

Extensible software for hierarchical modeling: using the NIMBLE platform to explore models and algorithms

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<http://r-nimble.org>

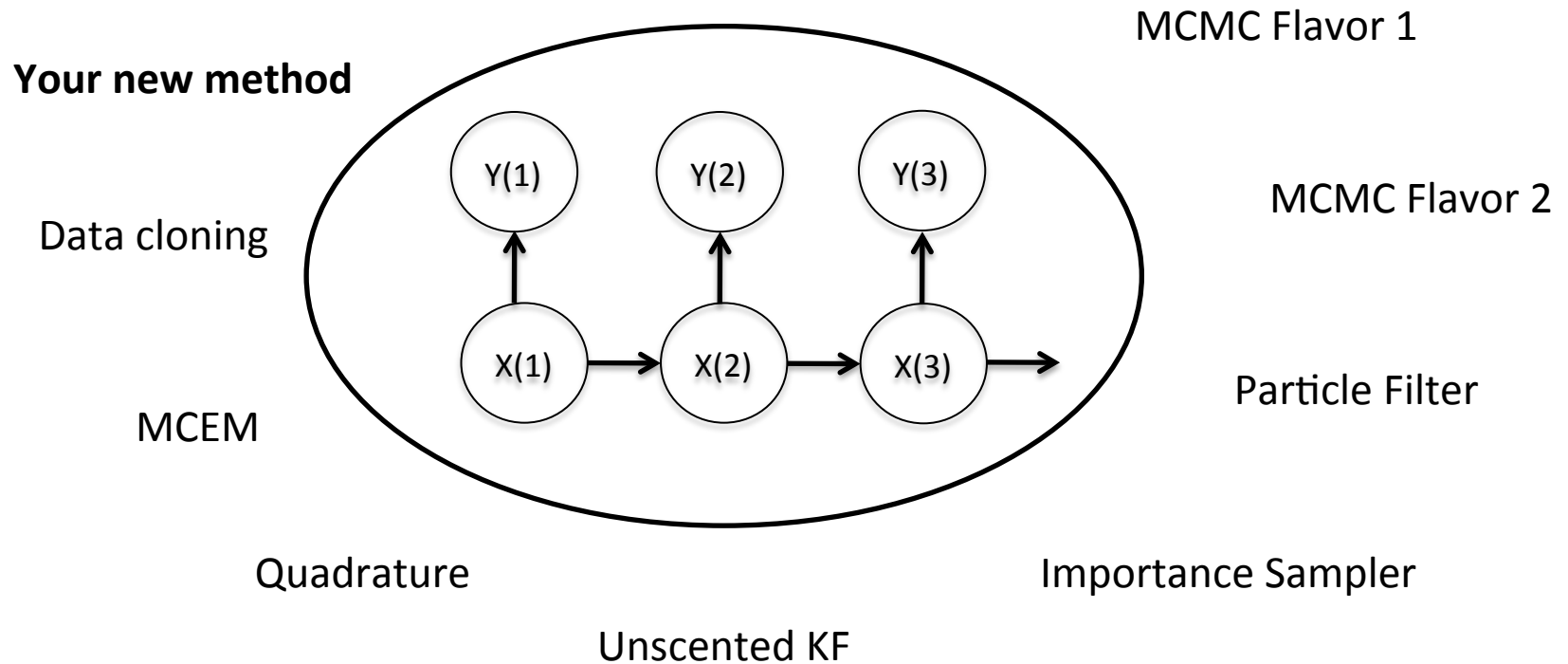
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Background and Goals

- Software for fitting hierarchical models has opened their use to a wide variety of communities
- Most software for fitting such models is either model-specific or algorithm-specific
- Software is often a black box and hard to extend
- Our goal is to divorce model specification from algorithm, while
 - Retaining BUGS compatibility
 - Providing a variety of standard algorithms
 - **Allowing developers to add new algorithms (including modular combination of algorithms)**
 - Allowing users to operate within R
 - Providing speed via compilation to C++, with R wrappers

Divorcing Model Specification from Algorithm



NIMBLE Design

- High-level processing in R (as much as possible)
 - Process BUGS language for declaring models (with some extensions)
 - Process model structure (node dependencies, conjugate relationships, etc.)
 - Generate and customize algorithm specifications
 - Generate model-specific C++ code to be compiled on the fly
 - Provide matching implementation in R for prototyping / debugging / testing
 - Some high-level algorithm control possible in R (adapting tuning parameters, monitoring convergence, high levels of iteration)
- Low-level processing in C++
 - Model and algorithm computations
 - “Run-time” parameters allow some modification of behavior without recompiling

User Experience: Creating a Model from BUGS

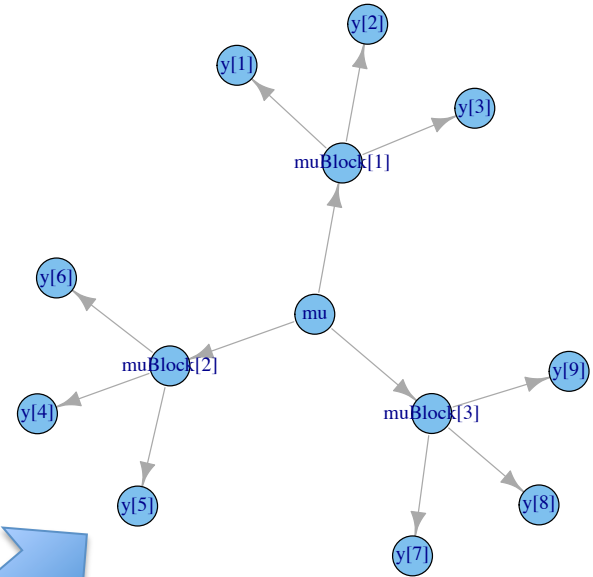
```
littersModelCode <- quote({
  for(j in 1:G) {
    for(l in 1:N) {
      r[l, j] ~ dbin(p[l, j], n[l, j]);
      p[l, j] ~ dbeta(a[j], b[j]);
    }
    mu[j] <- a[j]/(a[j] + b[j]);
    theta[j] <- 1.0/(a[j] + b[j]);
    a[j] ~ dgamma(1, 0.001);
    b[j] ~ dgamma(1, 0.001);
  }
})
```

1

Parse and process BUGS code.
Collect information in model object.

2

Use igraph plot method.



3

Provides variables and functions
(calculate, simulate) for algorithms to
use.

```
> littersModel <- nimbleModel(littersModelCode, constants = list(N = 16, G = 2), data = list(r = input$r))
> littersModel_cpp <- compileNimble(littersModel)
```

User Experience: Specializing an Algorithm to a Model

```
littersModelCode <- modelCode({
  for(j in 1:G) {
    for(l in 1:N) {
      r[i, j] ~ dbin(p[i, j], n[i, j]);
      p[i, j] ~ dbeta(a[j], b[j]);
    }
    mu[j] <- a[j]/(a[j] + b[j]);
    theta[j] <- 1.0/(a[j] + b[j]);
    a[j] ~ dgamma(1, 0.001);
    b[j] ~ dgamma(1, 0.001);
  })
```

```
sampler_slice <- nimbleFunction(
  setup = function((model, mvSaved, control) {
    calcNodes <- model$getDependencies(control$targetNode)
    discrete <- model$getNodeInfo()[[control$targetNode]]$isDiscrete()
    [...snip...]
  })
  run = function() {
    u <- getLogProb(model, calcNodes) - rexp(1, 1)
    x0 <- model[[targetNode]]
    L <- x0 - runif(1, 0, 1) * width
    [...snip....]
  })
  ...
```

```
> littersMCMCspec <- MCMCspec(littersModel)
> getUpdaters(littersMCMCspec)
[...snip...]
[3] RW sampler; targetNode: b[1], adaptive: TRUE, adaptInterval: 200, scale: 1
[4] RW sampler; targetNode: b[2], adaptive: TRUE, adaptInterval: 200, scale: 1
[5] conjugate_beta sampler; targetNode: p[1, 1], dependents_dbin: r[1, 1]
[6] conjugate_beta sampler; targetNode: p[1, 2], dependents_dbin: r[1, 2]
[...snip...]
> littersMCMCspec$addSampler('slice', list(targetNodes = c('a[1]', 'a[2]'), adaptInterval = 100))
> littersMCMCspec$addMonitor('theta')
> littersMCMC <- buildMCMC(littersMCMCspec)
> littersMCMC_Cpp <- compileNimble(littersMCMC, project = littersModel)

> littersMCMC_Cpp(20000)
```

User Experience: Specializing an Algorithm to a Model (2)

```
littersModelCode <- quote({
  for(j in 1:G) {
    for(l in 1:N) {
      r[i, j] ~ dbin(p[i, j], n[i, j]);
      p[i, j] ~ dbeta(a[j], b[j]);
    }
    mu[j] <- a[j]/(a[j] + b[j]);
    theta[j] <- 1.0/(a[j] + b[j]);
    a[j] ~ dgamma(1, 0.001);
    b[j] ~ dgamma(1, 0.001);
  })
```

```
buildMCEM <- nimbleFunction(
  while(runtime(converged == 0)) {
    ....
    calculate(model, paramDepDetermNodes)
    mcmcFun(mcmc.its, initialize = FALSE)
    currentParamVals[1:nParamNodes] <- getValues(model,paramNodes)
    op <- optim(currentParamVals, objFun, maximum = TRUE)
    newParamVals <- op$maximum
    ....
```

```
> littersMCEM <- buildMCEM(littersModel, latentNodes = 'p', mcmcControl = list(adaptInterval =
50), boxConstraints = list( list('a', 'b'), limits = c(0, Inf))), buffer = 1e-6)
> set.seed(0)
> littersMCEM(maxit = 50, m1 = 500, m2 = 5000)
```

Modularity (UNDER CONSTRUCTION):

One can plug any MCMC sampler into the MCEM, with user control of the sampling strategy, in place of the default MCMC.

Programmer Experience: NIMBLE Algorithm DSL

- Analogy: BUGS is a Domain-Specific Language (DSL) for models
- NIMBLE provides a DSL for algorithms
 - The DSL is a modified subset of R.
- We provide
 - Basic types (double, logical)
 - Basic (vectorized) math and distribution/probability calculations
 - Basic data storage classes (“modelValues”)
 - Control structures – for loops and if-then-else
 - Ability to define functions
 - Linear algebra (via the Eigen package)
 - Specific functions for a model: *calculate*, *simulate*
- Function definitions in the DSL include code for two steps:
 - A generic run-time function is written in the DSL for any model structure
 - When a model is provided, a set of one-time setup processing is executed in R based on the model structure to “specialize” algorithm to model
 - Run-time code can use information determined from the setup processing

Programmer Experience: Creating an Algorithm

```
myAlgorithmGenerator <- nimbleFunction (  
  
  setup = function(model, <otherSetupArguments>) {  
  
    # code that does the specialization of algorithm to model  
    # e.g., determine nodes to sample,  
    # initialize storage  
  
  },  
  
  run = function(<runtimeArguments>) {  
  
    # code that carries out the generic algorithm  
    # for example, iterations of an algorithm  
    # simulate into nodes, calculate log probability values  
    returnType(double())  
    return(x)  
  })
```



Two sections to a
NIMBLE function.

Usage:

```
specializedAlgo <- myAlgorithmGenerator(myModel, <setupArgs>)  
specializedAlgo(<runtimeArguments>)
```

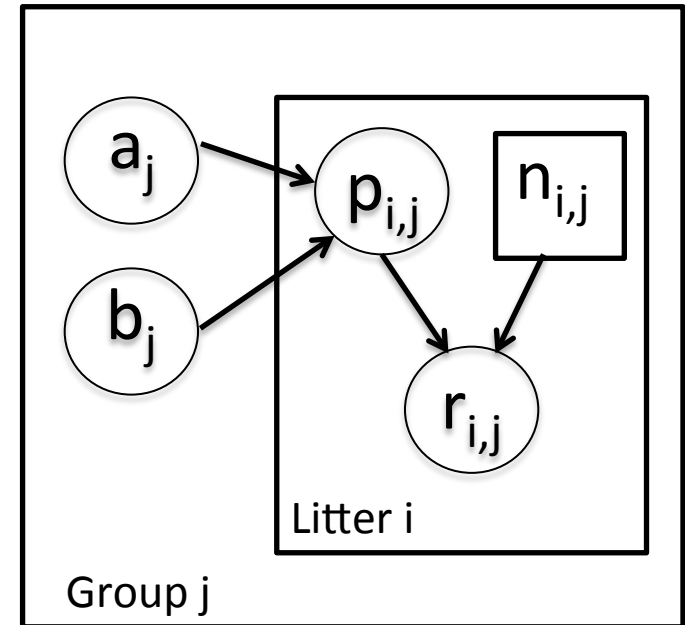
Programmer Experience: Slice Sampler Example

```
sampler_slice <- nimbleFunction( contains = sampler_BASE,  
  setup = function(model, mvSaved, control) {  
    targetNode  <- control$targetNode  
    adaptive    <- control$adaptive  
    ....  
    calcNodes <- model$getDependencies(targetNode)  
    ....  
    discrete   <- model$getNodeInfo()[[targetNode]]$isDiscrete()  
  },  
  run = function() {  
    u <- getLogProb(model, calcNodes) - rexp(1, 1)  
    x0 <- model[[targetNode]  
    L <- x0 - runif(1, 0, 1) * width  
    R <- L + width  
    maxStepsL <- floor(runif(1, 0, 1) * maxSteps)  
    maxStepsR <- maxSteps - 1 - maxStepsL  
    lp <- setAndCalculateTarget(L)  
    while(maxStepsL > 0 & !is.nan(lp) & lp >= u) {  
      L <- L - width  
      lp <- setAndCalculateTarget(L)  
      maxStepsL <- maxStepsL - 1  
    }  
    ....  
  }  
)
```

NIMBLE in Action: the Litters Example

Beta-binomial for clustered binary response data

```
littersModelCode <- quote({
  for(j in 1:G) {
    for(l in 1:N) {
      r[i, j] ~ dbin(p[i, j], n[i, j]);
      p[i, j] ~ dbeta(a[j], b[j]);
    }
    mu[j] <- a[j]/(a[j] + b[j]);
    theta[j] <- 1.0/(a[j] + b[j]);
    a[j] ~ dgamma(1, 0.001);
    b[j] ~ dgamma(1, 0.001);
  })
```

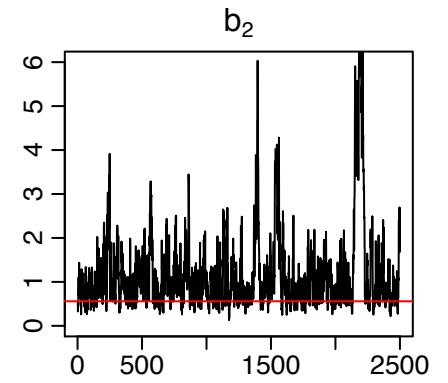
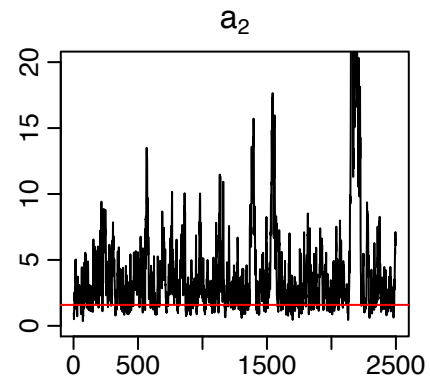
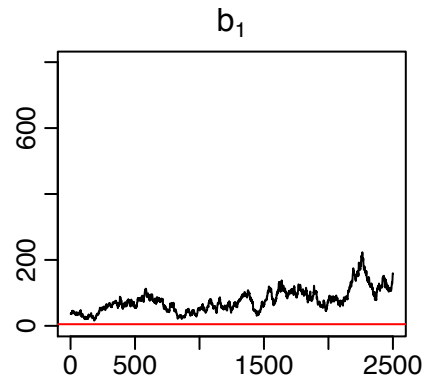
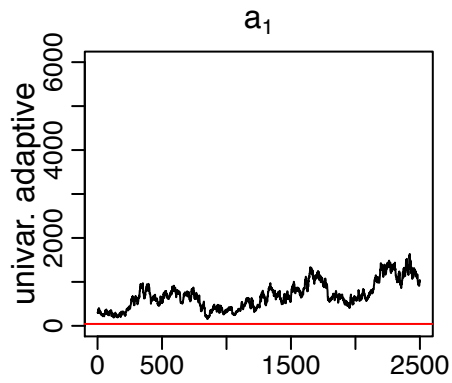


Challenges of the toy example:

- BUGS manual: “The estimates, particularly a_1 , a_2 suffer from extremely poor convergence, limited agreement with m.l.e.’s and considerable prior sensitivity. This appears to be due primarily to the parameterisation in terms of the highly related a_j and b_j , whereas direct sampling of μ_j and θ_j would be strongly preferable.”
- But that’s not all that’s going on. Consider the dependence between the p ’s and their a_j , b_j hyperparameters.
- And perhaps we want to do something other than MCMC.

Default MCMC: Gibbs + Metropolis

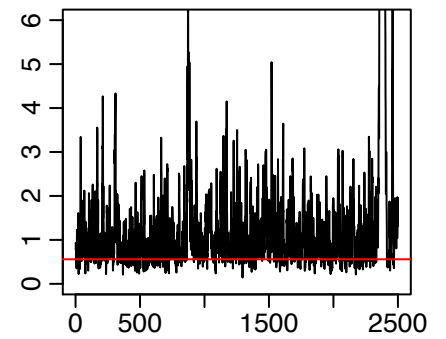
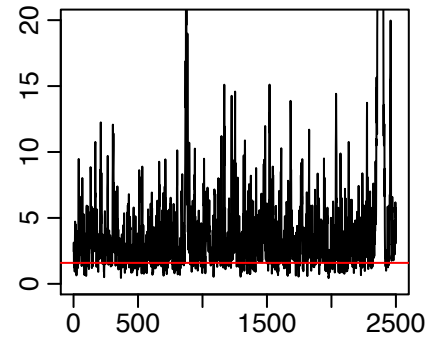
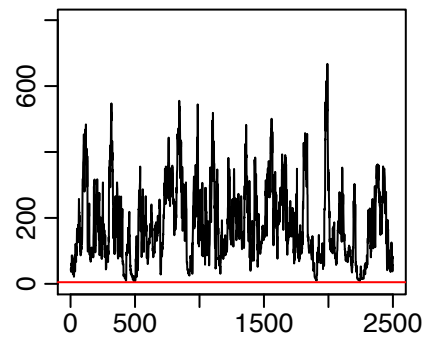
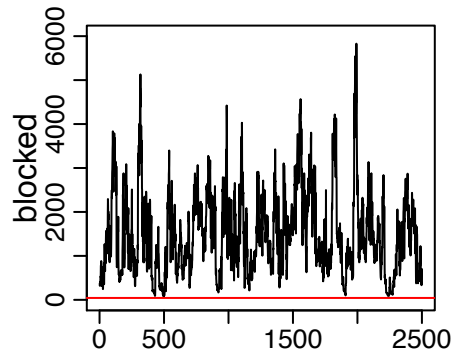
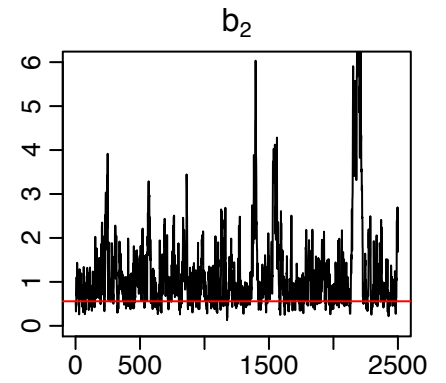
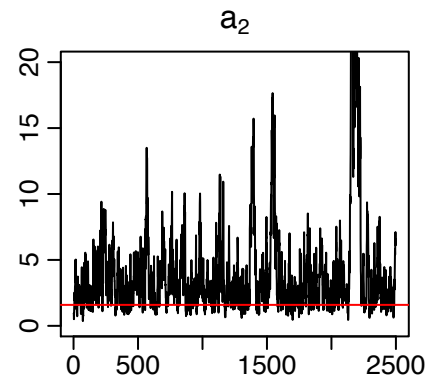
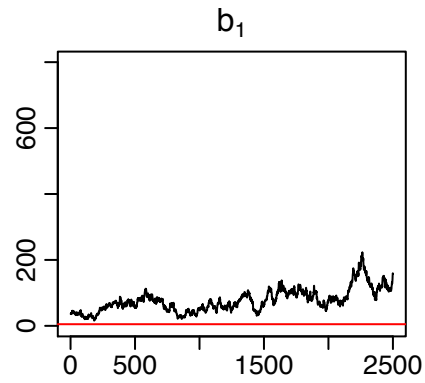
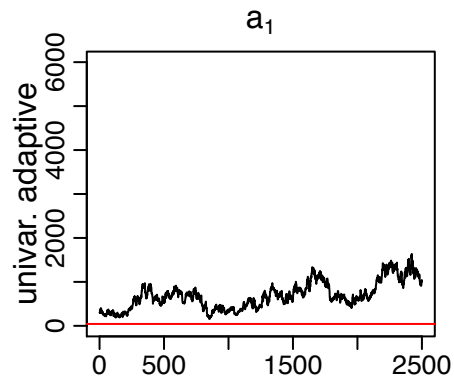
```
> littersMCMCspec <- MCMCspec(littersModel, list(adaptInterval = 100))  
> littersMCMC <- buildMCMC(littersMCMCspec)  
> littersMCMC_cpp <- compileNIMBLE(littersModel, project = littersModel)  
> littersMCMC_cpp(10000)
```



Red line is MLE

Blocked MCMC: Gibbs + Blocked Metropolis

```
> littersMCMCspec2 <- MCMCspec(littersModel, list(adaptInterval = 100))
> littersMCMCspec2$addSampler('RW_block', list(targetNodes = c('a[1]', 'b[1]'), adaptInterval = 100))
> littersMCMCspec2$addSampler('RW_block', list(targetNodes = c('a[2]', 'b[2]'), adaptInterval = 100))
> littersMCMC2 <- buildMCMC(littersMCMCspec2)
> littersMCMC2_cpp <- compileNIMBLE(littersMCMC2, project = littersModel)
> littersMCMC2_cpp(10000)
```



Blocked MCMC: Gibbs + Cross-level Updaters

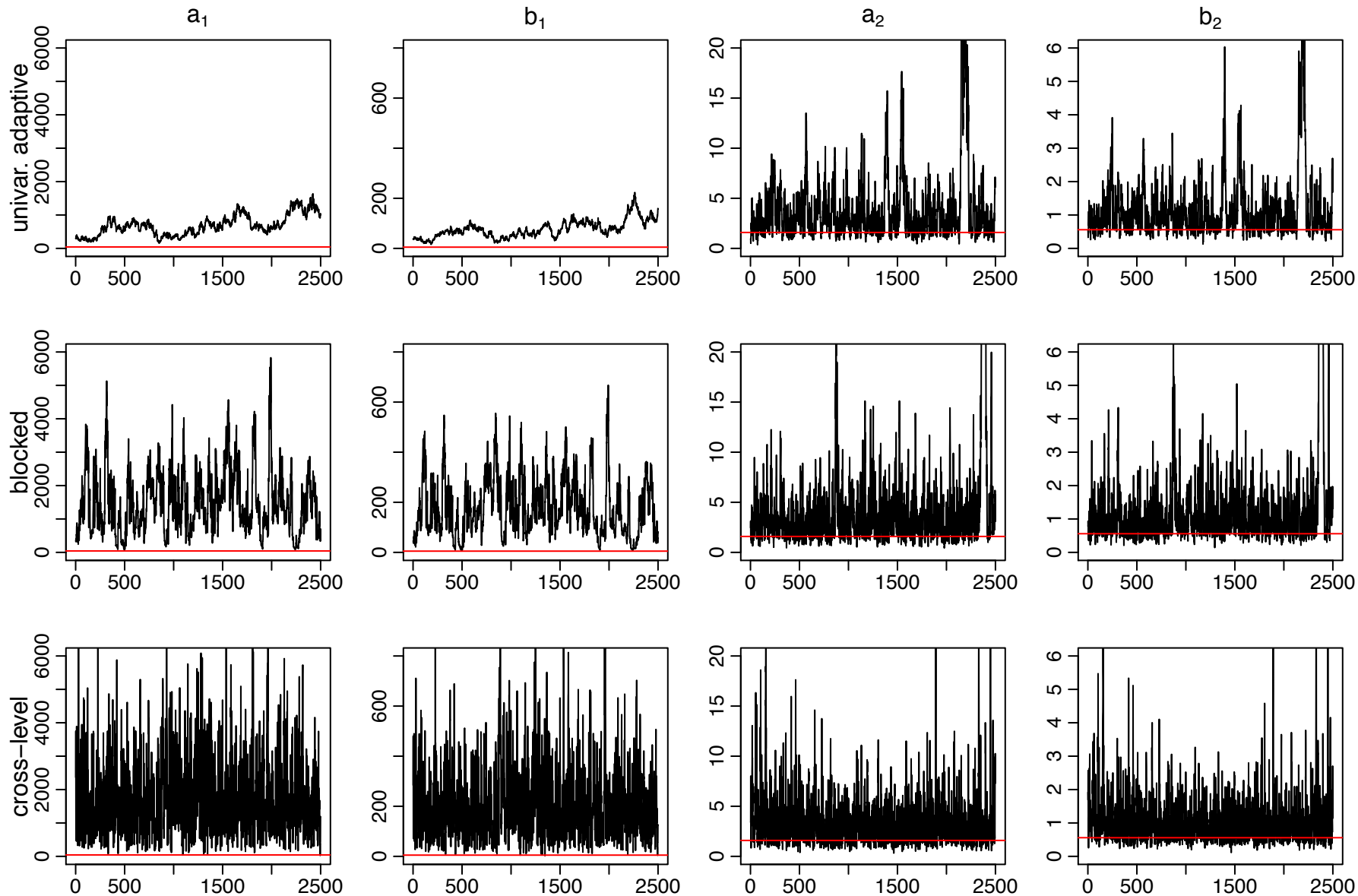
- Cross-level dependence is a key barrier in this and many other models.
- We wrote a new “cross-level” updater function using the NIMBLE DSL.
 - The updater is a blocked Metropolis random walk on a set of hyperparameters with conditional Gibbs updates on dependent nodes (provided they are in a conjugate relationship).
 - This is equivalent to (analytically) integrating the dependent (latent) nodes out of the model.
- We can then add this updater to an MCMC for a given model.....

```
> littersMCMCspec3 <- MCMCspec(littersModel, adaptInterval = 100)

> topNodes1 <- c('a[1]', 'b[1]')
> littersMCMCspec3$addSampler('crossLevel', list(topNodes = topNodes1, adaptInterval = 100)

> topNodes2 <- c('a[2]', 'b[2]')
> littersMCMCspec3$addSampler('crossLevel', list(topNodes = topNodes1, adaptInterval = 100)

> littersMCMC3 <- buildMCMC(littersMCMCspec3)
> littersMCMC3_cpp <- compileNIMBLE(littersMCMC3, project = littersModel)
> littersMCMC3_cpp(10000)
```

Litters MCMC: BUGS and JAGS

- BUGS gives similar performance to the default NIMBLE MCMC
 - Be careful – values of `$sim.list` and `$sims.matrix` in R2WinBUGS output are randomly permuted
 - Mixing for `a2` and `b2` modestly better than default NIMBLE MCMC
- JAGS slice sampler gives similar performance BUGS, but fails for some starting values with this (troublesome) parameterization
- NIMBLE provides user control and transparency.
 - NIMBLE is faster than JAGS on this example (if one ignores the compilation time).
 - Note: we're not out to build the best MCMC but rather a flexible framework for algorithms – we'd love to have someone else build a better default MCMC and distribute for use in our system.

Stepping outside the MCMC box: maximum likelihood/empirical Bayes via MCEM

```
> littersMCEM <- buildMCEM(littersModel, latentNodes = 'p')  
> littersMCEM(maxit = 500, m1 = 500, m2 = 5000)
```

- Gives estimates consistent with direct ML estimation (possible in this simple model with conjugacy for 'p') to 2-3 digits
- VERY slow to converge, analogous to MCMC mixing issues
- Stochasticity in the embedded MCMC makes this basic MCEM unstable; a more sophisticated treatment should help here

Many algorithms are of a modular nature/combine other algorithms, e.g.

- particle MCMC
- normalizing constant algorithms
- many, many others in the literature in the last 15 years

Status of NIMBLE and Next Steps

- First release was last week; lots to do, including:
 - Improve the user interface and speed up compilation
 - Refinement/extension of the DSL for algorithms
 - Enhance current algorithms provided (e.g., add multivariate conjugate updates for MCMC)
 - Additional algorithms written in NIMBLE DSL (e.g., particle MCMC)
 - Advanced features (e.g., auto. differentiation, paralleliz'n)
- Interested?
 - Announcements: [nimble-announce](#) Google site
 - User support/discussion: [nimble-users](#) Google site
 - Write an algorithm using NIMBLE!
 - Help with development of NIMBLE: email nimble.stats@gmail.com or see github.com/nimble-dev