, after accounting for temperature and abdominal pain. Carry out a hypothesis test to investigate this research question.

Comparing models

If deviance always decreases when I add additional variables, how can I assess whether including hematocrit substantially improves the model?

Option 1: Likelihood ratio test

- + Is the change in deviance bigger than we would expect if hematocrit doesn't really matter?

Option 2: AIC

AIC

In linear regression, what quantity did we use to compare models with different numbers of parameters?

AIC

In linear regression, what quantity did we use to compare models with different numbers of parameters?

Adjusted R^2

- + We can use something similar for logistic regression, called the *Akaike information criterion* (AIC)
- + Motivation: penalize the deviance based on the number of parameters

AIC

AIC: Suppose our model has p parameters (including the intercept). Then the AIC is

$$AIC = 2p + \text{deviance}$$

AIC

Model 1: (adding hematocrit)

```
## Null Deviance:          6956  
## Residual Deviance: 6745      AIC: 6755
```

Model 2: (no hematocrit)

```
## Null Deviance:          6956  
## Residual Deviance: 6914      AIC: 6922
```

Which model do we prefer, based on AIC?

Model comparison

Does the model improve when we add hematocrit (the proportion of red blood cells)?

- + **Likelihood ratio test:** $p\text{-value} \approx 0$
- + **AIC:** AIC is smaller when we add hematocrit

Conclusion: We have convincing evidence that adding hematocrit improves the model.

A new question...

You report your results to the hospital, and they ask a follow-up question:

How good is your model at predicting whether a patient has dengue?

Making predictions

- + For each patient in the data, we calculate $\hat{\pi}_i$
- + But, we want to decide which patients to treat. So we need to guess whether patient i has dengue ($Y_i = 1$) or doesn't ($Y_i = 0$)

How can we turn $\hat{\pi}_i$ into a dengue prediction?

Confusion matrix

		Actual	
		$Y = 0$	$Y = 1$
Predicted	$\hat{Y} = 0$	3957	1631
	$\hat{Y} = 1$	66	66

- + For 3957 patients, we correctly predicted they did not have dengue
- + For 66 patients, we correctly predicted they had dengue
- + For 1631 patients, we incorrectly predicted they did not have dengue

Did we do a good job at predicting?

Accuracy

		Actual	
		$Y = 0$	$Y = 1$
Predicted	$\hat{Y} = 0$	3957	1631
	$\hat{Y} = 1$	66	66

$$\begin{aligned}\text{Accuracy} &= \frac{\text{number of correct predictions}}{\text{number of observations}} \\ &= \frac{3957 + 66}{5720} \\ &= 0.703\end{aligned}$$

We correctly predict dengue status 70% of the time.

Class activity, Part II

https://sta214-f22.github.io/class_activities/ca_lecture_11.html

Class activity

		Actual	
		$Y = 0$	$Y = 1$
Predicted	$\hat{Y} = 0$	3957	1631
	$\hat{Y} = 1$	66	66

Are our predictions better for patients who actually have dengue, or for patients who don't have dengue?

Class activity

		Actual	
		$Y = 0$	$Y = 1$
Predicted	$\hat{Y} = 0$	3990	503
	$\hat{Y} = 1$	33	1194

What is the accuracy of the rapid test?

Class activity

Which method would you prefer -- our logistic regression model, or the rapid test?