Logistic regression with multiple predictors

Scenario: dengue fever in Vietnam

- Dengue fever: a mosquito-borne viral disease, which infects hundreds of millions of people a year. Common in tropical climates
- Researchers in Vietnam are interested in identifying the relationship between specific symptoms and the probability of having dengue

Data

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

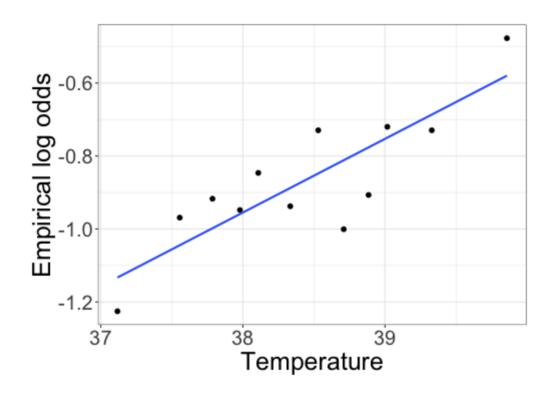
- → Dengue: whether the patient actually has dengue fever, based on a lab test (0 = no, 1 = yes)
- Temperature: patient's body temperature (in Celsius)
- Abdominal: whether the patient has abdominal pain (0 = no, 1 = yes)
- HCT: patient's hematocrit (proportion of red blood cells)
- Age: patient's age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

Research question

You are approached by the researchers to help analyze their data. Their initial question:

What is the relationship between temperature and the probability of dengue, and does the relationship differ if the patient also presents with abdominal pain?

How can we visualize the relationship between temperature and the probability of having dengue?



Based on the empirical log odds plot, what would be a reasonable model for the relationship between temperature and the probability a patient has dengue?

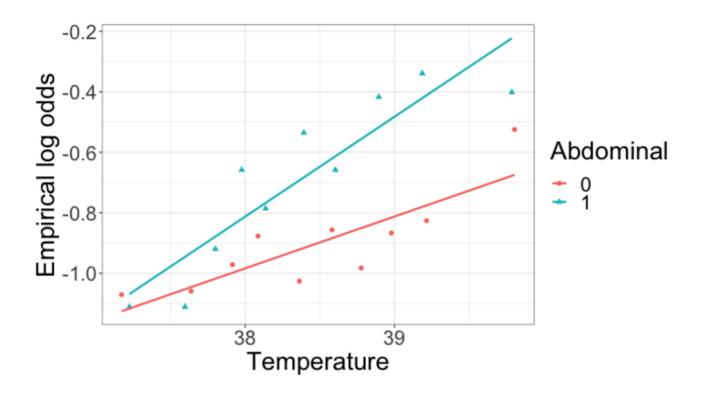
Initial model

$$Y_i \sim Bernoulli(\pi_i)$$

$$\log\!\left(rac{\pi_i}{1-\pi_i}
ight) = eta_0 + eta_1 \ Temperature_i$$

We're also interested in whether this relationship is different depending on abdominal pain.

What plot can I use to investigate this question?



Based on this plot, what would be a reasonable model that incorporates both temperature and abdominal pain?

Adding abdominal pain

$$Y_i \sim Bernoulli(\pi_i)$$

$$egin{split} \logigg(rac{\pi_i}{1-\pi_i}igg) &= eta_0 + eta_1 \ Temperature_i + eta_2 \ Abdominal_i \ &+ eta_3 \ Temperature_i \cdot Abdominal_i \end{split}$$

The interaction term allows the relationship between temperature and π_i to change depending on abdominal pain

https://sta214-f22.github.io/class_activities/ca_lecture_10.html

$$egin{split} \log\left(rac{\widehat{\pi}_i}{1-\widehat{\pi}_i}
ight) &= -7.745 + 0.178\ Temperature_i \ &- 6.129\ Abdominal_i \ &+ 0.166\ Temperature_i\cdot Abdominal_i \end{split}$$

What is the estimated probability of dengue for a patient with a temperature of 38C and abdominal pain?

$$egin{split} \log\left(rac{\widehat{\pi}_i}{1-\widehat{\pi}_i}
ight) &= -7.745 + 0.178\ Temperature_i \ &-6.129\ Abdominal_i \ &+0.166\ Temperature_i\cdot Abdominal_i \end{split}$$

For patients with abdominal pain, what is the estimated change in odds associated with an increase in temperature of 1C?

$$Y_i \sim Bernoulli(\pi_i)$$

$$egin{split} \logigg(rac{\pi_i}{1-\pi_i}igg) &= eta_0 + eta_1 \ Temperature_i + eta_2 \ Abdominal_i \ &+ eta_3 \ Temperature_i \cdot Abdominal_i \end{split}$$

I want to test whether there is a relationship between abdominal pain and the probability of dengue, after accounting for the relationship between temperature and the probability of dengue.

What are my null and alternative hypotheses?

Hypothesis testing

$$Y_i \sim Bernoulli(\pi_i)$$

$$egin{split} \logigg(rac{\pi_i}{1-\pi_i}igg) &= eta_0 + eta_1 \ Temperature_i + eta_2 \ Abdominal_i \ &+ eta_3 \ Temperature_i \cdot Abdominal_i \end{split}$$

Hypotheses:

$$H_0: \beta_2 = \beta_3 = 0$$

 H_A : at least one of $\beta_2, \beta_3 \neq 0$

Which type of test can I use to test these hypotheses (Wald test, likelihood ratio test, or either)?

Likelihood ratio test

$$Y_i \sim Bernoulli(\pi_i)$$

$$egin{split} \logigg(rac{\pi_i}{1-\pi_i}igg) &= eta_0 + eta_1 \ Temperature_i + eta_2 \ Abdominal_i \ &+ eta_3 \ Temperature_i \cdot Abdominal_i \end{split}$$

Hypotheses:

$$H_0: \beta_2 = \beta_3 = 0$$

 H_A : at least one of $\beta_2, \beta_3 \neq 0$

What are my full and reduced models?

Full and reduced models

Full model:

$$\log\!\left(rac{\pi_i}{1-\pi_i}
ight) = eta_0 + eta_1 \, Temperature_i + eta_2 \, Abdominal_i \ + eta_3 \, Temperature_i \cdot Abdominal_i$$

Reduced model:

$$\log\!\left(rac{\pi_i}{1-\pi_i}
ight) = eta_0 + eta_1 \ Temperature_i$$

Test statistic

G = deviance of reduced model - deviance of full model

Full model:

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

Reduced model:

```
## Null Deviance: 6956
## Residual Deviance: 6927 AIC: 6931
```

G =

How do I calculate a p-value for this test statistic?

p-value

$$G = 6927 - 6914 = 13 \quad \ G \sim \chi_k^2$$

where k= difference in number of parameters between full and reduced models.

What is k for this test?

p-value

$$G = 6927 - 6914 = 13 \quad \ G \sim \chi_k^2$$

where k= difference in number of parameters between full and reduced models.

```
pchisq(13, df=2, lower.tail=F)
```

[1] 0.001503439

Conclusion

- Question: Is there a relationship between abdominal pain and the probability of dengue, after accounting for the relationship between temperature and the probability of dengue?
- + Hypotheses:

$$H_0: \beta_2 = \beta_3 = 0$$

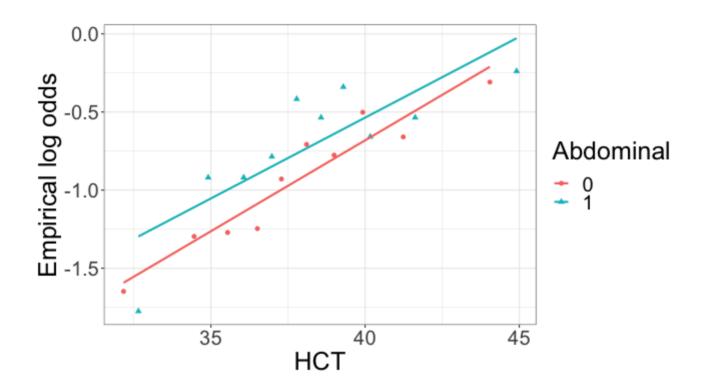
 H_A : at least one of $\beta_2, \beta_3 \neq 0$

- **p-value:** 0.0015
- Conclusion: There is strong evidence that there is a relationship between abdominal pain and the probability of dengue, after accounting for the relationship between temperature and the probability of dengue.

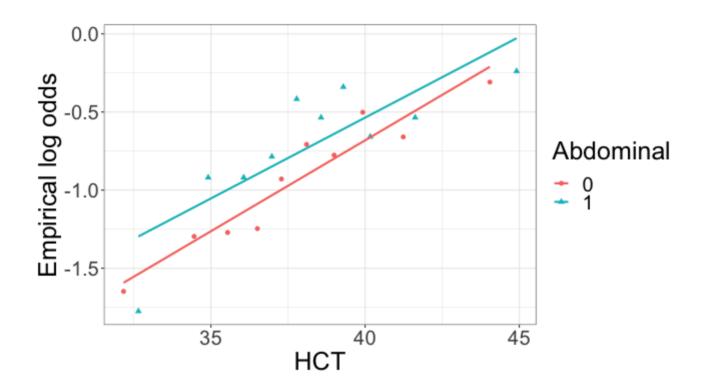
A new question...

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

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



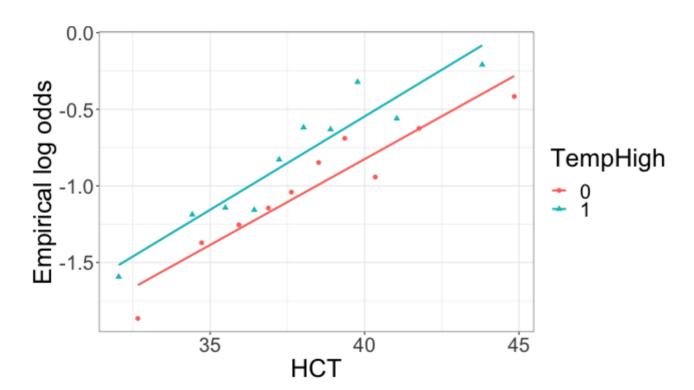
Does it look like we need an interaction between hematocrit (HCT) and abdominal pain?

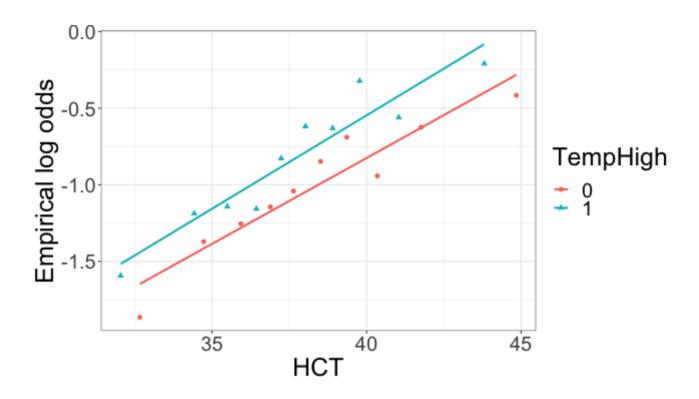


How can I check whether there might be an interaction between temperature and hematocrit?

Define TempHigh by

- $extbf{+} TempHigh_i = 1 \text{ if } Temperature_i > 38$
- $+ TempHigh_i = 0 \text{ if } Temperature_i <= 38$





Does it look like we need an interaction between temperature and hematocrit?

Model

$$Y_i \sim Bernoulli(\pi_i)$$

$$egin{split} \logigg(rac{\pi_i}{1-\pi_i}igg) &= eta_0 + eta_1 \ Temperature_i + eta_2 \ Abdominal_i \ &+ eta_3 \ Temperature_i \cdot Abdominal_i \ &+ eta_4 \ HCT_i \end{split}$$

 Note that while we binarized temperature for EDA, we use the original variable in the model here