Bayesian statistics with R 5. Markov chains Monte Carlo (MCMC)

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Get posteriors with Markov chains

Monte Carlo (MCMC) methods

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- Difficult, if not impossible to calculate!

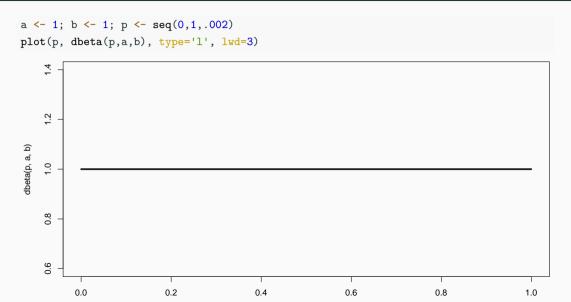
Brute force approach via numerical integration

Deer data

```
y <- 19 # nb of success
n <- 57 # nb of attempts
```

- Likelihood Binomial(57, θ)
- Prior Beta(a=1,b=1)

Beta prior



Apply Bayes theorem

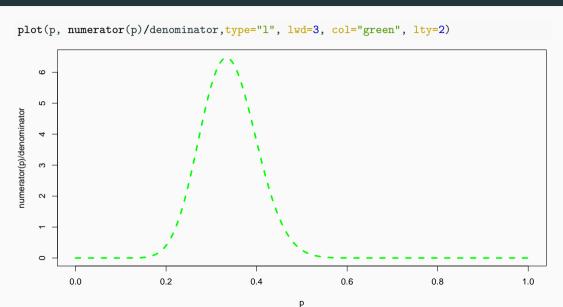
• Likelihood times the prior: $Pr(data \mid \theta) Pr(\theta)$

```
\texttt{numerator} \begin{tabular}{ll} \textbf{<-} & \textbf{function}(\texttt{p}) & \textbf{dbinom}(\texttt{y},\texttt{n},\texttt{p}) * \textbf{dbeta}(\texttt{p},\texttt{a},\texttt{b}) \\ \end{tabular}
```

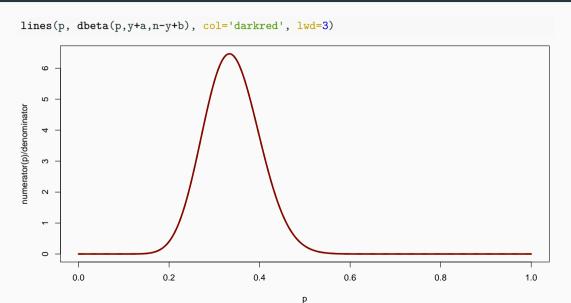
• Averaged likelihood: $Pr(data) = \int L(\theta \mid data) Pr(\theta) d\theta$

```
denominator <- integrate(numerator,0,1)$value</pre>
```

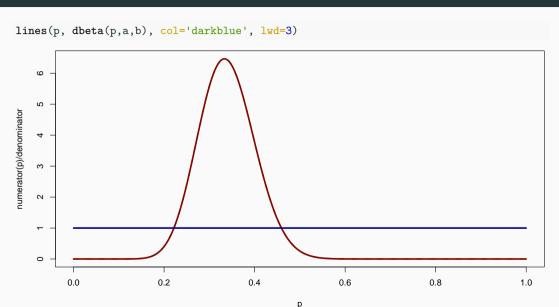
Posterior inference via numerical integration



Superimpose explicit posterior distribution (Beta formula)



And the prior



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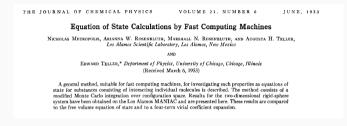
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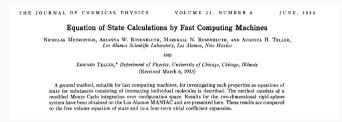
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Do we really wish to calculate a 3D integral?

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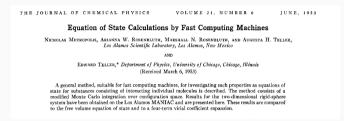


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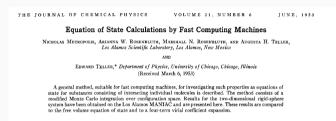
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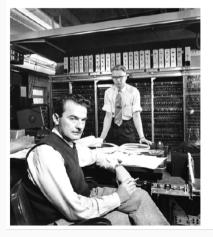
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- Instead, approximate posterior to arbitrary degree of precision by drawing large sample.
- Markov chain Monte Carlo = MCMC; boost to Bayesian statistics!

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MANIAC: Mathematical Analyzer, Numerical Integrator, and Computer



MANIAC: 1000 pounds 5 kilobytes of memory 70k multiplications/sec

Your laptop: 4–7 pounds 2–8 million kilobytes Billions of multiplications/sec

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- How to implement them in practice?!

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- We illustrate sampling from the posterior distribution of winter survival.
- We write functions in R for the likelihood, the prior and the posterior.

```
# deer data, 19 "success" out of 57 "attempts"
survived <- 19
released <- 57
# log-likelihood function
loglikelihood <- function(x, p){</pre>
  dbinom(x = x, size = released, prob = p, log = TRUE)
# prior density
logprior <- function(p){</pre>
  dunif(x = p, min = 0, max = 1, log = TRUE)
# posterior density function (log scale)
posterior <- function(x, p){</pre>
  loglikelihood(x, p) + logprior(p) # - log(Pr(data))
```

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- 5. We repeat 2-4 a number of times called **steps** (many steps).

```
# propose candidate value
move <- function(x, away = .2){
  logitx \leftarrow log(x / (1 - x))
  logit_candidate <- logitx + rnorm(1, 0, away)</pre>
  candidate <- plogis(logit_candidate)</pre>
  return(candidate)
# set up the scene
steps <- 100
theta.post <- rep(NA, steps)
set.seed(1234)
# pick starting value (step 1)
inits <- 0.5
theta.post[1] <- inits</pre>
```

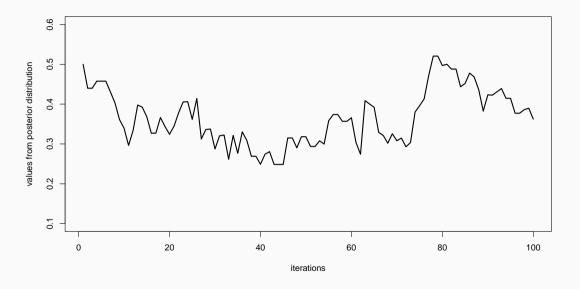
```
for (t in 2:steps){ # repeat steps 2-4 (step 5)
  # propose candidate value for prob of success (step 2)
  theta_star <- move(theta.post[t-1])</pre>
  # calculate ratio R (step 3)
  pstar <- posterior(survived, p = theta_star)</pre>
  pprev <- posterior(survived, p = theta.post[t-1])</pre>
  logR <- pstar - pprev</pre>
  R <- exp(logR)
  # decide to accept candidate value or to keep current value (step 4)
  accept \leftarrow rbinom(1, 1, prob = min(R, 1))
  theta.post[t] <- ifelse(accept == 1, theta star, theta.post[t-1])
```

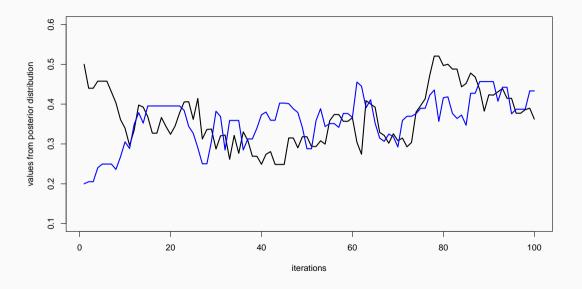
Starting at the value 0.5 and running the algorithm for 100 iterations.

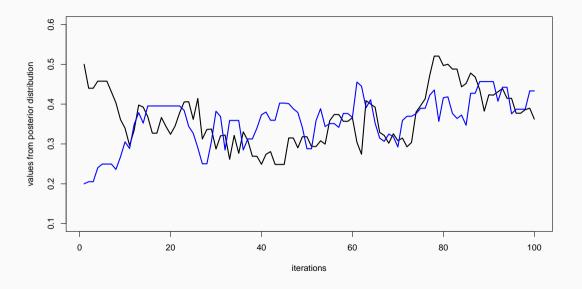
```
head(theta.post)

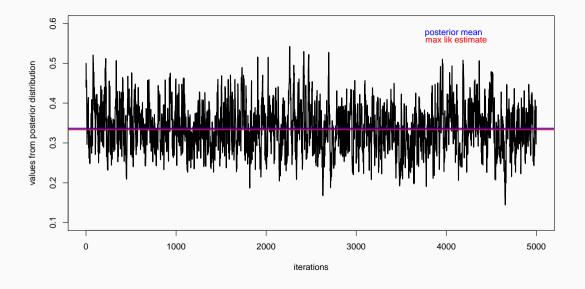
#> [1] 0.5000000 0.4399381 0.4399381 0.4577124 0.4577124 0.4577124
tail(theta.post)

#> [1] 0.4145878 0.3772087 0.3772087 0.3860516 0.3898536 0.3624450
```









Animating the Metropolis algorithm - 1D example

https://gist.github.com/oliviergimenez/5ee33af9c8d947b72a39ed1764040bf3

Animating the Metropolis algorithm - 2D example

https://mbjoseph.github.io/posts/2018-12-25-animating-the-metropolis-algorithm/signal and the state of the

The Markov-chain Monte Carlo Interactive Gallery

https://chi-feng.github.io/mcmc-demo/