Bayesian statistics with RBayesian analyses in R with the Jags software

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Bayes in practice

Software implementation (R compatible)

Oldies but goodies:

- WinBUGS, OpenBUGS: Where it all began.
- Jags: What we will use in this course.

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The new kids on the block:

- Nimble: What I'm going for these days.
- Stan: Entirely different algorithmic approach.
- Greta: Dunno anything about it.

Introduction to JAGS (Just Another Gibbs Sampler)

Martyn Plummer



Real example

Impact of climatic conditions on white stork breeding success



Assess effects of temperature and rainfall on productivity.

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- We need to build a model write down the likelihood.
- We need to specify priors for parameters.

Read in the data

```
nbchicks \leftarrow c(151,105,73,107,113,87,77,108,118,122,112,120,122,89,69,71,
              53.41.53.31.35.14.18)
nbpairs <- c(173,164,103,113,122,112,98,121,132,136,133,137,145,117,90,80,
            67,54,58,39,42,23,23)
```

```
temp \leftarrow c(15.1, 13.3, 15.3, 13.3, 14.6, 15.6, 13.1, 13.1, 15.0, 11.7, 15.3, 14.4, 14.4
          12.7.11.7.11.9,15.9,13.4,14.0,13.9,12.9,15.1,13.0)
```

```
rain < c(67,52,88,61,32,36,72,43,92,32,86,28,57,55,66,26,28,96,48,90,86,
           78,87)
```

datax <- list(N = 23, nbchicks = nbchicks, nbpairs = nbpairs,</pre>

tomp = (tomp - mosp(tomp))/ad(tomp)

Write down the model

[likelihood]	$nbchicks_i \sim Binomial(nbpairs_i, p_i)$
[linear model]	$logit(p_i) = a + b_{temp} \; temp_i + b_{rain} \; rain_i$
[prior for a]	$a \sim Normal(0, 1000)$
[prior for b_{temp}]	$b_{temp} \sim Normal(0, 1000)$
[prior for b_{rain}]	$b_{rain} \sim Normal(0, 1000)$

Build the model

```
{
# Likelihood
    for( i in 1 : N){
        nbchicks[i] ~ dbin(p[i],nbpairs[i])
        logit(p[i]) <- a + b.temp * temp[i] + b.rain * rain[i]
      }
# ...</pre>
```

Specify priors

```
# Priors
a ~ dnorm(0,0.001)
b.temp ~ dnorm(0,0.001)
b.rain ~ dnorm(0,0.001)
}
```

Warning: Jags uses precision for Normal distributions (1 / variance)

You need to write everything in a file

```
model <-
paste("
model
    for( i in 1 : N)
        nbchicks[i] ~ dbin(p[i],nbpairs[i])
        logit(p[i]) <- a + b.temp * temp[i] + b.rain * rain[i]</pre>
a \sim dnorm(0, 0.001)
b.temp \sim dnorm(0,0.001)
b.rain \sim dnorm(0.0.001)
```

Alternatively, you may write a R function

```
logistic <- function() {</pre>
    for( i in 1 : N)
        nbchicks[i] ~ dbin(p[i],nbpairs[i])
        logit(p[i]) <- a + b.temp * temp[i] + b.rain * rain[i]</pre>
# priors for regression parameters
a \sim dnorm(0,0.001)
b.temp ~ dnorm(0,0.001)
b.rain \sim dnorm(0, 0.001)
```

Let us specify a few additional things

```
# list of lists of initial values (one for each MCMC chain)
init1 \leftarrow list(a = -0.5, b.temp = -0.5, b.rain = -0.5)
init2 <- list(a = 0.5, b.temp = 0.5, b.rain = 0.5)
inits <- list(init1.init2)</pre>
# specify parameters that need to be estimated
parameters <- c("a","b.temp","b.rain")</pre>
# specify nb iterations for burn-in and final inference
nb.burnin < -1000
nb iterations <-2000
```

Run Jags

```
# load R2jags
library(R2jags)
# run Jags
storks <- jags(data = datax,
               inits = inits,
               parameters.to.save = parameters,
               model.file = "code/logistic.txt",
               # model.file = logistic, # if a function was written
               n.chains = 2.
               n.iter = nb.iterations,
               n.burnin = nb.burnin)
storks
```

Inspect parameter estimates

```
#> Compiling model graph
#>
     Resolving undeclared variables
     Allocating nodes
#>
  Graph information:
#>
     Observed stochastic nodes: 23
#>
     Unobserved stochastic nodes: 3
#>
     Total graph size: 181
#>
  Initializing model
#> Inference for Bugs model at "code/logistic.txt", fit using jags,
   2 chains, each with 2000 iterations (first 1000 discarded)
\# n.sims = 2000 iterations saved
           mu.vect sd.vect 2.5%
                                     25%
                                             50%
                                                    75% 97.5% Rhat n.eff
#>
             1.548 0.084 1.429 1.518 1.553 1.589 1.669 1.196
#> a
                                                                      2000
#> b.rain -0.153 0.061 -0.264 -0.197 -0.153 -0.110 -0.045 1.018
                                                                      2000
             0.028 0.062 -0.092 -0.011
                                           0.026
                                                  0.067
                                                          0.157 1.007
                                                                       230
#> b.temp
#> deviance 206.443 31.697 201.780 202.774 203.886 205.682 213.467 1.089
                                                                      2000
```

Your turn

Practical

- Run the stork analysis yourself.
- Does it seem like there is an effect of rainfall or temperature on breeding success?

Assess convergence

Reminder - MCMC Algorithm

 MCMC algorithms can be used to construct a Markov chain with a given stationary distribution (set to be the posterior distribution).

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- For the MCMC algorithm, the posterior distribution is only needed to be known up to proportionality.
- Once the stationary distribution is reached we can regard the realisations of the chain as a (dependent) sample from the posterior distribution (and obtain Monte Carlo estimates).
- We consider some important implementation issues.

MCMC - Proposal Distribution

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- To implement a MCMC algorithm, we often need to specify a proposal distribution from which we generate candidate value then accept/reject.
- This typically involves
 - specifying a given distribution family (e.g. normal, uniform), and then,
 - setting the parameters of the given distribution.
- Although the exact distribution specified is essentially arbitrary it will have a significant effect on the performance of the MCMC algorithm.

• If only small moves can be proposed, the acceptance probability is high, but it will take a long time to explore the posterior distribution.

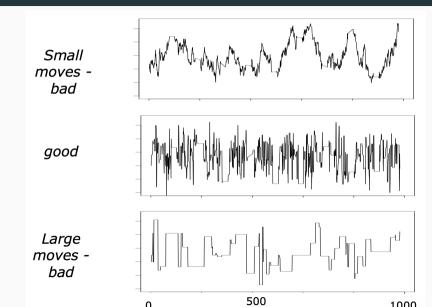
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- The movement around the parameter space is often referred to as mixing.

Good/Bad Traces



Autocorrelation functions

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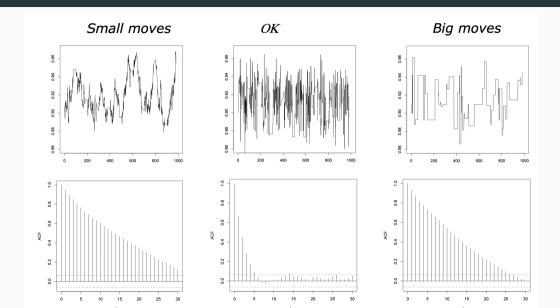
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Autocorrelation functions

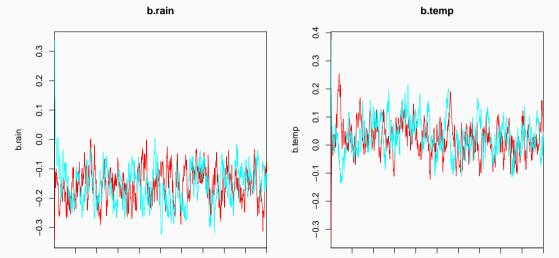
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- Autocorrelation function (ACF) plots are a convenient way of displaying the strength of autocorrelation in the given sample values.
- ACF plots provide the autocorrelation between successively sampled values separated by k iterations, referred to as lag, (i.e. $cor(\theta_t, \theta_{t+k})$) for increasing values of k.

ACFs

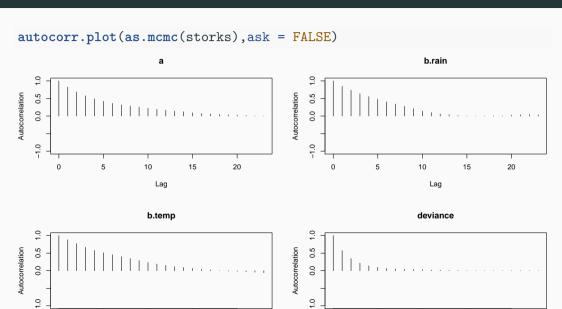


Traceplots for the storks

traceplot(storks,mfrow = c(1, 2), varname = c('b.rain', 'b.temp'), ask = FA



Autocorrelation for the storks



How do good chains behave?

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- Once there, explore efficiently: The post-convergence sample size required for suitable numerical summaries.

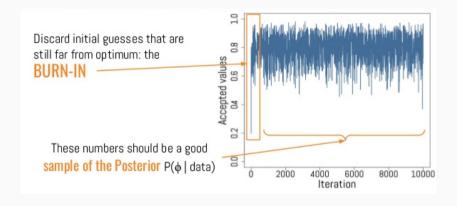
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- In practice, we must discard observations from the start of the chain and just use observations from the chain once it has converged.
- The initial observations that we discard are referred to as the **burn-in**.
- The simplest method to determine the length of the burn-in period is to look at trace plots.

Burn-in (if simulations cheap, be conservative)



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- The effective sample size (n.eff) measures chain length while taking into account the autocorrelation of the chain.
 - n.eff is less than the number of MCMC iterations.
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 - n.eff is less than the number of MCMC iterations.
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- We need n.eff \geq 100 independent steps.

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- Gelman-Rubin statistic \hat{R}
- Measures the ratio of the total variability combining multiple chains (between-chain plus within-chain) to the within-chain variability. Asks the question is there a chain effect? Very much alike the F test in an ANOVA.
- Values near 1 indicates likely convergence, a value of ≤ 1.1 is considered acceptable.
- Necessary condition, not sufficient; In other words, these diagnostics cannot tell you that you have converged for sure, only that you have not.

n.eff and \hat{R} for the storks

storks

- #> Inference for Bugs model at "code/logistic.txt", fit using jags,
- #> 2 chains, each with 2000 iterations (first 1000 discarded)

DTC info (using the rule nD = var(deviance)/2)

- # n.sims = 2000 iterations saved
- mu.vect sd.vect 2.5% 25% 50% 75% 97.5% Rhat #>

- #> a
- #> b.rain -0.153 0.061 -0.264 -0.197 -0.153 -0.110 -0.045 1.018
- #> b.temp 0.028 0.062 -0.092 -0.011 0.026 0.067 0.157 1.007
- #> deviance 206.443 31.697 201.780 202.774 203.886 205.682 213.467 1.089
- #>
- #> For each parameter, n.eff is a crude measure of effective sample size,
- #> and Rhat is the potential scale reduction factor (at convergence, Rhat= #>

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- Assume convergence when all chains reach same regime.
- Discard initial burn-in phase.
- Check autocorrelation, effective sample size and \hat{R} .

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- Something wrong with your model?
 - Start with a simpler model (remove complexities).
 - Use simulations.
- Change your sampler. Upgrade to Nimble or Stan.

MCMC makes you queens and kings

of the stats world

Get all values sampled from posteriors

```
res <- as.mcmc(storks) # convert outputs in a list
res <- rbind(res[[1]],res[[2]]) # put two MCMC lists on top of each other
head(res)
#>
                a b.rain b.temp
                                         deviance
#> [1,] 0.04622874 -0.3417552 -0.33815288 1272.3221
#> [2,] 0.39929350 -0.2120148 -0.24532150 793.1198
#> [3,] 0.62334979 -0.1868884 -0.14196832 559.0869
#> [4,] 0.80827838 -0.1216267 -0.07715738 420.4606
#> [5,] 0.95639600 -0.1104892 -0.02060528 336.1302
#> [6,] 1.04438951 -0.1639800 0.04484738 294.1633
tail(res)
#>
                   b.rain b.temp deviance
#> [1995,] 1.448762 -0.1244035 -1.459287e-02 206.1755
```

#\ [1006] 1 | F7597 _0 1567100 _2 0226080_00 206 2116

Compute a posteriori Pr(rain < 0)

```
# probability that the effect of rainfall is negative
mean(res[,'b.rain'] < 0)
#> [1] 0.994
```

Compute a posteriori Pr(temp < 0)

```
# probability that the effect of temperature is negative
mean(res[,'b.temp'] < 0)
#> [1] 0.325
```

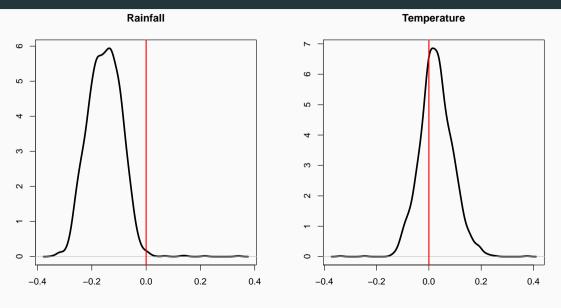
Get credible interval for the rain effect

```
quantile(res[,'b.rain'],c(0.025,0.975))
#> 2.5% 97.5%
#> -0.26365265 -0.04505914
```

Get credible interval for the temperature effect

```
quantile(res[,'b.temp'],c(0.025,0.975))
#> 2.5% 97.5%
#> -0.09205657 0.15735156
```

Graphical summaries



Your turn

A stupid question

• Get the posterior distribution of $b_{rain}^2 + \cos(b_{temp})$

Solution

Evaluate the function for each MCMC iteration

```
stupid_pd <- res[,'b.rain']^2 + cos(res[,'b.temp'])
head(stupid_pd)
#> [1] 1.060166 1.015010 1.024867 1.011818 1.011996 1.025884
```

Plot the distribution

```
plot(density(stupid_pd), xlab = '', main = '', lwd = 3)
```

