

Logistic regression

Inference & Model selection

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Risk of coronary heart disease

This dataset is from an ongoing cardiovascular study on residents of the town of Framingham, Massachusetts. We want to predict if a patient has a high risk of getting coronary heart disease in the next 10 years.

Response:

TenYearCHD:

- 0 = Patient is not high risk of having coronary heart disease in the next 10 years
- 1 = Patient is high risk of having coronary heart disease in the next 10 years

Predictors:

- **age**: Age at exam time.
- **currentSmoker**: 0 = nonsmoker; 1 = smoker
- **totChol**: total cholesterol (mg/dL)

Modeling risk of coronary heart disease

term	estimate	std.error	statistic	p.value	conf.low	conf.high
(Intercept)	-2.111	0.077	-27.519	0.000	-2.264	-1.963
ageCent	0.081	0.006	13.477	0.000	0.070	0.093
currentSmoker1	0.447	0.099	4.537	0.000	0.255	0.641
totCholCent	0.003	0.001	2.339	0.019	0.000	0.005

Hypothesis test for β_j

$$\log \left(\frac{\hat{\pi}}{1 - \hat{\pi}} \right) = \hat{\beta}_0 + \hat{\beta}_1 x_1 + \hat{\beta}_2 x_2 + \cdots + \hat{\beta}_p x_p$$

The test of significance for the coefficient β_j is

Hypotheses: $H_0 : \beta_j = 0$ vs $H_a : \beta_j \neq 0$

Test Statistic:

$$z = \frac{\hat{\beta}_j - 0}{SE(\hat{\beta}_j)}$$

P-value: $P(|Z| > |z|)$,

where $Z \sim N(0, 1)$, the Standard Normal distribution

Confidence interval for β_j

- We can calculate the **C\% confidence interval** for β_j using the following:

$$\hat{\beta}_j \pm z^* SE(\hat{\beta}_j)$$

where z^* is calculated from the $N(0, 1)$ distribution

We are $C\%$ confident that for every one unit change in x_j , the odds multiply by a factor of $\exp\{\hat{\beta}_j - z^* SE(\hat{\beta}_j)\}$ to $\exp\{\hat{\beta}_j + z^* SE(\hat{\beta}_j)\}$, holding all else constant.

Modeling risk of coronary heart disease

term	estimate	std.error	statistic	p.value	conf.low	conf.high
(Intercept)	-2.11059	0.07670	-27.51865	0.00000	-2.26360	-1.96285
ageCent	0.08150	0.00605	13.47651	0.00000	0.06973	0.09344
currentSmoker1	0.44743	0.09862	4.53705	0.00001	0.25461	0.64134
totCholCent	0.00254	0.00108	2.33919	0.01933	0.00040	0.00465

1. How is the test statistic for `currentSmoker1` calculated?
2. Is `totCholCent` a statistically significant predictor of the odds a person is high risk for coronary heart disease?
 - Justify your answer using the test statistic and p-value.
 - Justify your answer using the confidence interval.

Model Selection

Comparing Nested Models

- Suppose there are two models:
 - Reduced Model includes predictors $x_1, \dots, x_q$
 - Full Model includes predictors $x_1, \dots, x_q, x_{q+1}, \dots, x_p$
- We want to test the hypotheses

$$H_0 : \beta_{q+1} = \dots = \beta_p = 0$$

$$H_a : \text{at least 1 } \beta_j \text{ is not } 0$$

- To do so, we will use the **Drop-in-Deviance test** (also known as the Nested Likelihood test)

Deviance residual

- The **deviance residual** is a measure of how much the observed data differs from what is measured using the likelihood ratio
- The deviance residual for the i^{th} observation is

$$d_i = \text{sign}(y_i - \hat{\pi}_i) \sqrt{2 \left[y_i \log \left(\frac{y_i}{\hat{\pi}_i} \right) + (1 - y_i) \log \left(\frac{1 - y_i}{1 - \hat{\pi}_i} \right) \right]}$$

where $\text{sign}(y_i - \hat{\pi}_i)$ is positive when $y_i = 1$ and negative when $y_i = 0$.

Drop-in-Deviance Test

- The **deviance statistic for Model k** is $G_k^2 = \sum_{i=1}^n d_i^2$
- To test the hypotheses

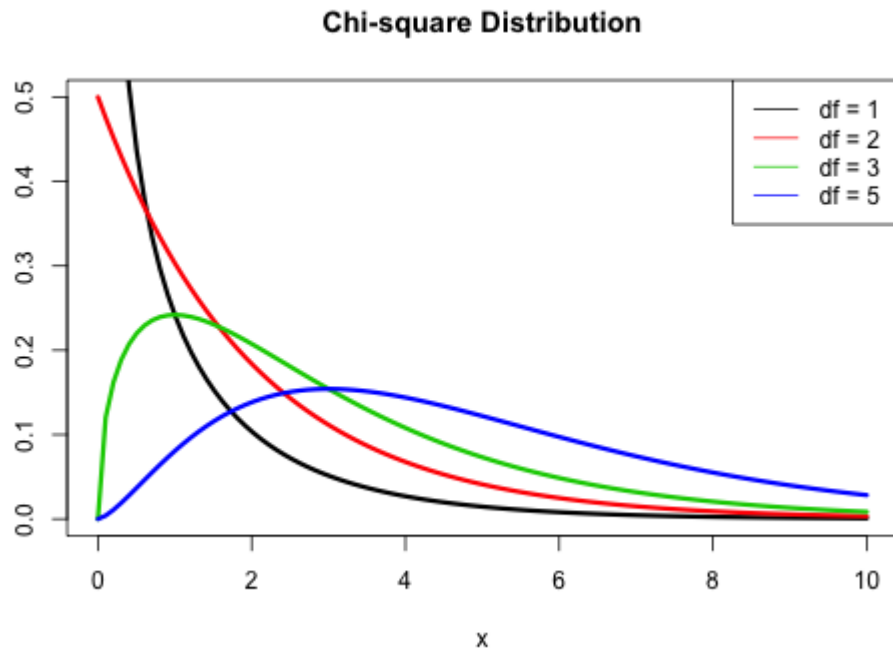
$$H_0 : \beta_{q+1} = \cdots = \beta_p = 0$$

$$H_a : \text{at least one } \beta_j \text{ is not } 0$$

the **drop-in-deviance statistic** is $G_{reduced}^2 - G_{full}^2$

The p-value for the test is calculated using a Chi-square distribution (χ^2) with degrees of freedom equal to the difference in the number of parameters in the full and reduced models

χ^2 distribution



Calculate p-value for the drop-in-deviance test as $P(\chi^2 > \text{test statistic})$

Should we add education to the model?

```
model_red <- glm(TenYearCHD ~ ageCent + currentSmoker + totChol,  
                 data = heart_data, family = binomial)  
model_full <- glm(TenYearCHD ~ ageCent + currentSmoker + totChol +  
                 education,  
                 data = heart_data, family = binomial)
```

term	estimate	std.error	statistic	p.value	conf.low	conf.high
(Intercept)	-2.617	0.277	-9.448	0.000	-3.161	-2.074
ageCent	0.078	0.006	12.701	0.000	0.066	0.091
currentSmoker1	0.449	0.099	4.540	0.000	0.255	0.643
totChol	0.003	0.001	2.503	0.012	0.001	0.005
education2	-0.266	0.119	-2.233	0.026	-0.502	-0.034
education3	-0.319	0.144	-2.215	0.027	-0.607	-0.041
education4	-0.116	0.160	-0.725	0.468	-0.436	0.192

Should we add education to the model?

```
# Deviances  
(dev_red <- glance(model_red)$deviance)
```

```
## [1] 2894.989
```

```
(dev_full <- glance(model_full)$deviance)
```

```
## [1] 2887.206
```

```
# Drop-in-deviance test statistic  
(test_stat <- dev_red - dev_full)
```

```
## [1] 7.783615
```

Should we add education to the model?

```
# p-value  
1 - pchisq(test_stat, 3) #3 = number of new model terms in model2  
  
## [1] 0.05070196
```

What is your conclusion for the test?

Drop-in-Deviance test in R

- We can use the **anova** function to conduct this test
 - Add **test = "Chisq"** to conduct the drop-in-deviance test

```
anova(model_red, model_full, test = "Chisq")
```

```
## Analysis of Deviance Table
##
## Model 1: TenYearCHD ~ ageCent + currentSmoker + totChol
## Model 2: TenYearCHD ~ ageCent + currentSmoker + totChol + education
##   Resid. Df Resid. Dev Df Deviance Pr(>Chi)
## 1      3654      2895.0
## 2      3651      2887.2  3    7.7836  0.0507 .
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```


Model selection

- Use AIC or BIC for model selection

$$AIC = -2 * L - n \log(n) + 2(p + 1)$$

$$BIC = -2 * L - n \log(n) + \log(n) \times (p + 1)$$

where $L = \sum_{i=1}^n [y_i \log(\hat{\pi}_i) + (1 - y_i) \log(1 - \hat{\pi}_i)]$

AIC from glance function

Let's look at the AIC for the model that includes ageCent, currentSmoker, and totCholCent

```
glance(model_red)$AIC
```

```
## [1] 2902.989
```

Calculating AIC

```
(L <- glance(model_red)$logLik)
```

```
## [1] -1447.495
```

```
- 2 * L + 2 * (3 + 1)
```

```
## [1] 2902.989
```

Should we add education to the model?

Recall:

- Reduced Model includes AgeCent, currentSmoker, totCholCent
- Full Model includes AgeCent, currentSmoker, totCholCent, education (categorical)

```
glance(model_red)$AIC
```

```
## [1] 2902.989
```

```
glance(model_full)$AIC
```

```
## [1] 2901.206
```

Based on the AIC, which model would you choose?

**What remaining questions do you have
about logistic regression?**