

Generalized Linear Models and their Applications

STAT 431/STAT 831*

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Contents

Topic 1a: Review of Linear Regression	3
Topic 1b: Review of Likelihood Methods	12
Topic 2a: Formulation of Generalized Linear Models	19
Topic 2a: Maximum Likelihood Estimation for Generalized Linear Models	24
Topic 3a: Binary Data and Odds Ratios	29

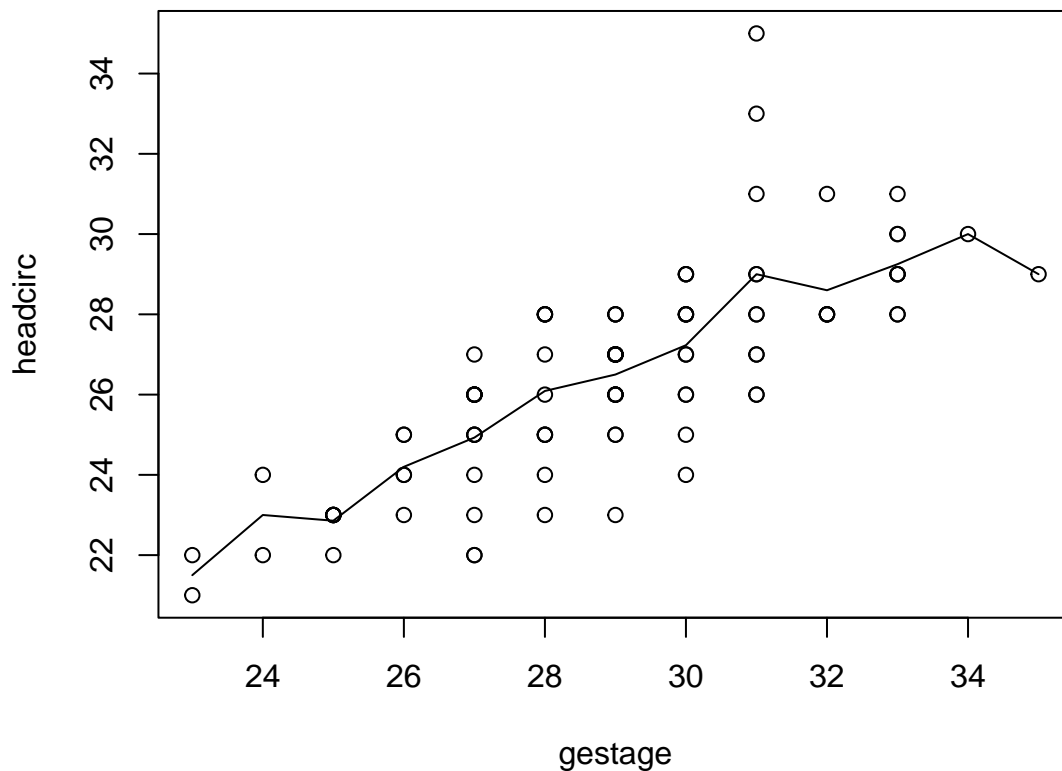
Topic 1a: Review of Linear Regression

EXAMPLE: LOW BIRTHWEIGHT INFANTS STUDY¹

A study was conducted at two teaching hospitals in Boston, Massachusetts, where the head circumference, gestational age and some other variables are recorded for 100 low birth weight infants.

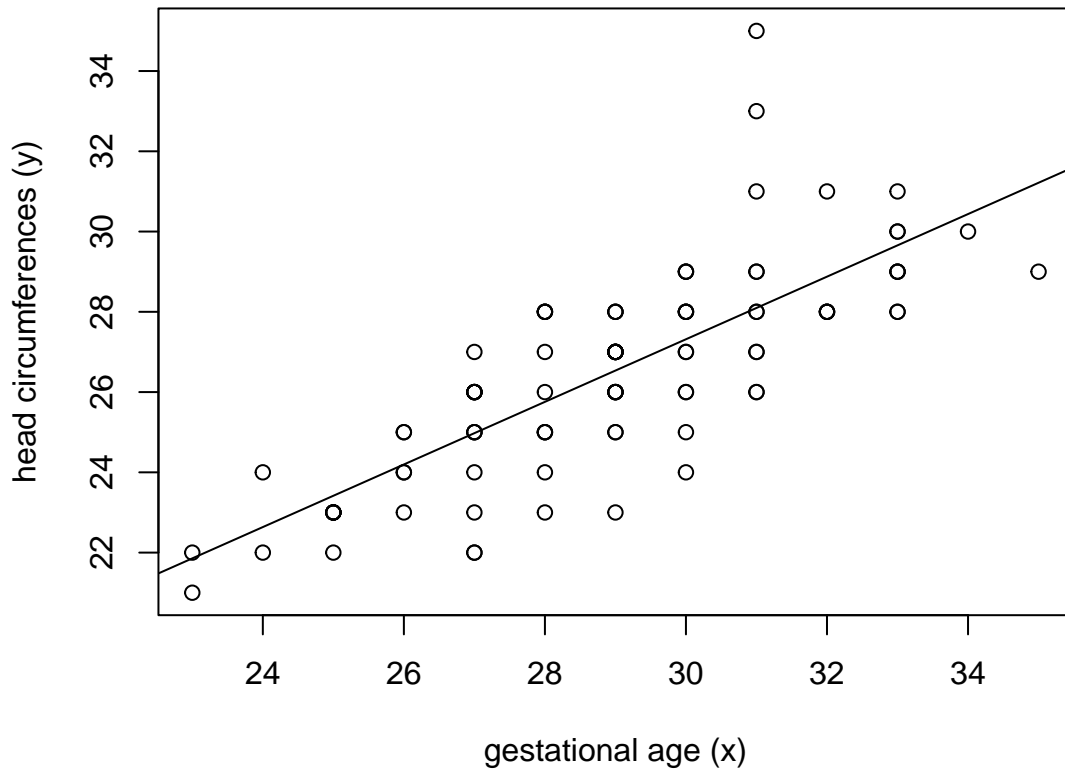
Question: what is the relationship between *gestational age* & *head circumference*?

A Scatterplot of the Data



We wish to model the relationship between *gestational age* and *head circumference* using a straight line!

¹Principles of Biostatistics 2nd Edition by Marcello Pagano, Kimberlee Gauvreau.



THE MODEL FITTING PROCESS

- ① **Model Specification:** select a probability distribution for the response variable and a linear equation linking the response to the explanatory variables.
- ② **Estimation:** finding the equation (the parameters of the model).
- ③ **Model checking:** how well does the model fit the data?
- ④ **Inference:** interpret the fitted model, calculate confidence intervals, conduct hypothesis tests.

① MODEL SPECIFICATION

Notation

For each subject $i = 1, \dots, n$ we have:

- Y_i = random variable representing the response, and
- $\mathbf{x}_i = (1, x_{i1}, \dots, x_{ip})^\top$ a vector of explanatory variables.

Specification for Multiple Linear Regression

- Linear regression equation:

$$Y_i = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip} + \varepsilon_i \text{ where } \varepsilon_i \stackrel{\text{iid}}{\sim} \mathcal{N}(0, \sigma^2).$$

- Equivalently, Y_i 's are independent $\mathcal{N}(\mu_i, \sigma^2)$ random variables or

$$\mu_i = \mathbb{E}[Y_i] = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip}.$$

- For convenience, we often write linear regression models in matrix form as

$$\mathbf{Y} = \mathbf{X}\boldsymbol{\beta} + \boldsymbol{\varepsilon},$$

where

$$\mathbf{Y} = \begin{bmatrix} Y_1 \\ Y_2 \\ \vdots \\ Y_n \end{bmatrix}, \quad \mathbf{X} = \begin{bmatrix} 1 & x_{11} & \cdots & x_{1p} \\ 1 & x_{21} & \cdots & x_{2p} \\ \vdots & \vdots & \ddots & \vdots \\ 1 & x_{n1} & \cdots & x_{np} \end{bmatrix}, \quad \boldsymbol{\beta} = \begin{bmatrix} \beta_0 \\ \beta_1 \\ \vdots \\ \beta_p \end{bmatrix}, \quad \boldsymbol{\varepsilon} = \begin{bmatrix} \varepsilon_1 \\ \varepsilon_2 \\ \vdots \\ \varepsilon_n \end{bmatrix}$$

and

$$\boldsymbol{\varepsilon} \sim \text{MVN}(\mathbf{0}, \sigma^2 \mathbf{I}).$$

2 ESTIMATION

Least Squares Method

We wish to minimize a loss function:

$$\begin{aligned} S(\boldsymbol{\beta}) &= \sum_{i=1}^n (y_i - \hat{y}_i)^2 \\ &= \sum_{i=1}^n (y_i - (\beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip}))^2 \\ &= (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta}). \end{aligned}$$

The least squares estimators (LSE) are the solutions to the equations:

$$\frac{\partial S}{\partial \boldsymbol{\beta}} = \frac{\partial}{\partial \boldsymbol{\beta}} (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta}) = 0.$$

Maximum Likelihood Method

The probability density function for Y_i is:

$$f(y_i) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp \left\{ -\frac{1}{2\sigma^2} (y_i - (\beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip}))^2 \right\}.$$

The log-likelihood function is therefore:

$$\begin{aligned}\ell(\boldsymbol{\beta}, \sigma^2) &= \log\left(\prod_{i=1}^n f(y_i)\right) \\ &= \sum_{i=1}^n \left(-\frac{1}{2} \log(2\pi\sigma^2) - \frac{1}{2\sigma^2} (y_i - (\beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip}))^2\right) \\ &= -\frac{n}{2} \log(2\sigma^2) - \frac{1}{2\sigma^2} (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta}).\end{aligned}$$

The maximum likelihood estimators (MLE) of $\boldsymbol{\beta}$ are obtained by solving:

$$\frac{\partial \ell}{\partial \boldsymbol{\beta}} = \frac{\partial}{\partial \boldsymbol{\beta}} \left[-\frac{1}{2\sigma^2} (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta})^\top (\mathbf{Y} - \mathbf{X}\boldsymbol{\beta}) \right] = 0.$$

- **Parameter Estimates:** For linear regression LSE and MLE of $\boldsymbol{\beta}$ are the same

$$\hat{\boldsymbol{\beta}} = (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \mathbf{Y}.$$

- **Fitted values:** $\hat{\mathbf{Y}} = \mathbf{X}\hat{\boldsymbol{\beta}}$.

- **Residuals:** $\hat{r}_i = (y_i - \hat{y}_i)$.

- **Variance estimates:**

- An unbiased estimate of σ^2 is:

$$\hat{\sigma}^2 = \frac{1}{n - (p + 1)} \sum_{i=1}^n \hat{r}_i^2.$$

- An estimate of the variance of $\hat{\boldsymbol{\beta}}$ is:

$$\hat{\mathbf{V}}(\hat{\boldsymbol{\beta}}) = \hat{\sigma}^2 (\mathbf{X}^\top \mathbf{X})^{-1}.$$

Low Birthweight Infant Data Example

- For $n = 100$ infants, we have observed Y_i = head circumference and x_i = gestational age for baby i , $i = 1, \dots, 100$.
- Consider a simple linear regression model:

$$Y_i = \beta_0 + \beta_1 x_i + \varepsilon_i.$$

- We can fit the model and obtain LSE/MSE using the `lm()` function in R.

```
lowbwt <- read.table("lowbwt.txt", header = T)
fit <- lm(headcirc ~ gestage, data = lowbwt)
summary(fit)
```

```
Call:
lm(formula = headcirc ~ gestage, data = lowbwt)
```

```
Residuals:
    Min       1Q   Median       3Q      Max
-3.5358 -0.8760 -0.1458  0.9041  6.9041
```

```

Coefficients:
              Estimate Std. Error t value Pr(>|t|)
(Intercept)  3.91426    1.82915    2.14   0.0348 *
gestage      0.78005    0.06307   12.37  <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 1.59 on 98 degrees of freedom
Multiple R-squared:  0.6095, Adjusted R-squared:  0.6055
F-statistic: 152.9 on 1 and 98 DF,  p-value: < 2.2e-16

```

- What is the interpretation of regression parameters β_0 and β_1 ?
 - β_0 (intercept): expected `headcirc` for a baby of a gestational age zero ($x = 0$).
 - β_1 (slope): expected change in `headcirc` associated with a one unit increase in gestational age.

③ MODEL CHECKING

Standardized Residuals:

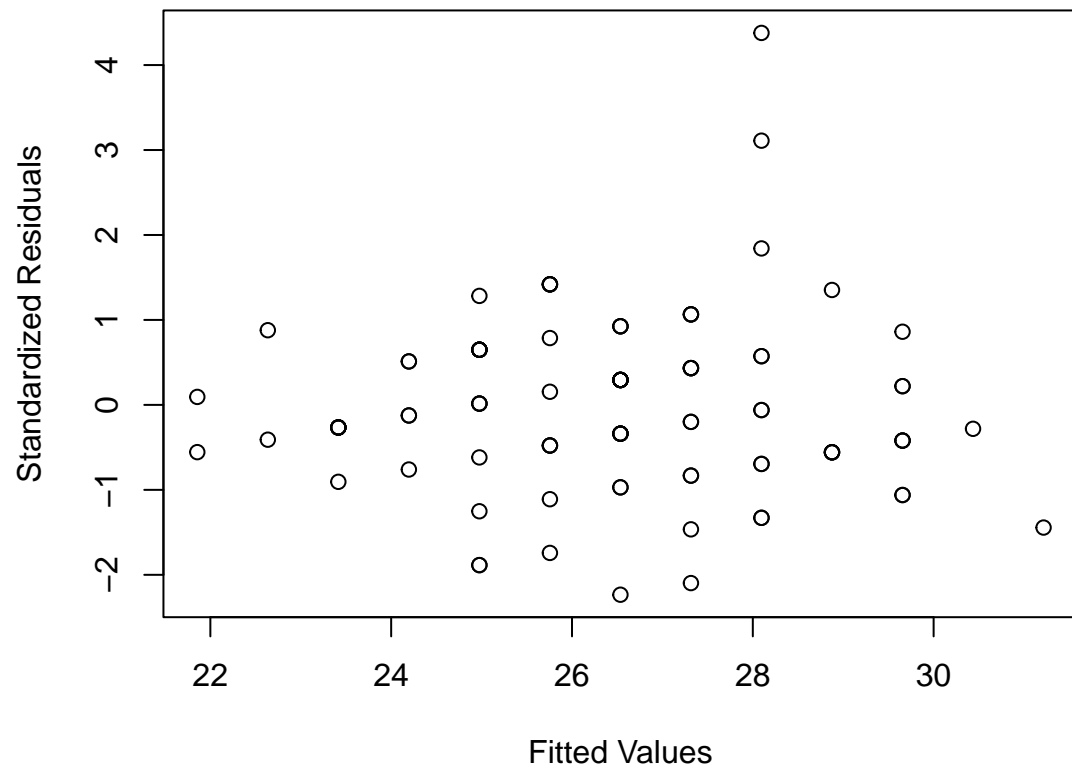
$$d_i = \frac{r_i}{\sqrt{\hat{\sigma}^2(1 - h_{ii})}},$$

where h_{ii} is the (i, i) element of $\mathbf{H} = (\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top$. By asymptotic theory, if the model provides a good fit to the data then we should expect that:

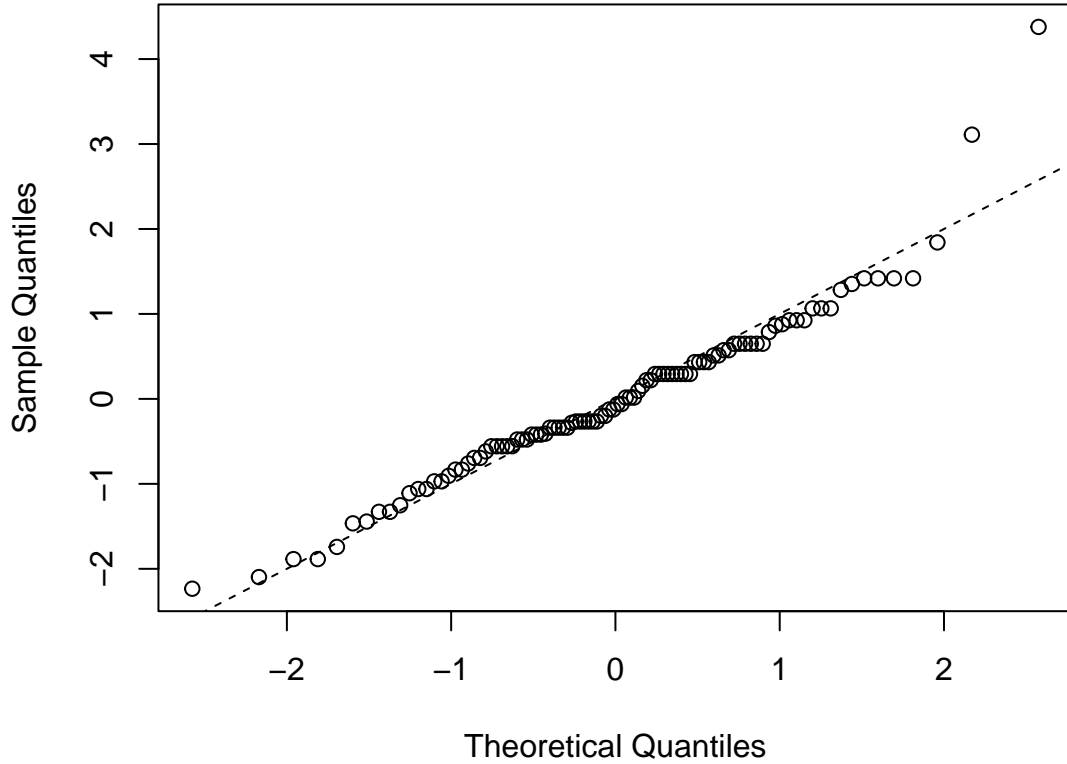
$$d_i \stackrel{\text{iid}}{\sim} \mathcal{N}(0, 1).$$

We visually check this by examining residual plots such as:

- Standardized residuals versus the fitted values.
- Standardized residuals versus the explanatory variable(s).
- Normal probability plot (QQ plot) of the standardized residuals.



Normal Q-Q Plot



④ INFERENCE

- Under suitable assumptions, the fitted regression parameters are asymptotically normally distributed:

$$\begin{aligned}\hat{\boldsymbol{\beta}} &\sim \text{MVN}(\boldsymbol{\beta}, \sigma^2(\mathbf{X}^\top \mathbf{X})^{-1}), \\ \hat{\beta}_j &\sim \mathcal{N}(\beta_j, \sigma^2 v_{jj}), \quad \text{where } v_{jj} = [(\mathbf{X}^\top \mathbf{X})^{-1}]_{(j,j)}.\end{aligned}$$

- Since σ^2 is generally unknown, we replace it with the unbiased estimate $\hat{\sigma}^2$, and obtain $\text{se}(\hat{\beta}_j) = \sqrt{\hat{\sigma}^2 v_{jj}}$.
- The inference is then based on the t -distribution result:

$$\frac{\hat{\beta}_j - \beta_j}{\text{se}(\hat{\beta}_j)} \sim t_{n-p-1}.$$

Low Birthweight Infant Data Example

- Is there a significant (linear) relationship between head circumference and gestational age?

We wish to test $H_0: \beta_1 = 0$ vs $H_A: \beta_1 \neq 0$.

$$t = \frac{\hat{\beta}_1 - (0)}{\text{se}(\hat{\beta}_1)} \sim t_{98},$$

if H_0 is true, and we reject H_0 if $|t| > t_{98,0.975} = 1.985$. Here we have $t = 0.78/0.063 = 12.37 \gg 1.985$, so we reject H_0 .

- What is the 95 % confidence interval for the expected increase in head circumference when the gestational age of a baby increases by 1 week?

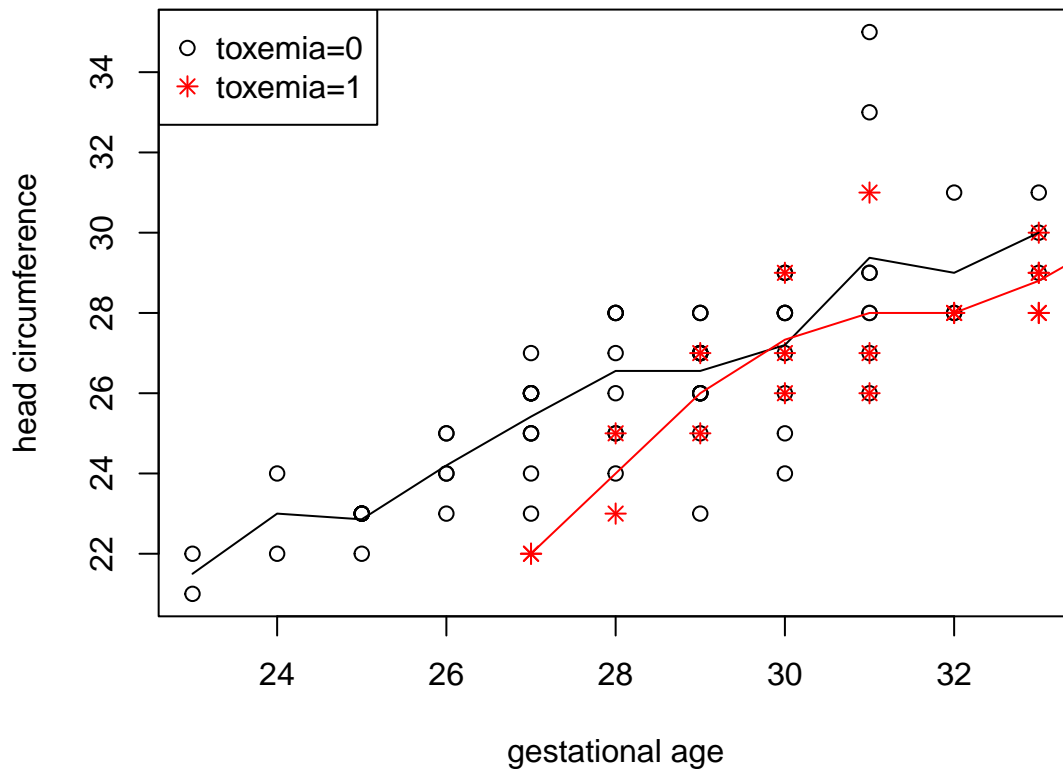
A 95 % CI for β_1 :

$$\hat{\beta}_1 \pm t_{98,0.975} \text{se}(\hat{\beta}_1) = 0.78 \pm 1.985(0.063) = (0.665, 0.905).$$

LINEAR MODELS WITH MULTIPLE PREDICTORS

Low Birthweight Infant Data Example

- *Toxemia*, a pregnancy complication characterized by high blood pressure and signs of damage to liver and kidneys, may also have an impact on the development of babies.



- Does *toxemia*, after adjustment for gestational age, also affect the head circumference?

```
fit <- lm(headcirc ~ gestage + factor(toxemia), data = lowbwt)
summary(fit)
```

Call:
lm(formula = headcirc ~ gestage + factor(toxemia), data = lowbwt)

Residuals:

Min	1Q	Median	3Q	Max
-----	----	--------	----	-----

```

-3.8427 -0.8427 -0.0525  0.8109  6.4092

Coefficients:
                Estimate Std. Error t value Pr(>|t|)
(Intercept)      1.49558    1.86799   0.801  0.42530
gestage          0.87404    0.06561  13.322 < 2e-16 ***
factor(toxemia)1 -1.41233    0.40615  -3.477  0.00076 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 1.507 on 97 degrees of freedom
Multiple R-squared:  0.6528, Adjusted R-squared:  0.6456
F-statistic: 91.18 on 2 and 97 DF,  p-value: < 2.2e-16

```

What is the interpretation of β_2 ?

$\hat{\beta}_3 = -1.41233$. After adjustment of gestational age, the babies whose mothers had toxemia have smaller (by 1.41 cm) than those whose mothers did not. This difference is significant (test $H_0: \beta_2 = 0$, p -value = 0.0076 < 0.05).

- Is the rate of increase of head circumference with gestational age the same for infants whose mothers with toxemia as those whose mother without it?

$$Y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_3 x_{i1} x_{i2} + \varepsilon_i.$$

```

fit <- lm(headcirc ~ gestage * factor(toxemia), data = lowbwt)
summary(fit)

Call:
lm(formula = headcirc ~ gestage * factor(toxemia), data = lowbwt)

Residuals:
    Min       1Q   Median       3Q      Max
-3.8366 -0.8366 -0.0928  0.7910  6.4341

Coefficients:
                Estimate Std. Error t value Pr(>|t|)
(Intercept)      1.76291    2.10225   0.839   0.404
gestage          0.86461    0.07390  11.700 <2e-16 ***
factor(toxemia)1 -2.81503    4.98515  -0.565   0.574
gestage:factor(toxemia)1  0.04617    0.16352   0.282   0.778
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 1.515 on 96 degrees of freedom
Multiple R-squared:  0.6531, Adjusted R-squared:  0.6422
F-statistic: 60.23 on 3 and 96 DF,  p-value: < 2.2e-16

```

What is the interpretation of β_3 ?

β_3 is the differences in slopes between the two groups (toxemia=1 vs toxemia=0). We want to test $H_0: \beta_3 = 0$, $t = 0.282$, p -value = 0.778 > 0.05. No evidence to reject H_0 .

LIMITATIONS OF LINEAR REGRESSION

Linear regression models can be very useful but may not be appropriate to use when response Y is not continuous and can not be assumed to be normally distributed, e.g.,

- Binary data ($Y = 0$ or $Y = 1$),
- Count data ($Y = 0, 1, 2, 3, \dots$).

Generalized Linear Models (GLM) extend the linear regression framework to address the above issue.

- Suitable for continuous and discrete data.
- Normal/Gaussian linear regression is a special case of GLM.
- Inference based on maximum likelihood methods (review next class — 431 Appendix, Stat 330 notes).

WEEK 2
13th to 17th September

Topic 1b: Review of Likelihood Methods

DISTRIBUTIONS WITH A SINGLE PARAMETER

Setup

- Suppose Y is a random variable with probability density (or mass) function $f(y; \theta)$, where $\theta \in \Omega$ is a continuous parameter.
- The true value of θ is unknown.
- We wish to make inferences about θ (i.e., we may want to estimate θ , calculate a 95 % CI or carry out tests of hypotheses regarding θ).

LIKELIHOOD FUNCTION

- The **Likelihood function** is any function which is proportional to the probability of observing the data one actually obtained, i.e.,

$$L(\theta; y) = cf(y; \theta) = cP(Y = y; \theta),$$

where c is a *proportionality constant* that does not depend on θ .

- $L(\theta; y)$ contains all the information regarding θ from the data.
- $L(\theta; y)$ ranks the various parameter values in terms of their consistency with the data.
- Since $L(\theta; y)$ is defined in terms of the random variable y , it is itself a random variable.

MAXIMUM LIKELIHOOD ESTIMATOR

- For the purposes of estimation we typically want to find θ value that makes the observed data the most likely (hence the term **maximum likelihood**).
- The **maximum likelihood estimator (MLE)** of θ is

$$\hat{\theta} = \arg \max_{\theta} L(\theta; y).$$

- Estimation becomes a simple optimization problem!

- It is often easier to work with the logarithm of the likelihood function, i.e., the **log-likelihood function**

$$\ell(\theta; y) = \log(L(\theta; y)).$$

- Equivalently, since the $\log(\cdot)$ function is monotonic, the value of θ that maximizes $L(\theta; y)$ also maximizes the log-likelihood $\ell(\theta; y)$.
- For simplicity, we drop the y and use $L(\theta) = L(\theta; y)$ and $\ell(\theta) = \ell(\theta; y)$.

A LIST OF IMPORTANT FUNCTIONS

- **Log-likelihood function:** $\ell(\theta) = \log(L(\theta))$.
- **Score function:** $S(\theta) = \frac{\partial \ell(\theta)}{\partial \theta} = \ell'(\theta)$.
- **Information function:** $I(\theta) = -\frac{\partial^2 \ell(\theta)}{\partial \theta^2} = -\ell''(\theta)$.
- **Fisher information function:** $\mathcal{I}(\theta) = \mathbb{E}[I(\theta)]$.
- **Relative likelihood function:** $R(\theta) = L(\theta)/L(\hat{\theta})$.
- **Log relative likelihood function:** $r(\theta) = \log(L(\theta)/L(\hat{\theta})) = \ell(\theta) - \ell(\hat{\theta})$.

MAXIMUM LIKELIHOOD ESTIMATION

- Want θ that maximizes $\ell(\theta)$, or equivalently solves $S(\theta) = 0$.
- Sometimes $S(\theta) = 0$ can be solved explicitly (easy in this case), but often we must solve iteratively.
- Check that the solution corresponds to a maxima of $\ell(\theta)$ by verifying the value of the second derivative at $\hat{\theta}$ is negative, or

$$I(\hat{\theta}) = -\ell''(\hat{\theta}) > 0.$$

- **Invariance property of MLEs:** if $g(\theta)$ is any function of the parameter θ , then the MLE of $g(\theta)$ is $g(\hat{\theta})$.

If $\hat{\theta}$ is the MLE of θ , then $e^{\hat{\theta}}$ is the MLE of e^{θ} .

EXAMPLE: BINOMIAL DISTRIBUTION

Example: Binomial Distribution

- A study was conducted to examine the risk for hormone use in healthy postmenopausal women.
- Suppose a group of n women received a combined hormone therapy, and were monitored for the development of breast cancer during 8.5 years followup.
- Let

$$Y_i = \begin{cases} 1 & \text{, if woman } i \text{ developed breast cancer,} \\ 0 & \text{, otherwise,} \end{cases}$$

for $i = 1, \dots, n$.

- Suppose $Y_i \stackrel{\text{iid}}{\sim} \text{Bernoulli}(\pi)$ where $\pi = P(Y_i = 1)$, then the total number of woman developed breast cancer is:

$$Y = \sum_{i=1}^n Y_i \sim \text{Binomial}(n, \pi).$$

- We wish to find the MLE of unknown parameter π (probability of cancer).

- **Likelihood function:**

$$L(\pi; y) = c P(Y = y; \pi) = \pi^y (1 - \pi)^{n-y},$$

where we take $c = 1/\binom{n}{y}$ to simplify the likelihood.

- **Log-likelihood function:**

$$\ell(\pi) = y \log(\pi) + (n - y) \log(1 - \pi).$$

- **Score function:**

$$S(\pi) = \frac{y}{\pi} - \frac{n - y}{1 - \pi}.$$

- **Maximum Likelihood Estimator:**

$$S(\pi) = 0 \implies \hat{\pi} = \frac{\sum_{i=1}^n y_i}{n} = \bar{y}.$$

- Second derivative test using **information function:**

$$I(\pi) = -\ell''(\pi) = \frac{y}{\pi^2} + \frac{n - y}{(1 - \pi)^2} > 0 \quad \forall \pi \in (0, 1).$$

Example: Hormone Therapy Data

- A group of $n = 8506$ postmenopausal women aged 50-79 received EPT and $Y = 166$ developed invasive breast cancer during the followup.
- Assume $Y \sim \text{Binomial}(n, \pi)$ with unknown parameter π .
- The **maximum likelihood estimate** of π is:

$$\hat{\pi} = \bar{y} = \frac{y}{n} = \frac{166}{8506} = 0.0195.$$

EXAMPLE: POISSON DISTRIBUTION

Suppose y_1, \dots, y_n is an iid sample from a Poisson distribution with probability mass function:

$$f(y; \lambda) = P(Y = y; \lambda) = \frac{\lambda^y e^{-\lambda}}{y!}, \quad \lambda > 0, y = 0, 1, 2, \dots$$

- **Likelihood function:**

$$L(\lambda; y_1, \dots, y_n) = \prod_{i=1}^n f(y_i; \lambda) = \frac{\lambda^{\sum y_i} e^{-n\lambda}}{\prod_i y_i!}.$$

- **Log-likelihood function:**

$$\ell(\lambda) = \left(\sum_i y_i \right) \log(\lambda) - n\lambda - \sum_{i=1}^n \log(y_i!).$$

- **Score function:**

$$S(\lambda) = \frac{\sum_i y_i}{\lambda} - n = 0 \implies \hat{\lambda} = \frac{\sum_{i=1}^n y_i}{n} = \bar{y}.$$

NEWTON RAPHSON ALGORITHM FOR FINDING MLE

- Sometimes, solving $S(\theta) = 0$ can be challenging and closed form solutions may not be obtained, iterative method need to be used to find the MLE.
- Recall **Taylor Series** expansion of a differentiable function $f(x)$ about a point a :

$$f(x) = f(a) + \frac{f'(a)}{1!}(x-a) + \frac{f''(a)}{2!}(x-a)^2 + \dots$$

- Now suppose we wish to find $\hat{\theta}$, the root of $S(\theta) = 0$ and $\theta^{(0)}$ is a guess that is “close” to $\hat{\theta}$.
- Consider the Taylor series expansion of $S(\theta)$ about $\theta^{(0)}$:

$$S(\theta) = S(\theta^{(0)}) + \frac{S'(\theta^{(0)})}{1!}(\theta - \theta^{(0)}) + \frac{S''(\theta^{(0)})}{2!}(\theta - \theta^{(0)})^2 + \dots$$

- For $|\theta - \theta^{(0)}|$ very small, the second and higher order terms can be dropped to a good approximation:

$$S(\theta) \simeq S(\theta^{(0)}) + S'(\theta^{(0)})(\theta - \theta^{(0)}).$$

$$S(\theta) \simeq S(\theta^{(0)}) - I(\theta^{(0)})(\theta - \theta^{(0)}).$$

- Then at $\theta = \hat{\theta}$,

$$S(\hat{\theta}) \simeq S(\theta^{(0)}) - I(\theta^{(0)})(\hat{\theta} - \theta^{(0)})$$

$$I(\theta^{(0)})(\hat{\theta} - \theta^{(0)}) \simeq S(\theta^{(0)})$$

$$(\hat{\theta} - \theta^{(0)}) \simeq I^{-1}(\theta^{(0)})S(\theta^{(0)})$$

$$\hat{\theta} \simeq \theta^{(0)} + I^{-1}(\theta^{(0)})S(\theta^{(0)}).$$

- This suggests a revised guess for $\hat{\theta}$ is:

$$\theta^{(1)} = \theta^{(0)} + I^{-1}(\theta^{(0)})S(\theta^{(0)})$$

Newton Raphson Algorithm for finding the MLE

- Begin with an initial estimate $\theta^{(0)}$.
- Iteratively obtain updated estimate by using:

$$\theta^{(i+1)} = \theta^{(i)} + I^{-1}(\theta^{(i)})S(\theta^{(i)}).$$

- Iteration continues until $\theta^{(i+1)} \simeq \theta^{(i)}$ within a specified tolerance.
- Then set $\hat{\theta} = \theta^{(i+1)}$, check that $I(\hat{\theta}) > 0$.

INFERENCE FOR SCALAR PARAMETERS θ

- So far we have discussed estimation of $\hat{\theta}$, next we want to conduct inference about θ , i.e., carry out hypothesis tests and construct confidence intervals of θ .
- Likelihood inference relies on the following **asymptotic distribution results**:

Useful asymptotic distributional results

- **(log) Likelihood ratio statistic:** $-2\log(R(\theta)) = -2r(\theta) \sim \chi^2_{(1)}$.
- **Score statistic:** $(S(\theta))^2/I(\theta) \sim \chi^2_{(1)}$.
- **Wald statistic:** $(\hat{\theta} - \theta)^2 I(\hat{\theta}) \sim \chi^2_{(1)}$ or $(\hat{\theta} - \theta)\sqrt{I(\hat{\theta})} \sim \mathcal{N}(0, 1)$ since $Z \sim \mathcal{N}(0, 1) \implies Z^2 \sim \chi^2_1$.

CONFIDENCE INTERVAL (CI)

Suppose we want a $100(1 - \alpha)\%$ confidence interval for θ .

- The **Likelihood ratio (LR)** based pivotal gives a confidence interval:

$$\{\theta : -2r(\theta) < \chi^2_1(1 - \alpha)\},$$

where $\chi^2_1(1 - \alpha)$ is the upper α percentage point of the χ^2_1 distribution.

- The **Wald**-based pivotal gives an interval:

$$\{\theta : (\hat{\theta} - \theta)^2 I(\hat{\theta}) < \chi^2_1(1 - \alpha)\},$$

or equivalently

$$\hat{\theta} \pm Z_{1-\alpha/2}(I(\hat{\theta}))^{-1/2},$$

where $Z_{1-\alpha/2}$ is the upper $\alpha/2$ percentage point of the standard normal.

EXAMPLE: HORMONE THERAPY DATA

Likelihood Ratio based 95 % CI: $\{\theta : -2r(\theta) < \chi^2_1(0.95)\}$ where $r(\theta) = \ell(\theta) - \ell(\hat{\theta})$.

- For the Binomial distribution: $\hat{\theta} = y/n$, and

$$r(\theta) = (y \log(\theta) + (n - y) \log(1 - \theta)) - \left(y \log\left(\frac{y}{n}\right) + (n - y) \log\left(1 - \frac{y}{n}\right) \right).$$

- To find the root of $-2r(\theta) = \chi^2_1(0.95)$:

```
y = 166
n = 8506
LRCI = function(theta, y, n) {
  -2 * (y * log(theta) + (n - y) * log(1 - theta) - y * log(y/n) -
    (n - y) * log(1 - y/n)) - qchisq(0.95, 1)
}
mle = y/n
uniroot(LRCI, c(0, mle), y = y, n = n)$root

[1] 0.01673867

uniroot(LRCI, c(mle, 1), y = y, n = n)$root

[1] 0.02260709
```


- The likelihood ratio based 95 % CI is (0.017, 0.023).

Wald based 95 % CI: $\hat{\theta} \pm Z_{0.975}(I(\hat{\theta}))^{-1/2}$.

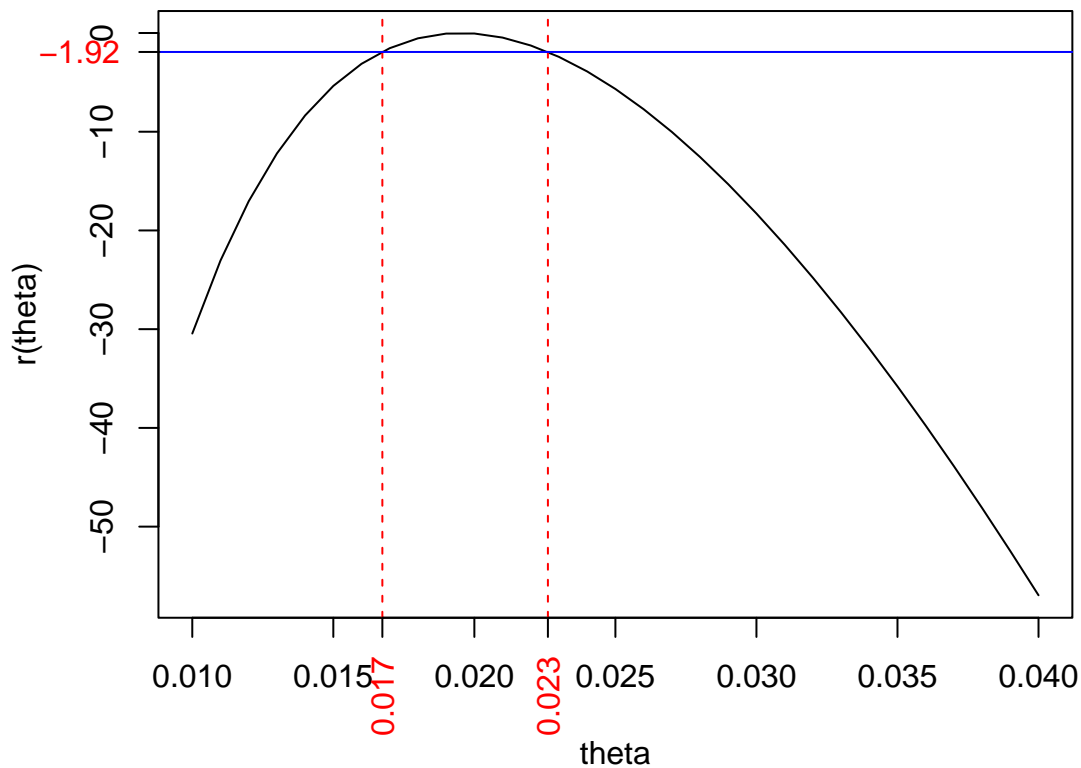
- For Binomial distribution $\hat{\theta} = y/n$ and

$$I(\hat{\theta}) = \frac{y}{\hat{\theta}^2} + \frac{n-y}{(1-\hat{\theta})^2} = n^2 \left(\frac{1}{y} + \frac{1}{n-y} \right).$$

- So we solve:

$$\begin{aligned} \hat{\theta} \pm 1.96(I(\hat{\theta}))^{-1/2} &= 0.0195 \pm 1.96(0.0015) \\ &= (0.017, 0.022). \end{aligned}$$

- The Wald based 95 % CI is: (0.017, 0.022).



HYPOTHESES TEST

Suppose we are interested in testing hypotheses:

$$H_0: \theta = \theta_0 \text{ vs } H_A: \theta \neq \theta_0.$$

- **Likelihood ratio (LR) test:** $p\text{-value} = P(\chi_1^2 > -2r(\theta_0)).$

- **Score test:** $p\text{-value} = P\left(\chi_1^2 > (S(\theta))^2 / I(\theta_0)\right)$.
- **Wald test:**

$$p\text{-value} = P\left(\chi_1^2 > (\hat{\theta} - \theta_0)^2 I(\hat{\theta})\right), \text{ or } p\text{-value} = P\left(|Z| > |\hat{\theta} - \theta_0| \sqrt{I(\hat{\theta})}\right).$$

EXAMPLE: HORMONE THERAPY DATA

Suppose we wish to test if women received EPT would have a risk of breast cancer same as that of the general population, say about 1.5 %.

$$H_0: \theta = 0.015 \text{ vs } H_A: \theta \neq 0.015.$$

- **Likelihood Ratio** based test:

$$\begin{aligned} r(\theta_0 = 0.015) &= \left(y \log(0.015) + (n - y) \log(1 - 0.015) \right) - \left(y \log\left(\frac{y}{n}\right) + (n - y) \log\left(1 - \frac{y}{n}\right) \right) \\ &= -5.3637. \end{aligned}$$

Thus, the p -value for the test is given by:

$$p = P\left(\chi_{(1)}^2 > -2r(0.015)\right) = P\left(\chi_{(1)}^2 > 10.7274\right) = 0.001.$$

Therefore, we *reject* H_0 and conclude that the risk of breast cancer for women received EPT is significantly different from 1.5 %.

NOTES ON ASYMPTOTIC INFERENCE

- Asymptotic results: approximation improves as sample size increases.
- Results are exact for a Normal linear model if θ is the mean parameter and σ^2 is known.
- **LR approach:**
 - Need to evaluate (log) likelihood at two locations.
 - Not always a closed form solution for a CI.
 - Usually the best approach.
- **Score approach:**
 - Usually the least powerful test.
 - Don't actually need to find MLE to use.
- **Wald's approach:**
 - Always get a closed form solution for a CI.
 - May not behave well for skewed likelihoods (transform?).
- All three are asymptotically equivalent!

LIKELIHOOD METHODS FOR PARAMETER VECTORS

Suppose $\boldsymbol{\theta} \in \Omega$ is a continuous $p \times 1$ parameter vector indexing a probability density (or mass) function $f(\mathbf{y}; \boldsymbol{\theta})$. The likelihood and log-likelihood functions are defined as before, but

- $\mathbf{S}(\boldsymbol{\theta}) = \frac{\partial \ell(\boldsymbol{\theta})}{\partial \boldsymbol{\theta}}$ is the $p \times 1$ **Score vector**, i.e.,

$$\mathbf{S}(\boldsymbol{\theta}) = \begin{bmatrix} \frac{\partial \ell(\boldsymbol{\theta})}{\partial \theta_1} \\ \vdots \\ \frac{\partial \ell(\boldsymbol{\theta})}{\partial \theta_p} \end{bmatrix}.$$

- $\mathbf{I}(\boldsymbol{\theta}) = -\frac{\partial^2 \ell(\boldsymbol{\theta})}{\partial \boldsymbol{\theta}^\top \partial \boldsymbol{\theta}}$ is the $p \times p$ **Information matrix**, i.e.,

$$\mathbf{I}(\boldsymbol{\theta}) = \begin{bmatrix} -\frac{\partial^2 \ell(\boldsymbol{\theta})}{\partial \theta_1^2} & -\frac{\partial^2 \ell(\boldsymbol{\theta})}{\partial \theta_1 \partial \theta_2} & \cdots & \frac{\partial^2 \ell(\boldsymbol{\theta})}{\partial \theta_1 \partial \theta_p} \\ & -\frac{\partial^2 \ell(\boldsymbol{\theta})}{\partial \theta_2^2} & \cdots & \frac{\partial^2 \ell(\boldsymbol{\theta})}{\partial \theta_2 \partial \theta_p} \\ & & \ddots & \frac{\partial^2 \ell(\boldsymbol{\theta})}{\partial \theta_p^2} \end{bmatrix}.$$

- The Newton Raphson algorithm applies as before, but with vectors and matrices as follows:

$$\boldsymbol{\theta}^{(i+1)} = \boldsymbol{\theta}^{(i)} + \mathbf{I}^{-1}(\boldsymbol{\theta}^{(i)})\mathbf{S}(\boldsymbol{\theta}^{(i)}).$$

- Again, we apply iteratively until we obtain convergence, but now check to see if $\mathbf{I}(\hat{\boldsymbol{\theta}})$ is a positive definite matrix.
- Analogs to the LR, Score and Wald results apply based on partitioning the Information matrix by $\boldsymbol{\theta} = (\boldsymbol{\alpha}, \boldsymbol{\beta})^\top$, where $\boldsymbol{\alpha}$ is a $p \times 1$ vector of nuisance parameters and $\boldsymbol{\beta}$ is a $q \times 1$ vector of parameters of interest:

$$\mathbf{I} = \mathbf{I}(\boldsymbol{\alpha}, \boldsymbol{\beta}) = \begin{pmatrix} \mathbf{I}_{\alpha\alpha}(\boldsymbol{\alpha}, \boldsymbol{\beta}) & \mathbf{I}_{\alpha\beta}(\boldsymbol{\alpha}, \boldsymbol{\beta}) \\ \mathbf{I}_{\beta\alpha}(\boldsymbol{\alpha}, \boldsymbol{\beta}) & \mathbf{I}_{\beta\beta}(\boldsymbol{\alpha}, \boldsymbol{\beta}) \end{pmatrix},$$

where $\mathbf{I}_{\alpha\alpha}(\boldsymbol{\alpha}, \boldsymbol{\beta}) = -\frac{\partial^2 \ell}{\partial \boldsymbol{\alpha} \partial \boldsymbol{\alpha}^\top}$ is $p \times p$, $\mathbf{I}_{\alpha\beta}(\boldsymbol{\alpha}, \boldsymbol{\beta}) = -\frac{\partial^2 \ell}{\partial \boldsymbol{\alpha} \partial \boldsymbol{\beta}^\top}$ is $p \times q$, $\mathbf{I}_{\beta\alpha}(\boldsymbol{\alpha}, \boldsymbol{\beta}) = -\frac{\partial^2 \ell}{\partial \boldsymbol{\beta} \partial \boldsymbol{\alpha}^\top}$ is $q \times p$, and $\mathbf{I}_{\beta\beta}(\boldsymbol{\alpha}, \boldsymbol{\beta}) = -\frac{\partial^2 \ell}{\partial \boldsymbol{\beta} \partial \boldsymbol{\beta}^\top}$ is $q \times q$.

Topic 2a: Formulation of Generalized Linear Models

THE EXPONENTIAL FAMILY

Definition (Exponential Family)

Consider a random variable Y with probability density (or mass) function $f(y; \theta, \phi)$, we say that the distribution is a member of the **exponential family** if we can write

$$f(y; \theta, \phi) = \exp \left\{ \frac{y\theta - b(\theta)}{a(\phi)} + c(y; \phi) \right\},$$

for some functions $a(\cdot)$, $b(\cdot)$, and $c(\cdot)$.

- The parameter θ is called the **canonical** parameter, and it is unknown.
- The parameter ϕ is called the **scale/dispersion** parameter, is constant, and assumed to be known.

Many well known distributions (continuous/discrete) can be shown to be a member of the exponential family.

EXAMPLES

- Poisson Distribution: $Y \sim \text{Poisson}(\lambda)$,

$$f(y; \lambda) = \frac{\lambda^y e^{-\lambda}}{y!}, \quad \lambda > 0, y = 0, 1, \dots$$

Show that Poisson is a member of exponential family and identify the canonical parameter and the functions $a(\cdot)$, $b(\cdot)$, and $c(\cdot)$.

Solution. $f(y; \lambda) = \exp\{\log(f(y; \lambda))\} = \exp\left\{\frac{y \log(\lambda) - \lambda}{1} - \log(y!)\right\}$. Therefore,

$$\begin{aligned} \theta &= \log(\lambda) && (\text{canonical/natural parameter}), \\ b(\theta) &= \lambda = e^\theta, \\ \phi &= 1, \\ a(\phi) &= 1, \\ c(y; \phi) &= -\log(y!). \end{aligned}$$

- Normal Distribution: $Y \sim \mathcal{N}(\mu, \sigma^2)$ and σ^2 known,

$$f(y; \theta, \phi) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left\{-\frac{(y - \mu)^2}{2\sigma^2}\right\}.$$

Show that this Normal distribution is a member of the exponential family.

Solution.

$$\begin{aligned} f(y; \mu, \sigma^2) &= \exp\left\{-\frac{y^2 - 2\mu y + \mu^2}{\sigma^2} - \frac{1}{2} \log(2\pi\sigma^2)\right\} \\ &= \exp\left\{\frac{y\mu - \mu^2/2}{\sigma^2} - \frac{y^2}{2\sigma^2} - \frac{1}{2} \log(2\pi\sigma^2)\right\}. \end{aligned}$$

Therefore,

$$\begin{aligned} \theta &= \mu, \\ \phi &= \sigma^2, \\ a(\phi) &= \phi = \sigma^2, \\ b(\theta) &= \frac{\mu^2}{2} = \frac{\theta^2}{2}, \\ c(y; \phi) &= -\frac{y^2}{2\sigma^2} - \frac{1}{2} \log(2\pi\sigma^2). \end{aligned}$$

PROPERTIES OF EXPONENTIAL FAMILY

Consider a single observation y from the exponential family.

$$L(\theta, \phi; y) = f(y; \theta, \phi) = \exp\left\{\frac{y\theta - b(\theta)}{a(\phi)} + c(y; \phi)\right\}.$$

$$\ell(\theta, \phi; y) = \log(f(y; \theta, \phi)) = \frac{y\theta - b(\theta)}{a(\phi)} + c(y; \phi).$$

$$S(\theta) = \frac{\partial \ell}{\partial \theta} = \frac{y - b'(\theta)}{a(\phi)}.$$

$$I(\theta) = -\frac{\partial^2 \ell}{\partial \theta^2} = \frac{b''(\theta)}{a(\phi)}.$$

$$\mathcal{I}(\theta) = \mathbb{E}\left[-\frac{\partial^2 \ell}{\partial \theta^2}\right] = I(\theta).$$

SOME GENERAL RESULTS FOR SCORE AND INFORMATION

Result # 1

The expectation of the score function is zero.

$$\mathbb{E}[S(\theta)] = 0.$$

Proof:

$$\begin{aligned} \int f(y; \theta, \phi) \, dy &= 1 \\ \frac{\partial}{\partial \theta} \int f(y; \theta, \phi) \, dy &= 0 \\ \int \frac{\partial}{\partial \theta} f(y; \theta, \phi) \, dy &= 0 \\ \int \left(\frac{\partial}{\partial \theta} \log(f(y; \theta, \phi)) \right) f(y; \theta, \phi) \, dy &= 0 \\ \int S(\theta) f(y; \theta, \phi) \, dy &= 0 \\ \mathbb{E}[S(\theta)] &= 0 \end{aligned} \tag{1}$$

Result # 2

The expectation of the score function squared is the expected information.

$$\mathbb{E}[S(\theta; y)^2] = \mathbb{E}[I(\theta; y)]$$

Proof: Differentiate (1) again,

$$\begin{aligned} \int \left(\frac{\partial}{\partial \theta} \log(f(y; \theta, \phi)) \right) f(y; \theta, \phi) \, dy &= 0 \\ \int \left(\frac{\partial^2}{\partial \theta^2} \log(f(y; \theta, \phi)) \right) f(y; \theta, \phi) \, dy + \int \left(\frac{\partial}{\partial \theta} \log(f(y; \theta, \phi)) \right) \frac{\partial}{\partial \theta} f(y; \theta, \phi) \, dy &= 0 \\ \int \frac{\partial^2}{\partial \theta^2} \log(f(y; \theta, \phi)) f(y; \theta, \phi) \, dy + \int \left(\frac{\partial}{\partial \theta} f(y; \theta, \phi) \right)^2 f(y; \theta, \phi) \, dy &= 0 \\ \int -I(\theta) f(y; \theta, \phi) \, dy + \int S(\theta)^2 f(y; \theta, \phi) \, dy &= 0 \\ \mathbb{E}[-I(\theta; y)] + \mathbb{E}[S(\theta; y)^2] &= 0 \end{aligned}$$

Now for the exponential family, we apply above results and obtain:

$$\begin{aligned} \mathbb{E}[S(\theta)] &= 0, \\ \mathbb{E}\left[\frac{Y - b'(\theta)}{a(\phi)}\right] &= 0, \\ \mathbb{E}[Y] &= b'(\theta), \\ \mathbb{E}[S(\theta)^2] &= \mathbb{E}[I(\theta)], \\ \mathbb{E}\left[\left(\frac{Y - b'(\theta)}{a(\phi)}\right)^2\right] &= \mathbb{E}\left[\frac{b''(\theta)}{a(\phi)}\right], \\ \frac{1}{a(\phi)^2} \mathbb{E}\left[(Y - \mathbb{E}[Y])^2\right] &= \frac{b''(\theta)}{a(\phi)}, \\ \text{Var}(Y) &= b''(\theta)a(\phi). \end{aligned}$$

Mean and Variance for the Exponential Family

- Mean: $\mathbb{E}[Y] = b'(\theta) = \mu$.
- Variance: $\text{Var}(Y) = b''(\theta)a(\phi)$.

Note that:

- $b'(\theta) = \mu$ tells the relationship between *canonical* parameter θ and μ .
- $b''(\theta)$ is a function of θ and hence can be also expressed as a function of μ .
- Thus, we write $b''(\theta) = V(\mu)$ and call $V(\mu)$ the **variance function**.
- Subsequently, we have:

$$\text{Var}(Y) = b''(\theta)a(\phi) = V(\mu)a(\phi),$$

which is the **mean-variance relationship** for the exponential family.

LINK FUNCTIONS

Definition (Link Function)

The **link function** relates the linear predictor $\eta = \mathbf{x}^\top \boldsymbol{\beta}$ to the expected value μ of the random variable Y , i.e.,

$$g(\mu) = \eta = \mathbf{x}^\top \boldsymbol{\beta},$$

where $g(\cdot)$ is the link function.

Definition (Canonical Link Function)

When Y is a member of the exponential family we define the **canonical link function** to be:

$$g(\mu) = \theta = \eta = \mathbf{x}^\top \boldsymbol{\beta}$$

(i.e., the choice of $g(\cdot)$ that sets canonical parameter = linear predictor).

EXAMPLES

Recall that Poisson(λ) is a member of exponential family,

$$f(y; \lambda) = \frac{\lambda^y e^{-\lambda}}{y!} = \exp\left\{\frac{y \log(\lambda) - \lambda}{1} - \log(y!)\right\}$$

where $\theta = \log(\lambda)$, $\phi = 1$, $b(\theta) = \lambda = e^\theta$, and $a(\phi) = 1$. Now to find the mean, variance function, and canonical link function:

- **Mean:** $E[Y] = b'(\theta) = e^\theta = \mu \implies \theta = \log(\mu)$.
- **Variance Function:** $V(\mu) = b''(\theta) = e^\theta \implies V(\mu) = \mu$.
- **Variance:** $\text{Var}(Y) = V(\mu)a(\phi) = \mu$ (mean-variance relationship).
- **Canonical link:** set $\theta = \eta$ using $\theta = \log(\mu) = \eta = \mathbf{x}^\top \boldsymbol{\beta}$, i.e., $g(\mu) = \log(\mu)$ where $\log(\cdot)$ is the canonical link.

Moving forward, we consider a log-linear model: $\log(\mu_i) = \mathbf{x}_i^\top \boldsymbol{\beta}$.

REMARKS ON LINK FUNCTION

- We can choose any function $g(\cdot)$ as the link function in theory.
- The canonical link is a special link function, we often choose to use canonical link for its good statistical properties.
- Context and goodness of fit should motivate the choice of link function in practice.

GENERALIZED LINEAR MODELS

Definition (Generalized Linear Model (GLM))

A **Generalized Linear Model (GLM)** is composed of three components:

- **Random Component:** The responses Y_1, \dots, Y_n are independent random variables and each Y_i is assumed to come from a parametric distribution that is a member of the exponential family.
- **Systematic Component** (or linear predictor):

$$\eta_i = \mathbf{x}_i^\top \boldsymbol{\beta},$$

a linear combination of explanatory variables \mathbf{x}_i and regression parameters $\boldsymbol{\beta}$.

- **Link function:**

$$g(\mu_i) = \eta_i = \mathbf{x}_i^\top \boldsymbol{\beta},$$

a function that relates the mean of response to the linear predictor.

TOPIC SUMMARY

1. Definition of the **Exponential Family**.
 - Exponential form of the probability density (or mass) function.
 - Derivation of Score and Information.
 - Properties of exponential family, mean-variance relationship.
 - Definition of canonical link.
2. Definition of a **Generalized Linear Model**.

Next Topic: 2b Estimation for Generalized Linear Models.

Topic 2b: Maximum Likelihood Estimation for Generalized Linear Models

GENERALIZED LINEAR MODELS

Suppose for each subject $i = 1, \dots, n$ in a random sample:

- Y_i is the response variable.
- x_{i1}, \dots, x_{ip} are explanatory variables associated with Y_i .

We consider a **Generalized Linear Model** (GLM) for the data, by definition the GLM is composed following three components:

① **Random Component:**

$$Y_i \sim \text{exponential family}, \quad Y_1, \dots, Y_n \text{ are independent.}$$

② **Systematic Component** (or linear predictor):

$$\eta_i = \mathbf{x}_i^\top \boldsymbol{\beta} = \beta_0 + \beta_1 x_{i1} + \dots + \beta_p x_{ip}.$$

- $\mathbf{x}_i = (1, x_{i1}, \dots, x_{ip})^\top$ is a covariate vector.
- $\boldsymbol{\beta} = (\beta_0, \beta_1, \dots, \beta_p)^\top$ is a vector of regression coefficients.

③ **Link function:** a function $g(\cdot)$ links $E[Y_i] = \mu_i$ to a linear prediction η_i :

$$g(\mu_i) = \eta_i = \mathbf{x}_i^\top \boldsymbol{\beta}.$$

EXAMPLE: A POISSON REGRESSION MODEL

Suppose $Y_i \stackrel{\text{ind}}{\sim} \text{Poisson}(\lambda_i)$ with mean $E[Y_i] = \lambda_i$, $i = 1, \dots, n$:

$$f(y_i) = \frac{e^{-\lambda_i} \lambda_i^{y_i}}{y_i!} = \exp\{y_i \log(\lambda_i) - \lambda_i - \log(y_i!)\}.$$

Poisson distribution is a member of exponential family with:

- Canonical parameter: $\theta_i = \log(\lambda_i)$.
- Canonical link: $\theta_i = \eta_i \implies \log(\lambda_i) = \mathbf{x}_i^\top \boldsymbol{\beta}$ (log link).

A Poisson regression model with the canonical link takes the form:

$$\log(\lambda_i) = \mathbf{x}_i^\top \boldsymbol{\beta} = \beta_0 + \beta_1 x_{i1} + \dots + \beta_p x_{ip} \quad (\text{log-linear model}).$$

EXAMPLE: A NORMAL REGRESSION MODEL

Assume $Y_i \stackrel{\text{ind}}{\sim} \mathcal{N}(\mu_i, \sigma^2)$ and σ^2 is known, $i = 1, \dots, n$:

$$\begin{aligned} f(y_i) &= (2\pi\sigma^2)^{-1/2} \exp\left\{-\frac{(y_i - \mu_i)^2}{2\sigma^2}\right\} \\ &= \exp\left\{\frac{y_i \mu_i - \mu_i^2/2}{\sigma^2} - \frac{1}{2}\left(\frac{y_i^2}{\sigma^2} + \log(2\pi\sigma^2)\right)\right\}. \end{aligned}$$

A Normal distribution (σ^2 known) is a member of exponential family with:

- Canonical parameter: $\theta_i = \mu_i$.
- Canonical link: $\theta_i = \eta_i \implies \mu_i = \mathbf{x}_i^\top \boldsymbol{\beta}$ (identity link).

A Normal regression model with the canonical link takes the form:

$$\mu_i = \mathbf{x}_i^\top \boldsymbol{\beta} = \beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip} \quad (\text{linear model}).$$

Linear regression model (STAT 331) is a Normal GLM using the canonical link!

LIKELIHOOD FOR GENERALIZED LINEAR MODELS

We wish to use likelihood methods for the estimation of the regression parameter $\boldsymbol{\beta}$ from the GLM: $g(\mu_i) = \mathbf{x}_i^\top \boldsymbol{\beta}$. Consider the log-likelihood for a *single* observation from the exponential family:

$$\ell(\theta, \phi; y) = \frac{y\theta - b(\theta)}{a(\phi)} + c(y; \phi).$$

- ℓ is a function of θ (assume that ϕ is known).
- θ is related to μ through the result:

$$\mu = b'(\theta).$$

- η can be expressed in terms of μ through the link function:

$$g(\mu) = \eta.$$

- $\boldsymbol{\beta}$ can be expressed in terms of η through the linear predictor:

$$\eta = \mathbf{x}^\top \boldsymbol{\beta}.$$

SCORE VECTOR

To find the maximum likelihood estimator for $\boldsymbol{\beta}$, we must solve $\mathbf{S}(\boldsymbol{\beta}) = \frac{\partial \ell}{\partial \boldsymbol{\beta}} = \mathbf{0}$. Consider taking derivative with respect to β_j using the chain rule:

$$\frac{\partial \ell}{\partial \beta_j} = \frac{\partial \ell}{\partial \theta} \frac{\partial \theta}{\partial \mu} \frac{\partial \mu}{\partial \eta} \frac{\partial \eta}{\partial \beta_j},$$

where

$$\frac{\partial \ell}{\partial \theta} = \frac{y - b'(\theta)}{a(\phi)},$$

$$\frac{\partial \theta}{\partial \mu} = \left(\frac{\partial \mu}{\partial \theta} \right)^{-1} = \frac{1}{b''(\theta)}$$

since $\mu = b'(\theta)$,

$$\frac{\partial \mu}{\partial \eta} = \frac{\partial \mu}{\partial \eta},$$

$$\frac{\partial \eta}{\partial \beta_j} = x_j$$

since $\eta = \beta_0 + \beta_1 x_1 + \cdots + \beta_j x_j + \cdots + \beta_p x_p$.

Hence, we have:

$$\begin{aligned}
\frac{\partial \ell}{\partial \beta_j} &= \frac{y - b'(\theta)}{a(\phi)} \frac{1}{b''(\theta)} \frac{\partial \mu}{\partial \eta} x_j \\
&= \frac{y - \mu}{\text{Var}(Y)} \frac{\partial \mu}{\partial \eta} x_j && \text{since } \mu = b'(\theta), \text{Var}(Y) = a(\phi)b''(\theta) \\
&= \frac{y - \mu}{\text{Var}(Y)} \left(\frac{\partial \mu}{\partial \eta} \right)^2 \frac{\partial \eta}{\partial \mu} x_j && \text{since } \frac{\partial \mu}{\partial \eta} \frac{\partial \eta}{\partial \mu} = 1 \\
&= (y - \mu) \left(\text{Var}(Y) \left(\frac{\partial \mu}{\partial \eta} \right)^2 \right)^{-1} \frac{\partial \eta}{\partial \mu} x_j \\
&= (y - \mu) W \frac{\partial \eta}{\partial \mu} x_j,
\end{aligned}$$

where $W^{-1} = \text{Var}(Y) \left(\frac{\partial \eta}{\partial \mu} \right)^2$. Note that generally $\frac{\partial \eta}{\partial \mu}$ is easier to calculate than $\frac{\partial \mu}{\partial \eta}$ since we define the link as $\eta = g(\mu)$.

For a random sample Y_1, \dots, Y_n from exponential family and each Y_i has a probability density function

$$f(y_i; \theta, \phi) = \exp \left\{ \frac{y_i \theta_i - b(\theta_i)}{a(\phi)} + c(y_i, \phi) \right\}.$$

We write likelihood and log-likelihood functions as:

$$\begin{aligned}
L &= \prod_{i=1}^n f(y_i; \theta_i, \phi) = \prod_{i=1}^n \exp \left\{ \frac{y_i \theta_i - b(\theta_i)}{a(\phi)} + c(y_i, \phi) \right\}, \\
\ell &= \sum_{i=1}^n \ell_i = \sum_{i=1}^n \frac{y_i \theta_i - b(\theta_i)}{a(\phi)} + c(y_i, \phi).
\end{aligned}$$

The **element of the score vector** is:

$$[\mathbf{S}(\boldsymbol{\beta})]_j = \frac{\partial \ell}{\partial \beta_j} = \sum_{i=1}^n \frac{\partial \ell_i}{\partial \beta_j} = \sum_{i=1}^n (y_i - \mu_i) W_i \frac{\partial \eta_i}{\partial \mu_i} x_{ij}$$

where $W^{-1} = \text{Var}(Y_i) \left(\frac{\partial \eta_i}{\partial \mu_i} \right)^2$, $g(\mu_i) = \eta_i = \mathbf{x}_i^\top \boldsymbol{\beta}$. In vector and matrix form we can write:

$$\mathbf{S}(\boldsymbol{\beta}) = \mathbf{X} \mathbf{W} \mathbf{A}(\mathbf{y} - \boldsymbol{\mu}),$$

where

- $\mathbf{y} = (y_1, \dots, y_n)^\top$ and $\boldsymbol{\mu} = (\mu_1, \dots, \mu_n)^\top$ are $n \times 1$ vectors,
- $\mathbf{X} = (\mathbf{x}_1, \dots, \mathbf{x}_n)$ is a $(p+1) \times n$ matrix,
- $\mathbf{W} = \text{diag}(W_1, \dots, W_n) = \begin{bmatrix} W_1 & 0 & \dots & 0 \\ \vdots & \ddots & \ddots & \vdots \\ 0 & \dots & 0 & W_n \end{bmatrix}$, and
- $\mathbf{A} = \text{diag} \left(\frac{\partial \eta_1}{\partial \mu_1}, \dots, \frac{\partial \eta_n}{\partial \mu_n} \right)$.

EXAMPLE: POISSON REGRESSION MODEL (PROBLEM 1.4)

For a random sample from Poisson distribution, $Y_i \sim \text{Poisson}(\lambda_i)$, $i = 1, \dots, n$,

$$\ell_i = \log(f(y_i; \lambda_i)) = (y_i \log(\lambda_i) - \lambda_i - \log(y_i!)).$$

Poisson regression with a log-link:

$$\log(\lambda_i) = \eta_i = \mathbf{x}_i^\top \boldsymbol{\beta}.$$

To write down the score vector for the regression coefficients $\boldsymbol{\beta}$, we may calculate the derivative using standard methods, i.e.,

$$\begin{aligned} [\mathbf{S}(\boldsymbol{\beta})]_j &= \sum_i \frac{\partial \ell_i}{\partial \beta_j} \\ &= \sum_i \frac{\partial}{\partial \beta_j} (y_i \log(\lambda_i) - \lambda_i - \log(y_i!)) \\ &= \sum_i (y_i x_{ij} - e^{\mathbf{x}_i^\top \boldsymbol{\beta}} x_{ij}). \end{aligned}$$

Or we can use the general results derived for the GLMs on the previous slides.

SOLVING $\mathbf{S}(\boldsymbol{\beta}) = \mathbf{0}$ FOR MLE

① Newton Raphson update equation is:

$$\hat{\boldsymbol{\beta}}^{(r+1)} = \hat{\boldsymbol{\beta}}^{(r)} + \mathbf{I}^{-1}(\hat{\boldsymbol{\beta}}^{(r)}) \mathbf{S}(\hat{\boldsymbol{\beta}}^{(r)}),$$

where \mathbf{I} is the observed information matrix.

- This requires us to find and repeatedly evaluate the information \mathbf{I} (possibly computationally intensive).
- Fisher suggested using the expected information matrix \mathcal{I} rather than the observed information matrix.

② Fisher Scoring update equation is:

$$\hat{\boldsymbol{\beta}}^{(r+1)} = \hat{\boldsymbol{\beta}}^{(r)} + \mathcal{I}^{-1}(\hat{\boldsymbol{\beta}}^{(r)}) \mathbf{S}(\hat{\boldsymbol{\beta}}^{(r)}).$$

INFORMATION MATRIX

Consider the (j, k) element of the Information matrix:

$$\begin{aligned} \mathbf{I}_{jk} &= -\frac{\partial^2 \ell}{\partial \beta_j \partial \beta_k} \\ &= -\frac{\partial}{\partial \beta_k} \frac{\partial \ell}{\partial \beta_j} \\ &= \sum_i -\frac{\partial}{\partial \beta_k} \left[(y_i - \mu_i) W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] \\ &= \sum_i -(y_i - \mu_i) \left\{ \frac{\partial}{\partial \beta_k} \left[W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] \right\} - W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \left(\frac{\partial}{\partial \beta_k} (y_i - \mu_i) \right) \\ &= \sum_i -(y_i - \mu_i) \left\{ \frac{\partial}{\partial \beta_k} \left[W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] \right\} + W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \frac{\partial \mu_i}{\partial \beta_k} \frac{\partial \eta_i}{\partial \mu_i} \\ &= \sum_i -(y_i - \mu_i) \frac{\partial}{\partial \beta_k} \left[W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] + x_{ij} W_i x_{ik}. \end{aligned}$$

FISHER INFORMATION

To get an element of the Expected/Fisher Information matrix:

$$\begin{aligned}
\mathcal{I}_{jk} &= \sum_i \mathbb{E} \left[-\frac{\partial^2 \ell}{\partial \beta_j \partial \beta_k} \right] \\
&= \sum_i \mathbb{E} \left[-(y_i - \mu_i) \frac{\partial}{\partial \beta_k} \left[W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] + x_{ij} W_i x_{ik} \right] \\
&= \sum_i -\mathbb{E}[(y_i - \mu_i)] \frac{\partial}{\partial \beta_k} \left[W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] + x_{ij} W_i x_{ik} \\
&= \sum_i x_{ij} W_i x_{ik}.
\end{aligned}$$

Therefore, we can write the (j, k) element of the Fisher information as:

$$\mathcal{I}_{jk} = \sum_{i=1}^n x_{ij} W_i x_{ik} = [\mathbf{X} \mathbf{W} \mathbf{X}^\top]_{jk}$$

where again, $\mathbf{W} = \text{diag}(W_1, \dots, W_n)$ and $W_i^{-1} = \text{Var}(Y_i) \left(\frac{\partial \eta_i}{\partial \mu_i} \right)^2$.

WHEN IS FISHER SCORING EQUIVALENT TO NEWTON RAPHSON?

Recall information matrix:

$$\mathbf{I}_{jk} = \sum_i -(y_i - \mu_i) \frac{\partial}{\partial \beta_k} \left[W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] + x_{ij} W_i x_{ik}.$$

Now examine:

$$\begin{aligned}
W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} &= \left(\text{Var}(Y_i) \left(\frac{\partial \eta_i}{\partial \mu_i} \right)^2 \right)^{-1} \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \\
&= \left(a(\phi) b''(\theta_i) \frac{\partial \eta_i}{\partial \mu_i} \right)^{-1} x_{ij} && \text{since } \text{Var}(Y_i) = a_i(\phi) b''(\theta_i) \\
&= \left(a(\phi) \frac{\partial \mu_i}{\partial \theta_i} \frac{\partial \eta_i}{\partial \mu_i} \right)^{-1} x_{ij} && \text{since } b'(\theta_i) = \mu_i, b''(\theta_i) = \frac{\partial \mu_i}{\partial \theta_i} \\
&= (a(\phi))^{-1} x_{ij} && \text{under the canonical link } \theta_i = \eta_i.
\end{aligned}$$

So under the **canonical link**,

$$\frac{\partial}{\partial \beta_k} \left[W_i \left(\frac{\partial \eta_i}{\partial \mu_i} \right) x_{ij} \right] = \frac{\partial}{\partial \beta_k} \left[(a(\phi))^{-1} x_{ij} \right] = 0,$$

therefore information matrix is same as the Fisher information:

$$\mathbf{I}_{jk} = \sum_i x_{ij} W_i x_{ik} = \mathcal{I}_{jk}$$

and Fisher Scoring is equivalent to Newton Raphson.

ITERATIVELY REWEIGHTED LEAST SQUARES

The Fisher Scoring is also called **iteratively reweighted least squares** (IRWLS). The reason is that the update equation can be rewritten as:

$$\hat{\beta}^{(r+1)} = \left(\mathbf{X} \mathbf{W}(\hat{\beta}^{(r)}) \mathbf{X}^\top \right)^{-1} \mathbf{X} \mathbf{W}(\hat{\beta}^{(r)}) \mathbf{Z}(\hat{\beta}^{(r)})$$

where \mathbf{Z} is a transformation of the response vector \mathbf{Y} such that:

$$\mathbf{Z} = \boldsymbol{\eta} + (\mathbf{Y} - \boldsymbol{\mu}) * \frac{\partial \boldsymbol{\eta}}{\partial \boldsymbol{\mu}}$$

- See manipulation in Section 1.2.3 of course notes.
- Same form as the weighted LS estimate of β with dependent variable \mathbf{Z} and weight matrix \mathbf{W} .
- \mathbf{Z} and \mathbf{W} are updated at each iteration.

TOPIC SUMMARY

2b Maximum Likelihood Estimation of Generalized Linear Models:

- When Y_i come from a distribution in the **exponential family**, we can use the theory of **Generalized Linear Models** to fit the regression equations of the form:

$$g(\mu_i) = \mathbf{x}_i^\top \boldsymbol{\beta}.$$

- The **link function** $g(\cdot)$ may be the canonical link, but its choice should come from model interpretation and fit.
- Can use Fisher Scoring (also known as IRWLS) to estimate the regression parameters β from any GLM based on general forms for $\mathbf{I}(\beta)$ and $\mathbf{S}(\beta)$.
- **PRACTICE**: Chapter 1 review problems.

Topic 3a: Binary Data and Odds Ratios

BINARY DATA SET-UP

Consider the simplest case with two *binary* variables:

- COVID-19: infected or not infected (response).
- Vaccination: yes or no (explanatory variable).

Use a 2×2 table to summarize the data:

Vaccination	COVID		
	infected	not infected	
yes	y_1	$m_1 - y_1$	m_1
no	y_2	$m_2 - y_2$	m_2
Total	y_\bullet	$m_\bullet - y_\bullet$	m_\bullet

Treat m_1 and m_2 as fixed, assume Y_1 and Y_2 are independent binomial r.v.'s

$$Y_k \sim \text{Bin}(m_k, \pi_k), \quad k = 1, 2,$$

where $\pi_k = P(\text{infection} \mid \text{group } k)$.

How do we measure the associate between COVID-19 infection and vaccination?

MEASURES OF ASSOCIATION

Definition (Odds Ratio)

The **Odds Ratio** (OR) is the ratio of the odds of an event occurring in one group to the odds of the event in another group (e.g., not vaccinated):

$$\text{Odds Ratio} = \frac{\pi_1/(1-\pi_1)}{\pi_2/(1-\pi_2)}.$$

Interpretation of OR:

$$\begin{array}{llll} \pi_1 = \pi_2 & \implies & \text{OR} = 1 & \implies \text{equal risk (no association)} \\ \pi_1 > \pi_2 & \implies & \text{OR} > 1 & \implies \text{higher risk in group 1} \\ \pi_1 < \pi_2 & \implies & 0 < \text{OR} < 1 & \implies \text{higher risk in group 2} \end{array}$$

Relative Risk (RR)

The **Relative Risk** (RR) is the ratio of the probability of an event occurring in one group versus another group:

$$\text{Relative Risk} = \frac{\pi_1}{\pi_2}$$

In the case of a **rare disease** (i.e., when π_1 and π_2 are very small),

$$\text{OR} = \frac{\pi_1/(1-\pi_1)}{\pi_2/(1-\pi_2)} = \frac{\pi_1}{\pi_2} \underbrace{\left(\frac{1-\pi_2}{1-\pi_1} \right)}_{\approx 1} \approx \frac{\pi_1}{\pi_2} = \text{RR},$$

then

$$\text{OR} \approx \text{RR}.$$

MAXIMUM LIKELIHOOD ESTIMATION OF ODDS RATIO

Goal: Estimate odds ratio $\psi = \frac{\pi_1/(1-\pi_1)}{\pi_2/(1-\pi_2)}$ using likelihood method. Based on “grouped” binomial data, $Y_k \sim \text{Bin}(m_k, \pi_k)$, $k = 1, 2$,

$$\begin{aligned} L(\pi_1, \pi_2) &= \binom{m_1}{y_1} \pi_1^{y_1} (1-\pi_1)^{m_1-y_1} \binom{m_2}{y_2} \pi_2^{y_2} (1-\pi_2)^{m_2-y_2} \\ &\propto \left(\frac{\pi_1/(1-\pi_1)}{\pi_2/(1-\pi_2)} \right)^{y_1} \left(\frac{\pi_2}{1-\pi_2} \right)^{y_2+y_1} (1-\pi_1)^{m_1} (1-\pi_2)^{m_2}. \end{aligned}$$

Note that $\pi_1, \pi_2 \in [0, 1]$ and odds ratio $\psi \in (0, \infty)$ are restricted, we consider re-parameterize:

$$\theta_1 = \log\left(\frac{\pi_1/(1-\pi_1)}{\pi_2/(1-\pi_2)}\right) = \log(\psi), \quad \theta_2 = \log\left(\frac{\pi_2}{1-\pi_2}\right),$$

and now $\theta_1, \theta_2 \in (-\infty, \infty)$.

Our re-parameterization implies:

$$\pi_1 = \frac{e^{\theta_1+\theta_2}}{1+e^{\theta_1+\theta_2}}, \quad \pi_2 = \frac{e^{\theta_2}}{1+e^{\theta_2}}.$$

Now the likelihood becomes:

$$\begin{aligned} L(\theta_1, \theta_2) &= (e^{\theta_1})^{y_1} (e^{\theta_2})^{y_1+y_2} (1+e^{\theta_1+\theta_2})^{m_1} (1+e^{\theta_2})^{-m_2}, \\ \ell(\theta_1, \theta_2) &= y_1\theta_1 + (y_1+y_2)\theta_2 - m_1 \log(1+e^{\theta_1+\theta_2}) - m_2 \log(1+e^{\theta_2}). \end{aligned}$$

The score vector is:

$$S(\theta_1, \theta_2) = \begin{pmatrix} \frac{\partial \ell}{\partial \theta_1} \\ \frac{\partial \ell}{\partial \theta_2} \end{pmatrix} = \begin{pmatrix} y_1 - m_1 \left(\frac{e^{\theta_1 + \theta_2}}{1 + e^{\theta_1 + \theta_2}} \right) \\ y_1 + y_2 - m_1 \left(\frac{e^{\theta_1 + \theta_2}}{1 + e^{\theta_1 + \theta_2}} \right) - m_2 \left(\frac{e^{\theta_2}}{1 + e^{\theta_2}} \right) \end{pmatrix}.$$

Solving $S(\theta_1, \theta_2) = \mathbf{0}$ gives us the MLEs:

$$\hat{\theta}_1 = \log \left(\frac{y_1 / (m_1 - y_1)}{y_2 / (m_2 - y_2)} \right), \quad \hat{\theta}_2 = \log \left(\frac{y_2}{m_2 - y_2} \right).$$

So by the invariance property of MLEs, we have:

$$\hat{\pi}_1 = \frac{y_1}{m_1}, \quad \hat{\pi}_2 = \frac{y_2}{m_2}, \quad \hat{\psi} = \frac{\hat{\pi}_1 / (1 - \hat{\pi}_1)}{\hat{\pi}_2 / (1 - \hat{\pi}_2)} = \frac{y_1 / (m_1 - y_1)}{y_2 / (m_2 - y_2)}.$$

INFERENCE FOR ODDS RATIO

In order to do inference we will need the Information Matrix:

$$\mathbf{I}(\theta_1, \theta_2) = \begin{bmatrix} I_{11} & I_{12} \\ I_{21} & I_{22} \end{bmatrix} \quad \text{where } I_{jk} = -\frac{\partial^2}{\partial \theta_j \partial \theta_k} \ell(\theta_1, \theta_2).$$

Here, we have:

$$\begin{aligned} I_{11} &= m_1 \left(\frac{e^{\theta_1 + \theta_2}}{(1 + e^{\theta_1 + \theta_2})^2} \right), \\ I_{12} &= I_{21} = m_1 \left(\frac{e^{\theta_1 + \theta_2}}{(1 + e^{\theta_1 + \theta_2})^2} \right), \\ I_{22} &= m_1 \left(\frac{e^{\theta_1 + \theta_2}}{(1 + e^{\theta_1 + \theta_2})^2} \right) + m_2 \left(\frac{e^{\theta_2}}{(1 + e^{\theta_2})^2} \right). \end{aligned}$$

We are interested in doing inference on $\theta_1 = \log(\psi)$ (while θ_2 is nuisance).

Recall the asymptotic distribution result of a **Wald statistic**:

Wald Statistic

For a vector $\boldsymbol{\theta} = (\theta_1, \theta_2)^\top$ where $\theta_1 = \log(\psi)$ is a scalar parameter of interest:

$$(\hat{\theta}_1 - \theta_1)^2 (I^{11}(\hat{\theta}_1, \hat{\theta}_2))^{-1} \sim \chi_{(1)}^2,$$

where I^{11} is the (1, 1) element of \mathbf{I}^{-1} evaluated at MLE $\hat{\theta}_1$ and $\hat{\theta}_2$.

- Calculation of I^{11} by using a general result:

$$\mathbf{I} = \begin{pmatrix} I_{11} & I_{12} \\ I_{21} & I_{22} \end{pmatrix}, \quad \mathbf{I}^{-1} = \begin{pmatrix} \textcolor{red}{I}^{11} & I^{12} \\ I^{21} & I^{22} \end{pmatrix}, \quad \textcolor{red}{I}^{11} = (I_{11} - I_{12} I_{22}^{-1} I_{21})^{-1}.$$

- We can use the Wald result to find a confidence interval for $\theta_1 = \log(\psi)$.

CONFIDENCE INTERVAL FOR ODDS RATIO

Here, we obtain:

$$I^{11}(\hat{\theta}_1, \hat{\theta}_2) = \frac{1}{y_1} + \frac{1}{m_1 - y_1} + \frac{1}{y_2} + \frac{1}{m_2 - y_2}.$$

Thus, a Wald-based 95 % confidence interval for $\theta_1 = \log(\psi)$ is:

$$\hat{\theta}_1 \pm 1.96 \sqrt{\frac{1}{y_1} + \frac{1}{m_1 - y_1} + \frac{1}{y_2} + \frac{1}{m_2 - y_2}} = (\hat{\theta}_{1L}, \hat{\theta}_{1U}).$$

A 95 % confidence interval for the Odds Ratio ψ is:

$$(\exp\{\hat{\theta}_{1L}\}, \exp\{\hat{\theta}_{1U}\}).$$

EXAMPLE: PRENATAL CARE FROM TWO CLINICS

Consider the data below for the relationship between:

- **Response:** Fetal Mortality.
- **Explanatory variable:** Level of Care.

Level of Care	<i>Fetal Mortality</i>		Total
	Died	Survived	
Intensive	20	316	336
Regular	46	373	419
	66	689	755

- Using the above data, we obtain MLE of odds ratio ψ :

$$\hat{\psi} = \frac{y_1/(m_1 - y_1)}{y_2/(m_2 - y_2)} = \frac{20/316}{46/373} = 0.51.$$

$\hat{\psi} = 0.51 < 1$, the risk of mortality is lower with intensive care.

- A 95 % CI for $\theta_1 = \log(\psi)$:

$$\log(0.51) \pm 1.96 \sqrt{\frac{1}{20} + \frac{1}{316} + \frac{1}{46} + \frac{1}{373}} = (-1.219, -0.127).$$

- A 95 % CI for odds ratio ψ :

$$(\exp\{-1.219\}, \exp\{-0.127\}) = (0.30, 0.89).$$

Note that the CI does not cover the value $\psi = 1$ (no association), so we reject the null hypothesis of no association between fetal mortality and level of care. In other words, there is evidence of association.

EXAMPLE: PRENATAL CARE FROM TWO CLINICS

There is an **additional explanatory variable**: Clinic (A vs B).

Prenatal Care Data Stratified by Clinic

Level of Care	<i>Clinic A</i>			<i>Clinic B</i>		
	Died	Survived	Total	Died	Survived	Total
Intensive	16	293	309	4	23	27
Regular	12	176	188	34	197	231
	28	469	497	38	220	258

- $\hat{\psi}_A = 0.80$ (0.37, 1.73) and $\hat{\psi}_B = 1.01$ (0.33, 3.10). These cover value 1, different from the results from the pooled analysis on the previous slide.
- These results do NOT agree with the results from the pooled analysis on the previous slide.

Association Between Clinic and Level of Care

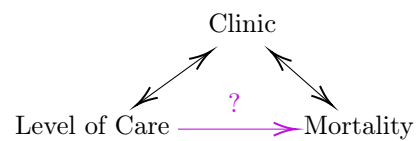
	A	B	
Intensive	309	27	336
Regular	118	231	419
	497	258	755

- $\hat{\psi} = 14.06$ (9.12, 21.76).

Association Between Clinic and Mortality

	A	B	
Died	28	38	66
Survived	469	220	689
	497	258	755

- $\hat{\psi} = 0.35$ (0.21, 0.58).
- The initial strong association between Level of Care and Infant Mortality disappeared when we stratified by clinic.



- Instead of having to examine multiple 2×2 tables we'd like to estimate the OR and compute associations using a multiple regression model.
- One way to do this is by fitting a Binomial GLM to the data.