Generalized Linear Models (GLMs/GLIMs)

STAT 757

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Model Framework

The GLM is described by three components:

- 1. The random component specifies the conditional distribution of $y_i | \vec{x_i}$ and is typically a member of the exponential family of distributions (Normal, binomial, Poisson, Negative-binomial, etc.) but other distributions are possible.
- 2. We call our linear sum of predictors the *linear predictor*, and denote it as $\eta_i = \beta_0 + \sum_{i=1}^{p} \beta_j x_{ij}$
- We call the transformation that links the expected response values μ_i = E(y_i|x̄) and the linear predictor η_i the link function: g(μ_i) = η_i. This link function is assumed to be smooth (differentiable) and invertible. It's inverse g⁻¹ is often called the *mean function* since μ_i = g⁻¹(η_i).

GLM vs MLR

Recall the MLR model with untransformed response values can be written as

$$y_i = \beta_0 + \sum_{j=1}^p \beta_j \, x_{ij} + \epsilon_i.$$

In that case, we model $E(g(y_i))$ as a linear sum of x_i values, and further assume Normal errors with constant variance.

Consider, for now, the simple untransformed case (i.e., g is the identity function).

One could pose the MLR model as a GLM (not to be confused with a General Linear Model) as follows:

- 1. The random component is Normally distributed.
- 2. The *linear predictor* is $\eta_i = \beta_0 + \sum_{j=1}^{p} \beta_j x_{ij}$ (nothing new here!)

3. The *link function* is the identity function: $g(E(y_i)) = E(y_i) = \eta_i$

GLM vs MLR

Note that transforming Y values under MLR is different than specifying a non-identity link function!

GLMs model $g(E(y_i))$, the transformed expecation of the response, using the linear predictor. This gives more flexibility to apply linearizing transformations without affecting the distribution about that trend. For example, compare the two models by comparing y_i values and inverse-transforms:

MLR:
$$y_i = g^{-1}(\eta_i + \epsilon_i)$$

GLM: $y_i = g^{-1}(\eta_i) + \epsilon_i$

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This distinction often makes GLMs preferrable over MLR.

Parameter estimation is done via Maximum Liklihood, and most of the diagnostics for multiple linear regression carry over to GLMs.

For more information, please see Ch. 15 of *Applied Regression Analysis & Generalized Linear Models* by John Fox. http://www.sagepub.com/sites/default/files/ upm-binaries/21121_Chapter_15.pdf

Example: Logistic Regression (Sheather, Ch. 8)

A common form of response data are counts of a particular type of outcome among m trials. For example, the number of individuals in a sample with a specific genotype. In such cases, the data are best modeled using a binomial distribution, not a Normal distribution, using *logistic regression*.

 $E(Y|x) \sim binom(m, \theta)$

Example: Logistic Regression (Sheather, Ch. 8)

Here the parameter of interest is p (or θ) – the probability of a *success* on each of our n (or m) trials. Since m is known and not a parameter that needs to be estimated, the goal is to estimate θ as a function of our linear predictor. In logistic regression, this is done by assuming

$$E(Y|X) = m\theta = m rac{\exp(\eta_i)}{1 + \exp(\eta_i)} = rac{m}{1 + \exp(-\eta_i)}$$

Thus, a little algebra gives that

$$\eta_i = \log\left(\frac{\theta(x)}{1-\theta(x)}\right)$$

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We call the right side of that equation the *logit* function, and $\theta/(1-\theta)$ the *odds*.

Example: Logistic Regression (Sheather, Ch. 8)

To see how this can be cast as a GLM, note that:

- 1. The distribution is binomial.
- 2. The relationship between the mean (let's use $E(y_i/m = \theta_i)$) and linear predictor η_i is given by the logit function

$$\eta_i = g(heta) = \log\left(rac{ heta(x)}{1- heta(x)}
ight)$$

Exercise: Work through the examples from Sheather, Ch. 8. (Are there alternative link functions?)