# Generalized linear models

Katie Pollard

BMI 206 docpollard.org/bmi206 October 5, 2016

# Generalized linear model (GLM)

If outcome is not quantitative, the linear model framework can be extended via data transformations, called link functions.

- Binary: logit (alternatives: probit, log-log)
- Counts: log (also known as log-linear model)

The covariates are still a linear combination.

- The parameters are estimated by numerical methods (e.g., Newton-Raphson).
- But the error has a different distribution.

## Link functions in GLMs

Link function systematically relates expected value of outcome (E[Y] =  $\mu$ ) to a linear combination of covariates (X):

 $g(\mu)=\beta'X$ 

- Identity link:  $g(\mu) = \mu$
- Log link:  $g(\mu) = \log(\mu)$
- Logit link:  $g(\mu) = \log(\mu/(1-\mu))$
- Log-log link:  $g(\mu) = \log(-\log(1-\mu))$
- Probit link:  $g(\mu) = Phi^{-1}(\mu)$

## Error distributions in GLMs

Different types of outcome variables require different error distributions, e.g.,

- Continuous (link=identity): Gaussian
- Binary (link=logit): Binomial
- Counts (link=log): Poisson

These are the random components.

They are examples of the exponential family:

 $f(y) = a(\mu)b(y)exp\{g(\mu)y\}$ 

#### Logistic regression parameters

Consider:

 $logit(\pi) = \beta_0 + \beta_1 X$ 

Interpretation of  $B_1$  is the expected change in logit for a unit increase in X. What is this?

If X is binary (e.g., 0=wild-type vs. 1=mutant):

 $odds(X=0) = exp\{\beta_0\}, odds(X=1) = exp\{\beta_0\}exp\{\beta_1\}$ 

Odds increase multiplicatively by  $exp\{B_1\}$  per unit X.

Odds ratio = odds(X=1)/odds(X=0) = exp{ $B_1$ }

## Logistic regression model

What is the distribution function?

Can we write it as an exponential family?

What is the canonical link?

What is the systematic component?

#### **Poisson regression parameters**

Consider:

$$\log(\mu) = \beta_0 + \beta_1 X$$

Interpretation of  $B_1$  is the expected change in log count for a unit increase in X.

Exponentiate to get back to count scale.

If X is binary (e.g., 0=wild-type vs. 1=mutant):

 $\mu(X=0) = \exp\{\beta_0\} \text{ and } \mu(X=1) = \exp\{\beta_0 + \beta_1\}$ 

Relative risk =  $\mu(X=1)/\mu(X=0) = \exp\{\beta_1\}$ 

## Poisson regression model

What is the distribution function?

Can we write it as an exponential family?

What is the canonical link?

What is the systematic component?

#### **Over-dispersion**

The Poisson distribution has the variance equal to the mean. Count data in bioinformatics frequently violates this assumption, e.g.,

- Gene expression via RNA-seq (read counts/transcript)
- Taxon abundance in metagenomics (reads counts/taxa)

Variance > mean is called "over-dispersion".

The negative binomial distribution is a good alternative:

mean = 
$$\mu$$
, variance =  $\mu + \mu^2/k$ 

#### Linear model as a GLM

What is the distribution function?

Can we write it as an exponential family?

What is the canonical link?

What is the systematic component?