STA 303/1002-Methods of Data Analysis II Sections L0101& L0201, Winter 2019

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STA 303/1002: Class 11- Binomial Logistic Regression

- ► Case Study IV: Island size and bird extinction
 - ► R syntax
 - Data visualization
 - Interpreting coefficients
 - Wald procedures
- Principle of the week: K-Keep, I-It, S-Simple, S-Stupid(US Navy, 1960)



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Suppose $Y \sim \mathsf{Binomial}(m, \pi)$

Y-binomial count of the number of "successes"

$$P(Y = y) = {m \choose y} \pi^{y} (1 - \pi)^{m-y}, \ y = 0, 1, ..., m$$

Link to Bernoulli: $Y = \sum_{i=1}^{m} X_i$ if X_i 's are independent Bernoulli (π) r.v.s. Assume that π is the same for each Bernoulli trial.

Mean: $E(Y)=m\pi$ Variance: $Var(Y)=m\pi(1-\pi)$

Suppose $Y \sim \mathsf{Binomial}(m, \pi)$

Consider modelling

$$\frac{Y}{m}$$

- the proportion of "successes" out of *m* independent Bernoulli trials.
- where,

$$\blacktriangleright \ \mathsf{E}\bigg(\frac{Y}{m}\bigg) = \pi$$

$$Var\left(\frac{Y}{m}\right) = \frac{\pi(1-\pi)}{m}$$

Case Study IV Data Example

@1949

▶ Data: counts of bird species for 18 Krunnit Islands off Finland.

| <u> </u> | | | | |
|-----------------|----------------|----------|-----------------------|--|
| i =1,,18 | X _i | m_i | y _i /e'las | |
| | area | nspecies | s nextinct | |
| ISLAND | AREA | ATRISK | EXTINCT | |
| Ulkokrunni | 185.8 | 75 | 5 | |
| Maakrunni | 105.8 | 67 | 3 | |
| Ristikari | 30.7 | 66 | 10 | |
| Isonkivenletto | 8.5 | 51 | 6 | |
| | | | | |
| Tiirakari | 0.2 | 40 | 13 | |
| Ristikarenletto | 0.07 | 6 | 3 | |
| | | | | |

- ► AREA- area of island in km^2 , x_i
- ▶ ATRISK- number of species on each island in 1949, m;
- **EXTINCT** number of species no longer found on each island in 1959, y_i

Case Study IV: Model

- 71 1st island
- π_i probability of 'extinction' for each island.

 Assume that this is the same for each species of bird on a particular island.
- Assume species survival is independent. Then

$$Y_i \sim Binomial(m_i, \pi_i)$$

▶ Unlike Case III- Donner party binary logistic example, we can estimate π_i from the data.

Case Study IV: Model

Observed response proportion:

$$\bar{\pi}_{i} = \frac{y_i}{m_i}$$

Observed or Empirical logits: (S-"saturated")

$$\log\left(\frac{\bar{\pi}_{S,i}}{1-\bar{\pi}_{S,i}}\right) = \log\left(\frac{y_i}{m_i - y_i}\right)$$

► Proposed Model:

$$\left|\log\left(rac{\pi_i}{1-\pi_i}
ight)=eta_0+eta_1Area_i,\ i=1,\ldots,18$$





- Learn how to create nature preserves that help endangered
 - species.
- Are large or small preserves better?

Case Study IV: Initial assessment of data

brgit (17) Area

- Plot observed logits versus area to see if a linear relationship seems appropriate.
- ▶ From that plot, we decide to look at log(Area) instead.
- The relationship between empirical logits and log(Area) seems linear.
- ► Hence, we fit

$$\left|\log\left(\frac{\pi_i}{1-\pi_i}\right)=eta_0+eta_1\log(Area_i), \right| \ i=1,\ldots,18$$

Case Study IV: R syntax

▶ In R, the model formula has the form:

$$\texttt{cbind}(\texttt{y}_{\texttt{i}}, \texttt{m}_{\texttt{i}} - \texttt{y}_{\texttt{i}}) \sim \texttt{log}(\texttt{Area})$$

Need to specify both:

- \triangleright y_i number of successes and
- $ightharpoonup (m_i y_i)$ number of failures

Case Study IV: Model Summary

- Number of observations: 18 islands
- Number of coefficients: 2

► Fitted model:

logit
$$(\hat{\pi}) = -1.196 - 0.297 \log(Area)$$

Case Study IV: Wald procedures

(Similar test as in binary logistic regression)

Hypotheses:

$$H_0: \beta_1 = 0 \text{ vs } H_a: \beta_1 \neq 0$$

► Test statistic:

$$z = \frac{-0.2971}{0.0549} = -5.42 \sim N(0, 1) \text{ or } z^2 = 29.3 \sim \chi_1^2$$

- ▶ P-value < 0.0001
- ► Conclusion: Strong evidence that coefficient of log(Area) is not zero. Evidence that extinction probabilities are associated with island area.
- ▶ 95% CI for β_1 :

Binomial Logistic Regression

$$-0.2971 \pm 1.96(0.0549) = (-0.40, -0.19)$$

not incl. 0

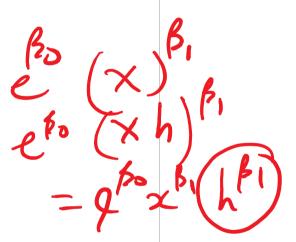


Case Study IV: Interpretation of β_1

► Model:

$$\operatorname{logit}(\pi) = eta_0 + eta_1 \operatorname{log}(x) \ \Longrightarrow \ rac{\pi}{1-\pi} = e^{eta_0} e^{eta_1 \operatorname{log}(x)} = e^{eta_0} x^{eta_1}$$

▶ Interpretation: Hence, changing x by a factor of h, changes the odds by a multiplicative factor of h^{β_1} .



Case Study IV: Interpretation of β_1

Example 1: Halving island area changes odds by a factor of $0.5^{-0.2971} = 1.23$.

Therefore, the odds of extinction on a smaller island are 123% of the odds of extinction on an island double its size.

In other words, halving of area is associated with an increase in the odds of extinction by an estimated 23%. An approximate 95% confidence interval for the percentage change in odds is 14% to 32%.

Example 2: Doubling island area changes odds by a factor of $2^{-0.2971} = 0.81$.

Therefore, the odds of extinction for an at-risk species on a larger island are only 81% of the odds of extinction for such a species on an island half its size.

メート メンニースト (一つ)



Case Study IV: Estimating probability of extinction



- Q: Estimate the probability of extinction for a species on the Ulkokrunni island.
- Fitted Model (M):

$$logit(\hat{\pi}_{M,i}) = -1.196 - 0.297 log(Area_i)$$

- For Ulkokrunni island, i = 1 and Area=185.5 km^2 , then $\log it(\hat{\pi}_{M,1}) = -1.176 0.277 (rg (185.5) = <math>\pi_{M,1} = 0.06 = 2$
- ▶ Compared to the response proportion, $\bar{\pi}_{S,1} = \frac{5}{75} = 0.067$.

Checking Model Assumptions

Model Assumptions for Binomial Logistic Regression

- 1. Underlying probability model for response is Binomial.
 - Variance is not constant; is a function of the mean.
- 2. Observations are independent.
 - 3. The form of the model is correct
 - Linear relationship between logits and explanatory variables
 - All relevant variables are included; irrelevant ones excluded
 - 4. Sample size is large enough for valid inference-tests and Cls. (Recall large-sample properties of MLEs.)
 - Check for outliers.

What is the SATURATED Model?

Observed response proportion:

$$\bar{\pi}_i = \frac{y_i}{m_i}$$

Observed or Empirical logits: (S-"saturated")

$$\log\left(\frac{\bar{\pi}_{S,i}}{1-\bar{\pi}_{S,i}}\right) = \log\left(\frac{y_i}{m_i - y_i}\right)$$

- Fits the model exactly with the data
- Most general model possible for the data.

Which Models are often compared?

Consider one explanatory variable, X with n unique levels for the outcome, $Y \sim (Bin(m, \pi))$

Saturated (FULL) Model: as many parameter coefficients as *n*

$$logit(\widehat{\pi}) = \widehat{\alpha}_0 + \widehat{\alpha}_1 \mathbb{1}_1 + \dots + \widehat{\alpha}_{n-1} \mathbb{1}_{n-1}$$

Fitted (REDUCED) Model: nested within a FULL model; has (p+1) parameters

$$logit(\widehat{\pi}) = \widehat{\beta}_0 + \widehat{\beta}_1 X$$

NULL Model: Intercept only model

$$logit(\widehat{\pi}) = \widehat{\gamma}_0$$



Checking model adequacy: Form of the model

Deviance Goodness -Of -Fit (G-O-F) Test

- ► To check model adequacy in binomial logistic regression, we can use the Deviance Goodness -Of -Fit (G-O-F) Test.
- ► Analogous to GOF test for comparing 2 models in Linear Regression.
- Form of hypotheses: H_0 : REDUCED model, H_a : FULL model
- The DEVIANCE GOF test compares the fitted model (M) to the saturated model (S).

$$H_0$$
: (Fitted) $logit(\widehat{\pi}) = \widehat{eta}_0 + \widehat{eta}_1 X$



$$H_a: (Saturated)logit(\widehat{\pi}) = \widehat{\alpha}_0 + \widehat{\alpha}_1 \mathbb{1}_1 + \cdots + \widehat{\alpha}_{n-1} \mathbb{1}_{n-1}$$

Compared to Saturated model: Deviance G-O-F test

- ▶ Uses LRT
- Sometimes called "Drop-in-Deviance" test
- as extra-sum-of-squares tests; based on the deviance residual
- Hypotheses:

$$H_0$$
: $logit(\pi) = \alpha_0 + \alpha_1 X$ (Fitted model fits data as well as Saturated model) H_a : $logit(\pi) = \beta_0 + \beta_1 \mathbb{1}_1 + \cdots + \beta_{n-1} \mathbb{1}_{n-1}$ (Saturated model is better)

► Test Statistic:

$$Deviance = -2\log\left(\frac{\mathcal{L}_R}{\mathcal{L}_F}\right) = -2\log\left(\frac{\mathcal{L}_M}{\mathcal{L}_S}\right)$$

- ▶ Under H_0 , Deviance \sim Chi-square distribution with n (p + 1) df.
- Warning: This is an asymptotic approximation, so it works better if each $m_i > 5$.)

Calculating the Deviance test statistic

Recall underlying model of $Y: Y_i \sim Binomial(m_i, \pi_i)$

$$P(Y_i = y_i) = \binom{m_i}{y_i} \pi_i^{y_i} (1 - \pi_i)^{m_i - y_i}, \ \ y_i = 0, 1, \dots, m_i$$

Hence the likelihood is:

$$\mathcal{L} = \prod_{i=1}^n \binom{m_i}{y_i} \pi_i^{y_i} (1 - \pi_i)^{m_i - y_i}$$

where

$$\pi_{i} = \frac{\exp(\beta_{0} + \beta_{1}X_{i1} + \ldots + \beta_{p}X_{ip})}{1 + \exp(\beta_{0} + \beta_{1}X_{i1} + \ldots + \beta_{p}X_{ip})} = \frac{2}{1 + \exp(\beta_{0} + \beta_{1}X_{i1} + \ldots + \beta_{p}X_{ip})}$$

Calculating the Deviance test statistic

Then the log-likelihood is:

$$\log \mathcal{L} = \sum_{i=1}^{n} [y_i \log(\pi_i) + (m_i - y_i) \log(1 - \pi_i) + \log\binom{m_i}{y_i}]$$

The deviance test statistic is based on a ratio of likelihoods.

$$\begin{aligned} Deviance &= -2\log\frac{\mathcal{L}_M}{\mathcal{L}_S} \\ &= -2(\log\mathcal{L}_M - \log\mathcal{L}_S) \\ &= 2(\log\mathcal{L}_S - \log\mathcal{L}_M) \end{aligned}$$

• Q: A Saturated Model has Deviance = 0 = 2 (mg hg - kg - kg

Calculating the Deviance test statistic

Deviance =
$$2(\log \mathcal{L}_{S} - \log \mathcal{L}_{M})$$

$$= 2\sum_{i=1}^{n} y_{i} \log \left(\frac{y_{i}}{m_{i}}\right) + (m_{i} - y_{i}) \log \left(\frac{m_{i} - y_{i}}{m_{i}}\right) + \log \binom{m_{i}}{y_{i}}$$

$$- y_{i} \log \left(\frac{\widehat{y}_{i}}{m_{i}}\right) - (m_{i} - y_{i}) \log \left(\frac{m_{i} - \widehat{y}_{i}}{m_{i}}\right) - \log \binom{m_{i}}{y_{i}}$$

$$= 2\sum_{i=1}^{n} \left(y_{i} \log(y_{i}) + (m_{i} - y_{i}) \log(m_{i} - y_{i})\right)$$

$$- y_{i} \log(\widehat{y}_{i}) - (m_{i} - y_{i}) \log(m_{i} - \widehat{y}_{i})$$

$$- y_{i} \log(\widehat{y}_{i}) - (m_{i} - y_{i}) \log(m_{i} - \widehat{y}_{i})$$

$$= 2\sum_{i=1}^{n} \left[y_{i} \log \left(\frac{y_{i}}{\widehat{y}_{i}}\right) + (m_{i} - y_{i}) \log \left(\frac{m_{i} - y_{i}}{m_{i} - \widehat{y}_{i}}\right)\right]$$

Case Study IV Exercise: Using Deviance

0.2

Using R output,

Q: Determine whether a saturated model is an improvement over the simpler model with linear function of log(Area). (In R, we get deviance of a model by using deviance('fittedmodel'))

Hypotheses:

Ho: Fitted Model: Ingit (7) ~ Log (Area) Pt Ho: Sachrated Deviance=12.062 In R: Residual deviance

► Test Statistic: Deviance=12.062

► Distribution of TS: \(\(\sigma \) \(\

► P-value: P(-1/2) > 12.062) =0.74

Ink: (-pdnisq(2-062))

Conclusion: The data are consistent with H_0 ; the simpler model with linear function of log(Area) is adequate (fits as well as the saturated model).

Binomial Logistic Regression: Interpreting Deviance

- Smaller deviance leads to larger p-value and vice versa.
- ► Large *p*-values means:
 - Fitted model is adequate, OR
 - ► Test is not powerful enough to detect inadequacies
- ► Small *p*-values means:
 - Fitted model is not adequate; consider a more complex model with more explanatory variables or higher order terms and so on, OR
 - Response distribution is not adequately modelled by the Binomial distribution, OR
 - There are severe outliers.

Can we do a Deviance GOF test in Binary case?

In Binary logistic regression case, $m_i = 1$ for all i, and $y_i = \begin{cases} 0 \\ 1 \end{cases}$ Then deviance becomes:

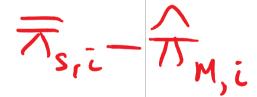
Deviance =
$$2\sum_{i=1}^{n} [y_i \log(y_i) + (1 - y_i) \log(1 - y_i) - y_i \log(\widehat{y}_i) - (1 - y_i) \log(1 - \widehat{y}_i)]$$

= $2\sum_{i=1}^{n} [-y_i \log(\widehat{y}_i) - (1 - y_i) \log(1 - \widehat{y}_i)].$

Notice that the terms that came from the saturated model, $\log \mathcal{L}_S$ are gone, so deviance is no longer useful to compare \mathcal{L}_M with \mathcal{L}_S .

Model assessment in Binomial Logistic Regression

- ▶ Is linear relationship appropriate?
 - Plot observed logit versus quantitative explanatory variable
- Is the form of the model correct?
- Use Wald or LRT tests
 Is saturated model better than fitted model?
 - Deviance GOF test
- Are there outliers?
 - Examine standardized residuals: Pearson and Deviance Residuals
- Consider other model fit statistics: AIC, BIC
- Other issues/concerns in model fitting



Residuals: Pearson and Deviance

duals: Pearson and Deviance

► Response (raw) residuals: (observed – fitted) proportion

$$(-1,1)$$

$$\widehat{\pi}_{S,i} - \widehat{\pi}_{M,i} = \frac{y_i}{m_i} - \widehat{\pi}_{M,i}$$

- Standardized residuals:
 - (1) Pearson Residuals: uses estimate of s.d. of Y (in denominator)

$$(-\infty,\infty)$$

$$P_{res,i} = \frac{y_i - m_i \widehat{\pi}_{M,i}}{\sqrt{m_i \widehat{\pi}_{M,i} (1 - \widehat{\pi}_{M,i})}}$$

Deviance Residuals: defined so that the sum of the squares of the residuals is the deviance $D_{res,i} = sign(y_i - m_i \widehat{\pi}_{M,i})$

$$(-\infty,\infty)$$

$$D_{res,i} = \operatorname{sign}(y_i - m_i \widehat{\pi}_{M,i})$$

$$e^2 = \sum_{i=1}^{n} D_i^2$$

$$2 - \sum_{i=1}^{n} \sum_{i=1}^{n} \sum_{i=1}^{n} \sum_{j=1}^{n} \sum_{j=1}^{n} \sum_{j=1}^{n} \sum_{i=1}^{n} \sum_{j=1}^{n} \sum_{j=1}^{n} \sum_{i=1}^{n} \sum_{j=1}^{n} \sum_{j=1}^{n}$$

Response, Pearson and Deviance Residuals in R

Response residuals Model object
residuals(fitbl, type="response")

► Pearson residuals

```
residuals(fitbl, type="pearson")
```

▶ Deviance residuals

```
residuals(fitbl, type="deviance")
```

Case Study IV Example: Were there outliers in the data?

| | Pearson, $P_{res,i}$ | Deviance, $D_{res,i}$ | |
|-----------------------------|--|-----------------------|--|
| Asymptotic Dist. | N(0,1) | N(0, 1) | |
| R code | pearson | deviance | |
| Possible outlier if | $ P_{res,i} > 2$ | $ D_{res,i} > 2$ | |
| Outlier if | $ P_{res,i} > 3$ | $ D_{res,i} > 3$ | |
| Under small <i>n</i> | D_{res} closer to $N(0,1)$ than P_{res} | | |
| • | | | |
| $\hat{\pi}$ close to 0 or 1 | P_{res} are unstable; related to instability of Wald | | |

E

Results: Both are < |2|, so no outliers

Case IV

Other Model Fit Statistics

- Useful for comparing models with same response and same data
- Two popular fit statistics: AIC and BIC; combines log-likelihood with a penalty
 - 1. Akaike's Information Criterion (AIC)

$$AIC = -2\log \mathcal{L} + 2(p+1)$$

2. Schwarz's (Bayesian Information) Criterion (BIC)

$$BIC = -2 \log \mathcal{L} + (p+1) \log N$$

where

- p-number of explanatory variables, and
- $N = \sum_{i=1}^n m_i$.
- Example: see AIC, BIC for Case IV model



- Extrapolation- don't make inferences/predictions outside range of observed data; model may no longer be appropriate.
- Multicollinearity- highly correlated explanatory variables; difficult to assess individual effects on response. Consequences include:
 - Unstable fitted equation
 - Coefficient that should be statistically significant is not
 - Coefficient may have the wrong sign
 - \blacktriangleright Sometimes, large s.e. of β
 - Sometimes numerical procedure to find MLEs does not converge

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 - Sometimes, large s.e. of $\widehat{\beta}$
 - Sometimes numerical procedure to find MLEs does not converge

- ▶ Influential points— an observation is influential if its removal substantially changes estimated coefficients (such as, fitted $\widehat{\beta}$'s, deviance)
- ► Model Building- choosing explanatory variables and their forms (eg. polynomial terms, interaction and transformations) tend to overfit the data; should build model on training data and test on test data (cross validation).

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Two problems specific to Logistic Regression

1. Extra-binomial variation



- variance of Y_i greater than $m_i\pi_i(1-\pi_i)$
- also called "over dispersion"
- does not bias $\widehat{\beta}$'s but s.e. of $\widehat{\beta}$'s will be too small (too small *p*-values, too narrow Cls)

SOLUTION: add one more parameter to the model, ψ dispersion parameter. Then $Var(Y_i) = \psi m_i \pi_i (1 - \pi_i)$.

Two problems specific to logistic regression

2. Complete and Quasi-complete separation

- Complete separation:
 - one or a linear combination of explanatory variables perfectly predict whether Y=1 or Y=0
 - In Binary response, when $y_i = 1$, $\hat{y}_i = 1$, then $\sum_{i=1}^{n} \{y_i \log(\hat{y}_i) + (1 y_i) \log(1 \hat{y}_i)\} = 0.$
 - MLE's cannot be computed
- Quasi-complete separation:
 - explanatory variables predict Y = 1 or Y = 0 almost perfectly (just a few points wrong)
 - MLE's are numerically unstable

SOLUTION: simplify the model. Other options- penalized maximum likelihood, exact logistic regression, bayesian methods