第四章 二项分布的统计推断

The binomial trial

- The setup:
 - 1. There are a fixed number of observations in the sample,m.
 - The m observations are all independent.
 - There are only two possible outcomes for each observation: success(1)or failure(0).
 - The probability of success,p,is constant for each observation.
- Then,the number(count)of success,Y,has a Binomial distribution with parameters m and p. We say

$$Y \sim B(m, p)$$



The binomial distribution

The pmf of Y at a value y is given by

$$f_Y(y) = {m \choose y} p^y (1-p)^{m-y}.$$

The moment generating function of Y is

$$M_{Y}(t) = (1 - p + pe^{t})^{m}$$

• Thus, the mean is

$$E(Y) = mp$$
,

and the variance is

$$var(Y) = mp(1 - p).$$



Large sample normal approximations

Asymptotically for large m, we have that

$$\frac{Y - E(Y)}{\sqrt{var(Y)}} = \frac{Y - mp}{\sqrt{mp(1 - p)}},$$

has a N(0,1) distribution, for any fixed value of p.

• Proof: Use the standard central limit theorem noting that $Y/m = \sum_{i=1}^{m} W_i/m$, where W_i are independent B(1,p) RVs (Bernoulli RVs).

Large sample normal approximations

a better approximation for smaller values of m is to use the continuity correction:

We use the fact that

$$Pr(Y \leq y) \rightarrow \Phi(\frac{y - mp + \frac{1}{2}}{\sqrt{mp(1-p)}})$$

and

$$Pr(Y \ge y) \rightarrow 1 - \Phi(\frac{y - mp - \frac{1}{2}}{\sqrt{mp(1 - p)}})$$

as $m \to \infty$ for **fixed** p.

思考题: 利用矩母函数证明当 $n \to \infty$ 时, 二项分布可以利用正态分布近似.

证明: 假定X 服从二项分布B(n,p), 则其矩母函数为

$$M_X(t) = E[e^{tX}] = \sum_{x=0}^{n} C_n^x \rho^x q^{n-x} e^{tx}$$

$$= \sum_{x=0}^{n} C_n^x (\rho e^t)^x q^{n-x}$$

$$= (q + \rho e^t)^n \sum_{x=0}^{n} C_n^x (\frac{\rho e^t}{q + \rho e^t})^x (\frac{q}{q + \rho e^t})^{n-x}$$

$$= (q + \rho e^t)^n$$

令 $\mathbf{Y} = \frac{\mathbf{X} - \mathbf{np}}{\sqrt{\mathbf{npq}}}$, 可以得到 \mathbf{Y} 的矩母函数

$$\begin{aligned} M_{Y}(t) = & E[e^{tY}] = E[e^{t(\frac{X-np}{\sqrt{npq}})}] \\ = & e^{-\frac{npt}{\sqrt{npq}}} E[e^{\frac{tX}{\sqrt{npq}}}] = e^{-\frac{npt}{\sqrt{npq}}} (q + pe^{\frac{t}{\sqrt{npq}}})^{n} \\ = & (qe^{-\frac{pt}{\sqrt{npq}}} + pe^{\frac{qt}{\sqrt{npq}}})^{n} \\ = & \left[q(1 - \frac{pt}{\sqrt{npq}} + \frac{pt^{2}}{2nq}) + p(1 + \frac{qt}{\sqrt{npq}} + \frac{qt^{2}}{2np} + o(\frac{t^{2}}{n})) \right]^{n} \\ = & \left[1 + \frac{t^{2}}{2n} + o(\frac{t^{2}}{n}) \right]^{n} \end{aligned}$$

$$\lim_{n \to +\infty} M_Y(t) = \lim_{n \to +\infty} \left[1 + \frac{t^2}{2n} + o(\frac{t^2}{n}) \right]^n$$

$$= \lim_{n \to +\infty} \left[1 + \frac{t^2}{2n} \right]^{\frac{2n}{\ell^2} \cdot \frac{t^2}{2}}$$

$$= e^{\frac{t^2}{2}}$$

当 $n \to \infty$ 时, $M_Y(t) \to e^{\frac{t^2}{2}}$, 即证二项分布近似正态分布.

思考题:利用密度函数证明当 $n \to \infty$ 时,二项分布近似正态分布.

证明:假定X服从二项分布B(n,p),则

$$P(X=m)=C_n^mp^mq^{n-m}$$

不妨假设m = np + d,则有

$$P(X = m) = P(X = np + d)$$

$$= C_n^{np+d} p^{np+d} q^{n-(np+d)}$$

$$= C_n^{np+d} p^{np+d} q^{nq-d}$$

$$= \frac{n!}{(np+d)!(nq-d)!} p^{np+d} q^{nq-d}$$

对上式利用Stirling公式 $n! = \sqrt{2\pi n} n^n e^{-n}$, 可得

$$P(X = np + d) = \frac{\sqrt{2\pi n} n^{n} e^{-n}}{\sqrt{2\pi (np + d)} (np + d)^{np+d} e^{-(np+d)}} \cdot \frac{1}{\sqrt{2\pi (nq - d)} (nq - d)^{nq-d} e^{-(nq-d)}} \cdot p^{np+d} q^{nq-d}$$

$$= \frac{1}{\sqrt{2\pi}} \cdot \left[\frac{n}{(np + d)(nq - d)} \right]^{\frac{1}{2}} \cdot \left(\frac{n}{np + d} \right)^{np+d}$$

$$\cdot \left(\frac{n}{nq - d} \right)^{nq-d} \cdot p^{np+d} q^{nq-d}$$

$$= \frac{1}{\sqrt{2\pi}} \cdot \left[\frac{(np + d)(nq - d)}{n} \right]^{-\frac{1}{2}} \cdot \left(\frac{np + d}{n} \right)^{-(np+d)}$$

$$\cdot \left(\frac{nq - d}{n} \right)^{-(nq-d)} \cdot p^{np+d} q^{nq-d}$$

$$\begin{split} &= \frac{1}{\sqrt{2\pi npq}} [(1 + \frac{d}{np})(1 - \frac{d}{nq})]^{-\frac{1}{2}} (p + \frac{d}{n})^{-(np+d)} \\ &\cdot (q - \frac{d}{n})^{-(nq-d)} p^{np+d} q^{nq-d} \\ &= \frac{1}{\sqrt{2\pi npq}} (1 + \frac{d}{np})^{-\frac{1}{2}} (1 - \frac{d}{nq})^{-\frac{1}{2}} (1 + \frac{d}{np})^{-(np+d)} (1 - \frac{d}{nq})^{-(nq-d)} \end{split}$$

$$In\left[(1+\frac{d}{np})^{-\frac{1}{2}}(1-\frac{d}{nq})^{-\frac{1}{2}}(1+\frac{d}{np})^{-(np+d)}(1-\frac{d}{nq})^{-(nq-d)}\right]$$

$$=-\frac{1}{2}In(1+\frac{d}{np})-\frac{1}{2}In(1-\frac{d}{nq})-(np+d)In(1+\frac{d}{np})$$

$$-(nq-d)In(1-\frac{d}{nq})$$

$$=-\frac{d}{2np}+\frac{d}{2nq}-(np+d)\left[\frac{d}{np}-\frac{1}{2}(\frac{d}{np})^2+\cdots\right]$$

$$-(nq-d)\left[-\frac{d}{nq}-\frac{1}{2}(\frac{d}{nq})^2-\cdots\right]$$

$$=-\frac{d^2}{2npq}+\frac{1}{6}\frac{d^3}{n^2p^2}-\frac{1}{6}\frac{d^3}{n^2q^2}-\frac{d}{2np}+\frac{d}{2nq}+\cdots$$

从而可得

$$P(X = m) = P(X = np + d) \approx \frac{1}{\sqrt{2\pi npq}} e^{-\frac{d^2}{2npq}}$$

思考题: Prove that when $n \to \infty$,binomial distribution is close to Poisson distribution.

证明: Assume that $X \sim B(n, p)$, $np = \lambda$

$$\binom{n}{k} p^{k} (1-p)^{n-k} = \frac{n(n-1)...(n-k+1)}{k!} (\frac{\lambda}{n})^{k} (1-\frac{\lambda}{n})^{n-k}$$
$$= \frac{\lambda}{k!} (1-\frac{1}{n})(1-\frac{2}{n})...(1-\frac{k-1}{n})(1-\frac{\lambda}{n})^{n-k}$$

for the given k

$$\lim_{n \to +\infty} (1 - \frac{\lambda}{n})^{n-k} = e^{-\lambda},$$

$$\lim_{n \to +\infty} (1 - \frac{1}{n})...(1 - \frac{k-1}{n}) = 1.$$

So,

$$\lim_{n\to+\infty}\binom{n}{k}p^k(1-p)^{n-k}=\frac{\lambda^k}{k!}e^{-\lambda}$$
.

The equation holds when $np \to +\lambda$,so when we calculate the binomial distribution b(n,p),when $n \to +\infty$ and p is small, $\lambda = np$.

$$\lim_{n\to+\infty} \binom{n}{k} p^k (1-p)^{n-k} = \frac{\lambda^k}{k!} e^{-\lambda}$$

is established.

采用另外一种方法证明: Using the Characteristic Function The Characteristic Function of the binomial distribution with P is

$$f_n(t) = (p_n e^{it} + q_n)^n = [1 + \frac{np_n(e^{it} - 1)}{n}].$$

when $n \to +\infty$, $np_n \to +\lambda$

$$\lim_{n\to +\infty} f_n(t) = \exp[\lambda(e^{it}-1)] = f(t).$$

Remarks on the normal limit

- To use the normal approximation for a fixed sample size m, we note that the approximation is best for p=1/2.
- Useful rule of thumb: use the normal approximation when

$$mp \geq 5$$
, and $m(1-p) \geq 5$.

(some people use 10 instead of 5)

Poisson limit

- Let $Y \sim B(m, p)$.
- Suppose that $m \to \infty$ and $p \to 0$ such that mp converges to (or is always equal to) some fixed constant λ .
- Then Y converges in distribution to a Poisson RV with parameter λ .



Inference for a binomial proportion

• Let $Y \sim B(m, p)$ and let the sample proportion be

$$\hat{p} = \frac{Y}{m}$$
.

• Then, \hat{p} is an unbiased estimator of p, with

$$var(\hat{p}) = \frac{p(1-p)}{m}.$$

- Asymptotically for large m, \hat{p} is normally distributed.
- The standard $100(1 \alpha)\%$ confidence interval(CI) for p is

$$\hat{\rho}\pm z_{1-\alpha/2}\sqrt{\frac{p(1-p)}{m}},$$

where $z_{1-\alpha/2}$ is the $(1-\alpha/2)$ th quantile of a N(0,1) RV.



Inference for p in practice

- In practice we do not know p in the variance for p̂.
- Some solutions:
 - 1. Use p=1/2. The estimated variance in this case is $(4m)^{-1/2}$. (Intervals are conservative).
 - 2. Estimate the variance by plugging in \hat{p} for p:

$$\widehat{var}(\hat{p}) = \sqrt{\frac{\hat{p}(1-\hat{p})}{m}}.$$

(Intervals are anti-conservative).

3. We "add two successes and add two failures". Let

$$\tilde{p} = \frac{Y+2}{m+2+2} = \frac{Y+2}{m+4}.$$

Then the adjusted CI is

Inference for p in practice(cont.)

4. In the Wilson CI we solve the equation

$$\frac{\hat{p}-p}{\sqrt{p(1-p)/m}}=\pm z_{\alpha/2}.$$

Letting $z = z_{\alpha/2}$, the resulting $100(1 - \alpha)\%$ CI for p is

$$\frac{Y+z^2/2}{m+z^2} \pm \sqrt{\frac{m\hat{p}(1-\hat{p})+z^2/4}{(m+z^2)^2}}.$$

(Intervals oscillate between conservative and anticonservative with increasing m).

5. You can ignore the normal limit, and use the exact binomial distribution of Y instead. (Intervals tends to be conservative).



思考题: 设 X_1, X_2, \dots, X_n 为取自二点分布 B(1, p) 的一个样本, 其中 $0 \le p \le 1$. 求p 的置信水平为 $1 - \alpha$ 的置信区间 $[\hat{p}_L, \hat{p}_U]$.

(1) 当样本量 n 充分大时,由中心极限定理得出 p 的置信水平近似为 $1-\alpha$ 的置信区间 $[\hat{p}_L, \hat{p}_U]$

$$\left[\bar{X} - U_{1-\frac{\alpha}{2}} \cdot \frac{\sqrt{\bar{X}(1-\bar{X})}}{\sqrt{n}}, \; \bar{X} + U_{1-\frac{\alpha}{2}} \cdot \frac{\sqrt{\bar{X}(1-\bar{X})}}{\sqrt{n}}\right]. \tag{1}$$

(2) 当样本精确分布已知且样本量 n 较小时,可基于充分统计量 $T(X) = \sum_{i=1}^{n} X_i$ 的分布函数 G(t,p) 构造 p 的置信区间.

p的精确置信区间构造过程:

(1) p的MLE为 \bar{X} ,且 p 的充分统计量为 $T(X) = \sum_{i=1}^{n} X_i$,其分布函数为

$$G(t,p) = P_p(T \le t) = \sum_{i=0}^{[t]} C_n^i \cdot p^i \cdot (1-p)^{n-i}.$$

其中 [t] 表示 $t(0 \le t \le n)$ 的整数部分.

(2) 判断分布函数 G(t,p) 的单调性:

$$\sum_{i=0}^k C_n^i \cdot p^i \cdot (1-p)^{n-i} = \frac{\Gamma(n+1)}{\Gamma(k+1) \cdot \Gamma(n-k)} \cdot \int_p^1 u^k \cdot (1-u)^{n-k-1} du,$$

令 k = [t],且 Be(x|m,n) 表示 Be(m,n) 的分布函数,则 G(t,p) = 1 - Be(p|k+1,n-k),可见,T(X) 的分布函数 G(t,p) 是 p 的连续、严格减函数。

(3) 如果 $G(t,\theta)$ 是 θ 的连续、严格减函数,那么 θ 的置信水平为 $1-\alpha$ 的置信区间为 $[\theta_L, \theta_U]$,且 θ_L 和 θ_U 分别是关于 θ 的方程 $G(T-0,\theta)=1-\alpha_1$ 和 $G(T,\theta)=\alpha_2$ 的解,其中 $\alpha_1+\alpha_2=\alpha$,且 $0\leq \alpha\leq 1$.

$$\left\{ \begin{array}{l} \frac{\Gamma(n+1)}{\Gamma(k)\cdot\Gamma(n-k+1)}\cdot\int_0^{\rho_L}u^{k-1}\cdot(1-u)^{n-k}\;du=\alpha_1,\\ \\ \frac{\Gamma(n+1)}{\Gamma(k+1)\cdot\Gamma(n-k)}\cdot\int_0^{\rho_U}u^k\cdot(1-u)^{n-k-1}\;du=1-\alpha_2, \end{array} \right.$$

$$\begin{cases}
Be(p_L|k, n-k+1) = \alpha_1, \\
Be(p_U|k+1, n-k) = 1 - \alpha_2,
\end{cases}$$
(2)

(4) 若随机变量 $B \sim Be(m,n)$, 则 $F = \frac{B}{1-B} \cdot \frac{n}{m} \sim F(2m,2n)$. 所以方程 (2) 可以等价变换为

$$\begin{cases}
F(\frac{p_L}{1-p_L} \cdot \frac{n-k+1}{k} \mid 2k, 2(n-k+1)) = \alpha_1, \\
F(\frac{p_U}{1-p_U} \cdot \frac{n-k}{k+1} \mid 2(k+1), 2(n-k)) = 1 - \alpha_2.
\end{cases}$$
(3)

$$\begin{cases}
\frac{\rho_{L}}{1-\rho_{L}} \cdot \frac{n-k+1}{k} = F_{\alpha_{1}}(2k, 2(n-k+1)), \\
\frac{\rho_{U}}{1-\rho_{U}} \cdot \frac{n-k}{k+1} = F_{1-\alpha_{2}}(2(k+1), 2(n-k)).
\end{cases} (4)$$

- (5) 解方程 (4) 得 p 的置信水平为 $1-\alpha$ 的置信区间 $[\hat{p}_L, \hat{p}_U]$
 - 当0<k<n时,</p>

$$\widehat{p}_{L} = \frac{k}{k + (n - k + 1) \cdot F_{1-\alpha_{1}}(2(n - k + 1), 2k)}$$

$$\widehat{p}_{U} = \frac{(k+1) \cdot F_{1-\alpha_{2}} (2(k+1), 2(n-k))}{(k+1) \cdot F_{1-\alpha_{2}} (2(k+1), 2(n-k)) + (n-k)}.$$

● 当 k = 0 时,

$$\widehat{p}_{L} = 0, \ \widehat{p}_{U} = \frac{F_{1-\alpha_{2}}(2, 2n)}{F_{1-\alpha_{2}}(2, 2n) + n}.$$

● 当 k = n 时,

$$\widehat{p}_U = 1, \ \widehat{p}_L = \frac{n}{n + F_{1-\alpha_1}(2, 2n)}.$$



Example:

从一批产品中随机抽查63件,发现有3件不合格品,求这批产品的不合格品率p的0.90置信区间.

Solution:

$$n = 63, k = 3, \hat{p} = 0.048, \alpha = 0.1.$$
 $F_{0.95}(122, 6) = 3.704, F_{0.95}(8, 120) = 2.016.$

$$\hat{p}_L = \frac{3}{3+61 \times F_{0.95}(122, 6)} = 0.013,$$

$$\hat{p}_U = \frac{4}{60+4 \times F_{0.95}(8, 120)} = 0.119.$$

故这批产品的不合格品率的 0.90 置信区间为 [0.013, 0.119].

Further problems with inference for p: An example

- Suppose a marketing analyst carries out a study to determine the proportion of people in a city who drink coffee regularly. In her random sample of 50 people, the analyst finds that 2 regularly drink coffee.
- What is a 95% CI for the population proportion of those who regularly drink coffee?

The logit transformation

- For certain combinations of m and y, regardless of the CI used, we obtain intervals that may not lie entirely in the region [0,1].
- To evade this problem, we define a 1-1 transformation of the probability p which maps the interval(0,1) to the whole real line. Consider the **logit** transformation. Define a new variable η by

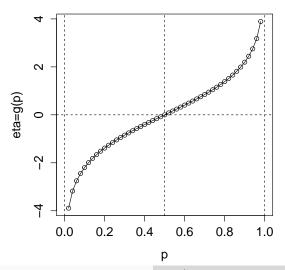
$$\eta = g(p) = \log(\frac{p}{1-p}).$$

- The key idea: build a CI for η and then transform back to obtain a CI for p.
- The inverse function of g(.) is

$$p=h(\eta)=\frac{e^{\eta}}{1+e^{\eta}}.$$



A plot of the transformation



Likelihood inference (revision)

The binomial log likelihood is

$$I(p) = log \binom{m}{y} + ylog(p) + (m-y)log(1-p).$$

The first derivative with respect to p is

$$\frac{dI}{dp} = \frac{y}{p} - \frac{m - y}{1 - p}.$$

• Solving the score equations, the MLE of p is $\hat{p} = \frac{y}{m}$

Likelihood inference (revision)

• The second derivative with respect to p is

$$\frac{d^2I}{dp^2} = -\frac{y}{p^2} - \frac{m-y}{(1-p)^2}.$$

and thus the Fisher information is

$$-E(\frac{d^2I}{dp^2}) = \frac{mp}{p^2} + \frac{m-mp}{(1-p)^2} = m(\frac{1}{p} + \frac{1}{1-p}).$$

Using the Fisher information

The inverse of the Fisher information is

$$I(p)^{-1} = [-E(\frac{d^2I}{dp^2})]^{-1} = \frac{p(1-p)}{m}.$$

• Using standard results for MLEs, an approximate $100(1 - \alpha)\%$ CI for p is as before,

$$\hat{p} \pm z_{1-\alpha/2} \sqrt{\frac{p(1-p)}{m}},$$

 If we replace the Fisher information by the observed information the CI becomes

$$\hat{p} \pm z_{1-\alpha/2} \sqrt{\frac{\hat{p}(1-\hat{p})}{m}}.$$



Using the delta method

• On the previous page we used the result that, as $m \to \infty$,

$$\sqrt{m}(\hat{p}-p) \stackrel{d}{\longrightarrow} N(0,I(p)^{-1}).$$

Using the Delta method it follows that

$$\sqrt{m}(g(\hat{p})-g(p)) \stackrel{d}{\longrightarrow} N(0,I(p)^{-1}(g^{'}(p))^{2}),$$

as $m o \infty$, where $g^{'}(.)$ is the derivative of the logit transform given by

$$g'(p) = (\frac{p}{1-p})^{-1} \frac{1}{(1-p)^2} = \frac{1}{p(1-p)}.$$

Calculating we find that

$$I(\eta)^{-1} \equiv I(p)^{-1}(g^{'}(p))^{2} = \frac{p(1-p)}{m}(\frac{1}{p(1-p)})^{2} = \frac{1}{mp(1-p)}.$$



CI for the logit

Now

$$g(\hat{p}) = log(\frac{\hat{p}}{1 - \hat{p}}) = log(\frac{y}{m - y}) = \hat{\eta};$$

$$g(p) = log(\frac{p}{1 - p}) = \eta.$$

$$I(\eta)^{-1} = \frac{(1 + e^{\eta})^2}{me^{\eta}}.$$

• An estimate of $I(\eta)^{-1}$ is

$$I(\hat{\eta})^{-1} = \frac{m}{y(m-y)} = \frac{1}{y} + \frac{1}{m-y}.$$

• Thus, an approximate 100(1 $-\alpha$)% CI for η is

$$\log(\frac{y}{m-y}) \pm z_{1-\alpha/2} \sqrt{\frac{1}{y} + \frac{1}{m-y}}.$$

that is, $[\eta_L, \eta_U]$, say.

• An approximate $100(1-\alpha)\%$ CI for p is then



The coffee example revisited

• A 95% CI for η is given by

$$log(\frac{y}{m-y}) \pm z_{0.975} \sqrt{\frac{1}{y} + \frac{1}{m-y}}$$

Hence, a 95% CI for p is

$$[h(\eta_L),h(\eta_U)]=[\frac{e^{\eta_L}}{1+e^{\eta_L}},\frac{e^{\eta_U}}{1+e^{\eta_U}}].$$

Generalized linear models for binary data

- Previously We considered statistical inference on a binomial proportion, p, for Y ~ B(m, p).
- In this section we will consider binary data:
 - -Each RV Y_i only has two possible values: 0 or 1.
 - -We assume that $Pr(Y_i = 0) = 1 p_i$ and $Pr(Y_i = 1) = p_i$.
 - -For each observation, *i*, we have a vector of **covariates** or **explanatory** variables

$$\mathbf{x}_{i}^{T} = (\mathbf{x}_{i,1}, \ldots, \mathbf{x}_{i,p})^{T}.$$

- -The RVs, $\{Y_1, \ldots, Y_n\}$ are independent.
- Our aim is to model the relationship between p_i and the explanatory variables x_i^T

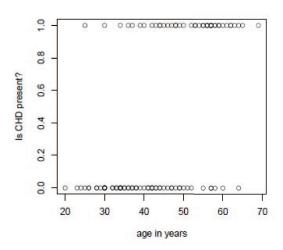


CHD example

- (Taken from Hosmer and Lemeshow (2000), "Applied Logistic Regression: Second Edition", Copyright John Wiley & Sons).
- In a study of Coronary Heart Disease the following data were collected on 100 individuals:
 - 1. Patient identification code.
 - 2. Age
 - 3. Whether they have Coronary Heart Disease (0=absent, 1=present).
- Investigators are interested in modeling the relationship between age and the presence or absence of Coronary Heart Disease.



Plotting the age versus CHD



Examining CHD proportion and age

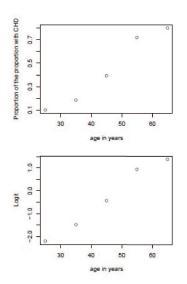
- Instead of age, consider age groups, (e.g.,20-29,30-39,...)
- Count the number of 0's and 1's in each age group.
- Now tabulate the proportion of 1's for each age group.

Age gr	oup	CHD absent	CHD present	CHD Proportion
20-2	29	9	1	0.100
30-3	9	22	5	0.185
40-4	.9	17	11	0.393
50-5	9	7	18	0.720
60-6	9	2	8	0.800

- Plot the midpoint of each age group versus the proportion.
- We can also plot the midpoint of each group versus the logit of the proportion.

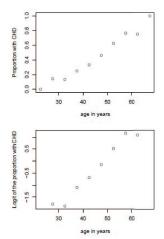


Examining CHD proportion and age (cont.)



Examining CHD proportion and age (cont.)

Now consider a different choice of age groups (every 5 years of age):



The simple linear logistic regression model

• Suppose that $Y_i (i = 1, ..., n)$ are n independent $B(1, p_i)$ RVs. We let

$$\eta_i = \log(\frac{p_i}{1 - p_i}) = \beta_0 + \beta_1 x_i,$$

Where $x_i (i = 1, ..., n)$ is some explanatory variable.

- For our example, $Y_i = CHD$ present/absent, and $x_i = age$.
- The above model equivalent to fitting $Y_i \sim B(1, p_i)$ where

$$p_i = \frac{e^{\beta_0 + \beta_1 x_i}}{1 + e^{\beta_0 + \beta_1 x_i}}$$

 This is a generalized linear model for binomial data with a logit link function.



To fit the simple linear logistic model:

```
Deviance Residuals:
   Min
            10 Median 30 Max
-1.9718 -0.8456 -0.4576 0.8253 2.2859
Coefficients:
          Estimate Std. Error z value Pr(>|z|)
(Intercept) -5.30945 1.13365 -4.683 2.82e-06 ***
          0.11092 0.02406 4.610 4.02e-06 ***
AGE
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 136.66 on 99 degrees of freedom
Residual deviance: 107.35 on 98 degrees of freedom
ATC: 111.35
```

Number of Fisher Scoring iterations: 4

We use the command anova(chd.model) to obtain the analysis of deviance table (here in R):

```
Analysis of Deviance Table

Model: binomial, link: logit

Response: CHD

Terms added sequentially (first to last)

Df Deviance Resid. Df Resid. Dev

NULL 99 136.66

AGE 1 29.31 98 107.35
```

• The observed value of $D(\hat{y}, \hat{\mu})$, is the "deviance residual".

The model test

- Asymptotically for large n, $D(y, \hat{\mu})$ has a χ_1^2 distribution.
- So the P-value for this model is

$$Pr(D(y, \hat{\mu}) > 29.31) < 0.001$$

To see this, note we obtain the P-value as follows.

$$1 - pchisq(29.31, 1)$$

$$6.167658e - 08$$

Conclusion:



Statistical inference

- Example: What is the estimated probability that a randomly chosen 50 year old subject has CHD present?
- Example: Produce a 95% CI for the estimated probability that a randomly chosen 50 year old subject has CHD present.

Using other link functions

- The logit is the most commonly used link function.
- Historically, it was not the first link function to be used in binomial regression. The probit link is defined to be

$$g_1(p)=\phi^{-1}(p),$$

where $\phi^{-1}()$ is the inverse cumulative distribution function for a standard normal random variable.

- Used traditionally for bioassay experiments (Finney, 1973).
 - Experiments in which a biological organism is used to test for chemical toxicity.

Beeetle example

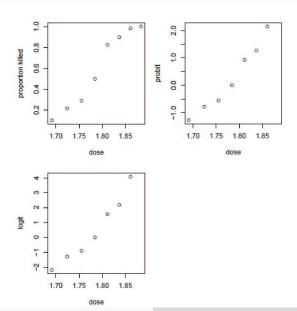
- (Taken from Table 7.2 of Dobson, 2000).
- The number of beetles that died after five hours exposure to a different number of concentrations of carbon disulfide(doses).
- Concentration units are log₁₀ CS₂ mg/l.

Dose	Total Number	Number Killed
1.6907	59	6
1.7242	60	13
1.7552	62	18
1.7842	56	28
1.8113	63	52
1.8369	59	53
1.8610	62	61
1.8839	60	60

£9 | 0

What link do we choose?

Examining proportion and dose



Common choices of link functions

 The four hours commonly used link functions(from most common to least common)are:

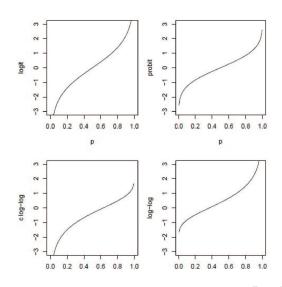
```
1.Logit: g_1(p) = \log(p/(1-p))
```

2.Probit:
$$g_2(p) = \Phi^{-1}(p)$$

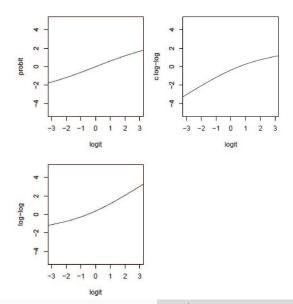
3.Complementary log-log:
$$g_3(p) = \log(-\log(1-p))$$

4.Log-log:
$$g_4(p) = -\log(-\log p)$$

Plots of the links



Comparing links(on the logit scale)



Comments on the links

- All the functions are increasing, continuous, and differentiable over 0
- The logit and probit are almost linearly related over the internal $p \in [0.1, 0.9]$.
- For small p, complementary log-log close to logit.
- For large p, log-log close to logit.
- The complementary log-log approaches infinity slower than any other link function.

- The data is represented in a compact form of counts for each combination of dose.
- We do not need to expand the dataset into 0s and 1s.
- Instead, we fit the model using the weights command:

Comments on the code

- Weights declares the number of individuals which make up each proportion and each factor combination.
- Binominal(link="probit") declares a binominal glm model with a probit link function.

(we can also uselogit or cloglog-log is not available in R)

```
Deviance Residuals:
```

```
Min 1Q Median 3Q Max
-1.5714 -0.4703 0.7501 1.0632 1.3449
```

Coefficients:

```
Estimate Std. Error z value Pr(>|z|)

(Intercept) -34.935 2.648 -13.19 <2e-16 *** dose 19.728 1.487 13.27 <2e-16 ***
```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 284.20 on 7 degrees of freedom Residual deviance: 10.12 on 6 degrees of freedom

AIC: 40.318

Number of Fisher Scoring iterations: 4

Comparing analysis of deviance tables

• Analysis of Deviance Table

```
Model: binomial, link: probit
```

Response: prob

Terms added sequentially (first to last)

```
Df Deviance Resid. Df Resid. Dev NULL 7 284.20 dose 1 274.08 6 10.12
```

Analysis of Deviance Table

```
Model: binomial, link: logit
```

Response: prob

Terms added sequentially (first to last)

```
Df Deviance Resid. Df Resid. Dev
NULL 7 284.202
dose 1 272.97 6 11.232
```

Which link do would you choose?



Interpreting glm models with different links

 Interpretation of the coefficients in each model follows by considering the inverse link function.

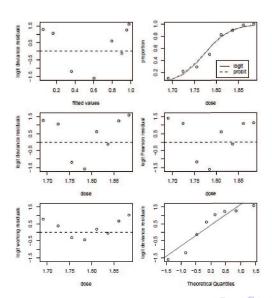
1.Logit:
$$h_1(\eta) = e^{\eta}/(1 - e^{\eta})$$

2.Probit:
$$h_2(\eta) = \Phi(\eta)$$

3. Complementary log-log:
$$h_3(\eta) = 1 - \exp(-e^{\eta})$$

4.Log-log:
$$h_4(\eta) = \exp(-e^{\eta})$$

Residuals analysis



Residuals analysis(cont.)

Conclusions?

思考题: X 和 Y 均为二分类变量, $X \in \{0,1\}, Y \in \{0,1\}$,则列联表分析的卡方检验和logistic模型对于系数 β_1 的检验是否一致?

 X/Y
 0
 1

 对于获取的样本 X_i , Y_i ($i=1,2,\cdots,n$), 整理可得
 0
 n_{00} n_{01}

 1
 n_{10} n_{11}

二维列联表的独立性检验

设有X, Y二个离散型随机变量,分别取r个值和c个值,对应地有二维列联表 $r \times c$,作n次观测,在(i,j)格的观测频数为 n_{ij} , i=1,2,...,r; j=1,2,...,c. 观测值落入(i,j)格的概率为 p_{ij} ,观测频数服从多项分布,其概率密度为

$$\frac{n!}{\prod_{i=1}^{r} \prod_{j=1}^{c} n_{ij}!} \cdot \prod_{i=1}^{r} \prod_{j=1}^{c} p_{ij}^{n_{ij}}$$

由于 $\sum_{i=1}^r\sum_{j=1}^c p_{ij}=1$,所以参数空间的独立参数个数为 $r\cdot c-1$, p_{ij} 的MLE 为 $\hat{p}_{ij}=n_{ij}/n$

考虑列联表的独立性检验,原假

设 $H_0: p_{ij} = p_{i.} \cdot p_{.j}, i = 1, 2, ..., r; j = 1, 2, ..., c$, 其中 $p_{i.}$ 和 $p_{.j}$ 分别是X 和Y 的边缘分布。原假设成立时, $p_{ij} = p_{i.} \cdot p_{.j}$,所以参数空间的独立参数个数为r + c - 2个。

这时, p_{ij} 的MLE为 $\hat{p}_{i.}\cdot\hat{p}_{.j}=(n_i/n)\cdot(n_j/n)$

该检验问题的似然比统计量为:

$$\Lambda(X) = \frac{\prod_{i=1}^{r} \prod_{j=1}^{c} (\frac{n_{ij}}{n})^{n_{ij}}}{\prod_{i=1}^{r} \prod_{j=1}^{c} (\frac{n_{i}}{n} \cdot \frac{n_{.j}}{n})^{n_{ij}}}$$

由于 $(r \cdot c - 1) - (r + c - 2) = (r - 1)(c - 1)$,故在原假设成立时, $2 \ln \Lambda$ 的极限分布为 $\chi^2((r - 1)(c - 1))$ 。在 $2 \ln \Lambda \ge \chi^2_{1-\alpha}((r - 1)(c - 1))$ 时,拒绝原假设。

logistic模型系数检验:

$$Y_i \sim B(1, p_i), \ \$$
密度函数为: $f(y_i, p_i) = p_i^{y_i} (1 - p_i)^{1 - y_i}.$

logistic 模型: $p_i = \frac{\exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)}$.

 $H_0: \beta_1 = 0 \ H_1: \beta_1 \neq 0$, 似然比检验统计量:

$$\Lambda = \frac{\prod_{i=1}^n \left(\frac{\exp(\hat{\beta}_0 + \hat{\beta}_1 x_i)}{1 + \exp(\hat{\beta}_0 + \hat{\beta}_1 x_i)}\right)^{y_i} \left(\frac{1}{1 + \exp(\hat{\beta}_0 + \hat{\beta}_1 x_i)}\right)^{1 - y_i}}{\prod_{i=1}^n \left(\frac{\exp(\tilde{\beta}_0)}{1 + \exp(\tilde{\beta}_0)}\right)^{y_i} \left(\frac{1}{1 + \exp(\tilde{\beta}_0)}\right)^{1 - y_i}}$$

其中 \hat{eta}_0 和 \hat{eta}_1 是在饱和模型下参数的极大似然估计值。 \tilde{eta}_0 是在原假设下参数的极大似然估计值, $2 \ln \Lambda \sim \chi^2(1)$ 。

证明两种方法的一致性
先计算
$$\hat{eta}_0$$
 和 \hat{eta}_1 :

$$L = \textstyle \prod_{i=1}^n \left(\frac{\exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)}\right)^{y_i} \left(\frac{1}{1 + \exp(\beta_0 + \beta_1 x_i)}\right)^{1 - y_i}$$

$$InL = \sum_{i=1}^{n} \left(y_i ln \frac{\exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)} + (1 - y_i) ln \frac{1}{1 + \exp(\beta_0 + \beta_1 x_i)} \right)$$

$$\frac{\partial \ln\!L}{\partial \beta_0} = \sum_{i=1}^n \left(y_i \frac{1}{1 + \exp(\beta_0 + \beta_1 x_i)} + (1 - y_i) \frac{-\exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)} \right)$$

令
$$\frac{\partial lnL}{\partial \beta_0} = 0$$
, 等价于

$$\sum_{i=1}^{n} \frac{y_i}{1 + \exp(\beta_0 + \beta_1 x_i)} = \sum_{i=1}^{n} \frac{(1 - y_i) \exp(\beta_0 + \beta_1 x_i)}{1 + \exp(\beta_0 + \beta_1 x_i)}$$

$$n_{01} \cdot \frac{1}{1 + \exp(\beta_0)} + n_{11} \cdot \frac{1}{1 + \exp(\beta_0 + \beta_1)} = n_{00} \cdot \frac{\exp(\beta_0)}{1 + \exp(\beta_0)} + n_{10} \cdot \frac{\exp(\beta_0 + \beta_1)}{1 + \exp(\beta_0 + \beta_1)}$$



$$\begin{split} &\frac{\partial lnL}{\partial \beta_{1}} = \sum_{i=1}^{n} \left(y_{i} \frac{x_{i}}{1 + \exp(\beta_{0} + \beta_{1} x_{i})} + (1 - y_{i}) x_{i} \frac{-\exp(\beta_{0} + \beta_{1} x_{i})}{1 + \exp(\beta_{0} + \beta_{1} x_{i})} \right) \\ & \diamondsuit \frac{\partial lnL}{\partial \beta_{1}} = 0, \quad \raisetate{0.5cm} \raisetate{0.5cm}$$

将 $\hat{\beta}_0$ 和 $\hat{\beta}_1$ 代入 Λ 分子部分的似然函数中:

$$L = \prod_{i=1}^{n} \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1}x_{i})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1}x_{i})} \right)^{y_{i}} \left(\frac{1}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1}x_{i})} \right)^{1-y_{i}}$$

$$= \left(\frac{1}{1 + \exp(\hat{\beta}_{0})} \right)^{n_{00}} \cdot \left(\frac{\exp(\hat{\beta}_{0})}{1 + \exp(\hat{\beta}_{0})} \right)^{n_{01}} \cdot \left(\frac{1}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}_{1})} \right)^{n_{10}} \cdot \left(\frac{\exp(\hat{\beta}_{0} + \hat{\beta}_{1})}{1 + \exp(\hat{\beta}_{0} + \hat{\beta}$$

计算
$$\tilde{\beta}_0$$
:

$$L = \prod_{i=1}^n \left(\frac{\exp(\beta_0)}{1 + \exp(\beta_0)} \right)^{y_i} \left(\frac{1}{1 + \exp(\beta_0)} \right)^{1 - y_i}$$

$$\frac{\partial \ln L}{\partial \beta_0} = \sum_{i=1}^n \left(y_i \frac{1}{1 + \exp(\beta_0)} + (1 - y_i) \frac{-\exp(\beta_0)}{1 + \exp(\beta_0)} \right)$$

令
$$\frac{\partial \ln L}{\partial \beta_0} = 0$$
, 等价 于 $\sum_{i=1}^n y_i = \sum_{i=1}^n (1 - y_i) \exp(\beta_0)$

将 $\tilde{\beta}_0$ 代入 Λ 分母部分的似然函数:

$$L = \textstyle \prod_{i=1}^n (\frac{n_{.1}}{n})^{y_i} (\frac{n_{.0}}{n})^{1-y_i} = (\frac{n.1}{n})^{n_{11}+n_{01}} \cdot (\frac{n.0}{n})^{n_{00}+n_{10}}$$

$$\Lambda = \frac{\left(\frac{n_{00}}{n_{0}}\right)^{n_{00}} \cdot \left(\frac{n_{01}}{n_{0}}\right)^{n_{01}} \cdot \left(\frac{n_{10}}{n_{1}}\right)^{n_{10}} \cdot \left(\frac{n_{11}}{n_{1}}\right)^{n_{11}}}{\left(\frac{n_{11}}{n}\right)^{n_{11}+n_{01}} \cdot \left(\frac{n_{0}}{n}\right)^{n_{00}+n_{10}}} \\
= \left(\frac{n_{00} \cdot n}{n_{0.} \cdot n_{.0}}\right)^{n_{00}} \cdot \left(\frac{n_{01} \cdot n}{n_{0.} \cdot n_{.1}}\right)^{n_{01}} \cdot \left(\frac{n_{10} \cdot n}{n_{1.} \cdot n_{.0}}\right)^{n_{10}} \cdot \left(\frac{n_{11} \cdot n}{n_{1.} \cdot n_{.1}}\right)^{n_{11}} \\
= \frac{\left(\frac{n_{00}}{n_{0.}}\right)^{n_{00}} \cdot \left(\frac{n_{01}}{n}\right)^{n_{01}} \cdot \left(\frac{n_{10}}{n}\right)^{n_{10}} \cdot \left(\frac{n_{11}}{n}\right)^{n_{11}}}{\left(\frac{n_{0.}}{n_{0.}} \cdot \frac{n_{.0}}{n}\right)^{n_{00}} \cdot \left(\frac{n_{0.}}{n_{0.}} \cdot \frac{n_{.1}}{n}\right)^{n_{01}} \cdot \left(\frac{n_{1.}}{n} \cdot \frac{n_{.0}}{n}\right)^{n_{10}} \cdot \left(\frac{n_{1.}}{n} \cdot \frac{n_{.1}}{n}\right)^{n_{11}}} \\
= \frac{\prod_{i \in \{0,1\}} \prod_{j \in \{0,1\}} \left(\frac{n_{ij}}{n} \cdot \frac{n_{ij}}{n}\right)^{n_{ij}}}{\prod_{i \in \{0,1\}} \prod_{j \in \{0,1\}} \left(\frac{n_{ij}}{n} \cdot \frac{n_{.j}}{n}\right)^{n_{ij}}}}$$

证毕。

问题

- 当X为四分类变量, X ∈ {1,2,3,4}.
- 例如, X表示螃蟹的颜色, X = 1表示light medium, X = 2表示medium, X = 3表示dark medium, X = 4表示dark.
- 如何检验X作为一个整体, 对Y是否有影响.

检验1:列联表

似然比统计量为:

$$\lambda(X) = \frac{\prod_{i=1}^{n} p(X_i; \hat{\theta})}{\prod_{i=1}^{n} p(X_i; \theta_0)}$$

$$2 ln \lambda(X) \sim \chi^2(3)$$

检验2: logistic模型似然比检验

引入3个虚拟变量构建logistic模型

$$logit(p_i) = \beta_0 + X_1\beta_1 + X_2\beta_2 + X_3\beta_3$$

原假设 H_0 : $\beta_1 = \beta_2 = \beta_3 = 0$

似然比统计量:

$$\frac{\prod_{i=1}^n f(y_i; \hat{p}_i)}{\prod_{i=1}^n f(y_i; p_i)}$$

分子: 饱和模型似然函数; 分母: Ho成立下模型的似然函数

$$2ln\lambda(X) \sim \chi^2(3)$$

More complicated binomial GLM models

- So far we have considered binomial GLMs in which we wish to model the dependence between a binomial proportion and a continuous explanatory variable.
- We shall now consider models which
 - incorporate factors.
 - incorporate multiple terms.
 - allow for interactions.

Horse shoe crabs example

(Taken from Agresti, 1996).

- Data from a study of nesting horse shoe crabs.
- Response variable:number of satellites (number of males residing nearby).
- Explanatory variables of interest:
 - color of the crab (1:light medium,2:medium,3:dark medium,4:dark).
 - spine condition (1:both good,2:one worn or broken,3:both worn or broken).
 - width of the crab (in cm).
 - weight of the crab (in kg).

A binomial response

- Instead of counting the number of satellites, consider a binomial response:
 - has satellite (0:no males reside nearby, 1:at least one male resides nearby).
- Scientific question of interest:
 Can we build a model to relate the probability of having a satellite nearby to the explanatory variables of interest?

Incorporating factors

- We start by relating the probability of having a satellite nearby to whether color is dark or not. Recode color as a variable 'is.dark': 0 not dark (light medium or medium), 1 dark (dark medium or dark).
- Summarizing the counts we have:

color	no satellites nearby	satellites nearby	total
not dark	29	78	107
dark	33	33	66

Expressed as proportions:

color	no satellites nearby	satellites nearby	
not dark	0.271	0.729	
dark	0.500	0.500	

Initial thoughts:



Estimating these proportions using a binomial GLM

Here is the code to fit the binomial glm model:

```
Deviance Residuals:
```

```
Min 1Q Median 3Q Max
-1.6159 -1.1774 0.7951 0.7951 1.1774
```

Coefficients:

```
Estimate Std. Error z value Pr(>|z|) (Intercept) 0.9894 0.2175 4.549 5.39e-06 *** is.darkTRUE -0.9894 0.3285 -3.012 0.0026 **
```

```
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: 216.53 on 171 degrees of freedom ATC: 220.53
```

Number of Fisher Scoring iterations: 4

Odds and odds ratio

- Commonly used concepts when working with probabilities
- Odds of an event
 - The probability that the event happens=p
 - Odds in favor of the event is given as $odds = p/(1 p) = \exp[logit(p)]$
 - Examples: 病情复发的odds;
- The odds ratio is the ratio of the odds of an event occurring in one group to the odds of it occurring in another group
 - Example: Use odds ratio to compare the effects of the treatments



Odds and odds ratio

- Let p denote the probability of relapse occurrence
- Odds of relapse occurrence is p/(1-p)
 - Smaller is better, indicating a relapse is less likely to occur
- The odds ratio of occurrence in drug A group against placebo group is used to compare the treatments
 - An odd ratio less than 1 indicates that relapse is less likely to occur in drug A group, i.e., drug A is better than placebo

Odds and odds ratio-Interpret the parameter of a continuous predictor

Assume a glm contains a single continuous predictor x. The mean structure is

$$logit(p) = \beta_0 + \beta_1 x$$

- When x = a, the odds is $exp(logit(p)) = exp(\beta_0 + \beta_1 a)$
- Increasing x = a by one unit to x = a + 1, the odds changes to $\exp(\beta_0 + \beta_1 a + \beta_1)$
- The odds ratio is

$$\frac{adds_{a+1}}{odds_a} = \frac{\exp(\beta_0 + \beta_1 a + \beta_1)}{\exp(\beta_0 + \beta_1 a)} = \exp(\beta_1)$$

• Then $[\exp(\beta_1) - 1] \times 100\%$ is the percent change in the odds of event occurrence when x is increased by one unit



Odds and odds ratio-Interpret the parameter of a categorical predictor

 Assume a glm contains a single categorical predictor x with two levels (A and B). The mean structure is

$$logit(p_i) = \begin{cases} eta_0, \\ eta_0 + eta_B, & \mathsf{x=B} \end{cases}$$

- When x = A the odds is $\exp(\beta_0)$
- When x = B the odds is $\exp(\beta_0 + \beta_B)$
- The odds ratio (B against A) is

$$\frac{odds_B}{odds_A} = \exp(\beta_B)$$

• Then $[\exp(\beta_B-1]\times 100\%$ tells the percent change in the odds in group B compared to group A

Understanding the model

• Our model is that $\{Y_{i,j}: i=1,2; j=i,\cdots,m_j\}$ are a set of independent RVs with $Y_{i,j} \sim B(1,p_i)$ and where p_i satisfies

$$\log\left(\frac{p_i}{1-p_i}\right) = \mu + \alpha_i$$

- In our example we have: $m_1 = 107, m_2 = 66$.
- To fit the model we assume that $\alpha_1 = 0$. This is the default in R.

An equivalent model

• Equivalently $\{Y_i : i = 1, 2\}$ are two independent RVs with $Y_i \sim B(m_i, p_i)$ and where p_i satisfies

$$\log\left(\frac{p_i}{1-p_i}\right) = \mu + \alpha_i$$

We can fit this model using the weights option in glm (see last lecture).

Interpreting the factor level effects

 Estimated logit for the two factors based glm model is not dark:
 dark:

-

 Estimated proportions are not dark:
 dark:

 Confidence intervals for these logits and proportions follow in a natural way.

Differences of logits

Example: What is the estimated difference between the 'not dark' and 'dark' logits? What is a 95% CI for this difference?
 (This difference is the log odds ratio).

Here is the analysis of deviance table:

```
Analysis of Deviance Table

Model: binomial, link: logit

Response: has.satellite

Terms added sequentially (first to last)

Df Deviance Resid. Df Resid. Dev

NULL 172 225.76
is.dark 1 9.2275 171 216.53
```

- Conclusions:
- Exercise: what is this deviance test equivalent to?

More factor levels

 Now consider a model relating the probability of having a satellite and the color (which has 4 factor levels).

The table of counts are:

color	no satellites nearby	satellites nearby	total
light medium	3	9	12
medium	26	69	9
dark medium	18	26	44
dark	15	7	22

Initial summary:



The code and model summary

The code is:

```
crabs=read.table("file:///F:/Xu_WL/crabs.dat.txt",
    header=T)
has.satellite=c(crabs$satellite>=1)
crabs.color=glm(has.satellite~factor(color),
    data=crabs,family="binomial")
summary(crabs.color)
anoya(crabs.color)
```

The code and model summary

The result is:

```
Deviance Residuals:
   Min 10 Median 30 Max
-1.6651 -1.3370 0.7997 0.7997 1.5134
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) 1.0986 0.6667 1.648 0.0994 .
factor(color)2 -0.1226 0.7053 -0.174 0.8620
factor(color)3 -0.7309 0.7338 -0.996 0.3192
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: 212.06 on 169 degrees of freedom
ATC: 220.06
Number of Fisher Scoring iterations: 4
```

Analysis of deviance:

Analysis of Deviance Table

```
Model: binomial, link: logit
```

Response: has.satellite

Terms added sequentially (first to last)

Df Deviance Resid. Df Resid. Dev NULL 172 225.76 factor(color) 3 13.698 169 212.06

Overall conclusions:

A warning and a discussion of asymptotics

- The limiting distribution of λ_{n-p}^2 for the deviance and Pearson statistics are based on the following assumptions:
 - 1. The sample of n observations are distributed independently with a $B(m_i, p_i)$ distribution.
 - 2. The overall sample size n remains fixed, but $m_i \to \infty$, such that $m_i p_i (1 p_i) \to \infty$ for each i.
- if *n* is large and $m_i p_i (1 p_i)$ remains bounded the theory breaks down:
 - 1. The λ^2 distribution no longer applies.
 - 2. $D(y, \hat{\mu})$ is not independent of \hat{p} .

Light at the end of the tunnel

- Suppose we fit two glm models. In the first model assume that $\hat{\mu}_0$ is the estimate of μ . In the second model we add one additional covariate to the first model. Let $\hat{\mu}_1$ be the estimate of μ in this case.
- Then $D(y; \hat{\mu}_0) D(y; \hat{\mu}_1)$ has an asymptotic λ_1^2 distribution under either of following settings.
 - 1. $n \to \infty$.
 - 2. The overall sample size n remains fixed, but $m_i \to \infty$, such that $m_i p_i (1 p_i) \to \infty$ for each i.

Multiple terms in the binomial GLM

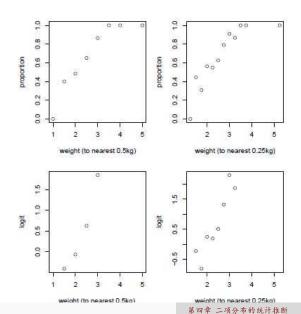
- We will continue our analysis of the horseshoe crabs dataset.
- We are now interested in model building, allowing for the possibility of incorporating more terms in the model.
 (We have already shown that the color is associated with the probability of having a satellite nearby).
- We start by producing pairwise summaries of the response variable (whether there is a satellite nearby) by each of the explanatory variables in the dataset (first the factor variables, and then the continuous variables).
- We could extend these comparisons to combinations of three or more variables.

Summary of having a satellite by weight

	counts		proportions	
color	no satellites nearby	satellites nearby	no satellites nearby	satellites nearby
light medium	3	9	0.250	0.750
medium	26	69	0.274	0.726
dark medium	18	26	0.409	0.591
dark	15	7	0.682	0.318

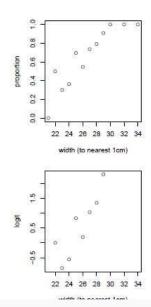
spine condition	counts		proportions	
	no satellites nearby	satellites nearby	no satellites nearby	satellites nearby
both good	11	26	0.297	0.703
one broken	8	7	0.533	0.467
both broken	43	78	0.355	0.645

2.2.Summary of having a satellite by weight





Summary of having a satellite by width



Our model building strategy

- We will consider a step-wise modeling strategy, starting with the simplest model and making our model more complicated.
- We are looking for a parsimonious model, which models/predicts the probability of having a satellite nearby.
- This is not the only way to choose a model!
- We use tests based on the analysis of deviance table to help us select a model.(We may also use residual plots, when appropriate).
- For brevity, I will demonstrate only a subset of the models we could consider.

Modeling the probability of having a satellite nearby in terms of the width

```
crabs=read.table("file:///F:/Xu WL/crabs.dat.txt".header=T)
has.satellite=c(crabs$satellite>=1)
crabs.width<-glm(has.satellite~width,
data=crabs,family="binomial")
summary(crabs.width)
anova(crabs.width)
Deviance Residuals:
   Min
             10 Median
                         30
                                      Max
-2.0281 -1.0458 0.5480 0.9066 1.6942
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
(Intercept) -12.3508 2.6287 -4.698 2.62e-06 ***
width
            0 4972
                       0 1017 4 887 1 02e-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: 194.45 on 171 degrees of freedom
AIC: 198.45
Number of Fisher Scoring iterations: 4
```

The analysis of deviance

```
Analysis of Deviance Table

Model: binomial, link: logit

Response: has.satellite

Terms added sequentially (first to last)

Df Deviance Resid. Df Resid. Dev

NULL 172 225.76
width 1 31.306 171 194.45
```

- We test:
- The P-value is:
- Conclusion:

Adding color to the model

Number of Fisher Scoring iterations: 4

```
crabs=read.table("file:///F:/Xu WL/crabs.dat.txt",header=T)
has.satellite=c(crabs$satellite>=1)
crabs.widcol<-qlm(has.satellite~width+factor(color),data=crabs.family="binomial")
summary(crabs.widcol)
anova(crabs.widcol)
Deviance Residuals:
            10 Median
   Min
                              30
                                     Max
-2.1124 -0.9848 0.5243 0.8513 2.1413
Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept) -11.38519 2.87346 -3.962 7.43e-05 ***
width
              0.46796 0.10554 4.434 9.26e-06 ***
factor(color)2 0.07242 0.73989 0.098 0.922
factor(color)3 -0.22380 0.77708 -0.288 0.773
factor(color)4 -1.32992 0.85252 -1.560 0.119
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: 187.46 on 168 degrees of freedom
ATC: 197 46
```

The analysis of deviance table

Analysis of Deviance Table

Model: binomial, link: logit

Response: has.satellite

Terms added sequentially (first to last)

Df Deviance Resid. Df Resid. Dev NULL 172 225.76 width 1 31.3059 171 194.45 factor(color) 3 6.9956 168 187.46

- We test:
- The P-value is:
- Conclusion:

Adding weight to the width model

```
Call:
qlm(formula = has.satellite ~ width + weight, family = "binomial",
   data = crabs)
Deviance Residuals:
   Min
       10 Median
                        30
                                    Max
-2.1127 -1.0344 0.5304 0.9006 1.7207
Coefficients:
          Estimate Std. Error z value Pr(>|z|)
(Intercept) -9.3547 3.5280 -2.652 0.00801 **
width 0.3068 0.1819 1.686 0.09177 .
weight 0.8338 0.6716 1.241 0.21445
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: 192.89 on 170 degrees of freedom
ATC: 198.89
Number of Fisher Scoring iterations: 4
```

The analysis of deviance table

```
Analysis of Deviance Table

Model: binomial, link: logit
```

Response: has.satellite

Terms added sequentially (first to last)

```
        Df
        Deviance Resid. Df
        Resid. Dev

        NULL
        172
        225.76

        width
        1 31.3059
        171
        194.45

        weight
        1 1.5608
        170
        192.89
```

- We test:
- The P-value is:
- Conclusion:

Further steps in the modeling strategy

- Continuing in a similar way we find that:
 Spine condition is **not significant** when added to the model which predicts the probability of having a satellite nearby using the width variable;
 The variable 'is dark' **not significant** when added to the width model.
- Instead of our current definition of 'is dark', consider the variable 'is very dark' which is:
 - 0 if the color is light medium, medium, or dark medium;
 - 1 if the color is dark.

```
crabs=read.table("file:///F:/Xu_WL/crabs.dat.txt",header=T)
has.satellite=c(crabs$satellite>=1)
is.dark=crabs$color<=3
crabs.widdark-glm(has.satellite~width+is.dark,data=crabs,family="binomial")
summary(crabs.widdark)
anova(crabs.widdark)</pre>
```

Adding "is.very.dark" to the width model

```
Call:
qlm(formula = has.satellite ~ width + is.dark, family = "binomial",
   data = crabs)
Deviance Residuals:
   Min
            10 Median
                        30
                                     Max
-2.0821 -0.9932 0.5274 0.8606 2.1553
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
(Intercept) -12.9795 2.7272 -4.759 1.94e-06 ***
width 0.4782 0.1041 4.592 4.39e-06 ***
is.darkTRUE 1.3005 0.5259 2.473 0.0134 *
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: 187.96 on 170 degrees of freedom
ATC: 193.96
Number of Fisher Scoring iterations: 4
```

The analysis of deviance table

Model: binomial, link: logit

Analysis of Deviance Table

Response: has.satellite

Terms added sequentially (first to last)

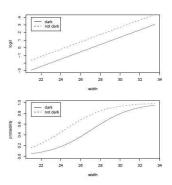
 NULL
 172
 225.76

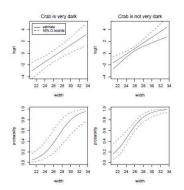
 width
 1
 31.3059
 171
 194.45

 is.dark
 1
 6.4948
 170
 187.96

- We test:
- The P-value is:
- Conclusion:

Demonstrating the model predictions





Incorporating interaction terms

- On the link scale (e.g., the logit), the interpretation of interactions is the same as for linear models.
 - -The interpretation is more complicated on the original scale.
- For our example, suppose we want to fit a different slope parameter for the 'width' variable according to whether the crabs are 'very dark' or 'not very dark'.
- Arrange the data into two groups, letting i = 1 if a crab is dark and i = 2 if a crab is not dark. Let the index j denote the crabs in group i.
- Let y_{ij} be 0 if no satellites nearby and 1 otherwise, with Y_{ij} denoting the associated RVs.
- Let w_{ij} be the widths of the *j*th crab in the *i*th color group.



Comparing the model with and without the interaction

• Without an interaction term, the model is $Y_{ij} \sim B(1; p_{ij})$ with

$$\eta_{ij} = g(p_{ij}) = \alpha + \beta w_{ij} + \gamma_i$$

To fit this model we assume that $\gamma_1 = 0$.

• Adding an interaction term, the model becomes $Y_{ij} \sim B(1; p_{ij})$ with

$$\eta_{ij} = g(p_{ij}) = \alpha + \beta w_{ij} + (\beta \gamma)_i w_{ij}$$

We assume that $\gamma_1 = 0$ and $(\beta \gamma)_1 = 0$.



The R model specifications

Without an interaction term, the model is:

has.satellite \sim width + weight.

Adding an interaction term, the model becomes:

has.satellite \sim width * is.very.dark

Another way to write this is:

 $has.satellite \sim width + is.very.dark + width: is.very.dark$

The resulting model summary

```
Call:
qlm(formula = has.satellite ~ width + is.dark + width * is.dark,
   family = "binomial", data = crabs)
Deviance Residuals:
   Min 10 Median 30 Max
-2.1366 -0.9344 0.4996 0.8554 1.7753
Coefficients:
               Estimate Std. Error z value Pr(>|z|)
(Intercept) -5.8538 6.6939 -0.874 0.382
width
                0.2004 0.2617 0.766 0.444
is.darkTRUE -6.9578 7.3182 -0.951 0.342
width:is.darkTRUE 0.3217 0.2857 1.126 0.260
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 225.76 on 172 degrees of freedom
Residual deviance: 186.79 on 169 degrees of freedom
ATC: 194 79
Number of Fisher Scoring iterations: 4
```

Analysis of Deviance Table

Analysis of Deviance Table

Model: binomial, link: logit

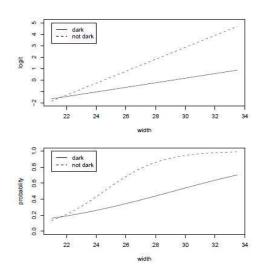
Response: has.satellite

Terms added sequentially (first to last)

	Df	Deviance	Resid. Df	Resid. Dev
NULL			172	225.76
width	1	31.3059	171	194.45
is.dark	1	6.4948	170	187.96
width:is.dark	1	1.1715	169	186.79

- We test:
- The P-value is:
- Conclusion:

Demonstrating the effect of the interactions



思考题: Logistic回归中,参数假设检验 $H_0: \beta_1=0, H_1: \beta_1\neq 0$,似然比检验与两因素t检验的关系。

- t统计量:根据Y的值将X分成两组,当y=1时, $x_i \sim N(\mu_1, \sigma^2), i=1,\ldots,n_1$, \bar{X}_1 服从 $N(\mu_1,\frac{\sigma^2}{n_1})$;当y=0时, $x_j \sim N(\mu_2,\sigma^2), j=1,\ldots,n_2$, \bar{X}_2 服从 $N(\mu_2,\frac{\sigma^2}{n_2})$ 。
- 等价于检验H₀: μ₁ = μ₂, H_a: μ₁ ≠ μ₂。
- 当 H_0 成立时, $t = rac{ar{x}_1 ar{x}_2}{\sqrt{(rac{1}{n_1} + rac{1}{n_2})\hat{\sigma^2}}} \sim t(n-2)$ 。

- 似然比检验: 当 H_0 成立时, 即 $\beta_1 = 0$
- $p = \frac{e^{\beta_0}}{1+e^{\beta_0}}$, 对数似然函数为:

$$l_0 = \sum_{i=1}^{n} [y_i \log \frac{e^{\beta_0}}{1 + e^{\beta_0}} + (1 - y_i) \log \frac{1}{1 + e^{\beta_0}}]$$

$$= n_1 \log \frac{e^{\beta_0}}{1 + e^{\beta_0}} + n_2 \log \frac{1}{1 + e^{\beta_0}}$$

$$= n_1 \beta_0 - n \log(1 + e^{\beta_0})$$

● 关于β₀求导数:

$$\frac{dI_0}{d\beta_0} = n_1 - n \frac{e^{\beta_0}}{1 + e^{\beta_0}} = 0$$

可以得到
$$\frac{\hat{\beta_0}}{1+e^{\hat{\beta_0}}} = \frac{n_1}{n}$$
, $I(\hat{\beta_0}) = n_1 \log \frac{n_1}{n} + n_2 \log \frac{n_2}{n}$



● 似然比检验: 当 H_0 不成立时, 即 $\beta_1 \neq 0$

•
$$p_i = \frac{e^{\beta_0 + \beta_1 x_i}}{1 + e^{\beta_0 + \beta_1 x_i}}$$
, 对数似然函数为:

$$I_{1} = \sum_{i=1}^{n} [y_{i} \log \frac{e^{\beta_{0} + \beta_{1} x_{i}}}{1 + e^{\beta_{0} + \beta_{1} x_{i}}} + (1 - y_{i}) \log \frac{1}{1 + e^{\beta_{0} + \beta_{1} x_{i}}}]$$

$$= \sum_{i=1}^{n_{1}} \log \frac{e^{\beta_{0} + \beta_{1} x_{i}}}{1 + e^{\beta_{0} + \beta_{1} x_{i}}} + \sum_{j=1}^{n_{2}} \log \frac{1}{1 + e^{\beta_{0} + \beta_{1} x_{j}}}$$

$$= \beta_{0} n_{1} + \beta_{1} \sum_{i=1}^{n_{1}} x_{i} - \sum_{i=1}^{n} \log(1 + e^{\beta_{0} + \beta_{1} x_{i}})$$

● 关于β₀求偏导:

$$\frac{\partial I_1}{\partial \beta_0} = n_1 - \sum_{i=1}^n \frac{e^{\beta_0 + \beta_1 x_i}}{1 + e^{\beta_0 + \beta_1 x_i}} = 0$$

$$\frac{\partial l_1}{\partial \beta_1} = \sum_{i=1}^{n_1} x_i - \sum_{i=1}^n \frac{x_i e^{\beta_0 + \beta_1 x_i}}{1 + e^{\beta_0 + \beta_1 x_i}} = 0$$

可以得到
$$\sum_{i=1}^n \hat{p}_i = n_1$$
, $\sum_{i=1}^{n_1} x_i = \sum_{i=1}^n \hat{p}_i x_i$ 《日本日本》 是 今年 第四章 二项分布的统计推断

可以得到
$$\sum_{i=1}^{n} \hat{p}_i = n_1$$
, $\sum_{i=1}^{n_1} x_i = \sum_{i=1}^{n} \hat{p}_i x_i$

● 由此可以得到deviance为:

$$D = 2[I_1 - I_0] = 2\left[\sum_{i=1}^{n_1} \log \hat{\rho}_i + \sum_{j=1}^{n_2} \log(1 - \hat{\rho}_j) - (n_1 \log \frac{n_1}{n} + n_2 \log \frac{n_2}{n})\right]$$

当
$$H_0$$
成立时, $D \sim \chi^2(1)$

思考题2,3

- 思考题:采用向前法、向后法和逐步回归法对逻辑回归进行变量选择
- 思考题:加入随机生成的四个随机变量,两个离散变量X₅、X₆,两个连续变量X₇、X₈,采用向前法、向后法和逐步回归法对逻辑回归进行变量选择

多元线性回归模型:
$$Y = X\beta + \varepsilon$$

原假设: $H_0: \beta = 0$

$$W = \sqrt{n}(\hat{\beta} - 0)^{T}(Cov(\sqrt{n}\hat{\beta}))^{-1}\sqrt{n}(\hat{\beta} - 0)$$
$$F = \frac{\sum_{i=1}^{k}(\hat{y}_{i} - \bar{y})^{2}/k}{\sum_{i=1}^{k}(y_{i} - \hat{y}_{i})^{2}/(n - k - 1)}$$

判断F统计量与W的关系

当 $Cov(\sqrt{n\beta})$ 已知时,W新近服从 $\chi^2(k)$ 当 $Cov(\sqrt{n\beta})$ 未知时,构造F统计量:

$$F = \sqrt{n(\hat{\beta} - 0)^T (\hat{Cov}(\sqrt{n\hat{\beta}}))^{-1}} \sqrt{n(\hat{\beta} - 0)^{\frac{n-k-1}{k}}}$$

$$F = \sqrt{n(\hat{\beta} - 0)^{T}(\hat{Cov}(\sqrt{n\hat{\beta}}))^{-1}} \sqrt{n(\hat{\beta} - 0)} \frac{n - k - 1}{k}$$

$$= \frac{\hat{\beta}^{T}(X^{T}X)\hat{\beta}/k}{\hat{\sigma}^{T}\hat{\sigma}/(n - k - 1)}$$

$$= \frac{Y^{T}X(X^{T}X)^{-1}(X^{T}X)(X^{T}X)^{-1}X^{T}Y/k}{\hat{\sigma}^{T}\hat{\sigma}/(n - k - 1)}$$

$$= \frac{Y^{T}X(X^{T}X)^{-1}X^{T}Y/k}{\hat{\sigma}^{T}\hat{\sigma}/(n - k - 1)}$$

$$= \frac{Y^{T}HY/k}{\hat{\sigma}^{T}\hat{\sigma}/(n - k - 1)}$$

$$= \frac{\sum_{i=1}^{k} \hat{y}_{i}^{2}/k}{\sum_{i=1}^{k} (y_{i} - \hat{y}_{i})^{2}/(n - k - 1)}$$

此时原假设为: $H_0: \beta = 0$

构造F统计量:

$$F = \frac{(\hat{\gamma} - \gamma_0)^T M^T [M(U^T U)^{-1} M^T]^{-1} M(\hat{\gamma} - \gamma_0)/k}{\hat{\sigma}^T \hat{\sigma}/(n-k-1)}$$

其中

$$[M(U^TU)^{-1}M^T]^{-1} = X^T[I_n - \frac{1}{n}J]X$$
$$\hat{\beta} = M(\hat{\gamma} - \gamma_0) = (X^T[I_n - \frac{1}{n}J]X)^{-1}X^T[I_n - \frac{1}{n}J]Y$$

此时F统计量为:

$$F = \frac{Y^T[I_n - \frac{1}{n}J]X(X^T[I_n - \frac{1}{n}J]X)^{-1}X^T[I_n - \frac{1}{n}J]Y/k}{\hat{\sigma}^T\hat{\sigma}/(n-k-1)}.$$

 X_1 、 X_2 是0-1变量,Y是连续随机变量,对Y与 X_1 、 X_2 建立线性模型,考虑 X_1 与 X_2 之间存在交互作用,求 β_{12} 的表达式,并解释 β_{12} 等于0所表示的意义。

X_1/X_2	0	1
0	<i>n</i> ₀₀	<i>n</i> ₀₁
1	<i>n</i> ₁₀	<i>n</i> ₁₁

线性模型无交互作用情况

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \epsilon$$

 $\epsilon \sim N(0,1)$, 似然函数:

$$L(\beta; y) \propto \sum_{i=1}^{n} -\frac{(Y_i - EY_i)^2}{2}$$
$$= -\sum_{i=1}^{n} \frac{(y_i - \beta_0 - \beta_1 X_{1i} - \beta_2 X_{2i})^2}{2}$$

似然函数求偏导:

$$\sum_{i=1}^{n} y_i - n\beta_0 - (n_{10} + n_{11})\beta_1 - (n_{01} + n_{11})\beta_2 = 0$$

$$\sum_{X_{1i}=1} y_i - (n_{10} + n_{11})\beta_0 - (n_{10} + n_{11})\beta_1 - n_{11}\beta_2 = 0$$

$$\sum_{X_{1i}=1} y_i - (n_{01} + n_{11})\beta_0 - n_{11}\beta_1 - (n_{01} + n_{11})\beta_2 = 0$$

最小二乘方法:

$$\hat{\beta} = (X^T X)^{-1} X^T Y$$

$$= \begin{bmatrix} n & n_{10} + n_{11} & n_{01} + n_{11} \\ n_{10} + n_{11} & n_{10} + n_{11} & n_{11} \\ n_{01} + n_{11} & n_{11} & n_{01} + n_{11} \end{bmatrix}^{-1} \begin{bmatrix} \sum_{i} y_i \\ \sum_{x_1=1} y_i \\ \sum_{x_2=1} y_i \end{bmatrix}$$

线性模型有交互作用情况

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_{12} X_1 X_2 + \epsilon$$

 $\epsilon \sim N(0,1)$, 似然函数:

$$L(\beta; y) \propto -\sum_{i=1}^{n} \frac{(y_{i} - \beta_{0} - \beta_{1}X_{1i} - \beta_{2}X_{2i} - \beta_{12}X_{1i}X_{2i})^{2}}{2}$$

似然函数求偏导:

$$\sum_{i=1}^{n} y_i - n\beta_0 - (n_{10} + n_{11})\beta_1 - (n_{01} + n_{11})\beta_2 - n_{11}\beta_{12} = 0$$

$$\sum_{X_{1i}=1} y_i - (n_{10} + n_{11})\beta_0 - (n_{10} + n_{11})\beta_1 - n_{11}\beta_2 - n_{11}\beta_{12} = 0$$

$$\sum_{X_{2i}=1} y_i - (n_{01} + n_{11})\beta_0 - n_{11}\beta_1 - (n_{01} + n_{11})\beta_2 - n_{11}\beta_{12} = 0$$

$$\sum_{X_{1i}=1, X_{2i}=1} y_i - n_{11}\beta_0 - n_{11}\beta_1 - n_{11}\beta_2 - n_{11}\beta_{12} = 0$$

极大似然估计:

$$\widehat{\beta}_{0} = \frac{1}{n_{00}} \sum_{X_{1i}=0, X_{2i}=0} y_{i}$$

$$\widehat{\beta}_{1} = \frac{1}{n_{10}} \sum_{X_{1i}=1, X_{2i}=0} y_{i} - \frac{1}{n_{00}} \sum_{X_{1i}=0, X_{2i}=0} y_{i}$$

$$\widehat{\beta}_{2} = \frac{1}{n_{01}} \sum_{X_{1i}=0, X_{2i}=1} y_{i} - \frac{1}{n_{00}} \sum_{X_{1i}=0, X_{2i}=0} y_{i}$$

$$\widehat{\beta}_{12} = \frac{1}{n_{11}} \sum_{X_{1i}=1, X_{2i}=1} y_{i} - \frac{1}{n_{10}} \sum_{X_{1i}=1, X_{2i}=0} y_{i} - \frac{1}{n_{01}} \sum_{X_{1i}=0, X_{2i}=1} y_{i}$$

$$+ \frac{1}{n_{00}} \sum_{X_{1i}=0, X_{2i}=0} y_{i}$$

当没有交互作用项时, $\hat{\beta}_{12}=0$

$$\frac{1}{n_{11}} \sum_{X_{1i}=1, X_{2i}=1} y_i - \frac{1}{n_{10}} \sum_{X_{1i}=1, X_{2i}=0} y_i$$

$$= \frac{1}{n_{01}} \sum_{X_{1i}=0, X_{2i}=1} y_i - \frac{1}{n_{00}} \sum_{X_{1i}=0, X_{2i}=0} y_i$$

$$\frac{1}{n_{11}} \sum_{X_{1i}=1, X_{2i}=1} y_i - \frac{1}{n_{01}} \sum_{X_{1i}=0, X_{2i}=1} y_i$$

$$= \frac{1}{n_{10}} \sum_{X_{1i}=1, X_{2i}=1} y_i - \frac{1}{n_{00}} \sum_{X_{1i}=0, X_{2i}=0} y_i$$

$$\frac{1}{n_{11}} \sum_{X_{1i}=1, X_{2i}=1} y_i + \frac{1}{n_{00}} \sum_{X_{1i}=0, X_{2i}=0} y_i$$

$$= \frac{1}{n_{01}} \sum_{X_{1i}=0, X_{2i}=1} y_i + \frac{1}{n_{10}} \sum_{X_{1i}=1, X_{2i}=0} y_i$$

方差分析

$$y_{ijl} = u + a_i + b_j + (ab)_{ij} + \epsilon_{ijl}, i = 0, 1; j = 0, 1; l = 1, ..., n_{ij}.$$

假设
$$\sum a_i = 0$$
, $\sum b_j = 0$, $\sum_i (ab)_{ij} = \sum_j (ab)_{ij} = 0$ 。

假设 $a_0=0$, $b_0=0$, $(ab)_{00}=(ab)_{01}=(ab)_{11}=0$, 则是我们上面讨论的情况。

方差分析的估计:

$$\begin{split} \widehat{\boldsymbol{u}} &= \overline{\boldsymbol{y}}, \quad \widehat{\boldsymbol{a}}_i = \overline{\boldsymbol{y}}_{i.} - \overline{\boldsymbol{y}}, \quad \widehat{\boldsymbol{b}}_j = \overline{\boldsymbol{y}}_{.j} - \overline{\boldsymbol{y}}, \\ \widehat{\boldsymbol{(ab)}}_{ij} &= \overline{\boldsymbol{y}}_{ij} - \overline{\boldsymbol{y}}_{i.} - \overline{\boldsymbol{y}}_{.j} + \overline{\boldsymbol{y}} \end{split}$$

注意: 无交互作用和有交互作用的估计值一致。

 X_1 、 X_2 、Y均为二分类变量,令 Y_{ij} 为 $X_1=i$, $X_2=j$ 时Y的第I次观测,则 $Y_{ij}\sim B(1,p_{ij})$,其中i=0,1,j=0,1, $l=1,...n_{ij}$. 令 Y_{ij} 为 $X_1=i$, $X_2=j$ 时Y=1的总次数,则 $Y_{ij}\sim B(n_{ij},p_{ij})$ 采用logistic模型,在无交互作用项时,有

$$logit(p_{ij}) = \beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2}$$

在有交互作用项时,有

$$logit(p_{ij}) = \beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2} + \beta_{12} x_{ij1} \cdot x_{ij2}$$

比较两个模型系数间的差异,及交互作用项的含义。

由于 $Y_{ij} \sim B(n_{ij}, p_{ij})$,采用极大似然估计,对数似然函数为:

$$L \propto \sum_{i} \sum_{j} y_{ij} ln p_{ij} + (n_{ij} - y_{ij}) ln (1 - p_{ij})$$

$$\frac{\partial I}{\partial p_{ij}} = \frac{y_{ij}}{p_{ij}} - \frac{n_{ij} - y_{ij}}{1 - p_{ij}} = 0$$

则 p_{ij} 的MLE为

$$\hat{p}_{ij} = \frac{y_{ij}}{n_{ij}}$$

在无交互作用项时,
$$logit(p_{ij}) = \beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2}$$

$$p_{ij} = \frac{exp(\beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2})}{1 + exp(\beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2})}$$

则有:

$$\begin{cases} \frac{y_{00}}{n_{00}} = \frac{exp(\beta_0)}{1 + exp(\beta_0)} \\ \frac{y_{01}}{n_{01}} = \frac{exp(\beta_0 + \beta_2)}{1 + exp(\beta_0 + \beta_2)} \\ \frac{y_{10}}{n_{10}} = \frac{exp(\beta_0 + \beta_1)}{1 + exp(\beta_0 + \beta_1)} \\ \frac{y_{11}}{n_{11}} = \frac{exp(\beta_0 + \beta_1 + \beta_2)}{1 + exp(\beta_0 + \beta_1 + \beta_2)} \end{cases}$$

化简得:

$$\exp(\beta_0) = \frac{y_{00}}{n_{00} - y_{00}} \quad \exp(\beta_0 + \beta_2) = \frac{y_{01}}{n_{01} - y_{01}}$$

$$\exp(\beta_0 + \beta_1) = \frac{y_{10}}{n_{10} - y_{10}} \quad \exp(\beta_0 + \beta_1 + \beta_2) = \frac{y_{11}}{n_{11} - y_{11}}$$

解得:

$$\begin{cases} exp(\beta_0) = \frac{y_{00}}{n_{00} - y_{00}} \\ exp(\beta_1) = \frac{y_{11}(n_{01} - y_{01})}{(n_{11} - y_{11})y_{01}} = \frac{y_{10}(n_{00} - y_{00})}{(n_{10} - y_{10})y_{00}} \\ exp(\beta_2) = \frac{y_{11}(n_{10} - y_{10})}{(n_{11} - y_{11})y_{10}} = \frac{y_{01}(n_{00} - y_{00})}{(n_{01} - y_{01})y_{00}} \end{cases}$$

在无交互作用项时:

$$\begin{cases} \exp(\beta_0) = \frac{p_{00}}{1 - p_{00}} \\ \exp(\beta_1) = \frac{p_{11}/(1 - p_{11})}{p_{01}/(1 - p_{01})} = \frac{p_{10}/(1 - p_{10})}{p_{00}/(1 - p_{00})} \\ \exp(\beta_2) = \frac{p_{11}/(1 - p_{11})}{p_{10}/(1 - p_{10})} = \frac{p_{01}/(1 - p_{01})}{p_{00}/(1 - p_{00})} \end{cases}$$

在有交互作用项时,

$$logit(p_{ij}) = \beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2} + \beta_{12} x_{ij1} \cdot x_{ij2}$$

$$\rho_{ij} = \frac{exp(\beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2} + \beta_{12} x_{ij1} \cdot x_{ij2})}{1 + exp(\beta_0 + \beta_1 x_{ij1} + \beta_2 x_{ij2} + \beta_{12} x_{ij1} \cdot x_{ij2})}$$

则有:

$$\begin{cases} \frac{y_{00}}{n_{00}} = \frac{\exp(\beta_0)}{1 + \exp(\beta_0)} \\ \frac{y_{01}}{n_{01}} = \frac{\exp(\beta_0 + \beta_2)}{1 + \exp(\beta_0 + \beta_2)} \\ \frac{y_{10}}{n_{10}} = \frac{\exp(\beta_0 + \beta_1)}{1 + \exp(\beta_0 + \beta_1)} \\ \frac{y_{11}}{n_{11}} = \frac{\exp(\beta_0 + \beta_1 + \beta_2 + \beta_{12})}{1 + \exp(\beta_0 + \beta_1 + \beta_2 + \beta_{12})} \end{cases}$$

化简得:

$$\exp(\beta_0) = \frac{y_{00}}{n_{00} - y_{00}} \quad \exp(\beta_0 + \beta_2) = \frac{y_{01}}{n_{01} - y_{01}}$$

$$\exp(\beta_0 + \beta_1) = \frac{y_{10}}{n_{10} - y_{10}} \quad \exp(\beta_0 + \beta_1 + \beta_2 + \beta_{12}) = \frac{y_{11}}{n_{11} - y_{11}}$$

解得:

$$\begin{cases} exp(\beta_0) = \frac{y_{00}}{n_{00} - y_{00}} \\ exp(\beta_1) = \frac{y_{10}(n_{00} - y_{00})}{(n_{10} - y_{10})y_{00}} \\ exp(\beta_2) = \frac{y_{01}(n_{00} - y_{00})}{(n_{01} - y_{01})y_{00}} \\ exp(\beta_{12}) = \frac{y_{11}(n_{10} - y_{10})(n_{01} - y_{01})y_{00}}{(n_{11} - y_{11})y_{10}y_{01}(n_{00} - y_{00})} \end{cases}$$

在有交互作用项时:

$$\begin{cases} exp(\beta_0) = \frac{p_{00}}{1 - p_{00}} \\ exp(\beta_1) = \frac{p_{10}/(1 - p_{10})}{p_{00}/(1 - p_{00})} \\ exp(\beta_2) = \frac{p_{01}/(1 - p_{01})}{p_{00}/(1 - p_{00})} \\ exp(\beta_{12}) = \frac{(p_{11}/(1 - p_{11})) \times (p_{00}/(1 - p_{00}))}{(p_{10}/(1 - p_{10})) \times (p_{01}/(1 - p_{01}))} \end{cases}$$

若无交互作用,则有 $exp(\beta_{12}) = 1$,即:

$$(p_{11}/(1-p_{11}))\times(p_{00}/(1-p_{00}))=(p_{10}/(1-p_{10}))\times(p_{01}/(1-p_{01}))$$

比值比

$$\exp(\beta_0) = \frac{\rho_{00}}{1 - \rho_{00}} \quad \exp(\beta_0 + \beta_2) = \frac{\rho_{01}}{1 - \rho_{01}}$$

$$\exp(\beta_0+\beta_1) = \frac{p_{10}}{1-p_{10}} \quad \exp(\beta_0+\beta_1+\beta_2+\beta_{12}) = \frac{p_{11}}{1-p_{11}}$$

则有:

$$exp(\beta_0) = OR_{00} = 1$$
 $exp(\beta_2) = OR_{01}$

$$exp(\beta_1) = OR_{10} \quad exp(\beta_{12}) = \frac{OR_{11}}{OR_{10} \times OR_{01}}$$



交互效应

$$logit(p) = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + \beta_{12} x_1 \cdot x_2$$

有:

$$logit(p) = \beta_0 + (\beta_1 + \beta_{12}x_2)x_1 + \beta_2x_2$$

或:

$$logit(p) = \beta_0 + \beta_1 x_1 + (\beta_2 + \beta_{12} x_1) x_2$$

即: X_1 对Y的作用被 X_2 影响或 X_2 对Y的作用被 X_1 影响,产生交互作用