ADIM: Bayesian GLM. MCMC methods

Master's Degree in Data Analysis, Process Improvement and Decision Support Engineering

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Heart Disease

- The study examines the relationship between:
- \circ the **myocardial infarction** (MI): y=1 if MI occurrence, or y=0 if No MI occurrence; and
- \circ **Age60**: Patients aged \geq 60 (1) versus < 60 (0).
- \circ Systolic blood pressure (SBP140): SBP ≥ 140 mmHg (1) versus < 140 mmHg (0).

Objective:

- Evaluate the association of age60 and sbp140 with MI probability.
- Interpret the odds ratio (OR) for both predictors.

Table: Summary of Data from Study

sbp140	>=140	<140	>=140	>=140	<140	<140
age60	09>	09=<	09>	09=<	09=<	09>
>	0	0	0	0	0	<u></u>

- Logistic regression is used to model MI's probability based on age60 and sbp140.
- Likelihood

$$y_i \sim \mathrm{Bernoulli}(\pi_i)\,, i=1,\ldots,400\,,$$

using logit link:

$$logit(\pi_i) = eta_0 + eta_1 age 60_i + eta_2 sbp 140_i$$

Prior distributions (weakly-informative):

$$\beta_0 \sim \mathcal{N}(0, 10^3), \; \beta_1 \; \sim \mathcal{N}(0, 10^3), \; \beta_2 \; \sim \mathcal{N}(0, 10^3),$$

Note: There are no conjugate priors available for the logistic regression model.

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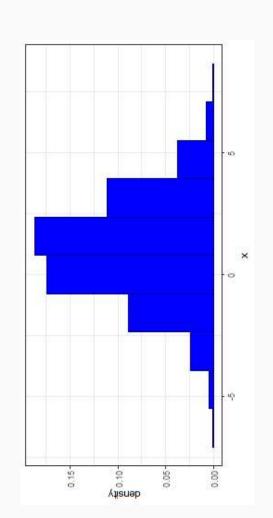
- 1. Bayesian computation. MCMC methods
- 2. Bayesian Software for MCMC



Monte Carlo Simulation

 Draw realizations of a random variable for which only its density function is (fully or partially) known.

$$x \leftarrow rnorm(1000, mean = 1, sd = 2)$$



Monte Carlo Integration

Computing the mean of a N(1, 2),

$$\circ \; E(X) = \int_{-\infty}^{\infty} x \cdot f(x) dx$$

- Using Monte Carlo integration:
- \circ Simulate from N(1, 2^2): ϕ^1, \ldots, ϕ^N .
- \circ Compute the mean of the simulated values: $E(X)pprox rac{1}{N}\sum_{i=1}^N\phi^i$
 - Doing summary of the simulation, we compute more measures:

Markov Chain Monte Carlo

- A Markov chain is a stochastic sequence of numbers where each value in the sequence depends only upon the last.
- If $\phi^1,\phi^2,\ldots,\phi^N$ is a sequence of numbers, then ϕ^2 is only a function of ϕ^1,ϕ^3 of ϕ^2 , etc.
- Under certain conditions, the distribution over the states of the Markov chain will converge to a stationary distribution.
- The stationary distribution is independent of the initial starting values specified for the
- AIM: construct a Markov chain such that the stationary distribution is equal to the posterior distribution $p(\theta \mid x)$.
- We combine Markov Chain with Monte Carlo simulation --> Markov chain Monte Carlo (MCMC).
- They were proposed by first time in the Statistics area by Gelfand and Smith (1990).

Posterior distribution

Estimating the probability to score a penalty

- Likelihood

$$p(oldsymbol{y}\mid\pi)=\pi^k(1-\pi)^{N-k}$$

- Prior distribution

$$p(\pi) = \pi^{a-1}(1-\pi)^{b-1}$$

- Posterior distribution

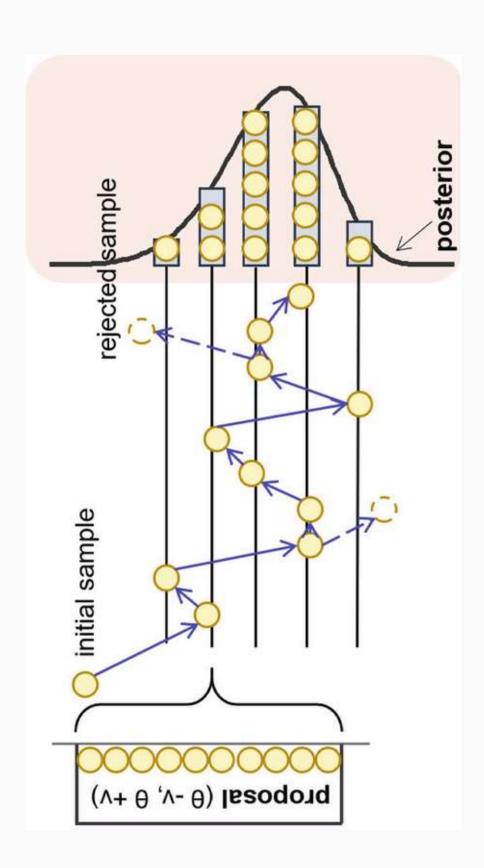
$$p(\pi \mid oldsymbol{y}) \propto p(oldsymbol{y} \mid \pi) imes p(\pi) \propto \pi^{k+a-1} (1-\pi)^{N-k+b-1} = p^*(\pi)$$

- 1. Starting value $\pi^{(0)}$
- 2. For $t=1,\ldots,T$
- We define a proposal distribution (Usually similar to the objective distribution). In this case, $q(\pi \mid \pi^{(t-1)}) \sim logit - N(\pi^{(t-1)}, \sigma = 0.5)$. Simulate $\pi^{(prop)}$ from it.
- Compute probability of acceptance:

$$lpha = \min \left(1, rac{p^*(\pi^{(prop)})q(\pi^{(t-1)} \mid \pi^{(prop)})}{p^*(\pi^{(t-1)})q(\pi^{(prop)} \mid \pi^{(t-1)})}
ight)$$

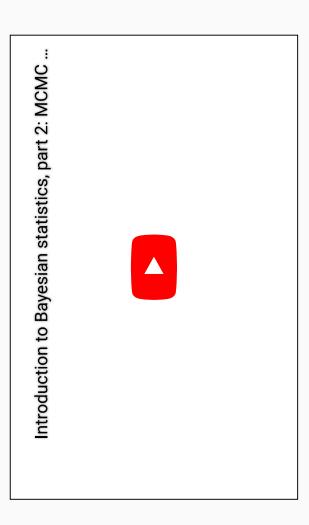
- \circ Generate a **random number** u from the Uniform(0, 1).
- ullet $\pi^{(t+1)}=\pi^{(prop)}$, if $u\geq lpha$,
- ullet $\pi^{(t+1)}=\pi^{(t)}$, if u<lpha
- 3. Finally, we obtain $\pi^0, \pi^1, \ldots, \pi^T$ which is a simulation of the posterior distribution.

MCMC: Metropolis-Hastings (MH)



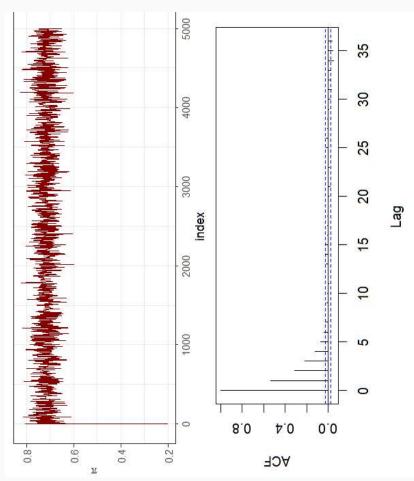
Approaching probability of score using MH

Visual Metropolis-Hastings

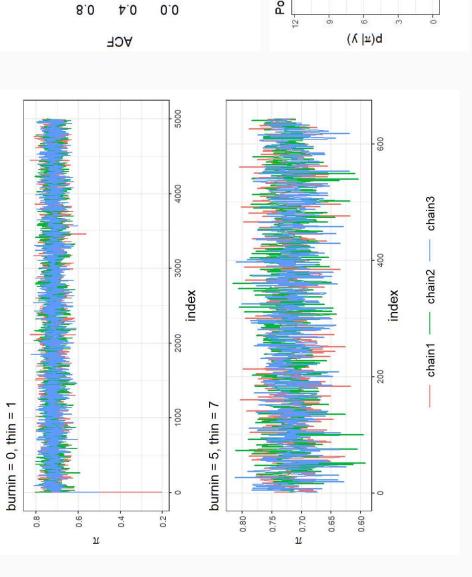


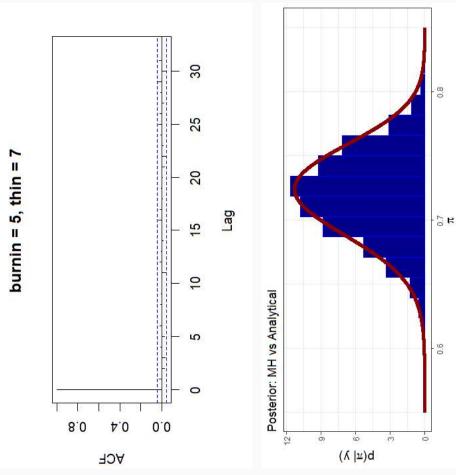
Play the video from minute 4:44.

Tracing the chain. Is the chain autocorrelated?



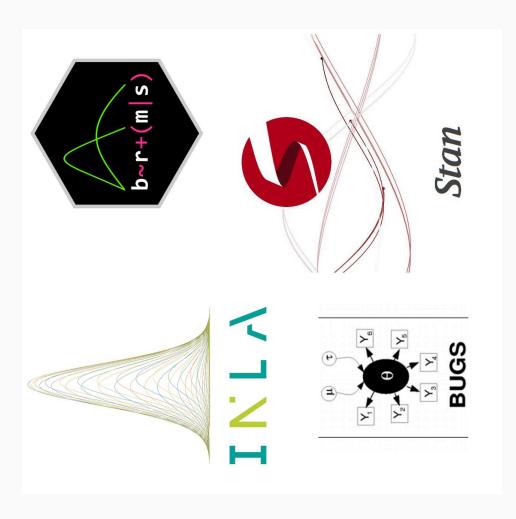
MCMC. Burnin and thin







- JAGS
- Stan
- brms: allow for easy Bayesian
 Inference using Hamiltonian Monte Carlo
- INLA (it does not use MCMC methods, but it is a very powerful tool)
- inlabru : facilitate Bayesian Inference in Spatio-temporal models
- MCMC pack (With this R-package, you can use MCMC methods with similar notation as usually use in R)
- Nimble



Bayesian Logistic Regression using JAGS

Likelihood

$$y_i \sim \mathrm{Bernoulli}(\pi_i) \,, i = 1, \dots, 400 \,,$$

using logit link:

$$logit(\pi_i) = eta_0 + eta_1 age 60_i + eta_2 sbp 140_i$$

Prior distributions (weakly-

informative):

$$eta_0 \sim \mathcal{N}(0, 10^3),$$

$$eta_1 \sim \mathcal{N}(0, 10^3),$$

$$eta_2 \sim \mathcal{N}(0, 10^3).$$

```
model_string ← "
model {
    for (i in 1:N) {
        y[i] ~ dbern(pi[i])
        logit(pi[i]) ← beta0 +
        beta1 * age60[i] +
        beta2 * sbp140[i]
    }

# Priors for regression coefficients
beta0 ~ dnorm(0, 0.001)
beta1 ~ dnorm(0, 0.001)
beta2 ~ dnorm(0, 0.001)
}
beta2 ~ dnorm(0, 0.001)
}.
```

Check S1-JAGS-heart_attack.Rmd for the complete solution

Bayesian Logistic Regression using brms

Likelihood

$$y_i \sim \mathrm{Bernoulli}(\pi_i)\,, i=1,\dots,400\,,$$

using logit link:

$$\operatorname{logit}(\pi_i) = \beta_0 + \beta_1 \operatorname{age60}_i + \beta_2 \operatorname{sbp140}_i$$

Prior distributions (weakly-

informative):

$$eta_0 \sim \mathcal{N}(0, 10^3),$$
 $eta_1 \sim \mathcal{N}(0, 10^3),$ $eta_2 \sim \mathcal{N}(0, 10^3).$

```
# Total number of
formula <- bf(y ~ age60 + sbp140, fami
                                                                                                                                                                                                # Number of itera
                                                                                                                                                                                                                       # Thinning interv
                                                                                                                                                                                                                                                Seed for reprod
                                                                                                                                             # Number of MCMC
                                                                                                                                                                                                                                                                                                                   # Summary of the fitted model
                                          # Fit the model using brms
                                                                  fit_brms ← brm(formula,
                                                                                            data = data_hattack,
                                                                                                                     prior = priors,
                                                                                                                                                                                                warmup = 1000,
                                                                                                                                                                     iter = 5000,
                                                                                                                                             chains = 3,
                                                                                                                                                                                                                                                seed = 123,
                                                                                                                                                                                                                        thin = 1,
                                                                                                                                                                                                                                                                                                                                          summary(fit)
```

Check S1-brms-heart_attack.Rmd for the complete solution

What we have learned so far

- The biggest challenge for Bayesian inference has always been the computational power that more complex models require.
- MCMC methods allow computationally estimating posterior distributions that are analytically intractable.
- computational techniques for computing a posteriori distributions. Today, there are many, many teams of researchers developing

References

Books

- Stan & R: Lambert, B. (2018). A Student's Guide to Bayesian Statistics. SAGE Publications.
- JAGS: Plummer, M. (2019). JAGS User Manual Version 4.3.0.
- OpenBUGS: Cowles, M. K. (2013). Applied Bayesian statistics: with R and OpenBUGS examples (Vol. 98). Springer Science & Business Media.
- WinBUGS: Ntzoufras, I. (2011). Bayesian modeling using WinBUGS. John Wiley & Sons.
- Stan: Andrew Gelman, John B. Carlin, Hal S. Stern, David B. Dunson, Aki Vehtari, Donald B. Rubin (2013). Bayesian Data Analysis. Chapman and Hall/CRC
- INLA: Gómez-Rubio, V. (2020). Bayesian inference with INLA. CRC Press.

References

Blogs

- http://wlm.userweb.mwn.de/R/wlmRcoda.htm
- https://rpubs.com/FJRubio/IntroMCMC
- https://darrenjw.wordpress.com/tag/mcmc/
- https://www.tweag.io/blog/2019-10-25-mcmc-intro1/

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