

4.5. Inference for GLMs

the likelihood equation

$$\sum_{i=1}^N \frac{(y_i - \mu_i) x_{ij}}{\text{var}(Y_i)} \frac{\partial \mu_i}{\partial \eta_i} = 0$$

y
 $\{E(Y)\}$
 μ
 $S \propto \mu$
 η
 $S \propto \beta$
 β

(β is there implicitly via $\mu_i = g^{-1}(\sum_j \beta_j x_{ij})$)

For most GLMs, it's nonlinear in β

Significance testing & interval estimation via
 (Score, Wald, L-R) methods apply to any GLM.

4.5.1 Deviance & Goodness of Fit

Let $\tilde{\theta}$ be the estimate of θ for the saturated model
 (corresponds to $\tilde{\mu}_i = y_i \forall i$)
 $\hat{\theta}, \hat{\mu}$ are ML estimates for a particular model.
 unsaturated

lack of fit = $-2 [L(\hat{\mu}; y) - L(y; y)]$

If $a(\phi) = \frac{\phi}{w_i}$

$$= 2 \sum [y_i \tilde{\theta}_i - b(\tilde{\theta}_i)] / a(\phi) - 2 \sum [y_i \hat{\theta}_i - b(\hat{\theta}_i)] / a(\phi)$$

$$= 2 \sum w_i [y_i (\tilde{\theta}_i - \hat{\theta}_i) - b(\tilde{\theta}_i) + b(\hat{\theta}_i)] / \phi$$

deviance $\rightarrow \frac{D(y; \hat{\mu})}{\phi}$: scaled deviance

4.5.2 Deviance for Poisson model

$\hat{\theta}_i = \log \hat{\mu}_i$ $b(\hat{\theta}_i) = \exp(\hat{\theta}_i) = \hat{\mu}_i$

$\tilde{\theta}_i = \log y_i$ $b(\tilde{\theta}_i) = y_i$ $a(\phi) = 1$

$\Rightarrow D(y; \hat{\mu}) = 2 \sum y_i \log \left(\frac{y_i}{\hat{\mu}_i} \right)$ ← same as G^2 for testing independence

4.5.3 Binomial models

binomial GLMs w/ sample proportions $\{y_i\}$
 based on $\{n_i\}$ ← note

(X $y_i \sim \text{bin}(n_i, \pi_i)$
 (O $y_i \times n_i \sim \text{bin}(n_i, \pi_i)$
 why?

$\hat{\theta}_i = \log \frac{\hat{\pi}_i}{1 - \hat{\pi}_i}$, $\tilde{\theta}_i = \log \frac{y_i}{1 - y_i}$

$b(\hat{\theta}_i) = \log [1 + \exp(\hat{\theta}_i)] = -\log(1 - \hat{\pi}_i)$

$$v_i = \frac{y_i}{1 - \hat{\pi}_i}, \quad \sigma_i = \frac{y_i}{1 - y_i} \quad \text{why?}$$

$$b(\hat{\theta}_i) = \log[1 + \exp(\hat{\theta}_i)] = -\log(1 - \hat{\pi}_i)$$

$$b(\tilde{\theta}_i) = -\log(1 - y_i)$$

$$a(\phi) = 1/n_i \quad (\text{thus } \phi = 1 \text{ \& } w_i = n_i)$$

$$D(\hat{y}, \hat{M}) = 2 \sum \text{observed} \log\left(\frac{\text{observed}}{\text{fitted}}\right)$$

* We can construct data files either as
 grouped data | ungrouped data
 counts of success at each X setting | individual 0-1 obs. for each subject

* χ^2 approximation holds only for grouped data. (why?)

4.5.4 L-R model comparisons using the deviance

Consider two models $(M_0 \text{ w/ } \hat{\mu}_0, M_1 \text{ w/ } \hat{\mu}_1)$
 that are nested, i.e. $M_0 \subset M_1$

$\Rightarrow L(\hat{\mu}_0; \underline{y}) \leq L(\hat{\mu}_1; \underline{y})$ why?
 $\Rightarrow D(\underline{y}; \hat{\mu}_1) \leq D(\underline{y}; \hat{\mu}_0)$

\uparrow
 simpler
 fewer parameters
 \downarrow
 smaller maximum
 \downarrow
 larger deviance

Assuming M_1 holds, testing $H_0: M_0$ uses

$$\begin{aligned} (*) &= D(\underline{y}; \hat{\mu}_0) - D(\underline{y}; \hat{\mu}_1) \\ &= -2 [L(\hat{\mu}_0; \underline{y}) - L(\hat{\mu}_1; \underline{y})] \end{aligned}$$

which is a LR test statistic.

Under regularity conditions,

$$(*) \sim \chi^2 (\# \text{ param of } M_1 - \# \text{ param of } M_0)$$

$(*)$ is large if M_0 fits poorly compared to M_1 .

4.5.5 Residuals for GLMs

If a GLM fits poorly

\Rightarrow examine where the fit is poor
 by looking at residuals

Note $D(y_i; \hat{\mu}_i) = \sum d_i$ where

$$d_i = 2w_i [y_i(\hat{\theta}_i - \tilde{\theta}_i) - b(\hat{\theta}_i) + b(\tilde{\theta}_i)]$$

the deviance residual

$$\sum d_{ij} = G^2$$

$$\sqrt{d_i} \operatorname{sgn}(y_i - \hat{\mu}_i)$$

The Pearson residual

$$e_i = \frac{y_i - \hat{\mu}_i}{[\widehat{\operatorname{var}}(Y_i)]^{1/2}}$$

e.g. $\operatorname{var}(Y_i) = \mu_i$
for Poisson

$\sum e_{ij}^2 = \chi^2$ in
Pearson χ^2 tests of
independence

Standardized Poisson residuals

$$r_i = \frac{e_i}{\sqrt{1 - \hat{h}_i}} \leftarrow \text{leverage of obs } i.$$