# Lecture 1

Recall the dengue data from last semester:

- Data on Vietnamese children admitted to hospital with possible dengue fever
- Variables include:
  - Age
  - White blood cell count (WBC)
  - Platelet count (PLT)
  - Dengue status ( $\emptyset$  = no dengue, 1 = dengue)

I want to model dengue status, with Age, WBC, and PLT as explanatory variables.

What does my model look like?

$$Y_i \sim Bernoulli(p_i)$$

$$\log\left(\frac{p_i}{1-p_i}\right) = \beta_0 + \beta_1 Age_i + \beta_2 WBC_i + \beta_3 PLT_i$$

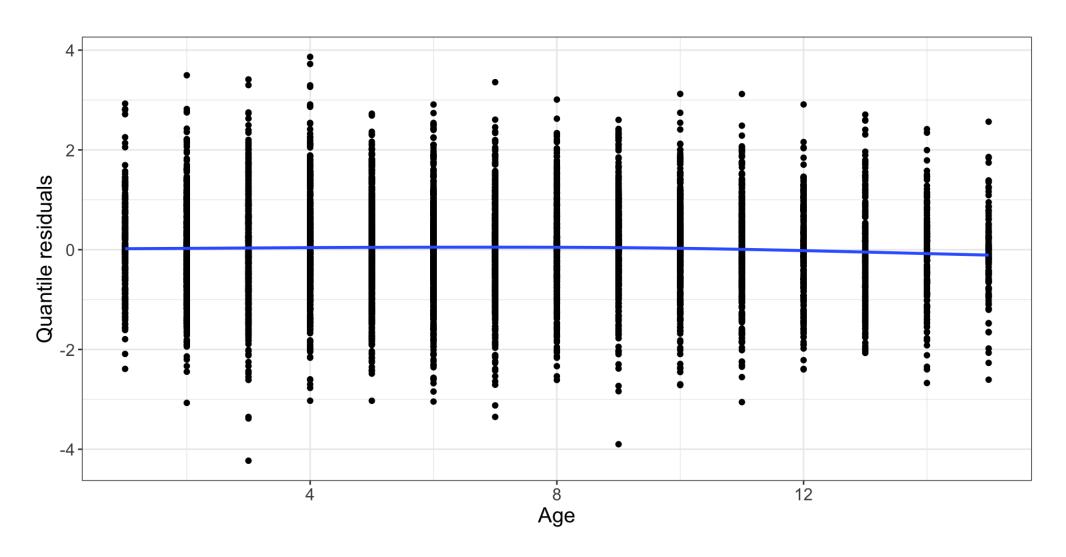
**Question:** How do I interpret a regression coefficient (e.g.  $\beta_1$ )?

$$Y_i \sim Bernoulli(p_i)$$

$$\log\left(\frac{p_i}{1-p_i}\right) = \beta_0 + \beta_1 Age_i + \beta_2 WBC_i + \beta_3 PLT_i$$

Question: What assumptions does this model make?

### Quantile residual plot:



$$Y_i \sim Bernoulli(p_i)$$

$$\log\left(\frac{p_i}{1-p_i}\right) = \beta_0 + \beta_1 Age_i + \beta_2 WBC_i + \beta_3 PLT_i$$

**Question:** I want to know whether there is relationship between Age and Dengue status, after accounting for WBC and PLT. How can I address this question?

```
1 m1 <- glm(Dengue ~ Age + WBC + PLT, data = dengue,
            family = binomial)
 3 summary(m1)
Call:
glm(formula = Dengue ~ Age + WBC + PLT, family = binomial, data =
dengue)
Coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept) 1.2525593 0.1548038 8.091 5.9e-16 ***
    0.1383186 \quad 0.0099763 \quad 13.865 \quad < 2e-16 ***
Age
WBC
    -0.2523294 0.0135371 -18.640 < 2e-16 ***
PLT -0.0060276 0.0006113 -9.860 < 2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

$$Y_i \sim Bernoulli(p_i)$$

$$\log\left(\frac{p_i}{1-p_i}\right) = \beta_0 + \beta_1 Age_i + \beta_2 WBC_i + \beta_3 PLT_i$$

Question: Suppose now I want to test  $H_0: \beta_2 = \beta_3 = 0$ . How do I carry out the likelihood ratio test?

```
1 m1 <- glm(Dengue ~ Age + WBC + PLT, data = dengue, family = binomial)
2 m1$deviance
[1] 5200.823

1 m2 <- glm(Dengue ~ Age, data = dengue, family = binomial)
2 m2$deviance
[1] 6272.458</pre>
```

#### Test statistic =

```
1 m1 <- glm(Dengue ~ Age + WBC + PLT, data = dengue, family = binomial)
2 m1$deviance
[1] 5200.823

1 m2 <- glm(Dengue ~ Age, data = dengue, family = binomial)
2 m2$deviance
[1] 6272.458</pre>
```

Test statistic =  $deviance_{reduced}$  -  $deviance_{full}$  = 1071.6 How do I calculate a p-value?

```
1 m1 <- glm(Dengue ~ Age + WBC + PLT, data = dengue, family = binomial)
2 m1$deviance
[1] 5200.823

1 m2 <- glm(Dengue ~ Age, data = dengue, family = binomial)
2 m2$deviance
[1] 6272.458</pre>
```

### Test statistic = deviance<sub>reduced</sub> - deviance<sub>full</sub> = 1071.6

```
1 pchisq(1071.6, df=2, lower.tail=F)
[1] 2.018443e-233
```

$$Y_i \sim Bernoulli(p_i)$$

$$\log\left(\frac{p_i}{1-p_i}\right) = \beta_0 + \beta_1 Age_i + \beta_2 WBC_i + \beta_3 PLT_i$$

**Question:** The researchers are interested in whether their model does a good job identifying patients with dengue. Do our hypothesis tests address that question?

### Rough course plan

- Logistic regression recap
- Prediction and model selection
- Supplementary skills (research papers, SAPs, simulation)
- Poisson regression and EDMs
- Mis-specified models (overdispersion, zero-inflation, etc.)
- Correlated data

### Course components

- Homework assignments (graded on completion)
- Challenge assignments (graded on mastery)
- Data analysis projects (graded on mastery)
- Semester research project (graded on mastery)
  - Group project
  - Involves written report, final presentation, and intermediate check-points
  - Due next Monday: group members and tentative topic

### Reading a research paper

Research papers in the sciences and social sciences typically contain:

- Abstract
- Introduction
- Methods
- Results
- Discussion
- Conclusion

### Reading a research paper

- Abstract: overview and key points
- Introduction: motivation, background, overview of work
- Methods: details on study design, data, statistical analysis
- **Results:** summary of results, including figures, tables, p-values, etc.
- Discussion: discussion of results in context of research question
- Conclusion: short summary of paper and key results;
   connection to broader research

### Class activity

Reading the original dengue paper:

https://sta712-f23.github.io/class\_activities/ca\_1.pdf

For next class:

- finish reading the paper and working through the class activity
- we will discuss the paper on Wednesday