## Lab 1: Generalized linear models

March 31, 2016

#### Generalized linear models

- Specify distribution for response variable
- Specify linear predictor
- Specify link function
  - Calculates expected response given linear predictor

## Example

Counts for local densities

$$c_i \sim Poisson(\lambda_i)$$
  
 $\log(\lambda_i) = \mathbf{x}_i \boldsymbol{\beta}$ 

## Common distributions for data

#### Discrete

Name	Notation	Domain	Range
Bernoulli	$B \sim Bernoulli(p)$	$0 \le p \le 1$	B = {0, 1}
Binomial	$N \sim Binomial(p, n)$	$0 \le p \le 1$	N = {0, 1,, n}
Poisson	$N \sim Poisson(\lambda)$	λ>0	N = {0, 1,,∞}
Negative binomial	$N \sim Negative Binomial(\lambda, \theta)$	λ>0 θ>0	N = {0, 1,,∞}
Conway-Maxwell- Poisson	$N\sim CMP(\mu,\nu)$	μ>0 ν>0	N = {0, 1,,∞}

## Common distributions for data

## Continuous

Name	Notation	Domain	Range
Normal	$Y \sim Normal(\mu, \sigma^2)$	$\sigma^2 > 0$	Unrestricted
Lognormal	$\log(Y) \sim Normal(\mu, \sigma^2)$	$\sigma^2 > 0$	Y > 0
Gamma	$Y \sim Gamma(\mu, CV)$	μ > 0 CV > 0	Y > 0
Beta	$p \sim Beta(\alpha, \beta)$	$\alpha > 0$ , $\beta > 0$	0 < p < 1

- Choice 1 is it continuous or discrete?
  - Continuous: normal, lognormal, beta, gamma
  - Discrete: Bernoulli, binomial, poisson, negative binomial
- Choice 2 what is the range of possible values?
  - E.g., if discrete:
    - If is is 0 or 1, then its Bernoulli
    - If its between 0 and N, where N is the number of trials, then its Binomial
- Choice 3 How flexible do you want it?

- Frequent null models:
  - 1. Binomial
  - 2. Poisson
  - 3. Normal

#### Binomial

– If you have one or more binary events:

$$B_i \sim \text{Bernoulli}(p)$$

Then the sum of successes...

$$N = \sum_{i=1}^{n_i} B_i$$

... follows a binomial distribution

$$N \sim Binomial(p, n)$$

- Characteristics:

$$\mathbb{E}(N) = pn$$

$$\mathbb{V}(N) = np(1-p)$$

#### Poisson

 If you have a lot of independent events, each with low probability:

$$N \sim \text{Binomial}(p, n)$$

where  $pn \gg 0$  and  $p \ll 1$ 

- Then the number of successes follows a Poisson distribution  $N \sim Poisson(np)$ 

– Characteristics:

$$\mathbb{E}(N) = np$$

$$\mathbb{V}(N) = np$$

- Normal
  - If you have one or more events:

$$B_i \sim g(\mathbf{\theta})$$

where  $g(\theta)$  is some unknown density function

Then the sum of outcomes ...

$$N = \sum_{i=1}^{n_i} b_i$$

... will converge on a normal distribution

$$N \sim \text{Normal}(\mu, \sigma_b^2)$$

... as the number of events gets large  $n_i \rightarrow \infty$ 

$$\mathbb{E}(N) = \mu = n_i \mathbb{E}(g(\mathbf{\theta}))$$

$$\mathbb{V}(N) = \sigma_b^2 = n_i^2 \mathbb{V}(g(\mathbf{\theta}))$$

#### Review:

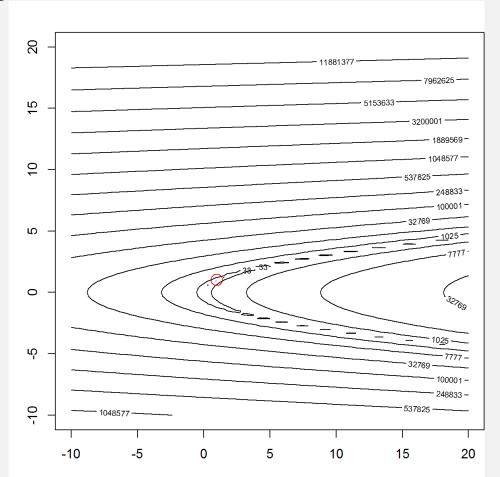
Maximum likelihood estimation (MLE)

$$\widehat{\boldsymbol{\theta}} = \operatorname{argmax}_{\boldsymbol{\theta}}(L(\boldsymbol{\theta}; \mathbf{y}))$$

- Where  $\widehat{m{\theta}}$  is the MLE estimate of parameters
- Where  $\arg\max_{\boldsymbol{\theta}}(L(\boldsymbol{\theta}; \mathbf{y}))$  is the maximum value for  $L(\boldsymbol{\theta}; \mathbf{y})$  that can be achieved for any value of  $\boldsymbol{\theta}$
- argmax is done using maximization algorithms

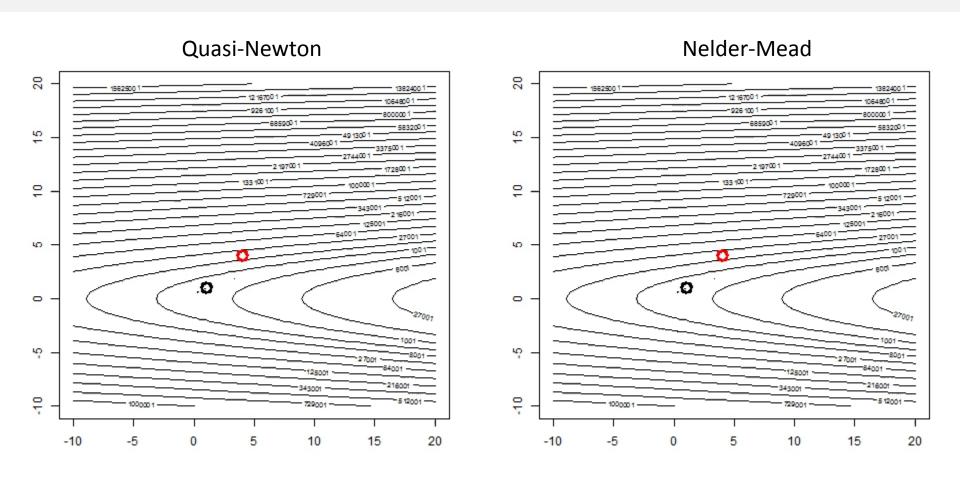
## How to maximize the likelihood function

- Nonlinear minimizers
- Test using Rosenbrook "Banana" function



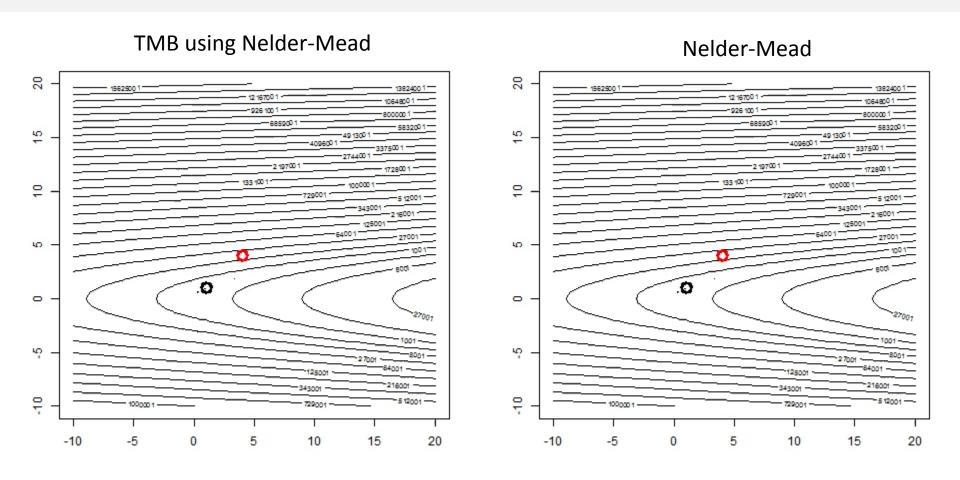
## How to maximize the likelihood function

Methods without gradients are slow



## How to maximize the likelihood function

- Methods with gradients are much faster!



# Example #1 – What is the mean density of canary rockfish in the California Current?

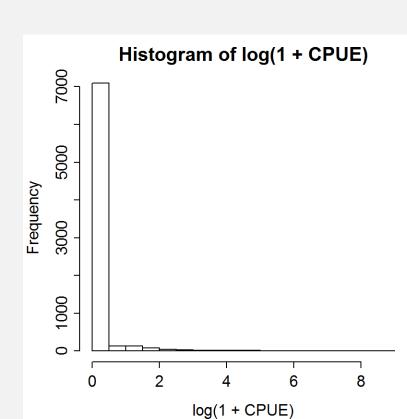
Define linear predictor matrix

$$x_i = 1$$

– i.e.,

$$X = 1$$

We call X an intercept matrix

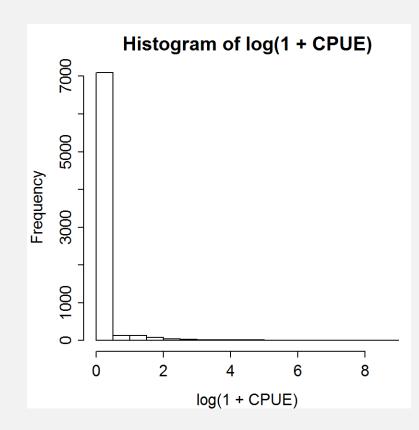


- Generalized linear models
  - Specify distribution for response variable
  - Specify function for expected value
- Canary catch rates

$$\log(\lambda_i) = \mathbf{x}_i \mathbf{\beta}$$

$$\Pr(C = c_i)$$

$$= \begin{cases} \theta_1 & \text{if } c_i = 0 \\ (1 - \theta_1) Lognormal(\lambda_i, \theta_2) & \text{if } c_i > 0 \end{cases}$$



Hint

If:

$$Pr(c_i) = \begin{cases} \theta_1 & \text{if } c_i = 0\\ (1 - \theta_1) Lognormal(\lambda_i, \theta_2) & \text{if } c_i > 0 \end{cases}$$

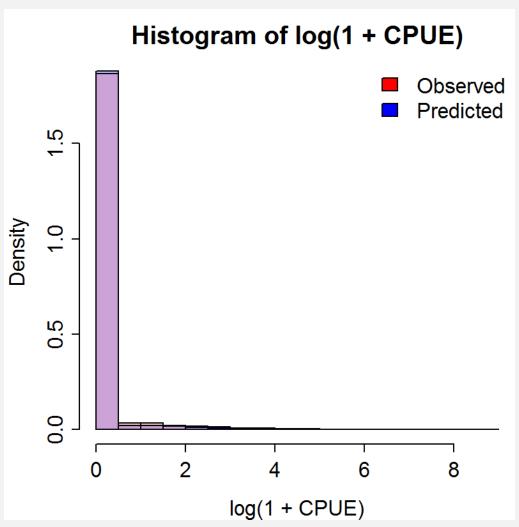
Then:

$$\begin{split} \log(\Pr(c_i)) \\ = \begin{cases} \log(\theta_1) & \text{if } c_i = 0 \\ \log(1 - \theta_1) + \log(Lognormal(\lambda_i, \theta_2)) & \text{if } c_i > 0 \end{cases} \end{split}$$

[Work on TMB code in groups of 2 for 20 minutes]

## Conclusion

Decent fit...



#### How do we assess fit?

- We want expected predictive loss
  - Assume there's a true "data-generating process" (DGP)

$$f(y_i)$$

- Where  $p(y|\theta) = L(\theta; y)$  is your specified probability distribution

predictive probability = 
$$\int p(y^*|\widehat{\boldsymbol{\theta}}) f(y^*) dy^*$$

- Where
  - y\* is some future data

Then

expected predictive log – probability = 
$$\sum_{j=1}^{J} \log(p(y_j|\widehat{\boldsymbol{\theta}}))$$

- Where
  - $y_i$  is some data that were "held out" when estimating parameters  $\widehat{m{\theta}}$

More reading: Gelman, A., Hwang, J. & Vehtari, A. (2014). Understanding predictive information criteria for Bayesian models. *Stat. Comput.*, 24, 997–1016.

#### How do we assess fit?

- K-fold crossvalidation
  - 1. Divide data set into *K* even partitions
  - 2. Calculate predictive probability for 1<sup>st</sup> partition
    - For each piece K, fit the model to all data except data in that partition
    - Calculate the predictive probability of data in partition K using this model
    - Record predictive probability
  - 3. Repeat step 2 for all K partitions
  - 4. Chose the model with the highest predictive probability

#### Confidence interval:

- Parameter estimates are normally distributed
- Computation

$$CI_{\chi\%}(\hat{\theta}) = \hat{\theta} \pm \widehat{SE}(\hat{\theta}) \times \Phi^{-1}(\frac{x}{2})$$

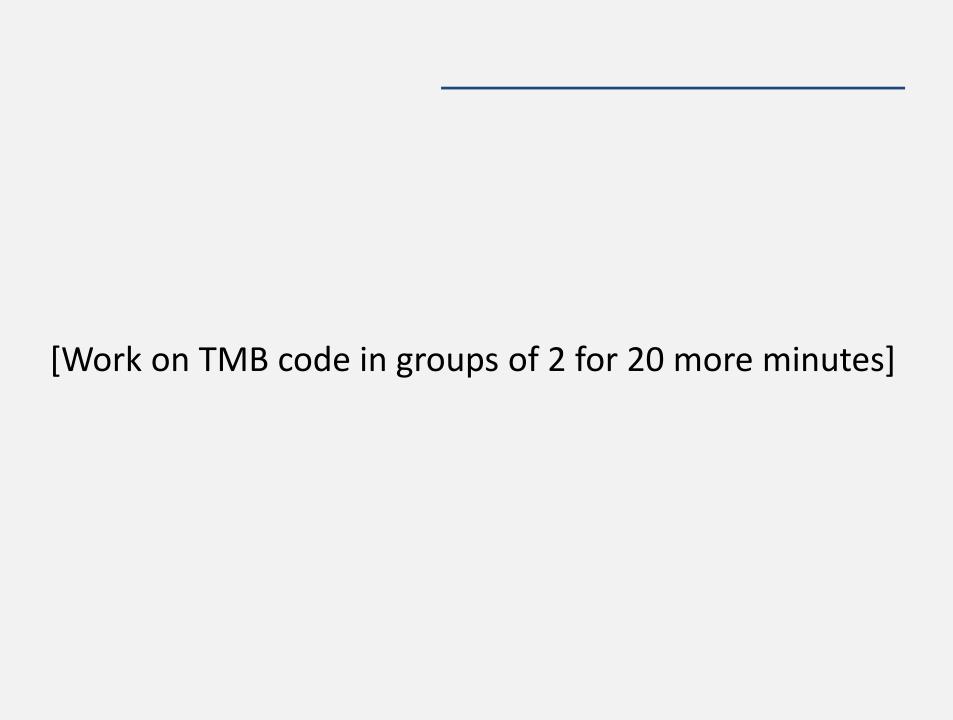
- Where  $CI_{x\%}$  contains the true value x% of the time if the model is correct
- $-\Phi^{-1}$  is the inverse cumulative distribution for a normal distribution
- $\hat{ heta}$  is the estimate for parameter heta
- $-\widehat{SE}(\widehat{\theta})$  is the estimated standard error for parameter  $\theta$

## Confidence interval coverage

Coverage – the expected proportion of times that an estimated x% confidence interval contains the true value given an estimation model and true "data-generating process"

#### **Estimation:**

- 1. Simulate data with a known value for parameter  $\theta$
- 2. Record true parameter values
- 3. Apply estimator
- 4. Record confidence interval  $CI_{x\%}(\widehat{\theta})$  for parameter  $\theta$
- 5. Repeat steps 1-4 hundreds of times
- 6. Compute the proportion of times where  $CI_{x\%}(\widehat{\theta})$  contains the true value for parameter  $\theta$



## Homework assignment:

- Due at beginning of Lab #2
- Must turn in your own code
- Cannot cut-paste any code from other students
  - You can hand-write your own code while working with someone else, or looking at my example code