

Lab 1: Generalized linear models

March 31, 2015

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- 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

$$c_i \sim \text{Poisson}(\lambda_i)$$

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

Common distributions for data

– Discrete

Name	Notation	Domain	Range
Bernoulli	$B \sim \text{Bernoulli}(p)$	$0 \leq p \leq 1$	$B = \{0, 1\}$
Binomial	$N \sim \text{Binomial}(p, n)$	$0 \leq p \leq 1$	$N = \{0, 1, \dots, n\}$
Poisson	$N \sim \text{Poisson}(\lambda)$	$\lambda > 0$	$N = \{0, 1, \dots, \infty\}$
Negative binomial	$N \sim \text{NegativeBinomial}(\lambda, \theta)$	$\lambda > 0$ $\theta > 0$	$N = \{0, 1, \dots, \infty\}$
Conway-Maxwell-Poisson	$N \sim \text{CMP}(\mu, \nu)$	$\mu > 0$ $\nu > 0$	$N = \{0, 1, \dots, \infty\}$

Common distributions for data

– Continuous

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

How to choose a distribution for data?

- 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 it is 0 or 1, then it's Bernoulli
 - If it's between 0 and N, where N is the number of trials, then it's Binomial
- Choice 3 – How flexible do you want it?

How to choose a distribution for data?

- Frequent null models:

1. Binomial
2. Poisson
3. Normal

How to choose a distribution for data?

- 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

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

- Characteristics:

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

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

How to choose a distribution for data?

- Poisson

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

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

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

- Then the number of successes follows a Poisson distribution

$$N \sim \text{Poisson}(np)$$

- Characteristics:

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

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

How to choose a distribution for data?

- Normal

- If you have one or more events:

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

where $g(\boldsymbol{\theta})$ 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 $n_i \rightarrow \infty$

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

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

Review:

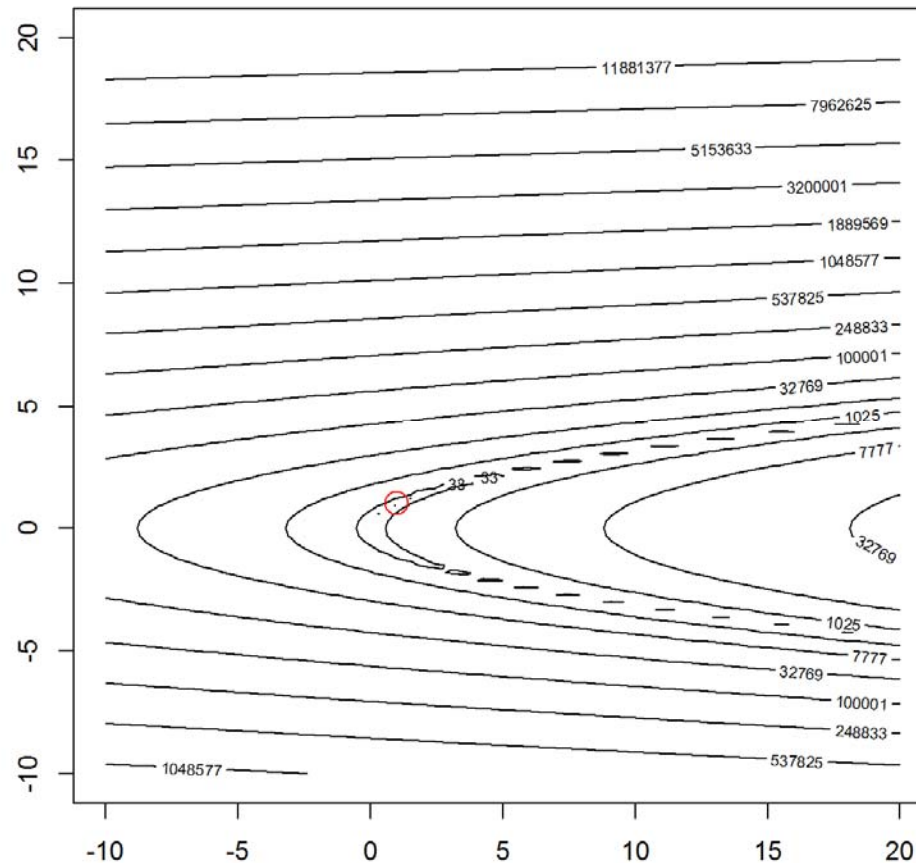
- Maximum likelihood estimation (MLE)

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

- Where $\hat{\boldsymbol{\theta}}$ is the MLE estimate of parameters
- Where $\operatorname{argmax}_{\boldsymbol{\theta}}(L(\boldsymbol{\theta}; \mathbf{y}))$ 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

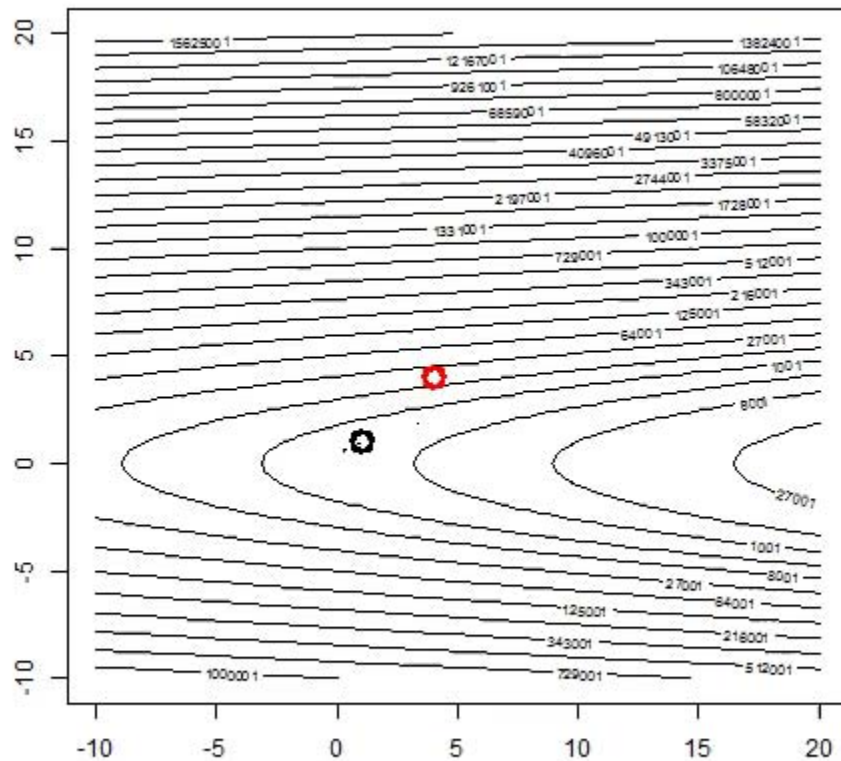
- Nonlinear minimizers
- Test using Rosenbrock “Banana” function



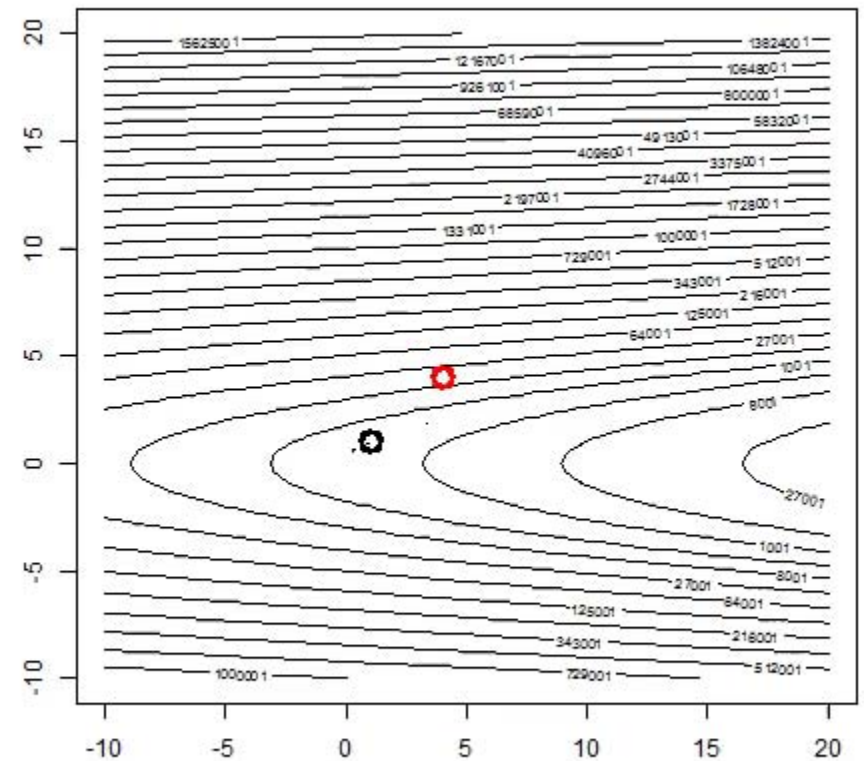
How to maximize the likelihood function

- Methods without gradients are slow

Quasi-Newton



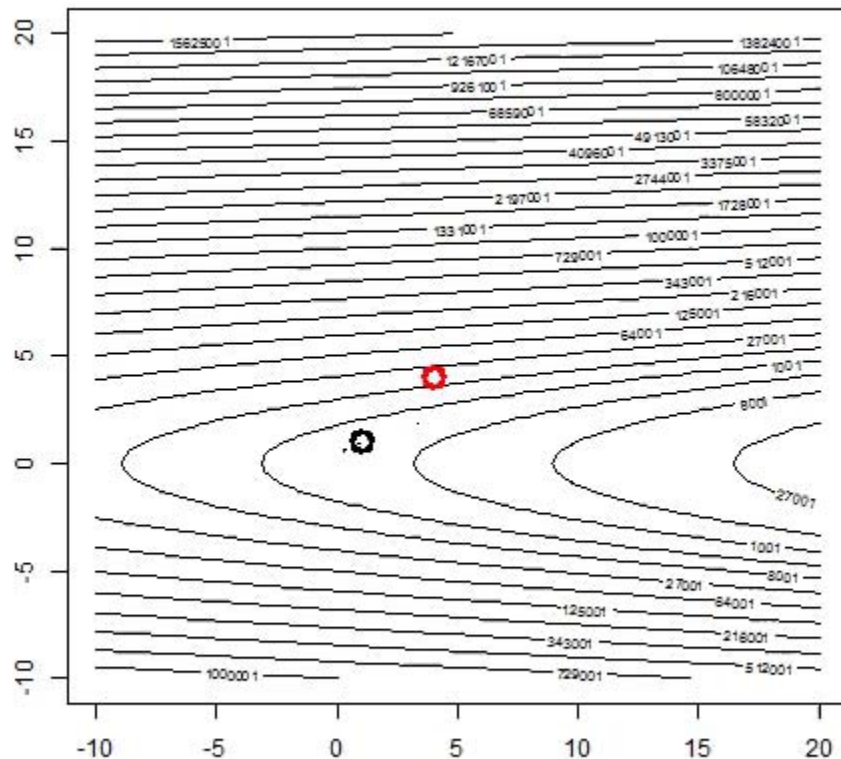
Nelder-Mead



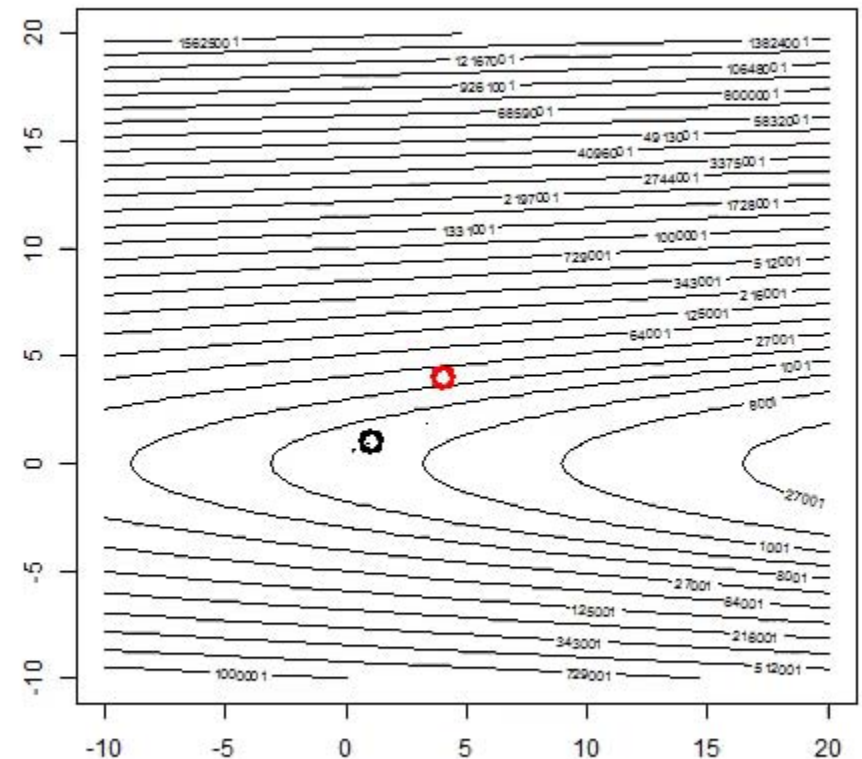
How to maximize the likelihood function

- Methods with gradients are much faster!

TMB using Nelder-Mead



Nelder-Mead



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

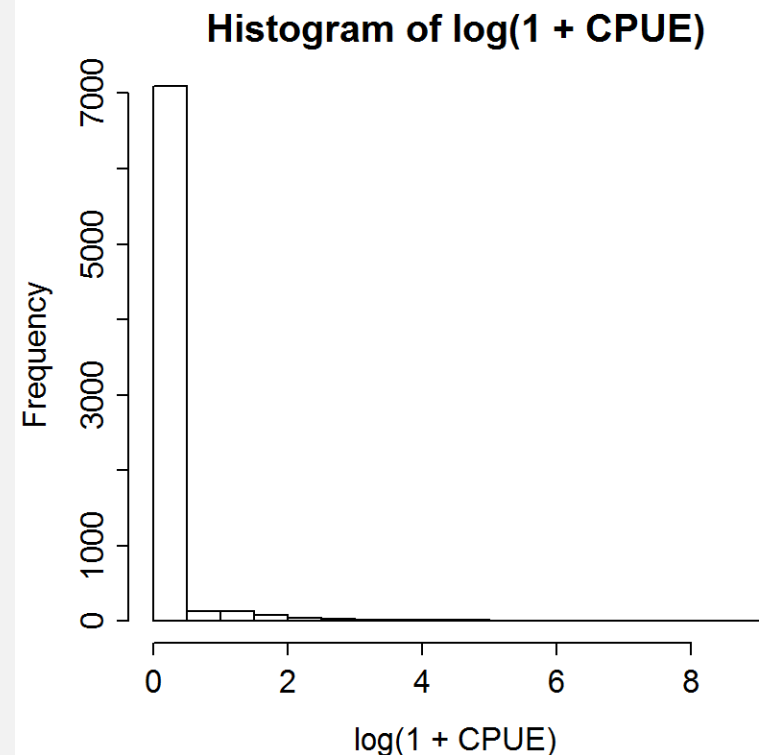
- Define linear predictor matrix

$$x_i = 1$$

– i.e.,

$$\mathbf{X} = \mathbf{1}$$

– We call \mathbf{X} an intercept matrix

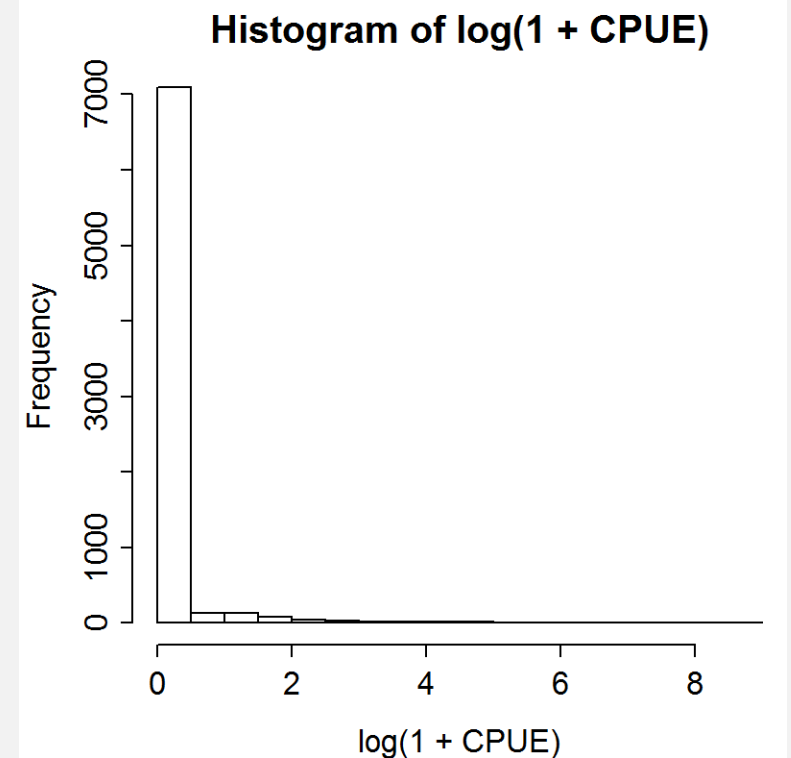


- Generalized linear models
 - Specify distribution for response variable
 - Specify function for expected value

- Canary catch rates

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

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



Hint

If:

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

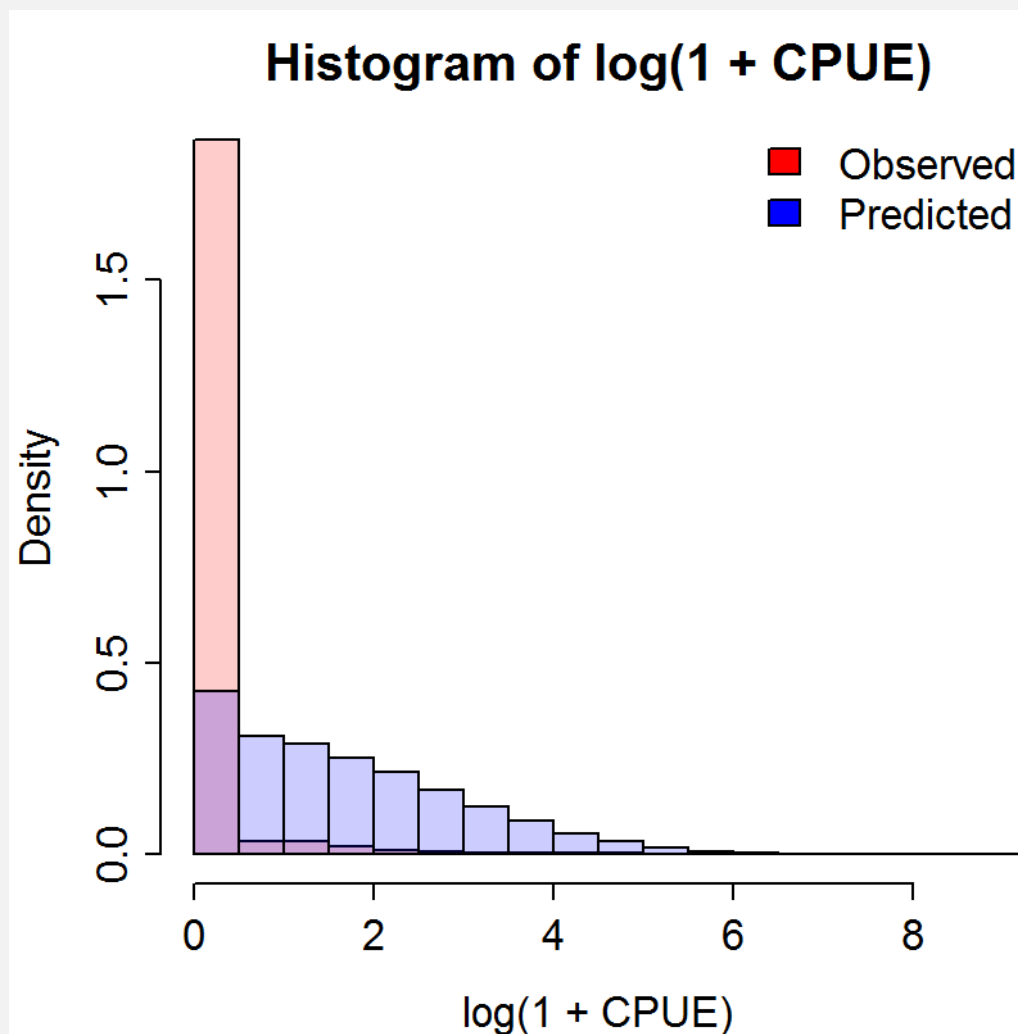
Then:

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

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

- Conclusion

- Still doesn't give very good fit!



How do we assess fit?

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

$$\Pr(y_i|\boldsymbol{\theta}_0) \sim f(y_i|\boldsymbol{\theta}_0)$$

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

$$\text{predictive score} \propto E \left(p(Y|\hat{\boldsymbol{\theta}}) \right) = \int f(y|\boldsymbol{\theta}_0) p(y|\hat{\boldsymbol{\theta}}) dy$$

- Where
 - $p(y^*|\hat{\boldsymbol{\theta}})$ is your predictive distribution for a new observation
 - $f(y^*|\boldsymbol{\theta}_0)$ is the true probability of

How do we assess fit?

- K-fold crossvalidation

1. Divide data set into K even partitions
2. Calculate predictive probability for 1st 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

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

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

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

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

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