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# **Research Article**

# BAYESIAN ANALYSIS OF POISSON RELIABILITY MODEL WITH R AND JAGS

# Tanwir Akhtar<sup>1\*</sup> and Athar Ali Khan<sup>2</sup>

<sup>1</sup>Department of Public Health, College of Health Sciences, Saudi Electronic University, Jeddah-23442, SA <sup>2</sup>Department of Statistics and Operations Research, Aligarh Muslim University, Aligarh-202002, India

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### **ABSTRACT**

In many practical situations, reliability data are generated in the form of counts, also called failure counts, which record the number of times that a component fails in a specified period of time. Such data may arise because of limitation of the data capture system or the way the data are reported. For example, a system may keep track of the monthly number of failures and repair them. In this paper, an attempt has been made to model such type of data using Poisson distribution in Bayesian paradigm using two different prior distributions. Moreover, for the purpose of Bayesian modelling, two important techniques, that is, approximate *EM* algorithm and MCMC method are implemented using R and JAGS software packages, respectively. Approximate *EM* algorithm is implemented using R functions for approximating the posterior densities of model parameters. Whereas, JAGS is used to approximate the posterior parameters using Gibbs sampling and the Metropolis algorithm. R and JAGS code are developed and provided. A real data set is used for the purpose of illustrations.

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### **INTRODUCTION**

In many practical situations, reliability data are generated in the form of counts, also called failure counts, which record the number of times that a component fails in a specified period of time. Such data may arise because of limitation of the data capture system or the way the data are reported. For example, a system may keep track of the monthly number of failures and repair them (Hamada et al., 2008). Experimenter wants to asses such types of reliability data. For modelling such type of data the Poisson model is used, and is appropriate when the probability of events occurring in disjoint time interval is independent and when the probability of events occurring in short time of interval is small (Hamada et al., 2008). In this paper an attempt has been made to model failure count data in Bayesian paradigm using different prior distributions like Student-t prior and weakly informative conjugate prior distributions. In many cases it has been seen that non-Bayesian analysis of failure count data is not an easy task, whereas it can be implemented in principle in Bayesian paradigm, provided simulation tools are used (Akhtar and Khan, 2014a; Akhtar and Khan, 2014b, Khan et al., 2016). Consequently, for the purpose of Bayesian modelling, two important techniques, that is, approximate EM algorithm and MCMC method are implemented using R (R Core Team, 2017) and JAGS (Plummer, 2003) software packages, respectively, from within

R. Approximate *EM* algorithm is implemented using arm package of R for approximating the posterior densities of model parameters. Whereas, JAGS approximates the posterior parameters using Gibbs sampling (Geman and Geman, 1984:, Gelfand and Smith, 1990; Casella and George, 1992) and the Metropolis algorithm (Metropolis *et al.*, 1953) to generate a Markov chain by sampling from full conditional distributions. A real data set is used in subsequent analysis for the purpose of illustrations. Thus, the Bayesian analysis of Poisson reliability model has been made with the objectives: first to define a Bayesian model, that is, specification of likelihood and prior distribution, second to write down R and JAGS code for approximating the posterior densities using approximate *EM* algorithm and simulation tools, and finally to illustrate numeric as well as graphic summaries of the posterior densities.

### The Poisson Reliability Model

**The Model and Likelihood:** Assume that we have a set of discrete reliability data generated in the form of counts  $y_i$  which follows Poisson distribution with mean  $\theta$ . Algebraically,

$$y_i \sim \text{Poisson}(\theta t) = \frac{(\theta t)^y e^{-\theta t}}{y!}, i = 0, 1, 2, ... n, \ \theta > 0$$
 (1)

where  $y_i$  (i = 0, 1, 2, ... n) is the observed number of failures,  $\theta$  is the mean number of failure per unit time, and t is the length of the specified time period. Note that equal mean and variance

<sup>\*</sup>Corresponding author: Tanwir Akhtar

(here  $\theta$ t) is the most limiting characteristic of the Poisson distribution (Hamada *et al.*, 2008).

For the Poisson model, the parameter of interest is the Poisson rate  $\theta$ , which we wish to estimate. Therefore, the likelihood as a function of  $\theta$  given the observed failure count y is given by,

$$p(y|\theta) = \prod_{i=1}^{n} \frac{e^{-\theta t_{i}} (\theta t_{i})^{y_{i}}}{y_{i}!} = \frac{e^{-\theta \sum_{i=0}^{n} t_{i}} \prod_{i=0}^{n} (\theta t_{i})^{y_{i}}}{y_{i}!}$$
(2)

$$p(y|\theta) \sim e^{-\theta \sum_{i=0}^{n} t_i} \theta^{\sum_{i=0}^{n} y_i}$$
(3)

ignoring factors that do not depend on  $\theta$ .

Bayesian Inference: In order to complete the Bayesian model, one must specify a prior distribution. Usually the target posterior distribution is not analytically tractable. In the past, intractability was avoided by using the conjugate prior distributions, which have the nice property of resulting to posteriors of the same distributional family (Ntzoufras, 2009). Furthermore, an advantage of conjugate prior is that it makes posterior calculation easy. Thus, to obtain posterior density for the Poisson model, let us consider a gamma prior for mean  $\theta$  with parameter  $\alpha$  and  $\beta$  having density function,

$$\theta \sim Gamma (\alpha, \beta)$$

$$p(\theta) = \frac{\beta^{\alpha}}{\Gamma_{\alpha}} \theta^{\alpha-1} e^{-\beta \theta} \propto \theta^{\alpha-1} e^{-\beta \theta}.$$

Then the resulting posterior distribution of  $\theta$  by applying the Bayes' theorem is given as

$$\begin{array}{l} p(\theta|y) \propto p(\theta)p(y|\theta) \propto \theta^{\alpha-1} \ e^{-\beta\theta} \ \times \theta^{\sum_{i=0}^n y_i} \ e^{-\theta \sum_{i=0}^n t_i} \\ \propto \theta^{\alpha+\sum_{i=0}^n y_i-1} e^{-\theta(\beta+\sum_{i=0}^n t_i)}, \end{array}$$
 which implies that

$$\theta | y \sim Gamma \ (\alpha + \sum_{i=0}^{n} y_i, \beta + \sum_{i=0}^{n} t_i)$$

where  $y=(y_1,y_2,y_3,...,y_n)$ , and  $\beta$  can be interpreted as a prior sample size in contrast with the data sample size  $\sum_{i=0}^n t_i$  where  $\alpha$  is the prior number of failures in contrast with the observed number of failures  $\sum_{i=0}^n y_i$ . The posterior mean and variance are, respectively,

$$E(\theta|y) = \frac{\alpha + \sum_{i=0}^{n} y_i}{\beta + \sum_{i=0}^{n} t_i} = \frac{\alpha + n\bar{y}}{\beta + \sum_{i=0}^{n} t_i}$$

$$V(\theta|y) = \frac{\alpha + \sum_{i=0}^{n} y_i}{(\beta + \sum_{i=0}^{n} t_i)^2} = \frac{\alpha + n\bar{y}}{(\beta + \sum_{i=0}^{n} t_i)^2}$$

where  $\bar{y}$  is the sample mean (maximum likelihood estimator).

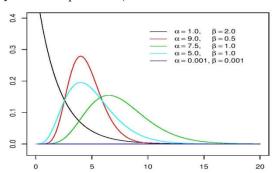


Figure 1 It is evident from the above plot that for shape  $\alpha=0.001$  and rate  $\beta=0.001$  the gamma distribution becomes almost uniform.

The usually selected low informative conjugate prior is a gamma distribution with low and equal prior parameters such as  $\alpha = \beta = 10^{-3}$ . This prior is convenient since its mean is equal to one while the variance is given by  $1/\alpha$ , which becomes large, expressing prior ignorance, for low value of  $\alpha$  (Ntzoufras, 2009). This fact is evident from the Figure 1.

### Implementation in Bayesian Paradigm

In order to implement the Bayesian analysis of Poisson model a super computer failure count data is taken from Hamada *et al.* (2008). Using these data different aspects of Bayesian analysis is discussed. For the purpose of fitting the model two different functions bayesglm and jags of R and JAGS, respectively, are used to estimate the parameter of interest,  $\theta$ .

#### Supercomputer Failure Count Data

The supercomputer failure count data discussed about the monthly number of failures of shared memory processors (SMPs) components of the Los Alamos National Laboratory Blue Mountain Supercomputer. The supercomputer consists of 47 identical SMPs and Table 1 represents monthly number of failures for the first month of operation. The supercomputer engineers expect that there should be no more than 10 failures for each component in the first month of operation.

**Table 1** Monthly number of failures for 47 supercomputer components (Hamada *et al.*, 2008).

| 1 | 5 | 1 | 4 | 2 | 3 | 1 | 3 | 6 | 4 | 4 | 4 | 2 | 3 | 2 | 2 |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| 4 | 5 | 5 | 2 | 5 | 3 | 2 | 2 | 3 | 1 | 1 | 2 | 5 | 1 | 4 | 1 |
| 1 | 1 | 2 | 1 | 3 | 2 | 5 | 3 | 5 | 2 | 5 | 1 | 1 | 5 | 2 |   |

### Analysis Using R

In the beginning, a Bayesian analysis of Poisson model using the bayesglm function from arm (Gelman and Su, 2014) package of R is conducted. This function is a Bayesian alteration of classical generalized linear model that uses an approximate EM algorithm to update the  $\beta$ s at each step using an augmented regression to represent the prior information (Gelman et al., 2008). This function uses Student-t prior distribution for the coefficients. Since, in our case, model involves intercept only, therefore, the prior distribution for the intercept term is t-distribution with 1 degree of freedom (i.e., Cauchy distribution), and prior mean is 0 whereas prior scale is 10. The main arguments of the bayesglm function are:

The first argument formula requires a symbolic description of the model to be fitted. For example, if a model involves intercept only, the formula of the model will be of the form y~1. The argument family requires a description of the error distribution and link function to be used in the model. The default error distribution is gaussian. The data argument needs an optional data frame, list, or environment containing the variables in the model. For more details see Gelman and Su (2014).

### Data Creation

Firstly, for Bayesian fitting of Poisson model for supercomputer data with the function bayesglm, the data are created in R format as follows.

$$n < -47$$

```
y<-c(1,5,1,4,2,3,1,3,6,4,4,4,2,3,2,2,4,5,5,2,5,3,2,2,3,1,1,2,5,1,4,1,1,1,2,1,3,2,5,3,5,2,5,1,1,5,2)
```

where n represents the total number of SMPs, and y represents the monthly number of failures for the first month of operations.

## **Model Specification**

The distribution assumed for count data is a Poisson, which applies when counted things are distributed independently and randomly. The Poisson model has a single parameter, the expected count  $\theta$ , that is often called intensity (Kery, 2010) and, in case of supercomputer data, represents monthly failure rate or mean number of failures. In contrast to the normal, the Poisson variance in not a free parameter but is equal to the mean,  $\theta$ .

Thus, for supercomputer failure count data, where the response  $(y_1, y_2, ..., y_n)$  is the monthly number of failures recorded for the SMPs, the model for these responses is

$$y_i \sim Poisson(\theta),$$
  $i = 0, 1, 2, ..., 47.$ 

Here, count response  $y_i$  for *i*th SMP is distributed as a Poisson random variable with mean number of failure  $\theta$ , for the first month of operation (i.e.,  $t_i = 1$  month). Moreover, for Bayesian analysis of Poisson model the function bayesglm uses default log-link function, that is,

$$log(\theta) = Linear \ predictor = Intercept$$

In this case, intercept is the only term in linear predictor. Therefore, after fitting the model, to get the results in original metric the exponential function <code>exp</code> is used to transform the results.

# Model Fitting

Finally, to fit the above defined model, the function bayesglm is called to perform the analysis and put its results into an object called Fit. The display function of arm package is used to print the summary of results. This generic function with its detail=TRUE argument gives a clean printout of the fitted object, focusing on the most pertinent pieces of information including p-value or z-value.

```
Fit<-bayesglm (y~1, family=poisson
(link="log"))
display (Fit, detail=TRUE)</pre>
```

In order to get the posterior simulations of the model parameter theta, a generic function of arm package called sim is used. To get the clear picture of parameter estimates, 1000 (n.sim=1000) independent draws are made. Moreover, to convert the results in original metric the function exp is used, and its output is reported and discussed in the next section.

```
set.seed (123)
## Simulation
Fit.sim<-sim (Fit,n.sim=1000)
coef.Fit.sim<-coef (Fit.sim)
exp(coef.Fit.sim)
## Estimate
apply(exp(coef.Fit.sim),2,mean)
## Standard Deviation
apply(exp(coef.Fit.sim),2,sd)</pre>
```

```
## Quantiles
quantile(exp(coef.Fit.sim),prob=c(0.025,0.
25,0.5,0.75,0.975))
```

### Summarizing Output

Posterior summaries and densities after being fitted the model with bayesglm are provided in Table 2. From this table, it is observed that, the estimate of theta is 2.82 with standard deviation 0.24, and that the 95% credible interval is [2.37, 3.35], which is statistically significant. The plot for the posterior density of theta can be seen in Figure 1.

**Table 2** From this output it is evident that posterior mean and median are very close, which is an indication of symmetric posterior density

| Parameter | Mean | SD   | 2.5% | 25%  | Median | 75%  | 97.5% |
|-----------|------|------|------|------|--------|------|-------|
| Theta     | 2.82 | 0.24 | 2.37 | 2.65 | 2.80   | 2.97 | 3.35  |

### Analysis Using JAGS

Let us consider the Bayesian analysis of the same failure count data with JAGS using its interface of R that is R2jags (Su and Yajima, 2014) package of R, which includes the posterior simulation and convergence diagnostic of the model. For modelling of these data in JAGS, one must specify the model to run, and to load data which is created in a separate file and the initial values of the model parameters for a specified number of Markov chains (Akhtar and Khan, 2017; Khan *et al.*, 2017). The R2jags package makes use of this feature and provides convenient functions to call JAGS directly from within R. Furthermore, it is possible to work with the results after importing them back into R again, for example, to create a posterior predictive simulations, or more generally, graphical displays of data and posterior simulations.

### Data Creation

The first part of modelling in JAGS is data definition, which must be defined in a list containing the name of each vector. In this case, the monthly number of failures are the response variable  $y_i$  (i = 0, 1, 2, ..., 47). These data can be created in R format as:

```
n <- 47
y <-
c(1,5,1,4,2,3,1,3,6,4,4,4,2,3,2,2,4,5,5,2,5,3,2,2,3,1,1,2,5,1,4,1,1,1,2,1,3,2,5,3,5,2,5,1,1,5,2)
jags.data <- list("n","y")</pre>
```

Here n is the total number of supercomputer components (SMPs), y is the response values of monthly number of failures, and these are combined in a list with object jags.data.

### **Model Specification**

For modelling these data, the monthly number of failures  $y_i$  is assumed to follow a Poisson distribution

```
y_i \sim Poisson(\theta), \qquad i = 0, 1, 2, ..., 47,
```

where  $\theta$  is the individual monthly failure rate for the SMPs. In this case, the monthly failure rate  $\theta$  is the unknown model parameter which we want to estimate. It is expected that there should be no more than 10 failures for each component. To represent this, the prior information for the parameter  $\theta$  is

assumed to be a conjugate gamma prior distribution with a mean of 5, that is,

```
\theta \sim Gamma (5, 1),
```

It is observed that, the probability that  $\theta$ , exceeds 10 for Gamma(5, 1) prior is 0.03, which is very small (Hamada *et al.*, 2008).

Thus, the specification of above defined model in JAGS language must be put in a separate file which is then read by JAGS. When working in R, this is most conveniently done using the cat function of R which behaves pretty much like paste with the exception that the results is not a character object but directly written to a file we specify (Akhtar and Khan, 2017; Khan *et al.*, 2017). Here is the JAGS code specifying the model using cat function to put in the file model2.jags: cat("model{"model}")

```
for(i in 1:n) {
  y[i]~dpois(theta)
}
theta~dgamma(5,1)
}", file="model2.jags.txt")
```

This model allows each observed response y to follow a Poisson distribution with mean theta, which is drawn from a gamma distribution with shape 5 and rate 1, that is, mean value 5.

#### Initial Values

The starting values used to initialize the chain are simply called the *initial values*. Usually it is necessary to specify a starting value for the chains to start the MCMC simulation. In most cases, JAGS will however be able to generate the initial values itself. In order to be able to monitor convergence, we will normally run several chains for each parameter. The starting value for the chains is a named list, names are the parameters used in the model. Each element of the list is itself a list of starting values for the JAGS model, or a function creating (possible random) initial values. In this case, there is only one parameter, called theta, in the model:

```
inits <- function() {list(theta =
rlnorm(1))}</pre>
```

### Model Fitting

Using the objects jags.data, inits, and model2.jags, we can now compile the model and run MCMC simulation to get estimate for theta by calling JAGS. For MCMC simulation 3 chains (n.chain=3) each with 5000 iterations (n.iter=5000) has been run, and parameter theta is monitored. Its results are assigned with object Fit, and can conveniently be printed by print (Fit), which prints details summary of the results and are summarized in the next section.

```
Fit <- jags(jags.data, inits,
parameter=c("theta"),n.iter=5000,
n.chain=3, model.file="model2.jags.txt",)
print(Fit)</pre>
```

### Summarizing Output

The output of the R function jags is a list which includes several components, most notable are the summary of the inference and convergence, and a list containing the simulation

draws of all the saved parameters. In this case, the jags call is assigned to the R object Fit, and so typing print (Fit) from the R console will display the summary of the fitted model shown below. The print method displays information on the mean, standard deviation, 95% credible interval (CI) estimates, the effective sample size, and potential scale reduction factor  $\hat{R}$  of the Brook-Gelman-Rubin (BGR) statistics (Gelman and Rubin, 1992; Brooks and Gelman, 1998). The BGR statistics is an analysis of variance (ANOVA)-type diagnostic that compares within- and among-chain variance (Kery, 2010). Values around 1 indicate good convergence, with 1.1 considered as acceptable limit (Gelman and Hill, 2007).

**Table 3** Posterior summary of JAGS simulation after being fitted to the Poisson model for the supercomputer failure count data.

| Parameter | Mean   | SD    | 2.5%   | Median | 97.5%  | Rhat  | n.eff |
|-----------|--------|-------|--------|--------|--------|-------|-------|
| theta     | 2.86   | 0.249 | 2.39   | 2.85   | 3.36   | 1.001 | 3800  |
| deviance  | 171.61 | 1.507 | 170.57 | 171.04 | 175.77 | 1.001 | 3800  |
| pD        |        | 1.1   |        | DIC    |        | 172.7 |       |

The first five columns of numbers of Table 3 give inference for model parameter. In this case, the posterior estimate for mean theta is 2.86 with standard deviation 0.249 and 95% credible interval is [2.39, 3.36], which is statistically significant. The median estimate is 2.85 and very close to mean estimate. At the bottom, pD shows the estimated effective number of parameters in the model, and DIC, the deviance information criterion, an estimate of predictive error. Finally, consider the right most columns of the output, where Rhat gives information about convergence of the algorithm. At convergence, the number at this column should be equal to 1. In this case, Rhat is very close to 1, indication of well mixing of the three chains and thus good convergence. The final column, n.eff is the effective sample size of the simulations.

Additionally, to see the complete picture of the results, a plot can be generated by plot(Fit), and the resulting plot is depicted in Figure 1. In this plot, the left column shows a quick summary of inference and convergence, that is, Rhat. The right column shows inference for the model parameter theta. Finally, the posterior density plot for parameter theta is depicted in Figure 2.

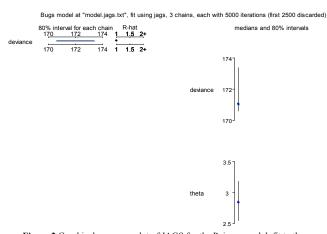
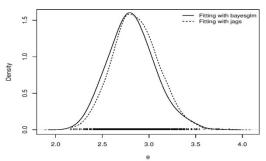


Figure 2 Graphical summary plot of JAGS for the Poisson model, fit to the supercomputer failure count data. R-hat is near to one for parameter theta, indicating good convergence.



**Figure 3** Posterior density plot of the model parameter theta. Results from different methods have different styles. The closeness of the two approaches look evident in the graphics.

# **DISCUSSION AND CONCLUSION**

Component reliability is the foundation of reliability assessment and refers to the reliability of single component. In this paper, Bayesian approach is used to model failure count component reliability data for intercept model with Poisson distribution. The log-link function is used to relate the model parameter and intercept term for linear Poisson model. Two different prior distributions, that is, Student-t and gamma prior distributions are used and implemented using R and JAGS software packages, respectively. For modelling of failure count data, the complete R and JAGS code are written and provided with detailed description. It is observed that the results obtained from approximate EM algorithm and simulation method using different software packages are very close to each other. The benefits of approximate EM algorithm and simulation method seem clear in the plot of posterior densities. Moreover, it is evident from the summaries of results that the Bayesian approach based on different priors is simpler to implement than the frequentist approach. Finally, the wealth of information provided in these numeric as well as graphic summaries is not possible in classical framework.

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