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Estimation and model selection for left-truncated and right-censored lifetime data with application to electric power transformers analysis

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In lifetime analysis of electric transformers, the maximum likelihood estimation has been proposed with the EM algorithm. However, it is not clear whether the EM algorithm offers a better solution compared to the simpler Newton-Raphson algorithm. In this paper, the first objective is a systematic comparison of the EM algorithm with the Newton-Raphson algorithm in terms of convergence performance. The second objective is to examine the performance of Akaike's information criterion (AIC) for selecting a suitable distribution among candidate models via simulations. These methods are illustrated through the electric power transformer dataset.

Keywords Akaike's information criterion; EM algorithm; lognormal distribution; Newton-Raphson algorithm; Weibull distribution; Reliability.

Mathematics Subject Classification 62N01, 62N02, 62N05

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1. Introduction

Electric power transformers have long lifetime, typically 30 - 40 years under normal operating conditions, due to their high level of reliability (Zhou 2013). Accordingly, researchers require long follow-up studies with certain observational constraints, which lead to left-truncation and right-censoring. Truncation and censoring are common in lifetime data as discussed in the books by Meeker and Escobar (1998) and Lawless (2003). Recently, Hong, Meeker and McCalley (2009) carry out lifetime analysis of electric power-transformer data in the US. Their lifetime data were left-truncated at the starting date of record keeping and right-censored at the ending date of the study. For this dataset, they propose likelihood inference and prediction analysis appropriately adjusted for truncation and censoring.

The lognormal and Weibull distributions would be the two most relevant statistical distributions to model the electric power-transformer data. They have been extensively used to model lifetime data in the literature. Readers are referred to the books by Crow and Shimizu (1988) for the lognormal and by Bryan (2006) for the Weibull. They are often fitted to lifetime data after discriminating between the Weibull and lognormal distributions (e.g., Kundu and Manglick 2004; Emura and Wang 2010).

The fitting of the Weibull distribution to the electric power-transformer data is considered in Hong et al. (2009). They propose a parametric likelihood analysis that properly adjusts for the sampling bias due to left-truncation and right-censoring. Their study provides a prediction analysis of the remaining lifetime of the power transformers. Zhou (2013) also considers the Weibull model for the power transformer data in the absence of truncation.

The fitting of the lognormal distribution to the electric power-transformer data is developed by Balakrishnan and Mitra (2011). In their paper, the EM algorithm (EM) for fitting the lognormal distribution to the left-truncated and right-censored data is described. Under the same model, confidence intervals and prediction intervals are developed using the EM-based missing information principle (Balakrishnan and Mitra, 2013). Their simulations

show that the EM-based interval has correct coverage rates and is comparable to the intervals based on the observed information matrix and the parametric bootstrap. The EM is also developed under the Weibull distribution by Balakrishnan and Mitra (2012). The EM-based likelihood inference for the gamma and the generalized gamma distribution is developed in the discussion paper of Balakrishnan and Mitra (2014). The generalized gamma distribution includes both the Weibull and lognormal distributions as special cases, and hence it provides a unified framework for performing the EM.

The EM requires mathematical derivation of the expected log-likelihood (E-step) and its numerical maximization (M-step). In fact, the EM proposed by Balakrishnan and Mitra (2011, 2012) is not simple as it requires some numerical approximations to the M-step. Nevertheless, the EM often provides stable results when appropriately used. Hence, it is important to clarify whether the EM offers better solution compared to the simpler Newton-Raphson algorithm.

Checking the adequacy of models is an important issue in parametric analyses. Hong et al. (2009) used a graphical model checking procedure to verify the Weibull assumption. However, they did not consider model selection among other candidate distributions. A recently published Ph.D. thesis of Mitra (2013) and the discussion paper of Balakrishnan and Mitra (2014) considered model selection via Akaike's information criterion (AIC) and Bayesian information criterion (BIC).

The first objective of this paper is to make a comparison between the Newton-Raphson (NR) method and EM algorithm (EM) under the lognormal and Weibull distributions via simulations and real data analysis. For the Weibull distribution, we also propose a simplified NR (called one-dimensional NR), which will be a better alternative to the usual NR and EM.

The second objective of this paper is to investigate Akaike's information criterion (AIC, Akaike, 1974) to select a suitable model among candidate models. The AIC allows one to compare several candidate models with different degree of freedoms and hence provides an objective criterion to select a model. During our study, we find that the application of AIC is

also discussed in the Ph.D. thesis of Mitra (2013) and the discussion paper of Balakrishnan and Mitra (2014). However, our work is conducted independently and hence supplements their work under different settings. In fact, the candidate models in our simulations are different from those in Mitra (2013) and Balakrishnan and Mitra (2014).

The rest of this paper is organized as follows. Section 2 describes the data structure and the likelihood function. Section 3 introduces the Newton-Raphson and the EM algorithms. Section 4 defines AIC for model selection. Section 5 presents simulations to compare the NR and EM algorithms and to examine the performance of AIC. Section 6 analyzes the electric power transformer datasets. Section 7 concludes the paper.

2. Likelihood construction with left-truncation and right-censoring

In this section, we review the data structures and likelihood construction in the presence of the left-truncation and right-censoring as considered in Hong et al. (2009) and Balakrishnan and Mitra (2011, 2012).

2.1 Left-truncated and right-censored data

Data collected on electric power transformers typically involves lower and upper observational limits, which produces left-truncation and right-censoring respectively. In an example considered in Hong et al. (2009), the lifetime of the US power transformers are recorded from 1980 to 2008, a 28 year-period. The interval is still shorter than the average lifetimes of power transformers of 30 - 40 years under normal operating conditions. Ignoring censoring and truncation leads to sampling bias.

Figure 1 gives an illustration of truncation and censoring. For Case 1, the installation year of the machine is between 1980 and 2008, and the machine is still in service even after 2008. Hence, Case 1 is right-censored on 2008. For Case 2, the machine installed before 1980, and it fails between 1980 and 2008. Hence, Case 2 is left-truncated on 1980. For Case 3, the

machine is installed before 1980, and it still works after 2008. Hence, Case 3 is both left-truncated and right-censored. For Case 2, the installation year of the machine and the failure time of the machine are both before 1980. There is no data available on the lifetime for Case 4 as the machine is completely missed out of the sampling protocol.

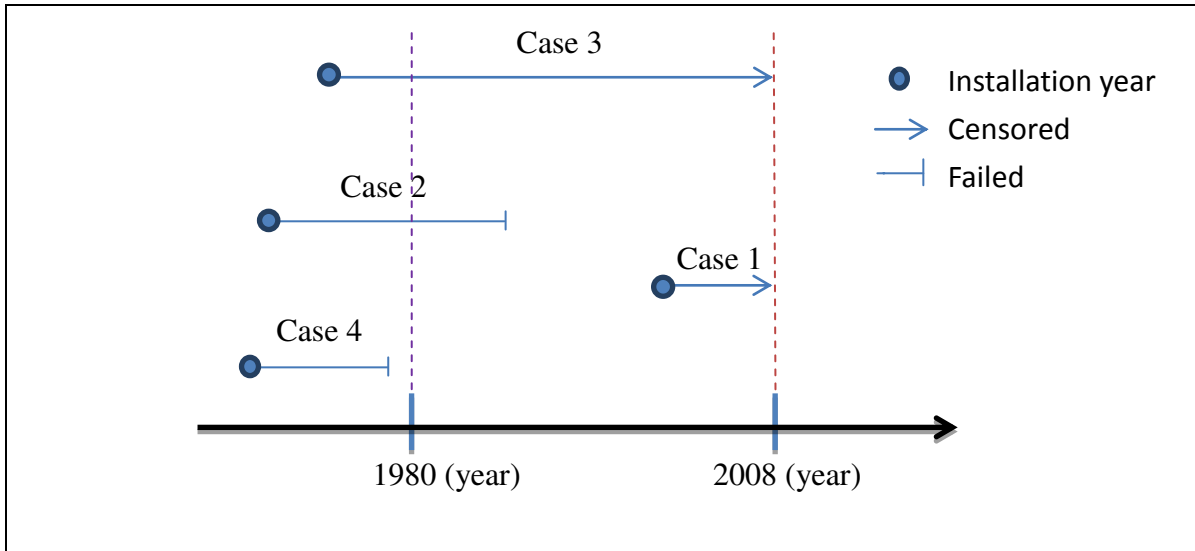


Figure 1 Example for left truncated and right censored data

2.2 Likelihood construction

To construct the likelihood, we follow the parametric likelihood approaches of Hong et al. (2009) and Balakrishnan and Mitra (2011, 2012). Note that their likelihoods suitably adjust for the bias due to left-truncation and right-censoring.

Let X be the original lifetime variable and $T = \log(X)$ be the log-transformed variable. For i -th machine, t_i denotes the observed value for T and c_i denotes the right-censored time. More precisely, for a machine still in service after 2008, c_i is the time between the year of installation and the censoring point of 2008 (Figure 2). Let c_i^L be the log-transformed right-censored time and δ_i be the censoring indicator, i.e.,

$$\delta_i = \begin{cases} 0, & \text{if the observation is censored,} \\ 1, & \text{if the observation is not censored.} \end{cases}$$

Let τ_i be the left-truncated time. More precisely, for a machine installed before 1980, τ_i is the time between the year of installation and the truncation point of 1980 (Figure 2). Let τ_i^L be the log-transformed left-truncated time and v_i be the truncation indicator, i.e.,

$$v_i = \begin{cases} 0, & \text{if the observation is truncated,} \\ 1, & \text{if the observation is not truncated.} \end{cases}$$

Further, let S_1 and S_2 be two index sets, where $S_1 = \{i; v_i = 1\}$ is the set of machines installed after 1980, and $S_2 = \{i; v_i = 0\}$ is the set of machines installed before 1980.

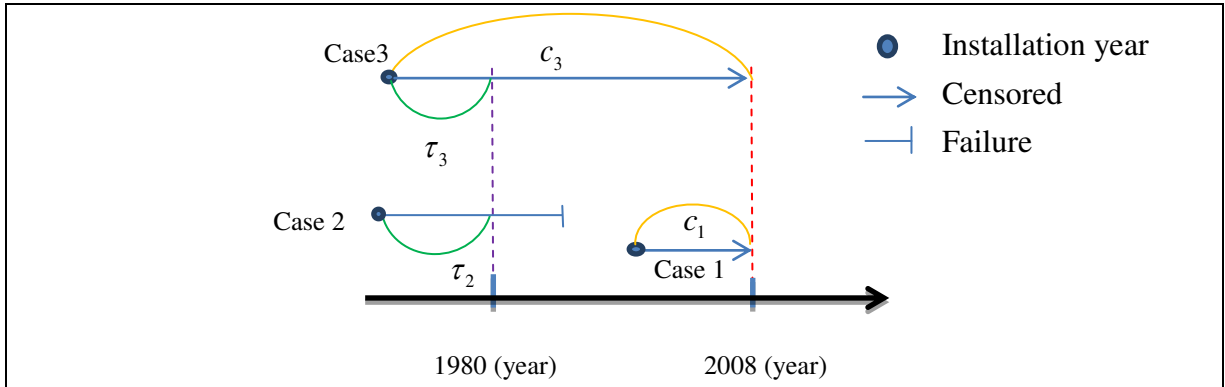


Figure 2 An example for truncated time (τ_i) and censored time (c_i).

Example 1: Lognormal distribution

We introduce the likelihood under the lognormal distribution. Assume that the log-transformed lifetime $T = \log X$ follows a normal distribution with mean μ and standard deviation $\sigma > 0$. Then, the likelihood function is

$$L(\mu, \sigma) = \prod_{i \in S_1} \left\{ \frac{1}{\sigma} \phi \left(\frac{y_i - \mu}{\sigma} \right) \right\}^{\delta_i} \left\{ 1 - \Phi \left(\frac{y_i - \mu}{\sigma} \right) \right\}^{1 - \delta_i} \times \prod_{i \in S_2} \left\{ \frac{\frac{1}{\sigma} \phi \left(\frac{y_i - \mu}{\sigma} \right)}{1 - \Phi \left(\frac{\tau_i^L - \mu}{\sigma} \right)} \right\}^{\delta_i} \left\{ \frac{1 - \Phi \left(\frac{y_i - \mu}{\sigma} \right)}{1 - \Phi \left(\frac{\tau_i^L - \mu}{\sigma} \right)} \right\}^{1 - \delta_i},$$

where $y_i = \min(t_i, c_i^L)$ and $\phi(\cdot)$ and $\Phi(\cdot)$ are the density and distribution function of the standard normal distribution. The log-likelihood function (without the constant term) is

$$\begin{aligned} & \log L(\mu, \sigma) \\ &= \sum_{i=1}^n \left[-\delta_i \left\{ \log \sigma + \frac{(y_i - \mu)^2}{2\sigma^2} \right\} + (1 - \delta_i) \log \left\{ 1 - \Phi \left(\frac{y_i - \mu}{\sigma} \right) \right\} \right] - \sum_{i \in S_2} \log \left\{ 1 - \Phi \left(\frac{\tau_i^L - \mu}{\sigma} \right) \right\} \\ &= \sum_{i=1}^n \left[-\delta_i \left\{ \log \sigma + \frac{(y_i - \mu)^2}{2\sigma^2} \right\} + (1 - \delta_i) \log \left\{ 1 - \Phi \left(\frac{y_i - \mu}{\sigma} \right) \right\} \right] - \sum_{i=1}^n (1 - v_i) \log \left\{ 1 - \Phi \left(\frac{\tau_i^L - \mu}{\sigma} \right) \right\}. \end{aligned}$$

□

Example 2: Weibull distribution

We introduce the likelihood under the Weibull distribution with the density of X ,

$$\left(\frac{\beta}{\eta} \right) \left(\frac{x}{\eta} \right)^{\beta-1} \exp \left[- \left(\frac{x}{\eta} \right)^\beta \right], \quad x \geq 0, \eta > 0, \beta > 0,$$

where η is the scale and β is the shape parameter. The likelihood function is

$$\begin{aligned} L(\eta, \beta) &= \prod_{i \in S_1} \left\{ \left(\frac{\beta}{\eta} \right) \left(\frac{x_i}{\eta} \right)^{\beta-1} \exp \left[- \left(\frac{x_i}{\eta} \right)^\beta \right] \right\}^{\delta_i} \left\{ \exp \left[- \left(\frac{x_i}{\eta} \right)^\beta \right] \right\}^{1 - \delta_i} \\ &\quad \times \prod_{i \in S_2} \left\{ \frac{\left(\frac{\beta}{\eta} \right) \left(\frac{x_i}{\eta} \right)^{\beta-1} \exp \left[- \left(\frac{x_i}{\eta} \right)^\beta \right]}{\exp \left[- \left(\frac{\tau_i}{\eta} \right)^\beta \right]} \right\}^{\delta_i} \left\{ \frac{\exp \left[- \left(\frac{x_i}{\eta} \right)^\beta \right]}{\exp \left[- \left(\frac{\tau_i}{\eta} \right)^\beta \right]} \right\}^{1 - \delta_i}, \end{aligned}$$

where $x_i = \exp(y_i)$ is the observed lifetime. The log-likelihood function is

$$\log L(\eta, \beta) = \sum_{i=1}^n \left[\delta_i \left\{ \log \beta - \log \eta + (\beta - 1) \log \left(\frac{x_i}{\eta} \right) \right\} - \left(\frac{x_i}{\eta} \right)^\beta \right] + \sum_{i=1}^n (1 - v_i) \left(\frac{\tau_i}{\eta} \right)^\beta. \quad \square$$

Example 3: Exponential distribution

We introduce the likelihood under the exponential distribution, which is a special case of the Weibull distribution with $\beta = 1$. Therefore, the log-likelihood function is given by

$$\log L(\eta) = \sum_{i=1}^n \left\{ -\delta_i \log \eta - \frac{x_i}{\eta} \right\} + \sum_{i=1}^n (1 - v_i) \left(\frac{\tau_i}{\eta} \right). \square$$

In the following, we will use the log-likelihood of each distribution to derive the MLE.

3. Newton-Raphson and EM algorithms

3.1 Newton-Raphson algorithm

If the first-order and second-order derivatives of the log-likelihood are available, one can maximize the likelihood function using the Newton-Raphson (NR) method. The NR is suitable to the present problem since all the required derivatives are analytically available.

Example 1: Lognormal distribution

The formulas for the first- and second-order derivatives of the log-likelihood with respect to the parameters are available in Balakrishnan and Mitra (2011). The MLE of (μ, σ) is obtained by sequentially updating the estimate with

$$\begin{bmatrix} \mu_{k+1} \\ \sigma_{k+1} \end{bmatrix} = \begin{bmatrix} \mu_k \\ \sigma_k \end{bmatrix} - J_f^{-1}(\mu_k, \sigma_k) \begin{bmatrix} f_1(\mu_k, \sigma_k) \\ f_2(\mu_k, \sigma_k) \end{bmatrix},$$

where $f_1(\mu, \sigma) = \partial \log L(\mu, \sigma) / \partial \mu$, $f_2(\mu, \sigma) = \partial \log L(\mu, \sigma) / \partial \sigma$, and

$$J_f(\mu, \sigma) = \begin{bmatrix} \partial f_1(\mu, \sigma) / \partial \mu & \partial f_1(\mu, \sigma) / \partial \sigma \\ \partial f_2(\mu, \sigma) / \partial \mu & \partial f_2(\mu, \sigma) / \partial \sigma \end{bmatrix}.$$

The iteration continues until $|\mu_{k+1} - \mu_k| < \varepsilon$ and $|\sigma_{k+1} - \sigma_k| < \varepsilon$, for some pre-fixed $\varepsilon > 0$.

We set $\varepsilon = 0.001$ as the stopping criterion for all the simulations.

Remark 1: Balakrishnan and Mitra (2011) do not study the numerical performance of the NR.

Instead, they compare their EM with the Fisher scoring algorithm.

Example 2: Weibull distribution

The first-order derivatives of the log-likelihood are given by

$$\frac{\partial}{\partial \eta} \log L(\eta, \beta) = \frac{\beta}{\eta} \sum_{i=1}^n \left\{ -\delta_i + \left(\frac{x_i}{\eta} \right)^\beta \right\} - \frac{\beta}{\eta} \sum_{i=1}^n (1-v_i) \left(\frac{\tau_i}{\eta} \right)^\beta,$$

$$\frac{\partial}{\partial \beta} \log L(\eta, \beta) = \sum_{i=1}^n \left\{ \delta_i \left[\frac{1}{\beta} + \log \left(\frac{x_i}{\eta} \right) \right] - \left(\frac{x_i}{\eta} \right)^\beta \log \left(\frac{x_i}{\eta} \right) \right\} + \sum_{i=1}^n \left\{ (1-v_i) \left(\frac{\tau_i}{\eta} \right)^\beta \log \left(\frac{\tau_i}{\eta} \right) \right\}.$$

Importantly, the equation $\partial \log L(\eta, \beta) / \partial \eta = 0$ leads to an explicit solution,

$$\hat{\eta}(\beta) = \left\{ \frac{\sum_{i=1}^n x_i^\beta - \sum_{i=1}^n (1-v_i) (\tau_i)^\beta}{\sum_{i=1}^n \delta_i} \right\}^{\frac{1}{\beta}}.$$

Therefore, given $\eta_k = \hat{\eta}(\beta_k)$, one can obtain a one-dimensional estimating function

$f(\beta) = \partial \log L(\eta_k, \beta) / \partial \beta = 0$ for β . We propose a one-dimensional NR to obtain $\hat{\beta}$ with

$$\beta_{k+1} = \beta_k - f(\beta_k) / f'(\beta_k),$$

where $f'(\beta)$ is defined as

$$\frac{\partial^2}{\partial \beta^2} \log L(\eta_k, \beta) = \sum_{i=1}^n \left\{ -\frac{\delta_i}{\beta^2} - \left(\frac{x_i}{\eta_k} \right)^\beta \left[\log \left(\frac{x_i}{\eta_k} \right) \right]^2 + (1-v_i) \left(\frac{\tau_i}{\eta_k} \right)^\beta \left[\log \left(\frac{\tau_i}{\eta_k} \right) \right]^2 \right\}.$$

The iteration stops if $|\beta_{k+1} - \beta_k| < \varepsilon$, for some $\varepsilon > 0$, where we set $\varepsilon = 0.001$ for all the

simulations. The MLE of $\hat{\eta}$ is explicitly obtained after finding $\hat{\beta}$. A similar procedure in

the absence of truncation is proposed by Zhou (2013).

Remark II: Although Balakrishnan and Mitra (2012) compare their EM with the NR, their NR is the usual two dimensional NR using R maxNR routine for their simulations. We rather propose the present one-dimensional NR due to its simplicity. \square

Example 3: Exponential distribution

Since the exponential distribution is a special case of the Weibull distribution with $\beta = 1$, we immediately find the solution to $\partial \log L(\eta) / \partial \eta = 0$ as

$$\hat{\eta} = \hat{\eta}(1) = \frac{\sum_{i=1}^n x_i - \sum_{i=1}^n (1-v_i)(\tau_i)}{\sum_{i=1}^n \delta_i}.$$

There is no need to use the NR. However, the second derivative of the log-likelihood is still useful to confirm that the solution $\hat{\eta}$ is indeed the maximum of the likelihood by $\partial^2 \log L(\hat{\eta}) / \partial \eta^2 < 0$ (Appendix I). \square

3.2 EM algorithm

In this section, we briefly introduce the EM algorithm proposed by Balakrishnan and Mitra (2011, 2012). Let $\mathbf{t} = (t_1, t_2, \dots, t_n)'$ be the complete log-transformed lifetimes. Since some of t_i 's are censored and hence not observed exactly, $\boldsymbol{\delta} = (\delta_1, \delta_2, \dots, \delta_n)'$ and $\mathbf{y} = (y_1, y_2, \dots, y_n)'$ are observed data. Let $\boldsymbol{\theta} = (\mu, \sigma)'$ be parameters to be estimated.

Example 1: Lognormal distribution

The complete data log-likelihood (without constant terms) is

$$\log L_c(\mathbf{t}, \boldsymbol{\theta}) = -n \log \sigma - \sum_{i=1}^n \left\{ \frac{(t_i - \mu)^2}{2\sigma^2} \right\} - \sum_{i=1}^n (1-v_i) \log \left[1 - \Phi \left(\frac{\tau_i^L - \mu}{\sigma} \right) \right].$$

The E-step calculates the conditional expectation of the complete data log-likelihood

$$Q(\boldsymbol{\theta}, \boldsymbol{\theta}_k) = E_{\boldsymbol{\theta}_k} [\log L_c(\mathbf{t}, \boldsymbol{\theta}) | \mathbf{y}, \boldsymbol{\delta}].$$

The M-step performs the maximization $\boldsymbol{\theta}_{k+1} = \arg \max_{\boldsymbol{\theta}} Q(\boldsymbol{\theta}, \boldsymbol{\theta}_k)$.

Balakrishnan and Mitra (2011) suggest a numerical approximation either by a Taylor expansion (EM1) or EM gradient algorithm (EM2) for the M-step. The iterations continue

until $|\mu_{k+1} - \mu_k| < \varepsilon$ and $|\sigma_{k+1} - \sigma_k| < \varepsilon$ for $\varepsilon = 0.001$ as specified for simulations. \square

Example 2: Weibull distribution

Let $\mu = \log(\eta)$ and $\sigma = 1/\beta$, where η is the scale parameter and β is the shape parameter as before. The complete data log-likelihood (without constant terms) is

$$\log L_c(\mathbf{t}, \boldsymbol{\theta}) = -n \log \sigma - \sum_{i=1}^n \left\{ \left(\frac{t_i - \mu}{\sigma} \right) - \exp \left(\frac{t_i - \mu}{\sigma} \right) \right\} + \sum_{i=1}^n (1 - v_i) \exp \left(\frac{\tau_i^L - \mu}{\sigma} \right).$$

The formula of the conditional expectation of the complete data log-likelihood $Q(\boldsymbol{\theta}, \boldsymbol{\theta}_k)$ is a complicated nonlinear function as given explicitly in Balakrishnan and Mitra (2012). They suggest a linear approximation based on the EM gradient algorithm. The MLE is obtained by repeatedly calculating $\boldsymbol{\theta}_{k+1} = \arg \max_{\boldsymbol{\theta}} Q(\boldsymbol{\theta}, \boldsymbol{\theta}_k)$. \square

4. Model selection

Akaike's information criterion or AIC (Akaike, 1974) is a measure of the relative goodness-of-fit for a given model penalized by the number of parameters in the model. In particular, AIC is given by $AIC = -2 \log \hat{L} + 2k$, where k is the number of unknown parameters in the model and \hat{L} is the maximized value of the likelihood function for the fitted model. One selects the model with the minimum AIC value.

Another commonly used method for model selection is the Bayesian information criterion or BIC (Schwarz, 1978) given by $BIC = -2 \log \hat{L} + k \log n$, where n is the sample size.

Although, AIC and BIC are both simple to apply, their empirical performances are often different. A comparison of AIC and BIC is given by Burnham and Anderson (2002). In the biological and social sciences and medicine, they argue that the AIC-type criteria are reasonable for the analysis of empirical data. BIC might find use in some physical sciences where a simple true model exists and where sample size is quite large. They recommend AIC

for general use in the model selection. It is well known that AIC is minimax-rate optimal for estimating the regression function, and BIC is consistent in selecting the true model (Yang, 2005). In other words, AIC selects the best fitted model and BIC selects the true model.

For a given data, we do not know the true model and even do not know whether the true model belongs to our candidate models or not. Therefore, we suggest AIC as a general way for model selection in order to find the best fitted model.

5. Simulations

5.1 Simulation design

We adopt the simulation design of Balakrishnan and Mitra (2011). We generate the installation years under the fixed percentage of truncation at 30 or 60%. The set of installation years are split into two parts: (1960-1979) and (1980-1995). Then, the installation years were simulated according to the sampling probabilities on Figure 3. For example, suppose that the sample size is $n = 100$, and the percentage of truncation is 30%. Since the truncation year is fixed at 1980 as in Hong et al. (2009), we generate the installation years that follow the probability of each year from the truncated part of (1960-1979) with 30 sample sizes, and the from un-truncated part of (1980-1995) with other 70 sample sizes (Figure 3).

Divide installation years into truncated and un-truncated cases									
↓					↓				
(1960-1979); Truncated part					(1980-1995); Un-truncated part				
Year	1960	1961	1962	1963	Year	1980	1981	1982	1983
Probability	0.1	0.1	0.1	0.1	Probability	0.15	0.15	0.15	0.15
Year	1964	1965	1966~1979		Year	1984	1985~1995		
Probability	0.1	0.1	0.4/14 (each)		Probability	0.15	0.25/11 (each)		

Figure 3 Sampling probabilities for generating the installation years.

Then the lifetimes of the machines, in years, are simulated from lognormal, Weibull or exponential distributions. Adding these lifetimes to the corresponding installation years, we obtain the failure years of the machines. If the failure year of a machine exceeds 2008, the machine is censored, where 2008 is the fixed censoring point.

There are some remarks to notice. For the left-truncated machines, if the year of failure is before 1980, one cannot detect the machine (Case 4 in Figure 1). Therefore, if the year of the failure is before 1980, we ignore this machine, and generate a new one with the new installation year and lifetime. The sample sizes used in our simulations are $n = 50, 100,$ and 200 . All the simulation results are based on 1000 Monte Carlo runs. We set the stopping criterion $\varepsilon = 0.001$ for all the simulations for both NR (Newton-Raphson) and EM (EM algorithms).

5.2 Results under the lognormal distribution

The lifetimes of the machines are simulated from the lognormal distribution with (μ, σ) being $(3.5, 0.5)$ or $(3.0, 0.2)$, the same values as Balakrishnan and Mitra (2011). We compare the three algorithms, namely EM1, EM2, NR, where EM1 corresponds to EM algorithm approximating the hazard function by a Taylor expansion, EM2 corresponds to the EM gradient algorithm, and NR corresponds to the Newton-Raphson method. The sample mean and sample standard deviation of y_i 's are used as initial values for μ and σ , respectively.

Table 1 compares the results of the three different methods. Overall, the three methods produce almost unbiased results and have small MSE. As the sample size increases, the bias and the MSE tend to decrease. This implies that the MLE obtained by the three methods all work well and the three methods are quite comparable in terms of the bias and MSE. However, the average number of iterations in the NR is smaller than both EM1 and EM2. This quick convergence may be regarded as the advantage of the NR over EM1 and EM2.

Under a different setting from Table 1, we find that occasional un-convergence occurs

especially for the NR under small sample sizes and high censoring percentages. In the following we pick up such a case.

Table 2 gives the separate simulation results under small sample sizes and high censoring percentage. It can be seen that the NR sometimes produces un-convergence. In spite of the problem in the NR, the EM1 always converges. Although the percentages of un-convergent runs in the NR are quite small, the problem may still occur as many engineering applications have small sample sizes with high censoring percentages.

Table 3 shows the results for the EM1 with $n = 50$, where the NR and EM2 occasionally fail to converge. We see that the EM1 always converges and has reasonable performance for the bias and MSE. Therefore, under this configuration, only the EM1 works properly.

From Tables 1-3, it can be concluded that, for moderate samples, the EM1, EM2, and NR perform very similarly in terms of the bias and MSE. However, the NR method converges more quickly than the EM1 and EM2. Nevertheless, under small sample sizes and high percentage of censoring, EM1 is the only one reliable method.

[Insert Tables 1-3]

5.3 Results under the Weibull distribution

The lifetimes of the machines are simulated from the Weibull distribution with (η, β) being $(35, 3)$ and $(40, 3)$, which corresponds to $(\mu, \sigma) = (3.55, 0.33)$ and $(3.69, 0.33)$, respectively. For the Weibull distribution, we denoted T as the log-transformed lifetime variable which follows an extreme value distribution with parameters μ and σ . Then $E(T) = \mu - \gamma\sigma$ and $\text{Var}(T) = \pi^2\sigma^2/6$, where $\gamma = 0.5772$ (approximately) is Euler's constant. Accordingly, we choose the initial values (μ_0, σ_0) such that

$$\sum_{i=1}^n y_i / n = \mu_0 - \gamma\sigma_0 \quad \text{and} \quad \sum_{i=1}^n (y_i - \bar{y})^2 / (n-1) = \pi^2\sigma_0^2 / 6.$$

Table 4 compares the performance of the NR and EM in terms of the bias, MSE, and

average number of iterations until convergence. It can be observed that the bias and MSE are very close to zero for both the NR and EM. Hence the accuracy of the EM and NR is virtually the same. Also, as the sample size increases, the MSE decreases for both methods. A remarkable difference is that the NR takes fewer steps until convergence than the EM. Unlike the lognormal distribution, the one-dimensional NR always converges under the Weibull distribution even for the small sample sizes ($n = 50$) and high censoring percentage (66.3%).

[**Insert Tables 4**]

5.4 Model selection performance

We examine the model selection performance of AIC. For instance, if the data is simulated from lognormal distribution, the MLEs of the three models (lognormal, Weibull and exponential) are calculated, and then AICs are computed under the three models. Finally, we select the model that has the smallest AIC among the three. In this case, one may expect that AIC is the smallest with the lognormal distribution.

Table 5 gives the model selection performance of AIC when the data is simulated from the lognormal distribution. As expected, the percentage that the lognormal is selected is higher than the other two. The result is consistent with the observation that the average AIC calculated under the lognormal distribution is the smallest among the three distributions.

Table 6 shows the performance of AIC when the data is generated from the Weibull distribution. Again, the percentage of selecting the Weibull model is the highest and the average AIC calculated under the Weibull model is the smallest among the three distributions.

It should be noted that the mean lifetimes of the data simulated from the lognormal and Weibull distributions are both near 30. Therefore, we choose the mean parameter η of the exponential distribution as 30. Table 7 shows the performance of AIC when the data is simulated from the exponential distribution. As expected, the percentage of selecting the exponential distribution is the highest and the average AIC is smallest under the exponential

distribution among the three distributions.

From Tables 5-7, we find that AIC can appropriately identify the correct model and the percentage of choosing the correct model increases as the sample size increases. Therefore, the model selection via AIC seems to have a model selection consistency.

[Insert Tables 5-7]

6. Data analysis

The power transformer lifetime data consist of 710 observations with 62 failures from manufactures (Hong, et al. 2009). Although the original data is not available, their paper provides a systematic subset of the data containing 286 observations with 39 failures, which is described in Appendix II.

Table 8 shows the successive steps of iterations of the NR and EM (EM1 and EM2) for fitting the lognormal distribution. We find that the NR diverges at the 2nd iteration step. The high censoring percentage (86.4%) explains this phenomenon as the NR occasionally diverges under small sample sizes and high censoring percentage in the simulations (Section 5). On the other hand, the two EM algorithms converge and their estimates are very close to one another. The EM1 converges more quickly than the EM2 does. In all cases, the initial values for the parameters μ and σ are taken as the sample mean and sample standard deviation of y_i 's .

Now, we reveal the detailed behavior of convergence using the EM and NR from Figure 4. Obviously, the NR moves a wrong way while EM gradually moves to the maximum of the likelihood. This may be because NR tends to have a big leap in one iteration step. While the big leap accelerates the convergence speed, it can increase the chance of divergence.

Based on the dataset, we estimate the parameters of the lognormal, Weibull, and exponential distributions and then compute AIC for the respective models. The resultant AIC values are 470.04 (lognormal), 472.29 (Weibull) and 470.67 (Exponential). Therefore, we choose the lognormal distribution to be the suitable model for this data.

[Insert Table 8]

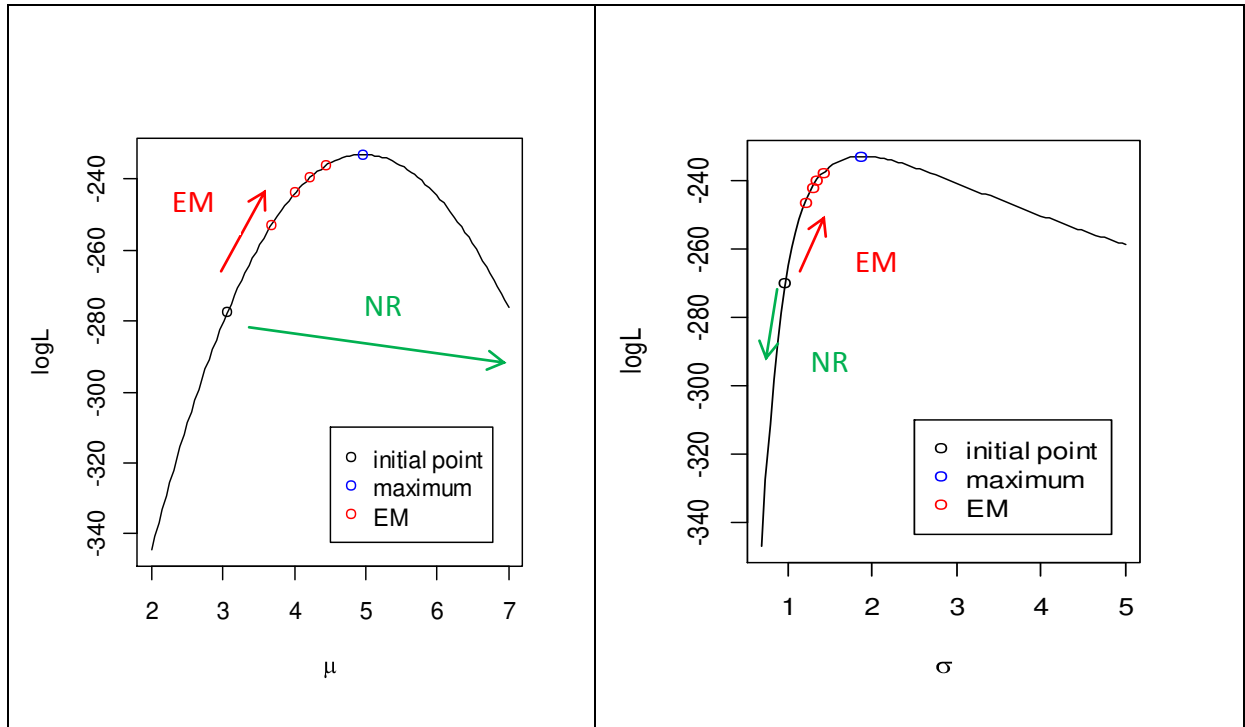


Figure 4 The directions of convergence using EM algorithm and NR (Newton-Raphson) method. The EM algorithm converges to the maximum while the NR algorithm diverges.

7. Conclusion and discussion

The first objective of this paper is to compare the performance of the EM algorithm and Newton-Raphson method based on the left-truncated and right-censored data. We summarize the highlights of our finding as follows:

- For the lognormal distribution, when the sample sizes are small and censoring percentage is high, the Newton-Raphson occasionally fails to converge. On the other hand, the EM algorithm with the approximation by a Taylor expansion still converges.
- The Newton-Raphson converges more quickly than the EM when both converge.
- The transformer lifetime data analysis demonstrates the real case where the EM algorithm converges but the Newton-Raphson diverges.

- For the Weibull distribution, the proposed one-dimensional Newton-Raphson provides faster convergence speed than the EM algorithms in any circumstance and the two algorithms converge to the virtually same limit. Therefore, the EM algorithm appears to have little advantage over the one-dimensional Newton-Raphson under the Weibull.

Although our comparison between the NR and EM algorithms is based on the left-truncated and right-censored data, our conclusion (EM is better for lognormal; NR is better for Weibull; NR is faster when it converges) may be generalized to other data structures that use EM algorithms for censored data. Obviously, there are many papers that utilize the EM algorithms for handling censored data. For instance, Ng, Chan, and Balakrishnan (2002) used EM algorithms to determine the maximum likelihood estimates of the lognormal and Weibull distributions when data are progressively Type II censored. Recently, Fan and Wang (2011) and Balakrishnan and Pal (2013) develop EM algorithms for the Weibull analysis under very general competing risks structures. Since the likelihood in their paper seems to be twice differentiable, the NR method may still apply. However, it is less clear to us whether our one-dimensional NR method is appropriate or not for such complicated data structure.

The second objective of this paper is to investigate Akaike's information criterion (AIC) for model selection. In the simulations, we have confirmed that AIC can correctly identify the true model among the candidate models. In addition, when the sample size gets large, the percentage of choosing the correct model increases. Therefore, AIC exhibits model selection consistency. Note that Barakrishnan and Mitra (2014) also consider AIC as well as BIC as a model selection tool. Our candidate models in the simulations are exponential, Weibull and lognormal distributions while those in Barakrishnan and Mitra (2014) are Weibull, lognormal, gamma, and generalized gamma distributions. Hence, our paper provides additional support for the performance of AIC under different simulation settings.

In future work, one may not only consider AIC for distribution choice, but also for

variable selection and grouping. For example, Hong et al. (2009) consider a regression model that includes manufacture ID, insulation class, and cooling system as explanatory variables. It is possible to apply AIC to select optimal sets of explanatory variables, though the numerical performance remains to be studied. One can also use AIC for grouping. Hong et al. (2009) first split the sample into “Old” and “New” groups, where the Old group mostly consists of truncated samples and the New group mostly consists of un-truncated samples. Then, they fit a Weibull model with different shape and scale parameters between the two groups. One may apply AIC to see whether this split results in a better fit. It is interesting to point out that the different parameters due to truncation can be associated with the concept of “dependent truncation” [see Emura and Wang (2012) and references therein; Bakoyannis and Touloumi, 2011]. This implies that truncation has some information about the lifetime. It is also interesting to consider the effect of “dependent censoring” [see Emura and Chen (2014) and references therein]. How to incorporate the dependent truncation/censoring information in model selection of power transformer lifetimes is an interesting topic for further investigation.

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Appendix I: Confirming that $\hat{\eta}$ is indeed the MLE

Under the exponential distribution, the solution to the likelihood equation $\partial \log L(\eta) / \partial \eta = 0$ is

$$\hat{\eta} = \left\{ \sum_{i=1}^n x_i - \sum_{i=1}^n (1 - v_i) \tau_i \right\} \left\{ \sum_{i=1}^n \delta_i \right\}^{-1} > 0.$$

Now we verify that the solution is indeed the maximum of the likelihood function by

$$\begin{aligned} \left. \frac{\partial^2}{\partial \eta^2} \log L(\eta) \right|_{\eta=\hat{\eta}} &= \frac{1}{\hat{\eta}^3} \left\{ \sum_{i=1}^n [x_i - (1-v_i)\tau_i] - 2 \sum_{i=1}^n [x_i - (1-v_i)\tau_i] \right\} \\ &= -\frac{1}{\hat{\eta}^3} \left\{ \sum_{i=1}^n [x_i - (1-v_i)\tau_i] \right\} < 0. \end{aligned}$$

The last inequality holds since $x_i \geq \tau_i \geq (1-v_i)\tau_i$ for all i .

Appendix II: The subset of data provided by Hong, Meeker, and McCalley (2009)

Table A1 is obtained by reading off numerical values from Fig. 1 of Hong, et al. (2009).

References

- Akaike, H. (1974). A new look at the statistical model identification, *IEEE Transactions on Automatic Control* 19 (6), 716–723.
- Bakoyannis G, Touloumi G. (2012). Practical methods for competing risks data: a review. *Statistical Method in Medical Research* 21, 257-272.
- Balakrishnan, N., Mitra, D. (2011). Likelihood inference for lognormal data with left truncation and right censoring with an illustration. *Journal of Statistical Planning and Inference* 141, 3536-3553.
- Balakrishnan, N., Mitra, D. (2012). Left truncated and right censored Weibull data and likelihood inference with an illustration. *Computational Statistics & Data analysis* 56, 4011-4025.
- Balakrishnan, N., Mitra, D. (2013). Some further issues concerning likelihood inference for left truncated and right censored lognormal data. *Communications in Statistics-Simulation and Computation* 43, 400-416.
- Balakrishnan, N., Pal, S. (2013). Expectation maximization-based likelihood inference for flexible cure rate models with Weibull lifetimes. *Statistical Methods in Medical Research*, doi: 10.1177/0962280213491641.
- Balakrishnan, N., Mitra, D. (2014). EM-based likelihood inference for some lifetime distributions based on left truncated and right censored data and associated model discrimination. *The South African Statistical Journal*, to appear.
- Bryan, D. (2006). *The Weibull Analysis Handbook*, Second Edition, Milwaukee: American Society for Quality, Quality Press.
- Burnham, K.P., Anderson, D.R. (2002). *Model Selection and Multimodel Inference: A Practical Information-Theoretic Approach*, Second ed.
- Crow, E.L., Shimizu, K. (1988). *Lognormal Distributions: Theory and Applications*. Marcel Dekker, New York.
- Emura, T, Wang, H. (2010). Approximate tolerance limits under the log-location-scale models in the presence of censoring, *Technometrics* 52(3), 313-323.
- Emura, T. and Wang, W. (2012), Nonparametric maximum likelihood estimation for dependent truncation data based on copulas, *Journal of Multivariate Analysis* 110, 171-188.
- Emura, T. and Chen, Y.H. (2014), Gene selection for survival data under dependent censoring: a copula-based

- approach, *Statistical Methods in Medical Research*, doi: 10.1177/0962280214533378.
- Fan, T.H., Wang, W.L. (2011). Accelerated Life Tests for Weibull Series Systems With Masked Data. *IEEE Transaction on Reliability* 60, No.3.
- Hong, Y., Meeker, W.Q., McCalley, J.D. (2009). Prediction of remaining life of power transformers based on left truncated and right censored lifetime data. *The Annals of Applied Statistics* 3, 857-879.
- Kundu, D, Manglick, A. (2004). Discriminating between the Weibull and log-normal distributions, *Naval Research Logistics* 5, 893-905.
- Lawless, J.F. (2003). *Statistical Models and Methods for Lifetime Data*, Second Edition, New York: John Wiley and Sons.
- Mitra, D. (2013). Likelihood inference for left truncated and right censored data. *Open Access Dissertations and Theses*. Paper 7599.
- Meeker, W.Q., Escobar, L.A. (1998). *Statistical Methods for Reliability Data*. John Wiley & Sons, New York.
- Ng, H.K.T., Chan, P.S., Balakrishnan, N. (2002). Estimation of parameters from progressively censored data using EM algorithm. *Computational Statistics & Data Analysis* 39, 371-386.
- Schwarz, G.E. (1978). Estimating the dimension of a model. *Annals of Statistics* 6 (2), 461–464.
- Yang, Y. (2005). Can the strengths of AIC and BIC be shared? A conflict between model identification and regression estimation, *Biometrika* 92, 937-950.
- Zhou, D. (2013). Comparison of two popular methods for transformer Weibull lifetime modeling, *International Journal of Advanced Research in Electrical, Electronics and Instrumentation Engineering* Vol. 2, Issue 4.

Table 1

Bias (B), mean square error (MSE), average number of iterations (AI) for the lognormal distribution using three different methods (EM1, EM2, and NR) in 1000 Monte Carlo simulations.

(μ, σ)	Trun.(%) Cen. (%)	Method	B($\hat{\mu}$)	B($\hat{\sigma}$)	MSE($\hat{\mu}$)	MSE($\hat{\sigma}$)	AI
<i>n</i> = 100							
(3.5, 0.5)	30; 62.3	EM1	0.00001	0.00028	0.00508	0.00445	14.0
		EM2	-0.00033	-0.00001	0.00507	0.00445	17.5
		NR	0.00216	0.00260	0.00517	0.00449	7.0
	60; 51.1	EM1	0.00703	-0.00102	0.00431	0.00363	10.3
		EM2	0.00662	-0.00134	0.00430	0.00362	13.5
		NR	0.00772	0.00026	0.00437	0.00364	6.4
(3.0, 0.2)	30; 17.4	EM1	-0.00103	-0.00134	0.00047	0.00025	3.9
		EM2	-0.00110	-0.00150	0.00047	0.00025	4.3
		NR	-0.00010	-0.00136	0.00047	0.00025	3.9
	60; 10.1	EM1	0.00107	-0.00142	0.00054	0.00025	3.7
		EM2	0.00096	-0.00140	0.00054	0.00025	3.7
		NR	0.00099	-0.00135	0.00054	0.00025	3.5
<i>n</i> = 200							
(3.5, 0.5)	30; 62.3	EM1	-0.00052	-0.00128	0.00256	0.00201	13.9
		EM2	-0.00084	-0.00155	0.00255	0.00201	17.4
		NR	0.00160	0.00103	0.00261	0.00202	7.0
	60; 51.1	EM1	0.00009	-0.00274	0.00188	0.00173	10.2
		EM2	-0.00031	-0.00306	0.00188	0.00173	13.3
		NR	0.00072	-0.00153	0.00190	0.00173	6.3
(3.0, 0.2)	30; 17.4	EM1	0.00036	-0.00037	0.00023	0.00012	3.9
		EM2	0.00028	-0.00054	0.00023	0.00012	4.2
		NR	0.00038	-0.00039	0.00023	0.00013	3.9
	60; 10.1	EM1	0.00099	-0.00055	0.00024	0.00013	3.7
		EM2	0.00087	-0.00052	0.00024	0.00013	3.6
		NR	0.00089	-0.00047	0.00024	0.00013	3.4

Trun. = Truncation percentage (%); Cens. = Censoring percentage (%);

EM1 = the EM algorithm method approximating the hazard function by the Taylor expansion;

EM2 = the EM gradient algorithm; NR is the Newton-Raphson method.

Table 2

The percentage of unconvergence for the lognormal distribution under three different methods (EM1, EM2, and NR) in 1000 Monte Carlo simulations.

Method	(μ, σ)	Sample size	Truncation (%)	Censoring (%)	Unconvergence (%)
EM1	(3.5, 0.5)	50	30	62.3	0
			60	51.1	0
		100	30	62.3	0
			60	51.1	0
EM2	(3.5, 0.5)	50	30	62.3	0.1
			60	51.1	0
		100	30	62.3	0
			60	51.1	0
NR	(3.5, 0.5)	50	30	62.3	1.9
			60	51.1	0.5
		100	30	62.3	0.2
			60	51.1	0

Unconvergence (%) = $100 \times (\text{the number of un-convergent runs}) / 1000$;

EM1 = the EM algorithm method approximating the hazard function by the Taylor expansion,

EM2 = the EM gradient algorithm,

NR = the Newton-Raphson method.

Table 3

Bias (B), mean square error (MSE), average number of iterations (AI) for the lognormal distribution under the EM1 method for a sample size of 50 in 1000 Monte Carlo simulations.

(μ, σ)	Trun.(%); Cen. (%)	Method	$B(\hat{\mu})$	$B(\hat{\sigma})$	$MSE(\hat{\mu})$	$MSE(\hat{\sigma})$	AI
<i>n</i> = 50							
(3.5, 0.5)	30; 62.3	EM1	0.00235	-0.00594	0.01038	0.00802	14.17
	60; 51.1	EM1	0.00372	-0.00712	0.00802	0.00713	10.33
(3.0, 0.2)	30; 17.4	EM1	0.00141	-0.00254	0.00089	0.00050	3.97
	60; 10.1	EM1	0.00102	-0.00248	0.00103	0.00052	3.77

EM1 = the EM algorithm method approximating the hazard function by the Taylor expansion.

Table 4

Bias (B), mean square error (MSE), and average number of iterations (AI) for the Weibull distribution under two different methods (EM and NR) in 1000 Monte Carlo simulations.

(μ, σ)	Trun. (%); Cens. (%)	Method	B($\hat{\mu}$)	B($\hat{\sigma}$)	MSE($\hat{\mu}$)	MSE($\hat{\sigma}$)	AI
<i>n</i> = 50							
(3.55, 0.33)	30; 56.8	EM	-0.00414	-0.00848	0.00569	0.00333	21.8
		NR	0.00028	-0.00610	0.00585	0.00333	5.3
	60; 42.3	EM	-0.00321	-0.00125	0.00415	0.00310	12.8
		NR	-0.00149	-0.00037	0.00421	0.00310	4.7
(3.69, 0.33)	30; 66.3	EM	-0.00722	0.00957	0.00463	0.00433	33.1
		NR	0.00114	-0.00002	0.00471	0.00442	6.0
	60; 53.3	EM	-0.00328	-0.00244	0.00528	0.00451	19.5
		NR	0.00055	0.00013	0.00546	0.00459	5.0
<i>n</i> = 100							
(3.55, 0.33)	30; 56.8	EM	-0.00474	-0.00308	0.00274	0.00187	21.4
		NR	-0.00053	-0.00076	0.00279	0.00188	5.3
	60; 42.3	EM	-0.00309	-0.00438	0.00196	0.00164	12.5
		NR	-0.00143	-0.00323	0.00199	0.00165	4.7
(3.69, 0.33)	30; 66.3	EM	-0.00878	-0.00764	0.00390	0.00220	32.1
		NR	-0.00096	-0.00350	0.00402	0.00220	5.9
	60; 53.3	EM	-0.00204	-0.00111	0.00244	0.00175	19.2
		NR	0.00162	0.00140	0.00253	0.00178	5.0
<i>n</i> = 200							
(3.55, 0.33)	30; 56.8	EM	-0.00478	-0.00532	0.00147	0.00082	21.1
		NR	-0.00069	-0.00305	0.00149	0.00081	5.3
	60; 42.3	EM	-0.00220	-0.00158	0.00101	0.00079	12.4
		NR	-0.00058	-0.00044	0.00102	0.00080	4.8
(3.69, 0.33)	30; 66.3	EM	-0.00538	-0.00362	0.00208	0.00111	31.3
		NR	0.00233	0.00055	0.00217	0.00112	5.8
	60; 53.3	EM	-0.00443	-0.00475	0.00124	0.00102	18.9
		NR	-0.00088	-0.00232	0.00127	0.00102	5.0

Trun. = Truncation percentage (%); Cens. = Censoring percentage (%)

EM = the EM algorithm

NR = the Newton-Raphson method

Table 5

The percentage of the model selected by AIC and the average of AIC when the data is simulated from the lognormal distribution with $(\mu, \sigma) = (3.5, 0.5)$.

Truncation (%); Censoring (%)	Sample size	Percentage (%)			Average AIC		
		LN	WB	EP	LN	WB	EP
30; 62.3	$n = 50$	76.9	20.4	0	172.2	173.8	190.6
	$n = 100$	86.0	14.0	0	344.4	347.6	381.4
	$n = 200$	91.6	8.4	0	684.9	691.3	759.7
60; 51.1	$n = 50$	79.9	20.1	0	216.6	218.2	232.9
	$n = 100$	85.6	14.4	0	431.5	434.6	465.3
	$n = 200$	92.4	7.6	0	860.8	866.7	928.1

Note: LN = lognormal; WB = Weibull; EP = Exponential;
We select the model that has the smallest AIC.

Table 6

The percentage of the model selected by AIC and the average of AIC when data is simulated from the Weibull distribution with $(\mu, \sigma) = (3.55, 0.33)$.

Truncation (%); Censoring (%)	Sample size	Percentage (%)			Average AIC		
		LN	WB	EP	LN	WB	EP
30; 56.8	$n = 50$	33.0	67.0	0	193.0	191.1	214.1
	$n = 100$	17.9	82.1	0	383.3	378.7	424.0
	$n = 200$	8.5	91.5	0	763.8	753.7	847.3
60; 42.3	$n = 50$	31.6	68.4	0	240.3	238.0	262.4
	$n = 100$	17.5	82.5	0	481.7	475.9	526.0
	$n = 200$	7.2	92.8	0	959.1	948.2	1049.1

Note: LN = lognormal; WB = Weibull; EP = Exponential;
We select the model that has the smallest AIC.

Table 7

The percentage of the model selected by AIC and the average of AIC when the data is simulated from the exponential distribution with $\eta = 30$.

Truncation (%); Censoring (%)	Sample size	Percentage (%)			Average AIC		
		LN	WB	EP	LN	WB	EP
30 ; 43.9	$n = 50$	19.0	11.3	69.7	250.8	249.2	248.3
	$n = 100$	13.9	12.2	73.9	503.6	499.3	498.3
	$n = 200$	8.7	12.8	78.5	989.2	967.3	966.8
60; 41.6	$n = 50$	22.0	10.5	67.5	260.0	258.7	257.7
	$n = 100$	17.6	11.8	70.6	518.3	515.0	514.0
	$n = 200$	9.0	12.2	78.8	1036.4	1028.3	1027.3

Note: LN = lognormal; WB = Weibull; EP = Exponential;
We select the model that has the smallest AIC.

Table 8.

The successive steps of iterations of the EM algorithms and Newton-Raphson (NR) method for the lognormal distribution.

EM1		EM2		NR	
Step	$(\hat{\mu}, \hat{\sigma})$	Step	$(\hat{\mu}, \hat{\sigma})$	Step	$(\hat{\mu}, \hat{\sigma})$
1	(3.065, 0.968)	1	(3.065, 0.968)	1	(3.065, 0.968)
2	(3.675, 1.209)	2	(4.255, 0.659)	2	(13.156,-3.395)
3	(4.015, 1.292)	3	(4.196, 0.769)		
4	(4.221, 1.339)	4	(4.164, 0.873)		
5	(4.355, 1.379)	5	(4.161, 0.966)		
6	(4.446, 1.418)	6	(4.182, 1.045)		
⋮	⋮	⋮	⋮		
45	(4.961, 1.875)	61	(4.957, 1.872)		
46	(4.962, 1.876)	62	(4.959, 1.873)		
47	(4.963, 1.877)	63	(4.960, 1.874)		
48	(4.964, 1.878)	64	(4.961, 1.875)		

EM1 = the EM algorithm method approximating the hazard function by the Taylor expansion;
EM2 = the EM gradient algorithm; NR is the Newton-Raphson method.

Table A1.

The systematic subset of the transformer lifetime data provided by Hong, Meeker, and McCalley (2009). (Unit: years).

Number of machines	Truncation indicator	Truncation time	Lifetime	Censoring indicator	Censoring time
2	1	*	1	1	*
14	1	*	3	0	3
23	1	*	4	0	4
2	1	*	5	1	*
10	1	*	6	0	6
10	1	*	8	0	8
9	1	*	10	0	10
8	1	*	11	0	11
2	1	*	12	1	*
2	1	*	13	1	*
5	1	*	14	1	*
3	1	*	15	1	*
2	1	*	16	1	*
3	0	7	17	1	*
5	1	*	19	0	19
2	0	14	20	1	*
6	1	*	22	0	22
2	1	*	23	0	23
3	1	*	24	0	24
3	1	*	25	0	25
8	1	*	26	0	26
1	0	4	27	1	*
1	0	14	28	1	*
2	0	10	29	1	*
1	0	14	30	1	*
1	0	10	31	1	*
2	0	12	32	1	*
2	0	12	33	1	*
11	0	5	35	0	35
16	0	7	36	0	36
1	0	12	37	1	*
3	0	22	38	1	*
16	0	12	40	0	40

Number of machines	Truncation indicator	Truncation time	Lifetime	Censoring indicator	Censoring time
22	0	12	41	0	41
1	0	18	42	1	*
19	0	16	44	0	44
5	0	18	46	0	46
5	0	18	47	0	47
9	0	20	48	0	48
3	0	20	49	0	49
7	0	22	51	0	51
7	0	24	53	0	53
8	0	26	55	0	55
10	0	28	57	0	57
3	0	30	59	0	59
2	0	32	60	0	60
1	0	32	61	0	61
1	0	38	61	1	*
1	0	38	66	0	66
1	0	40	69	0	69