

Lecture 18

Logistic Regression

STAT 8020 Statistical Methods II
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Logistic Regression
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18.1

Notes

A Motivating Example: Horseshoe Crab Mating [Brockmann, 1996, Agresti, 2013]



| sat | y | weight | width |
|-----|---|--------|-------|
| 8 | 1 | 3.05 | 28.3 |
| 0 | 0 | 1.55 | 22.5 |
| 9 | 1 | 2.30 | 26.0 |
| 0 | 0 | 2.10 | 24.8 |
| 4 | 1 | 2.60 | 26.0 |
| 0 | 0 | 2.10 | 23.8 |
| 0 | 0 | 2.35 | 26.5 |
| 0 | 0 | 1.90 | 24.7 |
| 0 | 0 | 1.95 | 23.7 |
| 0 | 0 | 2.15 | 25.6 |

Source: <https://www.britannica.com/story/horseshoe-crab-a-key-player-in-ecology-medicine-and-more>

In the rest of today's lecture, we are going to use this data set to illustrate **logistic regression**. The response variable is y : whether there are males clustering around the female

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Notes

Logistic Regression

Let $P(Y = 1) = \pi \in [0, 1]$, and x be the predictor (weight in the previous example). In SLR we have

$$\pi(x) = \beta_0 + \beta_1 x,$$

which will lead to invalid estimate of π (i.e., > 1 or < 0).

Logistic Regression

$$\log\left(\frac{\pi}{1-\pi}\right) = \beta_0 + \beta_1 x.$$

- $\log\left(\frac{\pi}{1-\pi}\right)$: the log-odds or the logit
- $\pi(x) = \frac{e^{\beta_0 + \beta_1 x}}{1 + e^{\beta_0 + \beta_1 x}} \in (0, 1)$

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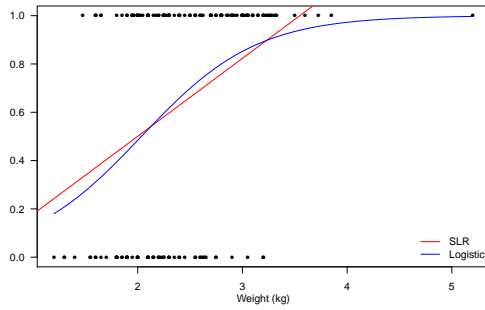
18.3

Notes

Logistic Regression Fit

$$\hat{\pi}(x) = \frac{e^{\hat{\beta}_0 + \hat{\beta}_1 x}}{1 + e^{\hat{\beta}_0 + \hat{\beta}_1 x}}, \quad \hat{\beta}_0 = -3.6947(0.8802),$$

$$\hat{\beta}_1 = 1.8151(0.3767).$$



Notes

Properties

- Similar to SLR, Sign of β_1 indicates whether $\pi(x) \uparrow$ or \downarrow as $x \uparrow$
- If $\beta_1 = 0$, then $\pi(x) = e^{\beta_0} / (1 + e^{\beta_0})$ is a constant w.r.t x (i.e., π does not depend on x)
- Curve can be approximated at fixed x by straight line to describe rate of change: $\frac{d\pi(x)}{dx} = \beta_1 \pi(x)(1 - \pi(x))$
- $\pi(-\beta_0/\beta_1) = 0.5$, and $1/\beta_1 \approx$ the distance of x values with $\pi(x) = 0.5$ and $\pi(x) = 0.75$ (or $\pi(x) = 0.25$)



Notes

Odds Ratio Interpretation

Recall $\log\left(\frac{\pi(x)}{1-\pi(x)}\right) = \beta_0 + \beta_1 x$, we have the odds

$$\frac{\pi(x)}{1 - \pi(x)} = \exp(\beta_0 + \beta_1 x).$$

If we increase x by 1 unit, the the odds becomes

$$\exp(\beta_0 + \beta_1(x + 1)) = \exp(\beta_1) \times \exp(\beta_0 + \beta_1 x).$$

$$\Rightarrow \frac{\text{Odds at } x+1}{\text{Odds at } x} = \exp(\beta_1), \forall x$$

Example: In the horseshoe crab example, we have $\hat{\beta}_1 = 1.8151 \Rightarrow e^{1.8151} = 6.14 \Rightarrow$ **Estimated odds of satellite multiply by 6.1 for 1 kg increase in weight.**



Notes

Parameter Estimation

In logistic regression we use **maximum likelihood estimation** to estimate the parameters:

- **Statistical model:** $Y_i \sim \text{Bernoulli}(\pi(x_i))$ where
$$\pi(x_i) = \frac{\exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)}.$$

- **Likelihood function:** We can write the joint probability density of the data $\{x_i, y_i\}_{i=1}^n$ as

$$\prod_{i=1}^n \pi(x_i)^{y_i} (1 - \pi(x_i))^{(1-y_i)}.$$

We treat this as a function of parameters (β_0, β_1) given data.

- **Maximum likelihood estimate:** The maximizer $\hat{\beta}_0, \hat{\beta}_1$ is the maximum likelihood estimate (MLE). This maximization can only be solved numerically.



Notes

Horseshoe Crab Logistic Regression Fit

```
> logitFit <- glm(y ~ weight, data = crab, family = "binomial")
> summary(logitFit)
```

```
Call:
glm(formula = y ~ weight, family = "binomial", data = crab)
```

```
Deviance Residuals:
    Min       1Q   Median       3Q      Max
-2.1108  -1.0749   0.5426   0.9122   1.6285
```

```
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept)  -3.6947     0.8802  -4.198 2.70e-05 ***
weight        1.8151     0.3767   4.819 1.45e-06 ***
---
Signif. codes:
  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
(Dispersion parameter for binomial family taken to be 1)
```

```
Null deviance: 225.76 on 172 degrees of freedom
Residual deviance: 195.74 on 171 degrees of freedom
AIC: 199.74
```

```
Number of Fisher Scoring iterations: 4
```



Notes

Inference: Confidence Interval

A 95% confidence interval of the parameter β_i is

$$\hat{\beta}_i \pm z_{0.025} \times \text{SE}_{\hat{\beta}_i}, \quad i = 0, 1$$

Horseshoe Crab Example

A 95% (Wald) confidence interval of β_1 is

$$1.8151 \pm 1.96 \times 0.3767 = [1.077, 2.553]$$

Therefore a 95% CI of e^{β_1} , the **multiplicative effect** on odds of 1-unit increase in x , is

$$[e^{1.077}, e^{2.553}] = [2.94, 12.85]$$



Notes

Inference: Hypothesis Test

Null and Alternative Hypotheses:

$H_0 : \beta_1 = 0 \Rightarrow Y$ is independent of $X \Rightarrow \pi(x)$ is a constant

$H_a : \beta_1 \neq 0$

Test Statistics:

$$z_{obs} = \frac{\hat{\beta}_1}{SE_{\hat{\beta}_1}} = \frac{1.8151}{0.3767} = 4.819.$$

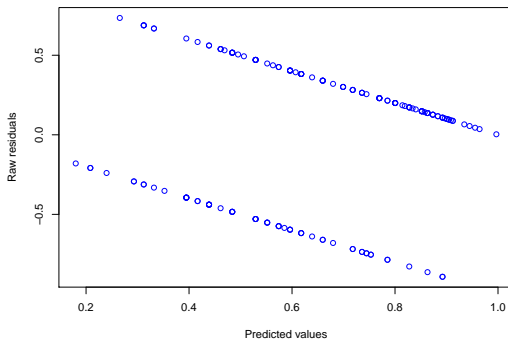
P-value = 1.45×10^{-6}

We have sufficient evidence that `weight` has positive effect on π , the probability of having satellite male horseshoe crabs



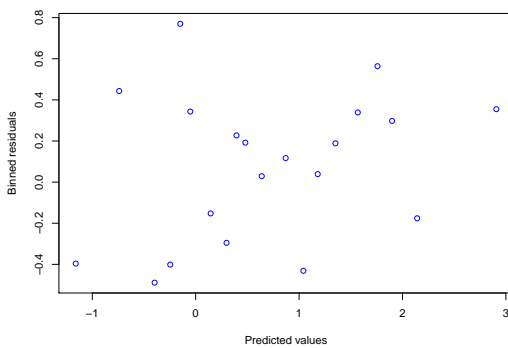
Notes

Diagnostic: Raw Residual Plot



Notes

Diagnostic: Binned Residual Plot



Notes

Model Selection

```
> logitFit2 <- glm(y ~ weight + width, data = crab, family = "binomial")
> step(logitFit2)
Start: AIC=198.89
y ~ weight + width

      Df Deviance   AIC
- weight 1  194.45 198.45
<none>    192.89 198.89
- width  1  195.74 199.74

Step: AIC=198.45
y ~ width

      Df Deviance   AIC
<none>    194.45 198.45
- width  1  225.76 227.76

Call: glm(formula = y ~ width, family = "binomial", data = crab)

Coefficients:
(Intercept)      width
   -12.3508      0.4972

Degrees of Freedom: 172 Total (i.e. Null); 171 Residual
Null Deviance:      225.8
Residual Deviance: 194.5      AIC: 198.5
```



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