# 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)

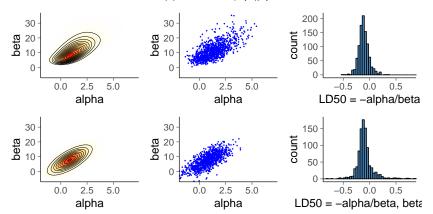
- Often posterior converges to normal distribution when  $n \to \infty$ 
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  - Laplace used this (before Gauss) to approximate the posterior of binomial model to infer ratio of girls and boys born

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

#### Multivariate Taylor series

Multivariate series expansion

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

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

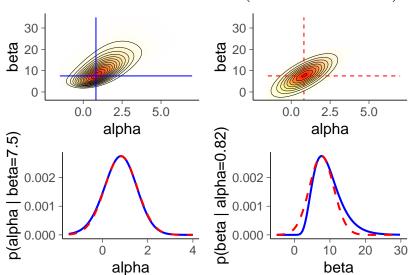
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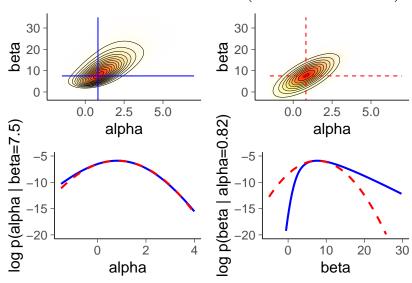
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 $Hessian H(\theta) = -I(\theta)$ 

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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

 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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  - 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, hessi thetahat <- optimres$par Sigma <- solve(optimres$hessian)
```

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  - uses autodiff for gradients
  - uses finite differences of gradients to compute Hessian

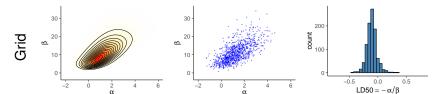
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    - second order autodiff in progress

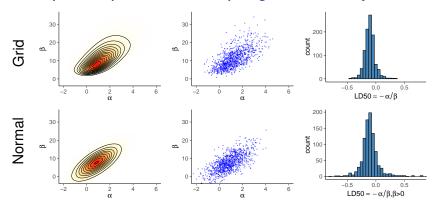
 Optimization and computation of Hessian requires usually much less density evaluations than MCMC

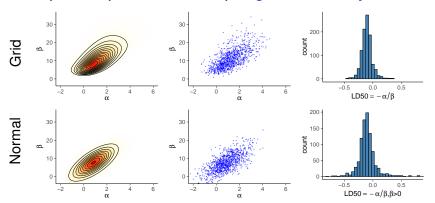
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  - Rasmussen & Williams: Gaussian Processes for Machine Learning
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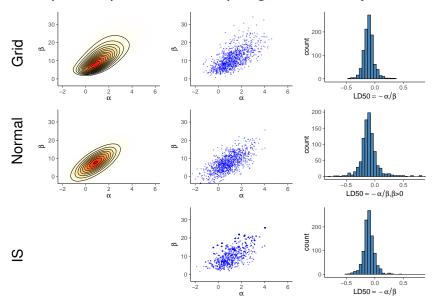
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- Accuracy can be improved by importance sampling (Ch 10)

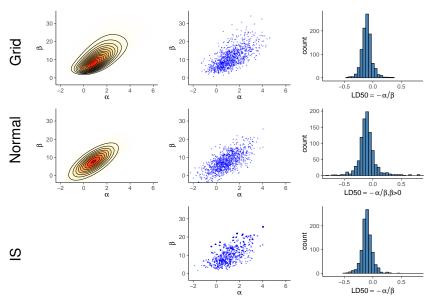






But the normal approximation is not that good here: Grid  $sd(LD50) \approx 0.1$ , Normal  $sd(LD50) \approx .75!$ 





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

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- CmdStan(R) has Laplace algorithm
  - since version 2.33 (2023)
    - + Pareto-k diagnostic via posterior package
    - + importance resampling (IR) via posterior package

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```
real<lower=0, upper=0> theta;
```

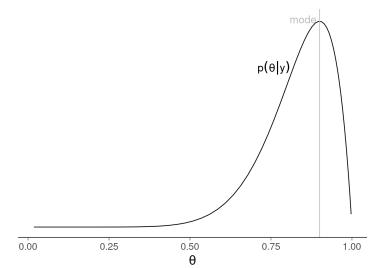
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  - density of the transformed parameter needs to include Jacobian of the transformation (BDA3 p. 21)

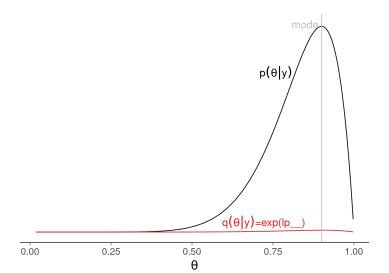
Binomial model  $y \sim Bin(\theta, N)$ , with data y = 9, N = 10

With Beta(1, 1) prior, the posterior is Beta(9 + 1, 1 + 1)



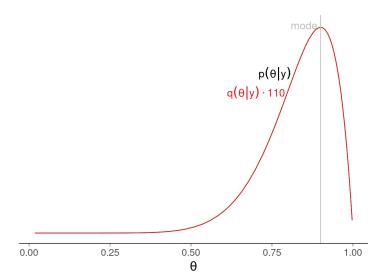
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Stan computes only the unnormalized posterior  $q(\theta|y)$ 



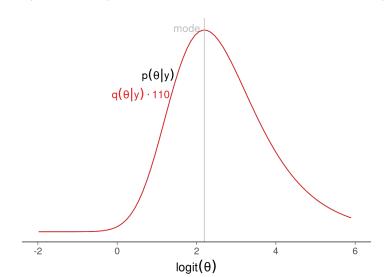
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For illustration purposes we normalize Stan result  $q(\theta|y)$ 

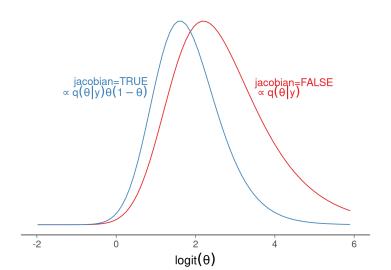


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Beta(9 + 1, 1 + 1), but x-axis shows the unconstrained  $logit(\theta)$ 

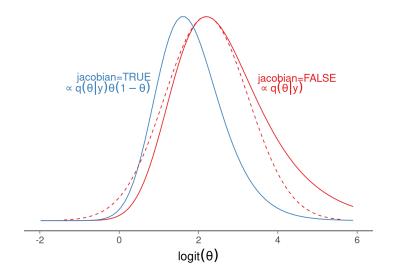


...but we need to take into account the absolute value of the determinant of the Jacobian of the transformation  $\theta(1-\theta)$ 



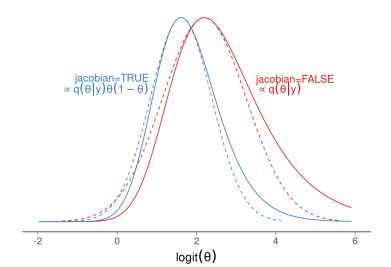
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Let's compare a wrong normal approximation...

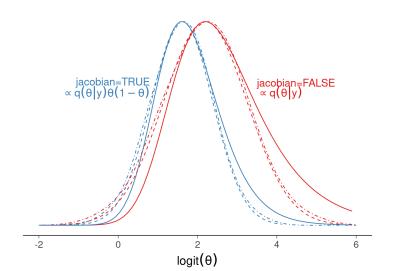


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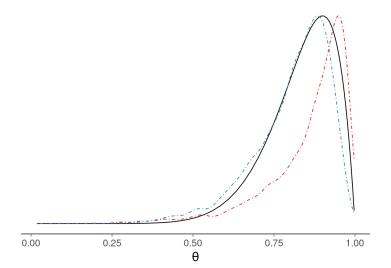
Let's compare a wrong normal approximation and correct one



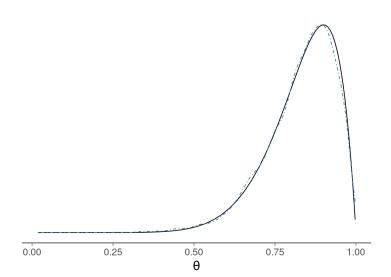
Let's compare a wrong normal approximation and correct one Sample from both approximations and show KDEs for draws



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



Laplace approximation can be further improved with importance resampling



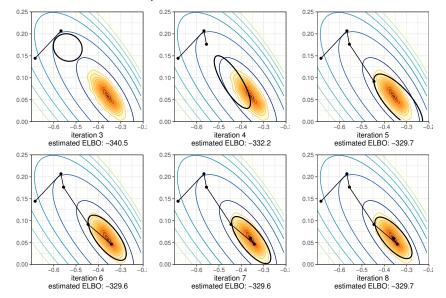
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- Instead of mode and Hessian at mode, e.g.
  - variational inference (Ch 13)
    - CS-E4820 Machine Learning: Advanced Probabilistic Methods
    - CS-E4895 Gaussian Processes
    - Stan has the ADVI algorithm (not very good implementation)
    - Stan has Pathfinder algorithm (CmdStanR, brms)
    - 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

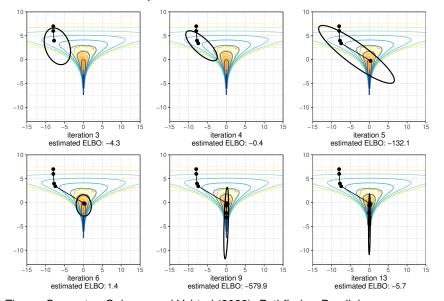
## Pathfinder: Parallel quasi-Newton variational inference.



qZhang Carpenter Gelman and Vehtari (2022) a Pathfinder: Parallel earch, 23(306):1–49.

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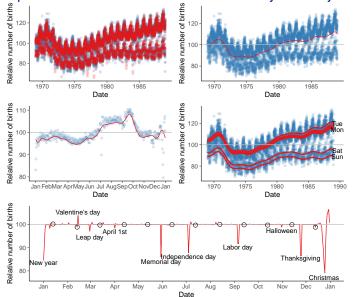
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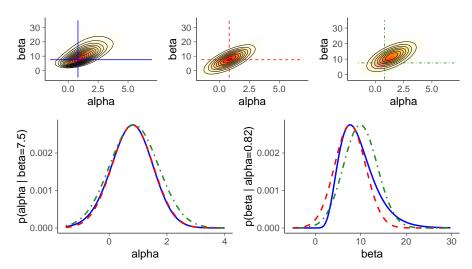
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Birthdays case study uses Pathfinder to speed up workflow https://users.aalto.fi/~ave/casestudies/Birthdays/birthdays.html



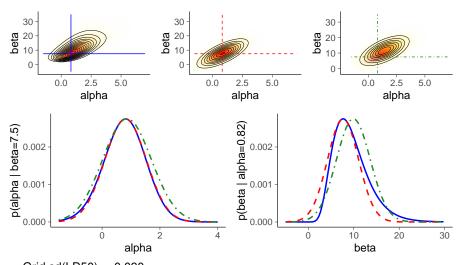
#### Distributional approximations

Exact, Normal at mode, Normal with variational inference



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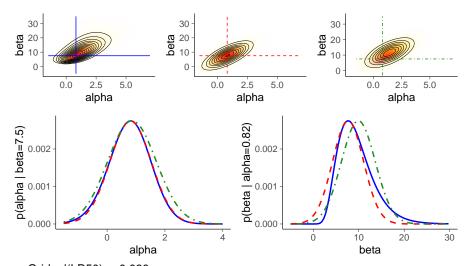
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  - with increasing number of posterior dimensions, the stochastic divergence estimate gets worse and flows have problems, too (Dhaka, Catalina, Andersen, Welandawe, Huggins, and Vehtari, 2021)

#### brms supports Laplace / Pathfinder / ADVI

These might be useful for initializing MCMC or big data. The ADVI implementation is not very good.

```
fit1 <- brm(..., algorithm = "laplace")
fit1 <- brm(..., algorithm = "pathfinder")
fit1 <- brm(..., algorithm = "meanfield")
fit1 <- brm(..., algorithm = "fullrank")</pre>
```