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An Alternative Method for Prediction Intervals of an Ordered Observation from Pareto value Distribution Based on Censored Sample

Hsiaoting Chiu, Hai-Lin Lu

Hsiao-Ting Chiu

Department of Information Management

Hai-Wen Lu

Department of Information Management

Chia-Nan University

Department of E-commerce

Chia-Nan University 886-6-3663994

886-6-3662019

Nanhua University 886-6-3663994

htchiu@mail.chna.edu.tw

huangi@mail.chna.edu.tw

hwlu@mail.nhu.edu.tw

ABSTRACT

Pareto Distribution is widely used in biostatistics and economics areas. In this paper, we provide some suitable pivotal quantities for estimating the prediction intervals of the jth future ordered observation in a sample of size n from the pareto distribution based on doubly type II censored samples. We also give simulation study to analyze its feasibility of pivotal quantities. Finally, two illustrative examples are also included.

Categories

G.3 [PROBABILITY AND STATISTICS]: Probabilistic algorithms---Reliability and life testing, Random number generation

General Terms

Reliability

Keywords: pivotal quantity, doubly type II censored samples, pareto distribution, approximation maximum likelihood estimation.

1. INTRODUCTION

In most researching of reliability, the Weibull distribution is widely used as a model of lifetime data (Bain and Engelhardt [2], Agresti [1). Let us consider the two-parameter Weibull distribution with probability density function (pdf)

$$w(t) = \frac{\beta}{\alpha} \left(\frac{t}{\alpha}\right)^{\beta - 1} \exp\left\{-\left(\frac{t}{\alpha}\right)^{\beta}\right\}, \quad t > 0,$$
 (1)

and cumulative distribution function(cdf)

$$W(t) = 1 - \exp\left\{-\left(\frac{t}{\alpha}\right)^{\beta}\right\},\tag{2}$$

where > 0 and > 0 are the shape and scale parameters, respectively.

Pareto distribution is similar to Weibull distribution. For convenience, we adopt Weibull distribution to explain the process. It is worth noting that if T is a random variable having the Weibull cdf given by formula (2), then the random variable X = lnT is distributed as a smallest Type I extreme value variate with pdf

Pareto distribution with probability density function (pdf)

$$f(x) = \frac{\beta}{\alpha} (1 + \frac{y - \mu}{\sigma})^{-(\beta + 1)} \qquad \begin{array}{c} \mu < y < \infty, \\ -\infty < \mu < \infty, \\ \sigma > 0, \beta > 0 \end{array}$$
(3)

and cumulative distribution function(cdf)

$$F(x) = 1 - (1 + \frac{y - \mu}{\sigma})^{-\beta}$$
 (4)

where μ and σ are location and scale parameters, respectively. The notation $Y \sim P(\mu, \sigma, \beta)$ will be used to indicate that a random variable X has c.d.f. (1).

In life testing studies, several lifetimes of units put on test may not be observed due to time limitations or money and material resources restrictions on data collection. Consider an experiment in which n identical components are placed on test simultaneously. Suppose the experiment was terminated when the (n-s)th component failed, thus censoring the last s components. Such a sample is called Type II right censored sample. If some initial r observations are also censored, it is called Type II doubly censored.

The studies of estimating the prediction intervals of the future

data are quite important and valuable in lifetime analysis. There have been several studies in the literature dealing with such problems. For the exponential distribution, Lawless[9] and Likes[12] estimated the prediction intervals based on the order statistics, $X_{(j)}(r < j \le n)$, of a sample while the first rdata of the sample were observed. Mann and Grubbs[15] proposed an alternative method to construct approximate prediction intervals. Kaminsky and Nelson[8] constructed prediction intervals by using the best linear unbiased estimates (BLUE) of the parameters as a pivotal statistic. For the Weibull distribution, Mann and Saunders[16] used three specially selected order statistics to predict the minimum of a single future sample. Engelhardt and Bain[5] constructed the prediction limits for the jth smallest of some set of future observations. Fertig et al.[6] provided Monte Carlo estimates of percentiles of the distribution of a statistics S for constructing prediction intervals of a future observation. Lawless[10] used a conditional method to obtain a prediction interval for the first order statistic of a set of future observations, based on previous

data; Hsieh[7] used the same technique to construct prediction

intervals for future observations. Mann and Fertig[14] constructed the tables for obtaining the best linear invariant

estimates (BLIE) of parameters. Balakrishnan and Cohen[3]

proposed an approximate maximum likelihood

(AMLE) of parameters. All these researches are under the scheme that the available data is either right censored or doubly censored.

It is well known that the Type II censored data, the right, left

and doubly censored data are all special cases of multiple censored data. In this paper, we consider the general case of the multiple Type II censored data scheme. Suppose n components are placed on test in life testing. The lifetime of the first r, the middle l, and the last s components are assumed unobserved or missing. That is, we assume $X_{(r+1)} < X_{(r+2)} < ... < X_{(r+k)}$ and $X_{(r+k+l+1)} < X_{(r+k+l+2)} < ... < X_{(n-s)}$ are observable and no others. In practice, multiple Type II censored problems may arise when some components failed between two points of observation with exact times of these

Balakrishnan [4]). In next section, following the ideas of Wu *et al.* [18], we present our method of constructing the prediction intervals of the future unknown observations for Type II censored data. We describe the procedure for calculating the percentiles of the distributions of the pivotal quantities, and the simulation results are compared with the existing method in section 3 and 4, respectively. In section 5, we illustrate our method with two examples. A brief discussion is presented in section 6.

failure unobservable components (Balasubramanian and

2. A GENERAL FORM OF PIVOTAL QUANTITY

The prediction intervals of our method for $X_{(j)}$ are based on a subset $\{X_{(n_i)}\}_{i=1}^c$ of $\{X_{(k)}\}_{k=1}^d$, where $1 \le n_1 < n_2 < \ldots < n_c \le d < j \le n$. Let $Y_i = \frac{X_i - \mu}{\sigma}$, then Y_i has extreme value distribution with $\mu = 0$ and $\mu = 1$. And $Y_{(i)} = \frac{X_{(i)} - \mu}{\sigma}$ is the ith order statistic of Y_i . We define some

pivotal quantities (proof shown in Appendix I) of the following general forms.

$$\widetilde{U}_h = \frac{X_{(j)} - X_{(n_c)}}{\widetilde{W}_i}, \quad h = 1, ..., 4, \quad n - s < j \le n,$$

where

$$\widetilde{W}_{1} = \sum_{i=2}^{c} \frac{g(E(Y_{(n_{i})}))}{\sum_{i=2}^{c} g(E(Y_{(n_{i})}))} (X_{(n_{i})} - X_{(n_{1})}), \tag{5}$$

$$\begin{split} \widetilde{W}_{2} &= \sum_{i=1}^{n_{1}-1} \frac{g(E(Y_{(i)}))}{\sum_{t=1}^{n} g(E(Y_{(t)}))} (X_{(n_{2})} - X_{(n_{1})}) \\ &+ \sum_{i=2}^{c} \frac{g(E(Y_{(i)}))}{\sum_{t=1}^{n} g(E(Y_{(t)}))} (X_{(n_{i})} - X_{(n_{1})}) \\ &+ \sum_{i=1}^{c-1} \sum_{j=n_{i}+1}^{n_{i+1}-1} \frac{g(E(Y_{(i)}))}{\sum_{t=1}^{n} g(E(Y_{(t)}))} \times \left(\frac{X_{(n_{i})} + X_{(n_{i+1})} - 2X_{(n_{1})}}{2} \right) \\ &+ \sum_{i=n_{c}+1}^{n} \frac{g(E(Y_{(i)}))}{\sum_{t=1}^{n} g(E(Y_{(t)}))} (X_{(n_{c})} - X_{(n_{1})}), \end{split}$$

 $\widetilde{W}_{3} = \prod_{i=0}^{c} (X_{(n_{i})} - X_{(n_{1})})^{\frac{g(E(Y_{(n_{i})}))}{\sum_{t=2}^{c} g(E(Y_{(n_{t})}))}},$ (7)

$$\widetilde{W}_{4} = \left(\prod_{i=1}^{n_{1}-1} (X_{(n_{2})} - X_{(n_{1})})^{\frac{g(E(Y_{(n_{i})}))}{\sum_{i=1}^{n} g(E(Y_{(n_{i})}))}}}\right)$$

$$\times \left(\prod_{i=2}^{c} (X_{(n_{i})} - X_{(n_{1})})^{\frac{g(E(Y_{(n_{i})}))}{\sum_{i=1}^{n} g(E(Y_{(n_{i})}))}}\right)$$

$$\times \prod_{i=1}^{c-1} \prod_{i=n_{i}+1}^{n_{i+1}-1} \left(\frac{X_{(n_{i})} + X_{(n_{i+1})} - 2X_{(n_{1})}}{2}\right)^{\frac{g(E(Y_{(i)}))}{\sum_{i=1}^{n} g(E(Y_{(i)}))}}$$

$$\times \left(\prod_{i=n_{c}+1}^{n} (X_{(n_{c})} - X_{(n_{1})})^{\frac{g(E(Y_{(i)}))}{\sum_{i=1}^{n} g(E(Y_{(i)}))}}\right)$$
(8)

 \widetilde{W}_1 and \widetilde{W}_2 come from the ideas of arithmetic means, and \widetilde{W}_{3} and \widetilde{W}_{4} follow the concepts of geometric means. The equations (5) to (8) are general forms. Wu et al. [18] consider the case where every datum of different position has the same weight. In the case of extreme value distribution, it seems reasonable to assume that the weight of each datum point should be different for different position. From the properties of the extreme value distribution, we suggest that the weighted factors are equal to $g(E(Y_{(n_i)})) / \sum_{i=1}^n g(E(Y_{(n_i)}))$ in \widetilde{W}_1 , where $E(Y_{(n_i)})$ is the expected values of $Y_{(n_i)}$. The $E\left(Y_{(n_i)}\right)$ is defined as $\int_{-\infty}^{\infty}Y_{(n_i)}q(Y_{(n_i)})dY_{(n_i)}$, where $q(Y_{(n)})$ is the pdf of the n_i th order statistic of $Y_{(n)}$. Since the parameters μ and σ in (5) will be cancelled (see the proof in Appendix I), without loss of generosity, we simply treat them as standard extreme value distribution. Therefore, $E(Y_{(n_i)})$ will be constants and set $g(z) = e^z e^{-e^z}$, $z = E(Y_{(n, \cdot)})$. It also showed that the weighted factors do not depend on the parameters μ and .

According to the general data scheme mentioned in section 1, the pivotal quantities of (5) to (8) are transformed to the following forms

$$\hat{U}_h = \frac{X_{(j)} - X_{(n-s)}}{\hat{W}_h}, \qquad h = 1, ..., 4, \qquad n - s < j \le n,$$

where

(6)

$$\hat{W}_{1} = \sum_{i=r+2}^{r+k} \frac{g(E(Y_{(i)}))}{S_{c}} (X_{(i)} - X_{(r+1)}) + \sum_{i=r+k+l+1}^{r+k+l+m} \frac{g(E(Y_{(i)}))}{S_{c}} (X_{(i)} - X_{(r+1)}),$$
(9)

$$\begin{split} \hat{W_2} &= \sum_{i=1}^{r} \frac{g(E(Y_{(i)}))}{S_n} (X_{(r+2)} - X_{(r+1)}) \\ &+ \sum_{i=r+2}^{r+k} \frac{g(E(Y_{(i)}))}{S_n} (X_{(i)} - X_{(r+1)}) \\ &+ \sum_{i=r+k+1}^{r+k+1} \frac{g(E(Y_{(i)}))}{S_n} \left(\frac{X_{(r+k)} + X_{(r+k+l+1)} - 2X_{(r+1)}}{2} \right) \\ &+ \sum_{i=r+k+l+1}^{n-s} \frac{g(E(Y_{(i)}))}{S_n} (X_{(i)} - X_{(r+1)}) \\ &+ \sum_{i=n-s+1}^{n} \frac{g(E(Y_{(i)}))}{S_n} (X_{(n-s)} - X_{(r+1)}), \end{split}$$
(10)

$$\hat{W}_{3} = \prod_{i=r+2}^{r+k} (X_{(i)} - X_{(r+1)})^{\frac{g(E(Y_{(i)}))}{S_{c}}} \times \prod_{i=r+k+l+1}^{r+k+l+m} (X_{(i)} - X_{(r+1)})^{\frac{g(E(Y_{(i)}))}{S_{c}}},$$
(11)

$$\begin{split} \hat{W_4} &= \prod_{i=1}^{r} \left(X_{(r+2)} - X_{(r+1)} \right)^{\frac{g(E(Y_{(i)}))}{S_n}} \\ &\times \prod_{i=r+2}^{r+k} \left(X_{(i)} - X_{(r+1)} \right)^{\frac{g(E(Y_{(i)}))}{S_n}} \\ &\times \prod_{i=r+k+1}^{r+k+l} \left(\frac{X_{(r+k)} + X_{(r+k+l+1)} - 2X_{(r+1)}}{2} \right)^{\frac{g(E(Y_{(i)}))}{S_n}} \\ &\times \prod_{i=r+k+l+1}^{n-s} \left(X_{(i)} - X_{(r+1)} \right)^{\frac{g(E(Y_{(i)}))}{S_n}} \\ &\times \prod_{i=n-s+1}^{n} \left(X_{(n-s)} - X_{(r+1)} \right)^{\frac{g(E(Y_{(i)}))}{S_n}}, \end{split}$$

where

$$S_{c} = \sum_{t=r+2}^{r+k} g(E(Y_{(t)})) + \sum_{t=r+k+l+1}^{r+k+l+m} g(E(Y_{(t)})),$$

$$S_{n} = \sum_{t=1}^{n} g(E(Y_{(t)})).$$

For comparison, the other pivotal quantity is \hat{U}_{a} . Let

$$\hat{U}_a = \frac{X_{(j)} - X_{(n-s)}}{\hat{W}_a}, \qquad n - s < j \le n, \qquad \text{where}$$

$$\hat{W}_a = \hat{\sigma}. \tag{13}$$

The $\hat{\sigma}$ is the AMLE of , which can be obtained from Balakrishnan and Cohen[3]. The percentiles data of \hat{U}_a distribution are listed in Table 9 (see Appendix II).

From equations (9) to (12), the distributions of \hat{U}_h depend only on n, r, k, l, m, s, j, but not on μ and . Then, we have

$$\begin{split} &1 - \alpha = P\{0 < \frac{X_{(j)} - X_{(n-s)}}{\hat{W}_h} < \hat{u}_h(1 - \alpha; n, r, k, l, m, s, j)\} \\ &= P\{X_{(n-s)} < X_{(j)} < X_{(n-s)} + \hat{u}_h(1 - \alpha; n, r, k, l, m, s, j) \times \hat{W}_h\}. \end{split}$$

Table 1. The properties of parameters $\widetilde{\mu}$ and $\widetilde{\sigma}$ of 60,000 random samples for each combination case.

Therefore, $(X_{(n-s)}, X_{(n-s)} + \hat{u}_h(1-\alpha; n, r, k, l, m, s, j) \times \hat{W}_h)$, $h = 1, \ldots, 4$ are one-sided 100(1- α)% prediction intervals of $X_{(j)}$ based on m+k observations.

3. CALCULATION AND ALGORITHM

The exact distributions of the pivotal quantities \hat{U}_h (h=1,..., 4) can not be derived algebraically, but we can approximate the distributions of \hat{U}_h (h=1,..., 4) by using large quantities of the Monte Carlo sampling with some programming algorithms to generate the percentiles of \hat{U}_h . All the simulations were run with the aid Microsoft Quick Basic 4.5 program and Foxbase database software package. The procedures for generating the percentiles of \hat{U}_h are as follows:

- a. Give and set μ =0, =1. (For providing the properties of parameters $\widetilde{\mu}$ and $\widetilde{\sigma}$ of the random samples generated by computer, 60,000 Monte Carlo runs are done for each combination of n, r, k, l, m, s, j (some selected cases). The results are presented in Table I. Using Table 5.3 in Mann *et al.*[17] for case of n=13 and Table 1 in Mann *et al.*[14] for remaining cases to obtain the necessary weights, we can calculate their BLIE's of μ and respectively. The $\widetilde{\mu}_{mean}$ and $\widetilde{\sigma}_{mean}$ of those random samples are very close to 0 and 1, respectively.)
- b. Calculate the following statistics: \hat{U}_1 in (9), \hat{U}_2 in (10), \hat{U}_3 in (11), \hat{U}_4 in (12).
- c. In the Step a and Step b, 600,000 replicates are used to compute the percentiles of \hat{U}_h (h=1,...,4) for each combination of n, r, k, l, m, s, j.
- d. Sort 600,000 results of each combination of *n*, *r*, *k*, *l*, *m*, *s*, *j* in ascending order.
- e. Retrieve the value of \hat{U}_h (h=1,...,4) under different significance levels of .

From the above procedures, we obtain the values of \hat{U}_h (h=1,...,4) according to the exact position of \hat{U}_h (h=1,...,4) in Step d.

In our simulation, 600,000 replicates are done for each combination of n, r, k, l, m, s, j. To save space, we only list part of the percentiles of \hat{U}_h (h=1,..., 4) in Table 5 to 8 (see Appendix II).

4. COMPARISON

In this section, we compare the performance of our method with \hat{U}_a . We calculate their average lengths of 95% prediction intervals, and coverage probabilities for some selected combinations of n, r, k, l, m, s, j. Referring to the data scheme mentioned in section 1, the simulation is computed by the following procedures:

- a. Give and set $\mu = 0$, =1.
- b. Generate *n* (*n*=10,13,20,40) random samples from the standard extreme value distribution.
- c. Calculate the values of \hat{W}_h (h=1,...,4,a), and then make a multiply of \hat{W}_h by \hat{U}_h (h=1,...,4,a) (from Table 5 to 9) for each combination of n, r, k, l, m, s, j.
- d. Repeat steps b to c, execute 10,000 runs and record all upper bounds of the confidence intervals of $X_{(n-s+1)}$ and $X_{(n-s+2)}$.
- e. From the results in steps c and d, calculate the average length of the 10,000 confidence intervals, and coverage probabilities for all methods

The results of simulation are listed in Table 2. It is clear that the 95% estimated expected lengths of \hat{U}_2 or \hat{U}_a are shorter. The difference among \hat{U}_1 , \hat{U}_2 , and \hat{U}_a is not significant (about 0% to 3%). It is also shown that the confidence intervals of \hat{U}_h (h=1,..., 4, a) have almost 95% coverage probabilities. It is interesting to note that if the sample size n is larger, then the difference of average lengths among \hat{U}_h (h=1,..., 4, a) will be smaller. And it also showed that simulation has the property of convergence.

Table 2. The average length of 95% prediction intervals , and coverage probabilities for $X_{(j)}$ by difference statistics: u=0 =1

5. EXAMPLES

5.1 Example 1

Consider the following 13 components were placed on test, and the test was terminated at the time of the 10th failure (Mann and Fertig[13]). The first 10 observations are given below:

0.22, 0.50, 0.88, 1.00, 1.32, 1.33, 1.54, 1.76, 2.50, 3.00. It is assumed that the 10 observed data are from the same Weibull distribution. We transform the data to extreme value form: the logs of the 10 observations are

-1.541, -0.693, -0.128, 0.000, 0.278, 0.285, 0.432, 0.565, 0.916, 1.099.

Table 3. The one-sided 90% prediction intervals, the percentiles of $\hat{U}_k(k=1,...,4,a)$, and CMd using MLE and BLIE (Hsieh[7]).

Note: CMd=Conditional Method, M=MLE

Table 4. The one-sided 95% prediction intervals and the percentiles of $\hat{U}_k(k=1,...,4,a)$

In this case, we have n = 13, r = 0, k = 10, l = 0, m = 0, and s = 3. Applying our method to estimate the one-sided 90% prediction intervals of $X_{(11)}$ and $X_{(12)}$. The results are presented in Table III. It is clear that the shorter prediction intervals are obtained by

the pivotal quantities \hat{U}_{2} and \hat{U}_{a} .

5.2 Example 2

The following are 10 observations data from Lawless[11]. -3.57, -2.55, -2.02, -1.66, -1.36, -1.15, -0.95, -0.77, -0.61, -0.45. It assumed that above data were obtained from a sample of 20, which are distributed according to extreme value distribution, and the last 50% data were censored. And only from the 3rd to the 6th failure times and from the 9th to the 10th failure times are available. In other words, this is the case of n = 20, r = 2, k = 4, k = 2, and k = 10, The one-sided 95% prediction intervals of k = 10, and k = 10, are listed in Table 4. It is obvious that the pivotal quantity k = 10, has the shortest prediction intervals.

6. DISCUSSION

From Table 2, the average length of prediction intervals of \hat{U}_1 , \hat{U}_2 , and \hat{U}_a are shorter than \hat{U}_3 and \hat{U}_4 . Since \hat{U}_3 and \hat{U}_4 are longer than \hat{U}_2 , \hat{U}_1 and \hat{U}_2 are preferred to both of them. The average lengths of prediction intervals of \hat{U}_3 and \hat{U}_4 are longer than \hat{U}_1 and \hat{U}_2 . It may be the reason that the power operations in geometric means will cause the results extended unexpectedly. Intuitively, our method produces good result because we have given different weight to each datum point. Following the algorithm of section 3, it is straightforward to construct prediction intervals for the future failure time by the pivotal quantities \hat{U}_1 and \hat{U}_2 . Note also that \hat{U}_1 and \hat{U}_2 can be applied to any kind of data scheme.

Comparing with the existing methods, it is true that calculation procedures of \hat{U}_a are simpler than \hat{U}_1 and \hat{U}_2 . But since the computations of \hat{U}_1 and \hat{U}_2 can be easily done by computers, it seems to be not an important consideration. Furthermore, following the some algorithm, it is not difficult for generating and simulating larger sample size n. Thus it makes this method to be potentially more useful than the existing ones. For further study, this simulation scheme can be easily applied to other family of location and scale distributions.

7. APPENDICES

7.1 Appendix I

Theorem: If \widetilde{W}_1 in (5) is an estimator of μ and based on multiple type II censored sample $X_{(n_0)} \leq X_{(n_1)} \leq \cdots \leq X_{(n)}$ from two-parameter Weibull distribution, then $\widetilde{U}_i = \frac{X_{(j)} - X_{(n_i)}}{\widetilde{W}_i}$ is a

pivotal (parameter-free) quantity

Proof

Define $Y_i = \frac{X_i - \mu}{\sigma}$, then Y_i does not depend on μ and . And $Y_{(i)} = \frac{X_{(i)} - \mu}{\sigma}$ is the *i*th order statistics of Y_i .

Let
$$Z_i = Y_{(j)} - Y_{(n_c)}$$
, $Z_2 = \sum_{i=2}^c \frac{g(E(Y_{(n_i)}))}{\sum_{i=2}^c g(E(Y_{(n_i)}))} (Y_{(n_i)} - Y_{(n_1)})$,

where

$$\sum_{i=2}^{c} \frac{g(E(Y_{(n_i)}))}{\sum_{i=2}^{c} g(E(Y_{(n_i)}))}$$
 are constant.

Define $Z_3 = Z_1/Z_2$. Both Z_1 and Z_2 are pivotal; therefore, Z_3 is also pivotal. It follows that

$$Z_{3} \text{ is also pivotal.} \quad \text{It follows that}$$

$$\frac{X_{(j)} - \mu}{\sigma} - \frac{X_{(n_{c})} - \mu}{\sigma}$$

$$\sum_{i=2}^{c} \frac{g(E(Y_{(n_{i})}))}{\sum_{i=2}^{c} g(E(Y_{(n_{i})}))} (\frac{X_{(n_{i})} - \mu}{\sigma} - \frac{X_{(n_{1})} - \mu}{\sigma})$$

$$= \frac{Y_{(j)} - Y_{(n_{c})}}{\sum_{i=2}^{c} \frac{g(E(Y_{(n_{i})}))}{\sum_{i=2}^{c} g(E(Y_{(n_{i})}))} (Y_{(n_{i})} - Y_{(n_{1})})} = \frac{Z_{1}}{Z_{2}} = Z_{3}.$$
Similarly, it is easy to show that the estimators \widetilde{X}

Similarly, it is easy to show that the estimators $\ \widetilde{U}_2$, $\ \widetilde{U}_3$, and $\ \widetilde{U}_4$ are also pivotal quantities.

7.2 Appendix II

Table 5. The percentiles data of \hat{U}_1 distribution for $X_{(i)}$

Table 6. The percentiles data of \hat{U}_2 distribution for $X_{(i)}$.

Table 7. The percentiles data of \hat{U}_3 distribution for $X_{(i)}$.

Table 8. The percentiles data of \hat{U}_4 distribution for $X_{(i)}$

Table 9. The percentiles data of \hat{U}_a distribution for $X_{(i)}$

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