# Likelihood ratio tests

### Last time

Data on the RMS *Titanic* disaster. We have data on 891 passengers on the ship, with the following variables:

- Passenger: A unique ID number for each passenger.
- Survived: An indicator for whether the passenger survived (1) or perished (0) during the disaster.
- Pclass: Indicator for the class of the ticket held by this passengers; 1 = 1st class, 2 = 2nd class, 3 = 3rd class.
- Sex: Binary Indicator for the biological sex of the passenger.
- Age: Age of the passenger in years; Age is fractional if the passenger was less than 1 year old.
- Fare: How much the ticket cost in US dollars.
- + + others

### Last time

Is there a relationship between passenger age and their probability of survival, after accounting for sex, passenger class, and the cost of their ticket?

$$egin{split} \logigg(rac{p_i}{1-p_i}igg) &= eta_0 + eta_1Age_i + eta_2Sex_i + \ eta_3Age_i \cdot Sex_i + eta_4\log(Fare_i+1) \end{split}$$

What hypotheses should we test to investigate this research question?

#### Likelihood ratio tests

```
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
  (Intercept) -1.40695
                          0.44682 -3.149 0.00164 **
         0.01107
                          0.01107 1.000 0.31730
## Age
## Sexmale -1.27467
                          0.41654 - 3.060 0.00221 **
## log(Fare + 1) 0.69449
                          0.11065 6.276 3.47e-10
                                                 ***
  Age:Sexmale -0.03638
                          0.01378 - 2.639
                                         0.00831 **
##
      Null deviance: 964.52 on 713 degrees of freedom
##
## Residual deviance: 697.21 on 709 degrees of freedom
```

What information replaces  ${\cal R}^2$  and  ${\cal R}^2_{adj}$  in the GLM output?

### **Deviance**

**Definition:** The *deviance* of a fitted model with parameter estimates  $\widehat{\beta}$  is given by

$$2\ell( ext{saturated model}) - 2\ell(\widehat{eta})$$

### Residual and null deviance

```
m1 <- glm(Survived ~ Age*Sex + log(Fare + 1),</pre>
          data = titanic, family = binomial)
summary(m1)
      Null deviance: 964.52 on 713 degrees of freedom
##
## Residual deviance: 697.21 on 709 degrees of freedom
m2 <- glm(Survived ~ Sex + log(Fare + 1),
          data = titanic, family = binomial)
summary(m2)
##
      Null deviance: 964.52 on 713 degrees of freedom
## Residual deviance: 708.04 on 711 degrees of freedom
                                                      7/11
```

Full model:

Hypotheses:

Reduced model:

**Test statistic:** 

#### Full model:

$$egin{split} \logigg(rac{p_i}{1-p_i}igg) &= eta_0 + eta_1 Sex_i + eta_2 \log(Fare_i+1) + \ eta_3 Age_i + eta_4 Age_i \cdot Sex_i \end{split}$$

#### Reduced model:

$$\logigg(rac{p_i}{1-p_i}igg) = eta_0 + eta_1 Sex_i + eta_2 \log(Fare_i+1)$$

$$G = 2\ell(\widehat{eta}) - 2\ell(\widehat{eta}^0)$$

Why is G always  $\geq 0$ ?

#### Full model:

$$egin{split} \logigg(rac{p_i}{1-p_i}igg) &= eta_0 + eta_1 Sex_i + eta_2 \log(Fare_i+1) + \ eta_3 Age_i + eta_4 Age_i \cdot Sex_i \end{split}$$

#### Reduced model:

$$\logigg(rac{p_i}{1-p_i}igg) = eta_0 + eta_1 Sex_i + eta_2 \log(Fare_i+1)$$

$$G = 2\ell(\widehat{eta}) - 2\ell(\widehat{eta}^0) = 10.83$$

If the reduced model is correct, how unusual is G=10.83?

## Likelihood ratio test