

ETC3580: Advanced Statistical Modelling

Week 4: Binomial and proportion responses

Outline

1 Binomial responses

2 Proportion responses

Binomial responses

Binomial distribution

Y is binomially distributed B(m, p) if

$$P(Y = y) = \binom{m}{y} p^{y} (1 - p)^{m-y}$$

Y = number of "successes" in m independent trials, each with probability p of success.

Likelihood

$$L = \prod_{i=1}^{n} {m_i \choose y_i} p_i^{y_i} (1 - p_i)^{m_i - y_i}$$

Binomial regression also uses a logit link

$$p_i = e^{\eta_i}/(1 + e^{\eta_i})$$

$$\eta_i = \beta_0 + \beta_1 x_{i,1} + \dots + \beta_q x_{i,q}$$

Binomial likelihood

Likelihood

$$L = \prod_{i=1}^{n} {m_i \choose y_i} p_i^{y_i} (1 - p_i)^{m_i - y_i}$$

$$\log L(\beta) = \sum_{i=1}^{n} \left[\log \binom{m_i}{y_i} + y_i \log(p_i) + (m_i - y_i) \log(1 - p_i) \right]$$

$$= \sum_{i=1}^{n} \left[\log \binom{m_i}{y_i} + y_i \eta_i - y_i \log(1 + e^{\eta_i}) - (m_i - y_i) \log(1 + e^{\eta_i}) \right]$$

$$= \sum_{i=1}^{n} \left[\log \binom{m_i}{y_i} + y_i \eta_i - m_i \log(1 + e^{\eta_i}) \right]$$

Binomial responses

In R:

glm needs a two-column matrix of success and failures. (So rows sum to m).

```
fit <- glm(cbind(successes, failures) ~
    x1 + x2,
    family=binomial, data=df)</pre>
```

Everything else works the same as for binary regression.

- If mean correctly modelled, but observed variance larger than model, we called the data "overdispersed". [Same for underdispersion.]
- Concept of overdispersion irrelevant for OLS and logistic regression because there cannot be any more variance than what is modelled.
- For binomial regression: $y_i \sim B(m_i, p_i)$, $E(y_i) = m_i p_i$, $Var(y_i) = m_i p_i (1 - p_i)$.
- If model correct, $D = -2 \log L \sim \chi_{n-q}^2$. So D > n - q indicates overdisperson.

- D > n q can also be the result of:
 - missing covariates or interaction terms
 - negligence of non-linear effects
 - large outliers
 - sampling from clusters
 - non-independence
 - \blacksquare m small (χ^2 approximation fails)

Solution 1: Drop strict binomial assumption and let $E(y_i) = m_i p_i$, $Var(y_i) = \phi m_i p_i (1 - p_i)$.

Pearson residuals

$$r_i = \frac{y_i - m_i \hat{p}_i}{\sqrt{m_i \hat{p}_i (1 - \hat{p}_i)}}$$

Simple estimate of dispersion parameter

Estimate

$$\hat{\phi} = \frac{1}{n} \sum_{i=1}^{n} r_i^2.$$

- OK for estimation and standard errors on coefficients.
- But no proper inference via deviance.

Solution 2: Define a "quasi-likelihood" that behaves like the log-likelihood but allows for $Var(y_i) = \phi m_i p_i (1 - p_i)$.

```
fit <- glm(cbind(successes, failures) ~
    x1 + x2,
    family=quasibinomial, data=df)</pre>
```

■ Must use F-tests rather than χ^2 tests for comparing models. (Approximation only.)

Inference for GLMs

Model	F-test	χ^2 test
Normal OLS	Exact	-
Binary Logistic	Approx	Better approx
Binary Probit	Approx	Better approx
Binomial Logistic	Approx	Better approx
Quasibinomial logistic	Approx	_

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Proportion responses: three approaches

Suppose $y \in [0, 1]$.

logitNormal model

$$\log(y/(1-y)) \mid \mathbf{x} \sim \mathsf{N}(\mathbf{x}'\boldsymbol{eta}, \sigma^2)$$

quasiBinomial model

$$y \mid \mathbf{x}$$
 has mean p and variance $\phi p(1-p)$
where $\log(p/(1-p)) = \mathbf{x}'\beta$

Beta model

$$y \mid \mathbf{x} \sim \text{Beta}(a, b)$$

where $E(y \mid \mathbf{x}) = a/(a+b) = e^{\mathbf{x}'\beta}/(1+e^{\mathbf{x}'\beta})$

logitNormal model

logitNormal model

$$\log(y/(1-y)) \mid \mathbf{x} \sim N(\mathbf{x}'\boldsymbol{\beta}, \sigma^2)$$

- Provided no empirical proportions are at either 0 or 1, we can compute a logit transformation of the observed proportions. log(y/(1-y))
- Then just fit a Gaussian linear regression using OLS.
- Back-transform the predictions using the inverse logit. $e^y/(1+e^y)$

```
lm(log(y/(1-y)) \sim x1 + x2, data=df)
```

Quasi-binomial model

quasiBinomial model

```
y \mid \mathbf{x} has mean p and variance \phi p(1-p)
where \log(p/(1-p)) = \mathbf{x}'\beta
```

- logit link keeps predicted proportions in (0, 1)
- Variance function $\phi p(1-p)$ makes sense for proportions as greatest variation around p=0.5 and least around p=0 and p=1.

```
glm(y ~ x1 + x2,
    family=quasibinomial, data=df)
```

Beta model

$$y \mid \mathbf{x} \sim \text{Beta}(a, b)$$

where $E(y \mid \mathbf{x}) = a/(a+b) = e^{\mathbf{x}'\beta}/(1+e^{\mathbf{x}'\beta})$

Beta density

$$f(y) = \frac{\Gamma(a+b)}{\Gamma(a)\Gamma(b)} y^{a-1} (1-y)^{b-1}$$
 where $y \in [0, 1]$ and $\Gamma(u) = \int_0^\infty x^{u-1} e^{-x} dx$.

$$\blacksquare E(y) = \frac{a}{a+b} \qquad Var(y) = \frac{ab}{(a+b)^2(a+b+1)}$$

Beta model

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E(y) =
$$\frac{a}{a+b}$$
 Var(y) = $\frac{ab}{(a+b)^2(a+b+1)}$

Reparameterize so $\mu = a/(a+b)$ and $\phi = a+b$.

Beta model

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- Reparameterize so $\mu = a/(a+b)$ and $\phi = a+b$.
- Then E(Y) = μ and Var(Y) = μ (1 μ)/(1 + ϕ).

Beta model

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■
$$E(y) = \frac{a}{a+b}$$
 $Var(y) = \frac{ab}{(a+b)^2(a+b+1)}$

- Reparameterize so $\mu = a/(a+b)$ and $\phi = a+b$.
- Then E(Y) = μ and Var(Y) = μ (1 μ)/(1 + ϕ).

```
mgcv::gam(y ~ x1 + x2, family=betar())
```