

# Yet another intro to MCMC

## Focusing on 2 main methods

Konstantinos Pateras

University of Thessaly, Department of Public and One Health

*kostas.pateras@gmail.com*

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# What to expect?

Go to [www.menti.com](http://www.menti.com)

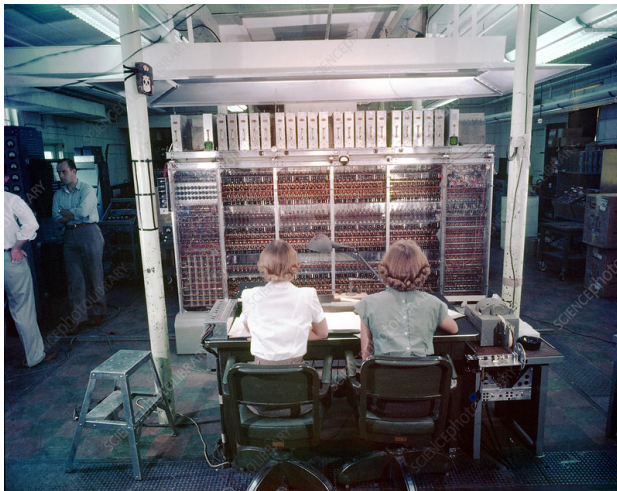
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# Why bother?

# Why bother?

- Monte Carlo: Physical sciences (Astrophysics), Engineering (Thermodynamics), Climate change, Computational biology (Phylogenetics), Applied statistics (Bayesian approach), Artificial intelligence, among many
- MCMC: Computational statistics, computational physics and computational biology among others. Hierarchical modelling in Bayesian statistics

# MANIAC I





# Charles III of Monaco



# Historical notes

## Prologue

Questions

History

Inv-CDF

## Warming up

Accept/Reject  
Markov Chains

## The 1st algorithm

Metropolis-  
Hastings  
Step-by-step  
implementation

## The 2nd algorithm

Gibbs sampler  
Step-by-step  
implementation

## JAGS and further reading

## Epilogue

- Monte carlo methods since 1920 (Enrico Fermi)
- Named after Nicholas Metropolis, Monte Carlo casino.
- Los Alamos labs 1940s (Stanislaw Ulam, John von Neumann, Nicholas Metropolis and others)
- Metropolis-Hastings 1970, Geman Geman Gibbs (1984)
- Mainstream in stats by Gelman and Rubin 1992 among others
- Mainstream software (BUGS 1996 (Gibbs), JAGS (Slice), Stan (Hamiltonian))



# Basic notions

- 1 How to generate completely random numbers (not easy). R uses pseudo-random number generators.
- 2 Main assumption: Be able to generate from a  $\text{Uniform}(0,1)$
- 3 Inverse transform method, inv-CDF,, (Rizzo pg 49)

## Acceptance-Rejection method

## Prologue

Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithm

Gibbs sampler

Step-by-step  
implementationJAGS and  
further  
reading

## Epilogue

To generate a random variable  $Y$ , suppose  $Y$  and  $X$  are random variables with density  $f$  and  $g$ . Also, there exists a constant  $c$  such that  $f(t)/g(t) \leq c, f(t) > 0$

## Acceptance-Rejection method

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Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithm

Gibbs sampler

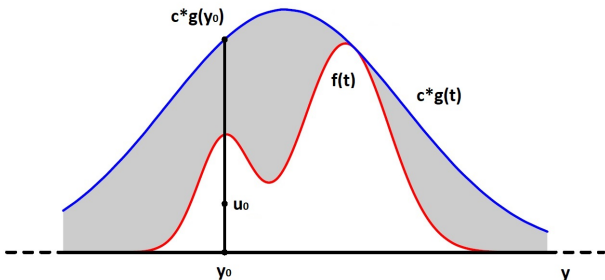
Step-by-step  
implementationJAGS and  
further  
reading

## Epilogue

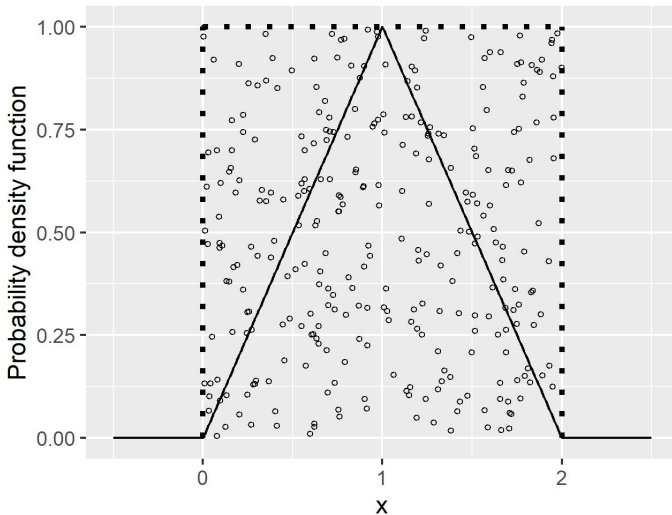
To generate a random variable  $Y$ , suppose  $Y$  and  $X$  are random variables with density  $f$  and  $g$ . Also, there exists a constant  $c$  such that  $f(t)/g(t) \leq c, f(t) > 0$

- 1 Find and generate  $Y$  from  $g$  for which
  - i  $f(t) \leq c \cdot g(t), f(t) > 0$ 
    - i Generate a random  $y^* \sim g$
    - ii Generate a random  $u \sim Uniform(0, 1)$
    - iii If  $u < f(y^*)/(c \cdot g(y^*))$  accept  $y^*$  as sample, otherwise reject  $y^*$ .
    - iv Repeat from step i.

# Acceptance-Rejection figure



# Acceptance-Rejection (quiz)



# Acceptance-Rejection method

## Example (Triangle probability density)

```
triangle.pdf = function(x){ # Definition of triangle pdf.
  ifelse((0 < x) & (x < 1), x,
  ifelse((1 <= x) & (x < 2), 2 - x, 0))
}

#simulates a sample of size (n) from the pdf f(x) via the acceptance/rejection algorithm
accept_reject = function(fx, n = 100, prop.val=1) {

  x = numeric(n);count = 0

  while(count < n) {
    temp.x <- runif(1, 0, 2)
    y <- runif(1, 0, 1)

    if (y < fx(temp.x)) {
      count = count + 1
      x[count] <- temp.x
    }
  }

  return(x)
}

sample = accept_reject(triangle.pdf, 50000)
length(sample)/50000 # Probability of acceptance
plot(density(sample), adjust=0.5)
```

Prologue

Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithmGibbs sampler  
Step-by-step  
implementationJAGS and  
further  
reading

Epilogue

# Markov Chains

## Some brief notions

- First order Markov Chains (Depend on last value)
- Irreducibility (Any state can reach another state)
- Aperiodicity
- (Positive) recurrent
- Stationary distribution

# Markov Chains Concept

## Prologue

Questions

History

Inv-CDF

## Warming up

Accept/Reject

Markov Chains

## The 1st algorithm

Metropolis-  
HastingsStep-by-step  
implementation

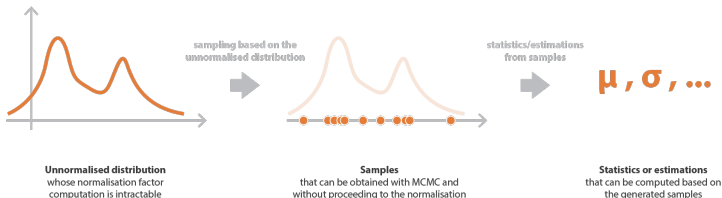
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## JAGS and further reading

## Epilogue



Source: Joseph Rocca, Bayesian inference problem, MCMC and variational inference



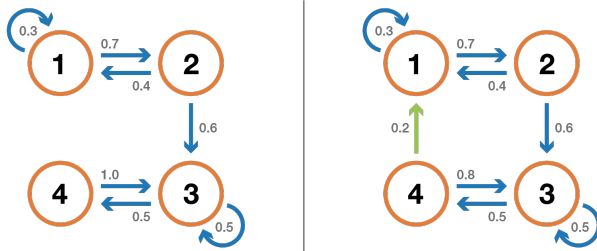
# Markov Chains Concept 2



Source: Joseph Rocca, Bayesian inference problem, MCMC and variational inference

# Markov Chains Assumption 1

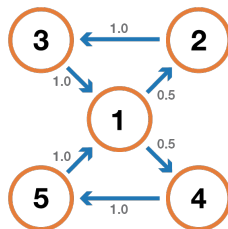
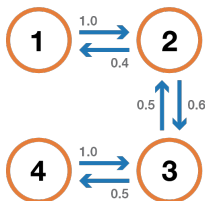
Illustration of the irreducibility property. The chain on the left is not irreducible: from 3 or 4 we can't reach 1 or 2. The chain on the right (one edge has been added) is irreducible: each state can be reached from any other state.



Source: Joseph Rocca, Bayesian inference problem, MCMC and variational inference

## Markov Chains Assumption 2

Illustration of the aperiodicity property. The chain on the left is 2-periodic: when leaving any state, it always takes a multiple of 2 steps to come back to it. The chain on the right is 3-periodic.



Source: Joseph Rocca, Bayesian inference problem, MCMC and variational inference

# Metropolis background

Metropolis with symmetric proposal (Random-walk metropolis) - If transition is reversible then target is invariant/stationary.

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- 1 Choose a proposal distribution

$q(\theta|\theta^*) = q(\theta^*|\theta) = N(\theta, s_\theta)$  [Irreducible, positive recurrent, aperiodic] and set initial  $\theta = \theta_0$

- 2 Repeat until chain has converged

## Metropolis background

## Prologue

Questions

History

Inv-CDF

Warming  
up

Accept/Reject

Markov Chains

The 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithm

Gibbs sampler

Step-by-step  
implementationJAGS and  
further  
reading

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- 2 Repeat until chain has converged
  - 2a Generate  $\theta^*$  from  $N(\theta, s_\theta)$ , what about  $s_\theta$ ?
  - 2b Generate  $U$  from  $\text{Uniform}(0,1)$

## Metropolis background

## Prologue

Questions

History

Inv-CDF

Warming  
up

Accept/Reject

Markov Chains

The 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithm

Gibbs sampler

Step-by-step  
implementationJAGS and  
further  
reading

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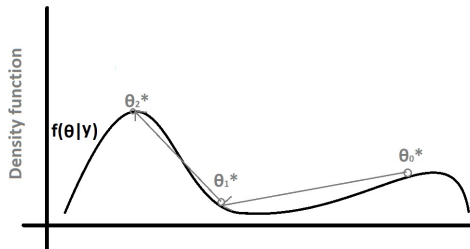
2a Generate  $\theta^*$  from  $N(\theta, s_\theta)$ , what about  $s_\theta$ ?

2b Generate  $U$  from Uniform(0,1)

2c If  $U \leq \frac{f(\theta^*|y)}{f(\theta|y)} = \frac{f(y|\theta^*)p(\theta^*)}{f(y|\theta)p(\theta)}$ , accept  $\theta^*$  and set  $\theta = \theta^*$ , otherwise set  $\theta = \theta$

2d Increment  $t$  and repeat

## Metropolis figure



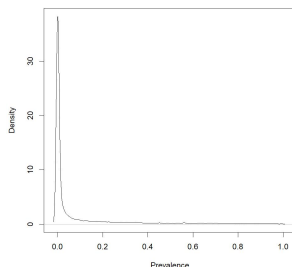


# Metropolis algorithm

- Based on expert beliefs the prior probability of an adverse effect can be summarized via a Beta distribution opinion  $\theta \sim \text{Beta}(\alpha = 0.1, \beta = 1)$ . Prior beliefs can be expressed as

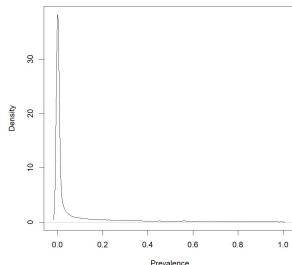
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- Toy example: Prevalence of moderate adverse events of Covid-19 vaccine. Observed 50 events in a sample of 10,000,  $y \sim B(50, 10^4)$ .

# Metropolis algorithm

## Prologue

Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithmGibbs sampler  
Step-by-step  
implementationJAGS and  
further  
reading

## Epilogue

## Example (Theorem Slide Code)

```
metropolis <- function(burnin = 0, sample = 10000,  
sigma = 0.05, initial_value = 0.05, plot=TRUE)  
  
  stopifnot(initial_value > 0, initial_value < 1)  
  stopifnot(sigma > 0)  
  burnin <- as.integer(burnin)  
  sample <- as.integer(sample)  
  stopifnot(burnin >= 0)  
  stopifnot(sample > 0)  
  
  # Redefine these to work on the log scale:  
  llikelihood_fun <- function(prevalence) dbinom(50, 10^4, prevalence, log=TRUE)  
  lprior_fun <- function(prevalence) dbeta(prevalence, 0.1, 1, log=TRUE)  
  
  parameters <- numeric(burnin+sample)  
  parameters[1] <- initial_value  
  current <- initial_value  
  
  post <- llikelihood_fun(current) + lprior_fun(current)
```

# Metropolis algorithm

## Example (Theorem Slide Code)

```
for(i in 2:(burnin+sample)){  
  
  proposal <- rnorm(1, current, sigma)  
  
  if(proposal > 0 && proposal < 1){  
    U=log(runif(1,0,1))  
    newpost <- llikelihood_fun(proposal) + lprior_fun(proposal)  
    accept <- U <= newpost-post  
  
    if(accept){  
      current <- proposal  
      post <- newpost  
    }  
    parameters[i] <- current  
  }  
}
```

## Metropolis results

## Prologue

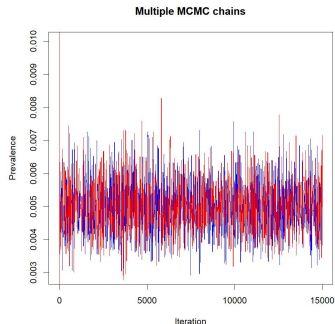
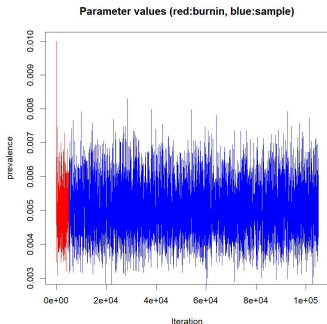
Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithmGibbs sampler  
Step-by-step  
implementationJAGS and  
further  
reading

## Epilogue



- ESS:  $10000/647.67 = 15.43$  iterations for an independent sample
- Geweke diagnostic: p-value=0.67
- Autocorrelation? How to fix this?

# Gibbs sampler background

Named after JW Gibbs, invented 80 years after his death.  
Why Gibbs? When it should be preferred?

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Why Gibbs? When it should be preferred?

- 1 Set initial  $\theta_0$  at time  $t = 0$
- 2 For each iteration  $t = 1, 2, \dots T$  repeat
  - 2a Set  $\theta = \theta^{(t-1)}$



## Gibbs sampler background

## Prologue

Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
Hastings  
Step-by-step  
implementationThe 2nd  
algorithmGibbs sampler  
Step-by-step  
implementationJAGS and  
further  
reading

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- 2 For each iteration  $t = 1, 2, \dots, T$  repeat
  - 2a Set  $\theta = \theta^{(t-1)}$
  - 2b For each parameter  $j = 1, 2, \dots, d$ 
    - 2bi Generate  $\theta_j$  from  $\theta_j \sim f(\theta_j | \theta_{-/j}, y)$

## Gibbs sampler background

## Prologue

Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
Hastings  
Step-by-step  
implementationThe 2nd  
algorithmGibbs sampler  
Step-by-step  
implementationJAGS and  
further  
reading

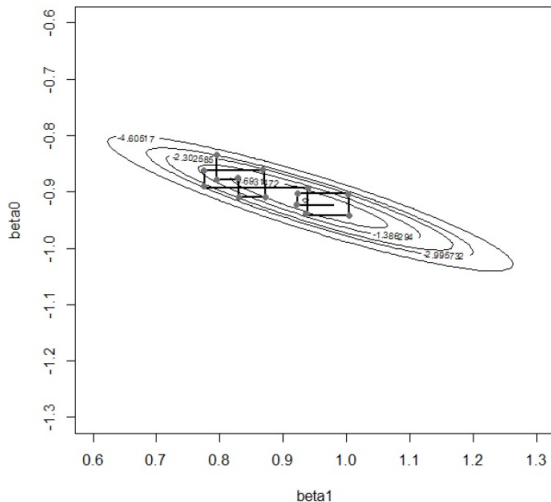
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    - 2bi Generate  $\theta_j$  from  $\theta_j \sim f(\theta_j | \theta_{-/j}, y)$
  - 2c Increase  $t$  and repeat

One component is updated at a time. Conditionally conjugate models and others.

# Gibbs sampler figure



## Gibbs sampler

## Prologue

Questions

History

Inv-CDF

Warming  
up

Accept/Reject

Markov Chains

The 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithm

Gibbs sampler

Step-by-step  
implementationJAGS and  
further  
reading

## Epilogue

- 1 Example with a normal variable indicating average temperature for July in a time-span of 12 years. (Ntzoufras 2009)

- $\mu \sim N(\mu_0, \sigma_0^2)$  and  $\sigma^2 \sim IG(a_0, b_0)$
- Compute conditional distributions  $f(\mu|\sigma^2, y)$  and  $f(\sigma^2|\mu, y)$
- $\mu|\sigma^2, y \sim N(w\tilde{y} + (1 - w)\mu_0, w\frac{\sigma^2}{n})$ ,  $w = \frac{\sigma_0^2}{\sigma^2/n + \sigma_0^2}$   
 $\sigma|\mu, y \sim IG[(a_0) + n/2, b_0 + 1/2 \sum_1^n (y_i - \mu)^2]$

## Prologue

Questions

History

Inv-CDF

Warming  
up

Accept/Reject

Markov Chains

The 1st  
algorithmMetropolis-  
HastingsStep-by-step  
implementationThe 2nd  
algorithm

Gibbs sampler

Step-by-step  
implementationJAGS and  
further  
reading

## Epilogue

## Example (Gibbs with non-conjugate priors)

```
y<-c(32,36,37,34,38,36,33,36,37,35,32,35)
bary<-mean(y); n<-length(y)  # Set up data
```

```
# Set number of iterations
Iterations<-5000
```

```
# Set prior parameters.
mu0<-0; s0<-100; a0<-0.001; b0<-0.001
```

```
# Initialize vectors of sample values mu and sigma
theta <- matrix(nrow=Iterations, ncol=2)
```

```
# Set initial current mu(0) and sigma(0)
cur.mu<-0; cur.tau<-1; cur.s<-sqrt(1/cur.tau)
```

## Gibbs sampler

## Example (Gibbs with non-conjugate priors)

```
for (t in 1:Iterations){ # Repeat for t=1,...,T iterations

  w<- s0^2/( cur.s^2/n+ s0^2) # Calculate w

  m <- w*bary + (1-w)*mu0 # Calculate m
  s <- sqrt( w * cur.s^2/n) # Calculate s

  cur.mu <- rnorm(1, m, s) # Calculate mu

  a <- a0 + 0.5*n # Calculate alpha
  b <- b0 + 0.5 * sum( (y-cur.mu)^2) # Calculate beta

  cur.tau <- rgamma( 1, a, b) # Calculate tau from gamma
  cur.s <- sqrt(1/cur.tau) # Calculate sigma (inversed tau)

  theta[t,]<-c( cur.mu, cur.s) # Set mu(t)=mu, sigma(t)=sigma
} # End of for loop
```

Prologue

Questions

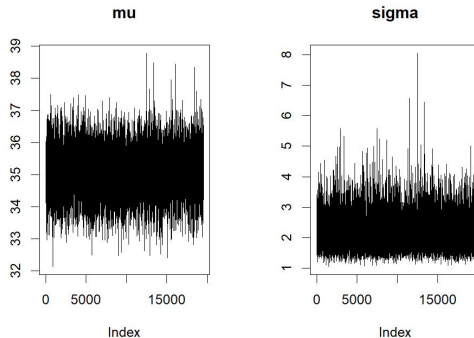
History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
Hastings  
Step-by-step  
implementationThe 2nd  
algorithmGibbs sampler  
Step-by-step  
implementationJAGS and  
further  
reading

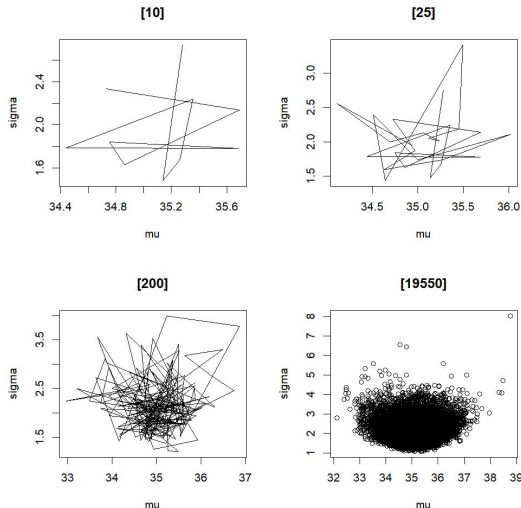
Epilogue

## Gibbs results [univariate]



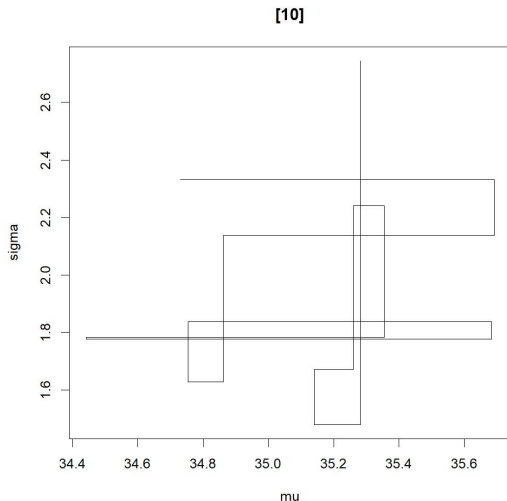
- ESS:  $20000/18757.47 = 1.07$  iterations for an independent sample of  $\mu$
- Geweke diagnostic: p-value = 0.49

# Gibbs results [both simultaneously]





## Gibbs results [both step by step]



JAGS/WinBUGS code -  
Examples

## Prologue

Questions

History

Inv-CDF

Warming  
upAccept/Reject  
Markov ChainsThe 1st  
algorithmMetropolis-  
Hastings  
Step-by-step  
implementationThe 2nd  
algorithmGibbs sampler  
Step-by-step  
implementationJAGS and  
further  
reading

## Epilogue

Metropolis example with  $y = 50$  and  $n = 10^4$

```
"model{
y ~ dbinom(main.ap, n)
main.ap ~ dbeta(0.1,1)
}"
```

Gibbs example with  $y$  temperatures

```
"model{
for (i in 1:12){
y[i] ~ dnorm (mu, prec)
}
mu ~ dnorm (0, 0.01)
prec ~ dgamma (0.001, 0.001)
s2 <- 1/prec
}
}"
```

## Prologue

Questions

History

Inv-CDF

## Warming up

Accept/Reject  
Markov Chains

## The 1st algorithm

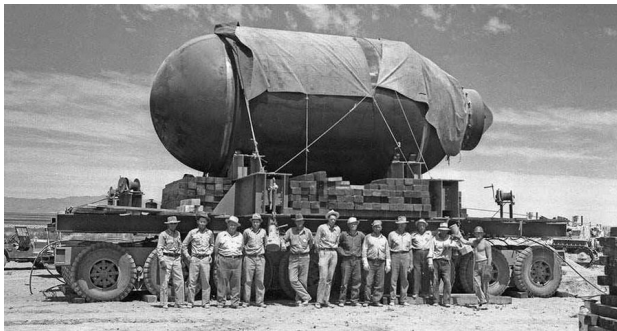
Metropolis-  
Hastings  
Step-by-step  
implementation

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Gibbs sampler  
Step-by-step  
implementation

## JAGS and further reading

## Epilogue



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# Bibliography

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