

Distributing OpenBUGs chains in parallel

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What is OpenBUGs?

- ▶ Software for **B**ayesian inference **U**sing **G**ibbs **S**ampling
- ▶ Uses Monte Carlo Markov Chains (MCMC) to model posterior distributions

Why use OpenBUGs?

- ▶ Free and open source software
- ▶ Integrates with R through R2OpenBugs/Coda packages
- ▶ Used in Bayesian analysis class at Iowa

What is being parallelized?

- ▶ Using MCMC requires drawing many samples from a converged chain
- ▶ Determining convergence is easier with multiple chains
- ▶ If chains converge to different spots, something is wrong
- ▶ Each chain is independent of the others, allowing for naive parallelization
- ▶ Just need to make sure they're seeded independently

Easy cluster/CPU parallization in R

- ▶ We used the Snow and Snowfall packages
- ▶ They let you easily set up a cluster to run R code on
- ▶ sfLapply works like regular apply, but distributes it over the cluster

Snow/Snowfall Example

- ▶

```
data = matrix(rnorm(1000 * 1000), ncol=100)
library(snow)
library(snowfall)
sfInit(parallel=TRUE, cpus=4)
print(system.time(sfLapply(data, mean)))
sfStop()
sfInit(parallel=TRUE, cpus=1)
print(system.time(sfLapply(data, mean)))
sfStop()
```
- ▶ Parallel: 2.61 seconds
- ▶ Sequential: 6.26 seconds

Parallel Bugs workflow

1. Define openbugs model as a string, data as an R list, and inits as a list
2. Call parallelBugs with these arguments, and the IP addresses of cluster machines
3. Use Coda or other R packages for analyzing MCMC output

Parallel Bugs workflow

1. Define openbugs model as a string, data as an R list, and inits as a list

```
model.str = "model
{
  tau ~ dgamma(0.0001, 0.0001)
  mu ~ dflat()
  sigma <- 1 / sqrt(tau)

  for(i in 1:N)
  {
    x.data[i]~dnorm(mu, tau)
  }
}"
```


Parallel Bugs workflow

1. Define openbugs model as a string, data as an R list, and inits as a list

```
# Simulated normal dataset
bugs.data = list(x.data=rnorm(1000, 12, 5), N=1000)

# Initial values of parameters. Note that you can use
# R random number functions here
init.list = list(
  chain1=list(mu=runif(1, -50, 50), tau=1),
  chain2=list(mu=-50, tau=10),
  chain3=list(mu=50, tau=.1)
)

# Parameters to collect statistics on
params <- c("mu", "sigma")
```

Does it help?

1. Parallel time for 100,000 iterations, 3 chains: 60 seconds
2. Sequential time: 115 seconds

```
system.time(bugs(data=bugs.data, inits=init.list,  
                 parameters.to.save=params, n.iter=100000,  
                 n.chains=3, n.burnin=1000, n.thin=1,  
                 model.file="model.txt", debug=FALSE,  
                 codaPkg=TRUE, OpenBUGS.pgm=OpenBUGS.exe,  
                 working.directory=folder,  
                 bugs.seed=as.integer(runif(1, 1, 14))))
```

```
user  system elapsed  
0.17   0.03   60.69
```

```
system.time(parallel.bugs('localhost', 3, model.str,  
                           bugs.data, 3, init.list, params,  
                           OpenBUGS.exe, n.iter=100000))
```

```
user  system elapsed  
0.42   0.00  115.22
```

Why Create an R Package?

- ▶ Functions that you write and use often can be easily loaded and used on your local machine
- ▶ Can be uploaded to CRAN, allowing other R users to utilize the functions you wrote
- ▶ Provides documentation, making your code easily understandable for other users

How to Create an R Package?

Packages that you will first need to download:

- ▶ devtools

```
install.packages("devtools")
```

- ▶ roxygen2

```
devtools::install_github(klutometis/roxygen)
```

Creating Your Package Directory

- ▶ Navigate to your desired parent directory
- ▶ Create the R package
`create("packagename")`
- ▶ The package you created will include:
 - ▶ Description document
 - ▶ man folder
 - ▶ R folder

Adding Functions to R Package

- ▶ Write your R function(s) and save them in to a .R file
- ▶ If you have multiple functions, you can either save them in multiple .R files or write them sequentially in one .R file
- ▶ Copy your .R file(s) to the R folder

Install Your New R Package!

- ▶ Make sure to navigate back to the parent directory
- ▶ Install package
`install("packagename")`
- ▶ Your R package is now ready to use!

References

- ▶ Luke Tierney, A. J. Rossini, Na Li and H. Sevcikova (2016). snow: Simple Network of Workstations. R package version 0.4-2. <https://CRAN.R-project.org/package=snow>
- ▶ Jochen Knaus (2015). snowfall: Easier cluster computing (based on snow).. R package version 1.84-6.1. <https://CRAN.R-project.org/package=snowfall>
- ▶ Martyn Plummer, Nicky Best, Kate Cowles and Karen Vines (2006). CODA: Convergence Diagnosis and Output Analysis for MCMC, R News, vol 6, 7-11