BAYESIAN ANALYSIS USING MONTE CARLO INTEGRATION WITH AN EXAMPLE OF THE ANALYSIS OF SURVIVAL DATA

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SUMMARY

Bayesian analysis using Monte Carlo integration is an effective approach for handling rich multiparameter families of distributions; nonconjugate priors; censored data; extrapolation uncertainty; and the computation of posterior distributions for parameters or predictions of interest. In the example, posterior percentile curves for a rate function are computed from survival data.

<u>Keywords</u>: Bayesian statistics; Monte Carlo integration; posterior distributions; censored data; survival analysis.

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1. INTRODUCTION

Bayesian statistics and numerical methods form a particularly effective partnership. There are at least two reasons for this:

- o It is often easy to compute the values of the likelihood function and the prior density (and therefore the posterior density up to a multiplicative constant) on any specified set of points in the parameter space. This is true even for many models and problems that would be very difficult to handle with conventional methods.
- The principal computational problem in making a Bayesian inference or decision is the evaluation of integrals involving the posterior density.

Thus the solution to many difficult inference and decision problems can be obtained by numerical integration using only the values of the likelihood function and the prior density on a finite set of parameter points. The suitable choice of this set is the most important, and sometimes the most difficult, aspect of the computation.

2. BAYESIAN ANALYSIS USING MONTE CARLO INTEGRATION

Bayesian inference and decision making usually require the computation of posterior cumulative distribution functions and/or posterior moments of functions of the parameter vector

 $\boldsymbol{\Theta} = (\boldsymbol{\Theta}_1, \boldsymbol{\Theta}_2, \dots, \boldsymbol{\Theta}_K)' \boldsymbol{\varepsilon} \boldsymbol{\Theta}$

This requires evaluation of integrals of the form

where h is a real valued function. The posterior density $\xi(\varphi|D)$ is given by Bayes' theorem,

 $\xi(e|D) \propto \xi(e) L(e|D)$

where $\xi(\varphi)$ is the prior density and $L(\varphi|D)$ is the likelihood function corresponding to the observed data D.

The Monte Carlo approximation to I is given by

$$\hat{I} = \sum_{m=1}^{M} h(\theta_{m}) W(\theta_{m}) / \sum_{m=1}^{M} W(\theta_{m})$$
(2)

The "weights" $W(\theta_m)$, m = 1, 2, ..., M, are computed from

$$W(\Theta) = \xi(\Theta) L(\Theta|D)/g^{*}(\Theta)$$
(3)

The θ_m , m = 1, 2, ..., M, are generated independently from the K-variate density $g^*(\theta)$. This is known as "importance sampling." The density $g^*(\theta)$ is called the "importance function" or "generating density" and is usually chosen to approximate the posterior density, with the restriction that its form must be such that the θ_m 's can be easily generated.

Details about the techniques used in applying this methodology and discussions of Monte Carlo integration error and its reduction can be found in Stewart 1968, 1970, 1977, 1979, 1983; Stewart and Johnson 1971, 1972; Johnson and Stewart 1971; Kloek and van Dijk 1975, 1978; van Dijk and Kloek 1978, 1980, 1983; McGhee and Walford 1965, 1967, 1968; Heiberger 1976; Zellner and Rossi 1982.

3. EXAMPLE

Consider data of the type shown in Fig. 1. Subjects enter the study at different ages and remain in the study until death, withdrawal or the termination of the study. (There may be some subjects who would have been in the study but died before their entry date. Since these occurrences are not included in the data, the term "left truncation" is sometimes used with this type of data.)

Let the subscript i denote the ith subject and

 $E_{j}^{}$ -age at entering the study $X_{j}^{}$ - age at death $C_{j}^{}$ - age at exit from study due to withdrawal or termination of study. \sim

For each subject we observe E_i and min (X_i, C_i)

We define a rate function r(x) by:

$$r(x) dx = P[x < X_{j} < x + dx] E_{j} \leq x < X_{j}, x + dx \leq C_{j}]$$
(4)



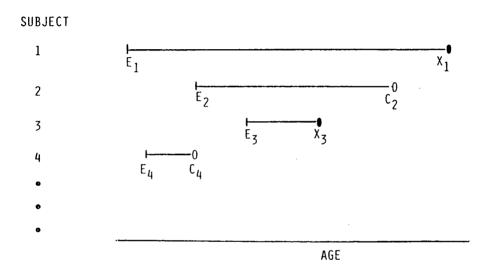


Fig. 1

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and assume that r(x) does not depend on i or E_i . (We also assume that the mechanisms that determine E_i and C_i are non-informative for inferences about r(x).)

r(x) is the hazard rate (or force of mortality) function corresponding to a density f(x) and cumulative distribution function, F(x). They are related by:

$$\mathbf{r}(\mathbf{x}) = \frac{\mathbf{f}(\mathbf{x})}{1 - \mathbf{F}(\mathbf{x})} \tag{5}$$

The conditional density of X given E and that $C = \infty$ is

$$f(x|E; c=\infty) = \frac{f(x)}{1-F(E)} x > E$$
 (6)

The likelihood function given the data D is

$$L(\boldsymbol{\theta}|\boldsymbol{D}) = \prod_{i:X_{i} < C_{i}} \begin{bmatrix} f(X_{i}|\boldsymbol{\theta}) \\ 1-F(E_{i}|\boldsymbol{\theta}) \end{bmatrix} \prod_{j:X_{j} > C_{j}} \begin{bmatrix} 1-F(C_{j}|\boldsymbol{\theta}) \\ 1-F(E_{j}|\boldsymbol{\theta}) \end{bmatrix}$$
$$= \frac{\prod_{i:X_{i} < C_{i}} f(X_{i}|\boldsymbol{\theta}) \prod_{j:X_{j} > C_{j}} [1-F(C_{j}|\boldsymbol{\theta})]}{\prod_{i} [1-F(C_{j}|\boldsymbol{\theta})]}$$
(7)

The data used in this example were from a survival study of the male residents of Channing House, a retirement center located in Palo Alto. The data and further information are given in Hyde 1977, 1980. Of the total of 97 men in the study, 46 died, 5 withdrew and 46 survived to the end of the study. The ages at entry varied from 62 to 89 years.

The prior distribution and the eight parameter (K=8) state space of rate functions were defined in a manner similar to Example 1 of Stewart 1979 but with $K^* = 16$ as defined in Section 3.1. An analyst with experience derived from similar situations may have chosen a prior with a different structure, but this should cause no problems because of the great flexibility allowed in choosing priors when using Monte Carlo integration.

Posterior percentile curves for the values of the rate functon, r(x), are shown in Fig. 2. Also shown are two dashed curves, designated by M and X, which are, respectively, the posterior mean and the r(x) that maximizes $\xi(e|D)$. To obtain these curves, Monte Carlo integration was used to compute the posterior distribution of the value of the function at each of the 15 evenly spaced values of X. Each posterior was evaluated at 250 points along the vertical axis. This required the simultaneous Monte Carlo evaluation of 3765 integrals. The same set $\{\theta_m, H(\theta_m): m=1, 2, \ldots, M=2000\}$ was used for all of the integrals.

Displaying a sample of rate functions from the posterior is another way of illustrating the properties of the posterior. Twenty rate functions generated from the posterior are displayed in Fig. 3. The sample was generated by the "acceptance-rejection" method described in Stewart 1983.

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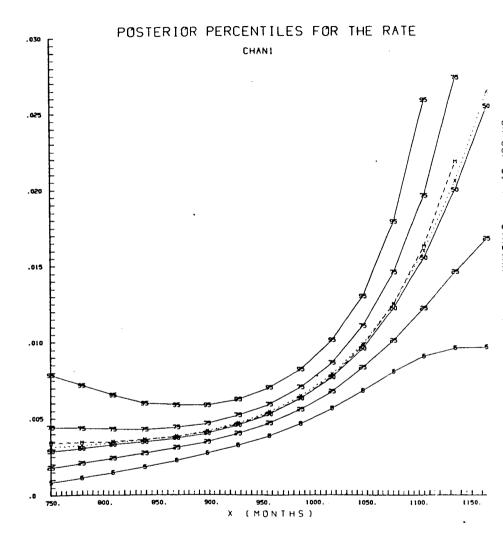
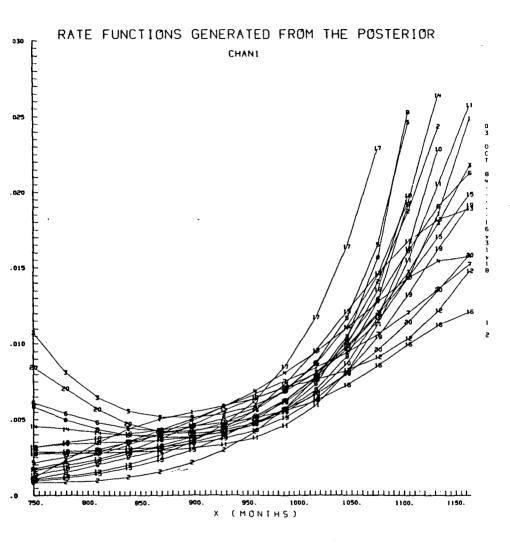


Fig. 2

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