80 051 2001

NUREG/CR-1401 EGG-EA-5112

# Estimators for the Binomial Failure Rate Common Cause Model

Propared by C.L. Atwood

EG&G Idaho, Inc.

Prepared for U.S. Nuclear Regulatory Commission

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# Estimators for the Binomial Failure Rate Common Cause Model

Manuscript Completed: March 1980 Date Published: April 1980

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Prepared for Probabilistic Analysis Staff Office of Nuclear Regulatory Research U.S. Nuclear Regulatory Commission Washington, D.C. 20555 NRC FIN No. A6292

#### ABSTRACT

In Vesely's binomial failure rate model, a system of m components is hit by random shocks which may cause components simultaneously to fail, each component with equal probability. Individual components may also fail when no shock has occurred. The data possibilities considered are that causes of single failures are identifiable (as shock or not) or not identifiable. Given data from such a system, non-Bayesian and Bayesian point and interval estimators are found for the various quantities of interest. Residual analyses and hypothesis tests are presented for checking the model assumptions. An example is worked out.

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#### ACKNOWLEDGEMENTS

I am grateful to W. E. Vesely for many very helpful comments and suggestions, and to N. D. Cox for valuable discussion. W. J. Suitt wrote the extensive computer coding for performing the work of Section 7.

# ESTIMATORS FOR THE BINOMIAL FAILURE RATE COMMON CAUSE MODEL

#### 1. SUMMARY

Vesely<sup>1</sup> proposes a binomial failure rate (BFR) model for modeling common cause failures in a system. In this model, each component of the system has a constant failure rate  $\lambda$ . In addition, a "common cause shock," or "secondary event" in the terminology of reliability analysts, can occur with constant occurrence rate  $\mu$ . If a shock occurs, the components fail independently of each other, each with probability p.

This paper presents results concerning the BFR model. There are six or seven related, unknown quantities of interest. The exact number depends on whether the causes of single failures are identifiable (i.e., due to shock or just individual failures) or not identifiable. Maximum likelihood estimators are given for all of these quantities. One of the likelihood equations requires numerical solution. Confidence intervals are given, which are sharp for some quantities and conservative for others. Confidence regions, not all rectangular, are given for estimating the quantities simultaneously.

Bayes estimators (posterior modes and means) and Bayes probability intervals are given for all the quantities of interest, as is the threedimensional probability region for the three basic parameters. All the Bayesian results, except finding the modes, require numerical integration. However, in almost all cases, only integrals with respect to a single variable need to be done numerically. Depending on the parameters of the prior distribution, the integrand may contain an infinite series.

Residual analyses and hypothesis tests are presented for checking the model assumptions. Most of the hypothesis tests require a large sample size.

The results are applied to some of the boiling water nuclear reactor control rod data given by  $Vesely^1$ .

#### 2. BASIC CALCULATIONS

Let there be a system with m components operating for time t. (if there are several such systems with the same m, each operating for time  $t_i$ , consider them as one system with  $t = \Sigma t_i$ .) Assume m > 2. (In Section 6, where the causes of single failures are identifiable, we will relax this assumption to  $m \ge 2$ .) Each component has an exponentially distributed lifetime with parameter (failure rate)  $\lambda$ , and the failures are mutually independent. In addition, a common cause shock may hit the system, with the shocks mutually independent and the time between shocks exponentially distributed with parameter  $\mu$ . If a shock hits the system, the components of the system fail independently of each other, each with probability p. All failures are discovered and repaired as soon as they occur. Define q = 1 - p.

Let  $N_i$  be the number of occurrences of i simultaneous failures. (We will follow the convention of using capital letters to denote random variables and using the corresponding small letters to denote specific values of the random variables.) Then, basic calculations (see Reference 1 or Mann, Schafer, and Singpurwalla,<sup>2</sup> Section 4.2) yield that

N; are independent Poisson( $\lambda_i$  t)

with

 $\lambda_1 = m\lambda + \mu r_1$ 

and

 $\lambda_{i} = \mu r_{i}, \quad i = 2, ..., m.$ 

Here, r is defined by

$$r_i = \binom{m}{i} p^i q^{m-i}.$$

(2.2)

(2.1)

The special expression for  $\lambda_1$  comes from the fact that a single failure may be simply an individual failure or it may come from a common cause shock that caused only one component to fail. (Reliability analysts would refer to a common cause shock that failed fewer than two components as a "potential common cause.")

The model has been formulated in terms of  $\lambda$ ,  $\mu$ , and p. Also of interest, indeed perhaps of great interest, are  $\lambda_1$ , defined by (2.1), and  $\lambda_+$ , defined by

$$\lambda_{+} = \sum_{i=2}^{m} \lambda_{i} = \mu(1 - r_{0} - r_{1}).$$
(2.3)

The quantity  $\lambda_+$  is the rate of common cause failure occurrences. It must be distinguished from  $\mu$ , the rate of common cause shocks. A last quantity of interest is  $\mu p^k$  for  $2 \le k \le m$ . This is the rate at which some specific k components fail simultaneously, and is the relevant rate corresponding to a k-element "AND" gate in a fault tree, or to a k-element cut set.

It may be that the causes of single failures are identifiable as either individual failure or common cause shock. Then,  $N_1$  can be decomposed into  $N_{I} + N_{C}$ , where  $N_{I}$  is the number of individual failures, distributed Poisson(mat), and  $N_{C}$  is the number of single failures due to common cause shocks, distributed Poisson( $\mu r_1 t$ ). It may be very difficult to extract the necessary information for this decomposition from failure reports. For example, in nuclear industry Licensee Event Reports, the information may simply be unavailable: the cause is reported "unknown." Or, it may be difficult to interpret: is "dirt in valve" a common cause shock that happened to fail only one valve? Finally, the report may be untrustworthy: in one report, 96 control rods failed to insert fully. A possible cause reported was that oil leaked past 96 independent piston seals. We do not believe in such a coincidence, but assume some common cause unrecognized by the reporter. But do we then classify a reported single oil leakage past a piston seal as an individual failure, or as a common cause shock that failed only one component?

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There are data sets in which causes of single failures are identifiable. For example, failures due to personnel error may all be regarded as due to common cause shock. However, the major portion of this paper assumes that  $N_I$  and  $N_C$  are not separately available. Everything simplifies if they are available, so the results are summarized in Section 6.

Since t is fixed, basic calculations show that the probability distribution of a set of failure data depends only on the number of failures, not on the times of the failures (Reference 2, page 180). Define

$$N_{+} = \sum_{i=2}^{m} N_{i} .$$

Then

$$P[N_{1} = n_{1}, ..., N_{m} = n_{m}]$$
  
=  $P[N_{1} = n_{1}] P[N_{+} = n_{+}] P[N_{2} = n_{2}, ..., N_{m} = n_{m}|N_{+} = n_{+}]$ 

symbolically written as

$$L = L_1 \times L_2 \times L_3$$

This decomposition is useful because

$$N_{1} \sim Poisson(\lambda_{1}t)$$

$$N_{+} \sim Poisson(\lambda_{+}t)$$

$$N_{2}, \dots, N_{m} \mid N_{+} = n_{+} \sim multinomial(n_{+}, z_{2}, \dots, z_{m})$$

where

 $z_i = r_i / (1 - r_0 - r_1)$ .

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The multinominal probability  $L_3$  can be written out as

$${}^{P}\left[N_{2} = n_{2}, \dots, N_{m} = n_{m} | N_{+} = n_{+}\right] = \frac{n_{+}!}{n_{2}! \cdots n_{m}!} \prod_{i=2}^{m} z_{i}^{n_{i}}$$

$$= \frac{n_{+}!}{n_{2}! \cdots n_{m}!} \frac{p^{s} q^{mn_{+}} - s}{(1 - q^{m} - mp q^{m-1})^{n_{+}}} \prod_{i=2}^{m} (m_{i}^{n})^{n_{i}}$$
(2.6)

where s is defined as

$$s = \sum_{i=2}^{m} i n_i$$
,

the total number of components failing in multiple failures.

Note that  $2n_+ \leq s \leq mn_+$ . Note also that  $(N_1, N_+, S)$  is a sufficient statistic.

#### 3. NON-BAYESIAN INFERENCE

#### 3.1 Maximum Likelihood Estimators

It will be most convenient to parametrize the model in terms of  $(\lambda_1, \lambda_+, p)$ . From estimators of these quantities, estimators of  $\lambda$ ,  $\mu$ , and  $\mu p^k$  will also be developed. The parameters  $\lambda_1$ ,  $\lambda_+$ , and p are related by (2.1) through (2.3). If we set  $\lambda \ge 0$  in (2.1), we obtain that the parameters satisfy the constraint

$$\lambda_{1} \geq \lambda_{+} \left[ m p q^{m-1} / (1 - q^{m} - m p q^{m-1}) \right].$$
(3.1)

This can be abbreviated  $\lambda_1 \ge \lambda_+ z_1$ , in the notation of (2.5).

From (2.4), the logarithm of the likelihood is

$$\log L = \log L_1 + \log L_2 + \log L_3$$

where

$$\log L_1 = -\lambda_1 + n_1 \log \lambda_1 - \log (n_1!)$$

 $\log L_2 = -\lambda_+ + n_+ \log \lambda_+ - \log (n_+!)$ 

and L<sub>3</sub> is given by (2.6) and does not depend on  $\lambda_1$  or  $\lambda_+$ .

Let us first maximize log L ignoring the constraint (3.1). The maximizing values of  $\lambda_1$  and  $\lambda_+$  are

$$\hat{\lambda}_1 = n_1/t$$

$$\hat{\lambda}_{+} = n_{+}/t.$$

The maximizing value of p, denoted  $\hat{p}$ , must now be found.

If m = 2 or if  $n_{+} = 0$ , then  $L_{3}$  is identically 1. We have assumed throughout that m > 2. Now assume also that  $n_{+} > 0$ . Differentiation of log  $L_{3}$  is straightforward:

$$\frac{\partial \log L_3}{\partial p} = \frac{s}{p q} - m n_+ \frac{1 - q^{m-1}}{q(1 - q^m - m p q^{m-1})} .$$
(3.2)

It is shown in Section 8.2 that if  $s = 2n_+$ , then  $\hat{p} = 0$ . If  $s = m n_+$ , then  $\hat{p} = 1$ . And otherwise,  $q(a \log L_3/ap)$  is strictly decreasing in p and changes sign between 0 and 1; that is,  $\hat{p}$  is the unique solution of

$$s = m n_{+} p \frac{1 - q^{m-1}}{1 - q^{m} - m p q^{m-1}}$$
 (3.3)

If m = 3, the smallest value that allows p to be estimated, then (3.3) has an explicit solution:  $\hat{p} = 3(s - 2n_+)/(2s - 3n_+)$ . For larger m, the solution must be found numerically.

The naive estimate of p would be  $s/m n_+$ , the average proportion of failed components in multiple failure occurrences. We would expect this naive estimate to be biased upwards, since it is based on only those common cause shocks that happen to result in at least two failures. And, in fact,  $s/m n_+$  is somewhat larger than  $\hat{p}$ , the MLE. The naive estimator  $s/m n_+$  can be used as an initial guess for solving (3.3).

It is interesting to note that an estimator based on the conditional first moment of S is the same as  $\hat{p}$ , because the equation

$$s = E \left[ S | N_{+} = n_{+} \right]$$

reduces to (3.3). This can be shown directly or derived from (3.2) and the well-known fact that, under regularity conditions,  $E(a \log L_3/ap) = 0$ . The equality of the two estimators is not mere coincidence, but follows from the fact that the truncated binomial distribution is in the exponential family. If  $s/n_+ >> 2$ , then the naive estimate and the MLE are approximately equal. For in this case, the data show many failures per observed shock, indicating that virtually all common cause shocks result in at least two failures. If  $s/n_+$  is close to 2, then the two estimates differ appreciably. The quantity  $S/N_+$  will cend to be large if mp is large and small if mp is small.

Sometimes m is not known exactly. Then it is important to know how  $\hat{p}$  varies as a function of m. It is shown in Section 8.1 that the right side of (3.3) is an increasing function of m, for fixed  $\hat{p}$ . Therefore, as m increases, p must be decreased to preserve equality in (3.3). That is, the MLE  $\hat{p}$  decreases as m increases.

Once  $\hat{\lambda}_1$ ,  $\hat{\lambda}_+$ , and  $\hat{p}$  are found,  $\hat{\lambda}$  and  $\hat{\mu}$  follow from (2.1) through (2.3), and the MLE for  $\mu p^k$  is  $\hat{\mu} \hat{p}^k$ .

It may be that the above procedure makes  $\hat{\lambda}$  negative! This is because log L was maximized without any contraints, whereas it should have been maximized subject to the constraint (3.1); i.e.,  $\lambda \ge 0$ . In this case, proceed as follows.

As a function of  $\lambda_1$ ,  $\lambda_+$ , and p, log L has a unique local maximum. Therefore, the local maximum is still unique when log L is written as a function of  $\lambda$ ,  $\mu$ , and p. If this local maximum occurs at  $\hat{\lambda} < 0$ , then the maximum subject to  $\lambda \ge 0$  must occur at  $\lambda = 0$ . So, set  $\lambda = 0$  in (2.4), and set  $\hat{\lambda} = 0$ . Define

$$N = \sum_{i=1}^{m} N_i$$

Then

 $L = P[N = n] P[N_1 = n_1, \dots, N_m = n_m | N = n]$ 

where

$$N \sim Poisson(\lambda' t), \quad \lambda' = \mu(1 - r_0)$$

and

$$P[N_{1} = n_{1}, \dots, N_{m} = n_{m} | N = n]$$

$$= \frac{n!}{n_{1}! \cdots n_{m}!} \frac{p^{s'} q^{m} n - s'}{(1 - q^{m})^{n}} \prod_{i=1}^{m} {m \choose i}$$

with

$$s' = \sum_{i=1}^{m} i n_i$$
.

The MLE's are found by setting

 $\hat{\lambda}' = n/t$ 

and letting  $\hat{p}$  be the solution of

 $s' = m n p/(1 - q^m)$ .

This  $\hat{p}$  is also the estimator obtained by setting

s' = E[S'|N = n].

It decreases as m increases. Then  $\hat{\mu}$  follows from  $\hat{\lambda}$ ',  $\hat{p}$ , and the definition of  $\lambda$ ', as does  $\hat{\mu}\hat{p}^k$ .

#### 3.2 Confidence Intervals

(3.4)

3.2.1 Terminology. Since the data are discrete, all confidence regions are necessarily inexact, in the sense that

 $P[region contains parameter] > 1 - \alpha$ 

for all values of the parameter, with strict inequality for some values of the parameter. This is a well-known characteristic of discrete data. A  $(1 - \alpha)$  confidence region will be called <u>sharp</u> and the inequality (3.4) will be called <u>sharp</u> if the infimum over all values of the parameter satisfies

inf P[region contains parameter] =  $1 - \alpha$ .

[We do not simply say that equality is attained in (3.4) for some p, since attainment of equality depends on whether open or closed confidence intervals are used and whether the parameter space is open or not. Use of the infimum avoids that difficulty.] The confidence region will be called conservative if (3.4) holds but is not sharp.

3.2.2 Intervals for  $\lambda_1$ ,  $\lambda_+$ , and p. Sharp confidence intervals for  $\lambda_1$  and  $\lambda_+$  are completely standard since N<sub>1</sub> and N<sub>+</sub> are independent Poisson variables. Expressions for the lower and upper limits for  $\lambda_1$  that are easy to compute from readily accessible tables, are

$$\lambda_{1L} = (X^{2} 2n_{1}, \alpha/2)/2t$$
$$\lambda_{1U} = (X^{2} 2(n_{1}+1), 1-\alpha/2)/2t$$

The interval for  $\lambda_{+}$  is of the same form, using  $n_{+}$  instead of  $n_{1}$ . For details, see Johnson and Kotz<sup>3</sup>, Vol. 1, Sec. 4.6.2. A sharp confidence interval ( $p_L$ ,  $p_U$ ) for p can be based on the conditional distribution of S given N<sub>+</sub>, if N<sub>+</sub> > 0. Let  $p_L$  and  $p_U$  satisfy

$$P[S \ge s | N_{+} = n_{+}, p = p_{L}] = \alpha/2$$
 (3.5)

$$P[S \le S | N_{+} = n_{+}, p = p_{U}] = \alpha/2$$
 (3.6)

Depending on whether s is large or small, it will be more convenient to rewrite one of these equations as

$$P[S < s | N_{+} = n_{+}, p = p_{|}] = 1 - \alpha/2$$
 (3.5')

$$P[S > S|N_{+} = N_{+}, p = p_{U}] = 1 - \alpha/2$$
 (3.6')

This choice of the confidence interval has a mathematical statistics justification, given in Section 8.2.

To use (3.5) and (3.6), the conditional distribution of S given N<sub>+</sub> must be found. This involves some complexity, and more notation is needed. Let  $v = (v_2, ..., v_m)$  be any vector of nonnegative integers. Define

$$v_{+} = \sum_{i=2}^{m} v_{i}.$$

Define the set  $T_k$  by

$$T_{k} = \left\{ v \mid v_{+} = n_{+}, \begin{array}{c} m \\ \Sigma & iv_{i} = k \end{array} \right\}.$$

$$i=2$$

$$c(v) = \frac{v_{+}!}{v_{2}! \cdots v_{m}!} \prod_{i=2}^{m} {\binom{n}{i}}^{i}.$$

Then for  $2n_{+} \leq k \leq m n_{+}$ ,

Let

$$P[S = k | N_{+} = n_{+}] = \sum_{v \in T_{k}} P[N_{2} = v_{2}, ..., N_{m} = v_{m} | N_{+} = n_{+}]$$

$$= \frac{m n_{+} - k}{(1 - q^{m} - m p q^{m-1})^{n_{+}}} \sum_{v \in T_{k}} c(v) \qquad (3.7)$$

from (2.6). The summation in (3.7) can be evaluated for the relevant values of k. Then (3.5) and (3.6) can be solved numerically for  $p_L$  and  $p_U$ .

The above procedure gives a sharp confidence interval for p, conditional on N<sub>+</sub> having any specified positive value. That is, the probability that the interval contains p, conditional on N<sub>+</sub>, is  $\geq 1 - \alpha$ , and the inequality is sharp. There is no unconditional confidence level, because when N<sub>+</sub> = 0 no interval has been defined. To remedy this, when N<sub>+</sub> = 0 the interval [0,1] can be used with confidence level 1. Let I(S,N<sub>+</sub>) denote the interval for p, depending on S and N<sub>+</sub>. Then the unconditional confidence level is

$$P\left[p \in I(S, N_{+})\right]$$

$$= \sum_{\substack{n_{+}=0}}^{\infty} P\left[p \in I(S, n_{+}) \mid N_{+} = n_{+}\right] P\left[N_{+} = n_{+}\right]$$

$$\geq \sum_{\substack{n_{+}=1}}^{\infty} (1 - \alpha) P\left[N_{+} = n_{+}\right] + 1 P\left[N_{+} = 0\right]$$

 $\geq$  (1 - a) .

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The first inequality results from the inexactness of the conditional confidence intervals.

Is the unconditional confidence interval sharp? For any fixed  $\lambda_+$ ,  $P[p \in I(S,N_+)]$  is strictly greater than 1- $\alpha$ . However, as  $n_+ \neq \infty$ , the distribution of S given  $n_+$  approaches a continuous distribution (S given  $n_+$  is asymptotically normal, by the central limit theorem). So for any fixed p,  $P[p \in I(S,n_+)|N_+ = n_+] \neq (1-\alpha)$  as  $n_+ \neq \infty$ . Therefore  $P[p \in I(S,N_+)] \geqslant (1-\alpha)$  as  $\lambda_+ t \neq \infty$ , so

$$\inf_{\lambda_+,p} P[p \in I(S,N_+)] = (1-\alpha).$$

In this sense, the unconditional confidence interval is sharp. This sense of sharpness may not be what the user really wants, since  $\lambda_{+}t$  may really be moderately small. It may perhaps be possible to shorten the conditional confidence intervals when  $n_{+}$  is small, and still maintain an unconditional confidence level  $(1-\alpha)$ . However if this is possible, the details are certainly quite complicated.

3.2.3 Joint Region for  $(\lambda_1, \lambda_+, p)$ . Confidence intervals for  $\lambda_1$ ,  $\lambda_+$ , and p have been given separately. Now consider how to find a confidence region for the three parameters taken together.

Let  $\alpha_1$ ,  $\alpha_+$ , and  $\alpha_p$  be between 0 and 1. Let  $I_1(N_1)$  denote the  $1-\alpha_1$  confidence interval for  $\lambda_1$ , and  $I_+(N_+)$  the  $1-\alpha_+$  confidence interval for  $\lambda_+$ . Let  $I_p(S,N_+)$  be the  $1-\alpha_p$  conditional confidence interval for p if  $N_+ > 0$ , and let it equal [0,1] if  $N_+ = 0$ . Let us tentatively use the product of the three intervals as a joint confidence region for the three parameters. Then

$$\begin{split} & P\left[\lambda_{1} \in I_{1}(N_{1}), \ \lambda_{+} \in I_{+}(N_{+}), \ p \in I_{p}(S,N_{+})\right] \\ &= P\left[\lambda_{1} \in I_{1}(N_{1})\right] P\left[\lambda_{+} \in I_{+}(N_{+}), \ p \in I_{p}(S,N_{+})\right] \\ &\geq (1-\alpha_{1}) \sum_{n_{+}=0}^{\infty} P\left[\lambda_{+} \in I_{+}(n_{+}), \ p \in I_{p}(S,n_{+}) | N_{+} = n_{+}\right] P\left[N_{+} = n_{+}\right]. \end{split}$$

The inequality results from the inexactness of  $I_1$ , and is sharp. Now let  $\delta(\lambda_+, n_+) = 1$  if  $\lambda_+ \in I_+(n_+)$ , and let  $\delta(\lambda_+, n_+) = 0$  otherwise. The last expression equals

$$\begin{array}{l} (1-\alpha_{1}) & \sum \limits_{n_{+}=0}^{\infty} \delta(\lambda_{+}, n_{+}) \ P\left[p \ \epsilon \ I_{p} \ (S, n_{+})\right] N_{+} = n_{+}\right] \ P\left[N_{+} = n_{+}\right] \\ \\ & \geq (1-\alpha_{1}) & \sum \limits_{n_{+}=1}^{\infty} \delta(\lambda_{+}, n_{+}) \ (1-\alpha_{p}) \ P\left[N_{+} = n_{+}\right] + (1-\alpha_{1}) \ \delta(\lambda_{+}, 0) \ P\left[N_{+} = 0\right] \\ \\ & \geq (1-\alpha_{1}) \ (1-\alpha_{p}) \ (1-\alpha_{+}) \ . \end{array}$$

As  $\lambda_{+}t \neq \infty$ , the last two inequalities approach equality, so the confidence region is sharp.

In fact, this product of the three intervals may be larger than necessary because a portion of it may not satisfy the constraint (3.1). That portion can be deleted as impossible. To visualize this, think of the product of the three intervals as a block with rectagular cross sections. For any p, the cross section is a rectangle parallel to the  $(\lambda_1, \lambda_+)$ plane. Invocation of (3.1) typically cuts a corner off of the rectangle. Several cross sections for the example of Section 7 are shown in Figure 7.

3.2.4 Joint Regions for Any Two of  $\lambda_1$ ,  $\lambda_+$ , and p. If we are interested in two of  $\lambda_1$ ,  $\lambda_+$ , and p, the constraint (3.1) does not apply, so the region is simply the product of the two intervals. Work parallel to that of the last section shows that the unconditional confidence level is the iduct of the two confidence levels for the intervals.

3.2.5 <u>Confidence Regions for Quantities Other than</u>  $\lambda_1$ ,  $\lambda_+$ , <u>and</u> p. If we are interested in three independent parameters other than  $\lambda_1$ ,  $\lambda_+$ , and p, for example if we want a confidence region for  $(\lambda, \mu, p)$ , the region for  $(\lambda_1, \lambda_+, p)$  can be rewritten in terms of the three desired parameters. The results are not especially neat, but they are straightforward.

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It is not apparent how to get sharp confidence intervals for  $\lambda$ ,  $\mu$ , or  $\mu \mu^k$ , or a sharp two-dimensional region for, say,  $(\lambda, \mu)$ . One conservative region for  $(\lambda, \mu)$  is the set of all  $(\lambda, \mu)$  corresponding to any point in the three-dimensional region. Its confidence coefficient is at least that of the three-dimensional region.

A conservative region for  $\mu$  is based on

$$\mu = \lambda_{+} / (1 - q^{m} - m p q^{m-1}).$$

If  $(\lambda_+, p)$  is in a confidence region with some confidence level, then the resulting maximum and minimum values of  $\mu$  form a confidence interval with at least that confidence level. Similarly, a conservative interval for  $\mu p^k$  is based on

$$\mu p^{k} = \lambda_{+} p^{k} / (1 - q^{m} - m p q^{m-1}).$$

The right hand side is monotone in p for  $k \ge 2$ , so conservative upper and lower values for  $\mu p^k$  can be obtained from the upper and lower values of  $\lambda_+$  and p.

A conservative one-sided interval for  $\lambda$  can be based on

$$\lambda \leq \lambda_1/m$$
.

A conservative two-sided interval can be based on

$$\lambda = \lambda_1 / m - \lambda_+ \left[ p q^{m-1} / (1 - q^m - m p q^{m-1}) \right].$$

This last equation requires maximization or minimization over all three confidence intervals, and has coefficient at least  $(1-\alpha_1)(1-\alpha_p)(1-\alpha_p)$ . In the example of Section 7, the first method gives a smaller 95% confidence interval.

If the sample size is large, approximate confidence regions can be constructed using the Fisher information matrix. (A reference is Cox and Hinkley,<sup>4</sup> Sec. 9.2.iii.) The Fisher information for  $\lambda_1$  or  $\lambda_+$  is  $t/\lambda_1$  or  $t/\lambda_+$ , respectively. To get the information for p, write the right hand side of (3.2) as

$$\frac{s}{pq} - m n_+ \frac{A}{B}$$
.

Then

$$I(p) \equiv -E \frac{a^2}{ap^2} \log L$$
  
=  $-\frac{E(S)(p-q)}{p^2q^2} + m n_+ \frac{BA' - AB'}{B^2}$   
=  $m n_+ \left[\frac{A(q-p)}{pqB} + \frac{BA' - AB'}{B^2}\right]$ .

The off-diagonal elements of the information matrix are 0. So, if the sample size is large,  $\hat{\lambda}_1$ ,  $\hat{\lambda}_+$ , and  $\hat{p}$  are approximately independent normal with means  $\lambda_1$ ,  $\lambda_+$ , and p, and variances  $\lambda_1/t$ ,  $\lambda_+/t$ , and 1/I(p). From this, approximate confidence regions can be found for any subset of  $\{\lambda_1, \lambda_+, p\}$ . A conservative interval for  $\lambda$ ,  $\mu$ , or  $\mu p^k$  can be based on the intervals for  $\lambda_1$ ,  $\lambda_+$ , and p.

(3.8)

#### 4. BAYESIAN INFERENCE

#### 4.1 Prior and Posterior Distributions

The constraint (3.1) makes the Bayesian problem inherently more complicated than the non-Bayesian one. Any prior distribution must be consistent with (3.1). The posterior distribution cannot be factored neatly as could the likelihood (2.4).

A suitable class of prior distributions must be selected. For greatest usefulness, this class should include the noninformative prior of Box and Tiao<sup>5</sup>, Secs. 1.3.4-7. Box and Tiao argue that a noninformative prior for a parameter should be proportional to the square root of the Fisher information for the parameter. Therefore, the (improper) noninformative prior density for  $\lambda_1$ , resp.  $\lambda_+$ , is proportional to  $\lambda_1^{-1/2}$ , resp.  $\lambda_+^{-1/2}$ . These distributions are in the class of (improper) gamma distributions. In this paper,  $\lambda_1$  and  $\lambda_+$  will have gamma prior distributions, possibly improper.

The information for p is the complicated expression (3.8). Therefore, the noninformative prior density for p is proportional to  $I^{1/2}(p)$ , the square root of (3.8). It may be approximated in several ways by a beta distribution, as follows. The beta (1/2, 1/2) distribution would be noninformative if the common cause failures were not restricted so that only multiple failures are observable. This is a first approximation to the noninformative prior. Better approximations are obtained by adjusting the parameters of a beta (c,d) distribution. If c = 1/2, then  $I^{1/2}(p)$  and the beta density will both be asymptotic to  $p^{-1/2}$  for p near 0. Then d may be chosen so that the two distributions have the same mean. For a still better overall approximation, c and d may be chosen so that the two distributions have the same mean and the same variance. Tables 1 and 2 show the correct values c and d for these approximations, for many values of m. Figures 1 through 3 show the noninformative prior cumulative distribution and the three approximations, for m = 5, 20, and 100. In the example considered in Table 7, the third beta approximation appears adequate. In this paper it will be assumed that p has a beta prior distribution, possible one of these approximations to the noninformative distribution.

	d
3	0.2893
4	0.3148
5	0.3323
6	0.3455
7	0.3560
8	0.3645
9	0.3716
10	0.3777
12	0.3877
14	0.3956
16	0.4020
18	0.4073
20	0.4119
25	0.4209
30	0.4276
40	0.4372
60	0.4486
80	0.4555
100	0.4602
150	0.4676
200	0.4720
300	0.4773
80	0.5000

TABLE 1. BINOMIAL WITH O AND 1 TRUNCATED: APPROXIMATELY NONINFORMATIVE d FOR BETA(1/2,d)

_ <u>m_</u>	C	d
3	0.6386	0.3695
4	0.6641	0.4181
5	0.6762	0.4495
6	0.6813	0.4708
7	0.6824	0.4858
8	0.6814	0.4967
9	0.6791	0.5048
10	0 6762	0.5108
12	6695	0.5191
14	0.6626	0.5242
16	0.6560	0.5274
18	0.6498	0.5294
20	0.6442	0.5307
25	0.6321	0.5321
30	0.6223	0.5322
40	0.6075	0.5312
60	0.5888	0.5283
80	0.5772	0.5259
100	0.5692	0.5239
150	0.5567	0.5206
200	0.5492	0.5185
300	0.5404	0.5159
80	0.5000	0.5000

TABLE 2. BING 1 TAUNCATED: APPROXIMATELY NONINFORMATIVE PARAMETERS FOR BETA(c,d)







1 truncated, m = 20.



Since  $\lambda_1$ ,  $\lambda_+$ , and p must satisfy the constraint (3.1), we will take the joint prior density to be proportional to the product of the individual prior densities in the region where (3.1) is satisfied. (See also Reference 5, pp. 56-58 and Section 1.5.) In summary, the following class of prior densities will be considered:

 $L(\lambda_1, \lambda_+, p)$ 

 $= C U(\lambda_{1}, \lambda_{+}, p) e^{-b_{1}\lambda_{1}} \lambda_{1}^{a_{1}-1} e^{-b_{+}\lambda_{+}} \lambda_{+}^{a_{+}-1} p^{c-1} q^{d-1}$ (4.1)

where C is a constant, and U( $\lambda_1$ ,  $\lambda_+$ , p) equals 1 where (3.1) holds, and equals 0 elswhere.

The parameters should be restricted as follows: require  $a_1 > 0$  and  $a_+ > 0$  in order to guarantee that the posterior density, given by (4.2) below, has finite integral for  $n_1 \ge 0$  and  $n_+ \ge 0$ . Require  $b_1 \ge 0$  and  $b_+ \ge 0$ , to force the posterior density to have finite integral for all t > 0. Require c > 0 and d > 0. For, if  $c \le 0$  and  $s = 2n_+ + 1$ , then the posterior density would be positive at p = 0, and if  $d \le 0$  and  $s = mn_+ - 1$ , the posterior density would be positive at p = 1. Either of these possibilities defies any reasonable interpretation of a posterior density, so restrict c and d to prevent it.

If  $a_1 = a_+ = 1/2$  and  $b_1 = b_+ = 0$ , the prior distribution is noninformative for  $\lambda_1$  and  $\lambda_+$ .

The posterior density of  $(\lambda_1, \lambda_+, \gamma)$  is obtained by multiplying (2.4) by (4.1) and adjusting the constant. It will be convenient to write it as follows:

 $L(\lambda_1, \lambda_+, p|data) = C U(\lambda_1, \lambda_+, p) V(\lambda_1, \lambda_+, p)$ (4.2)

where

$$V(\lambda_{1}, \lambda_{+}, p) = \frac{(b_{1}^{+}t)^{a_{1}^{+}n_{1}}}{\Gamma(a_{1}^{+}n_{1}^{-})} e^{-(b_{1}^{+}t)\lambda_{1}} \lambda_{1}^{a_{1}^{+}n_{1}^{-1}}$$

$$\cdot \frac{(b_{+}^{+}t)^{a_{+}^{+}n_{+}}}{\Gamma(a_{+}^{+}n_{+})} e^{-(b_{+}^{+}t)\lambda_{+}} \lambda_{+}^{a_{+}^{+}n_{+}^{-1}}$$

$$\cdot \frac{p^{s+c-1} q^{mn_{+}^{-s+d-1}}}{\Gamma(1-q^{m}-mpq^{m-1})^{n_{+}^{-1}}}.$$
(4.3)

Here C is that constant such that  $\iiint L(\lambda_1, \lambda_+, p \text{ data}) = 1$ .

#### 4.2 Bayes Point Estimators

4.2.1 <u>Posterior Modes</u>. One estimate of  $(\lambda_1, \lambda_+, p)$  is the mode of the posterior density. If the constraint (3.1) is ignored, the maximizing values for  $\lambda_1$  and  $\lambda_+$  are the maximizing values for (4.3),

$$\lambda_{1} = (a_{1} + n_{1} - 1)/(b_{1} + t)$$
  
$$\lambda_{+} = (a_{+} + n_{+} - 1)/(b_{+} + t).$$

To find the maximizing p, compute

$$q \frac{\partial}{\partial p} \log L(\lambda_1, \lambda_+, p | data) = \frac{s + c - 1}{p} - \left\{ c + d - 2 + mn_+ \frac{1 - q^m - 1}{1 - q^m - mpq^{m-1}} \right\}.$$

It is shown in Section 8.1 that this quantity either is always positive, or is always negative, or else is decreasing and equals zero at just one point. The possible cases are tabulated in Section 8.1. Therefore, there is a unique posterior mode p. The corresponding values of  $\lambda$ ,  $\mu$ , and  $\mu p^k$  follow from the defining relations (2.1) through (2.3). Suppose now that the above values of  $\lambda_1$ ,  $\lambda_+$ , and p do not satisfy constraint (3.1). Expression (4.3) is unimodal, so a maximum of (4.2) must occur where equality occurs in (3.1). Substitute  $\lambda_{1=\lambda_+}z_1$  in (4.2) and maximize the result with respect to  $\lambda_+$ , say at  $\lambda_+$ . Then

 $L(\lambda_{+}^{*}z_{1}, \lambda_{+}^{*}, p|data)$ 

$$= C \frac{z_1^{a_1+n_1-1}}{\left[ (b_1+t)z_1+(b_++t) \right]^{a_1+n_1+a_++n_+-2}} \frac{p^{s+c-1}q^{mn_+-s+d-1}}{(1-q^m-mpq^{m-1})}$$
  
= C  $\frac{p^{s+a_1+n_1+c-2}q^{m(n_++n_1+a_1-1)+d-1-(s+a_1+n_1-1)}}{\left[ (b_1+t)z_1+(b_++t) \right]^{a_1+n_1+a_++n_+-2}(1-q^m-mpq^{m-1})^{n_++n_1+a_1-1}}$ 

Here C is some constant. This must be maximized with respect to p. If  $b_1 = b_+$ , the expression simplifies to the following, with  $s'=s+a_1+n_1-1$ ,  $n'=n_++n_1+a_1-1$ :

$$C \frac{p^{s'+c-1} q^{mn'-s'+d-1}}{(1-q^{m})} \left[ \frac{1-q^{m}}{1-q^{m}-mpq^{m-1}} \right]^{a_{+}-1}$$

This must be maximized numerically with respect to p. (Note, if  $c=d=a_1=a_+=1$ , then this was the expression which was maximized to get the MLE.)

With this p used to calculate  $z_1$ , the maximizing  $\lambda_+$  is  $\lambda_+ = (a_1 + a_+ + n_1 + n_+ - 2)/[(b_1 + t)z_1 + (b_+ + t)]$ , and then  $\lambda_1 = \lambda_+ z_1$ .

4.2.2 <u>Posterior Means</u>. A more difficult estimate to obtain is the triple of posterior means. The difficulty arises from having to integrate (4.3) over the region satisfying (3.1) rather than over the region  $\lambda_1 \ge 0$ ,  $\lambda_+ \ge 0$ ,  $0 \le p \le 1$ .

The integrals needed are usually of the following forms. For A > 0 and B > 0, define

$$I(A,B,x) = \frac{B^{A}}{\Gamma(A)} \int_{0}^{x} e^{-Bt} t^{A-1} dt$$
 (4.4)

the gamma(A,B) cumulative distribution function. Note that  $I(A,B,\infty) = 1$  for any A and B. Now assume that A, B, A', and B' are all positive, and define

$$J(A,B,A',B',x) = \int_{0}^{x} \int_{s}^{\infty} \frac{B^{A}}{\Gamma(A)} e^{-Bt} t^{A-1} \frac{(B')^{A'}}{\Gamma(A')} e^{-B's} s^{A'-1} dt ds.$$
(4.5)

In Section 8.3, formulas are given for evaluating I(A,B,x) and J(A,B,A',B',x) as finite sums or infinite series. The formulas are simplest if A and A' are integers, somewhat narder to evaluate if A and A' are integers plus 1/2, and hardest to evaluate if A and A' are arbitrary. In practice, the formulas for evaluating I and J should be written into computer subprograms.

(4.6)

To express the results compactly, let us also define

$$f(p) = C_0 \frac{p^D q^E}{(1-q^m - mpq^{m-1})^F}$$

where

$$D = s+c-1$$

$$E = mn_{+}-s+d-1$$

$$F = n_{+}$$

$$C_{0} = normalizing constant.$$

Let

$$A_1 = a_1^{+n} A_+ = a_+^{+n} A_+$$

 $B_1 = (b_1^{+t})z_1$  $B_+ = b_+^{+t}$ .

With this notation defined, the results are easily expressed. Let us begin by obtaining L(p|data). This is done by integrating (4.2), first with respect to  $\lambda_1$  (making the change of variables  $u = \lambda_1/z_1$ ), and then with respect to  $\lambda_+$ . Integration is over the region satisfying (3.1). Then

$$L(p|data) = f(p) J(A_1, B_1, A_+, B_+, \infty).$$
(4.7)

The quantity J depends on p through  $B_1$ . The constant  $C_0$  in f(p) is such that  $\int L(p|data) dp = 1$ . It must be found numerically.

Once  $C_0$  is found, p L(p|data) can be integrated numerically to get the posterior mean of p. To find the mean of  $\lambda_1$ ,  $\lambda_+$ ,  $\mu$ , or  $\mu p^k$ , multiply (4.2) by the appropriate quantity and integrate. Integration with respect to  $\lambda_1$  and  $\lambda_+$  resembles the integration to obtain (4.7), and gives an expression of the form

$$C_1 f(p) J(A_1, B_1, A_+, B_+, \infty)$$
 (4.8)

with redefined values for some of the constants. The changes in the constants are

for  $\lambda_1 = a_1 + n_1 + 1$ 

for  $\lambda_{+} = a_{+} + n_{+} + 1$ 

for  $\mu$  A<sub>+</sub> = a<sub>+</sub> + n<sub>+</sub> + 1, F = n<sub>+</sub> + 1

for  $\mu p^k A_+ = a_+ + n_+ + 1$ , D = s+c-1+k,  $F = n_+ + 1$ .

The multiplicative constant  $C_1$  is  $(a_1+n_1)/(b_1+t)$  for  $\lambda_1$ , and  $(a_++n_+)/B_+$  for  $\lambda_+$ ,  $\mu$ , and  $\mu p^k$ . The constant  $C_0$  remains the same throughout, the

value needed to make  $\int L(p|data)dp = 1$ . The desired posterior mean is found by integrating (4.8) numerically with respect to p.

The posterior mean of  $\lambda$  is given by

$$\mathbb{E}\left[\lambda_{1}/m\right] - \mathbb{E}\left[\lambda_{+} p q^{m-1}/(1 - q^{m} - m p q^{m-1})\right].$$

E  $\lambda_1$  has just been found. The second term is found in the same way, using

 $A_{+} = a_{+} + n_{+} + 1, \quad D = s + c,$ 

 $E = mn_{+} - s + d + m - 2$ ,  $F = n_{+} + 1$ ,

and multiplicative factor  $C_1 = (\alpha_+ + n_+)/B_+$ .

#### 4.3 Bayes Probability Regions

4.3.1 Joint Region for Three Parameters. A three-dimensional region R is sought such that  $P[(\lambda_1, \lambda_+, p) \in R|data]$  equals some specified value. There are many such regions R. One approach is to try to treat the three parameters one at a time, so that R will, to some extent, resemble the confidence region of Section 3. Another approach, following Box and Tiao<sup>5</sup>, Sec. 2.8, is to choose R such that the posterior density is greater at any point inside R than at any point outside R. This is called the highest posterior density (HPD) region. While this second approach is natural in principle, it requires numerical triple integration, so is awkward in practice. Moreover, if such a region were found it would be shaped like a highly distorted ellipsoid and so would be difficult to describe. In particular, it would not be defined by any simple set of equations. For these reasons, the first approach is the only one which will be followed for a three-dimensional region.

The posterior density of p is given by (4.7), so for any desired  $\alpha_p$  between 0 and 1, an interval  $I_p$  can be found numerically such that  $P[p \in I_p | data] = 1 - \alpha_p$ . This interval may be the interval with equal tail probabilities, or the HPD interval for p, or another interval.
Now use

$$L(\lambda_{+}|p, data) = \int L(\lambda_{1}, \lambda_{+}, p|data) d\lambda_{1}/L(p|data)$$

integrating over  $\lambda_1$  satisfying (3.1), to obtain

$$L(\lambda_{+}|p, data) = \frac{B_{+}^{A_{+}}}{\Gamma(A_{+})} e^{-B_{+}\lambda_{+}} \lambda_{+}^{A_{+}-1} \left[1 - I(A_{1}, B_{1}, \lambda_{+})\right] / J(A_{1}, B_{1}, A_{+}, B_{+}, \infty).$$

The constants  $A_1$ ,  $B_1$ ,  $A_+$ , and  $B_+$  are as defined below (4.6). Therefore, for any p, an interval  $I_+(p)$  can be found numerically such that  $P[\lambda_+ \in I_+(p)|p, data] = 1 - \alpha_+$ . This interval may be such that there is equal probability (conditional on p and the data) that  $\lambda_+$  is on each side of  $I_+(p)$ . Or, the interval may be the conditional analogue of an HPD interval, chosen now so that  $L(\lambda_+|p, data)$  is highest in  $I_+(p)$ . Or the interval may be chosen in some other way.

Finally, from (4.2), the conditional density of  $\lambda_1$  is

$$L(\lambda_{1}|\lambda_{+}, p, data) = \frac{(b_{1}^{+t})^{A_{1}}}{\Gamma(A_{1})} e^{-\lambda_{1}(b_{1}^{+t})} \lambda_{1}^{A_{1}-1} / \left[1 - I(A_{1}, B_{1}, \lambda_{+})\right]$$

for  $\lambda_1 \ge \lambda_+ z_1$ .

So an interval  $I_1(\lambda_+, p)$  can be found such that  $P[\lambda_1 \in I_1(\lambda_+, p) | \lambda_+, p, data] = 1 - \alpha_1$ .

The resulting three-dimensional region has posterior probability  $(1-\alpha_1) (1-\alpha_p)$ .

If desired, this three-dimensional region can be re-expressed as a region for  $(\lambda, \mu, p)$ .

4.3.2 Intervals for Single Parameters. Let us find a probability interval for  $\mu p^k$ . For any  $c \ge 0$ ,

$$\begin{split} & \mathbb{P}\left[\mu p^{k} \leq c \left| \text{data} \right] = \mathbb{P}\left[\frac{\lambda_{+} p^{k}}{(1 - r_{0} - r_{1})} \leq c \left| \text{data} \right] \\ & = \mathbb{P}\left[\lambda_{1} \geq z_{1} \lambda_{+}, \lambda_{+} \leq c(1 - r_{0} - r_{1})/p^{k}, 0$$

Denote  $(1-r_0-r_1)/p^k$  by u(p). Make the change of variables  $t = \lambda_1/z_1$ . Then, using the notation (4.5) and (4.6), the above probability equals

$$\int_{0}^{1} J[A_{1}, B_{1}, A_{+}, B_{+}, cu(p)] f(p) dp.$$
(4.9)

The expression J may be evaluated using (8.10) through (8.13). Even if the evaluation requires summing an infinite series, this is probably faster than performing a double or triple integral numerically. So an interval [c,d] can be found numerically such that  $P[c \leq \mu p^k \leq d|data]$  equals any desired value 1- $\alpha$  between 0 and 1. The interval can be chosen so that

$$P[\mu p^{k} < c|data] = P[\mu p^{k} > d|data] = \alpha/2.$$

This gives the interval with equal tail probabilities. If instead the HPD interval is desired, the posterior density of  $\mu p^{k}$  must be found by differentiating (4.9) with respect to c. By the definitions (4.5) and (4.4),

$$L_{\mu p} k (c|data) = \int_{0}^{1} \frac{\partial}{\partial c} J[A_{1}, B_{1}, A_{+}, B_{+}, cu(p)] f(p)$$
  
= 
$$\int_{0}^{1} \frac{B_{+}^{A_{+}}}{\Gamma(A_{+})} e^{-B_{+}cu(p)} [cu(p)]^{A_{+}} [1 - I(A_{1}, B_{1}, cu(p))] u(p) f(p) dp.$$

Evaluation of this density requires use of one of (8.5) through (8.7) followed by numerical integration, but when this is done the HPD interval for  $\mu p^{k}$  can be found.

An interval for  $\mu$  follows from the above by setting k = 0. An interval for  $\lambda_{+}$  is obtained by replacing u(p) by 1 in (4.9). An interval for  $\lambda_{1}$  can be found by using

$$P[\lambda_{1} \leq c | data]$$

$$= P[z_{1} \lambda_{+} \leq \lambda_{1} \leq c, 0 
$$= P[\lambda_{+} \leq c/z_{1} | data]$$

$$- P[\lambda_{1} > c, \lambda_{+} \leq c/z_{1}, 0 
$$= \int_{0}^{1} J(A_{1}, B_{1}, A_{+}, B_{+}, c) f(p) dp$$

$$- \int_{0}^{1} [1 - I(A_{1}, B_{1}, c/z_{1})] I(A_{+}, B_{+}, c/z_{1}) f(p) dp.$$$$$$

Numerical integration can be used to evaluate this.

An interval for  $\lambda$  can be found using

$$\begin{split} & P[\lambda \leq c|data] \\ &= P[\lambda_{1} - \lambda_{+} | z_{1} \leq mc|data] \\ &= 1 - P[\lambda_{1}/z_{1} > \lambda_{+} + mc/z_{1}, | 0$$

This must be evaluated by numerical double integration.

Two-dimensional probability regions for pairs of the parameters will not be given.

### 5. DIAGNOSTIC CHECKS ON THE MODEL

### 5.1 Poisson Parameters

Suppose that there are I "sources" of data. These sources could be plants, vendors, time periods, etc. Let us investigate whether the sources all have the same values of  $\lambda_1$  or  $\lambda_+$ . The methods below do not use the binomial distribution assumptions at all. The only assumption is that the single failures and multiple failures form two Poisson processes.

5.1.1 <u>Graphical Methods</u>. Denote the  $\lambda_+$  corresponding to the ith source by  $\lambda_{+i}$ . Suppose that all the sources but the ith have a common  $\lambda_+$ , denoted  $\lambda_{+,-i}$ . Let  $N_{+i}$  and  $t_i$  be the number of multiple failures and the operating time for the ith source.

To test the null hypothesis  $\lambda_{+i} = \lambda_{+,-i}$ , a uniformly most powerful similar test is based on  $N_{+i}$  given  $N_{+}$ . The conditional distribution of  $N_{+i} | N_{+} = n_{+}$  is binomial $(n_{+}, t_{i}/t)$  under the null hypothesis. (These assertions are all shown in Cox and Hinkley<sup>4</sup>, pp. 136-7.)

One can therefore examine either residuals or the corresponding significance levels. The ith standardized residual is defined as

$$R_{i} = \frac{N_{i} - n_{t}t_{i}/t}{\left[n_{t} \frac{t_{i}}{t} \left(1 - \frac{t_{i}}{t}\right)\right]^{1/2}}$$

If  $\lambda_{+i}$  is the same for all i, then for a given  $n_+$ , the  $R_i$ 's all have mean 0 and variance 1. Define the ith significance level  $\alpha_i$  as

$$Min(2P[N_{+j} \ge n_{+j} | N_{+}=n_{+}], 2P[N_{+j} \le n_{+j} | N_{+}=n_{+}], 1).$$

That is, the ith significance leve! is  $2P[N_{+i} \ge n_{+i} | N_{+}=n_{+}]$  if the observed  $n_{+i}$  is greater than the median, and it is similarly defined if  $n_{+i}$  is less than the median or equal to the median. The tail probability

is doubled to reflect the fact that  $N_{+j}$  could be, a priori, either large or small.

Outliers correspond to large values of  $R_i$  or to small values of  $a_i$ . The standardized residuals can be graphed, but they have a skewed distribution, so a glance at the plot will not necessarily pick out the values which are significantly large or small. For this, the individual significance levels are preferable.

These methods can be used to identify sources for which  $\lambda_{+i}$  seems to be substantially different from the others. They cannot be used for an overall test of whether all the  $\lambda_{+i}$ 's are equal, for if the number of sources is large, then random variability alone will produce some apparently extreme values of  $N_{+i}$ . Testing is discussed below.

For investigating  $\lambda_1$ , exactly the same techniques work, replacing  $N_{+i}$  and  $n_+$  by  $N_{1i}$  and  $n_1$ .

5.1.2 Hypothesis Tests. Let us test

$$H_0: N_{+i} \sim Poisson(\lambda_+ t_i), i = 1,..,I$$

H<sub>1</sub>: N<sub>+i</sub> ~ Poisson( $\lambda_{+i}$ ), i = 1,..., I, with no relation among the  $\lambda_{+i}$ 's.

If the sample size  $n_{+}$  is large, the generalized likelihood ratio test can be used. (See Mood, Graybill, and Boes<sup>6</sup>, Sec. IX.5.1.) The test statistic is

$$-2 \log \Lambda = 2 \sum_{i=1}^{I} N_{i} \log \left( \frac{N_{i}}{N_{i}} \frac{t}{t_{i}} \right).$$
 (5.1)

The null hypothesis should be rejected if (5.1) is greater than the  $1-\alpha$  point of a  $\chi^2$  distribution with I-1 degrees of freedom.

This is exactly the same test as would be used to test

$$H_{0}: (N_{+1},..,N_{+1}) | n_{+} \sim multinomial(n_{+}, t_{1}/t,..,t_{1}/t).$$
(5.2)

If  $n_{+}$  is not large, the test statistic (5.1) can still be used, but since  $\lambda_{+}$  is unknown the critical point should be derived from the multinomial distribution (5.2). The exact distribution can be calculated for small I and  $n_{+}$  by using the program POLYPOW, presented by Atwood<sup>7</sup>. This works because the multinomial probabilities are the terms in the algebraic expansion of

$$(t_1/t + ... + t_1/t)^{n_+}$$
,

and POLYPOW finds these terms.

The  $x^2$  test statistic

$$\sum \frac{(N_{+i} - n_{+}t_{i}/t)^{2}}{n_{+}t_{i}/t}$$

could be used instead of the likelihood ratio test statistic.

For studying  $\lambda_1$ , simply replace N<sub>+i</sub> and N<sub>+</sub> by N<sub>1i</sub> and N<sub>1</sub>.

5.2 Binomial Parameter

Suppose again that there are I sources of data, which may be plants, etc., as before, but may now also be the  $n_+$  individual observations of multiple failures. Let us investigate whether the I sources have the same p, and whether the data as a whole seem consistent with the assumption of a binomial distribution. 5.2.1 <u>Graphical Methods</u>. Let the random variable X be the number of failed components in a multiple failure. (We have been assuming that X is truncated binomial(m,p), truncated so that  $X \ge 2$ .) Let  $S_i$  denote S (i.e., the total number of failed components in multiple failures) based on observations from only the ith source. So, conditional on  $n_{+i}$ ,  $S_i$  is a sum of  $n_{+i}$  independent observations of X. Denote S -  $S_i$  by  $S_{-i}$ . Then, conditional on  $n_{+i}$ , and  $n_{+,-i}$ ,  $S_i/n_{+i} - S_{-i}/n_{+,-i}$  has expectation 0 and variance

 $[1/n_{+i} + 1/n_{+,-i}]$  var X.

Define

$$R_{i} = \left[S_{i}/n_{+i} - S_{-i}/n_{+,-i}\right] / \left[1/n_{+i} + 1/n_{+,-i}\right]^{1/2} .$$
(5.3)

If all sources correspond to the same distribution of X (in particular, if all sources obey the BFR assumptions with the same p), then, conditional on the  $n_{+i}$ 's, the  $R_i$  are identically distributed with mean 0 and variance = var X. If X is truncated binomial(m,p) with X  $\geq$  2, then direct calculation yields

$$EX = \frac{m p(1 - q^{m-1})}{1 - q^m - mpq^{m-1}}$$

$$EX(X-1) = \frac{m (m - 1) p^2}{1 - q^m - mpq^{m-1}}$$

and the variance of X is

$$\sigma_{\chi}^2 = EX(X-1) - EX(EX-1).$$

This may be estimated by substituting some estimate for p. The standardized residuals are then obtained by dividing  $R_i$  by  $\hat{\sigma}_{\chi}$ , the estimate of  $\sigma_{\chi}$ .

They are not independent, but a plot of i versus  $R_i/\hat{\sigma}_{\chi}$  will help point out sources with unusually high or low p.

The above method was used to identify sources with high or low numbers of failures. The following method investigates whether the distribution within a source is really truncated binomial. To simplify the notation. assume that there is a single data source, with  $n_{+}$  multiple failure occurrences. For  $2 \le i \le m$ , conditional on  $n_{+}$ , the number of instances of i failures,  $N_{i}$ , is binomial $(n_{+}, z_{i})$ , with  $z_{i}$  defined by (2.5) and (2.2). Let  $\hat{z}_{i}$  be the estimate based on the MLE  $\hat{p}$ , i.e.,

$$\hat{z}_{i} = {m \choose i} \hat{p}^{i} \hat{q}^{m-i} / (1 - \hat{q}^{m} - m\hat{p}\hat{q}^{m-1})$$
.

Then

$$U_{i} = \frac{N_{i} - n_{+} \hat{z}_{i}}{\left[n_{+} \hat{z}_{i}(1 - \hat{z}_{i})\right]^{1/2}}$$
(5.4)

has mean and variance approximately 0 and 1. Large or small values, or strong patterns, indicate that the distribution is not truncated binomial.

5.2.2 <u>Hypothesis Tests</u>. Let us test the null hypothesis that X is a truncated binomial(m,p) random variable truncated such that  $X \ge 2$ . Under this hypothesis,  $P[X = i] = z_i$ , in the notation of (2.5), for i=2,...,m.

Possible alternate hypotheses are that p varies from source to source, and that the distribution of X is the same for all sources but is not truncated binomial(m,p).

If the sample size is large, the generalized likelihood ratio test can be used. For testing against the alternate hypothesis that the sources are BFR but with possibly different p's, the test statistic is

$$-2 \log \Lambda = -2 L_0^+ 2 \sum_{i=1}^{I} L_i$$

where

$$L_{o} = s \log (\hat{p}/\hat{q}) + n_{+} \log \left[ \hat{q}^{m}/(1 - \hat{q}^{m} - m \hat{p} \hat{q}^{m-1}) \right]$$
(5.5)

and

$$L_{i} = s_{i} \log (\hat{p}_{i}/\hat{q}_{i}) + n_{+i} \log \left[ \hat{q}_{i}^{m} / (1 - \hat{q}_{i}^{m} - m \hat{p}_{i} \hat{q}_{i}^{m-1}) \right].$$
(5.6)

Here  $\hat{p}$  is the MLE based on all the data, and  $\hat{p}_i$  is the MLE based on only the ith source. Under H<sub>0</sub>, -2 log  $\Lambda$  is asymptotically  $\chi^2(I-1)$ .

To test against the alternate hypothesis that X is not truncated binomial(m,p), the test statistic is

$$-2 \log \Lambda = 2 \sum_{i=2}^{m} n_i \log (n_i/(n_+ \hat{z}_i))$$

where  $\hat{z}_i$  was defined above (5.4). Under  $H_0$ , the asymptotic distribution of -2 log  $\Lambda$  is  $\chi^2(m-4)$ . Another possible test in this case is the ordinary chi-square goodness of fit test. The test statistic is

$$\sum_{i=2}^{m} (n_{i} - n_{i}\hat{z}_{i})^{2}/n_{i}\hat{z}_{i}$$

and its asymptotic distribution under  $H_0$  is  $\chi^2(m-4)$ . If the sample size is only moderate rather than large, then the cells can be grouped and the degrees of freedom adjusted in the usual way.

## 6. RESULTS IF CAUSES OF SINGLE FAILURES ARE IDENTIFIABLE

### 6.1 Definitions

The data consist of  $(n_I, n_C, n_2, ..., n_m)$ , where  $n_I$  is the number of single failures that were not due to common cause shocks, and  $n_C$  is the number of single failures due to common cause shocks. Now m may be as small as 2. Define

$$n_{+}' = n_{C} + n_{+}$$

$$\lambda_{+}' = \mu r_{1} + \lambda_{+} = \mu (1 - r_{0})$$

$$s' = n_{C} + s$$

$$z_{i}' = r_{i} / (1 - r_{0}) \text{ for } i = 1, ..., m.$$

$$P[N_{I} = n_{I}, N_{C} = n_{C}, N_{2} = n_{2}, ..., N_{m} = n_{m}]$$
  
= 
$$P[N_{I} = n_{I}] P[N_{+}' = n_{+}'] P[N_{C} = n_{C}, N_{2} = n_{2}, ..., N_{m} = n_{m}|N_{+}' = n_{+}']$$

where

N<sub>I</sub> ~ Poissi mt)

 $N_{+}' \sim Poisson(\lambda_{+}'t)$ 

 $N_{C},..., N_{m} | N_{+}' = n_{+}' \sim multinomial(n_{+}', z_{1}',..., z_{m}').$ 

Inference follows the pattern of Sections 3 through 5, but is much simpler because the most easily estimable parameters,  $\lambda$ ,  $\lambda_+$ ', and p, are not constrained by any analogue of (3.1).

#### 6.2 Non-Bayesian Inference

The Poisson parameters  $\lambda$  and  $\lambda_+$ ' can be estimated, both by MLE's and by confidence intervals, in the standard way. The MLE for p is the solution of

 $s' = m n_{+}' p/(1 - q^{m}).$ 

[In the special case m = 2, the solution becomes simply  $\hat{p} = 2(s'-n_+')/s'$ .] Maximum likelihood estimates for any other parameters- $\lambda_1$ ,  $\lambda_+$ ,  $\mu$ , or  $\mu p^k$ -follow from substitution of  $\lambda$ ,  $\lambda_+$ , and p into the appropriate defining equations involving the desired quantities.

If  $n_{\star}' > 0$ , a confidence interval for p can be based on the conditional distribution of S' given  $N_{\star}'$ , using analogues of (3.5) and (3.6). This conditional distribution is now given. Let  $v = (v_1, ..., v_m)$ be any vector of nonnegative integers. Define

$$v_{+}' = \sum_{i=1}^{m} v_{i}$$

Define the set  $T_k$  by

$$T_{k} = \left\{ v \middle| v_{+}' = n_{+}', \begin{array}{c} m \\ \Sigma \\ i=1 \end{array} \right\}, \quad k \left\}.$$

Let

$$c(v) = \frac{v_{+}'!}{v_{1}!, \dots, v_{m}!} \prod_{i=1}^{m} {\binom{m}{i}}^{v_{i}}.$$

Then, for  $n_{+} \leq k \leq m n_{+}$ ,

$$P\left[S' = k | N_{+}' = n_{+}'\right] = \sum_{v \in T_{k}} P\left[N_{C} = v_{1}, N_{2} = v_{2}, \dots, N_{m} = v_{m} N_{+}' = n_{+}'\right]$$

$$= \frac{ps' q}{(1 - q^m)^{n+'}} \sum_{v \in T_k} c(v) .$$

If the interval [0,1] is used when  $N_{+}' = 0$ , then the procedure has unconditional confidence level  $(1-\alpha)$ .

A joint confidence region for more than one of  $\lambda$ ,  $\lambda_{+}$ ', and p is simply the product of the confidence intervals. Conservative intervals for  $\mu$  and  $\mu p^{k}$  can be based on the intervals for  $\lambda_{+}$ ' and p in the obvious ways. For  $\lambda_{1}$  and  $\lambda_{+}$ , conservative intervals can be based on the intervals for  $\lambda$ ,  $\lambda_{+}$ ', and p. Alternatively, the confidence intervals of Section 3.2 can still be used for  $\lambda_{1}$  and  $\lambda_{+}$ .

If  $n_{I}$  and  $n_{+}$ ' are large, then  $\hat{\lambda}$ ,  $\hat{\lambda}_{+}$ ', and  $\hat{p}$  are asymptotically independent normal with means  $\lambda$ ,  $\lambda_{+}$ ', and p, and with variances  $\lambda/mt$ ,  $\lambda_{+}$ '/t and 1/I(p), where

$$I(p) = \frac{mn_{+}}{pq} \frac{1 - q^{m} - mpq^{m-1}}{(1 - q^{m})^{2}}.$$
 (6.1)

Approximate confidence intervals can be based on this asymptotic distribution.

### 6.3 Bayesian Inference

We will consider the class of prior distributions:  $\lambda \sim \Gamma(a,b)$ ,  $\lambda_{+}' \sim \Gamma(a_{+}',b_{+}')$ ,  $p \sim beta(c,d)$ , with  $\lambda$ ,  $\lambda_{+}'$  and p independent. If  $a=a_{+}'=1/2$ ,  $b=b_{+}'=0$ , then the (improper) priors are noninformative for  $\lambda$ and  $\lambda_{+}'$ . The noninformative prior advocated by Box and Tiao<sup>5</sup> for p is proportional to  $I^{1/2}(p)$ , with I(p) given in (6.1). If c=d=1/2, the beta(c,d) distribution approximates the noninformative prior. Tables 3 and 4 give values of c and d that provide better approximations, and Figures 4 through 6 show these cumulative distributions for m = 5, 20, and 100.

	d
2	0.3541
3	0.3776
4	0.3923
5	0.4027
6	0.4105
7	0.4167
8	0.4217
9	0.4260
10	0.4296
12	0.4355
14	0.4401
16	0.4439
18	0.4470
20	0.4496
25	0.4549
30	0.4588
40	0.4642
60	0.4708
80	0.4748
100	0.4775
150	0.4817
200	0.4842
300	0.4872
00	0.5000

TABLE 3. BINOMIAL WITH O TRUNCATED: APPROXIMATELY NONINFORMATIVE d FOR BETA(1/2, d)

	m	C	d
	2	0.5902	0.4179
	3	0.6051	0.4569
	4	0.6101	0.4786
	5	0.6108	0.4919
	6	0.6096	0.5004
	7	0.6075	0.5063
	8	0.6050	0.5103
	9	0.6024	0.5132
	10	0.5998	0.5154
	12	0.5949	0.5181
	14	0.5904	0.5197
	16	0.5864	0.5205
	18	0.5828	0.5210
	20	0.5795	0.5212
	25	0.5728	0.5211
	30	0.5675	0.5207
	40	0.5597	0.5196
	60	0.5497	0.5177
	80	0.5436	0.5162
1	.00	0.5394	0.5151
1	.50	0.5327	0.5132
2	200	0.5287	0.5120
3	00	0.5240	0.5106
	80	0.5000	0.5000

# TABLE 4. BINOMIAL WITH O TRUNCATED: APPROXIMATELY NONINFORMATIVE PARAMETERS FOR BETA(c,d)



truncated, m = 5.

44





truncated, m = 100.

The posterior density factors into the product

 $L(\lambda, \lambda_{+}', p|data) = L(\lambda|data) L(\lambda_{+}'|data) L(p|data)$ 

where

$$\lambda$$
 data ~  $\Gamma(a + n_1, b + mt)$ 

$$\lambda_{+}' | data \sim \Gamma(a_{+}' + n_{+}', b_{+}' + t)$$

and

$$L(p|data) = C \frac{n_{+}' !}{n_{C}', \dots, n_{m}'} {\binom{m}{1}}^{n_{C}} \frac{m}{\prod_{i=2}^{m}} {\binom{m}{i}}^{n_{i}} \frac{p^{s'+c-1} q^{m_{+}'-s'+d-1}}{(1-q^{m})^{n_{+}'}}.$$

Bayes point and interval estimation of  $\lambda$ ,  $\lambda_{+}$ ', and p are routine, using numerical integration to treat p. Point estimates of  $\lambda_{1}$ ,  $\lambda_{+}$ ,  $\mu$ , and  $\mu p^{k}$  corresponding to the mode of the posterior density follow from the equations relating the parameters, and the posterior means of these quantities are obtained by straightforward integration.

A probability interval for  $\mu p^k$  can be found as in Section 4.3.2:

$$F\left[2 \leq up^{k} \leq d \; data\right]$$

$$= \int_{0}^{1} P\left[c(1-q^{m})/p^{k} \leq \lambda_{+}' \leq d \; (1-q^{m})/p^{k} \mid p, data\right] L(p|data) \; dp$$

$$= \int_{0}^{1} P\left[cu(p) \leq \lambda_{+}' \leq d \; u(p) \mid p, data\right] L(p|data) \; dp$$

with  $u(p) = (1-q^m)/p^k$ . In the notation of (4.4), this equals

$$\int_{0}^{1} \left[ I(A_{+}', B_{+}', d u(p)) - I(A_{+}', B_{+}', c u(p)) \right] L(p|data) dp$$

with  $A_{+}' = a_{+}' + n_{+}'$  and  $B_{+}' = b_{+}' + t$ . The integrand may be evaluated using (8.6) through (8.8), and the integral may be evaluated numerically.

For  $\mu$ , the method is the same, with k=G. For

$$\lambda_{+} = \lambda_{+}' (1 - q^{m} - mpq^{m-1})/(1 - q^{m}),$$

the method is the same as for  $\mu p^k$ , but now set  $u(p) = (1-q^m)/(1-q^m-mpq^{m-1})$ .

For

$$\lambda_1 = m\lambda + \lambda_+' (mpq^{m-1})/(1-q^m),$$

the method is similar, but integration is with respect to  $\lambda$  and  $\lambda_+$ '. The details are messy and will not be given. If N<sub>I</sub> and N<sub>C</sub> are identifiable, then  $\lambda_1$  is a less natural parameter than  $\lambda$  and may not be of interest to the user.

### 6.4 Diagnostic Checks

The checks for the Poisson parameters in Section 6 all carry over if  $N_{+i}$ ,  $n_+$ ,  $\lambda_{+i}$ , and  $\lambda_+$  are replaced by  $N_{+i}$ ',  $n_+$ ',  $\lambda_{+i}$ ', and  $\lambda_+$ ', and if  $N_{1i}$ ,  $n_1$ ,  $\lambda_{1i}$ , and  $\lambda_1$  are replaced by  $N_{1i}$ ,  $n_1$ ,  $\lambda_i$ , and  $\lambda$ .

The checks for the binomial parameter p carry over under the following translation. Now, X takes values from 1 to m. Primes should be given to S,  $S_i$ ,  $S_{-i}$ ,  $n_{+i}$ ,  $n_{+,-i}$ , and  $\hat{z}_i$ . Equations (5.5) and (5.6) must be replaced by

$$\begin{split} & L_{o} = s' \log (\hat{p}/\hat{q}) + n_{+}' \log \left[ \hat{q}^{m}/(1-\hat{q}^{m}) \right] \\ & L_{i} = s_{i}' \log (\hat{p}_{i}/\hat{q}_{i}) + n_{+i}' \log \left[ \hat{q}_{i}^{m}/(1-\hat{q}_{i}^{m}) \right]. \end{split}$$

To test against the alternate hypothesis that X is not truncated binomial, the likelihood ratio test statistic is

$$-2 \log \Lambda = 2n_{C} \log \left[ n_{C} / (n_{+}, \hat{z}_{1}) \right] + 2 \prod_{i=2}^{M} n_{i} \log \left[ n_{i} / (n_{+}, \hat{z}_{i}) \right]$$

where

$$\hat{z}_{i}' = \binom{m}{i} \hat{p}^{i} \hat{q}^{m-i} / \left[1 - \hat{q}^{m}\right].$$

Under  $H_0$ , the asymptotic distribution is  $\chi^2(m-3)$ . The  $\chi^2$  test statistic

$$\frac{(n_{C} - n_{+}' \hat{z}_{1}')^{2}}{n_{+}' \hat{z}_{1}'} + \sum_{i=2}^{m} \frac{(n_{i} - n_{+}' \hat{z}_{i}')^{2}}{n_{+}' \hat{z}_{i}'}$$

may be used instead of the likelihood ratio test statistic.

### 7. ILLUSTRATIVE EXAMPLE

### 7.1 Data Used

Vesely<sup>1</sup> gives data from 20 U.S. commercial boiling water reactors. The components are control rods, and a failure may be defined in at least two ways: failure to insert past notch 04, or any failure to meet technical specifications. The first definition includes failures that are serious enough to affect safety. The second also includes incidents such as slow rod insertion.

This data set is used for illustrative purposes only! Much more data has become available since the data set of Reference 1 was published. The treatment presented here is also noive in that it ignores the event descriptions given in the reports. These descriptions might suggest qualitative differences among the events, which would lead a careful analyst to consider portions of the data separately. Finally, the diagnostic checks point out inhomogeneity among the plants, and one failure occurrence which is a clear outlier. These are not investigated further here. A more thorough analysis of much more extensive data is now underway at the Idaho National Engineering Laboratory. Those who are interested in the answers, not merely in the method, must refer to the INEL reports that will appear. The numerical results given below serve only to illustrate the method.

The model of this paper assumes that m, the number of components, is constant. In Reference 1 the number of rods in a plant varies from 32 to 185. We will consider plants with 177 or 185 rods. The data for these eight plants are summarized in Table 5. The effect of pooling data with these two values of m will be discussed below.

The causes of single failures will be considered as not identifiable.

Reactor	Number of Rods (m)	Total Operating Months (t) <sup>a</sup>	Failures to Insert Past Notch 04	All Incidents
Dresden 2	177	41	n <sub>3</sub> =1	n <sub>96</sub> =1
				n <sub>2</sub> =2
				n <sub>1</sub> =1
Dresden 3	177	61	n <sub>1</sub> =1	n <sub>3</sub> =1
Quad-Cities 1	177	51	None	None
Quad-Cities 2	177	49	None	None
Peach Bottom 2	185	30	None	None
Peach Bottom 3	185	25	None	None
Browns Ferry 1	185	20	None	None
Browns Ferry 2	185	13	None	None
		290		

### TABLE 5. SUMMARY OF FAILURE DATA

a. Operating time in hours assumes 720 hours per month.

### 7.2 Analysis of Failures to Insert Past Notch 04

For this data set,  $n_1 = 1$ ,  $n_+ = 1$ , s = 3, t = 208800 hours, and estimates will be calculated using both m = 177 and m = 185. Estimates will be found for  $\lambda_1$ ,  $\lambda_+$ ,  $\lambda$ ,  $\mu$ , p, and  $\mu p^2$ . The last quantity is an example of  $\mu p^k$ .

Maximum likelihood estimates and 95% confidence intervals are given in Table 6. All the confidence intervals are two-sided (probability 0.025 for

	<u>.</u>	m = 177		m = 185
Parameter	MLE	Confidence Interval	MLE	Confidence Interval
р	0.0122	(0.00043, 0.0483)	0.0117	(0.00041, 0.0462)
$10^6 \times \lambda_1$	4.79	(0.121, 26.7)	4.79	(0.121, 26.7)
$10^6 \times \lambda_+$	4.79	(0.121, 26.7)	4.79	(0.121, 26.7)
10 <sup>8</sup> х х	1.65	(0, 12.8)	1.58	(0, 12.8)
10 <sup>6</sup> × μ	7.51	(0.061, 42248)	7.51	(0.061, 42468)
$10^9 \times \mu p^2$	1.12	(0.0040, 87.4)	1.02	(0.0037, 80.1)

TABLE 6. MAXIMUM LIKELIHOOD ESTIMATES AND 95% CONFIDENCE INTERVALS

each tail), except for the interval for  $\lambda$ . The interval for  $\lambda$  is the conservative one-sided interval based on  $\lambda \leq \lambda_1/m$  (probability 0.05 for the upper tail). The two-sided interval for  $\lambda$  that is based on  $\lambda_1$ ,  $\lambda_+$ , and p, and is described in Section 3.2.5, turns out also to have its lower end point at 0, and the interval is strictly larger than the interval of Table 6. The intervals for  $\mu$  and  $\mu p^2$  are based on two-sided intervals for  $\lambda_+$  and p with  $(1-\alpha)(1-\alpha_p) = 0.95$  and  $\alpha_+ = \alpha_p$ .

Bayesian point and interval estimates are given in Table 7. The intervals all have posterior probability 0.025 in each tail. Highest posterior density intervals are not shown because the computer programming is not yet complete. An interval for  $\lambda$  is not given because of the lengthy computation required. The prior distribution is noninformative for  $\lambda_1$  and  $\lambda_+$ . The first two portions of the table use a beta prior distribution uses a beta(0.5, 0.5) distribution.

A comparison of the first and third sections of Table 7 shows that the entries change very little, whether the prior beta distribution has

	m =	177, p ~ beta(0.552,	0.519)
Parameter	Posterior Mode	Posterior Mean	Interval
p	0.00769	0.0189	(0.00418, 0.0439)
$10^6 \times \lambda_1$	2.39	8.29	(0.923, 23.7)
$10^6 \times \lambda_+$	2.39	6.07	(0.411, 19.6)
10 <sup>8</sup> × λ	0.156	4.05	
10 <sup>6</sup> × µ	6.06	9.01	(0.626, 29.3)
$10^9 \times \mu p^2$	0.358	3.55	(0.0836, 18.4)
	m =	185, p ~ beta(0.551,	0.519)
р	0.00734	0.0180	(0.00399, 0.0420)
10 <sup>6</sup> × × <sub>1</sub>	2.39	8.30	(0.923, 23.7)
10 <sup>6</sup> x x <sub>+</sub>	2.39	6.07	(0.411, 19.6)
10 <sup>8</sup> х х	0.145	3.88	흔, 그 국가학
10 <sup>6</sup> × µ	6.08	9.01	(0.626, 29.3)
$10^9 \times \mu p^2$	0.327	3.25	(0.0765, 16.8)
	n	n = 177, p ~ beta(0.5,	0.5)
р	0.00710	0.0186	(0.00404, 0.0435)
$10^6 \times \lambda_1$	2.40	8.33	(0.937, 23.8)
$10^6 \times \lambda_+$	2.40	6.04	(0.407, 19.5)
10 <sup>8</sup> x x	0.00	4.05	-
10 <sup>6</sup> x µ	6.68	9.08	(0.630, 29.6)
$10^9 \times \mu p^2$	0.337	3.46	(0.0804, 18.0)

TABLE 7. BAYESIAN POINT ESTIMATES AND 95% INTERVALS

approximately noninformative parameters or parameters 0.5 and 0.5. As shown in Figures 1 through 3, the approximately roninformative beta distribution is "closer" to the noninformative prior than to the beta(0.5, 0.5) distribution. It therefore seems unlikely that the table entries would change much if the prior distribution were the noninformative prior based on the information matrix. This indicates that the beta approximation to the noninformative prior is adequate. Investigation of this point when m is small will be carried out at a later date.

In Section 3.1, it was mentioned that the MLE p decreases as m increases, so a more conservative (larger) estimate p will result from using m = 177, rather than m = 185. Tables 6 and 7 show that in this example, it is more conservative by any criterion to use m = 177 rather than m = 185, since doing so gives larger estimates of  $\lambda$ , p, and  $\mu p^2$ , and identical or virtually identical estimates of  $\lambda_1$ ,  $\lambda_+$ , and  $\mu$ . The maximum likelihood estimates and posterior means for p and for  $\lambda$  are approximately proportional to 1/m.

In every case, the maximum likelihood estimate lies between the Bayes posterior mode and the Bayes posterior mean. In problems with a single binomial or Poisson parameter, it can be shown directly that this relation holds when the noninformative prior is used. It is interesting that it also holds for every parameter in the present multiparameter problem.

For  $\lambda_1$ ,  $\lambda_+$ , and p, the Bayes intervals are shorter than the corresponding confidence intervals. In problems with a single binomial or Poisson parameter, it can be shown directly that the Bayesian interval based on the noninformative prior must be strictly shorter than the corresponding confidence interval. This is because confidence intervals based on discrete data are inexact, with the (unknown) true confidence level being greater than or equal to the nominal level. In the present multiparameter problem, the same relation holds between the sizes of the confidence interval and the Bayes interval, presumably for the same reason. For  $\mu$  and  $\mu p^2$ , the Bayes intervals are also shorter than the confidence intervals. This is due both to the reason just mentioned and to the fact

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that these confidence intervals are conservative rather than sharp. The Bayes interval for  $\lambda$  is not shown because it requires lengthy computation.

Figure 7 shows a 90% confidence region for  $(\lambda_+, \lambda_1)$ , given three values of p. The values of p are the MLE and the ends of the 95% confidence interval for p. For each p, the portion of the square which is above the corresponding line is that portion satisfying the constraint (3.1). Figures 8 through 11 show 90% Bayes regions for  $(\lambda_+, \lambda_1)$ , given various values of p. The values of p are the posterior mode, the posterior mean, and the end points of the 95% probability interval for p. The Bayesian and non-Bayesian regions are not directly comparable, since they assume different values of p. However, in all the figures, the estimates of  $\lambda_+$  and  $\lambda_1$  are positively correlated, due to the constraint (3.1), and the correlation is strongest for small p.



Figure 7. Ninety percent confidence region for  $(\lambda_+, \lambda_1)$ , given p. For each p, the region is that portion of the rectangle above the line corresponding to p.

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Figure 10. Ninety percent Bayes probability region for  $(\lambda_+, \lambda_1)$ , given p = 0.0189 = poster or mean.



### 7.3 Diagnostic Checks

In order to have enough data to illustrate the methods for diagnostic checks, we will consider all five failure incidents.

Let us combine single and multiple failures, and investigate  $\lambda_1 + \lambda_+$ . Table 8 shows the standardized residuals and significance levels for the eight plants, and for the parameter  $\lambda_1 + \lambda_+$ . The first plant shows a small significance level, indicating that the plant seems to be anomalous. However, one would expect random variability alone to produce some apparent anomalies among many plants. To test whether the plants all have the same value of  $\lambda_1 + \lambda_+$ , the likelihood ratio statistic (5.1) can be calculated to be 13.76. If all the plants have the same value of  $\lambda_1 + \lambda_+$ , then

 $P(-2 \log \Lambda \ge 13.76 | N_1 + N_+ = 5) = 0.016.$ 

This probability is exact rather than asymptotic, and indicates that the plants did not have the same value of  $\lambda_1 + \lambda_+$ .

Plant	ti	Observed Number of Failures	Standardized Residual	Significance Level
Dresden 2	41	4	4.23	0.0035
Dresden 3	61	1	-0.06	1.0
Quad-Cities 1	51	0	-1.03	0.76
Quad-Cities 2	49	0	-1.01	0.79
Peach Bottom 2	30	0	-0.76	1.0
Peach Bottom 3	25	0	-0.61	1.0
Browns Ferry 1	20	0	-0.69	1.0
Browns Ferry 2	13	0	-0.48	1.0

TABLE 8. STANDARDIZED RESIDUALS AND SIGNIFICANCE LEVELS FOR  $\lambda_1 + \lambda_+$ 

As an illustration of diagnostic checks for the assumptions about p, let us consider the four multiple failure occurrences as the "sources." Then the values of  $n_{+i}$  and  $n_{+,-i}$  are 1 and 3 for each source. For this data set, the MLE  $\hat{p}$  is 0.14548. This agrees to five places with the estimate s/mn<sub>+</sub>, which would be obtained if the observed data were treated as binomial rather than truncated binomial. Since the truncation effect is negligible, an approximate 95% confidence interval for p has upper end at

$$\hat{p} + 2 \left[ \hat{p}\hat{q}/(mn_{+}) \right]^{1/2} = 0.1720.$$

The standardized residuals from (5.3) are given in Table 9, based on p = 0.1455 and p = 0.1720. Plausible magnitudes of standardized residuals are, say, less than 3. The magnitudes in Table 9 are so much greater than 3, whichever p is used, that it seems conclusive that the data do not come from a single BFR model.

Number of Failed	R <sub>i</sub> /Ĝ <sub>x</sub>		
Components $(=S_i)$	<u>p = 0.1455</u>	<u>p = 0.1720</u>	
96	17.29	16.16	
2	-5.85	-5.46	
2	-5.85	-5.46	
3	-5.60	-5.23	
103			

TABLE 9. STANDARDIZED RESIDUALS FOR p Each multiple failure treated as data source, using formula (5.3)

Even more dramatic results are seen if the data are treated as coming from a single source, and  $U_i$  is calculated from (5.4). The values are summarized in Table 10. Truly enormous quantities are obtained because of observed failure numbers with extremely small estimated probabilities.

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Number of Failed Components i	Number of Occurrences N <sub>i</sub>	$U_i = (N_i - n_+ \hat{z}_i) / [n_+ \hat{z}_i (1 - \hat{z}_i)]^{1/2}$
2	2	51914.0
3	1	8237.0
4	0	-0.000
5	0	-0.001
24	0	-0.595
25	0	-0.608
26	0	-0.607
27	0	-0.591
95	0	-0.000
96	1	5.72 x 10 <sup>16</sup>
97	0	-0.000
177	0	-0.000

TABLE 10. STANDARDIZED RESIDUALS FOR BINOMIAL DISTRIBUTION Data treated as coming from single source, using formula (5.4)

In fact, in this case, it is possible to test

H\_: the four multiple failures come from a single BFR model.

Under  $H_0$ , given that 103 components fail in four occurrences, the conditional distribution of the four failure counts is multinomial(103, 1/4, 1/4, 1/4, 1/4). The distribution can then be found subject to the additional condition that each failure occurrence involves at least two failed components. Somewhat tedious but direct combinatorial calculations yield

 $P\left[-2 \log \Lambda \ge observed | H_0, n_+ = 4, s = 103, each failure count \ge 2\right]$ = 5 x 10<sup>-49</sup>.

Therefore, the occurrence with 96 failures does not come from the same BFR model as the other failure occurrences.
# 8. TECHNICAL DETAILS

### 8.1 Maximization of Likelihoods

Consider

$$L(p) = C \frac{p^{s+c-1} q^{mn_{+}-s+d-1}}{(1 - q^{m} - mpq^{m-1})^{n_{+}}}$$

If c > 0 and d > 0, this is the posterior density of p from a beta prior. If c = d = 1, it is the likelihood.

If m = 2 or  $n_{+} = 0$ , then s can only take the one value  $2n_{+}$ , and the data contain no information about p. So assume m > 2 and  $n_{+} > 0$ .

We will show that L(p) has a unique maximum (possibly infinite), at some  $p_0$  which is located according to the following seven cases and subcases.

If	Then
2n <sub>+</sub> < s < m n <sub>+</sub>	0 < p <sub>0</sub> < 1
$s = m n_+$	
d > 1	0 < p <sub>0</sub> < 1
d <u>≤</u> 1	p = ]
s = 2n <sub>+</sub>	
c > 1	0 < p <sub>0</sub> < 1
c < 1	$p_0 = 0$
c = 1	
$-\frac{n_{+}(m-2)}{3} - d + 1 > 0$	0 < p <sub>0</sub> < 1
$-\frac{n_{+}(m-2)}{3} - d + 1 \le 0$	$p_0 = 0$

Moreover, if  $0 < p_0 < 1$ , it is the value at which expression (8.1) equals zero.

Since s is an integer and c > 0,  $s + c - 1 - 2n_{+}$  can be negative if and only if  $s = 2n_{+}$  and c < 1. In this case L(p) is finite for 0 $and L(p) <math>\Rightarrow \infty$  as  $p \Rightarrow 0$ . So  $p_{0} = 0$ .

Assume now that  $s + c - 1 - 2n_+ \ge 0$ . Let us show that (a/ap)L(p) is zero at at most one point, and that the sign change can only be from positive to negative. The cases then follow from considering whether (a/ap)L(p)changes sign or not. Since (a/ap)L(p) has the same sign as  $q(a/ap)\log L(p)$ , it is sufficient to show that

$$q \frac{\partial}{\partial p} \log L(p) = \frac{s+c-1}{p} - \left\{ c+d-2 + mn_{+} \frac{1-q^{m-1}}{1-q^{m}-mpq^{m-1}} \right\}$$
 (8.1)

is strictly decreasing in p for 0 . Add the constant c+d-2 to both sides of (8.1) and write the result as

$$\frac{s+c-1-2n_{+}}{p} + n_{+} \left\{ \frac{2}{p} - m \frac{1-q^{m-1}}{1-q^{m}-mpq^{m-1}} \right\}.$$
(8.2)

Since  $s+c-1-2n_+ \ge 0$  by assumption, the first term of (8.2) is decreasing in p. Therefore it is sufficient to show that the expression in brackets is strictly decreasing in p for 0 and <math>m > 2.

The derivative of the expression in brackets in (8.2) is

$$\frac{m(m-1) p^2 q^{m-2} (1-q^m-mp) - 2(1-q^m-mpq^{m-1})^2}{p^2 (1-q^m-mpq^{m-1})^2}.$$

Denote the numerator by  $A_m(p)$ . It is not obvious that  $A_m(p)$  is negative, and we will show it by obtaining successively simpler expressions

 $B_m(p)$ ,  $C_m(p)$ , and  $D_m(p)$ . The outline below omits the tedious but direct algebra. Each assertion is verified in the obvious way.

$$A_2(p) = 0$$
, for  $0 .$ 

$$A_{m+1}(p) - A_{m}(p)$$

$$= -mp^{2}q^{m-2} \left\{ 6 - (4m + 6)p + (m+1)^{2}p^{2} + q^{m} \left[ -6 - (2m - 6)p + (m-1)p^{2} \right] \right\}$$

$$= -mp^{2}q^{m-1} B_{m}(p)$$

defining  $B_m(p)$ . Setting m=2 and q=1-p yields

$$\begin{split} B_{2}(p) &= p^{4} > 0, \quad \text{for } 0 0, \quad \text{for } 0 0, \quad \text{for all } p. \\ D_{m+1}(p) - D_{m}(p) &= p^{2}q^{m} \left[ 2m+3 - (m+2)p \right] > 0, \quad \text{for } m \ge 2, \ 0$$

Working up the above lines, we obtain successively  $D_m(p) > 0$  for  $m \ge 2$ ,  $0 , then <math>C_m(p) > 0$  for  $m \ge 2$ ,  $0 , then <math>B_m(p) > 0$  for  $m \ge 2$ ,  $0 , and finally <math>A_m(p) < 0$  for m > 2, 0 . This showsthat (8.2), and therefore (8.1), is strictly decreasing. So L(p) is maxi $mized at a unique <math>p_0$ .

To locate  $p_0$ , consider first (8.1) as  $p \ge 1$ . The limit is

$$s - mn_{+} - d + 1$$
,

which is negative if and only if

 $s < mn_+$ 

or

$$s = mn_+$$
 and  $d > 1$ .

So these are the cases for which  $p_0 < 1$ .

Now consider (8.1) as  $p \ge 0$ . Since

$$1 - q^{m-1} = (m-1)p - (\frac{m-1}{2})p^2 + O(p^3)$$

and

$$1 - q^{m} - mpq^{m-1} = {\binom{m}{2}}p^{2} q^{m-2} + {\binom{m}{3}}p^{3} + O(p^{4}),$$

we obtain after manipulation, that as  $p \ge 0$  (8.1) equals

$$\frac{s + c - 1 - 2n_{+}}{p} - \frac{n_{+}(m-2)}{3} - (c+d-2) + O(p).$$
(8.3)

Recall that s+c-1-2n<sub>+</sub> is assumed to be  $\geq 0$ . Expression (8.3) is positive if

 $s > 2n_{+}$ 

or

$$s = 2n_{+}, c > 1$$

or

 $s = 2n_{+}, c=1, -n_{+}(m-2)/3 - d + 1 > 0.$ 

So these are the cases for which  $p_0 > 0$ .

All the assertions made at the start of this section about the maximization of L(p) have now been proved.

We now verify the claim made in Section 3.1 that the right side of (3.3) is increasing in m. Let

$$G(m) = mp \frac{1 - q^{m-1}}{1 - q^m - mpq^{m-1}}$$
.

To show that G(m) is increasing in m, let us show that G(m+1) - G(m) is positive for p > 0. Algebraic manipulation shows that G(m+1) - G(m) is positive if and only if

$$(m+1)(1-q^m)(1-q^m - mpq^{m-1}) - m(1-q^{m-1})(1-q^{m+1} - (m+1)pq^m)$$

is positive. This quantity equals

$$(1-q^m)^2 - m^2 p^2 q^{m-1} = A$$

defining A. At p = 0, A is zero. The derivative  $\partial A/\partial p$  equals

$$mq^{m-2}[2q(1-q^m) - 2mpq + m(m-1)p^2] = mq^{m-2} B,$$

defining B. At p = 0, B and its first derivative are both zero. The second derivative is

 $2m(m+1)(1-q^{m-1}) > 0.$ 

Therefore, B is positive for all p > 0. Therefore, so is A, and therefore, G(m+1) > G(m) for p > 0.

# 8.2 Confidence Interval for p

In Section 3.2, a confidence interval for p was given, based on the conditional distribution of S given  $N_+$ . The choice of this particular confidence interval is now justified. The argument uses the properties of similar tests and Neyman structure (Ferguson<sup>8</sup>, pp. 226-7, or Reference 4, pp. 134-5) and monotone likelihood ratios (Reference 6, p. 423, or Reference 8, p. 208).

By the factorization (2.4), the problem may be formulated in terms of  $(\lambda_+, p)$  and the sufficient statistic  $(N_+, S)$ , ignoring  $\lambda_1$  and  $N_1$ . Suppose that we were testing

 $H_0: p \leq p_0$ 

 $H_1: p > p_0$ 

for some  $\boldsymbol{p}_{0}$  and some desired level  $\boldsymbol{\alpha}.$  A "similar" test is one with

$$P\left[\text{reject } H_0 \mid p = p_0\right] = \alpha$$

regardless of the value of  $\lambda_+$ . On the boundary

$$\left\{ (\lambda_{+}, p) | p = p_{0} \right\}$$

 $N_{+}$  is sufficient for  $\lambda_{+}$ . So, a test such that

 $P\left[(S, N_{+}) \text{ in critical region} | N_{+} = n_{+}\right] = \alpha \text{ for all } n_{+}$ (8.4)

has "Neyman structure." Since a Poisson random variable is boundedly complete, every similar test has Neyman structure, and any test which is uniformly most powerful (UMP) among tests satisfying (8.4) is UMP similar.

The conditional distribution of S given  $N_{+}$  is written in (3.7). So the likelihood ratio is

$$P\left[S = k \mid N_{+} = n_{+}; p_{1}\right] / P\left[S = k \mid N_{+} = n_{+}; p_{2}\right]$$
$$= \left(\frac{p_{2}}{q_{2}} \frac{q_{1}}{p_{1}}\right)^{k} \frac{q_{2}^{mn_{+}} (1 - q_{1}^{m} - mp_{1}q_{1}^{m-1})^{n_{+}}}{q_{1}^{mn_{+}} (1 - q_{2}^{m} - mp_{2}p_{2}^{m-1})^{n_{+}}}$$

which is monotone increasing in k for fixed  $p_1$  and  $p_2$ ,  $p_1 < p_2$ . Therefore, a UMP test among tests satisfying (8.4) is to reject  $H_0$  if and only if S is greater than some  $c(n_+)$ , with possible randomization if  $S=c(n_+)$ . This test is UMP similar.

Since a UMP similar test of a one-sided hypothesis has a one-sided critical region, good tests of two-sided hypotheses have two-sided critical regions, and one convenient such test assigns equal probabilities to the two tails. The corresponding confidence interval is given by (3.5) and (3.6).

#### 8.3 Integrals

Assume that A > 0 and B > 0. We repeat definition (4.4) here.

$$I(A, B, x) = \frac{B^{A}}{\Gamma(A)} \int_{0}^{x} e^{-Bt} t^{A-1} dt$$
 (4.4)

Then  $I(A,B,\infty) = 1$  for any A and B. For  $x < \infty$  the following three results hold. [See also Johnson and Kotz<sup>3</sup>, Vol. 2, Ch. 17, equations (23.1) through (24).]

If A is a positive integer,

$$I(A, B, x) = 1 - e^{-Bx} \frac{A-1}{\sum_{j=0}^{\Sigma} \frac{(Bx)^j}{j!}}.$$
 (8.5)

If A = k + 1/2 for integer k > 0,

$$I(A, B, x) = -e^{-Bx} \frac{k-1}{\Sigma} \frac{(Bx)^{j+1/2}}{\Gamma(j+3/2)} + 2 \Phi\left[(2Bx)^{1/2}\right] -1$$
(8.6)  
$$j=0$$

where  $\phi$  is the standard normal cumulative distribution function.

For arbitrary A > 0,

$$I(A, B, x) = e^{-Bx} \sum_{j=0}^{\infty} \frac{(Bx)^{A+j}}{\Gamma(A+j+1)} .$$
 (8.7)

To prove (8.5) and (8.6), integrate by parts and use induction. The proof of (8.6) is completed by observing that  $I(1/2, B, x) = P[X \le x]$  where  $X \sim \Gamma(1/2, B)$ . But this equals  $P[2BX \le 2Bx]$  where  $2BX \sim \chi^2(1)$ . Finally, this equals  $2\Phi[(2Bx)^{1/2}] - 1$ . To prove (8.7), observe that the two sides have identical derivatives and are equal at x=0.

Assume now that A, B, A', and B' are all positive. Define

$$= \int_{0}^{x} \int_{s}^{\infty} \frac{B^{A}}{\Gamma(A)} e^{-Bt} t^{A-1} \frac{(B')^{A'}}{\Gamma(A')} e^{-B's} s^{A'-1} dt ds.$$
(4.5)

Let

W(A, B, A', B') = 
$$\left(\frac{B}{B+B^{T}}\right)^{A} \left(\frac{B'}{B+B^{T}}\right)^{A'} \frac{\Gamma(A+A')}{\Gamma(A+1)}$$

Note the slight asymmetry of A and A' in W. The following four results hold.

If A is an integer,

$$J(A, B, A', B', x) = \sum_{j=0}^{A-1} W(j, B, A', B') I(A'+j, B+B', x).$$
(8.8)  
If A' is an integer,  

$$J(A, B, A', B', x) = I(A, B, x) + I(A', B', x) - I(A, B, x) I(A', B', x)$$

$$= I(A, B, x) + I(A', B', x) - I(A, B, x) I(A', B', x)$$

$$= I(A, B, x) + I(A', B', x) - I(A, B, x).$$
(8.9)  
If A and A' are arbitrary positive numbers,  

$$J(A, B, A', B', x) = I(A', B', x)$$

$$= I(A', B', x)$$

$$= I(A', B', x) [1 - I(A, B, x)]$$

$$+ \sum_{j=0}^{\infty} W(A'+j, B', A, B) I(A'+A'+j, B+B', x).$$
(8.10)

Equation (8.8) follows from (4.5) and use of (8.5). So does (8.9), after reversing the order of integration in (4.5). Equations (8.10) and (8.11) follow from (8.7), using both possible orders of integration in (4.5).

j=0

Of special interest is  $J(A, B, A', B', \infty)$ . If A or A' is an integer, then (8.8) and (8.9) give finite sums for  $J(A, B, A', B', \infty)$ . If A and A' are both integers plus 1/2, then there is also a finite expression for  $J(A, B, A', B', \infty)$ :

If A=k+1/2 and A'=k'+1/2, for nonnegative integers k and k', then

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$$J(A, B, A', B', \infty) = 1 + \sum_{j=0}^{k-1} W(j+1/2, B, A', B') - \sum_{j=0}^{k'-1} W(j+1/2, B', 1/2, B) - \sum_{j=0}^{k'-1} W(j+1/2, B', 1/2, B)$$

$$- \frac{2}{\pi} \arctan\left[ (B/B')^{1/2} \right]. \qquad (8.12)$$

To prove this last assertion, use (8.6) on the inner integral of (4.5). Note that  $I(A, B, \infty) = 1$  for any A and B, and obtain

$$J(A, B, A', B', \infty) = 2 + \sum_{j=0}^{k-1} W(j+1/2, B, A', B') - 2 \int_{0}^{\infty} \Phi[(2Bs)^{1/2}] \frac{(B')^{A'}}{\Gamma(A')} e^{-B's} s^{A'-1} ds.$$

Integrate the last integral by parts k' times, obtaining

$$J(A, B, A', B', \infty)$$

$$= 2 + \sum_{j=0}^{k-1} W(j+1/2, B, A', B') - \sum_{j=0}^{k'-1} W(j+1/2, B', 1/2, B_j)$$

$$- 2 \int_0^{\infty} \Phi\left[ (2Bs)^{1/2} \right] \frac{(B')^{1/2}}{\Gamma(1/2)} e^{-B's} s^{-1/2} ds.$$

The last integral equals  $P\left[T \le (2BS)^{1/2}\right]$ , where  $T \sim N(0,1)$  and  $S \sim \Gamma(1/2, B')$ . The distribution of 2B'S is  $X^2(1)$ , so the integral equals  $P\left[T \le (B/B')^{1/2}|Z|\right]$ , where T and Z are independent N(0,1). This probability is  $\left\{1 + \frac{2}{\pi} \arctan\left[(B/B')^{1/2}\right]\right\}/2$ , by the spherical symmetry of the bivariate normal distribution. This proves (8.12).

If J(A, B, A', B',  $\infty$ ) must be evaluated for arbitrary A and A', then one of (8.10) and (8.11) may converge much faster than the other. To decide which series to use, define

$$Q = B/(B+B')$$

Q' = B'/(B+B').

The ratio of term j+1 of the infinite series to term j is

 $\frac{A + A' + j}{A + j + 1} Q$ 

for (8.10), and

$$\frac{A + A' + j}{A' + j + 1} Q'$$

for (8.11). These expressions are monotone in j (decreasing if A > 1 and A' > 1), so are bounded by

$$\max(1, \frac{A+A'}{A+1}) Q$$
 (8.13)

and

$$\max(1, \frac{A + A'}{A' + 1}) Q'$$
 (8.14)

At least one of (8.13) and (8.14) is less than 1. For they are strictly bounded, respectively, by

$$\frac{Q}{A/(A + A')}$$
(8.15)

and

$$\frac{Q'}{A'/(A + A')} = \frac{1 - Q}{1 - A/(A + A')}$$
(8.16)

and the numerator of (8.15) is greater than the denominator if and only if the numerator of (8.16) is less than the denominator. So, reasonably fast convergence is assured by using (8.10) if (8.15) < (8.16) and using (8.11)otherwise.

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17. KEY WORDS AND DOCUMENT ANALYSIS

model assumptions. An example is worked out.

17a DESCRIPTORS

17b IDENTIFIERS/OPEN ENDED TERMS		
18. AVAILABILITY STATEMENT	19. SECURITY CLASS (This report) Unclassified	21 NO. OF PAGES
Unlimited	20 SECURITY CLASS (This page)	22 PRICE

NRC FORM 335 (7 77)

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