<table>
<thead>
<tr>
<th>Title</th>
<th>Test for homogeneity in gamma mixture models using likelihood ratio</th>
</tr>
</thead>
<tbody>
<tr>
<td>Author(s)</td>
<td>Wong, ST; Li, WK</td>
</tr>
<tr>
<td>Citation</td>
<td>Computational Statistics &amp; Data Analysis, 2014, v. 70, p. 127-137</td>
</tr>
<tr>
<td>Issued Date</td>
<td>2014</td>
</tr>
<tr>
<td>URL</td>
<td><a href="http://hdl.handle.net/10722/194985">http://hdl.handle.net/10722/194985</a></td>
</tr>
<tr>
<td>Rights</td>
<td>NOTICE: this is the author’s version of a work that was accepted for publication in Computational Statistics &amp; Data Analysis. Changes resulting from the publishing process, such as peer review, editing, corrections, structural formatting, and other quality control mechanisms may not be reflected in this document. Changes may have been made to this work since it was submitted for publication. A definitive version was subsequently published in Computational Statistics &amp; Data Analysis, 2014, v. 70, p. 127-137. DOI: 10.1016/j.csda.2013.09.001; This work is licensed under a Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License.</td>
</tr>
</tbody>
</table>
Test for homogeneity in gamma mixture models using likelihood ratio

Tony Siu Tung Wong*, Wai Keung Li

Department of Statistics and Actuarial Science,
The University of Hong Kong, Hong Kong

Abstract

A testing problem of homogeneity in gamma mixture models is studied. It is found that there is a proportion of the penalized likelihood ratio test statistic that degenerates to zero. The limiting distribution of this statistic is found to be the chi-bar-square distributions. The degeneration is due to the negative-definiteness of a complicated random matrix, depending on the shape parameter under the null hypothesis. In light of this dependency, bounds on the distribution are introduced and a weighted average procedure is proposed. Simulation suggests that the results are accurate and consistent, and that the asymptotic result applies to the maximum likelihood estimator, obtained via an Expectation-Maximization algorithm.

Keywords: Chi-bar-square distributions, gamma mixture, likelihood ratio, maximum likelihood, negative definite

1. Introduction

In recent years, gamma mixture models have seen a surge of applications in many fields. Craig and Strassels (2010) examined the out-of-pocket prices of analgesic medications using a two-component gamma mixture model. See also Mayrose et al. (2005) for applications in bioinformatics and the references in Liu et al. (2003). Due to their importance, developing effective and handy statistical procedures for gamma mixture models is an imperative

*Corresponding author. Tel.: +852 28592469; fax: +852 28589041.
Email addresses: wongtonyst@hku.hk (Tony Siu Tung Wong), hrntlwk@hku.hk (Wai Keung Li)

Preprint submitted to Computational Statistics and Data Analysis April 9, 2014
task, in particular for the test of homogeneity. An obvious way of approaching the problem is to use the ordinary likelihood ratio test (LRT). One of the few results available is Liu et al. (2003). The authors showed that when the range of some parameters is unbounded, the LRT statistic diverges to infinity at a rate of $\log \log n$ and that its asymptotic behaviour is of extreme-value type through a highly complex piece of stochastic analysis. However, their simulation results suggested that the limiting distribution is far from converging to the extreme value distribution and that a possible solution is to simulate the finite-sample null distribution. The peculiar behaviour of the statistic arises because the maximum likelihood estimator (MLE) of some parameters may not be consistent. See, for example, the asymptotic result for $R_n(\varepsilon; I)$ in Chen and Chen (2001). Related problems in general mixture models were also addressed by Ghosh and Sen (1985), Dacunha-Castelle and Gassiat (1999), Chen and Chen (2001) and Liu and Shao (2003). In particular, Ghosh and Sen (1985) and Chen and Chen (2001) showed that the asymptotic distribution involves the supremum of a Gaussian process. See also Liu and Shao (2004) in normal mixture models. However, there are several shortfalls of the above results. Firstly, the results lose their appeal because the supremum of a Gaussian process is difficult to compute (Chen et al., 2001). Secondly, the divergence to infinity is so slow that it is not detected in simulation (Liu and Shao, 2004). The convergence of the test statistic, normalized by $\log \log n$, to the extreme value distribution is hardly detectable (Liu et al., 2003). Lastly, Hall and Stewart (2005) provided a theoretical analysis on the reduction of power against alternative hypotheses regarding the above results.

In light of the peculiar behaviour of LRT, a resampling approach (McLachlan, 1987; McLachlan and Peel, 2000; McLachlan and Khan, 2004) can be carried out. However, when some of the regularity conditions are restored, especially consistency of the estimator, it is of great theoretical significance to further investigate the likelihood ratio.

The consistency of the MLE in the test for homogeneity has not been solved until the introduction of a clever penalized procedure proposed by Chen et al. (2001). The authors innovated the modified likelihood ratio test (MLRT) by incorporating a penalty function. The MLRT was also developed by Chen and Kalbfleisch (2005) in normal mixture models and further extended to an EM-test by Li et al. (2009) and Chen and Li (2009). Exact theoretical results on the asymptotic null distribution have been obtained in some special cases. For densities with a single parameter of interest, the
MLRT statistic has the limiting distribution $0.5\chi_0^2 + 0.5\chi_1^2$ (Chen et al., 2001; Li et al., 2009). For the normal mixture model, the statistic has $\chi_2^2$ when the means and the variances are unequal and unknown (Chen and Li, 2009). Conceivably, the MLRT falls into the type II likelihood ratio problem (Lindsay, 1995, Section 4.4) which generates the chi-bar-square distributions of which some are parameter-dependent limiting null distributions. The above result in the normal mixture models returns to the $\chi_2^2$ distribution due to loss of strong identifiability (Chen and Li, 2009, Example 1). Qin and Smith (2006) investigated an extension of the MLRT in multivariate normal mixture models. The authors showed the asymptotic null distribution being a mixture of distributions and suggested it must be found using numerical methods. For models with multidimensional parameters, Zhu and Zhang (2004) analysed the asymptotic properties of LRT and Niu et al. (2011) considered an EM test. Although the problem of estimator consistency has been solved in MLRT and the EM-test, in many other mixture models, such as the gamma mixture models, the results $0.5\chi_0^2 + 0.5\chi_1^2$ or $\chi_2^2$ cannot be applied directly without theoretical justifications. The general testing problem has not been completely solved and remains as a long-standing open problem. Charnigo and Sun (2004) acknowledged the generalization of the MLRT to higher dimensional problems and suggested that the null distribution can be obtained by simulation. However, the extension is not at all straightforward as presented in this paper and simulation of the null distribution in the absence of a closed-form expression should no longer be tolerated. A clear guideline has been long overdue for practitioners in the rejection or retention of the homogeneity assumption. The purposes of the paper are to fill this research gap in gamma mixture models and to explore how the limiting null distribution depends on the parameters.

Motivated by the above needs and the importance of the gamma mixture models, this paper aims at investigating the limiting distribution of the MLRT statistic. We obtain the condition under which the MLRT statistic degenerates to zero and determine the proportion of degeneration. Then, we can show that the asymptotic null distribution has parameter-dependent chi-bar-square distributions. This subsequently establishes a foundation for quick model selection using the $\chi_3^2$ distribution in practice. Moreover, in light of the popular Expectation-Maximization (EM) algorithm for parameter estimation in finite mixture models, we demonstrate through intensive simulation studies that our results can be applied to the likelihood ratio statistic evaluated at the MLE obtained via the EM algorithm.
The article is organized as follows. In Section 2, we present the asymptotic results. Section 3 lists a number of considerations in the applications of the results. The asymptotic analysis is supplemented by simulation in Section 4. Section 5 presents two data examples and Section 6 gives a conclusion.

2. Asymptotic Results

We consider a two-parameter gamma density function

\[ f(x; \alpha, \beta) = \frac{1}{\Gamma(\alpha)} \beta^\alpha x^{\alpha-1} e^{-\beta x}, \quad x > 0, \]

where \( \alpha > 1 \) and \( \beta > 0 \) are shape and scale parameters, respectively. Given a set of independent and identically distributed data, we are interested in testing the homogeneity hypothesis \( H_0 \) against the alternative hypothesis of a two-component gamma mixture model \( H_1 \) where

\[
H_0 : f(x) = f(x; \alpha, \beta); \\
H_1 : f(x) = \pi f(x; \alpha_1, \beta_1) + (1 - \pi) f(x; \alpha_2, \beta_2),
\]

and \( 0 < \pi < 1 \) is a mixing proportion. In this paper, we study a very general testing problem that the parameters under the hypotheses are all unknown and unequal. This is completely different from the setting in Liu et al. (2003). For parametric hypothesis testing problems it is customary to use the ordinary LRT based on the statistic which is defined as

\[
L_{Rn} = 2 \left\{ L\left(\hat{\pi}, \hat{\alpha}_1, \hat{\beta}_1, \hat{\alpha}_2, \hat{\beta}_2 \right) - L\left(0.5, \hat{\alpha}, \hat{\beta}, \hat{\alpha}, \hat{\beta} \right) \right\},
\]

where

\[
L(\pi, \alpha_1, \beta_1, \alpha_2, \beta_2) = \sum_{i=1}^{n} \log \left\{ \pi f(x_i; \alpha_1, \beta_1) + (1 - \pi) f(x_i; \alpha_2, \beta_2) \right\} \quad (1)
\]

is the log-likelihood function and \( \left(\hat{\pi}, \hat{\alpha}_1, \hat{\beta}_1, \hat{\alpha}_2, \hat{\beta}_2 \right) \) is the MLE of parameter \( \left(\pi, \alpha_1, \alpha_2, \beta_1, \beta_2 \right) \). It is well known that the consistency of the MLE, obtained by maximizing (1) directly, is not guaranteed. See for example Ghosh and Sen (1985); Hathaway (1985); Chen and Chen (2001). This motivates a penalized procedure coined by Chen et al. (2001) based on a modified log-likelihood function

\[
L^p(\pi, \alpha_1, \beta_1, \alpha_2, \beta_2) = L(\pi, \alpha_1, \beta_1, \alpha_2, \beta_2) + c \log \left\{ 4\pi (1 - \pi) \right\}, \quad (2)
\]
where \( c \) is a positive constant corresponding to the level of modification. Denote by \( \left( \hat{\pi}^p, \hat{\alpha}_1^p, \hat{\beta}_2^p \right) \) the penalized MLE of \( (\pi, \alpha_1, \alpha_2, \beta_1, \beta_2) \) obtained by maximizing (2) given a suitable value of \( c \). Adding a penalty function to the log-likelihood regains the consistency of the penalized MLE (Chen et al., 2008; Chen and Li, 2009). The MLRT statistic is

\[
LR_n^p = 2 \left\{ LP \left( \hat{\pi}^p, \hat{\alpha}_1^p, \hat{\beta}_2^p \right) - L \left( 0.5, \hat{\alpha}, \hat{\beta} \right) \right\}. \tag{3}
\]

We study the asymptotic distribution of \( LR_n^p \), which can be expressed as

\[
LR_n^p = LR_{1n}^p - LR_{0n}
\]

in terms of the true parameter \( (\alpha_0, \beta_0) \) under \( H_0 \), where

\[
LR_{0n} = 2 \left\{ L \left( 0.5, \hat{\alpha}, \hat{\beta}, \hat{\alpha}, \hat{\beta} \right) - L \left( 0.5, \alpha_0, \beta_0, \alpha_0, \beta_0 \right) \right\};
\]

\[
LR_{1n}^p = 2 \left\{ LP \left( \hat{\pi}^p, \hat{\alpha}_1^p, \hat{\beta}_2^p \right) - L \left( 0.5, \alpha_0, \beta_0, \alpha_0, \beta_0 \right) \right\}.
\]

An immediate asymptotic approximation for \( LR_{0n} \) is

\[
\left( n^{-1/2} \sum_{i=1}^{n} Y_i^T \right) \left( n^{-1} \sum_{i=1}^{n} Y_i Y_i^T \right)^{-1} \left( n^{-1/2} \sum_{i=1}^{n} Y_i \right) + o_p(1),
\]

where \( Y_i \) is a random vector given by

\[
Y_i = \begin{cases} 
- \Gamma^{(1)}(\alpha_0) + \log \beta_0 + \log X_i \\
\alpha_0 \beta_0^{-1} - X_i
\end{cases}
\]

and \( \Gamma^{(k)}(\alpha) = d^k \ln \Gamma(\alpha)/d\alpha^k \). In Appendix A, we derive the following asymptotic approximation for \( LR_{1n}^p \)

\[
\left( n^{-1/2} \sum_{i=1}^{n} Y_i^T \right) \left( n^{-1} \sum_{i=1}^{n} Y_i Y_i^T \right)^{-1} \left( n^{-1/2} \sum_{i=1}^{n} Y_i \right) + o_p(1)
\]

(5)

if \( n^{-1/2} \sum_{i=1}^{n} U_i \) is non-negative-definite, where

\[
W_i = \left( \sum_{i=1}^{n} Z_i \tilde{\gamma}_2 Y_i^T \right) \left( \sum_{i=1}^{n} Y_i Y_i^T \right)^{-1} Y_i - Z_i \tilde{\gamma}_2,
\]
\( \hat{\gamma}_2 \neq 0 \) is the solution to \((\sum_{i=1}^{n} W_i W_i^T)^{-1} \sum_{i=1}^{n} U_i = I_2 \), \( I_2 \) is the two-dimensional identity matrix,

\[
U_i = Z_i - V_i Z_i, \quad V_i = \left( \sum_{j=1}^{n} Y_j^T \right) \left( \sum_{j=1}^{n} Y_j^T Y_j \right)^{-1} Y_i, \tag{6}
\]

and \( Z_i \) is a symmetric random matrix whose elements on the top left, top right and bottom right are, respectively

\[
\begin{align*}
Z_{i[11]} &= -\Gamma^{(2)}(\alpha_0) + \{ -\Gamma^{(1)}(\alpha_0) + \log \beta_0 + \log X_i \}^2; \\
Z_{i[12]} &= \beta_0^{-1} + \{-\Gamma^{(1)}(\alpha_0) + \beta_0 + \log X_i \} \left( \alpha_0 \beta_0^{-1} - X_i \right); \\
Z_{i[22]} &= -\alpha_0 \beta_0^{-2} + \left( \alpha_0 \beta_0^{-1} - X_i \right)^2. \tag{7}
\end{align*}
\]

The random quantity \( V_i \) is scalar. If \( n^{-1/2} \sum_{i=1}^{n} U_i \) is negative-definite, \( W_i \) are taken to be zero resulting in \( LR_n^p = \alpha_p (1) \). Under \( H_0 \), \( Y_i \) and \( Z_i \) are random with mean zero. Then, by the central limit theorem, \( n^{-1/2} \sum_{i=1}^{n} W_i \) converges to a bivariate normal random vector with mean zero. We summarize the result in Theorem 1.

**Theorem 1.** Under \( H_0 \), the asymptotic distribution of \( LR_n^p \) degenerates to zero with a weight \( 0 < p < 1 \) and has a \( \chi^2 \) distribution with a weight \( 1 - p \) where \( p \) is the probability that the random matrix to which the matrix \( n^{-1/2} \sum_{i=1}^{n} U_i \) converges as \( n \to \infty \) is negative-definite and \( U_i \) is defined in (6). That is,

\[
LR_n^p \sim p \chi^2_2 + (1 - p) \chi^2_2 \tag{8}
\]

for large \( n \), where \( \chi^2_2 \) is a degenerate distribution with all its mass at zero and the notation \( \sim \) means ‘is distributed like’.

The limiting distribution in (8) is known as the chi-bar-square distributions (Johnson et al., 1994, pg. 454). A more precise expression for \( p \) will be derived in Section 3.2. Hence, the above result will be restated by (10) indicating clearly the dependency on the shape parameter.

### 3. Practical Considerations

#### 3.1. Estimating \( p \)

From the definition of \( U_i \) in (6), we observe its dependence on the random vector \( Y_i \) and the random matrix \( Z_i \) given by (4) and (7), respectively, which
Figure 1: Plots of weight $p_s(\alpha_0, \beta_0, n)$ against $\alpha_0$. The left panel shows three series of $\beta_0 = 2$ ($\bigtriangleup$), $\beta_0 = 6$ ($\square$) and $\beta_0 = 10$ ($\triangle$). The right panel shows three series of $n = 100$ ($\circ$), $n = 500$ ($\square$) and $n = 1000$ ($\triangle$). The solid line depicts the asymptotic weight $p(\alpha_0)$.

are related to the parameter $(\alpha_0, \beta_0)$ under $H_0$. In addition, the estimate of $p$ may also depend on $n$ as the random matrix concerned involves a summation over $n$ random matrices. As a rough visualization of the relations between these variables, we simulate $n$ random variables from $f(x; \alpha_0, \beta_0)$, compute $Y_i, Z_i$ and $U_i$, and evaluate the proportions in 10000 replications that $n^{-1/2} \sum_{i=1}^{n} U_i$ is negative-definite. Denote by $p_s(\alpha_0, \beta_0, n)$ such a proportion. Fig. 1 displays two plots of $p_s(\alpha_0, \beta_0, n)$ at some selected values of $\alpha_0$, $\beta_0$ and $n$. The left panel shows three series of $p_s(\alpha_0, \beta_0, n)$ against $\alpha_0$ at $n = 1000$, each series corresponding to different values of $\beta_0$. There is a decreasing trend of $p_s(\alpha_0, \beta_0, n)$ as $\alpha_0$ increases, this trend being invariant in $\beta_0$. The right panel shows another three series of $p_s(\alpha_0, \beta_0, n)$ against $\alpha_0$ at $\beta_0 = 2$, each series corresponding to different values of $n$. A similar decreasing trend of $p_s(\alpha_0, \beta_0, n)$ against $\alpha_0$ is observed. In addition, the values of $p_s(\alpha_0, \beta_0, n)$ get lower at larger sample sizes and seem to converge to some certain level as $n$ grows. Overall, $p_s(\alpha_0, \beta_0, n)$ seems to decreases as $\alpha_0$ increases, but remains constant as $\beta_0$ varies. Its possible convergence as $n$ tends to infinity motivates further investigation. Last, it is worth pointing out some merits of the simulation technique. Apart from quick and easy construction of the weight estimate, its use in the construction of a lower bound in a finite samples will be outlined in Section 4.
3.2. Asymptotic \( p \)

We require some general conditions on \( Y_i \), \( Z_i \) and the products of their elements. In particular,

\[
n^{-1} \sum_{i=1}^{n} Y_i Y_i^T \to M, \quad n^{-1} \sum_{i=1}^{n} Y_{ij} Z_i \to v_j
\]

in probability for \( j = 1, 2 \), where \( Y_{ij} \) denotes the \( j \)th element of vector \( Y_i \). The expression of each of the elements in matrices \( M \) and \( v \) are given in Appendix B. Denote by \( U_{i[11]} \), \( U_{i[12]} \) and \( U_{i[22]} \), respectively the elements on the top left, top right and bottom right of \( U_i \). By the central limit theorem, the vector on the left-hand side below

\[
n^{-1/2} \left( \sum_{i=1}^{n} U_{i[11]} \right) \to N_3 \left\{ 0, \begin{pmatrix} \sigma_{11} & \sigma_{12} & \sigma_{13} \\ \sigma_{12} & \sigma_{22} & \sigma_{23} \\ \sigma_{13} & \sigma_{23} & \sigma_{33} \end{pmatrix} \right\}
\]

converges, as \( n \to \infty \), to a random vector denoted by \( S^T = (S_1, S_2, S_3) \) having a trivariate normal distribution \( N_3 \) with zero mean vector and covariance matrix whose elements are

\[
\begin{align*}
\sigma_{11} & = 2 \{ \Gamma(2) (\alpha_0) \}^2 + \Gamma(4) (\alpha_0) + \alpha_0 \{ \Gamma(3) (\alpha_0) \}^2 \left\{ 1 - \alpha_0 \Gamma(2) (\alpha_0) \right\}^{-1}; \\
\sigma_{12} & = \left[ -2 \Gamma(2) (\alpha_0) + \Gamma(3) (\alpha_0) \left\{ 1 - \alpha_0 \Gamma(2) (\alpha_0) \right\}^{-1} \right] \beta_0^{-1}; \\
\sigma_{13} & = \left[ 2 + \alpha_0 \Gamma(3) (\alpha_0) \left\{ -1 + \alpha_0 \Gamma(2) (\alpha_0) \right\}^{-1} \right] \beta_0^{-2}; \\
\sigma_{22} & = \left[ -1 + \Gamma(2) (\alpha_0) \left\{ -1 + \alpha_0 \Gamma(2) (\alpha_0) \right\} \left\{ -1 + \alpha_0 \Gamma(2) (\alpha_0) \right\}^{-1} \beta_0^{-2} \\
\sigma_{23} & = \left[ -2 \alpha_0 + \left\{ -1 + \alpha_0 \Gamma(2) (\alpha_0) \right\}^{-1} \right] \beta_0^{-3}; \\
\sigma_{33} & = \alpha_0 \left\{ 2 + 2 \alpha_0 \left\{ 1 - \alpha_0 \Gamma(2) (\alpha_0) \right\}^{-1} \right\} \beta_0^{-4}.
\end{align*}
\]

Hence, the negative-definiteness condition implies that \( p \) can be obtained by the following probability

\[
P \left( \{ S_1 < 0 \} \cap \{ S_1 S_3 - S_2^2 > 0 \} \right) = \int_{-\infty}^{0} \int_{-\infty}^{\infty} \int_{-\infty}^{s_3} g (s_1, s_2, s_3) \, ds_3 \, ds_2 \, ds_1,
\]

where \( g (s_1, s_2, s_3) \) is the density function of the above trivariate normal distribution. The probability can be easily evaluated by numerical integration using, for example, Wolfram Mathematica®. It is important to observe that
the integral is independent of $\beta_0$. This can be easily verified by simple transformation in the integration. Hence, the probability may precisely be denoted by $p(\alpha_0)$ and the result in (8) is more appropriately written as

$$LR_n^p \sim p(\alpha_0) \chi^2_0 + \{1 - p(\alpha_0)\} \chi^2_2.$$  

(10)

A more precise description of the relation of $p(\alpha_0)$ as $\alpha_0$ varies can be obtained by (9). Fig. 1 overlays a curve of $p(\alpha_0)$ against $\alpha_0$ for $1 \leq \alpha_0 \leq 10$. Its match with the simulated weight $p_\alpha(\alpha_0, \beta_0, n)$ suggests that the mysterious dependency of the proportion of degeneration on the shape parameter is further illustrated. An astounding observation is that $p(\alpha_0)$ is small at some large values of $\alpha_0$. An example $p(12)=0.0497$ suggests that the $\chi^2$ distribution is quite accurate to approximate the asymptotic null distribution. In addition, as an empirical rule of thumb, we may use that $p(\alpha_0) \leq p(1) =0.1345$ and $p(\alpha_0) > 0$ to develop a lower bound $D_L$ and an upper bound $D_U$ for the statistic $LR_n^p$

$$D_L \leq LR_n^p \leq D_U, \quad D_L \sim 0.1345 \chi^2_0 + 0.8655 \chi^2_2, \quad D_U \sim \chi^2_2$$  

(11)

as a quick guideline. Given a significance level, $H_0$ is retained if the observed MLRT statistic falls below the critical value evaluated by the above lower bound, and is rejected if it is above the critical value based on the upper bound.

### 3.3. Weighted Average Procedure

The previous subsections demonstrate the theoretical analysis to the testing problem. However, practical implementation of (10) encounters a drawback in that the value $\alpha_0$ is unknown. A possible solution is to substitute this value by the parameter estimate, for instance, the maximum likelihood estimate $\hat{\alpha}$ under $H_0$. Then, the weight $p(\alpha_0)$ is estimated by $\hat{\alpha}$ through (9) and the asymptotic null distribution is established as in (10). However, the substitution may suffer a certain degree of bias because all prior beliefs are placed on $\hat{\alpha}$. Lindsay (1995) suggested the use of the least favourable null distribution by employing the least favourable critical value within a confidence interval for $\alpha_0$. However, the problem remains unsolved if the observed test statistic falls below this least favourable critical value.

In light of the above difficulties, we propose a weighted average procedure to accommodate the estimation error. It is well-known that $\hat{\alpha} - \alpha_0$ is asymptotically normal with mean zero and variance $v(\alpha_0) = n^{-1} \alpha_0 \left\{ -1 + \alpha_0 \Gamma^{(2)}(\alpha_0) \right\}^{-1}$.
Table 1: Weight $p$ against $\alpha$ using weighted average procedure $p_w(\alpha, r)$ and direct substitution $p(\alpha)$. Different numbers of candidates $r$ and the effect of sample size $n$ for the weighted average procedure are shown.

<table>
<thead>
<tr>
<th>$\alpha$</th>
<th>$p_w(\alpha, 10)$</th>
<th>$p_w(\alpha, 20)$</th>
<th>$p(\alpha)$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$n = 100$</td>
<td>$n = 1000$</td>
<td>$n = 1000$</td>
<td>$n = 1000$</td>
</tr>
<tr>
<td>2</td>
<td>0.1023</td>
<td>0.1023</td>
<td>0.1023</td>
</tr>
<tr>
<td>4</td>
<td>0.0744</td>
<td>0.0743</td>
<td>0.0743</td>
</tr>
<tr>
<td>6</td>
<td>0.0612</td>
<td>0.0611</td>
<td>0.0613</td>
</tr>
<tr>
<td>8</td>
<td>0.0540</td>
<td>0.0536</td>
<td>0.0536</td>
</tr>
<tr>
<td>10</td>
<td>0.0509</td>
<td>0.0497</td>
<td>0.0497</td>
</tr>
</tbody>
</table>

Then, $r$ candidates of $\alpha_{0,k}$ can be obtained from the normal distribution through

$$
\frac{k}{r+1} = \int_{-\infty}^{\alpha_{0,k}} \left\{ \frac{1}{2\pi \sqrt{v(\hat{\alpha})}} \right\} \exp \left\{ -\frac{(x-\hat{\alpha})^2}{2v(\hat{\alpha})} \right\} dx, \quad k = 1, \ldots, r
$$

provided that $\alpha_{0,k} \geq 1$. Each of these $\alpha_{0,k}$ forms an asymptotic null distribution given by (8). The assignment of an equal weight to each $\alpha_{0,k}$ leads to the asymptotic null distribution

$$
p_w(\hat{\alpha}, r) \chi^2_0 + \left\{ 1 - p_w(\hat{\alpha}, r) \right\} \chi^2_2, \quad p_w(\hat{\alpha}, r) = \frac{1}{r} \sum_{k=1}^{r} p(\alpha_{0,k}).
$$

As illustrated in Fig. 1 the convexity of the weight in the shape parameter, the weighted average procedure will give a weight slightly larger than the direct substitution does. The effect of this finite-sample refinement is illustrated in Table 1. The weights $p_w(\hat{\alpha}, r)$ using $n = 100$ are slightly larger than those using $n = 1000$ which are very close to the value obtained by direct substitution $p(\hat{\alpha})$. Hence, this procedure tends to yield a smaller $p$-value than the method of direct substitution leading to a conclusion which is less favourable to the null hypothesis when information from the sample is scarce. Moreover, the input $r$ seems less important compared to the sample size. We shall fix $r = 10$ in data analysis in Section 5.2.

3.4. MLE Obtained via EM Algorithm

Mixture models are getting popular in the statistics literature because of its wide range of applications, including examination of homogeneity of populations, assessment of unimodality and identifications of clusters or outliers.
The introduction of the EM algorithm has further pushed up its popularity. Frühwirth-Schnatter (2006) commented that the EM algorithm is the most common method for parameter estimation in finite mixture models nowadays. However, the penalized procedure will not be considered for parameter estimation of a mixture model, except only when a test of homogeneity is conducted. Given the homogeneous and mixture models, the latter one is fitted to the data by the EM algorithm. The goodness-of-fit may be justified by comparing the values of the log-likelihood. Hence, the EM algorithm has retained the convenience of the ordinary LRT. We will investigate through simulation whether the goodness-of-fit justification is appropriate and under what circumstances it can be applied. Another problem inherited in the MLRT is the possible reduction of power under $H_1$. In the twelve cases under study in the simulation, the power is not seriously affected but the reduction in log-likelihood value due to the penalty function should not be overlooked. In light of this, we may use the following conventional likelihood ratio as an alternative statistic

$$LR_{itr,n}^{EM} = 2 \left\{ L \left( \hat{\alpha}_{1}^{EM}, \hat{\beta}_{1}^{EM}, \hat{\alpha}_{2}^{EM}, \hat{\beta}_{2}^{EM} \right) - L \left( 0.5, \hat{\alpha}, \hat{\beta} \right) \right\},$$

where $\left( \hat{\alpha}_{1}^{EM}, \hat{\beta}_{1}^{EM}, \hat{\alpha}_{2}^{EM}, \hat{\beta}_{2}^{EM} \right)$ is the MLE of $(\pi_1, \beta_1, \alpha_2, \beta_2)$ obtained via the EM algorithm and $itr$ is the number of EM algorithm iterations given a suitable initial guess. This statistic not only preserves the convenience as the ordinary LRT does, but part of it is also very common in the formation of AIC and BIC in mixture model selection.

The limiting distribution of $LR_{itr,n}^{EM}$ will be given after a brief discussion of the asymptotic characteristics of the EM estimators. In the rest of this subsection, we assume without loss of generality that $\pi \geq 0.5$. The argument in Chen and Chen (2001) points to the problem that in the ordinary LRT under $H_0$, the products $(1 - \hat{\pi}^{EM}) \hat{\alpha}_{2}^{EM}$ and $(1 - \hat{\pi}^{EM}) \hat{\beta}_{2}^{EM}$ are consistent but not $\hat{\alpha}_{2}^{EM}$ and $\hat{\beta}_{2}^{EM}$. The EM algorithm suffers a similar problem except that it can never reach the boundary point of $\pi$ and that the iterations will either slowly merge $\hat{\alpha}_{1}^{EM}$ with $\hat{\alpha}_{2}^{EM}$ and $\hat{\beta}_{1}^{EM}$ with $\hat{\beta}_{2}^{EM}$, or force $\hat{\pi}^{EM}$ towards one (Lindsay, 1995, Section 3.4). Denote by $E_I$ the former event that individual estimators are consistent and by $E_{II}$ the latter event that individual estimators are not consistent. The advantage of $LR_{itr,n}^{EM}$ is on the extremely slow convergence of the EM algorithm under $H_0$. The occurrence of $E_I$ or $E_{II}$ can be easily observed as the iterations proceed. If $E_{II}$ is observed, we may retain $H_0$ in the absence of a tolerable significance level;
otherwise, large values of $LR_{itr,n}^{EM}$ may suggest rejection of $H_0$ according to (10) conditional on $E_I$. Precisely,

$$LR_{itr,n}^{EM} | E_I \sim p(\alpha_0) \chi_0^2 + \{1 - p(\alpha_0)\} \chi_2^2$$  \hspace{1cm} (13)

as $n \to \infty$. The number of iterations $itr$ may be determined based on some stopping rules as outlined in Lindsay (1995). Our simulation results suggest that when a suitable initial guess, such as the penalized maximum likelihood estimate, is adopted, the increase in the likelihood function is not significant as the iterative run proceeds. Hence, practitioners may pick a number of $LR_{itr,n}^{EM}$ values after the EM algorithm has changed insignificantly and conclude whether to retain or to reject $H_0$ if these values yield consistent results. The above suggestions essentially preserve the convenience in the use of likelihood-ratio-type tests and avoid power deterioration in applications. The arguments and suggestions in this subsection will be supplemented by the material lifetimes example in Section 5.2.

4. Simulation

We have conducted an extensive simulation study to evaluate the accuracy of the results. Due to the dependency of $p(\alpha_0)$ on $\alpha_0$, it is interesting to conduct simulations using different values of $\alpha_0$ and holding $\beta_0 = 1$ with a number of sample sizes. The first statistic under study is $LR_n^p$ given by (3). It is the MLRT statistic with $c = \log 50$ in the penalty function in accordance with the recommendations in Chen et al. (2001). The second statistic $LR_{itr,n}^{EM}$ given by (12) uses the likelihood ratio evaluated at the MLE obtained via the EM algorithm. The extremely slow convergence in the EM algorithm makes simulation studies tedious. Lindsay (1995) pointed out that the solution of the likelihood equations can depend greatly on the initial values. Therefore, we use the penalized MLEs as initial guesses and carry out ten iterations.

We report the empirical sizes obtained from 10000 replications. Two sets of simulations are illustrated in Table 2. Other sets using different values of $\alpha_0$ share similar results and hence are not reported. The agreement between the theoretical results and the simulation results is obvious. Improvements are generally obtained when we increase the sample size. The simulation also shows the dependency of $p(\alpha_0)$ on $\alpha_0$. In Table 3, we report the proportions of zero statistics obtained from the simulation and the weight $p_s(\alpha_0, \beta_0, n)$ obtained from simulation in Section 3.1, and the asymptotic weight $p(\alpha_0)$ is

12
Table 2: Simulation results at selected nominal levels of 0.1, 0.05 and 0.01 using two homogeneous gamma models.

<table>
<thead>
<tr>
<th>n</th>
<th>( LR_p^n )</th>
<th>( LR_{10,n}^{EM} )</th>
<th>( LR_p^n )</th>
<th>( LR_{10,n}^{EM} )</th>
<th>( LR_p^n )</th>
<th>( LR_{10,n}^{EM} )</th>
</tr>
</thead>
<tbody>
<tr>
<td>100</td>
<td>0.1144</td>
<td>0.1285</td>
<td>0.0598</td>
<td>0.0701</td>
<td>0.0138</td>
<td>0.0176</td>
</tr>
<tr>
<td>200</td>
<td>0.1057</td>
<td>0.1117</td>
<td>0.0524</td>
<td>0.0562</td>
<td>0.0112</td>
<td>0.0135</td>
</tr>
<tr>
<td>500</td>
<td>0.1025</td>
<td>0.1034</td>
<td>0.0559</td>
<td>0.0569</td>
<td>0.0111</td>
<td>0.0114</td>
</tr>
<tr>
<td>1000</td>
<td>0.0972</td>
<td>0.0974</td>
<td>0.0471</td>
<td>0.0476</td>
<td>0.0103</td>
<td>0.0104</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>n</th>
<th>( LR_p^n )</th>
<th>( LR_{10,n}^{EM} )</th>
<th>( LR_p^n )</th>
<th>( LR_{10,n}^{EM} )</th>
<th>( LR_p^n )</th>
<th>( LR_{10,n}^{EM} )</th>
</tr>
</thead>
<tbody>
<tr>
<td>100</td>
<td>0.1073</td>
<td>0.1207</td>
<td>0.0563</td>
<td>0.0662</td>
<td>0.0121</td>
<td>0.0158</td>
</tr>
<tr>
<td>200</td>
<td>0.1044</td>
<td>0.1114</td>
<td>0.0560</td>
<td>0.0600</td>
<td>0.0143</td>
<td>0.0161</td>
</tr>
<tr>
<td>500</td>
<td>0.1053</td>
<td>0.1064</td>
<td>0.0532</td>
<td>0.0540</td>
<td>0.0093</td>
<td>0.0097</td>
</tr>
<tr>
<td>1000</td>
<td>0.1047</td>
<td>0.1052</td>
<td>0.0535</td>
<td>0.0541</td>
<td>0.0106</td>
<td>0.0107</td>
</tr>
</tbody>
</table>

in the caption. First, it is interesting that both statistics \( LR_p^n \) and \( LR_{10,n}^{EM} \) result in the same figures. This implies that the EM algorithm no longer increases the likelihood value under the occurrence of degeneration. Second, it is obvious that the asymptotic analysis leading to \( p(\alpha_0) \) agrees quite well with the simulation results of \( LR_p^n \) and \( LR_{10,n}^{EM} \). This consistently justifies one of the main results of this paper that the degeneration arises from the negative-definiteness of the random matrix. The relatively weak approximation in the sample size of 100 can be explained by the relatively weak second-order approximation given by (5). Lastly, the value \( p^s(\alpha_0, \beta_0, n) \) is the largest when the sample size is less than 1000. We may replace the lower bound given by (11) by \( p^s(\alpha_0, \beta_0, n) \) if being smaller as a more prudent benchmark in a finite-sample situation.

Some insights on the power of the tests can be gained. We consider a number of gamma mixture models which are either entirely different in mixing proportion, shape and scale parameters or with some of these parameters being equal. Each of the following alternative hypotheses is formulated to
Table 3: Proportions of zero statistics and simulated weights $p_s(\alpha_0, \beta_0, n)$ using four homogeneous gamma models. The asymptotic weights for the four cases are $p(2) = 0.1023$, $p(6) = 0.0576$, $p(8) = 0.0536$ and $p(12) = 0.0497$.

<table>
<thead>
<tr>
<th>$n$</th>
<th>$LR_n^p$</th>
<th>$LR_{10,n}^EM$</th>
<th>$p_s(2, 1, n)$</th>
<th>$LR_n^p$</th>
<th>$LR_{10,n}^EM$</th>
<th>$p_s(6, 1, n)$</th>
</tr>
</thead>
<tbody>
<tr>
<td>100</td>
<td>0.1195</td>
<td>0.1195</td>
<td>0.1636</td>
<td>0.0516</td>
<td>0.0516</td>
<td>0.0925</td>
</tr>
<tr>
<td>200</td>
<td>0.1258</td>
<td>0.1258</td>
<td>0.1596</td>
<td>0.0484</td>
<td>0.0484</td>
<td>0.0839</td>
</tr>
<tr>
<td>500</td>
<td>0.1206</td>
<td>0.1206</td>
<td>0.1382</td>
<td>0.0542</td>
<td>0.0542</td>
<td>0.0763</td>
</tr>
<tr>
<td>1000</td>
<td>0.1225</td>
<td>0.1225</td>
<td>0.1301</td>
<td>0.0545</td>
<td>0.0545</td>
<td>0.0649</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>$n$</th>
<th>$LR_n^p$</th>
<th>$LR_{10,n}^EM$</th>
<th>$p_s(8, 1, n)$</th>
<th>$LR_n^p$</th>
<th>$LR_{10,n}^EM$</th>
<th>$p_s(12, 1, n)$</th>
</tr>
</thead>
<tbody>
<tr>
<td>100</td>
<td>0.0465</td>
<td>0.0465</td>
<td>0.0731</td>
<td>0.0349</td>
<td>0.0349</td>
<td>0.0578</td>
</tr>
<tr>
<td>200</td>
<td>0.0470</td>
<td>0.0470</td>
<td>0.0674</td>
<td>0.0324</td>
<td>0.0324</td>
<td>0.0439</td>
</tr>
<tr>
<td>500</td>
<td>0.0384</td>
<td>0.0384</td>
<td>0.0592</td>
<td>0.0336</td>
<td>0.0336</td>
<td>0.0357</td>
</tr>
<tr>
<td>1000</td>
<td>0.0444</td>
<td>0.0444</td>
<td>0.0473</td>
<td>0.0297</td>
<td>0.0297</td>
<td>0.0259</td>
</tr>
</tbody>
</table>

Every simulation experiment consists of 10000 replications, each of sample size 1000. We find that the upper bound $\chi^2_2$ given by (11) is extremely efficient in the testing process. Almost all simulated test statistics of $LR_n^p$ and $LR_{10,n}^EM$ are greater than the critical values of the $\chi^2_2$ distribution. The powers are all equal to one at significance levels 0.1, 0.05 and 0.01 except for the test of $H_{104}$. In this particular case, the statistic $LR_n^p$ gives powers of 0.9999, 0.9997 and 0.9985 at the corresponding significance levels, whereas $LR_{itr,n}^EM$ yields powers of 0.9999, 0.9997 and 0.9987. Both methods seem to be equally powerful. However, we should point out that the EM algorithm increases the likelihood value at each cycle in the iterative sequence (Dempster et al., 1977). Meanwhile, the penalty function in the MLRT may reduce the log-likelihood.

In Table 4, we report the average test statistics in 10000 replications for each of the alternative hypotheses. That the averages of $LR_{10,n}^EM$ are always higher
Table 4: Average values of test statistics under alternative hypotheses.

<table>
<thead>
<tr>
<th></th>
<th>$H_{101}$</th>
<th>$H_{102}$</th>
<th>$H_{103}$</th>
<th>$H_{104}$</th>
<th>$H_{105}$</th>
<th>$H_{106}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$LR^E_{10}$</td>
<td>576.1</td>
<td>487.5</td>
<td>780.4</td>
<td>43.54</td>
<td>61.82</td>
<td>61.82</td>
</tr>
<tr>
<td>$LR^M_{10,n}$</td>
<td>884.3</td>
<td>783.7</td>
<td>781.2</td>
<td>43.79</td>
<td>76.23</td>
<td>76.21</td>
</tr>
<tr>
<td></td>
<td>$H_{107}$</td>
<td>$H_{108}$</td>
<td>$H_{109}$</td>
<td>$H_{110}$</td>
<td>$H_{111}$</td>
<td>$H_{112}$</td>
</tr>
<tr>
<td>$LR^E_{10}$</td>
<td>470.8</td>
<td>129.0</td>
<td>341.4</td>
<td>62.63</td>
<td>124.9</td>
<td>61.82</td>
</tr>
<tr>
<td>$LR^M_{10,n}$</td>
<td>558.5</td>
<td>140.8</td>
<td>342.2</td>
<td>62.72</td>
<td>125.6</td>
<td>74.64</td>
</tr>
</tbody>
</table>

implies that the use of $LR^M_{10,n}$ may achieve higher power given some extremely small significance level or in some special cases that have not been considered in the simulation.

5. Data Example

5.1. Danish Fire Loss

This example is based on the Danish fire loss data set which consists of 2157 losses exceeding one million Danish Krone from the years 1980 to 1990 inclusive. It is well known that the data set has a heavy right tail in the extreme value literature (Embrechts et al., 1997). The adequacy of the homogeneous gamma model which has a moderate tail is suspected. We apply the proposed results and methods to see if a two-component gamma mixture model will improve the fitting with further verifications, justified by some goodness-of-fit measures. McNeil (1997) provided a time series plot to check for clustering of large losses and a sample mean excess function to determine heavy-tailed behaviour. The results suggest the validity of the independence assumption and the possibility in modelling excesses over high thresholds using the generalized Pareto distribution. Recently, Wong and Li (2010) proposed a threshold model incorporating the generalized Pareto distribution for excesses and a Weibull distribution for the rest of the observations. This threshold model flexibly gives a global fit and an appropriate tail modelling. These two findings suggest that the loss data are likely to be independent but from a heterogeneous population.

The maximum likelihood estimate of a gamma model is $(\hat{\alpha}, \hat{\beta}) = (1.299, 0.382)$ with a corresponding maximized log-likelihood of -4752. In the gamma mix-
ture model, the penalized procedure and the EM algorithm give

\[
\begin{align*}
\left( \hat{\alpha}^P, \hat{\beta}_1^P, \hat{\beta}_2^P \right) &= (0.5098,15.68,10.09,1.226,0.2040) ; \\
\left( \hat{\alpha}^{EM}, \hat{\beta}_1^{EM}, \hat{\beta}_2^{EM} \right) &= (0.2816,1.256,0.1619,10.19,6.036)
\end{align*}
\]

which yield the values of the test statistic of 1829 and 1978, respectively. The evidence for the mixture model is overwhelming as both statistics are far greater than the critical values of the \(\chi^2\) distribution at any reasonable significance level. Further support for this is a goodness-of-fit assessment based on probability-probability plots as shown in Fig. 2. The gamma mixture model provides a much better fit as the plot exhibits obviously a straight line pattern. This example lends further support to the asymptotic distribution in (10), improvement in power through the use of \(LR_{itr,n}^{EM}\), and demonstrates its simplicity in implementation.

5.2. Material Lifetimes

Gamma distributions give useful representations of a number of physical situations such as random processes in time. We consider a set of 101 observations for the lifetime of an aluminium sheet under maximum stress of 21,000 psi. A brief description and the data listed in increasing order are available in Birnbaum and Saunders (1958). The authors demonstrated a realistic adjustment to exponential models in representing lifetimes in a life-testing situation. Therefore, it is interesting to check the redundancy of a mixture structure in the representation. More insight may be gained by applying our results in studying the data set.

Figure 2: The probability-probability plots. The left panel shows the plot of the fitted gamma model and the right panel shows that of the fitted gamma mixture model using the EM algorithm. A 45-degree straight line is given for reference.
Table 5: Estimation results of the EM algorithm for the material lifetimes data. The $p$-values are calculated using (13) based on (a) direct substitution and (b) weighted average procedure.

<table>
<thead>
<tr>
<th>$itr$</th>
<th>Parameter estimate</th>
<th>$LR^\text{EM}_{itr,n}$</th>
<th>$p$-value (a)</th>
<th>$p$-value (b)</th>
</tr>
</thead>
<tbody>
<tr>
<td>10</td>
<td>$(0.5048, 22.73, 0.01452, 10.25, 0.008307)$</td>
<td>2.389</td>
<td>0.2879</td>
<td>0.2872</td>
</tr>
<tr>
<td>50</td>
<td>$(0.5222, 22.20, 0.01425, 10.15, 0.008262)$</td>
<td>2.397</td>
<td>0.2867</td>
<td>0.2860</td>
</tr>
<tr>
<td>100</td>
<td>$(0.5446, 21.53, 0.01389, 9.999, 0.008175)$</td>
<td>2.408</td>
<td>0.2851</td>
<td>0.2845</td>
</tr>
</tbody>
</table>

The parameter estimate of a gamma model is $(\hat{\alpha}, \hat{\beta}) = (11.86, 0.008462)$. In the gamma mixture model, the penalized maximum likelihood estimate of $(\hat{\pi}^p, \hat{\alpha}_1^p, \hat{\beta}_1^p, \hat{\alpha}_2^p, \hat{\beta}_2^p) = (0.5001, 22.82, 0.01457, 10.26, 0.008307)$

leads to a value of the MLRT statistic of 2.387. A lower bound of the $p$-value evaluated through (11) is 0.2624, larger than any reasonable size of a statistical test. In the absence of $H_0$, direct substitution and the weighted average procedure yields $p(\hat{\alpha})=0.0495$ and $p_w(\hat{\alpha}, 10)=0.0518$, respectively. The corresponding $p$-values are 0.2882 and 0.2875. The use of the penalized estimates initiates the EM iterative sequence. A series of 1000 iteration steps seems to indicate the occurrence of event $E_{II}$ that individual estimators are not consistent as shown in the left panel of Fig. 3 in which $\hat{\pi}^\text{EM}$ increases slowly to one as the iteration moves on. Evidence in favour of $H_0$ is obvious. On the other hand, the behaviour of $LR^\text{EM}_{itr,n}$ is agonizing, in particular as $\hat{\pi}^\text{EM}$ is closer to one that a jump in the test statistic is observed in the right panel of Fig. 3. The AIC criterion starts to reject $H_0$ in the 833rd iteration whereas the BIC criterion and the statistic $LR^\text{EM}_{itr,n}$ at the 5% significance level consistently suggest retention of $H_0$ in all 1000 iterations. However, the insignificant increase of the statistic in the first 600 iterations suggests early termination of the EM algorithm. Therefore, the suggestions in Section 3.4 are useful. We can consider a number of the test statistics in different iterative steps and apply (13). The results reported in Table 5 consistently suggest the retention of $H_0$ in agreement with the method of MLRT.
6. Conclusion

We investigate the modified likelihood ratio test for homogeneity in two-component gamma mixture models. We have found that the limiting distribution of the test statistic is the parameter-dependent chi-bar-square distributions given by a degeneration to zero with weight \( p(\alpha_0) \) and a chi-square distribution with two degrees of freedom with weight \( 1 - p(\alpha_0) \). This weight is related to the negative-definiteness of a complicated random matrix dependent on the shape parameter of the homogeneous gamma model. An asymptotic approximation using a trivariate normal probability has been developed for \( p(\alpha_0) \). All these theoretical results have been revealed through an extensive simulation to be very accurate and reliable.

In applications, the shape parameter is unknown. Based on the behaviour of \( p(\alpha_0) \), we have developed a lower bound for the retention of the homogeneous hypothesis and an upper bound for the rejection. The bounds have been proved to be extremely useful in simulation and in two real examples. On the rare occasion that the observed test statistic falls between the bounds or if practitioners require an evaluation of the \( p \)-value, we recommend the weighted average procedure which takes into account the estimation error of the shape parameter. This procedure has yielded consistent results in a study of the material lifetimes data.

Due to the popularity of the EM algorithm in the analysis of mixture models, we recommend that the likelihood ratio test statistic be evaluated at the maximum likelihood estimates obtained via the EM algorithm. There are some appealing advantages including the preservation of the convenience of the conventional likelihood ratio test procedure and in the prevention of
power reduction. The fact that the EM iterative sequence converges slowly allows the selection of a number of observed test statistics. A decision may therefore be based on these statistics by comparing the derived asymptotic null distribution conditional on the consistency of individual estimators. Its simplicity and convenience have been illustrated in the real examples.

A number of interesting insights have been obtained on the form of the asymptotic null distribution and on its practical implementation. We believe many other mixture models share similar characteristics and this deserves further research and discussion. In particular, the parameter-dependent structure of the limiting distribution may not be as simple as our situation in which only the shape parameter is involved. Developing simple decision criteria such as bounds appears to be very challenging.

Acknowledgement

Partial support by the Area of Excellence Scheme under the University Grants Committee of the Hong Kong Special Administration Region, China (Project AoE/P-04/2004 and GRF grant HKU7036/06P) is acknowledged. We thank Dr. G. Li of The University of Hong Kong, the Co-Editor, an Associate Editor and two reviewers for their constructive comments and suggestions that greatly led to the improvement of the paper.

Appendix A. Approximation for $LR_{1n}^p$

Define

$$Y_i = \frac{1}{f(X_i; \alpha_0, \beta_0)} \left\{ \frac{\partial f(X_i; \alpha_0, \beta_0)}{\partial \alpha} \right\} ; \quad Z_i = \frac{1}{f(X_i; \alpha_0, \beta_0)} \left\{ \frac{\partial^2 f(X_i; \alpha_0, \beta_0)}{\partial \alpha^2} \frac{\partial^2 f(X_i; \alpha_0, \beta_0)}{\partial \alpha \partial \beta} \frac{\partial^2 f(X_i; \alpha_0, \beta_0)}{\partial \beta^2} \right\}$$

whose expressions for the gamma model are given by (4) and (7), respectively. The penalty function regains the consistency and efficiency of the penalized maximum likelihood estimators (Chen et al., 2008; Chen and Li, 2009). Following Section 2.3 of Chen et al. (2000), but in a bivariate context below, the resulting characterization of $LR_{1n}^p$ involves the maximum of the following function

$$2 \sum_{i=1}^{n} \delta_i - \sum_{i=1}^{n} \delta_i^2$$

19
plus $o_p(1)$, where

$$
\delta_i = \pi \left( \frac{\alpha_1 - \alpha_0}{\beta_1 - \beta_0} \right)^T Y_i + \pi \left( \frac{\alpha_1 - \alpha_0}{\beta_1 - \beta_0} \right)^T \left( \frac{\alpha_2 - \alpha_0}{\beta_2 - \beta_0} \right)^T Y_i + (1 - \pi) \left( \frac{\alpha_2 - \alpha_0}{\beta_2 - \beta_0} \right)^T \left( \frac{\alpha_2 - \alpha_0}{\beta_2 - \beta_0} \right)^T Z_i.
$$

A re-parameterization using vector parameters $\gamma_1$ and $\gamma_2$, where

$$
\gamma_1 = \left\{ \pi \left( \frac{\alpha_1 - \alpha_0}{\beta_1 - \beta_0} \right) + (1 - \pi) \left( \frac{\alpha_2 - \alpha_0}{\beta_2 - \beta_0} \right) \right\}; \quad \gamma_2 = \left\{ 0.5 \pi (1 - \pi) \right\}^{1/2} \left( \frac{\alpha_1 - \alpha_2}{\beta_1 - \beta_2} \right)
$$

leads to

$$
LR^{p}_{1n} = \max_{\gamma_1, \gamma_2} q(\gamma_1, \gamma_2) + o_p(1),
$$

where

$$
q(\gamma_1, \gamma_2) = 2 \sum_{i=1}^{n} \left( \gamma_1^T Y_i + \frac{1}{2} \gamma_1^T Z_i \gamma_1 + \gamma_2^T Z_i \gamma_2 \right) - \sum_{i=1}^{n} \left( \gamma_1^T Y_i + \frac{1}{2} \gamma_1^T Z_i \gamma_1 + \gamma_2^T Z_i \gamma_2 \right)^2.
$$

From Lemma 1 in Charnigo and Sun (2004) and Lemma A2 in Li et al. (2009) that $\hat{\gamma}_1^p = O_p(n^{-1/2})$ and by the strong law of large numbers that

$$
n^{-1} \sum_{i=1}^{n} Z_i = o_p(1).$$

It follows that

$$
LR^{p}_{1n} = \max_{\gamma_1, \gamma_2} q^*(\gamma_1, \gamma_2) + o_p(1),
$$

where

$$
q^*(\gamma_1, \gamma_2) = 2 \sum_{i=1}^{n} \left( \gamma_1^T Y_i + \gamma_2^T Z_i \gamma_2 \right) - \sum_{i=1}^{n} \left( \gamma_1^T Y_i + \gamma_2^T Z_i \gamma_2 \right)^2.
$$

The maximum value of $q^*(\gamma_1, \gamma_2)$ is (5) excluding the term $o_p(1)$.

**Appendix B. Asymptotic weight $p(\alpha_0)$**

Denote the matrices $M$ and $v_j$ by

$$
M = \begin{pmatrix} m_{11} & m_{12} \\ m_{12} & m_{22} \end{pmatrix}^{-1}; \quad v_j = \begin{pmatrix} v_{j11} & v_{j12} \\ v_{j12} & v_{j22} \end{pmatrix},
$$

20
where
\[ m_{11} = \frac{\alpha_0}{1 + \alpha_0 \Gamma(\sigma(\alpha_0))}; \quad m_{12} = \frac{\beta_0}{1 + \alpha_0 \Gamma(\sigma(\alpha_0))}; \quad m_{22} = \frac{\beta_0^2}{1 + \alpha_0 \Gamma(\sigma(\alpha_0))}; \]
\[ v_{111} = \Gamma(3)(\alpha_0); \quad v_{112} = 0; \quad v_{122} = \frac{1}{\beta_0^2}; \quad v_{222} = -\frac{2\alpha_0}{\beta_0^3}. \]

Then, we express \( U_{i[11]}, U_{i[12]} \) and \( U_{i[22]} \) by
\[ U_{i[11]} = Z_{i[11]} - Y_{i[1]} (m_{11}v_{111} + m_{12}v_{211}) - Y_{i[2]} (m_{12}v_{111} + m_{22}v_{211}); \]
\[ U_{i[12]} = Z_{i[12]} - Y_{i[1]} (m_{11}v_{112} + m_{12}v_{212}) - Y_{i[2]} (m_{12}v_{112} + m_{22}v_{212}); \]
\[ U_{i[22]} = Z_{i[22]} - Y_{i[1]} (m_{11}v_{122} + m_{12}v_{222}) - Y_{i[2]} (m_{12}v_{122} + m_{22}v_{222}). \]

The result in Section 3.2 follows from the central limit theorem and the covariance matrix is obtained by
\[
\begin{pmatrix}
\sigma_{11} & \sigma_{12} & \sigma_{13} \\
\sigma_{12} & \sigma_{22} & \sigma_{23} \\
\sigma_{13} & \sigma_{23} & \sigma_{33}
\end{pmatrix}
= \begin{pmatrix}
E(U_{i[11]}U_{i[11]}) & E(U_{i[11]}U_{i[12]}) & E(U_{i[11]}U_{i[22]}) \\
E(U_{i[11]}U_{i[12]}) & E(U_{i[12]}U_{i[12]}) & E(U_{i[12]}U_{i[22]}) \\
E(U_{i[11]}U_{i[22]}) & E(U_{i[12]}U_{i[22]}) & E(U_{i[22]}U_{i[22]})
\end{pmatrix}.
\]

References


