# The R Package Management System: Bayesian Change Point Analysis

### John W. Emerson

http://www.stat.yale.edu/~jay/ Associate Professor of Statistics, Yale University (Professor Emerson prefers to be called "Jay")

I would like to thank Chandra Erdman (Yale GRD '08) for her collaboration on the Bayesian change point analysis and package **bcp**. Chandra is now at the Census Bureau.

#### Please feel free to ask questions along the way!

http://www.stat.yale.edu/~jay/Brazil/Campinas/bcpANDpackages/http://www.stat.yale.edu/~jay/RPC

### **Outline**

- Overview
- Bayesian Change Point Analysis
- The R Package Management System
- 4 The C/C++ Interface
- Parallel Programming

### Why R?

R is the *lingua franca* of statistics:

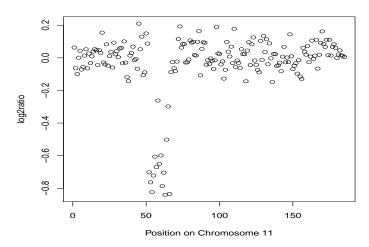
- The syntax is simple and well-suited for data exploration and analysis.
- It has excellent graphical capabilities.
- It is extensible, with over 2500 packages available on CRAN alone.
- It is open source and freely available for Windows/MacOS/Linux platforms.

This talk emphasizes the importance of the package management system. Much of the success of R should be attributed to:

- Ross & Robert's early decision to go open-source and encourage collaboration, and
- the growth of CRAN and the success of the package management system.



### Example: Coriell cell lines (raw data)

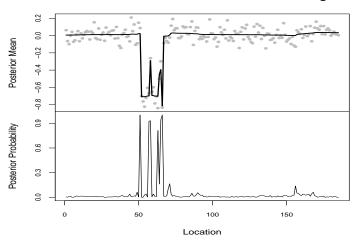


# Example: Coriell cell lines (bcp analysis)

See http://cran.r-project.org/web/packages/bcp/.

### Example: Coriell cell lines (bcp output)

#### Posterior Means and Probabilities of a Change



### The Bayesian change point model

- ho unknown partition into continuous blocks, with the transition between blocks being the "change points."
- p probability of a change point at position i, independently for all i.
- $X_i$  observations assumed independent  $N(\mu_i, \sigma^2)$ , where in this notation the  $\mu_i$  are equal for all i within a block.
- $\mu_{jk}$  mean of block from position j+1 to k, with prior  $N(\mu_0, \sigma_0^2/(k-j))$ . Note: larger deviations from  $\mu_0$  are expected for shorter blocks, but weak signals can be detected when sufficient data are available.
- w defined as  $\sigma^2/(\sigma^2 + \sigma_0^2)$  for convenience.

Possible point of notational confusion: conditional on  $\rho$  (the partition into blocks),  $\mu_i \equiv \mu_{jk}$  for all i in block jk.

### The Bayesian change point priors

$$\pi(\mu_0) = 1, -\infty \le \mu_0 \le \infty$$

$$\pi(\sigma^2) = 1/\sigma^2, \ 0 \le \sigma^2 \le \infty$$

$$\pi(\rho) = 1/\rho_0, \ 0 \le \rho \le \rho_0$$

$$\pi(w) = 1/w_0, \ 0 \le w \le w_0$$

$$\pi(\rho) = \frac{1}{\rho_0} \left[ \int_0^{\rho_0} \rho^{b-1} (1-\rho)^{n-b} d\rho \right]$$

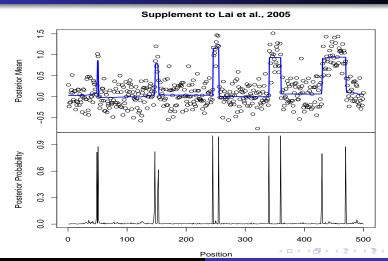
where  $p_0$  and  $w_0$  are pre-selected in [0, 1] (values of 0.2 are the default in package **bcp** and work well in a wide range of cases), and b is the number of blocks in  $\rho$ .

### Notes on Bayesian change point analysis

- Barry and Hartigan (1993): "A Bayesian analysis for change point problems" in JASA, 88, 309-319.
- Erdman and Emerson (2008): "A fast Bayesian change point analysis for the segmentation of microarray data" in Bioinformatics, 24 (19), 2143-2148.
- Erdman and Emerson (2007): "bcp: an R package for performing a Bayesian analysis of change point problems" in JSS, 23 (3).
- An exact implementation of the Bayes procedure is possible but the calculations would be  $O(n^3)$ .
- Package bcp provides a fast O(n) MCMC implementation:
  - inefficient MCMC would be O(n²)
  - solves some nasty numerical problems with large data
  - supports parallel MCMC
  - currently being extended for multivariate series with a common change point structure



# Second example: simulated abberations of length 2, 5, 10, 20, and 40



### Building an R build environment

If you want to build R packages, you'll need the full R build environment (not just the pre-compiled R binary that most of us use from CRAN). See

http://www.stat.yale.edu/~jay/RPC/RPackages.pdf

# A simple package: function babywhatis()

```
> 1s()
[1] "bcp.11" "chrom11" "coriell"
> rm(list = ls())
> 1s()
character (0)
 babywhatis <- function(x) {
      if (!is.data.frame(x)) {
+
          x < - data.frame(x)
+
+
          warning("Object coerced to a data frame.\n")
+
      return(unlist(lapply(x, class)))
+
+
> 1s()
   "babywhatis"
```

### A simple package: the package skeleton

```
> package.skeleton("MyToolkit")
Creating directories ...
Creating DESCRIPTION ...
Creating Read-and-delete-me ...
Saving functions and data ...
Making help files ...
Done.
Further steps are described in
   './MyToolkit/Read-and-delete-me'.
```

Let's go investigate together; we'll explore the package structure, make minor modifications, and will check/build/install it. More information is available in

http://www.stat.yale.edu/~jay/RPC/RPackages.pdf



# Package bcp

At this point, we'll just glance at package **bcp** quickly. I want to introduce the C/C++ interface with a simple example before looking more closely at **bcp**.

### A simple example: column minima

This material wouldn't display well in slides. Again see

http://www.stat.yale.edu/~jay/RPC/RPackages.pdf

and, specifically, materials in

http://www.stat.yale.edu/~jay/RPC/MyToolkitWithC/

# Package bcp

Package bcp uses the .C() interface instead of .Call(), though I'd like to change this in a new version.

### foreach

The user may register any one of several "parallel backends" like **doMC** or **doSNOW**, or none at all. The code will either run sequentially or will make use of the parallel backend, if specified, without code modification.

```
> library(foreach)
> library(doMC)
 registerDoMC(2)
>
> a < -10
 ans \leftarrow foreach (i = 1:5, .combine = c) %dopar%
          a + i^2
+
>
 ans
 [1]
       11
             14
                   19
                       26 35
```

### Parallel MCMC in package bcp

- An older version of bcp used NetWorkSpaces for parallel MCMC; this was very difficult to install and use, and the code was not portable to other parallel environments.
- The new bcp uses Steve Weston's foreach package, and the user may choose from a variety of parallel backends.
- I strongly recommend foreach() for parallel programming, to both users and package developers.
- Again, see supplementary materials for more information.

### **Thanks**

- Bell Laboratories (Rick Becker, John Chambers and Allan Wilks), for development of the S language
- Ross Ihaka and Robert Gentleman, for their work and unselfish vision for R
- The R Core team
- John Hartigan, for years of teaching and mentoring
- John Emerson (my father, Middlebury College), for getting me started in statistics and computer science
- All my students (and Chandra Erdman in particular on this project) for their willingness argue with me

