## 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 $n \rightarrow \infty$
- bounded, non-singular, the number of parameters don't grow with $n$
- we can then approximate $p(\theta \mid y)$ with normal distribution


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- Laplace used this (before Gauss) to approximate the posterior of binomial model to infer ratio of girls and boys born


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## Taylor series

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p(\theta \mid y) \approx \frac{1}{\sqrt{2 \pi} \sigma_{\theta}} \exp \left(-\frac{1}{2 \sigma_{\theta}^{2}}(\theta-\hat{\theta})^{2}\right)
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- i.e. $\log$ posterior $\log p(\theta \mid y)$ can be approximated with a quadratic function

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f(\theta)=f(\hat{\theta})+f^{\prime}(\hat{\theta})(\theta-\hat{\theta})+\frac{f^{\prime \prime}(\hat{\theta})}{2!}(\theta-\hat{\theta})^{2}+\frac{f^{(3)}(\hat{\theta})}{3!}(\theta-\hat{\theta})^{3}+\ldots
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- if $\hat{\theta}$ is at mode, then $f^{\prime}(\hat{\theta})=0$
- often when $n \rightarrow \infty, \frac{f^{(3)}(\hat{\theta})}{3!}(\theta-\hat{\theta})^{3}+\ldots$ is small


## Multivariate Taylor series

- Multivariate series expansion

$$
f(\theta)=f(\hat{\theta})+{\frac{d f\left(\theta^{\prime}\right)}{d \theta^{\prime}}}_{\theta^{\prime}=\hat{\theta}}(\theta-\hat{\theta})+\frac{1}{2!}(\theta-\hat{\theta})^{T}{\frac{d^{2} f\left(\theta^{\prime}\right)}{d \theta^{\prime 2}}}_{\theta^{\prime}=\hat{\theta}}(\theta-\hat{\theta})+\ldots
$$

## Normal approximation

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

$$
\log p(\theta \mid y)=\log p(\hat{\theta} \mid y)+\frac{1}{2}(\theta-\hat{\theta})^{T}\left[\frac{d^{2}}{d \theta^{2}} \log p\left(\theta^{\prime} \mid y\right)\right]_{\theta^{\prime}=\hat{\theta}}(\theta-\hat{\theta})+\ldots
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p(\theta \mid y) \approx \mathrm{N}\left(\hat{\theta},[I(\hat{\theta})]^{-1}\right)
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where $I(\theta)$ is called observed information

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

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- $I(\hat{\theta})$ 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 $\theta$ 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


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- 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)
```


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- uses finite differences of gradients to compute Hessian


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- second order autodiff in progress


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- In some cases accuracy for a conditional distribution is sufficient (Ch 13)
- e.g. Gaussian latent variable models, such as Gaussian processes (Ch 21) and Gaussian Markov random fields
- Rasmussen \& Williams: Gaussian Processes for Machine Learning
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- Accuracy can be improved by importance sampling (Ch 10)


## Example: Importance sampling in Bioassay





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But the normal approximation is not that good here: Grid $\operatorname{sd}($ LD50 $) \approx 0.1$, Normal sd(LD50) $\approx .75$ !

## Example: Importance sampling in Bioassay



## Example: Importance sampling in Bioassay



Grid $\mathrm{sd}(\mathrm{LD} 50) \approx 0.1, \mathrm{IS} \operatorname{sd}(\mathrm{LD} 50) \approx 0.1$

## Normal approximation

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- in Bioassay example $k=0.57$, which is ok
- CmdStan(R) has Laplace algorithm
- since version 2.33 (2023)
+ Pareto-k diagnostic via posterior package
+ importance resampling (IR) via posterior package


## Normal approximation and parameter transformations

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- for this, Stan does the inference in unconstrained space using logit transformation
- density of the transformed parameter needs to include Jacobian of the transformation (BDA3 p. 21)


## Normal approximation and parameter transformations

Binomial model $y \sim \operatorname{Bin}(\theta, N)$, with data $y=9, N=10$ With $\operatorname{Beta}(1,1)$ prior, the posterior is $\operatorname{Beta}(9+1,1+1)$


## Normal approximation and parameter transformations

With $\operatorname{Beta}(1,1)$ prior, the posterior is $\operatorname{Beta}(9+1,1+1)$
Stan computes only the unnormalized posterior $q(\theta \mid y)$


## Normal approximation and parameter transformations

With $\operatorname{Beta}(1,1)$ prior, the posterior is $\operatorname{Beta}(9+1,1+1)$
For illustration purposes we normalize Stan result $q(\theta \mid y)$


## Normal approximation and parameter transformations

With $\operatorname{Beta}(1,1)$ prior, the posterior is $\operatorname{Beta}(9+1,1+1)$
$\operatorname{Beta}(9+1,1+1)$, but $x$-axis shows the unconstrained $\operatorname{logit}(\theta)$


## Normal approximation and parameter transformations

...but we need to take into account the absolute value of the determinant of the Jacobian of the transformation $\theta(1-\theta)$


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...but we need to take into account Jacobian $\theta(1-\theta)$
Let's compare a wrong normal approximation and correct one


## Normal approximation and parameter transformations

Let's compare a wrong normal approximation and correct one
Sample from both approximations and show KDEs for draws


## Normal approximation and parameter transformations

Let's compare a wrong normal approximation and correct one Inverse transform draws and show KDEs


## Normal approximation and parameter transformations

Laplace approximation can be further improved with importance resampling


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- Split-normal and split-t by Geweke (1989) 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
- CS-E4895-Gaussian Processes
- Stan has the ADVI algorithm (not very good implementation)
- Stan has Pathfinder algorithm (CmdStanR github version)
- instead of normal, methods with flexible flow transformations
- expectation propagation (Ch 13)
- speed of these is usually between optimization and MCMC
- stochastic variational inference can be even slower than MCMC


## Pathfinder: Parallel quasi-Newton variational inference.


estimated ELBO: -340.5

estimated ELBO: -329.6

estimated ELBO: -332.2

estimated ELBO: -329.6

estimated ELBO: -329.7


Zhang, Carpenter, Gelman, and Vehtari (2022). Pathfinder: Parallel quasi-Newton variational inference. Journal of Machine Learning Research, 23(306):1-49.

## Pathfinder: Parallel quasi-Newton variational inference.


estimated ELBO: -4.3



estimated ELBO: -579.9

estimated ELBO: -132.1


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Exact, Normal at mode, Normal with variational inference






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Grid $\operatorname{sd}($ LD50 $) ~ \approx 0.090$,
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Exact, Normal at mode, Normal with variational inference






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

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- with increasing number of posterior dimensions, the stochastic divergence estimate gets worse and flows have problems, too (Dhaka, Catalina, Andersen, Welandawe, Huggins, and Vehtari, 2021)


## Large sample theory

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- eventually likelihood dominates the prior
- the higher order terms in Taylor series increase slower than the second order term
- see counter examples


## Large sample theory

- Assume "true" underlying data distribution $f(y)$
- observations $y_{1}, \ldots, y_{n}$ 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 at it has certain regularity conditions


## Large sample theory

- Consistency
- if true distribution is included in the parametric family, so that $f(y)=p\left(y \mid \theta_{0}\right)$ for some $\theta_{0}$, then posterior converges to a point $\theta_{0}$, when $n \rightarrow \infty$


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- If true distribution is not included in the parametric family, then there is no true $\theta_{0}$
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## 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 $\theta_{0}$ where posterior would converge


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- 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

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\binom{u}{v} \sim N\left(\binom{0}{0},\left(\begin{array}{ll}
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- Problem also for other inference methods like MCMC


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- But a finite data from this data generating process may lack the joint height and weight observations, and thus the the finite data likelihood doesn't have information about $\rho$
- If the likelihood is weakly informative for some parameters, priors and integration are more important


## 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 $y_{t} \sim \mathrm{~N}\left(\theta_{t}, \sigma^{2}\right)$ and $\theta_{t}$ has prior in time
- posterior of $\theta_{t}$ does not converge to a point, if additional observations do not bring enough information


## Large sample theory - counter examples

- Aliasing (valetoisto in Finnish)
- special case of under-identifiability where likelihood repeats in separate points
- e.g. mixture of normals

$$
p\left(y_{i} \mid \mu_{1}, \mu_{2}, \sigma_{1}^{2}, \sigma_{2}^{2}, \lambda\right)=\lambda \mathrm{N}\left(\mu_{1}, \sigma_{1}^{2}\right)+(1-\lambda) \mathrm{N}\left(\mu_{2}, \sigma_{2}^{2}\right)
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if $\left(\mu_{1}, \mu_{2}\right)$ are switched, $\left(\sigma_{1}^{2}, \sigma_{2}^{2}\right)$ are switched and replace $\lambda$ 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


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- asymptotic results assume that probability sums to 1
- e.g. Binomial model, with Beta $(0,0)$ prior and observation $y=n$
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- Should have a positive prior probability/density where needed


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- e.g. $y_{i} \sim \mathrm{~N}(\theta, 1)$ with a restriction $\theta \geq 0$ and assume that $\theta_{0}=0$
- posterior of $\theta$ 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


## Frequency evaluations

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- 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 very bad intervals for all possible values.


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- Maximum likelihood (often) fulfills asymptotic frequency properties
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- Requirement of unbiasedness may lead to higher variance or silly estimates
- unbiased estimate for strictly positive parameter can be negative
- Confidence interval is defined to have true value inside the interval in $\alpha \%$ cases of repeated data generation from the data generating mechanism
- doesn't need be useful to have perfect calibration


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- Bayesian inference
- easier for complex, e.g. hierarchical, models
- easier when model changes
- a consistent way to add prior information
- A lot of machine learning is not pure frequentist or Bayesian, but there is often a probabilistic flavor

