

Goodness-of-Fit Tests for Binary GLMs in the Case of No/Few Replicates

STAT 851 Final Project

Nikola Surjanovic, Peter Tea

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Introduction / Motivation

Why Do We Need GOF Tests?

We should check if GLM assumptions are correct

- Goodness-of-fit (GOF) tests

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However... with enough data, we will always reject the null hypothesis!

- What's the point then?

Why Do We Need GOF Tests?

Let's make sure our GOF tests have “stable” performance properties

- Probably don't want to blindly use these tests
- Still, make sure that the theory behind them makes sense!
- Use as an aid

$$Y_i | X_i = x_i \sim \text{Bin}(n_i, \pi_i), i = 1, \dots, J$$

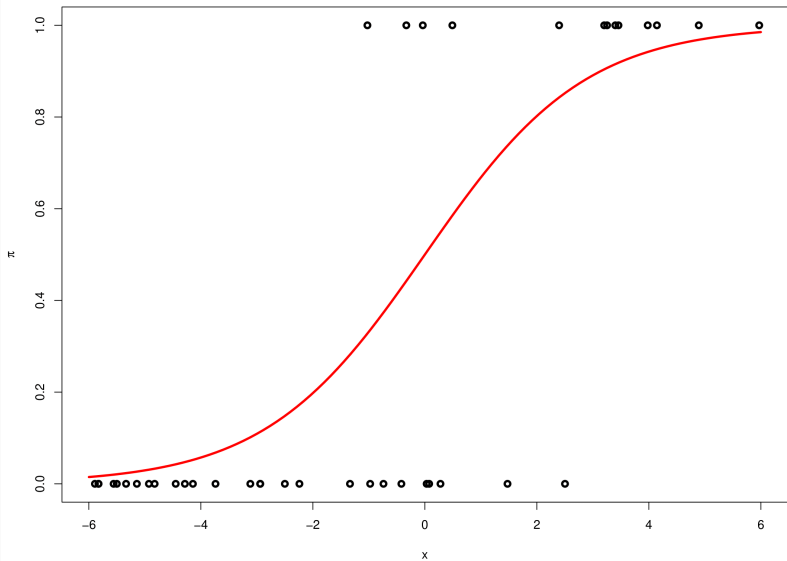
x_i 's are p -dimensional

$$q(\pi_i) = \beta^\top x_i,$$

for some p -dimensional vector β

$q(\cdot)$ is specified in advance

- E.g. could be $\text{logit}(\cdot)$



Consider $Y_i \sim \text{Bin}(n_i, \pi_i)$, $i = 1, \dots, J$

The Pearson chi-squared statistic:

$$\chi^2 = \sum_{i=1}^J \frac{(y_i - n_i \hat{\pi}_i)^2}{n_i \hat{\pi}_i (1 - \hat{\pi}_i)}$$

If J is *fixed*, and each $n_i \rightarrow \infty$,

$$\chi^2 \xrightarrow{d} \chi^2_{J-p},$$

where p is the number of estimated parameters.

Consider $Y_i \sim \text{Bin}(1, \pi_i)$, $i = 1, \dots, n$

The Pearson chi-squared statistic:

$$\chi^2 = \sum_{i=1}^n \frac{(Y_i - \hat{\pi}_i)^2}{\hat{\pi}_i(1 - \hat{\pi}_i)}$$

As $n \rightarrow \infty$,

$$X^2 \xrightarrow{d} \chi^2_{n-p} \quad (?)$$

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How can we converge to something that is “increasing”?

$$X^2 \xrightarrow{d} \chi_{n-p}^2 \quad (?)$$

Idea: Maybe

$$\frac{X^2 - \mu_n}{\sigma_n} \xrightarrow{d} N(0, 1) \quad (?),$$

for some μ_n, σ_n^2 by a generalization of CLT

What's Special About Few/No Replicates?

Few or no replicates?

This can happen, for example, when:

- A continuous explanatory variable is present
- A discrete variable that can take on infinitely many values

Test Statistics

1. Osious-Rojek test
2. Hosmer-Lemeshow test
3. C_n test statistic
4. Stukel test
5. Other tests

Consider $Y_i \sim \text{Bin}(1, \pi_i)$, $i = 1, \dots, n$, again

The Pearson chi-squared statistic:

$$\chi^2 = \sum_{i=1}^n \frac{(Y_i - \hat{\pi}_i)^2}{\hat{\pi}_i(1 - \hat{\pi}_i)}$$

In general (no replicates, few replicates, many replicates, etc.),

$$\frac{X^2 - \hat{\mu}}{\hat{\sigma}} \xrightarrow{d} N(0, 1),$$

for some $\hat{\mu}, \hat{\sigma}^2$ (see Osius and Rojek, 1992)

What are μ, σ^2 for Pearson X^2 ?

$$\hat{\mu} = \mu = J \quad (\text{number of covariate patterns})$$

$\hat{\sigma}^2$ can be obtained through the residual sum of squares from a weighted least squares regression on a transformed set of data

Question: Is this a one-sided or a two-sided test?

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If $J = n$, should treat as a two-sided test.

- Can be viewed as a type of score test

Results derived for a generalized distance between Y_i 's and $n_i\hat{\pi}_i$'s

Essentially formed infinitely many test statistics indexed by a parameter, λ

- $\lambda = 1$ for Pearson
- $\lambda = 0$ for deviance
- See Read and Cressie (1988)

Not clear how to generalize to other GLMs

- Should be a multinomial response (i.e. binomial is OK)

Idea: Group by fitted values, rather than by covariate pattern

- 1980: David Hosmer and Stanley Lemeshow
- Partition units into g subgroups based on $\hat{\pi}_i$'s

Hosmer-Lemeshow Test

Test Statistic Presentation

	1	2	...	g	
y = 0	1	0		3	
y = 1	0	2		5	

- J = # of distinct covariate patterns
- n_i = # of replicates, with the j th covariate pattern ($i = 1, 2, \dots, J$)
- y_i = # of successes among the n_i units

Hosmer-Lemeshow Test

$$\hat{C} = \sum_{k=1}^g \left[\frac{(o_{1k} - \hat{e}_{1k})^2}{\hat{e}_{1k}} + \frac{(o_{0k} - \hat{e}_{0k})^2}{\hat{e}_{0k}} \right] \quad (1)$$

$$o_{1k} = \sum_{j=1}^{c_k} y_j, \quad o_{0k} = \sum_{l_k} (m_j - y_j)$$
$$\hat{e}_{1k} = \sum_{l_k} m_j \hat{\pi}_j, \quad \hat{e}_{0k} = \sum_{l_k} m_j (1 - \hat{\pi}_j)$$

l_k is the collection of indices of observations that fall in the k th group

Distribution of \hat{C}

- Simulation study by Hosmer and Lemeshow (1980) found that when $J = n$,

$$\hat{C} \sim \chi^2_{g-2}$$

Approximately true when $J \approx n$

Strategies to split units into meaningful subgroups

- I. Percentiles of estimated probabilities (a.k.a. “deciles-of-risk” if $g = 10$)
- II. Fixed endpoints for the estimated probabilities

Method I is commonly used

Choice of g impacts results

“Why I Don’t Trust the Hosmer-Lemeshow Test for Logistic Regression” - Paul Allison (UPenn)

<https://statisticalhorizons.com/hosmer-lemeshow>

Based on sum of squared Pearson residuals

$$\sum_{i=1}^n \frac{(Y_i - \hat{\pi}_i)^2}{\hat{\pi}_i (1 - \hat{\pi}_i)}$$

Can be standardized to be distributed $N(0, 1)$ asymptotically

Unstable variance with extreme values of p_i (Chen et al., 2018)

Stabilized variance (see Chen et al., 2018)

$$C_n = \sum_{i=1}^n \frac{Y_i - \hat{\pi}_i}{\sqrt{\hat{\pi}_i (1 - \hat{\pi}_i)}}$$

Under some regularity assumptions,

$$\tilde{C}_n = \frac{n^{-\frac{1}{2}} C_n}{\sigma_n} \xrightarrow{d} N(0, 1)$$

Conduct 2-sided GOF test

No replicates?

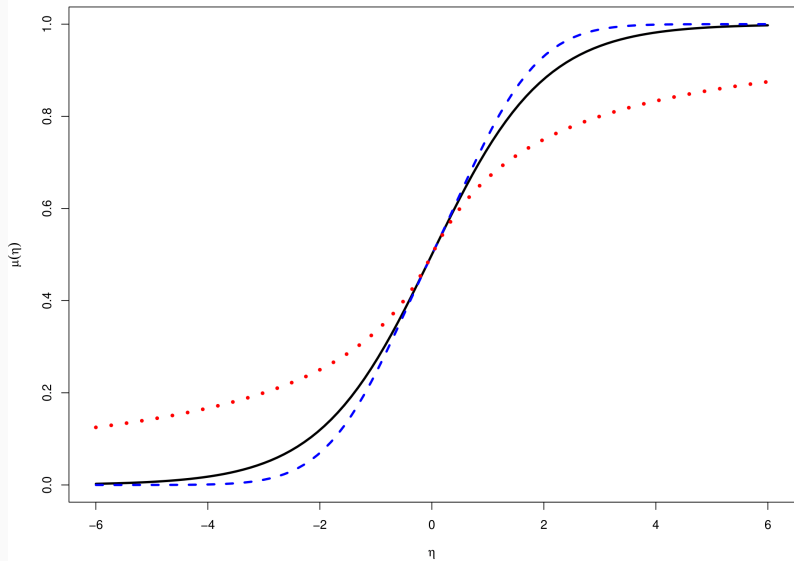
One regularity assumption:

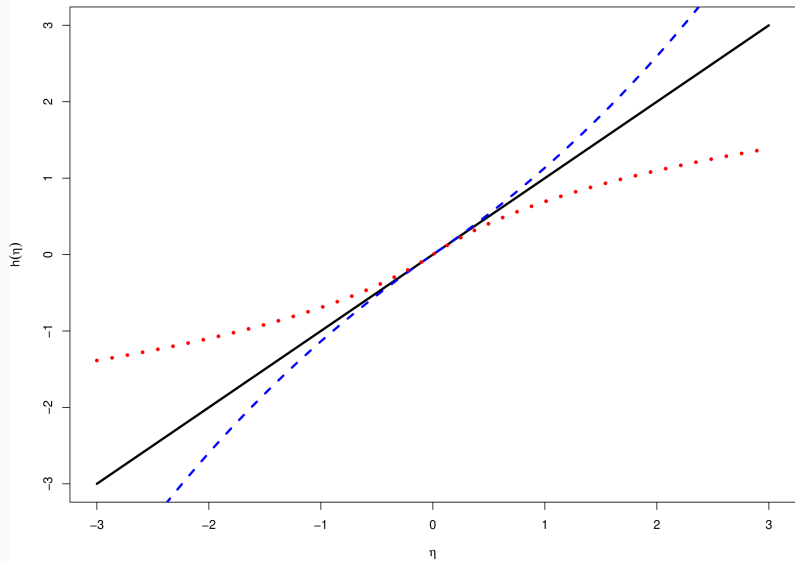
- There exists a finite number M such that $\|x_i\| \leq M, \forall i \in \mathbb{N}$ and $\lim_{n \rightarrow \infty} \sum_i x_i x_i^T$ is a finite non-singular matrix (x_i represents the i th row of the model matrix as a column vector)

Stukel test: Essentially a test of the link function

- A missing quadratic term might be (approximately) captured by a modified link function

Create a family of link functions





A family of link functions, parameterized by $\alpha = (\alpha_1, \alpha_2)^\top$

$$\pi_\alpha(\eta) = \frac{\exp(h_\alpha(\eta))}{1 + \exp(h_\alpha(\eta))}$$

For fixed α , we still have a GLM, with link $h_\alpha^{-1}(\text{logit}(\cdot))$

- We estimate α , though

When $\alpha = (0, 0)$ we have the regular logit link

$\alpha \approx (.165, .165)$ yields probit

$\alpha \approx (.62, -.037)$ yields complementary log-log

$$H_0 : \alpha = 0 \text{ vs. } H_1 : \alpha \neq 0$$

Score test essentially tests logit link

- Could probably test $\alpha = (.165, .165)$ (probit), etc.

There are many other tests

1. For example, see Chen et al. (2018) and Hosmer et al. (1997)

Information matrix statistic: A good test to keep in mind

- Very general
- Based on comparing two estimates of the Fisher information

Under some regularity conditions,

$$\mathbb{E} \left[\left(\frac{\partial}{\partial \theta} \log(f(X; \theta)) \right) \left(\frac{\partial}{\partial \theta} \log(f(X; \theta)) \right)^{\top} \right] = - \mathbb{E} \left[\left(\frac{\partial^2}{\partial \theta \partial \theta} \log(f(X; \theta)) \right) \right]$$

Estimates differ **if model is misspecified**

Roughly speaking, check if

$$|\text{Estimate 1} - \text{Estimate 2}| > 0$$

See White (1982)

Application



Novak Djokovic holding his 8th Australian Open Trophy (17th Grand Slam)

Context: Observe tennis serve speeds (continuous), and outcome of service point (binary)

- 70% of all points are won in 4 shots or less
- All points begin with a serve
- Fit binary logistic regression, $\text{Win Point} \sim \text{Serve Speed} + \text{Opponent (7 matches total)}$

Discussion

Pros

1. Osious-Rojek: Easy to compute, not restricted to logit link
2. Hosmer-Lemeshow: Simple and intuitive
3. C_n statistic: Also simple
4. Stukel: Tests link function directly

Cons

1. Osious-Rojek: Doesn't generalize to non-binary responses
2. Hosmer-Lemeshow: Depends on g
3. C_n statistic: Only works without replicates
4. Stukel: Generalized link function not exact for probit, cloglog, etc.

Summary of performances from Chen et al. (2018)

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Thank You!

Questions?

Bonus Material

For $\eta \geq 0$,

$$h_\alpha = \begin{cases} 1/\alpha_1(\exp(\alpha_1|\eta|) - 1) & \alpha_1 > 0 \\ \eta & \alpha_1 = 0 \\ -\alpha_1 \log(1 - \alpha_1|\eta|) & \alpha_1 < 0 \end{cases}$$

For $\eta < 0$,

$$h_\alpha = \begin{cases} -1/\alpha_2(\exp(\alpha_2|\eta|) - 1) & \alpha_2 > 0 \\ \eta & \alpha_2 = 0 \\ \alpha_2 \log(1 - \alpha_2|\eta|) & \alpha_2 < 0 \end{cases}$$

Generalized Distance Between Y_i and $n_i \hat{\pi}_i$

$$\begin{aligned} a_1(Y_i, \hat{\pi}_i) &= \frac{(Y_i - n_i \hat{\pi}_i)^2}{n_i \hat{\pi}_i} \\ &= \frac{2Y_i}{1(1+1)} \cdot \left[\left(\frac{Y_i}{n_i \hat{\pi}_i} \right)^1 - 1 \right] - \frac{2}{1+1} (Y_i - n_i \hat{\pi}_i) \end{aligned}$$

$$a_\lambda(Y_i, \hat{\pi}_i) = \frac{2Y_i}{\lambda(\lambda+1)} \cdot \left[\left(\frac{Y_i}{n_i\hat{\pi}_i} \right)^\lambda - 1 \right] - \frac{2}{\lambda+1}(Y_i - n_i\hat{\pi}_i)$$

Test is derived by

1. Taylor series expansion to replace $\hat{\pi}$'s with π_0 's
2. Lyapunov CLT (for independent but not identically distributed Y_i 's)

Hosmer-Lemeshow Test

Alternate form of Hosmer-Lemeshow test statistic:

$$\hat{C} = \sum_{k=1}^g \frac{(o_{1k} - n'_k \bar{\pi}_k)^2}{n'_k \bar{\pi}_k (1 - \bar{\pi}_k)} \quad (2)$$

where

$$\bar{\pi}_k = \frac{1}{n'_k} \sum_{j=1}^{c_k} m_j \hat{\pi}_j$$

and n'_k is the number of units in the k th subgroup