

Generalized linear models

29 Nov 2023

```
## it's nice to include packages at the top
## (and NOT automatically install them)
## try not to carry over packages you don't use
library(ggplot2); theme_set(theme_bw())
## diagnostics
library(performance)
library(DHARMA)
## downstream model evaluation
library(broom)
library(dotwhisker)
library(emmeans)
library(effects)
library(marginaleffects)
library(parameters)
## library(ggeffects)
```

Basics

- assume $\mathbf{y}_i \sim \text{Dist}(g^{-1}((\mathbf{X}\beta)_i))$
- g = **link function**
- $\eta = \mathbf{X}\beta$ = **linear predictor**
- **link scale, data or response scale**
- GLMs inverse-transform η , they don't transform y
- allows:
 - separate control of heteroscedasticity and nonlinearity
 - almost as convenient/efficient as LMs
 - equivalent to MLE in many cases
- in practice almost all GLMs are logistic (binary data) or Poisson

- lots of inference, diagnostics, etc. inherited from LM framework

Exponential family

- $f(x|\theta) = h(x)g(\theta) \exp(\eta(\theta)T(x))$
- e.g. Poisson: $f(x|\theta) = \theta^x \exp(-\theta)/x! = (1/x!) \exp(-\theta) \exp(x \log(\theta))$
- $h(x) = 1/x!; g(\theta) = \exp(-\theta); \eta(\theta) = \log(\theta); T(x)$
- η is the **canonical link** function for the family (nice mathematical properties)
- binomial, Poisson, Gamma (inverse Gaussian, von Mises distribution ...)

Mean-variance relations

- can show that all we need for computation is the link function and the **variance function** $V = f(\mu)$ (may also depend multiplicatively on a **scale** or **dispersion parameter**, e.g. $V = \mu$ for Poisson, $V = \sigma^2$)

Link functions

- canonical doesn't always work best (e.g. Gamma/inverse link)
- probit vs logit; not much difference
- cloglog; *log-hazard* scale
- inverse link: linear changes in the *rate* of events

Log-hazards and log-hazard offsets

- if hazard is h , probability is $1 - \exp(-h)$
- $C(\mu) = \log(-\log(1 - \mu))$
- $C^{-1}(\eta) = 1 - \exp(-\exp(\eta))$
- $C^{-1}(\eta + \log(\Delta t)) = 1 - \exp(-\exp(\eta) \cdot \Delta t)$
- $\rightarrow 1 - (1 - \mu_0)^{\Delta t}$

Computation

- iteratively reweighted least squares
- needs starting values, but almost always robust to them

in R

- “family” functions contain all of the components needed for GLM fitting, prediction, etc.
- some of the components are weird (e.g. `$aic`)
- canonical link is used by default

```
names(binomial)
```

NULL

Offsets

- allow for differential search effort, ratios, etc.
- typically add $\log(e)$
- e.g. $y \sim \text{Poisson}(\mathbf{X}\beta + \log(A))$ is equivalent to modeling the response y/A , but without messing up the mean-variance relationship

Offset/link tricks

- fit an exponential curve with constant variance: `family = gaussian(link = "log")`
- Ricker function $y = ax \exp(-bx)$: `log-link, y ~ x + offset(log(x))`
- Michaelis-Menten $y = ax/(b + x) \rightarrow 1/y = (b/a) \cdot (1/x) + 1/a$: `inverse-link, y ~ I(1/x)`

Model interpretation, visualization, testing

Parameter interpretation

- log scale: easy
- logit scale: $\approx \log$ for low baseline, $\approx \log(1 - x)$ for high baseline, slope $\beta/4$ for intermediate values
- cloglog: **log-hazard** scale

Inference

- Wald tests (no finite-size corrections!)
- approximate Wald CIs (compute then back-transform)
- profile CIs

Overdispersion

- too much variance
- SSQ of Pearson residuals $\sim \chi^2(n - p)$
 - quasi-likelihood (also handles **underdispersion**)
 - compounded models (negative binomial, beta-binomial)
 - observation-level random effects

Extended distributions

- VGAM, glmmTMB packages

Complete separation

- there is some linear combination of predictors that separates 0 from 1 responses (or 0 from non-zero responses in the case of count models)
- infinite MLE
- Hauck-Donner effect screws up Wald tests
- likelihood ratio tests still OK (sort of)
- Firth logistic regression (brglm2 package), Bayesian priors (arm::bayesglm)

Zero-inflation/hurdle models

- finite mixture models

Most common GLM problems

- binomial/Poisson models with non-integer data
- failing to specify family (default Gaussian: \rightarrow linear model); using glm() for linear models (unnecessary)
- predictions on effect scale
- using (k, N) rather than $(k, N - k)$ with family=binomial
- back-transforming SEs rather than CIs
- neglecting overdispersion
- Poisson for *underdispersed* responses
- equating negative binomial with binomial rather than Poisson
- worrying about overdispersion unnecessarily (binary/Gamma)
- ignoring random effects

Overdispersion in Bernoulli models?

I think many analysts read that binary models cannot be overdispersed and just do not question it. This happened with the deviance dispersion being the appropriate statistic to measure count model extra-dispersion. Some analysts simply took this on faith, so to speak. But they were mistaken.

[Joseph Hilbe \(2013\)](#)

References