

# Linear and logistic regression

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## Last time: parameter estimation for linear regression

$$\begin{bmatrix} Y_1 \\ Y_2 \\ \vdots \\ Y_n \end{bmatrix} = \begin{bmatrix} 1 & X_{11} & X_{12} & \cdots & X_{1k} \\ 1 & X_{21} & X_{22} & \cdots & X_{2k} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & X_{n1} & X_{n2} & \cdots & X_{nk} \end{bmatrix} \begin{bmatrix} \beta_0 \\ \vdots \\ \beta_k \end{bmatrix} + \begin{bmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \vdots \\ \varepsilon_n \end{bmatrix}$$

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon}$$

$$SSE(\boldsymbol{\beta}) = (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})^T (\mathbf{y} - \mathbf{X}\boldsymbol{\beta})$$

$$\frac{\partial}{\partial \boldsymbol{\beta}} SSE(\boldsymbol{\beta}) = -2 \mathbf{X}^T (\mathbf{y} - \mathbf{X}\boldsymbol{\beta}) \stackrel{\text{set}}{=} \mathbf{0}$$

## Parameter estimation for linear regression

$$\frac{\partial}{\partial \beta} SSE(\beta) = -2\mathbf{X}^T(\mathbf{y} - \mathbf{X}\beta) \stackrel{\text{set}}{=} \mathbf{0}$$

$$\Rightarrow \mathbf{X}^T(\mathbf{y} - \mathbf{X}\beta) = \mathbf{0}$$

$$\Rightarrow \mathbf{X}^T\mathbf{y} - \mathbf{X}^T\mathbf{X}\beta = \mathbf{0}$$

$$\Rightarrow \mathbf{X}^T\mathbf{X}\beta = \mathbf{X}^T\mathbf{y}$$

$$\Rightarrow \hat{\beta} = (\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T\mathbf{y}$$

$\hat{\beta}$   
least squares estimator of  $\beta$

# Warmup

Work on the warmup activity:

[https://sta711-s26.github.io/class\\_activities/ca\\_02.html](https://sta711-s26.github.io/class_activities/ca_02.html)

Submit your work on Canvas.

## Warmup

```
library(Stat2Data)
data("FirstYearGPA")

lm(GPA ~ HSGPA + SATM, data = FirstYearGPA) |> coef()
```

```
##      (Intercept)          HSGPA          SATM
## 0.7579761649 0.5305150632 0.0007984613
```

```
y <- FirstYearGPA$GPA
X <- cbind(1, FirstYearGPA$HSGPA, FirstYearGPA$SATM)
solve(t(X) %*% X) %*% t(X) %*% y
```

```
##              [,1]
## [1,] 0.7579761649
## [2,] 0.5305150632
## [3,] 0.0007984613
```

## Regression assumptions

$$\text{GPA}_i = \beta_0 + \beta_1 \text{HSGPA}_i + \beta_2 \text{SATM}_i + \varepsilon_i$$

**Question:** What assumptions do we often make when fitting a linear regression model?

# Regression assumptions

$$\text{GPA}_i = \beta_0 + \beta_1 \text{HSGPA}_i + \beta_2 \text{SATM}_i + \varepsilon_i$$

Some common assumptions:

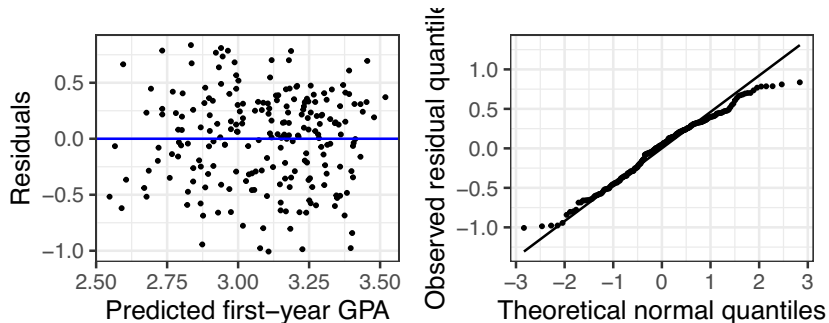
$$\varepsilon_i \stackrel{\text{iid}}{\sim} N(0, \sigma^2)$$

- ▶ Shape
- ▶ Constant variance (variance of  $\varepsilon_i$  is the same for all observations)
- ▶ Normality ( $\varepsilon_i$  comes from a normal distribution)
- ▶ Independence

**Question:** How do we assess these assumptions?

- normality : Q-Q plot
- independence: think about how data were generated
- shape & constant variance: residual plot

## Diagnostic plots



**Question:** Do the regression assumptions seem reasonable here?

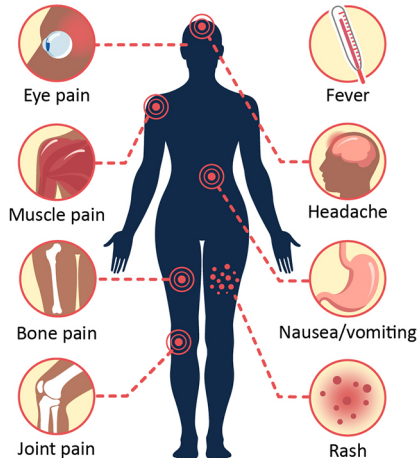


## Another motivating example: Dengue fever

**Dengue fever:** a mosquito-borne viral disease affecting 400 million people a year

### Dengue Symptoms

Fever with any of the following



## Motivating example: Dengue data

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

- ▶ *Sex*: patient's sex (female or male)
- ▶ *Age*: patient's age (in years)
- ▶ *WBC*: white blood cell count
- ▶ *PLT*: platelet count
- ▶ other diagnostic variables. . .
- ▶ *Dengue*: whether the patient has dengue (0 = no, 1 = yes)

**Research goal:** Predict dengue status using diagnostic measurements

## Fitting a model: initial attempt

What if we try a linear regression model?

$Y_i$  = dengue status of  $i$ th patient  
(1 = dengue, 0 = no dengue)

$$Y_i = \beta_0 + \beta_1 WBC_i + \varepsilon_i \quad \varepsilon_i \stackrel{iid}{\sim} N(0, \sigma^2)$$

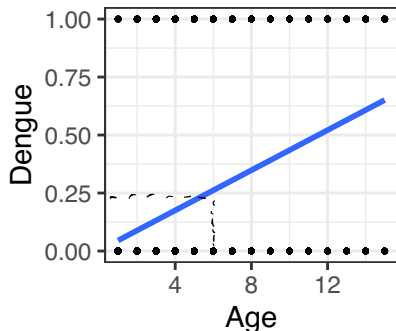
What are some potential issues with this linear regression model?

$$Y_i \in \{0, 1\}$$

$$\beta_0 + \beta_1 WBC_i + \varepsilon_i \in (-\infty, \infty) \quad (\text{potentially})$$

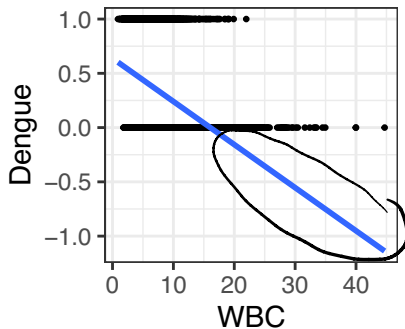
(continuous & possibly unbounded)

## Don't fit linear regression with a binary response



Age=6  $\rightarrow$   $\hat{\text{Dengue}} = 0.25$

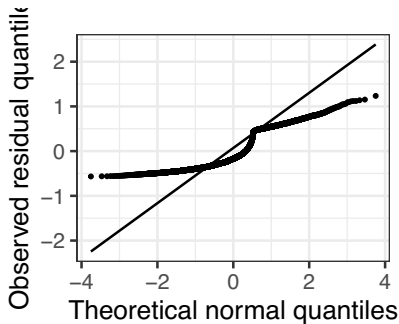
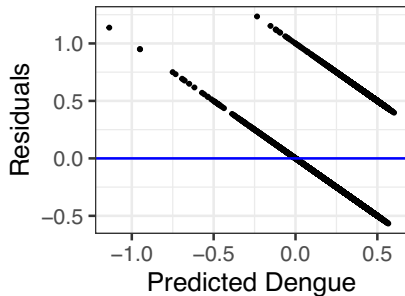
maybe this is a  
25% chance of  
having Dengue?



if  $\text{WBC} > 15$ ,  
 $\hat{\text{Dengue}} < 0$  ??

# Don't fit linear regression with a binary response

Diagnostic plots:



## Second attempt

$$\text{if } v \sim N(\mu, \sigma^2) \quad a, b \in \mathbb{R} \\ a + bv \sim N(a + b\mu, b^2\sigma^2)$$

Let's rewrite the linear regression model:

$$y_i = \beta_0 + \beta_1 \text{WBC}_i + \varepsilon_i \quad \varepsilon_i \stackrel{\text{iid}}{\sim} N(0, \sigma^2)$$

$$\Rightarrow y_i | \text{WBC}_i \sim N(\beta_0 + \beta_1 \text{WBC}_i, \sigma^2)$$

or written in two pieces:

describes distribution of  $y_i | \text{WBC}_i$   $\rightarrow y_i | \text{WBC}_i \sim N(\mu_i, \sigma^2)$  (random component)

$\mu_i = \beta_0 + \beta_1 \text{WBC}_i$  (systematic component)

specifies how distribution depends on  $\text{WBC}_i$

Problem:  $y_i \in \{0, 1\} \Rightarrow y_i | \text{WBC}_i$  is not normal!  
use Bernoulli instead!