

Chapter 4

- 4.1 Normal approximation (Laplace's method)
- 4.2 Large-sample theory
- 4.3 Counter examples
 - includes examples of difficult posteriors for MCMC, too
- 4.4 Frequency evaluation*
- 4.5 Other statistical methods*

Normal approximation (Laplace approximation)

- Often posterior converges to normal distribution when $n \rightarrow \infty$

- If posterior is unimodal and close to symmetric
 - we can approximate $p(\theta|y)$ with normal distribution

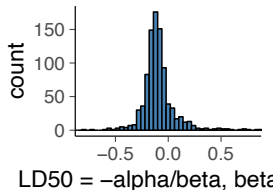
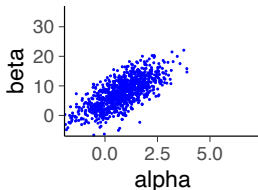
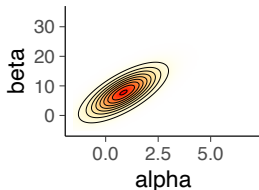
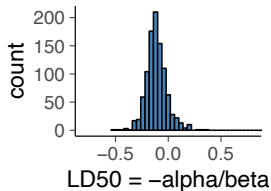
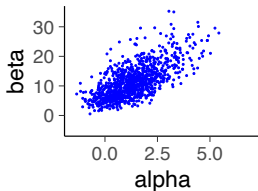
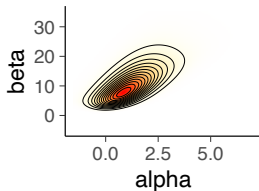
$$p(\theta|y) \approx \frac{1}{\sqrt{2\pi}\sigma_\theta} \exp\left(-\frac{1}{2\sigma_\theta^2}(\theta - \hat{\theta})^2\right)$$

- Laplace used this (before Gauss) to approximate the posterior of binomial model to infer ratio of girls and boys born
- A most strict proof by LeCam in 1950's

Normal approximation (Laplace approximation)

- Often posterior converges to normal distribution when $n \rightarrow \infty$
- If posterior is unimodal and close to symmetric
 - we can approximate $p(\theta|y)$ with normal distribution

$$p(\theta|y) \approx \frac{1}{\sqrt{2\pi}\sigma_\theta} \exp\left(-\frac{1}{2\sigma_\theta^2}(\theta - \hat{\theta})^2\right)$$



Taylor series

- We can approximate $p(\theta|y)$ with normal distribution

$$p(\theta|y) \approx \frac{1}{\sqrt{2\pi}\sigma_\theta} \exp\left(-\frac{1}{2\sigma_\theta^2}(\theta - \hat{\theta})^2\right)$$

- i.e. log posterior $\log p(\theta|y)$ can be approximated with a quadratic function

$$\log p(\theta|y) \approx \alpha(\theta - \hat{\theta})^2 + C$$

- Univariate Taylor series expansion around $\theta = \hat{\theta}$

$$f(\theta) = f(\hat{\theta}) + f'(\hat{\theta})(\theta - \hat{\theta}) + \frac{f''(\hat{\theta})}{2!}(\theta - \hat{\theta})^2 + \frac{f^{(3)}(\hat{\theta})}{3!}(\theta - \hat{\theta})^3 + \dots$$

- if $\hat{\theta}$ is at mode, then $f'(\hat{\theta}) = 0$
- often when $n \rightarrow \infty$, $\frac{f^{(3)}(\hat{\theta})}{3!}(\theta - \hat{\theta})^3 + \dots$ is small

Multivariate Taylor series

- Multivariate series expansion

$$f(\theta) = f(\hat{\theta}) + \frac{df(\theta')}{d\theta'} \Big|_{\theta'=\hat{\theta}} (\theta - \hat{\theta}) + \frac{1}{2!} (\theta - \hat{\theta})^T \frac{d^2f(\theta')}{d\theta'^2} \Big|_{\theta'=\hat{\theta}} (\theta - \hat{\theta}) + \dots$$

Normal approximation

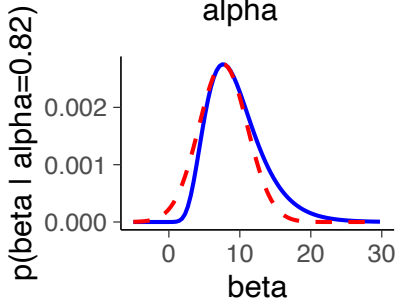
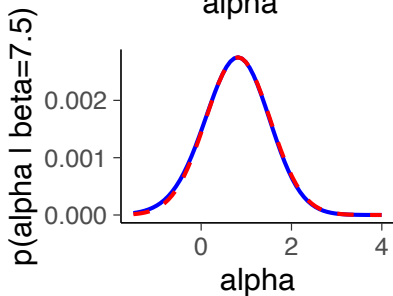
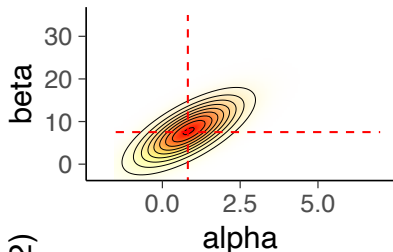
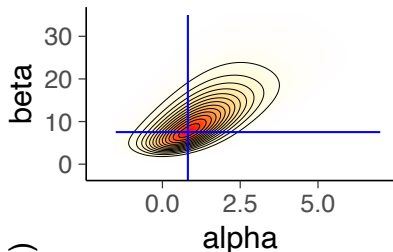
- Taylor series expansion of the log posterior around the posterior mode $\hat{\theta}$

$$\log p(\theta|y) = \log p(\hat{\theta}|y) + \frac{1}{2}(\theta - \hat{\theta})^T \left[\frac{d^2}{d\theta^2} \log p(\theta'|y) \right]_{\theta'=\hat{\theta}} (\theta - \hat{\theta}) + \dots$$

- Multivariate normal $\propto |\Sigma|^{-1/2} \exp \left(-\frac{1}{2}(\theta - \hat{\theta})^T \Sigma^{-1} (\theta - \hat{\theta}) \right)$

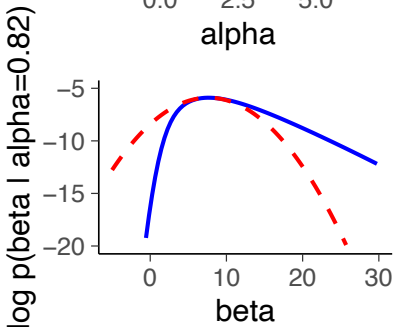
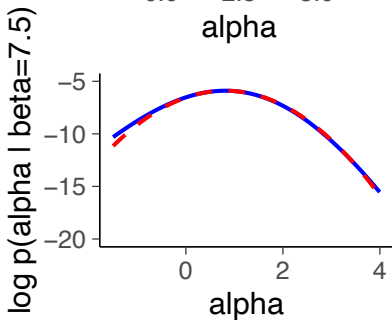
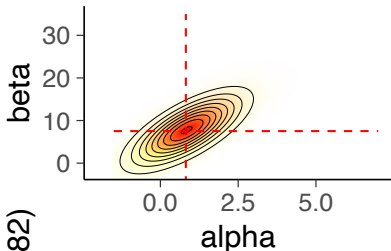
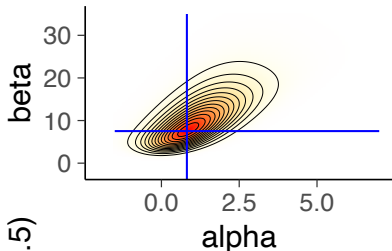
Normal approximation

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Normal approximation

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Normal approximation

- Taylor series expansion of the log posterior around the posterior mode $\hat{\theta}$

$$\log p(\theta|y) = \log p(\hat{\theta}|y) + \frac{1}{2}(\theta - \hat{\theta})^T \left[\frac{d^2}{d\theta^2} \log p(\theta'|y) \right]_{\theta'=\hat{\theta}} (\theta - \hat{\theta}) + \dots$$

- Multivariate normal $\propto |\Sigma|^{-1/2} \exp \left(-\frac{1}{2}(\theta - \hat{\theta})^T \Sigma^{-1} (\theta - \hat{\theta}) \right)$
- Normal approximation

$$p(\theta|y) \approx N(\hat{\theta}, [I(\hat{\theta})]^{-1})$$

where $I(\theta)$ is called *observed information*

$$I(\theta) = -\frac{d^2}{d\theta^2} \log p(\theta|y)$$

Normal approximation

- $I(\theta)$ is called *observed information*

$$I(\theta) = -\frac{d^2}{d\theta^2} \log p(\theta|y)$$

- $I(\hat{\theta})$ is the second derivatives at the mode and thus describes the curvature at the mode
- if the mode is inside the parameter space, $I(\hat{\theta})$ is positive
- if θ is a vector, then $I(\theta)$ is a matrix

Normal approximation

- BDA3 Ch 4 has an example where it is easy to compute first and second derivatives and there is easy analytic solution to find where the first derivatives are zero

Normal approximation – example

- Normal distribution, unknown mean and variance
 - uniform prior $(\mu, \log \sigma)$
 - normal approximation for the posterior of $(\mu, \log \sigma)$

$$\log p(\mu, \log \sigma | y) = \text{constant} - n \log \sigma - \frac{1}{2\sigma^2} [(n-1)s^2 + n(\bar{y} - \mu)^2]$$

first derivatives

$$\begin{aligned} \frac{d}{d\mu} \log p(\mu, \log \sigma | y) &= \frac{n(\bar{y} - \mu)}{\sigma^2}, \\ \frac{d}{d(\log \sigma)} \log p(\mu, \log \sigma | y) &= -n + \frac{(n-1)s^2 + n(\bar{y} - \mu)^2}{\sigma^2}, \end{aligned}$$

from which it is easy to compute the mode

$$(\hat{\mu}, \log \hat{\sigma}) = \left(\bar{y}, \frac{1}{2} \log \left(\frac{n-1}{n} s^2 \right) \right)$$

Normal approximation – example

- Normal distribution, unknown mean and variance
first derivatives

$$\frac{d}{d\mu} \log p(\mu, \log \sigma | y) = \frac{n(\bar{y} - \mu)}{\sigma^2},$$

$$\frac{d}{d(\log \sigma)} \log p(\mu, \log \sigma | y) = -n + \frac{(n-1)s^2 + n(\bar{y} - \mu)^2}{\sigma^2}$$

second derivatives

$$\frac{d^2}{d\mu^2} \log p(\mu, \log \sigma | y) = -\frac{n}{\sigma^2},$$

$$\frac{d^2}{d\mu d(\log \sigma)} \log p(\mu, \log \sigma | y) = -2n \frac{\bar{y} - \mu}{\sigma^2},$$

$$\frac{d^2}{d(\log \sigma)^2} \log p(\mu, \log \sigma | y) = -\frac{2}{\sigma^2} ((n-1)s^2 + n(\bar{y} - \mu)^2)$$

Normal approximation – example

- Normal distribution, unknown mean and variance
second derivatives

$$\frac{d^2}{d\mu^2} \log p(\mu, \log \sigma | y) = -\frac{n}{\sigma^2},$$

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$$\frac{d^2}{d(\log \sigma)^2} \log p(\mu, \log \sigma | y) = -\frac{2}{\sigma^2} ((n-1)s^2 + n(\bar{y} - \mu)^2)$$

matrix of the second derivatives at $(\hat{\mu}, \log \hat{\sigma})$

$$\begin{pmatrix} -n/\hat{\sigma}^2 & 0 \\ 0 & -2n \end{pmatrix}$$

Normal approximation – example

- Normal distribution, unknown mean and variance posterior mode

$$(\hat{\mu}, \log \hat{\sigma}) = \left(\bar{y}, \frac{1}{2} \log \left(\frac{n-1}{n} s^2 \right) \right)$$

matrix of the second derivatives at $(\hat{\mu}, \log \hat{\sigma})$

$$\begin{pmatrix} -n/\hat{\sigma}^2 & 0 \\ 0 & -2n \end{pmatrix}$$

normal approximation

$$p(\mu, \log \sigma | y) \approx N \left(\begin{pmatrix} \mu \\ \log \sigma \end{pmatrix} \middle| \begin{pmatrix} \bar{y} \\ \log \hat{\sigma} \end{pmatrix}, \begin{pmatrix} \hat{\sigma}^2/n & 0 \\ 0 & 1/(2n) \end{pmatrix} \right)$$

Normal approximation – numerically

- Normal approximation can be computed numerically
 - iterative optimization to find a mode (may use gradients)
 - autodiff or finite-difference for gradients and Hessian
 - e.g. in R, demo4_1.R:

```
bioassayfun <- function(w, df) {  
  z <- w[1] + w[2]*df$x  
  -sum(df$y*(z) - df$n*log1p(exp(z)))  
}
```

```
theta0 <- c(0,0)  
optimres <- optim(w0, bioassayfun, gr=NULL, df1, hessian=T)  
thetahat <- optimres$par  
Sigma <- solve(optimres$hessian)
```


Normal approximation – numerically

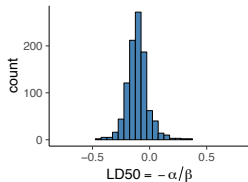
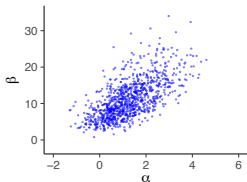
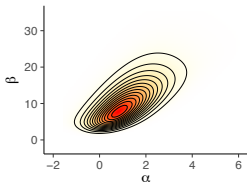
- Normal approximation can be computed numerically
 - iterative optimization to find a mode (may use gradients)
 - autodiff or finite-difference for gradients and Hessian
- RStanARM has an option `algorithm='optimizing'`
 - uses L-BFGS quasi-Newton optimization algorithm for finding the mode
 - uses autodiff for gradients
 - uses finite differences of gradients to compute Hessian
 - second order autodiff coming to Stan

Normal approximation

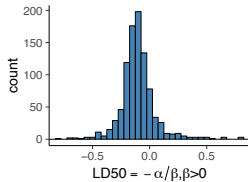
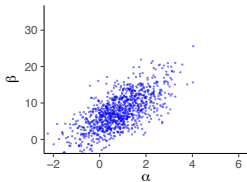
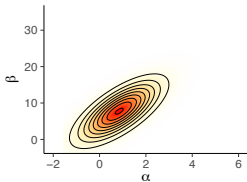
- Optimization and computation of Hessian requires usually much less density evaluations than MCMC
- In some cases accuracy is sufficient
- In some cases accuracy for a conditional distribution is sufficient (Ch 13)
 - e.g. Gaussian latent variable models, such as Gaussian processes (Ch 21)
 - Rasmussen & Williams: Gaussian Processes for Machine Learning
- Accuracy can be improved by importance sampling (Ch 10)

Example: Importance sampling in Bioassay

Grid



Normal

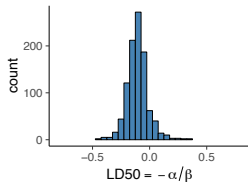
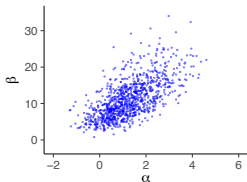
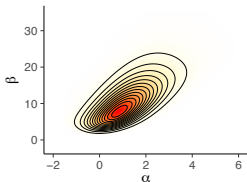


But the normal approximation is not that good here:

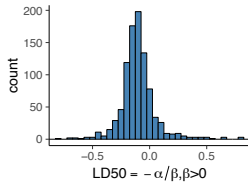
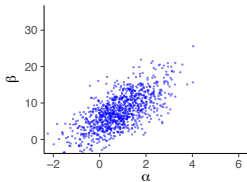
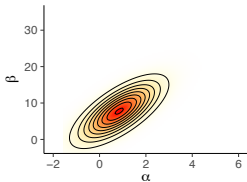
Grid $sd(LD50) \approx 0.1$, Normal $sd(LD50) \approx .75!$

Example: Importance sampling in Bioassay

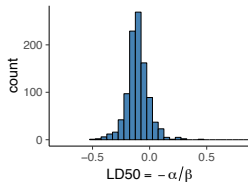
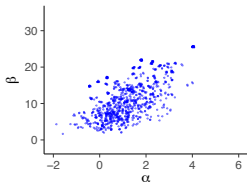
Grid



Normal



IS



Grid $sd(LD50) \approx 0.1$, IS $sd(LD50) \approx 0.1$

Normal approximation

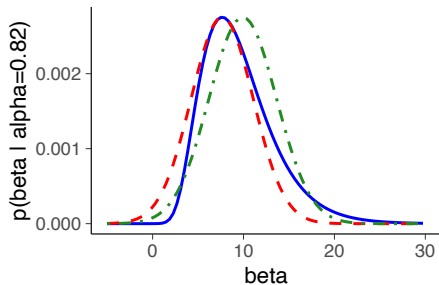
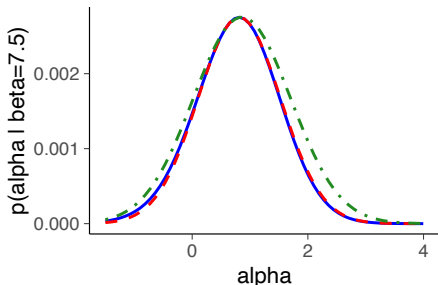
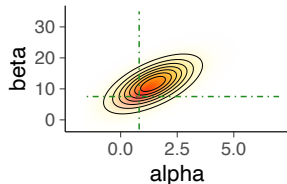
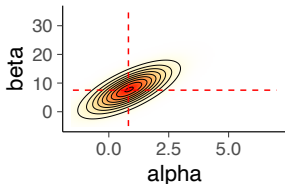
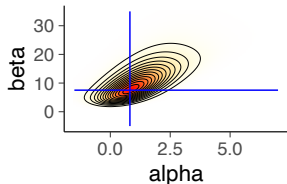
- Accuracy can be improved by importance sampling
- Pareto- k diagnostic of importance sampling weights can be used for diagnostic
 - in Bioassay example $k = 0.57$, which is ok
- RStanARM has an option `algorithm='optimizing'`
 - since version 2.19.2 (2019-10-03)
 - + Pareto- k diagnostic
 - + importance resampling (IR)

Other distributional approximations*

- Higher order derivatives at the mode can be used
- Split-normal and split- t by Geweke use additional scaling along different principal axes
- Other distributions can be used (e.g. t -distribution)
- Instead of mode and Hessian at mode, e.g.
 - variational inference (Ch 13)
 - CS-E4820 - Machine Learning: Advanced Probabilistic Methods
 - Stan has an experimental ADVI algorithm
 - expectation propagation (Ch 13)
 - speed of these is usually between optimization and MCMC

Distributional approximations

Exact, Normal at mode, Normal with variational inference



Grid sd(LD50) ≈ 0.090 ,
Normal sd(LD50) $\approx .75$, Normal + IR sd(LD50) ≈ 0.096 (Pareto- $k = 0.57$)
VI sd(LD50) ≈ 0.13 , VI + IR sd(LD50) ≈ 0.095 (Pareto- $k = 0.17$)

Large sample theory

- Asymptotic normality
 - as n the number of observations y_i increases the posterior converges to normal distribution

Large sample theory

- Assume "true" underlying data distribution $f(y)$
 - observations y_1, \dots, y_n are independent samples from the joint distribution $f(y)$
 - "true" data distribution $f(y)$ is not always well defined
 - in the following we proceed as if there were true underlying data distribution
 - for the theory the exact form of $f(y)$ is not important as long as it has certain regularity conditions

Large sample theory

- Consistency

- if true distribution is included in the parametric family, so that $f(y) = p(y|\theta_0)$ for some θ_0 , then posterior converges to a point θ_0 , when $n \rightarrow \infty$
- a point doesn't have uncertainty
- same result as for maximum likelihood estimate
- If true distribution is not included in the parametric family, then there is no true θ_0
 - true θ_0 is replaced with θ_0 which minimizes the Kullback-Leibler divergence from $f(y)$

$$H(\theta_0) = \int f(y_i) \log \left(\frac{f(y_i)}{p(y_i|\theta_0)} \right) dy_i$$

- this point doesn't have uncertainty, but it's a wrong point!
- same result as for maximum likelihood estimate

Large sample theory – counter examples

- Under- and non-identifiability
 - a model is under-identifiable, if the model has parameters or parameter combinations for which there is no information in the data
 - then there is no single point θ_0 where posterior would converge
 - e.g. if the model is

$$y \sim N(a + b + cx, \sigma)$$

- posterior would converge to a line with prior determining the density along the line
- e.g. if we never observe u and v at the same time and the model is

$$\begin{pmatrix} u \\ v \end{pmatrix} \sim N \left(\begin{pmatrix} 0 \\ 0 \end{pmatrix}, \begin{pmatrix} 1 & \rho \\ \rho & 1 \end{pmatrix} \right)$$

then correlation ρ is non-identifiable

- e.g. u and v could be length and weight of a student; if only one of them is measured for each student, then ρ is non-identifiable
- Problem also for other inference methods like MCMC

Large sample theory – counter examples

- If the number of parameter increases as the number of observation increases
 - in some models number of parameters depends on the number of observations
 - e.g. time series models $y_i \sim N(\theta_i, \sigma^2)$ and θ_i has prior in time
 - posterior of θ_i does not converge to a point, if additional observations do not bring enough information

Large sample theory – counter examples

- Aliasing (FI: [valettoisto](#))

- special case of under-identifiability where likelihood repeats in separate points
- e.g. mixture of normals

$$p(y_i | \mu_1, \mu_2, \sigma_1^2, \sigma_2^2, \lambda) = \lambda N(\mu_1, \sigma_1^2) + (1 - \lambda) N(\mu_2, \sigma_2^2)$$

if (μ_1, μ_2) are switched, (σ_1^2, σ_2^2) are switched and replace λ with $(1 - \lambda)$, model is equivalent; posterior would usually have two modes which are mirror images of each other and the posterior does not converge to a single point

- For MCMC makes the convergence diagnostics more difficult, as it is difficult to identify aliasing from other multimodality

Large sample theory – counter examples

- Unbounded (FI: [rajoittamaton](#)) likelihood
 - if likelihood is unbounded it is possible that there is no mode in the posterior
 - e.g. previous normal mixture model; assume λ to be known (and not 0 or 1); if we set $\mu_1 = y_i$ for any i and $\sigma_1^2 \rightarrow 0$, then likelihood $\rightarrow \infty$
 - if prior for σ_1^2 does not go to zero when $\sigma_1^2 \rightarrow 0$, then the posterior is unbounded
 - when $n \rightarrow \infty$ the number of likelihood modes increases
- Problem for any inference method including MCMC
 - can be avoided with good priors
 - note that a prior close to a prior allowing unbounded posterior may produce almost unbounded posterior

Large sample theory – counter examples

- Improper posterior
 - asymptotic results assume that probability sums to 1
 - e.g. Binomial model, with $\text{Beta}(0, 0)$ prior and observation $y = n$
 - posterior $p(\theta|n, 0) = \theta^{n-1}(1 - \theta)^{-1}$
 - when $\theta \rightarrow 1$, then $p(\theta|n, 0) \rightarrow \infty$
- Problem for any inference method including MCMC
 - can be avoided with proper priors
 - note that prior close to a improper prior may produce almost improper posterior

Large sample theory – counter examples

- Prior distribution does not include the convergence point
 - if in discrete case $p(\theta_0) = 0$ or in continuous case $p(\theta) = 0$ in the neighborhood of θ_0 , then the convergence results based on the dominance of the likelihood do not hold
- Should have a positive prior probability/density where needed

Large sample theory – counter examples

- Convergence point at the edge of the parameter space
 - if θ_0 is on the edge of the parameter space, Taylor series expansion has to be truncated, and normal approximation does not necessarily hold
 - e.g. $y_i \sim N(\theta, 1)$ with a restriction $\theta \geq 0$ and assume that $\theta_0 = 0$
 - posterior of θ is left truncated normal distribution with $\mu = \bar{y}$
 - in the limit $n \rightarrow \infty$ posterior is half normal distribution
- Can be easy or difficult for MCMC

Large sample theory – counter examples

- Tails of the distribution
 - normal approximation may be accurate for the most of the posterior mass, but still be inaccurate for the tails
 - e.g. parameter which is constrained to be positive; given a finite n , normal approximation assumes non-zero probability for negative values

Frequency evaluations

- Bayesian theory has epistemic and aleatory probabilities
- Frequency evaluations focus on frequency properties given aleatoric repetition of an observation and modeling
 - Consistency
 - Asymptotic unbiasedness
 - not that important in Bayesian inference, small and decreasing error more important
 - Asymptotic efficiency
 - no other point estimate with smaller squared error

Frequency evaluations

- Bayesian theory has epistemic and aleatory probabilities
- Frequency evaluations focus on frequency properties given aleatoric repetition of an observation and modeling
 - Consistency
 - Asymptotic unbiasedness
 - not that important in Bayesian inference, small and decreasing error more important
 - Asymptotic efficiency
 - no other point estimate with smaller squared error
 - Calibration
 - $\alpha\%$ -posterior interval has the true value in $\alpha\%$ cases
 - $\alpha\%$ -predictive interval has the true future values in $\alpha\%$ cases
 - approximate calibration with shorter intervals for likely true values more important than exact calibration with bad intervals for all possible values.