### Generalized linear models:

General linear models +
Non-normal residuals=
Generalized linear models

Negative binomial regression, beta regression and more...

### Generalized linear models

#### Main ideas:

- 1) A transformation of the expectation of the response is expressed as a linear combination of covariates rather than the mean response directly.
- 2) For the random part of the model, distributions other than the Normal can be chosen, e.g., Poisson, Binomial or gamma.

### Generalized linear models

#### GLMs are made up of three components:

- 1) A *statistical distribution* used to describe the random variation in the response *y*; this is the stochastic part of the system description
- 2) A *linear predictor*, i.e., a linear combination of covariate effects that are thought to make up *E(y)*; this is the systematic or deterministic part of the system description.
- 3) A *link function* that is applied to the E(y), expectation of the response

## Objectives

- So far, we've modeled data with 3 distributions: normal, Poisson, binomial
- Discuss modeling data with other distributions, namely the negative binomial and beta distributions
  - Understand when to use each distribution

# GLMs .... So far

Distribution	Dependent data type	When to use it?
Normal	Continuous $(-\infty, \infty)$	In cases where errors around means are normal
Poisson	Discrete positive values $(0, \infty)$	Modeling counts under the assumption that mean=variance
Binomial	Discrete positive values with an upper bound (0, N)	Modeling the probability of C successes in N trials

# Introducing the Gamma distribution

### Gamma distribution

A continuous distribution with a positive support – exponential and  $\chi^2$  are special cases

• Denoted:  $x \sim \Gamma(k, \theta)$   $k, \theta > 0$ 

• Mean:  $k\theta$ 

• Variance:  $k\theta^2$ 

• Support:  $x \in (0, \infty)$ 

### Gamma distribution

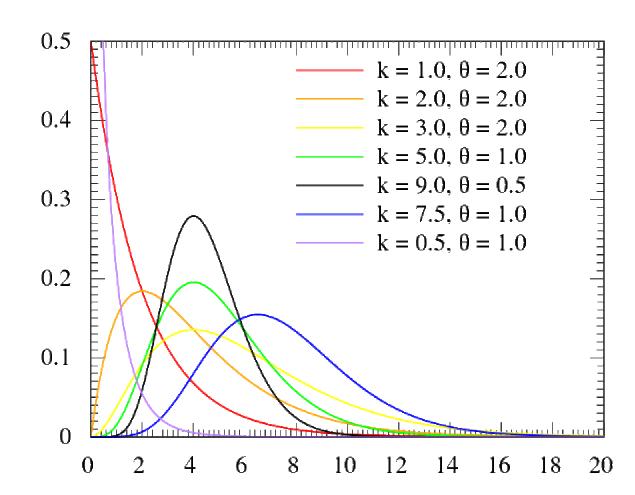
# Probability density function:

$$p(x \mid k, \theta)$$

$$= \frac{1}{\Gamma(k)\theta^k} x^{k-1} e^{-\frac{x}{\theta}}$$

#### Where

$$\Gamma(k) = (k-1)!$$
$$= \int_0^\infty x^{k-1} e^{-x} dx$$



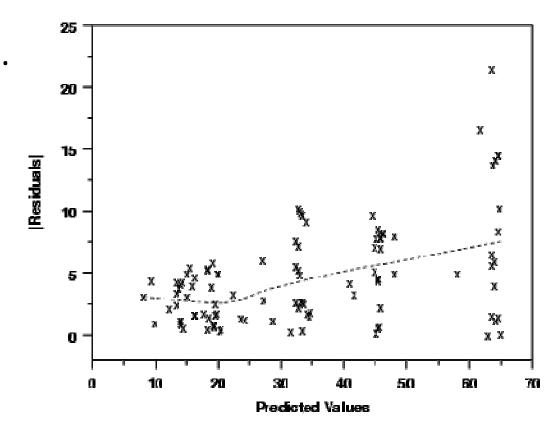
There might be an instance when you want to model something according to the gamma...

 In the instance of a continuous positive value that is heavily skewed, such that the error around the residuals is not normal

# There might be an instance when you want to model something according to the gamma... but it doesn't occur that often

- However, the gamma distribution is useful for understanding other kinds of modeling
- Also useful as a prior in Bayesian analyses

- Poisson distribution commonly used for modeling counts but.... mean=variance assumption can be quite restrictive
- In practice, most data won't mean this assumption
- Data are frequently overdispersed



# GLMs: stochastic part Negative binomial Distribution

An important *discrete* distribution that is useful for modeling counts

• Denoted: 
$$x \sim NB(r, p)$$

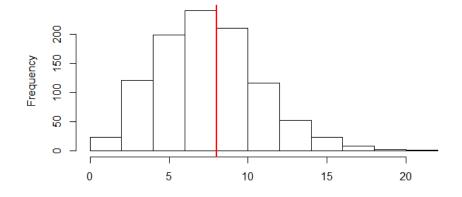
• Mean: 
$$r(1-p)/p$$

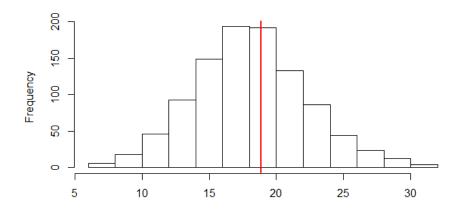
• Variance: 
$$r(1-p)/p^2$$

• Support: 
$$x \in \{0,1,2,3,...\}$$

# GLMs: stochastic part Negative binomial Distribution

- Classical examples: Number of individuals in a flock of birds or school of fish. (e.g., small probabilities of large values)
- Varieties: Mixture of the Poisson and gamma distributions (i.e., an overdispersed Poisson)
- Mathematical description: 2
   parameters: success probability p, and
   "size", r. Consider a number of Bernoulli
   trials in which the probability of
   success = p. Observe this sequence
   until a predefined number r of
   failures. The number of successes, x,
   has a negative binomial distribution.





# GLMs: stochastic part Negative binomial Distribution

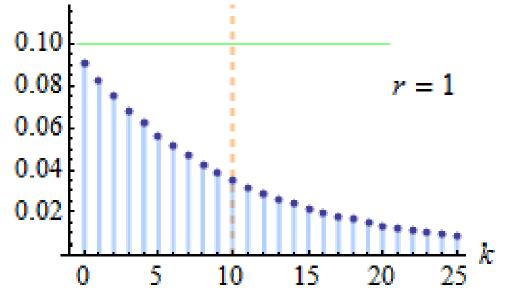
#### Probability mass function:

$$p(x = k \mid r, p) = \frac{(k + r - 1)!}{k! (r - 1)!} p^{r} (1 - p)^{k}$$

#### Likelihood function:

$$L(r, p | X) = \prod_{i=1}^{n} \frac{(x_i + r - 1)!}{x_i! (r - 1)!} p^r (1 - p)^{x_i}$$

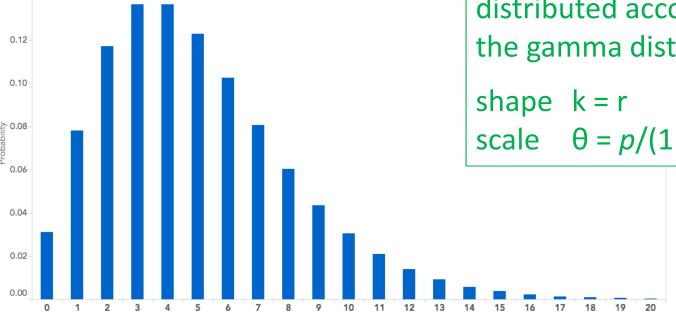
Where 
$$X = \{x_1, x_2, ..., x_n\}$$



# GLMs: stochastic part Negative binomial Distribution

#### Probability mass function:

$$p(x = k \mid r, p) = \frac{\Gamma(k+r)}{\Gamma(r)k!} p^r (1-p)^k$$



Number of Failures

Can view the negative binomial as a Pois(λ) distribution where  $\lambda$  is a random variable distributed according to the gamma distribution:

scale  $\theta = p/(1-p)$ 

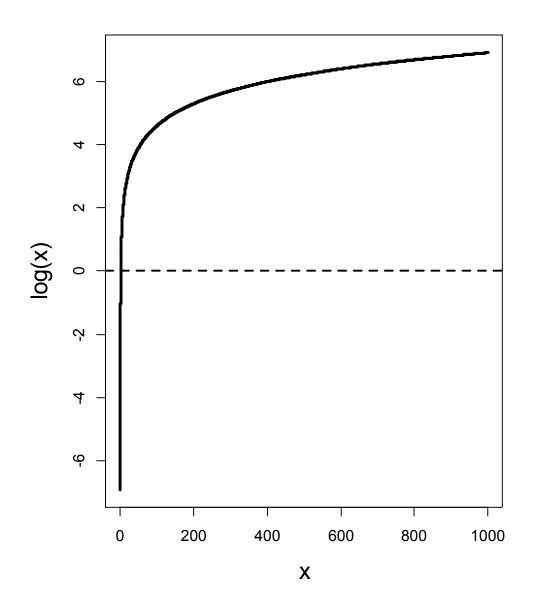
## Which link function to use?

- Modeling counts (whole numbers)
- Expected count must be greater than or equal to zero

#### Log(x)

- Range: x > 0
- Range:

$$-\infty > \log(x) > \infty$$





Estimate the abundance of sea lions relative to local human abundances and amount of prey.

- Data collection:
  - $-n_i$  the number of sea lions counted at location i.
  - $-human_i$  = people per sq km within 100 km of location *i*.
  - $-prey_i$  = the estimated amount of available prey at i.

Is human presence negatively correlated with sea lion abundance?

What is the distribution, link function, and linear predictor we should use?

Distribution:

Link function:

What is the distribution, link function, and linear predictor we should use?

Distribution:  $n_i \sim NB(r_i, p_i)$ 

Link function:

What is the distribution, link function, and linear predictor we should use?

Distribution:  $n_i \sim NB(r_i, p_i)$ 

Link function:  $\log (\mu_i) = \log (r_i(1 - p_i)/p_i)$ 

What is the distribution, link function, and linear predictor we should use?

Distribution:  $n_i \sim NB(r_i, p_i)$ 

Link function:  $\log (\mu_i) = \log (r_i(1 - p_i)/p_i)$ 

Linear predictor:

Note: We want to model the covariates relative to  $\mu$ , the mean, (on the log scale) not r or p.

What is the distribution, link function, and linear predictor we should use?

Distribution:  $n_i \sim NB(r_i, p_i)$ 

Link function:  $\log (\mu_i) = \log (r_i(1 - p_i)/p_i)$ 

Linear predictor:  $\alpha + \beta 1 * human_i + \beta 2 * prey_i$ 

Note: We want to model the covariates relative to  $\mu$ , the mean, (on the log scale) not r or p.

# Moving on to the beta distribution.....

## Beta regression

- Suppose you want to model proportions (e.g., % forest cover relative to tree basal area; allele frequencies)
- How should we one perform a regression analysis in which the dependent variable is restricted to the standard unit interval such as *rates* and *proportions*?

# GLMs: stochastic part Beta Distribution

A *continuous* distribution that is useful for modeling values between 0 and 1

• Denoted: 
$$Beta(a,b)$$
  $a,b>0$ 

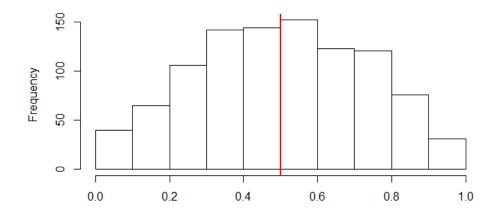
• Mean: 
$$a/(a+b)$$

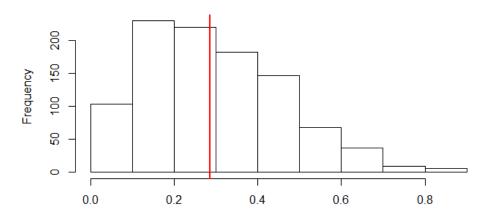
• Variance: 
$$ab/[(a+b)^2(a+b+1)]$$

• Support: 
$$x \in (0,1)$$

# GLMs: stochastic part Beta Distribution

- Sampling situation: Modeling the random behavior of percentages, rates, and proportions
- Classical examples: Allele frequency; Genetic distance between two populations; variability of soil properties; site connectivity
- Varieties: 1) The continuous uniform distribution between 0 and 1 is a special case where a = b = 1;
   2) The Balding-Nichols model is an alternative parametrization used in population genetics.
- Mathematical description: 2 shape parameters: a and b, both greater than zero





# GLMs: stochastic part Beta Distribution

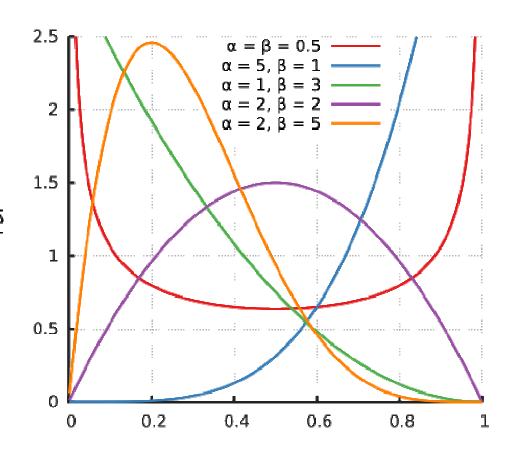
#### Probability density function:

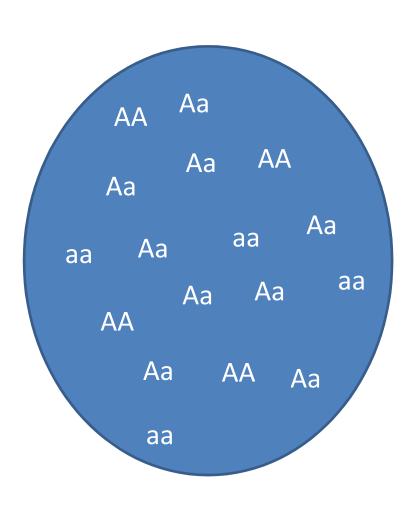
$$p(x \mid a, b) =$$

$$\frac{\Gamma(a+b)}{\Gamma(a)\Gamma(b)} x^{a-1} (1-x)^{b-1} \stackrel{\text{\tiny b}}{=}$$

#### Where

$$\Gamma(k) = \int_0^\infty x^{k-1} e^{-x} dx$$





Estimate whether there is a difference in gene frequency (proportion of aa) in populations that come from one of two ancestors.

- Data collection:
  - $-y_i$  the proportion of aa genotypes in population i.
  - $ancestor_i$  = is an indictor vector where the value = 0 if the ancestor was from the first ancestor or 1 if from the second for population *i*.

What is the distribution, link function, and linear predictor we should use?

Distribution:

Link function:

What is the distribution, link function, and linear predictor we should use?

Distribution:  $y_i \sim Beta(a_i, b_i)$ 

Link function:

What is the distribution, link function, and linear predictor we should use?

Distribution:  $y_i \sim Beta(a_i, b_i)$ 

Link function:  $logit (\mu_i)$ 

What is the distribution, link function, and linear predictor we should use?

Distribution:  $y_i \sim Beta(a_i, b_i)$ 

Link function:  $logit (\mu_i)$ 

Linear predictor:  $\alpha + \beta * ancestor_i$ 

What is the distribution, link function, and linear predictor we should use?

Distribution:  $y_i \sim Beta(a_i, b_i)$ 

Link function:  $logit (\mu_i)$ 

Linear predictor:  $\alpha + \beta * ancestor_i$ 

Note: We want to model the covariates relative to  $\mu$ , the mean, (on the logit scale) not a or b.

$$logit(\mu_i) = logit(a_i/(a_i + b_i))$$

## Generalized linear models

Lab: Negative binomial regression