

Applications of interval analysis to selected topics in statistical computing

by

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Chapter 1

GENERAL INTRODUCTION

1.1 Background

Since the beginning of mathematics, error has been a part of calculations, and reducing error has been an important component of research in many fields. Even with digital computers, error is still present. There are two common types of error in statistical calculations: error from rounding and algorithmic error.

Error from rounding occurs because the computer has only a finite set of numbers to approximate the whole real line. Even simple fractions are often impossible to represent exactly as a decimal on a computer. An additional complication arises from the requirement that calculations be performed in base 2 instead of base 10, further increasing the potential for error during base conversion of numbers.

Algorithmic error typically occurs when an infinite algorithm is terminated after a finite amount of time. For example, infinite series are frequently used to calculate functions such as \exp , \sin , \cos , etc. Truncating the infinite series to a finite number of terms will result in some error, regardless of how many digits of precision are used for the calculations.

Given that error exists, logical questions that follow are ones like “How large is the error” and “Is the error likely to be a problem?” Consider the system of linear equations $Ax = b$ where

$$A = \begin{pmatrix} 64919121 & -159018721 \\ 41869520.5 & -102558961 \end{pmatrix} \text{ and } b = \begin{pmatrix} 1 \\ 0 \end{pmatrix}. \quad (1.1)$$

The true solution to this set of equations is

$$\begin{pmatrix} x_1 \\ x_2 \end{pmatrix} = \begin{pmatrix} 205117922 \\ 83739041 \end{pmatrix}. \quad (1.2)$$

As reported by [Bohlender \(1990\)](#), even using IEEE double-precision arithmetic yields the following answer:

$$\begin{pmatrix} x_1 \\ x_2 \end{pmatrix} = \begin{pmatrix} 102558961 \\ 41869520.5 \end{pmatrix}, \quad (1.3)$$

which is completely wrong.

As a second example, [Hammer et al. \(1993a\)](#) consider an example with two real vectors,

$$\begin{aligned}x &= (10^{20}, 1223, 10^{18}, 10^{15}, 3, -10^{12}) \\y &= (10^{20}, 2, -10^{22}, 10^{13}, 2111, 10^{16}).\end{aligned}$$

The dot product of these two vectors is $x \cdot y = 10^{40} + 2446 - 10^{40} + 10^{28} + 6333 - 10^{28} = 8779$. On *all* standard computers, the result of this dot product is zero. In contrast, the use of interval arithmetic (introduced later) guarantees that the scalar product is somewhere in the interval $[0, 1.93429e + 25]$, possibly an acceptable answer, but likely to spur further investigation into the reasons behind the wide interval. It would be possible in this example to use a computer with more mantissa digits and obtain an accurate answer, but that assumes that one knows of the need to use more precision, and in any case, no matter how many digits of accuracy are maintained, it is always possible for actual computations to exceed this accuracy. Clearly, there is a need for additional understanding of error and ways to control the error. One of the tools which takes a step in the direction of these goals is interval analysis.

1.2 Introduction to Interval Analysis

A suitable introduction to interval analysis can be found in [Moore \(1979\)](#). The basic idea of interval analysis is to work with an interval not only as a set of numbers, but also as a number by itself. A real interval \mathbf{x} is defined $\mathbf{x} = [\underline{x}, \bar{x}]$ where \underline{x} and \bar{x} are real numbers with $\underline{x} \leq \bar{x}$. Let $\mathbf{x} = [\underline{x}, \bar{x}]$ and $\mathbf{y} = [\underline{y}, \bar{y}]$. Arithmetic operations for intervals are defined:

$$\mathbf{x} * \mathbf{y} = \{x * y : x \in \mathbf{x}, y \in \mathbf{y}\} \text{ for } * \in \{+, -, \cdot, \div\}.$$

Closed-form expressions of the basic interval arithmetic operators also exist and are the means by which rational expressions are actually computed. The closed-form expressions are:

$$\begin{aligned}\mathbf{x} + \mathbf{y} &= [\underline{x} + \underline{y}, \bar{x} + \bar{y}] \\ \mathbf{x} - \mathbf{y} &= [\underline{x} - \bar{y}, \bar{x} - \underline{y}] \\ \mathbf{x} \cdot \mathbf{y} &= [\min(\underline{x}\underline{y}, \underline{x}\bar{y}, \bar{x}\underline{y}, \bar{x}\bar{y}), \max(\underline{x}\underline{y}, \underline{x}\bar{y}, \bar{x}\underline{y}, \bar{x}\bar{y})] \\ 1/\mathbf{y} &= [1/\bar{y}, 1/\underline{y}], \quad 0 \notin \mathbf{y} \\ \mathbf{x}/\mathbf{y} &= \mathbf{x} \cdot (1/\mathbf{y}), \quad 0 \notin \mathbf{y}\end{aligned}$$

Examples of interval arithmetic:

$$\begin{aligned}[1, 1] + [-2, 5] &= [-1, 6] \\ [-2, 3] \cdot [1, 4] &= [-8, 12].\end{aligned}$$

From these basic definitions, the following properties are observed to hold for intervals:

$$\mathbf{x} + (\mathbf{y} + \mathbf{z}) = (\mathbf{x} + \mathbf{y}) + \mathbf{z}$$

$$\mathbf{x} \cdot (\mathbf{y} \cdot \mathbf{z}) = (\mathbf{x} \cdot \mathbf{y}) \cdot \mathbf{z}$$

$$\mathbf{x} + \mathbf{y} = \mathbf{y} + \mathbf{x}$$

$$\mathbf{x} \cdot \mathbf{y} = \mathbf{y} \cdot \mathbf{x}$$

for any intervals \mathbf{x} , \mathbf{y} , and \mathbf{z} . Interval addition and multiplication is therefore both associative and commutative for both addition and multiplication. One of the ways in which interval arithmetic differs from real arithmetic is that interval arithmetic is not distributive, but subdistributive only. As a consequence, interval subtraction and division are not the inverse of interval addition and multiplication. For example:

$$[0, 1] - [0, 1] = [-1, 1]$$

$$[1, 2]/[1, 2] = [1/2, 2].$$

For interval arithmetic, the subdistributive law states that for intervals \mathbf{x} , \mathbf{y} , and \mathbf{z} ,

$$\mathbf{x} \cdot (\mathbf{y} + \mathbf{z}) \subseteq \mathbf{x} \cdot \mathbf{y} + \mathbf{x} \cdot \mathbf{z}.$$

Further basic definitions and properties of intervals can be found in [Ratscheck and Rokne \(1984\)](#).

1.3 Interval Inclusion of Functions

Extending the ideas of real functions to interval functions is not straightforward. Several approaches are possible. An *interval function* is defined as an interval-valued function of one or more interval arguments.

A function $\mathbf{f}(\mathbf{x}_1, \dots, \mathbf{x}_n)$ is said to be an *interval extension* of $f(x_1, \dots, x_n)$ if

$$\mathbf{f}([x_1, x_1], \dots, [x_n, x_n]) = f(x_1, \dots, x_n)$$

for all $x_i, i = 1, \dots, n$.

Let $f(x_1, \dots, x_n)$ be a real-valued function of n real variables. For x_1, \dots, x_n defined over the intervals $\mathbf{x}_1, \dots, \mathbf{x}_n$ respectively, the *united extension* of f over $\mathbf{x}_1, \dots, \mathbf{x}_n$ is given by

$$\mathbf{f}(\mathbf{x}_1, \dots, \mathbf{x}_n) = \{f(x_1, \dots, x_n) : x_1 \in \mathbf{x}_1, \dots, x_n \in \mathbf{x}_n\}.$$

The united extension of a function is unique, however, it need not be an interval.

One kind of interval extension that is fairly logical is the so-called *natural interval extension* which is defined this way: let $f(x_1, \dots, x_n)$ be a rational function of n variables. Consider any sequence of arithmetic steps which serve to evaluate f with given arguments x_1, \dots, x_n . Suppose the arguments x_i are replaced by corresponding intervals $\mathbf{x}_i, (i = 1, \dots, n)$ and the arithmetic steps in the sequence used to evaluate f are replaced by the corresponding interval arithmetic steps. The result will be an interval $\mathbf{f}(\mathbf{x}_1, \dots, \mathbf{x}_n)$. This interval contains the value of $f(x_1, \dots, x_n)$ for all $x_i \in \mathbf{x}_i, (i = 1, \dots, n)$.

An interval-valued function \mathbf{f} is said to be *inclusion monotonic* if $\mathbf{f}(\mathbf{x}) \subseteq \mathbf{f}(\mathbf{y})$ whenever $\mathbf{x} \subseteq \mathbf{y}$. A fundamental theorem from interval analysis states that rational interval functions are inclusion monotonic.

It should be noted that an interval extension need not be unique, but can depend on the form of the real function. For example, while the following three expressions all represent the same real function,

$$\begin{aligned}f_1(x) &= x^2 - x + 1 \\f_2(x) &= (x - \frac{1}{2})^2 + \frac{3}{4} \\f_3(x) &= x \cdot (x - 1) + 1,\end{aligned}$$

the corresponding natural interval extensions

$$\begin{aligned}\mathbf{f}_1(\mathbf{x}) &= \mathbf{x}^2 - \mathbf{x} + 1 \\ \mathbf{f}_2(\mathbf{x}) &= (\mathbf{x} - \frac{1}{2})^2 + \frac{3}{4} \\ \mathbf{f}_3(\mathbf{x}) &= \mathbf{x} \cdot (\mathbf{x} - 1) + 1\end{aligned}$$

do *not* represent the same function. For illustration of how these functions differ, consider the following results:

$$\begin{aligned}\mathbf{f}_1([0, 2]) &= [-1, 5] \\ \mathbf{f}_2([0, 2]) &= \left[\frac{3}{4}, 3\right] \\ \mathbf{f}_3([0, 2]) &= [-1, 3].\end{aligned}$$

Thus, attention and care must be given to the choice of an interval extension in order to obtain the narrowest possible interval result. As shown in figure 1.1, the true range of f over $[0, 2]$ is $[\frac{3}{4}, 3]$, which is precisely that computed by the interval function \mathbf{f}_2 . As shown by Hansen (1997), this is because \mathbf{x} appears in the expression of \mathbf{f}_2 only once. In general, when a given interval argument appears only once in a function, the evaluation of the interval function produces a *sharp* interval that exactly matches the true range of the function over the given interval. When a given interval argument appears more than once in a function, the evaluation of the function may produce an interval wider than the sharp enclosure. This characteristic of interval analysis is referred to as the *dependency problem*. In some cases the dependency problem can be eliminated by simply changing a definition. For example, suppose $\mathbf{x} = [-1, 2]$. Then thinking of the square operation as simply multiplying two intervals together, $\mathbf{x}^2 = \mathbf{x} \cdot \mathbf{x} = [-1, 2] \cdot [-1, 2] = [-2, 4]$. This is not a sharp interval and contains negative real numbers—obviously not a desirable feature. It is possible to fix this problem with an appropriate definition of \mathbf{x}^2 as $\mathbf{x}^2 = \{x^2 : x \in \mathbf{x}\}$. Then, for example, $[-1, 2]^2 = [0, 4]$.

With the basic aspects of interval analysis thus defined, how should interval analysis be applied to particular problems? Walster (1988) presents a set of principles to guide the thinking of interval analysis:

1. Interval algorithms should bound all sources of error

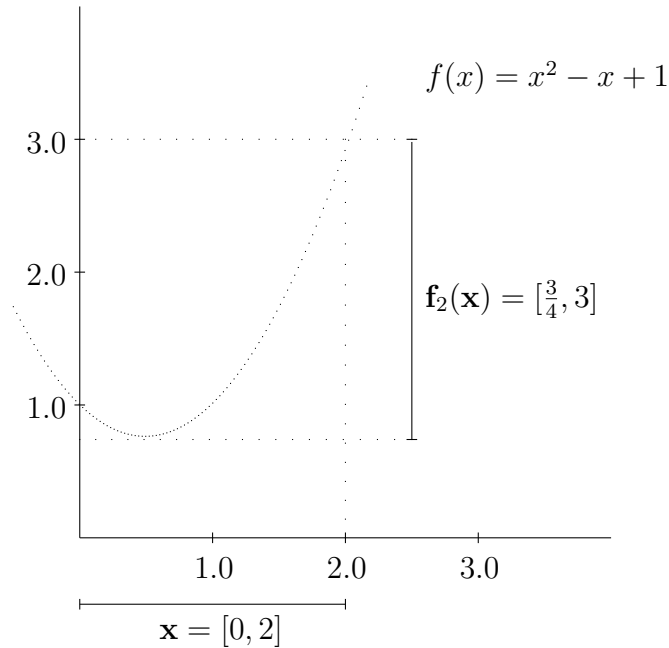


Figure 1.1: Example of an interval extension.

2. Interval input/output conventions should be consistent with people's normal interpretation of numerical accuracy
3. The application of interval algorithms should be universal
4. Where interval algorithms currently do not exist, we should get to work developing them rather than abandoning the principle of universal applicability.

1.4 Interval Analysis and Digital Computing

Efforts to produce accurate numerical results have been around nearly as long as number systems. The current widelyused approach for representation of numbers on computer systems dates back to the late 1970s with the establishment of a task force by the IEEE. From that work, and from the work of a second task force, emerged two standards [ANSI/IEEE \(1985, 1987\)](#). A brief overview and history of the hardware aspects of the standards can be found in [Cody \(1988\)](#). The widelyused Intel 80x87 series of chips conform to this standard as do the numerical processors in many workstations. The standards focus mainly on the representation of floating point numbers, but also include rules for handling NaNs, overflow, underflow, rounding, etc. As regards to rounding floating point numbers, the first standard [ANSI/IEEE \(1985\)](#) specifies that the floating point processor support rounding to zero, nearest, $+\infty$, and $-\infty$. Software support for allowing user access to the various rounding modes, while available, is generally not as easily available

as would be desired. As one simple example, Fortran 77 has no direct support for control of the rounding mode. Programmers wishing to utilize the special rounding modes need specialized software packages and/or detailed knowledge of a computer's inner workings. Examples of such packages include programming languages like PascalXSC [Hammer et al. \(1993b\)](#), ACRITHXSC [Walter \(1993\)](#), FortranSC [Walter \(1988\)](#); [Metzger \(1988\)](#), and CXSC [Lawo \(1993\)](#). A comparison of these environments (and others) can be found in [Kearfott \(1996a\)](#). These scientific computing languages typically have some combination of support for interval data types, dynamic vectors and arrays, dot product expressions, rounding control, a large set of standard mathematical functions, operator overloading and userdefined operators.

For example, consider a software package with functions RoundUp and RoundDown that are used to control the rounding mode. Imagine using these functions on a computer which has three digits in floatingpoint numbers. The interval enclosure of $1/3$ would be computed as

$$[\text{RoundDown}(1/3); \text{RoundUp}(1/3)] = [0.333; 0.334] = [0.33_3^4].$$

Use of the directed rounding rigorously guarantees that the resulting interval contains the true value. By using the monotonicity property of rational interval functions and by using other properties, more complicated functions can be constructed. The exponential function, for example, can be computed via a series expression,

$$\exp(x) = \sum_{n=0}^{\infty} \frac{x^n}{n!}$$

which converges for all real x . For $x \leq 0$, the inequality

$$\left| \exp(x) - \sum_{n=0}^N \frac{x^n}{n!} \right| \leq \frac{x^{N+1}}{(N+1)!}$$

can be used to calculate and enclosure of $\exp(x)$ by rearranging the expression as

$$\sum_{n=0}^N \frac{x^n}{n!} - \frac{x^{N+1}}{(N+1)!} \leq \exp(x) \leq \sum_{n=0}^N \frac{x^n}{n!} + \frac{x^{N+1}}{(N+1)!}$$

and using interval arithmetic to calculate the rational functions which bound $\exp(x)$ above and below, then taking the hull of these intervals.

An example from [Krämer \(1993\)](#) illustrates a completely different approach to interval enclosures of functions, in this case the natural logarithm. Beginning with $x_0 > 0$ and $y_0 > 0$, the sequences

$$\begin{aligned} x_{n+1} &:= \sqrt{x_n \frac{x_n + y_n}{2}} \\ y_{n+1} &:= \sqrt{x_n \frac{y_n + y_n}{2}} \end{aligned}$$

both converge to $\log(\frac{x_0}{y_0})$. At each step,

$$\lim_{n \rightarrow \infty} x_n = \lim_{n \rightarrow \infty} y_n \in [\min(x_n, y_n), \max(x_n, y_n)].$$

To compute $\log(x)$, set $x_0 = x$ and $y_0 = 1$. The sequences x_n and y_n can either be computed with scalars (and careful use of rounding modes) or with intervals, in which case $\log(x) \in \text{Hull}(\mathbf{x}_n; \mathbf{y}_n)$.

An extensive set of algorithms for computing standard functions and inverse standard functions for interval arguments can be found in [Brane \(1988\)](#) and [Krämer \(1988\)](#).

1.5 Automatic Differentiation

Automatic differentiation is a method for computing derivatives of functions without resorting to symbolic manipulation or numerical approximations, but instead using standard differentiation rules and propagation of numerical values. No explicit expression for the derivatives is required.

Automatic differentiation frequently arises in the context of interval analysis algorithms. Algorithms in interval analysis which use derivatives of functions need an accurate and efficient means by which to compute the derivative of the function. Numerical derivatives are generally insufficiently accurate for interval calculations. Symbolic derivatives can be difficult to calculate efficiently and may be difficult to include in an algorithm, either by programming symbolic manipulations from scratch, or by linking to an external program (e.g. Maple or Mathematica) that actually calculates the symbolic derivatives. [Corliss \(1988\)](#) makes the surprising claim that with automatic differentiation, “on most computers, it is less expensive to evaluate the first few derivatives of f than it is to evaluate f itself.”

For simplicity of expression, the techniques of automatic differentiation typically use Taylor coefficients instead of derivatives directly. For the Taylor coefficients of a real function f , the following notation is used:

$$(f)_k := \frac{1}{k!} f^{(k)}(x_0) := \frac{1}{k!} \frac{d^k f}{dx^k} \Big|_{x=x_0}, k = 0, 1, \dots$$

Using this notation, $f(x_0) = (f)_0$, $f'(x_0) = (f)_1$, and $f^{(k)}(x_0) = k!(f)_k$. The rules from calculus for differentiation of basic arithmetic functions lead to the following Taylor coefficients:

$$\begin{aligned} (f + g)_k &= (f)_k + (g)_k \\ (f - g)_k &= (f)_k - (g)_k \\ (f \cdot g)_k &= \sum_{j=0}^k (f)_j \cdot (g)_{k-j} \\ (f/g)_k &= \frac{1}{(g)_0} \left((f)_k - \sum_{j=0}^{k-1} (f/g)_j \cdot (g)_{k-j} \right) \end{aligned}$$

For the exponential function,

$$(w)_0 = \exp((f)_0)$$

$$(w)_k = \frac{1}{k} \sum_{j=0}^{k-1} (k-j) \cdot (w)_j \cdot (f)_{k-j}, k \geq 1$$

Similar rules exist for other functions (e.g. \sin , \cos).

1.6 Previous Applications of Interval Analysis to Statistics

Applications of interval analysis to statistical problems have previously appeared in the areas listed below.

1.6.1 Score Model

One early application of interval arithmetic to statistics is due to [Walster \(1988\)](#), where consideration is given to a score model $x_i = t + \epsilon_i$ where $\epsilon_i \sim (0, \sigma_\epsilon)$ with the observations being independent. The usual estimate of t is \bar{x} , which has the desirable properties of being an unbiased estimate of t and of having an increasing precision as the number of observations, n , increases, i.e. $Var(\bar{x}) = \sigma_\epsilon^2/n$. If it is possible to bound the errors ϵ_i , then for some finite δ , $-\delta < \epsilon_i < \delta$ for each i and it is possible to construct interval data as $x_i = [x_i - \delta, x_i + \delta]$. One interval estimate of t is to compute the mean of the interval data, $\bar{x} = [\bar{x} - \delta, \bar{x} + \delta]$. Unfortunately, this estimator has a fixed width, regardless of the amount of data collected. As an alternative, since the errors are bounded, it is the case that $t \in [x_i - \delta, x_i + \delta]$ for every i and thus

$$\mathbf{t} = \bigcap_{i=1}^n [x_i - \delta, x_i + \delta] = [\max(x_i) - \delta, \min(x_i) + \delta]$$

can also be used to estimate t . This estimator has at least the potential of decreasing in width as more data is collected.

1.6.2 Probability Distributions

From random number simulation to critical points of hypothesis tests, statistical distributions are an important part of statistical computing. [Wang and Kennedy \(1990\)](#) present a selfvalidated method for the calculation of bivariate Normal probabilities over rectangular regions using interval analysis. Their method served as the basis for comparing several different methods for computing probabilities of a bivariate Normal distribution when issues of speed and accuracy are of concern.

In dimensions higher than two, [Wang and Kennedy \(1992\)](#) use a Taylor series expansion of the multivariate Normal integral and automatic differentiation to calculate

interval enclosures of probabilities over rectangular regions. The method is used to conclude that the Taylor series approximation gives more accurate results than an algorithm by Schervish.

More recently, [Wang and Kennedy \(1994a\)](#) developed various intervalbased ways for obtaining selfvalidated probabilities and percentiles of several univariate distributions, including Normal, Incomplete Gamma, Incomplete Beta, and Noncentral Chi-Square. The work was expanded to include the Central and Noncentral F distributions in [Wang and Kennedy \(1995\)](#).

Using a slightly different approach, [Wang \(1994\)](#) used a MasPar (Massively Parallel) machine for selfvalidated probabilities from multivariate Normal and multivariate t distributions.

1.6.3 Least Squares

In the traditional Least Squares methodology, dependent variables y_1, y_2, \dots, y_n are assumed to be related to independent variables X_1, X_2, \dots, X_n by the relation

$$y = X\beta + e$$

where the errors are independent and distributed as $N(0, \sigma^2)$. When X has full rank, there is a unique b that minimizes the sum of the squares of the errors, $\|y - X\beta\|_2$. This unique solution,

$$b = (X'X)^{-1}X'y,$$

has many desirable properties, such as unbiasedness and being the maximum likelihood estimator of β .

If $\mathbf{X} = [\underline{X}, \overline{X}]$ and $y = [\underline{y}, \overline{y}]$, then an interval least squares solution provides an enclosure of

$$\mathbf{B} = \mathbf{X}'\mathbf{y} = \{X'y : X \in \mathbf{X}, y \in \mathbf{y}\}.$$

Interval least squares was researched by [Gay \(1988\)](#) with the conclusion (p. 203): “Interval leastsquares estimates can furnish rigorous and reasonably tight bounds on the effect of errors in the independent variables on forecasts and, as special cases, on parameter estimates.”

1.6.4 Optimization

Optimization techniques are an essential tool in statistical computing. Some of the areas which require optimization methods include:

1. Maximum likelihood
2. Optimal experimental design
3. Projection pursuit
4. Nonlinear least squares
5. Minimum volume ellipsoids

Traditional optimization methods used for these problems include Steepest Descent, NewtonRaphson, DUD (Doesn't Use Derivatives), and Simulated Annealing. The more widely used methods are discussed in the books by [Thisted \(1988\)](#) and [Kennedy and Gentle \(1980\)](#). Simulated Annealing has been applied to optimal design problems in a paper by [Bohachevsky et al. \(1986\)](#), while DUD is discussed in a nonlinear leastsquares context by [Ralston and Jennrich \(1978\)](#). Simulated Annealing has perhaps the best chance of the methods listed here for avoiding being trapped in a local optimum and has proven effective for solving some problems, but requires parameters that can be difficult to tune so that a global optimum is located with any degree of certainty.

Interval analysis can also be used for global optimization. [Hansen \(1988\)](#) and [Kearfott \(1996a\)](#) are two monographs on this topic. A short description of the basic idea of interval global optimization is now given in terms of maximization. (Minimization is very similar). Interval analysis can be used to enclose the range of a function over a given domain. A noninterval optimization method can be used to quickly locate a local optimum y^* , which has interval enclosure $\mathbf{f}(y^*)$. The maximum value of the function f then must be at least as large as the lower bound of the interval enclosure $\underline{\mathbf{f}}(y^*)$. Starting with an initial interval box \mathbf{y}_0 , the region is bisected repeatedly into ever smaller boxes. If a box \mathbf{y}_i satisfies $\mathbf{f}(\mathbf{y}_i) < \underline{\mathbf{f}}(y^*)$, then it is known that \mathbf{y}_i does not contain a global maximum and can be discarded.

[Wang \(1994\)](#) uses interval global optimization algorithms on a MasPar machine to solve several problems from nonlinear regression, optimal design, and maximum likelihood for a moving average model.

1.7 Example Applications of Interval Analysis to Statistics

This section describes several ways (that have not previously been published) in which interval analysis can be utilized in statistics. These are simple examples (and so are not presented in great detail) that serve as demonstrations.

1.7.1 Experimental Design

Table [1.1](#) from [Montgomery \(1991\)](#) (page 151) illustrates an experimental design with missing data in the response variable. In this case there is one missing value. A traditional approach to analysis of this data is to minimize the error sum of squares with respect to x_m . As stated by [Montgomery \(1991\)](#),

$$SS_E = \sum_{i=1}^a \sum_{j=1}^b y_{ij}^2 - \frac{1}{b} \sum_{i=1}^a \left(\sum_{j=1}^b y_{ij} \right)^2 - \frac{1}{1} \sum_{j=1}^b \left(\sum_{i=1}^1 y_{ij} \right)^2 + \frac{1}{ab} \left(\sum_{i=1}^a \sum_{j=1}^b y_{ij} \right)^2$$

or in the case of one missing value,

$$SS_E = x_m^2 - \frac{1}{b}(y'_{i.} + x_m)^2 - \frac{1}{a}(y'_{.j} + x_m)^2 + \frac{1}{ab}(y'_{..} + x_m)^2 + R$$

Table 1.1: Randomized complete block design with one missing value

Type of Tip	Block			
	1	2	3	4
1	-2	-1	1	5
2	-1	2	x_m	4
3	-3	1	0	2
4	2	1	5	7

where R includes terms not involving x_m . Solving this yields

$$x_m = \frac{ay'_{i.} + by'_{.j} - y'_{..}}{(a-1)(b-1)}$$

for the missing data. For the data presented here, the estimate of the missing data is $\hat{x}_m = 1.22$. The analysis of variance is performed with the estimated value and a reduction in the error degrees of freedom by one.

This is a useful approach for analysis of variance, but offers little help in the way of sensitivity analysis for the estimates of the parameters in the design. Designating the observed data by Y and the appropriate design matrix by X , the model can be written $Y = X\beta$. Using the value of 1.22 as an estimate of the missing data, $\hat{\beta} = (X'X)^{-1}X'Y$ gives

$$\hat{\beta} = \begin{bmatrix} 7.11125 \\ -5.5 \\ -5.25 \\ -2.695 \\ -3 \\ -3.195 \\ -4.25 \end{bmatrix}$$

as the best linear unbiased estimator.

Suppose the missing value x_m is replaced by an interval value \mathbf{x}_m . The choice of an estimate for \mathbf{x}_m can be made in several ways. The experimenter may possess knowledge about the response (dependent) variable which provides obvious bounds. If, for example, the response variable is a proportion, then $[0, 1]$ must be the widest possible estimate of \mathbf{x}_m . Depending on which response value is missing, it may be possible to use another approach. For illustration purposes, \mathbf{x}_m is here estimated by $[-3, 7]$, which encompasses the range of values observed in the response variable. Let Y_I be interval data equal to the original data Y with the exception that Y_I uses \mathbf{x}_m in place of the missing data x_m .

Setting $\hat{\beta}_I = (X'X)^{-1}X'Y_I$ yields

$$\hat{\beta}_I = \begin{bmatrix} [6.75, 7.375] \\ [-5.5, -5.5] \\ [-5.25, -5.25] \\ [-3.75, -1.25] \\ [-3, -3] \\ [-4.25, -1.75] \\ [-4.25, -4.25] \end{bmatrix}$$

as an interval estimate of β .

It is now possible to see immediately and at a glance which values and to what extent the values of $\hat{\beta}_I$ are influenced by the interval estimate of x_m .

1.7.2 Kolmogorov Smirnov Distribution

Tables for estimating the goodness of fit of empirical distributions were published by [Smirnov \(1948\)](#). The distribution used is

$$F(x) = 1 - 2 \sum_{i=1}^{\infty} (-1)^{i-1} e^{-i^2 x^2}. \quad (1.4)$$

Let \mathbf{F}_k denote the interval extension of (1.4) truncated after k terms. Since the series in (1.4) is alternating, $F(x) \in \text{Hull}(\mathbf{F}_k, \mathbf{F}_{k+1})$ for each k . The series is calculated until no change in $\text{Hull}(\mathbf{F}_k, \mathbf{F}_{k+1})$ is observed from one iteration of k to the next. Examples of the selfvalidated values for the distribution function appear in tables 1.2 and 1.3. The tables validate the results published in [Smirnov \(1948\)](#) with the exceptions of some values in the earlier published work that appear to be typographical and/or rounding errors.

1.7.3 A Bivariate F Distribution

Let $X \sim \chi^2(2n)$, $Y \sim \chi^2(2n)$, and $X_0 \sim \chi^2(2m)$ be independent random variables. The random variable $V = \frac{\min(X,Y)/n}{X_0/m}$ is known as the smaller of bivariate correlated F variables, or studentized minimum Chi-square variable. Some references to this distribution are given by [Hamdy et al. \(1988\)](#).

The density of V is given by

$$h(v) = 2 \sum_{i=0}^{n-1} \frac{\binom{n+i-1}{i} (n/m)^{i+n} v^{n+i-1}}{\beta(n+i, m) (1 + 2vn/m)^{n+m+i}} I(v > 0), \quad (1.5)$$

[Hamdy et al. \(1988\)](#) give an algorithm for finding c in $1 - \alpha = \int_c^\infty h(v) dv$. The heart of the algorithm involves computation of

$$q(h) = \sum_{i=0}^{n-1} \sum_{j=0}^{m-1} \binom{n+i-1}{i} \binom{n+m+i-1}{j} \left(\frac{1}{2}\right)^{n+i-1} h^j (1-h)^{n+m+i-1-j} \quad (1.6)$$

Table 1.2: Guaranteed bounds for distribution points of Smirnov's distribution

x	$F(x)$	x	$F(x)$
0.4	0.00280767322270 ₁₅ ²⁹	1.5	0.977782037383474 ₈ ⁹
0.5	0.03605475633512 ₄₄ ⁵⁶	1.6	0.988047956760803 ₄ ⁵
0.6	0.13571722094939 ₅₂ ⁶²	1.7	0.993822569365555 ₈ ⁹
0.7	0.28876480497031 ₀₂ ¹²	1.8	0.996932378652420 ₂ ³
0.8	0.45585758842580 ₁₆ ²⁴	1.9	0.998536395162812 ₆ ⁷
0.9	0.607269292059345 ₄ ⁹	2.0	0.999329074744220 ₃ ⁴
1.0	0.730000328322645 ₃ ⁷	2.1	0.99970450327953 ₀₆₉ ⁷
1.1	0.82228180739359 ₈₈ ⁹⁰	2.2	0.999874956992450 ₃ ⁴
1.2	0.887750333329275 ₀ ¹	2.3	0.999949161306967 ₅ ⁶
1.3	0.931907778155233 ₅ ⁷	2.4	0.999980140991388 ₂ ³
1.4	0.960318120461885 ₆ ⁷	2.5	0.999992546693655 ₈ ⁹

Table 1.3: Guaranteed bounds for critical points of Smirnov's distribution

x	$F(x)$
1.223847870217082 ₁ ⁵	0.90
1.3580986393225 ₅₀₀ ⁵⁰⁵	0.95
1.6276236115189 ₄₈₄ ⁵¹⁷	0.99

where $h = (1 + 2nc/m)^{-1}$ $0 \leq h \leq 1$. Since $q(h)$ is a rational function, its interval extension $\mathbf{q}(\mathbf{h})$ is immediately obtainable. For a given α , the determination of c proceeds via some suitable root-finding method. For the present work, derivative-free bracket-secant and bisection methods were used. Some percentile points of this distribution are tabulated below in table 1.4. This tabulation is a subset of the tables in Gupta and Sobel (1962). Table 1.4 verifies the results in Gupta and Sobel (1962) with the exception of a few values in the earlier work that appear to be rounding errors.

1.8 Internet Resources for Interval Analysis

The world wide web serves as an extremely valuable resource for researchers interested in interval computations. A pair of starting points for searching the WWW are given here.

Development of some interval software packages (including BIAS/PROFIL) is pro-

Table 1.4: Percentile points of V with degrees of freedom $m = n$

m	$1 - \alpha = 0.75$	$1 - \alpha = 0.90$
1	0.166666666666666 ₅ ⁹	0.055555555555555 ₄₇ ⁶⁹
2	0.31674887432700 ₆₉ ⁷⁶	0.16295234436871 ₇₆ ⁸¹
3	0.40404581771562 ₁₀ ¹⁹	0.241694741845574 ₃ ⁸
4	0.46279003617582 ₂₄ ³⁵	0.30016544276898 ₄₄ ⁵³
5	0.50596585743400 ₄₉ ⁶⁶	0.34570359411078 ₁₆ ²⁶
6	0.53953488035766 ₆₉ ⁸⁷	0.38253359804302 ₁₈ ²⁹
7	0.566666603181305 ₁₄ ³⁶	0.41318079377463 ₂₅ ⁴⁰
8	0.58922356764353 ₅₆ ⁶⁰	0.43924707520903 ₁₈ ³³
9	0.60838773909400 ₃₇ ⁷⁵	0.4618026569781 ₇₈₄ ⁸⁰⁷
10	0.6249483525127 ₃₉₈ ⁴³²	0.48159369356954 ₅₁ ⁷³
11	0.63945734273308 ₁₂ ⁴⁷	0.49915892547182 ₄₁ ⁶¹
12	0.6523140778575 ₀₇₈ ¹¹⁶	0.51489877713092 ₃₀ ⁵⁵
13	0.66381591094938 ₂₃ ⁶⁵	0.52911819748746 ₄₁ ⁷⁶
14	0.67418959284749 ₁₆ ⁶⁵	0.54205427500907 ₀₄ ³⁵
15	0.68361157764423 ₃₂ ⁸¹	0.5538946419155 ₂₉₆ ³²⁹
16	0.69222159319099 ₂₇ ⁷⁶	0.56479009787345 ₀₆ ⁴⁴
17	0.7001319730934 ₁₆₅ ²²⁷	0.57486348822727 ₃₃ ⁷⁵
18	0.70743423464199 ₃₁ ⁸⁹	0.58421608660013 ₂₉ ⁷⁴
19	0.7142038168150 ₂₈₅ ³⁵⁰	0.59293227318832 ₁₁ ⁵⁶
20	0.720503558738 ₆₉₄ ⁷⁰¹	0.6010830234646 ₀₇₇ ¹²⁸
21	0.7263862971195 ₆₃₉ ⁷⁰⁸	0.60872855022104 ₂₈ ⁷⁹
22	0.73189683545944 ₁₁ ⁷⁹	0.61592033239380 ₂₄ ⁸²
23	0.7370734575331 ₅₂₅ ⁶⁰³	0.6227026926850 ₆₇₃ ⁷³¹
24	0.74194910507571 ₀₄ ⁸³	0.62911403840910 ₃₈ ⁹⁰
25	0.7465523045448 ₈₂₉ ⁹¹³	0.6351878476747 ₁₈₅ ²⁴⁹

ceeding at Technische Universitat HamburgHarburg. See <http://www.ti3.tuharburg.de/indexEnglisch.html> for details.

The URL for the *Interval Computations* journal homepage is <http://cs.utep.edu/intervalcomp/main.html>. This site contains numerous links to bibliographies, software, homepages of interval computations centers and individuals, etc.

1.9 Dissertation Organization

The remainder of this dissertation is divided into several chapters, each of which applies methods of interval analysis to a separate problem in statistics. The following three chapters are written as manuscripts for submission to scientific journals. This arrangement means there may be some duplicity and some differences in notation between chapters, but also means that each chapter can be read independently of the others.

The chapter which follows this one addresses calculating critical points and tail probabilities for several bivariate Chi-square distributions. Since locating the critical point for a given tail probability of a distribution typically involves rootfinding, there is also development of an interval secant algorithm (with Illinois modification). Some tables of verified percentile points of the distributions are presented.

The third chapter takes a similar approach, but considers a class of bivariate F distributions. Series expansions are used in the calculations together with selfvalidated numerical quadrature rules.

The fourth chapter considers a different type of problem altogether. Several ways of using interval analysis together with the EM algorithm are considered. One of the ways in which these two have successfully been combined is to consider an enclosure of the gradient of the loglikelihood and eliminate portions of the parameter space where the gradient is not zero. The methodology for this technique is presented along with an algorithm for the procedure. The methodology is applied to several examples.

Conclusions are presented at the end of the dissertation.

Chapter 2

SELF-VALIDATED COMPUTATIONS FOR THE PROBABILITIES OF THE CENTRAL BIVARIATE CHI-SQUARE DISTRIBUTION

Abstract

Self-validated computations using interval arithmetic produce results with a guaranteed error bound. This article presents methods for self-validated computation of probabilities and percentile points of the bivariate chi-square distribution. For the computation of critical points (c_1, c_2) in $P(Y_1 \leq c_1, Y_2 \leq c_2) = 1 - \alpha$, the case $c_1 = c_2$ is considered. A combination of interval bracket-secant and bisection algorithms is developed for finding enclosures of the percentile points of the distribution.

2.1 Introduction

Several applications in statistical inference rely on the existence of a bivariate chi-square distribution. As mentioned in [Jensen and Howe \(1968\)](#), such areas include simultaneous inferences for variances, simultaneous tests in analysis of variance, simultaneous tests for goodness of fit, and the distribution of the larger of correlated Chi-square -variates. [Gunst and Webster \(1973\)](#) show how the bivariate chi-square distribution can be applied to the density function of a linear combination of independent chi-square random variables. [Jensen and Jones \(1969\)](#) further discuss simultaneous confidence intervals for variances while [Tuprah and Woodall \(1986\)](#) present a related application to bivariate dispersion quality control charts. For example, consider a manufacturing process which is characterized by two random variables, X and Y . Suppose σ_1 and σ_2 are the respective process standard deviations of the quality characteristics. It is desirable to detect shifts

in the process standard deviations away from specified target values σ_{1_0} and σ_{2_0} . If the two random variables are not independent, then the bivariate chi-square distribution can be used to construct bounds for determining when (σ_1, σ_2) is significantly off target.

Computations on digital computers should not be undertaken without giving some thought to error analysis. Examples of erroneous results obtained through naive computations appear often enough in scientific literature to cause concern. The techniques of interval analysis pioneered by [Moore \(1966, 1979\)](#) can be used to provide guaranteed error bounds for the results of mathematical computations. Guaranteed error bounds, provided they are sufficiently tight, can be used, to assess the accuracy of tabled values or to evaluate the quality of results produced by scalar algorithms. Interval analysis has been successfully employed in statistical areas, e.g. [Wang and Kennedy \(1994b\)](#), but has not yet had wide exposure to the statistical community.

The goals of this paper are:

1. To present truncation error bounds for infinite series related to some bivariate chi-square distributions;
2. To apply these bounds using interval analysis;
3. To develop and apply intervalized secant-bracket methods for root-finding to the location of critical points; and
4. To compare results obtained with previously published results and promote the utility of interval analysis.

2.2 Interval Analysis

An interval \mathbf{x} is defined to be a closed, bounded set of real numbers, $\mathbf{x} = [\underline{x}, \bar{x}]$. Throughout this paper, boldface is used to indicate intervals. To assist the reader in immediately grasping the accuracy of numerical values, a shortened form of interval notation which will sometimes be used to represent intervals is $2.3_{33}^{67} = [2.333, 2.367]$.

Let $\mathbf{x} = [\underline{x}, \bar{x}]$ and $\mathbf{y} = [\underline{y}, \bar{y}]$ be two intervals. The interval arithmetic operations are defined as

$$\mathbf{x} \circ \mathbf{y} = \{x \circ y : x \in \mathbf{x}, y \in \mathbf{y}\},$$

where $\circ \in \{+, -, \cdot, /\}$ and division is undefined for $0 \in \mathbf{y}$. Interval arithmetic operations can be expressed in closed form using real arithmetic operations. For example,

$$\mathbf{x} \cdot \mathbf{y} = [\min(\underline{x}\underline{y}, \underline{x}\bar{y}, \bar{x}\underline{y}, \bar{x}\bar{y}), \max(\underline{x}\underline{y}, \underline{x}\bar{y}, \bar{x}\underline{y}, \bar{x}\bar{y})].$$

Similar expressions exist for $+$, $-$, and $/$.

An *interval function* is an interval-valued function of one or more interval arguments. A function $\mathbf{f}(\mathbf{x}_1, \dots, \mathbf{x}_n)$ is said to be an *interval extension* of $f(x_1, \dots, x_n)$ if

$$\mathbf{f}([x_1, x_1], \dots, [x_n, x_n]) = f(x_1, \dots, x_n)$$

for all $x_i, i = 1, \dots, n$. An interval-valued function \mathbf{f} is said to be *inclusion monotonic* if $\mathbf{f}(\mathbf{x}) \subset \mathbf{f}(\mathbf{y})$ whenever $\mathbf{x} \subset \mathbf{y}$. A fundamental theorem from interval analysis states that rational interval functions are inclusion monotonic.

In this paper, the *natural interval extension* of a real function will be used. This is an interval-valued function in which intervals and interval operations are substituted for scalars and scalar operations. The value of the interval extension of a function is dependent on the form of the real function. For example, let $f_1(x) = xx - 2x$ and $f_2(x) = x(x - 2)$. Let \mathbf{f}_1 and \mathbf{f}_2 be the corresponding natural interval extensions and let $\mathbf{x} = [-1, 2]$. Then $\mathbf{f}_1(\mathbf{x}) = [-6, 6]$ and $\mathbf{f}_2(\mathbf{x}) = [-6, 3]$ which both contain $[-1, 3]$, the true range of f over \mathbf{x} . This feature of interval computations to sometimes overestimate the range of a function is referred to as *interval dependency*. Attention must be given to the exact expression of a function to reduce the effect of interval dependency.

When implementing interval arithmetic calculations on computers, care must be taken to ensure that rounding errors do not invalidate the inclusion monotonicity of interval results. One way to achieve this is through the use of directed rounding modes in the floating-point calculations. When calculating the lower endpoint of an interval result, the floating-point processor is set to round all results *down*. For calculation of the upper endpoint of an interval result, all calculations are rounded *up*. Using the symbols ∇ and Δ to denote downward and upward rounding respectively, the actual computer implementation of interval addition is $\mathbf{x} + \mathbf{y} = [\nabla(\underline{x} + \underline{y}), \Delta(\bar{x} + \bar{y})]$. Correct use of the rounding modes guarantees that the computed result contains the true answer.

2.3 Bivariate Chi-Square Distributions

Consider two random variables with variances σ_1^2 and σ_2^2 . Let s_1^2 and s_2^2 be estimates of σ_1^2 and σ_2^2 such that $\nu_i s_i^2 / \sigma_i^2$ follows a chi-square distribution with ν_i degrees of freedom. The joint distribution of $\nu_1 s_1^2 / \sigma_1^2$ and $\nu_2 s_2^2 / \sigma_2^2$ is referred to as a bivariate chi-square distribution. For example, suppose σ_1 and σ_2 are the process standard deviations for two characteristics that have a bivariate normal distribution. The sample variances s_1^2 and s_2^2 can be used along with a region determined by a bivariate chi-square distribution to simultaneously detect shifts in the process standard deviations away from specified target values.

Consideration will be given here to three cases of a bivariate chi-square distribution which are distinguished by degrees of freedom and the number of non-zero canonical correlations.

Case I. In the first case, let $\{(Z_{1i}, Z_{2i}), i = 1, \dots, m\}$ be independent random variables, $Z_{ij} \sim N(0, 1)$ with (canonical) correlation between Z_{1i} and Z_{2i} of ρ . Then $Y_i = \sum_{j=1}^m Z_{ij}^2, i = 1, 2$ are chi-square random variables, each with m degrees of freedom, and with m non-zero canonical correlations ρ . The joint density of Y_1 and Y_2 is given by

$$f(y_1, y_2) = (1 - \rho^2)^{m/2} \sum_{j=0}^{\infty} \frac{\Gamma(\frac{m}{2} + j) \rho^{2j}}{j! \Gamma(\frac{m}{2})} \times \frac{(y_1 y_2)^{(m/2)+j-1} \exp[-(y_1 + y_2)/2(1 - \rho^2)]}{[2^{(m/2)+j} \Gamma(\frac{m}{2} + j) (1 - \rho^2)^{(m/2)+j/2}]^2} \quad (2.1)$$

and the distribution is given by

$$P[Y_1 \leq d_1, Y_2 \leq d_2] = (1 - \rho^2)^{m/2} \times \sum_{j=0}^{\infty} \frac{\Gamma(\frac{m}{2} + j)}{j! \Gamma(\frac{m}{2})} \rho^{2j} \gamma\left(\frac{m}{2} + j, d_1^*\right) \gamma\left(\frac{m}{2} + j, d_2^*\right), \quad (2.2)$$

where $\gamma(\alpha, d)$ is the incomplete gamma function,

$$\gamma(\alpha, d) = \int_0^d \frac{1}{\Gamma(\alpha)} x^{\alpha-1} e^{-x} dx$$

and $d_j^* = d_j / (1 - \rho_{12}^2)$. When the infinite series in (2.2) is truncated after $t + 1$ terms, a bound on the truncation error R_t given by Krishnaiah (1980a) is

$$R_t \leq 1 - (1 - \rho^2)^{m/2} \sum_{j=0}^t \frac{\Gamma(\frac{m}{2} + j)}{j! \Gamma(\frac{m}{2})} \rho^{2j}. \quad (2.3)$$

For completeness, the derivation of this bound is given here. Let

$$k_j = (1 - \rho^2)^{m/2} \frac{\Gamma(\frac{m}{2} + j)}{j! \Gamma(\frac{m}{2})} \rho^{2j}.$$

Since $\gamma(\cdot, \cdot) \leq 1$ and the k_j are the density of a Negative Binomial distribution, the truncation error R_t satisfies

$$R_t = \sum_{j=t+1}^{\infty} k_j \gamma\left(\frac{m}{2} + j, d_1^*\right) \gamma\left(\frac{m}{2} + j, d_2^*\right) \leq \sum_{j=t+1}^{\infty} k_j = 1 - \sum_{j=0}^t k_j.$$

Let P_t represent the result when the first $t + 1$ terms of (2.2) are used, and let \mathbf{p}_t and \mathbf{r}_t represent the natural interval extensions of P_t and R_t respectively. Then

$$P[Y_1 \leq d_1, Y_2 \leq d_2] = P_t + R_t \in [p_t, \bar{p}_t + \bar{r}_t]$$

for all t . The stopping value of t used depends on machine and software precision. In practice, \mathbf{r}_t is computed successively and iteration stops when $\mathbf{r}_{t-1} = \mathbf{r}_t$ or when the width of \mathbf{r}_t is less than a specified tolerance.

Case II. Now consider the case when Y_1 and Y_2 follow chi-square distributions with m and n degrees of freedom respectively and have m non-zero canonical correlations ρ . The joint distribution of Y_1 and Y_2 given by Gunst and Webster (1973) is

$$P[Y_1 \leq d_1, Y_2 \leq d_2] = (1 - \rho^2)^{(m+n)/2} \times \sum_{j=0}^{\infty} \sum_{k=0}^{\infty} \frac{\Gamma(\frac{m}{2} + j)}{j! \Gamma(\frac{m}{2})} \frac{\Gamma(\frac{n}{2} + k)}{k! \Gamma(\frac{n}{2})} \rho^{2(j+k)} \gamma\left(\frac{m}{2} + j, d_1^*\right) \gamma\left(\frac{n}{2} + k, d_2^*\right). \quad (2.4)$$

Since a bound for the truncation error has not previously been published, one is given here. Because the method of derivation is completely analogous to the previous case, only the result is stated:

$$R_{t_1, t_2} \leq 1 - (1 - \rho^2)^{m/2+n/2} \sum_{j=0}^{t_1} \sum_{k=0}^{t_2} \frac{\Gamma(\frac{m}{2} + j)}{j! \Gamma(\frac{m}{2})} \frac{\Gamma(\frac{n}{2} + k)}{k! \Gamma(\frac{n}{2})} \rho^{2(j+k)}. \quad (2.5)$$

Let \mathbf{p}_{t_1, t_2} be the result obtained when the first $(t_1 + 1, t_2 + 1)$ terms of the natural interval extension of (2.4) are used, and let \mathbf{r}_{t_1, t_2} be the result obtained from the natural interval extension of (2.5). Then

$$P[Y_1 \leq d_1, Y_2 \leq d_2] \in [\underline{p}_{t_1, t_2}, \bar{p}_{t_1, t_2} + \bar{r}_{t_1, t_2}]$$

for all pairs (t_1, t_2) .

Case III. In the final case, Y_1 and Y_2 have chi-square distributions with $m + n$ and $m + p$ degrees of freedom, respectively, and there are m non-zero canonical correlations. The joint distribution given by [Gunst and Webster \(1973\)](#) is:

$$P[Y_1 \leq d_1, Y_2 \leq d_2] = (1 - \rho^2)^{(m+n+p)/2} \sum_{j=0}^{\infty} \sum_{k=0}^{\infty} \sum_{l=0}^{\infty} \frac{\Gamma(\frac{m}{2} + j)}{j! \Gamma(\frac{m}{2})} \times \frac{\Gamma(\frac{n}{2} + k)}{k! \Gamma(\frac{n}{2})} \frac{\Gamma(\frac{p}{2} + l)}{l! \Gamma(\frac{p}{2})} \rho^{2(j+k+l)} \gamma\left(\frac{m}{2} + \frac{n}{2} + k + j, d_1^*\right) \gamma\left(\frac{m}{2} + \frac{p}{2} + j + l, d_2^*\right). \quad (2.6)$$

The bound on the truncation error is again derived in a manner completely analogous to the first case:

$$R_t \leq 1 - (1 - \rho^2)^{m/2+n/2+p/2} \times \sum_{j=0}^{t_1} \sum_{k=0}^{t_2} \sum_{l=0}^{t_3} \frac{\Gamma(\frac{m}{2} + j)}{j! \Gamma(\frac{m}{2})} \frac{\Gamma(\frac{n}{2} + k)}{k! \Gamma(\frac{n}{2})} \frac{\Gamma(\frac{p}{2} + l)}{l! \Gamma(\frac{p}{2})} \rho^{2(j+k+l)}. \quad (2.7)$$

Similar to before,

$$P[Y_1 \leq d_1, Y_2 \leq d_2] \in [\underline{p}_{t_1, t_2, t_3}, \bar{p}_{t_1, t_2, t_3} + \bar{r}_{t_1, t_2, t_3}]$$

for all triples (t_1, t_2, t_3) and the choice of (t_1, t_2, t_3) is determined by machine/software limitations or a tolerance level.

2.4 Interval Bracket-Secant and Bisection Root-Finding

For a distribution function $F(x)$, the $100p^{th}$ percentile x_p is the solution of the equation $F(x_p) - p = 0$. For the bivariate chi-square distributions considered above, finding percentiles will involve solving

$$P(Y_1 \leq x_p, Y_2 \leq x_p) - p = 0,$$

which motivates a short discussion of interval root-finding techniques. Intervalized Newton methods for finding zeros of functions exist and could be used. However, interval Newton methods require an interval extension of both the function of interest and its derivative, which in this case are the distribution and density functions. Using a derivative-free search algorithm eliminates the need to obtain an enclosure of the density function. The technique of automatic differentiation presented in [Moore \(1979\)](#) could be used to obtain an enclosure of the derivative of a function, but only at the cost of

(sometimes considerable) loss of precision due to dependency and additional computing time.

The algorithm ZERO used here begins with an intervalized secant-bracket method using the Illinois modification. See [Thisted \(1988\)](#) for a complete explanation of this algorithm in the scalar case. Consider the general real equation $F(x) = 0$ as depicted by the curved line in figure 2.1. Let $\mathbf{F}(x) \equiv \mathbf{F}([x, x])$ (i.e. a scalar argument x is interpreted by the function as the interval $[x, x]$) be the interval extension of F . The algorithm begins with the user-specified interval $[x_{i-1}, x_i]$, which bounds the zero of the function. Using an intervalized secant method, the algorithm finds the point x_{i+1} and decides whether $[x_{i-1}, x_{i+1}]$ or $[x_{i+1}, x_i]$ now contains the zero of F . At some iteration $i + 1$ of the secant portion of the algorithm, $0 \in \mathbf{F}(x_{i+1})$ and it is not then known whether the zero of the function is to the left or the right of x_{i+1} . Figure 2.1 shows this condition.

The secant algorithm is successful in narrowing the enclosure of the zero of the function for the initial iterations, but may stop while the interval is wider than desired.

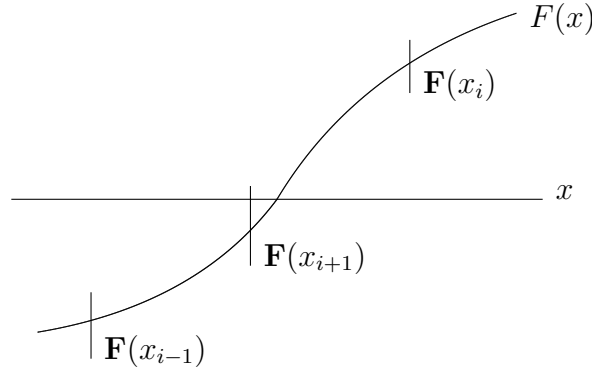


Figure 2.1: Termination of the bracket-secant portion of the algorithm. The curved line represents the real function and vertical line segments denote interval enclosures of the function.

After the bracket-secant portion of the algorithm terminates, let x_L and x_U represent the last two iterates x_{i-1} and x_i . A bisection algorithm is called twice, once each on the lower $[x_L, x_{i+1}]$ and upper $[x_{i+1}, x_U]$ intervals, to tighten the enclosure of the zero as much as possible, i.e. until (but not including) $0 \in \mathbf{F}(x_L), 0 \in \mathbf{F}(x_U)$. Figure 2.2 illustrates conditions at the termination of algorithm ZERO.

ALGORITHM Zero(MACHINE_REAL x_0, x_1 ; INTERVAL_FUNCTION \mathbf{F})

REM Bound the root of an increasing function F . Return $[l, u]$ to user.

REM Check that $F(x_0) < 0 < F(x_1)$ or $F(x_1) < 0 < F(x_0)$

REM Bold letters denote intervals

$l := x_0$

$u := x_1$

$\mathbf{FL} := \mathbf{F}([l, l])$

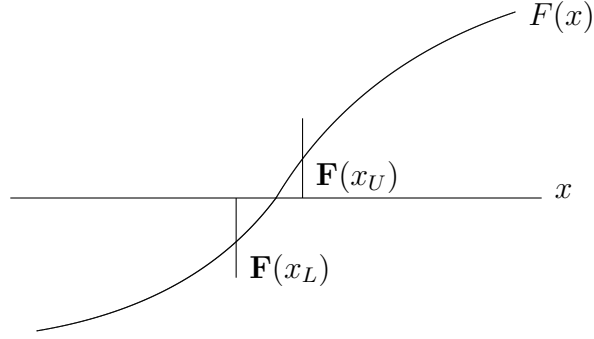


Figure 2.2: Termination of the bisection portion of the algorithm. The interval $[x_L, x_R]$ bounds the root of the function $F(x) = 0$.

```

FU := F([u, u])
REPEAT
  Xc :=  $u - (u - l)/(1 - \mathbf{FL}/\mathbf{FU})$     REM Find the secant intercept
   $x_{i+1} := (\mathbf{X}_c + \overline{\mathbf{X}}_c)/2$     REM Use the midpoint for the next iterate
  Fi+1 := F([xi+1, xi+1])
  IF Fi+1 > 0 THEN
    REM The zero is between l and xi+1
    u := xi+1
    FU := Fi+1
    IF Fi > 0 THEN FL := FL/2    REM Illinois modification
  ELSE IF Fi+1 < 0 THEN
    REM The zero is between xi+1 and u
    l := xi+1
    FL := Fi+1
    IF Fi < 0 THEN FU := FU/2    REM Illinois modification
  ELSE
    REM Cannot determine if the zero is to the left or right of xi+1
    Done := TRUE
  IF 0 ∈ (FU - FL) THEN
    REM In the next iteration, the denominator would contain zero
    Done := TRUE
  Fi := Fi+1
UNTIL (Done = TRUE)
REM Use bisection to tighten the upper endpoint
q := l
REPEAT
  prevq := q
  prevu := u

```

```

 $x := (q + u)/2$ 
 $\mathbf{FX} := \mathbf{F}([x, x])$ 
IF  $\mathbf{FX} > 0$  THEN  $u := x$ 
ELSE IF  $\overline{\mathbf{FX}} < 0$  THEN  $q := l := x$ 
ELSE  $q := x$ 
UNTIL (  $(q = \text{prev}q)$  AND  $(u = \text{prev}u)$  )
REM Use bisection to tighten the lower endpoint
 $q := u$ 
REPEAT
     $\text{prev}q := q$ 
     $\text{prev}l := l$ 
     $x := (l + q)/2$ 
     $\mathbf{FX} := \mathbf{F}([x, x])$ 
    IF  $\mathbf{FX} > 0$  THEN  $u := x$ 
    ELSE IF  $\overline{\mathbf{FX}} < 0$  THEN  $l := x$ 
    ELSE  $q := x$ 
UNTIL (  $(q = \text{prev}q)$  AND  $(l = \text{prev}l)$  )
RETURN  $[l, u]$ 

```

2.5 Numerical Results

Computer languages supporting overloaded operators are ideally suited to the implementation of interval arithmetic routines. For this reason, calculations in this paper made use of the BIAS/PROFIL (Basic Interval Arithmetic Subroutines/ Programmer's Optimized Fast Interval Library) C++ package developed by Knüppel (1993a,b) and were programmed on DEC 5000 and DEC Alpha workstations. Computing times for a single critical point varied from a few seconds in Case I to a few minutes in Case III. Routines to compute an enclosure of the Incomplete Gamma function were drawn from work by Wang and Kennedy (1994b) and from source code by Gessner (1992).

Since the expressions (2.2), (2.4), and (2.6) depend on ρ only through ρ^2 , tables need only include nonnegative values of ρ . For the first case, tables 2.1 and 2.2 illustrate critical points c for $P(Y_1 \leq c, Y_2 \leq c) = 1 - \alpha$ where $Y_1 \sim \chi^2(m)$, $Y_2 \sim \chi^2(m)$.

For the second case, $Y_1 \sim \chi^2(m)$, $Y_2 \sim \chi^2(m + n)$, $n > 0$ and there are m nonzero canonical correlations. Examples of critical values c for $P(Y_1 \leq c, Y_2 \leq c) = 1 - \alpha$ are given in table 2.3.

In the third case, $Y_1 \sim \chi^2(m + n)$, $Y_2 \sim \chi^2(m + p)$, $n > 0, p > 0$ and there are m nonzero canonical correlations. Table 2.4 illustrates critical values for this case.

The illustrative tables presented here are limited to $c_1 = c_2$ when computing the values of (c_1, c_2) in $P(Y_1 \leq c_1, Y_2 \leq c_2) = 1 - \alpha$. Other schemes are possible, fixing c_1 for example and calculating c_2 , or by adding a constraint. Distinct values of c_1 and c_2 can be entered when $1 - \alpha$ is the quantity to be computed.

Tables for the approximate critical points of the bivariate chi-square distribution have previously appeared in Gunst (1973), Gunst and Webster (1973), and Krishnaiah (1980a).

Table 2.1: Upper 0.05 percentile points of the bivariate chi-square distribution: Case I

ρ	m = 2	m = 12	m = 40
0.1	7.348735242636 ₆₂ ⁹⁴	23.291675614644 ₃ ⁹	59.27375898559 ₈₃ ⁹³
0.2	7.337736654468 ₅₂ ⁷³	23.279893907495 ₀ ⁵	59.25865809054 ₁₈ ³³
0.3	7.318116097295 ₀₀ ³³	23.257752618706 ₃ ⁸	59.2298092064 ₂₉₉ ³¹⁵
0.4	7.28777721964 ₁₉₄ ²³¹	23.22124105410 ₁₃ ²⁰	59.1811443051 ₂₉₀ ³¹⁶
0.5	7.243389878426 ₀₃ ³⁵	23.16405309924 ₇₂ ⁸⁰	59.10283493307 ₁₁ ³⁵
0.6	7.179739084402 ₀₀ ⁴⁷	23.07634122520 ₀₆ ¹⁹	58.9791571922 ₇₈₂ ⁸²⁶
0.7	7.088168635581 ₂₁ ⁷⁴	22.94173918779 ₀₉ ²⁶	58.78357869885 ₀₄ ⁶⁵
0.8	6.95217862545 ₄₆₂ ⁵⁶³	22.72905146810 ₀₀ ³⁰	58.4651869030 ₇₁₀ ⁸³⁰
0.9	6.73002568707 ₄₉₂ ⁶⁹⁹	22.3595746799 ₁₉₅ ²⁶³	57.8954062111 ₀₄₂ ²⁵⁹

Jensen and Howe (1968) determine the probability content over certain square and rectangular regions for which the marginal probabilities are specified. Dutt and Soms (1976) describe an alternative method for calculating multivariate chi-square probabilities using integral representations. The very narrow interval enclosures obtained using interval analysis can be useful for checking the accuracy of previously tabulated values. Indeed, Table I of Gunst (1973), *Upper 100 α % Critical Points*, is discovered to be widely accurate to only two decimal places (three are given), and the values in Table II, *Lower 5% Critical Points*, appear to be wholly incorrect. As noted in the text which accompanies the two tables of Gunst (1973), the critical points for $\rho = 0.10$ are nearly identical to the univariate critical points for $\alpha^* = 1 - (1 - \alpha)^{(1/2)}$. That this observation holds for values obtained in the current research and does not hold for the previously published values supports the correctness of the current research. The values in Table 6 of Krishnaiah (1980a), *Percentage Points of the bivariate chi-square distribution*, should be multiplied by two to obtain the correct values, and are then accurate to only two decimal places. If, for example, the incorrect values had been used to construct a confidence ellipsoid for the distribution of the variances of two random variables, the ellipsoid would be far too small to achieve the desired confidence.

The tables which appear in this paper are included to demonstrate the very high precision and the guarantee of accuracy which are obtained via the use of interval arithmetic. Except in certain cases, use of more digits in the table than significant digits in available data should be discouraged. The great value of such high-quality numbers is more likely in knowing that the second digit of a percentile point is guaranteed to be accurate than in knowing what the eighth digit is.

Table 2.2: Upper 0.01 percentile points of the bivariate chi-square distribution: Case I

ρ	m = 2	m = 12	m = 40
0.1	10.5901634351 ⁸⁰¹ ₇₈₈	28.29092305136 ³⁹ ₁₆	66.75375793961 ⁶⁸ ₃₁
0.2	10.58532786052 ⁶³ ₄₂	28.28686089292 ²⁴ ₀₀	66.74911374475 ⁸¹ ₂₁
0.3	10.5756426989 ⁴⁹⁷ ₄₈₁	28.27821885663 ²⁸ ₀₄	66.739131842 ³⁰³⁹ ₂₉₇₀
0.4	10.558524765 ²⁰¹¹ ₁₉₉₃	28.26172939228 ⁵⁵ ₂₁	66.719658957 ¹¹⁰⁶ ₀₉₆₆
0.5	10.52993690134 ⁸⁶ ₇₀	28.2317690519 ⁵²⁶ ₄₉₀	66.6830938986 ³⁸⁵ ₂₈₂
0.6	10.48358637908 ⁹¹ ₆₈	28.17882820433 ⁸¹ ₂₁	66.6158600156 ²⁵⁷ ₀₆₇
0.7	10.40898564766 ⁶⁴ ₃₈	28.0861355790 ⁷²⁴ ₆₅₅	66.4929962883 ⁸⁷⁷ ₆₁₉
0.8	10.28602680729 ⁸⁰ ₃₀	27.9204710072 ⁵⁴⁹ ₄₁₈	66.263713351 ⁴⁴⁵¹ ₃₉₄₅
0.9	10.063591724 ²⁰⁷⁷ ₁₉₇₄	27.5960985268 ⁶⁷⁴ ₃₇₈	65.795086198 ⁹⁴³⁹ ₈₅₃₄

2.6 Conclusions

Even in algorithms where theoretical error analysis bounds error terms, computer arithmetic rounding errors and cancellation can have catastrophic effects. In this paper, interval analysis techniques have been successfully applied to bivariate chi-square distributions to produce tables of critical values with guaranteed error bounds. The results obtained revealed inaccuracies and limitations of earlier published tables. Improved tables are useful in and of themselves, but it is expected that this research will be of more value in the long run by promoting the use of automatically verified computations and by providing new methodologies to achieve that end. For example, the techniques developed here have been modified slightly by the authors to produce tables of self-validated critical points of a bivariate F distribution. Interval analysis does have limitations in applications, but further research in applying interval techniques to statistical sciences should produce fruitful results.

Persons interested in obtaining a copy of the software described here may contact either of the authors.

Table 2.3: Upper 0.05 percentile points of the bivariate chi-square distribution: Case II

m	$m + n$	$\rho = 0.2$	$\rho = 0.4$
8	10	19.25562949840 ⁸³ ₅₇	19.2145575272 ¹¹⁸ ₀₅₁
8	12	21.43987191019 ³¹ ₀₁	21.4161617406 ³¹⁴ ₂₄₃
8	14	23.8527709314 ¹²⁵ ₀₉₃	23.8412410690 ¹⁵⁵ ₀₈₅
8	16	26.3613220302 ⁷⁰⁰ ₆₆₅	26.356264614 ⁸⁰²⁹ ₇₉₄₁
8	18	28.89382780550 ⁹⁴ ₅₇	28.8917353286 ⁷⁶⁹ ₆₅₅
m	$m + n$	$\rho = 0.6$	$\rho = 0.8$
8	10	19.1173225955 ⁴⁶⁵ ₂₆₈	18.90189596 ³⁰³⁴³ ₂₉₅₀₂
8	12	21.3604107620 ⁶²⁶ ₄₅₉	21.243609643 ⁹¹⁶⁶ ₈₃₈₆
8	14	23.8140331030 ⁸⁸⁷ ₆₅₃	23.759702671 ⁶⁴⁶² ₅₂₃₃
8	16	26.3442149104 ⁷³⁷ ₄₁₇	26.321129600 ³⁴⁰³ ₂₃₃₄
8	18	28.8866926661 ⁷³³ ₄₇₈	28.877401302 ⁶²⁴⁵ ₄₆₄₁

Table 2.4: Upper 0.05 percentile points of the bivariate chi-square distribution: Case III

m	n	p	$\rho = 0.4$	$\rho = 0.6$
7	1	11	28.892120984 ⁷²³⁶ ₆₄₈₆	28.887920316 ⁷⁴⁶¹ ₃₄₅₀
6	2	12	28.8924903043 ⁹²⁹ ₁₈₉	28.88907530 ⁴²⁹¹⁴ ₃₉₀₀₅
5	3	13	28.892843220 ⁴²³⁵ ₃₅₀₅	28.890155051 ⁴⁶⁴¹ ₁₁₀₅
4	4	14	28.893179676 ⁸⁵⁶³ ₇₈₅₀	28.891157355 ⁶¹⁹⁶ ₁₂₇₉
3	5	15	28.893499628 ⁹¹⁶² ₈₄₉₇	28.89208036 ⁹³²⁶⁹ ₈₈₆₃₂
2	6	16	28.893803042 ⁸⁷⁶⁹ ₇₉₅₉	28.892922583 ³⁸⁸⁶ ₀₀₂₃
1	7	17	28.894089895 ⁹⁰⁴⁷ ₈₀₅₄	28.89368281 ¹⁴⁰³⁸ ₀₉₉₂₆

Chapter 3

SELF-VALIDATED CRITICAL POINTS OF A BIVARIATE F DISTRIBUTION

Abstract

Self-validated computations using interval arithmetic produce results with a guaranteed error bound. This article presents methods for self-validated computation of probabilities and percentile points of a bivariate F distribution. For the computation of critical points (c_1, c_2) in $P(Y_1 \leq c_1, Y_2 \leq c_2) = 1 - \alpha$, the case $c_1 = c_2$ is considered. A combination of interval bracket-secant and bisection algorithms is utilized for finding enclosures of the percentile points of the distribution.

3.1 Introduction

A multivariate F distribution, though not common, has useful applications in statistics. [Schuurmann et al. \(1975\)](#) point out its use in hypothesis testing under fixed-effects models, in certain two-way classification models, and in simultaneous testing of no treatment and block effects in symmetrical balanced incomplete block designs. Tables for percentage points of a multivariate F distribution have appeared most recently in [Krishnaiah \(1980b\)](#).

Frequently implicit in the calculation of tables for critical points of statistical distributions is the assumption that numerical error does not invalidate the results. By using the techniques of interval analysis, it is possible to produce tables which are guaranteed to be free of rounding and certain other kinds of numerical error. This paper seeks to develop the techniques of interval analysis to a statistical application and promote the further utility of interval analysis to the statistical community.

3.2 Interval Analysis

Interval analysis first saw fruitful development in the 1960s, beginning with work published by [Moore \(1966\)](#). Since that time, interval analysis has seen extensive research a variety of applications, but has not seen wide exposure in statistical areas. This paper continues the development of interval analysis applications to statistical distributions as in [Wang and Kennedy \(1994b\)](#) and [Wright and Kennedy \(1999\)](#).

An interval \mathbf{x} is a closed, real interval $[\underline{x}, \overline{x}]$, where the underscore and overscore are used to denote the lower and upper endpoints of the interval. In this paper, boldface is used to distinguish intervals from real numbers. Fundamental arithmetic operators can be defined for intervals. Let \mathbf{x} and \mathbf{y} be intervals. For $*$ $\in \{+, -, \times, /\}$, define $\mathbf{x} * \mathbf{y} = \{x * y : x \in \mathbf{x}, y \in \mathbf{y}\}$, with division defined only when $0 \notin \mathbf{y}$. Closed-form expressions for the results of these operators exist, providing for easy computation of results, e.g.

$$\mathbf{x} \times \mathbf{y} = [\min\{\underline{x}\underline{y}, \underline{x}\overline{y}, \overline{x}\underline{y}, \overline{x}\overline{y}\}, \max\{\underline{x}\underline{y}, \underline{x}\overline{y}, \overline{x}\underline{y}, \overline{x}\overline{y}\}].$$

An *interval function* is understood to be a function with interval arguments and an interval result. An interval function \mathbf{f} is said to be *inclusion monotonic* if $\mathbf{x} \subset \mathbf{y}$ implies $\mathbf{f}(\mathbf{x}) \subset \mathbf{f}(\mathbf{y})$. A fundamental theorem from interval analysis states that rational interval functions are inclusion monotonic. When a rational interval function is evaluated on a computer, care must be taken to preserve the interval monotonicity of the function. This can be achieved by controlling the rounding mode of the CPU's floating-point processor. Processors which are compliant with the IEEE floating-point specifications ([ANSI/IEEE \(1987\)](#)) can be set to round down or round up, among other modes. To maintain inclusion monotonicity, the results of a lower endpoint computation are always rounded down and the results of an upper endpoint computation are always rounded up. For example, on a hypothetical three-digit computer and using directed triangles to indicate the appropriate rounding, the real fraction $1/6$ is computed as $[1, 1]/[6, 6] = [\nabla(1/6), \triangle(1/6)] = [.166, .167]$. When intervals with many decimal digits are displayed, an easily-understood representation of intervals is, for example, $[\underline{.166}, \overline{.167}]$. Most rounding-mode control can be made transparent to the programmer with the aid of appropriate software packages, such as the C++ libraries BIAS and PROFIL developed by [Knüppel \(1993a,b\)](#). With the definition of interval data types and overloaded operators for interval types, programming interval computations can be as simple as $z := x + y$.

3.3 Bivariate F Distribution

The multivariate F distribution considered by [Schuurmann et al. \(1975\)](#) and [Krishnaiah \(1980b\)](#) is reconsidered here. Let $S = (s_{ij})$ be a Wishart random matrix with m degrees of freedom and $E(S) = m\Sigma = m(\sigma_{ij})$. The joint distribution of s_{11}, \dots, s_{pp} , the diagonal elements of S , is a multivariate χ^2 distribution with m degrees of freedom. The matrix Σ is the covariance matrix of the underlying multivariate normal random variable. Let $F_i = \frac{s_{ii}\sigma^2/m}{s^2\sigma_{ii}/n}$, $i = 1, \dots, p$ where s^2/σ^2 is independently distributed as a χ^2 random variable

with n degrees of freedom. Then the joint distribution of F_1, \dots, F_p is a multivariate F distribution with (m, n) degrees of freedom. When $p = 2$, $\rho = \sigma_{12}/(\sigma_{11}\sigma_{22})^{1/2}$ is the correlation between standard normal random variables that underlie the bivariate χ^2 distribution. The bivariate distribution of F_1 and F_2 , first introduced by [Krishnaiah \(1965\)](#), is given by

$$f(x_1, x_2) = \frac{n^{n/2}(1-\rho^2)^{(m+n)/2}}{\Gamma(\frac{m}{2})\Gamma(\frac{n}{2})} \sum_{i=0}^{\infty} \frac{\rho^{2i}\Gamma(m+\frac{n}{2}+2i)m^{m+2i}(x_1x_2)^{m/2+i-1}}{i!\Gamma(\frac{m}{2}+i)[n(1-\rho^2)+m(x_1+x_2)]^{m+n/2+2i}} \quad (3.1)$$

The distribution function can be expressed as

$$P(F_1 \leq d_1, F_2 \leq d_2) = (1-\rho^2)^{m/2} \sum_{j=0}^{\infty} \frac{\Gamma(\frac{m}{2}+j)}{j!\Gamma(\frac{m}{2})} \rho^{2j} B_j \quad (3.2)$$

where

$$B_j = \int_0^{\infty} \frac{e^{-z/2} z^{n/2-1}}{2^{n/2}\Gamma(n/2)} I_{1j} I_{2j} dz$$

and

$$I_{kj} = \frac{1}{2^{m/2+j}\Gamma(m/2+j)} \int_0^{\frac{d_k m z}{2n(1-\rho^2)}} e^{-u/2} u^{m/2+j-1} du = \Gamma\left(\frac{m}{2}+j, \frac{d_k m z}{2n(1-\rho^2)}\right) \quad (3.3)$$

where $\Gamma(\alpha, d)$ is the incomplete gamma function,

$$\Gamma(\alpha, d) = \int_0^d \frac{1}{\Gamma(\alpha)} x^{\alpha-1} e^{-x} dx.$$

3.4 Computation of Tables

[Krishnaiah \(1980b\)](#) actually gives expressions for probability integrals over arbitrary rectangular regions, $P(c_1 \leq F_1 \leq d_1, c_2 \leq F_2 \leq d_2)$, but in all published tables of critical points, $c_1 = c_2 = 0$. The challenging aspect of computing an interval enclosure of a critical point for the bivariate F distribution is to find an appropriate rational interval function which gives reasonably tight bounds for the enclosure.

When the infinite series in (3.2) is truncated after $t+1$ terms, a bound on the truncation error R_t given by [Schuurmann et al. \(1975\)](#) is:

$$R_t \leq 1 - (1-\rho^2)^{m/2} \sum_{j=0}^t \frac{\Gamma(\frac{m}{2}+j)}{j!\Gamma(\frac{m}{2})} \rho^{2j}. \quad (3.4)$$

Let P_t denote the series in (3.2) truncated after $t+1$ terms. Also, let \mathbf{p}_t and \mathbf{r}_t denote interval enclosures of P_t and R_t respectively. Then

$$P[Y_1 \leq d_1, Y_2 \leq d_2] = P_t + R_t \in [\underline{p}_t, \bar{p}_t + \bar{r}_t]$$

for all t . The stopping value of t which is used will depend on machine and/or software precision. In practice, \mathbf{r}_t is computed successively and iteration stops when $\mathbf{r}_{t-1} = \mathbf{r}_t$.

Termination could also be specified to occur when the width of \mathbf{r}_t is less than a specified tolerance.

A guaranteed enclosure of the incomplete gamma function is available by using methods found in [Wang and Kennedy \(1994b\)](#). To compute an enclosure for B_j via a rational interval function, ideas similar to those of [Amos and Bulgren \(1972\)](#) are used. The integral is split into three pieces

$$\int_0^\infty \cdot dz = \int_0^{\varepsilon_1} \cdot dz + \int_{\varepsilon_1}^{\varepsilon_2} \cdot dz + \int_{\varepsilon_2}^\infty \cdot dz \quad (3.5)$$

The left and right tails of B_j can be bound in the following ways:

$$0 \leq \int_0^{\varepsilon_1} \cdot dz \leq \varepsilon_1$$

$$0 \leq \int_{\varepsilon_2}^\infty \cdot dz \leq 1 - \Gamma\left(\frac{n}{2}, \frac{\varepsilon_2}{2}\right)$$

The middle integrand in the right-hand side of (3.5) covers a finite domain, over which the second derivative exists, and is computed via the use of first-order Newton-Cotes quadrature. Some references to numerical quadrature with automatic result verification appear in [Kelch \(1993\)](#). The general form of Newton-Cotes quadrature is

$$\int_a^b f(x)dx = h \left(\frac{1}{2}f(a) + f(a+h) + f(a+2h) + \dots + f(a+(m-1)h) + \frac{1}{2}f(a+hm) \right) + E \quad (3.6)$$

where $h = (b-a)/m$ and the error has the form

$$E = -\frac{(b-a)^3}{12m^2} f''(\xi)$$

for some $\xi \in (a, b)$. The interval extension of E involves the computation of $\mathbf{f}''([a, b])$. If f'' is a rational function, as in this problem, then the interval extension \mathbf{f}'' is inclusion monotonic and the width of $\mathbf{f}''([a, b])$ is likely to be greater than $\mathbf{f}''([a', b'])$ for $[a', b'] \subset [a, b]$. Minimizing the width of the error term is one of the steps in achieving highly accurate final results. For this reason, the middle integral in the right-hand side of (3.5) is actually computed as the sum of a series of subintegrals, each of which is evaluated by numerical quadrature.

$$\int_{\varepsilon_1}^{\varepsilon_2} \cdot dz = \int_{\varepsilon_1}^{\varepsilon_1+\nu} \cdot dz + \int_{\varepsilon_1+\nu}^{\varepsilon_1+2\nu} \cdot dz + \dots + \int_{\varepsilon_1+(k-1)\nu}^{\varepsilon_1+k\nu} \cdot dz + \int_{\varepsilon_1+k\nu}^{\varepsilon_2} \cdot dz \quad (3.7)$$

Tuning the parameters $(\varepsilon_1, \varepsilon_2, k, \nu, h, m)$ of the method used here is not an immediately straightforward matter. Generally speaking, increasing the number of quadrature points will increase the accuracy (i.e. narrowness) of the final answer. A limit is reached, however, when increasing the number of quadrature points becomes counter-productive. This happens because each interval function evaluation at a quadrature point results in

a slight amount of overestimation and underestimation of the true value. Increasing m results in a narrower enclosure of the error E for each integrand, but eventually this gain is nullified by the sum of the overestimated and underestimated function values. A similar phenomenon occurs in deciding how many subintegrals to use in evaluating $\int_{\epsilon_1}^{\epsilon_2} \cdot dz$.

It is common in interval analysis to use so-called automatic derivatives for evaluation of the derivatives of a function. See [Moore \(1979\)](#) for an overview. In this research, computed enclosures of f'' were found to be much narrower when a hand-derived expression for f'' was coded into the software. The resulting expression involves evaluation of $x^{m/2-2}$ over an interval with a lower endpoint of 0. This limits the degrees of freedom to $m \geq 5$.

Tables of critical values d were computed for

$$P(F_1 \leq d, F_2 \leq d) = \alpha. \quad (3.8)$$

Critical values for non-rectangular regions could easily be computed uniquely by specifying an additional constraint. Solving the integral equation (3.8) for d requires the use of an iterative algorithm for finding roots, e.g. Newton-Raphson. This paper uses an intervalized bracket-secant algorithm, switching to an intervalized bisection algorithm to further narrow the enclosure of the critical point after the bracket-secant algorithm terminates. Complete details of the method appear in [Wright and Kennedy \(1999\)](#).

3.5 Conclusions

The computation of self-validated critical values for this bivariate F distribution is very time-consuming. The probability content of a rectangular region can be computed in a few minutes, but this is likely to be prohibitively costly for implementation of real-time computation of critical values. Since determination of critical points involves finding the roots of an equation, each entry in the table required several hours to compute on a DEC 5000 workstation. (Less time would be required for wider enclosures.) For this distribution, the real utility of interval analysis is the verification and guarantee of accuracy of previously published tables. The tables published in [Schuurmann et al. \(1975\)](#) and [Krishnaiah \(1980b\)](#) are generally quite accurate, but do have slight errors in the last (hundredths) digit which are likely due to rounding, exactly the kind of error which interval analysis can eliminate. Tables 3.1 and 3.2 are examples of the accuracy achieved by the software developed for this research.

Table 3.1: Upper 0.05 percentile points of the bivariate F distribution

m	$n = 10$			
	$\rho = 0.1$	$\rho = 0.3$	$\rho = 0.5$	$\rho = 0.7$
2	5.31734 ⁵⁸ ₁₆	5.2768 ¹³ ₀₈	5.18758 ⁹⁶ ₇₄	5.02529 ⁷⁸ ₃₄
4	4.3110 ⁸¹ ₇₇	4.28342 ³⁶ ₀₂	4.22239 ⁶³ ₂₈	4.11107 ⁵² ₁₂
6	3.89658 ⁸¹ ₄₉	3.87377 ⁴⁷ ₁₄	3.82350 ⁵⁵ ₂₀	3.73205 ⁷⁹ ₃₈
8	3.66318 ⁴⁹ ₃₄	3.64307 ⁸¹ ₄₈	3.59884 ⁹⁵ ₅₉	3.51864 ⁴⁵ ₀₁
10	3.51093 ⁹⁴ ₆₀	3.49262 ⁹⁹ ₆₅	3.4524 ²³ ₁₈	3.37971 ⁸⁴ ₃₅

Table 3.2: Upper 0.01 percentile points of the bivariate F distribution

m	$n = 10$			
	$\rho = 0.1$	$\rho = 0.3$	$\rho = 0.5$	$\rho = 0.7$
2	9.3014 ⁹⁹ ₈₁	9.2532 ²⁴ ₀₉	9.1430 ⁶¹ ₄₆	8.9312 ¹⁷ ₀₁
4	7.1869 ⁹⁸ ₈₇	7.1526 ⁷⁰ ₅₈	7.0749 ³⁶ ₂₄	6.9271 ⁹⁴ ₈₃
6	6.3616 ⁹⁹ ₇₇	6.3325 ⁹³ ₈₃	6.26714 ⁹⁶ ₀₃	6.1440 ⁶⁷ ₅₈
8	5.9094 ⁸³ ₆₁	5.8833 ²⁸ ₁₉	5.8248 ²³ ₁₄	5.7156 ⁷⁵ ₆₆
10	5.6197 ⁶⁰ ₃₈	5.5955 ⁹⁶ ₈₇	5.5417 ⁵⁶ ₄₇	5.4419 ³⁸ ₂₉

Chapter 4

AN INTERVAL ANALYSIS APPROACH TO THE EM ALGORITHM

Abstract

The EM algorithm is widely used in incomplete-data problems (and some complete-data problems) for parameter estimation. One limitation of the EM algorithm is that upon termination, it is not always near a global optimum. As reported by [Wu \(1982\)](#), when several stationary points exist, convergence to a particular stationary point depends on the choice of starting point. Furthermore, convergence to a saddle point or local minimum is also possible. In the EM algorithm, although the loglikelihood is unknown, an interval containing the gradient of the EM q function can be computed at individual points using interval analysis methods. By using interval analysis to enclose the gradient of the EM q function (and, consequently, the loglikelihood), an algorithm is developed which is able to locate all stationary points of the loglikelihood within any designated region of the parameter space. The algorithm is applied to several examples. In one example involving the t distribution, the algorithm successfully locates (all) seven stationary points of the loglikelihood.

Key Words: Interval arithmetic, Optimization, Interval EM

4.1 Introduction

This article explores a variation of the EM algorithm which uses techniques of interval analysis to locate multiple stationary points of a loglikelihood.

Interval analysis can be used to compute an interval which encloses the range of a function over a given domain. By using interval analysis to compute an enclosure of the gradient of the loglikelihood over specific regions, those regions where the enclosure of the gradient does not contain zero can be ruled out from containing any stationary points.

The algorithm locates stationary points by repeatedly dividing into smaller regions precisely those regions which have not been ruled out.

The structure of this paper proceeds as follows. Section 4.2 presents an introduction to interval analysis sufficient to understand this paper. Some of the differences between calculations with real numbers and interval numbers are noted, along with some comments about performing interval arithmetic on digital computers. Section 4.3 briefly states the traditional EM algorithm, then Section 4.4 presents a new approach to the EM algorithm using interval analysis. Section 4.5 presents several examples of the algorithm applied to different problems. These examples demonstrate both the accuracy which interval arithmetic can provide and the ability of the algorithm to locate multiple stationary points. Section 4.6 provides some conclusions.

4.2 Interval Analysis

A good introduction to Interval Analysis can be found in monographs by Hansen (1992) and Moore (1979). Some of the fundamental concepts of interval analysis are now presented.

In this paper, intervals will be indicated by superscript I and vectors will be denoted by boldface. An interval $x^I = [\underline{x}, \bar{x}]$ is a closed and bounded set of real numbers. For two intervals x^I and y^I , interval arithmetic operators are defined in the following manner:

$$x^I \circ y^I = \{x \circ y : x \in x^I, y \in y^I\}$$

where $\circ \in \{+, -, *, /\}$ and division is undefined for $0 \in y^I$. For these four interval arithmetic operators, closed-form expressions can be obtained for direct calculation of results of the operations. For example, if $x^I = [\underline{x}, \bar{x}]$ and $y^I = [\underline{y}, \bar{y}]$, then $x^I + y^I = [\underline{x} + \underline{y}, \bar{x} + \bar{y}]$. The *Hull* of a set of intervals x_1^I, \dots, x_n^I is the smallest interval containing x_1^I, \dots, x_n^I , i.e. $Hull(x_1^I, \dots, x_n^I) = [\inf\{x : x \in x_i^I; i = 1, \dots, n\}, \sup\{x : x \in x_i^I; i = 1, \dots, n\}]$. An *interval vector* or *box* is simply a vector of intervals. An *interval function* is an interval-valued function of one or more interval arguments. In this paper, capital letters are used to denote interval functions. An interval function $F(x_1^I, \dots, x_n^I)$ is said to be an *interval extension* or *interval enclosure* of $f(x_1, \dots, x_n)$ if $F([x_1, x_1], \dots, [x_n, x_n]) = f(x_1, \dots, x_n)$ for all $x_i; i = 1, \dots, n$. An interval function F is said to be *inclusion monotonic* if $F(x^I) \subset F(y^I)$ whenever $x^I \subset y^I$. A fundamental property of interval analysis is that rational interval functions are inclusion monotonic.

In this paper, the *natural interval extension* of a real function is used. This is an interval extension in which intervals and interval operations are substituted for scalars and scalar operations. The value of any interval extension of a function is dependent on the form of the real function. For example, let $f_1(x) = (x - 1)(x + 1)$ and $f_2(x) = xx - 1$. Let F_1 and F_2 be the corresponding natural interval extensions and let $x^I = [-2, 1]$. Then $F_1(x^I) = [-6, 3]$ and $F_2(x^I) = [-3, 3]$ which both contain $[-1, 3]$, the true range of f_1 and f_2 over x^I . This feature of interval computations to sometimes overestimate the range of a function is referred to as *interval dependency*. Attention must be given to the exact expression of an interval function to reduce the effect of interval dependency. Hansen (1992, 1997) presents some results regarding this topic.

When implementing interval arithmetic calculations on computers, care must be taken to ensure that rounding errors do not invalidate the inclusion monotonicity of interval results. One way to achieve this is through the use of directed rounding modes in the floating-point calculations. When calculating the lower endpoint of an interval result, the floating-point processor is set to round all results *down*. For calculation of the upper endpoint of an interval result, all calculations are rounded *up*. Using the symbols ∇ and Δ to denote downward and upward rounding respectively, the actual computer implementation of interval addition is $x^I + y^I = [\nabla(\underline{x} + \underline{y}), \Delta(\bar{x} + \bar{y})]$. Correct use of the rounding modes guarantees that the computed result contains the true interval answer. Some programming languages and software packages are able to work with interval data types and interval operators. INTLIB_90 and C-XSC are among the more widely referenced tools. Some mention of other software environments is found in [Kearfott \(1996b\)](#). For the research in this paper, the computations were done using the BIAS/PROFIL package in C++ developed by [Knüppel \(1993c\)](#).

4.3 The EM Algorithm

The present-day incarnation of the EM algorithm first appeared in a landmark paper by [Dempster et al. \(1977\)](#). The EM algorithm is a general iterative algorithm for maximum likelihood estimation in incomplete-data problems. The EM algorithm has not only been successfully applied in obvious incomplete-data problems, but also in many situations where the data appears to be complete, but can be viewed as incomplete by introducing latent variables. The intuitive idea behind the EM algorithm is to iterate the following two steps:

Expectation step: Replace missing values (sufficient statistics) by estimated values.

Maximization step: Estimate parameters as if no data were missing.

Formally, starting with a parameter estimate ϕ_p , the E-step calculates the conditional expectation of the complete-data log likelihood, $\log L_c(\phi)$, as $q(\phi|\phi_p) = E_{\phi_p}\{\log L_c(\phi)\}$ and then the M-step chooses ϕ_{p+1} to be any value of $\phi \in \Omega$ that maximizes $q(\phi|\phi_p)$, i.e. $q(\phi_{p+1}|\phi_p) \geq q(\phi|\phi_p)$ for any $\phi \in \Omega$.

4.4 Interval Arithmetic and EM

A method is now presented which uses certain properties of the EM algorithm and of interval arithmetic to locate all stationary points of the likelihood inside of a given region of the parameter space. Briefly, from the EM algorithm it is known that the q function has a gradient which is equal to the gradient of the loglikelihood at stationary points of the loglikelihood. Using interval arithmetic, it is possible to derive interval vectors which enclose values of the gradient of the q function even over regions which do not contain a stationary point.

The complete method is presented below, followed by a summary outline and additional comments. Some numerical results are presented in Section 4.5.

4.4.1 Enclosing the Gradient of the Log Likelihood

The fundamental task for the method being proposed will be to eliminate regions of the parameter space where it can be determined that a stationary point of the likelihood does not exist. This can be accomplished by finding a box which encloses the range of the gradient of the loglikelihood over a region. If, for example, the interval enclosure of the set of all values of the gradient of the loglikelihood $\ell(\phi)$ over the box ϕ^I ,

$$\left\{ \frac{\partial \ell(\phi)}{\partial \phi} \Big|_{\phi=\phi_p} : \phi_p \in \phi^I \right\},$$

does not contain zero in one or more of its coordinates, then the gradient of the log likelihood is nonzero over ϕ^I and $\ell(\phi)$ does not contain a stationary point inside the box ϕ^I . A more thorough explanation of how this is accomplished is now presented by deriving an interval enclosure for the gradient of the log likelihood. The first part of this derivation is similar to the development in Dempster et al. (1977).

Denote the complete data (which includes missing values) by \mathbf{x} and the observed (incomplete) data by \mathbf{y} , where $\mathbf{y} = \mathbf{y}(\mathbf{x})$. Let the density function of \mathbf{x} be $f(\mathbf{x}|\phi)$, where $\phi \in \Omega$. From this, the density function for \mathbf{y} is

$$g(\mathbf{y}|\phi) = \int_{\mathbf{x}(\mathbf{y})} f(\mathbf{x}|\phi) d\mathbf{x}.$$

For simplicity and tractability, the maximization step would ideally be accomplished over ϕ in $\log f(\mathbf{x}|\phi)$. However, since \mathbf{x} is unobservable, replace $\log f(\mathbf{x}|\phi)$ by its conditional expectation. To that end, let $k(\mathbf{x}|\mathbf{y}, \phi) = f(\mathbf{x}|\phi)/g(\mathbf{y}|\phi)$ be the conditional density of \mathbf{x} given \mathbf{y} and ϕ . Using this, the log-likelihood can be written

$$\ell(\phi) = \log g(\mathbf{y}|\phi) = \log f(\mathbf{x}|\phi) - \log k(\mathbf{x}|\mathbf{y}, \phi).$$

Taking the conditional expectation (using ϕ_p as an estimate for ϕ),

$$\ell(\phi) = E_{\phi_p} [\log f(\mathbf{x}|\phi)|\mathbf{y}] - E_{\phi_p} [\log k(\mathbf{x}|\mathbf{y}, \phi)|\mathbf{y}].$$

For simplicity, this is often written $\ell(\phi) = q(\phi|\phi_p) - h(\phi|\phi_p)$. To find values of $\phi \in \Omega$ which maximize $\ell(\phi)$, solutions to

$$\frac{\partial \ell(\phi)}{\partial \phi} = \frac{\partial q(\phi|\phi_p)}{\partial \phi} - \frac{\partial h(\phi|\phi_p)}{\partial \phi} = 0$$

are needed.

Now, it is easy to show that $h(\phi|\phi_p) \leq h(\phi_p|\phi_p)$ for any $\phi \in \Omega$, i.e. ϕ_p maximizes $h(\phi|\phi_p)$ with respect to ϕ , and so $\frac{\partial h(\phi|\phi_p)}{\partial \phi} \Big|_{\phi=\phi_p} = 0$. It is therefore sufficient

when searching for maxima of $\ell(\phi)$ to limit consideration to $\frac{\partial q(\phi|\phi_p)}{\partial \phi}$, specifically, to an enclosure of the gradient of the q function over the box ϕ^I ,

$$\left\{ \frac{\partial q(\phi|\phi_p)}{\partial \phi} \Big|_{\phi=\phi_p} : \phi_p \in \phi^I \right\}$$

for arbitrary $\phi^I \in \Omega$. Let $Q'(\phi|\phi^I) = [\underline{Q}'(\phi|\phi^I), \overline{Q}'(\phi|\phi^I)]$ be an interval extension of $\frac{\partial q(\phi|\phi_p)}{\partial \phi}$ for interval ϕ^I and $\phi_p \in \phi^I$. Note that $Q'(\phi|\phi^I)$ is *not* $\frac{\partial Q(\phi|\phi^I)}{\partial \phi}$ where $Q(\phi|\phi^I)$ is the interval extension of $q(\phi|\phi_p)$. Also, let $Q'_2(\phi^I|\phi^I)$ be an interval extension of $Q'(\phi|\phi^I)$.

At each $\phi_p \in \phi^I$, the enclosure of the gradient of the log likelihood can be obtained by

$$\frac{\partial \ell(\phi)}{\partial \phi} \Big|_{\phi=\phi_p} = \frac{\partial q(\phi|\phi_p)}{\partial \phi} \Big|_{\phi=\phi_p} \in [\underline{Q}'(\phi_p|\phi^I), \overline{Q}'(\phi_p|\phi^I)]$$

and

$$\left\{ \frac{\partial \ell(\phi)}{\partial \phi} \Big|_{\phi=\phi_p} : \phi_p \in \phi^I \right\} \subset Q'_2(\phi^I|\phi^I).$$

If zero is not contained in $Q'_2(\phi^I|\phi^I)$, then the box ϕ^I cannot contain a local maximizer of $\ell(\phi)$ and may therefore be excluded from further consideration. Thus $Q'_2(\phi^I|\phi^I)$ is an interval-valued function which encloses the union of the ranges of a class of interval functions $q(\phi|\phi^I)$ indexed by $\phi \in \phi^I$.

After a user of this method specifies an initial box $\phi^I \in \Omega$, locating optima of the loglikelihood proceeds by conducting a bisection search by dividing ϕ^I into successively smaller boxes and evaluating the enclosure of the gradient of the log likelihood over each box. Boxes which do not contain a stationary point are discarded. The initial box ϕ^I will frequently be quite large so as to (hopefully) enclose all stationary points of $\ell(\phi)$. At a certain point in this process, typically when the box size becomes smaller than a specified size, the subdividing stops and a list \mathcal{G} of boxes from the grid search is output along with the enclosure of the gradient and the enclosure of the range of q functions over each box. These boxes contain all the stationary points of $\ell(\phi)$ that exist within the initial interval box ϕ^I .

4.4.2 Definitions for Interval EM

In this section an interval EM algorithm is defined. A few necessary definitions are stated and then utilized in the interval EM method being presented.

Definition. An *interval EM algorithm* on an interval vector Φ in a parameter space Ω is an iterative method which employs a sequences of intervals $\phi_0^I \rightarrow \phi_1^I \rightarrow \cdots \phi_p^I \rightarrow$ with respect to interval enclosures $Q(\phi|\phi_0^I), Q(\phi|\phi_1^I), \dots, Q(\phi|\phi_p^I)$ of sets of functions $q(\phi|\phi_0), q(\phi|\phi_1), \dots, q(\phi|\phi_p)$ so that $q(\phi|\phi_p) \in Q(\phi|\phi_p^I)$ where $\phi_p \in \phi_p^I \subset \Phi$ for each p . The interval ϕ_{p+1}^I contains at least one value of ϕ_{p+1} which maximizes a $q(\phi|\phi_p)$ for at least one $\phi_p \in \phi_p^I \subset \Phi$. Moving from ϕ_p^I to ϕ_{p+1}^I is referred to as an *interval EM step*.

Definition. An *interval GEM algorithm* is an interval EM algorithm except instead of maximizing $q(\phi|\phi_p^I)$ with respect to ϕ , the interval ϕ_{p+1}^I contains as least one value ϕ_{p+1} such that $q(\phi_{p+1}|\phi_p) \geq q(\phi_p|\phi_p)$, where $\phi_{p+1} \in \phi_{p+1}^I$, $\phi_p \in \phi_p^I$. Moving from ϕ_p^I to ϕ_{p+1}^I is referred to as an *interval GEM step*.

The current implementation of the method may be more easily understood by referring to Figure 4.1, which graphically illustrates an interval EM step in a hypothetical one-dimensional case. In Figure 4.1, the dotted lines $q(\phi|\phi_i)$ and $q(\phi|\phi_j)$ are two separate scalar q functions that might be encountered in different iterations of a scalar EM algorithm. The solid lines $\underline{Q}(\phi|\phi_k^I)$ and $\overline{Q}(\phi|\phi_k^I)$ denote the extent of an interval-valued function $Q(\phi|\phi_k^I)$ which encloses all the scalar q functions $q(\phi|\phi_i)$ and $q(\phi|\phi_j)$ indexed by $\phi_i \in \phi_k^I$, $\phi_j \in \phi_k^I$. Finally, the vertical line segments $Q(\phi_{k+1,1}^I|\phi_{k+1,1}^I)$ and $Q(\phi_{k+1,2}^I|\phi_{k+1,2}^I)$ denote enclosures of $Q(\phi|\phi^I)$ evaluated for $\phi \in \phi_{k+1,1}^I$ and $\phi \in \phi_{k+1,2}^I$, respectively.

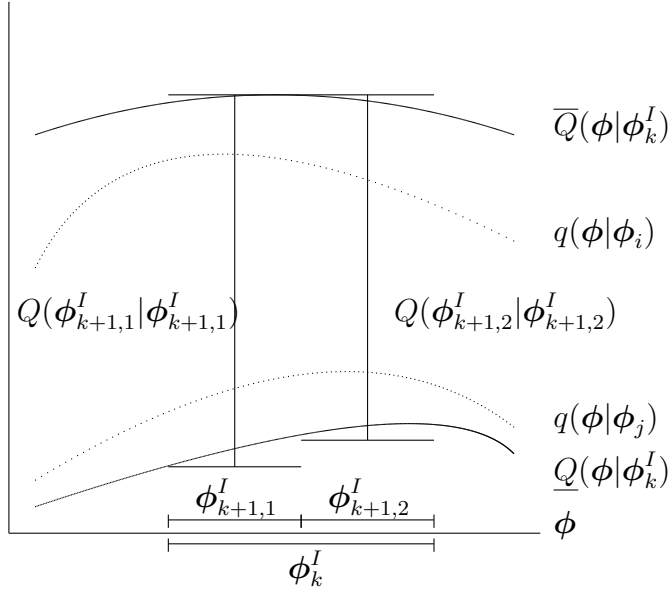


Figure 4.1: One interval EM step. $\overline{Q}(\phi|\phi_k^I)$ and $\underline{Q}(\phi|\phi_k^I)$ bound the extent of the interval-valued function $Q(\phi|\phi_k^I)$, while $q(\phi|\phi_i)$ and $q(\phi|\phi_j)$ are examples of two of the scalar functions contained within the interval function.

4.4.3 Full Bisection Search

The bisection algorithm starts with an initial box ϕ_0^I in a list of boxes \mathcal{G} . The method proceeds by simply bisecting boxes from \mathcal{G} until no boxes are left or until all boxes have reached a certain size. Let m be the dimension of ϕ and initialize $i := 1$. Proceed by removing and bisecting each box of \mathcal{G} in the i^{th} coordinate. Discard any boxes which do not contain zero in at least one direction of the enclosure of the gradient. Return all remaining boxes to \mathcal{G} and increase i by 1, resetting $i := 1$ when $i > m$. Repeat as

necessary until the diameter of every box is small. If at any point \mathcal{G} becomes empty, print a message stating that no stationary points were contained in the initial region ϕ_0^I .

The bisection algorithm differs from traditional EM in that there are no expectation and maximization steps. The only use of the EM theory was to obtain an enclosure for the gradient of the loglikelihood of ϕ . Still, the bisection search can in some way be viewed as many simultaneous interval GEM algorithms. In making an interval GEM step from ϕ_k^I to $\phi_{k+1,1}^I$ and from ϕ_k^I to $\phi_{k+1,2}^I$, there will be a nondecreasing change in the lower bound of the enclosure of the q functions, i.e. $\underline{Q}(\phi_k^I|\phi_k^I) \leq \underline{Q}(\phi_{k+1,i}^I|\phi_{k+1,i}^I)$ for $i = 1, 2$.

- **ALGORITHM: Bisection Interval EM Search**

Input an initial interval box ϕ_0^I and place it as the only element of the list \mathcal{G} .

i := 0

REPEAT

 i := (i + 1) mod m

 FOR j = 1 TO Length(\mathcal{G})

 Remove the first box from \mathcal{G} . Call it ϕ^I

 Bisect ϕ^I along the i^{th} direction, creating ϕ_1^I and ϕ_2^I

 If $0 \in Q'_2(\phi_k^I|\phi_k^I)$, append ϕ_k^I to \mathcal{G} , $k = 1, 2$

 NEXT

UNTIL \mathcal{G} is empty or maximum diameter of boxes $\leq \epsilon$

The method described above will not, of course, find any global optima which lie outside of the initial box $\phi_0^I \subset \Omega$. In practice this is often not of concern, primarily because the observed data places practical limitations on the portion of the parameter space of interest. Also, in a manner similar to that observed by Hansen (1992), it is often possible to make the parameter space exceedingly large without significantly increasing the computing time to search for global optima.

Because the algorithm uses intervals instead of real numbers, measurement error in data and floating-point approximations can immediately be incorporated. For example, one might use $\pi^I = [3.14, 3.15]$ to indicate uncertainty in known constants. Even more useful is the ability to represent data as intervals, e.g. $x_i^I = [x_i - \delta, x_i + \delta]$, where x_i is the observed value and δ is a bound on the measurement error.

4.4.4 Quick Search

For reasons of speed, memory, or accuracy considerations, the method described above may not always be optimally suitable. Let m be the dimension of ϕ . Bisection of just one box from \mathcal{G} has the potential to create 2^m additional boxes that will be added to \mathcal{G} . This might happen in situations where a region contains many stationary points or where the loglikelihood is relatively flat and the gradient is near zero. Since interval arithmetic

sometimes calculates an interval wider than optimal, it may be the case that the gradient is nonzero in every direction, but the enclosure of the gradient contains zero in at least one direction. If some combination of high dimensionality and/or fairly flat likelihood occurs, the length of \mathcal{G} can grow exponentially.

A variation on the algorithm given above is now presented as a faster, smaller alternative. The variation comes about simply as a matter of which order the boxes of \mathcal{G} are added to the list.

As before, boxes are removed from the start of the list \mathcal{G} . After removal, the box ϕ is bisected in the coordinate of the maximum width of ϕ . Boxes for which the enclosure of the gradient does not contain zero are discarded. If only one half of ϕ remains, prepend it to the list \mathcal{G} . If both halves remain, then evaluate and enclosure of the q function, $Q(\phi_i^I|\phi_i^I)$, for each half. Prepend both halves to \mathcal{G} , with the half that has the greater lower bound of $Q(\phi_i^I|\phi_i^I)$ added last. (In Figure 4.1, ϕ_2 is prepended after ϕ_1 .) When the first box of \mathcal{G} has reached a user-specified tolerance size, the algorithm stops and prints out only the first box of \mathcal{G} .

This approach will ensure that at least one box ϕ at the start of the list \mathcal{G} is made as small as possible in as short of time as possible. Of course, the gain in speed comes at the price of losing the guarantee that a stationary point is contained in the output. It could happen that if a smaller tolerance was used, at the next step the box would be bisected and both halves discarded. All that can be said is that in the final box output, the enclosure of the gradient over the box contains zero in each direction. Nonetheless, given the speed advantages, this algorithm is potentially useful. It may provide a quick answer which can suggest a location for a fairly small region to feed into the full bisection search given above.

- **ALGORITHM: Quick Interval EM Search**

Input an initial box ϕ_0^I and place it as the only element of the list \mathcal{G} .

Input ϵ

REPEAT

Remove the first box from \mathcal{G} . Call it ϕ^I

Bisect ϕ^I along the direction of maximum width, creating ϕ_1^I and ϕ_2^I

If $0 \in Q_2'(\phi_k^I|\phi_k^I)$, prepend ϕ_k^I to \mathcal{G} , $k = 1, 2$.

UNTIL \mathcal{G} is empty or the first box of \mathcal{G} has maximum diameter $\leq \epsilon$

4.5 Examples

Several examples are now presented to illustrate use of the method described above. Note that the following examples each have an algebraic, real expression for $q(\phi|\phi_k)$. This is consistent with traditional EM notation. Though not shown, a person would then determine an expression for the gradient of this function with respect to ϕ , $q'(\phi|\phi_k)$,

and then express $q'(\phi_k|\phi_k)$ in as simple a way as possible. This is coded in the program as $Q'_2(\phi_k|\phi_k)$.

When numerical results are reported, sub/superscript notation will sometimes be used to simplify the representation of an interval, e.g. $[2.33, 2.35] = 2.3_3^5$.

4.5.1 Multinomial Example

The following example from [Dempster et al. \(1977\)](#) is frequently used to introduce the EM algorithm. Consider a set of 197 animals which are classified into four categories. The observed classification counts are $\mathbf{y} = (y_1, y_2, y_3, y_4) = (125, 18, 20, 34)$. The classification of the random variable Y is modeled as following a multinomial distribution:

$$Y \sim \text{Multinomial} \left(197, \frac{1}{2} + \frac{1}{4}p, \frac{1}{4} - \frac{1}{4}p, \frac{1}{4} - \frac{1}{4}p, \frac{1}{4}p \right)$$

where p is unknown and to be estimated. There is no missing data in this problem and p is easily estimated by a maximum likelihood approach. For illustration purposes, the problem is reformulated with missing data. Suppose the first classification category Y_1 is split into two categories and a new random variable X is modeled:

$$X \sim \text{Multinomial} \left(197, \frac{1}{2}, \frac{1}{4}p, \frac{1}{4} - \frac{1}{4}p, \frac{1}{4} - \frac{1}{4}p, \frac{1}{4}p \right).$$

The incomplete data vector \mathbf{x} is $(x_1, x_2, x_3, x_4, x_5)$ and thus \mathbf{y} can be written $\mathbf{y}(\mathbf{x}) = (x_1 + x_2, x_3, x_4, x_5)$. Here x_1 and x_2 are unobserved except through their sum $x_1 + x_2$. It can be shown that

$$q(p|p_k) = k(\mathbf{x}) + \left[125 \frac{\frac{p_k}{4}}{\frac{1}{2} + \frac{p_k}{4}} + x_5 \right] \frac{1}{p} - (x_3 + x_4) \frac{1}{1-p} \quad (4.1)$$

where $k(\mathbf{x})$ does not depend on p and can be ignored in the maximization step. Figure 4.2 shows a plot of the corresponding interval extension, $Q(p|p_k^I)$. An accurate interpretation of this interval-valued function can be had in this case by actually overlaying plots of $q(p|p_k)$ for various $p \in p_k^I$, in this case $p = 0.1(0.1)0.9$.

The initial interval selected is $p_0^I = [.00001, .99999]$. While a wider interval can be used, the maximum likelihood estimate of p is certainly contained in $[.00001, .99999]$. Furthermore, the values of $p = 0$ and $p = 1$ are excluded by equation (4.1). If the user selected an inappropriate value for the initial interval, such as $p^I = [0.1, 0.2]$, then the algorithm terminates with the message:

```
Gradient of Q(Phi|Phi_k) = ([152.262,411.414])
Gradient of likelihood does not contain zero.
No stationary point in ([0.1,0.2])
```

The bisection algorithm applied to this problem using initial interval p_0^I produces a list \mathcal{G} which contains two interval boxes,

$$y_1 = 0.626821497870982_3^4$$

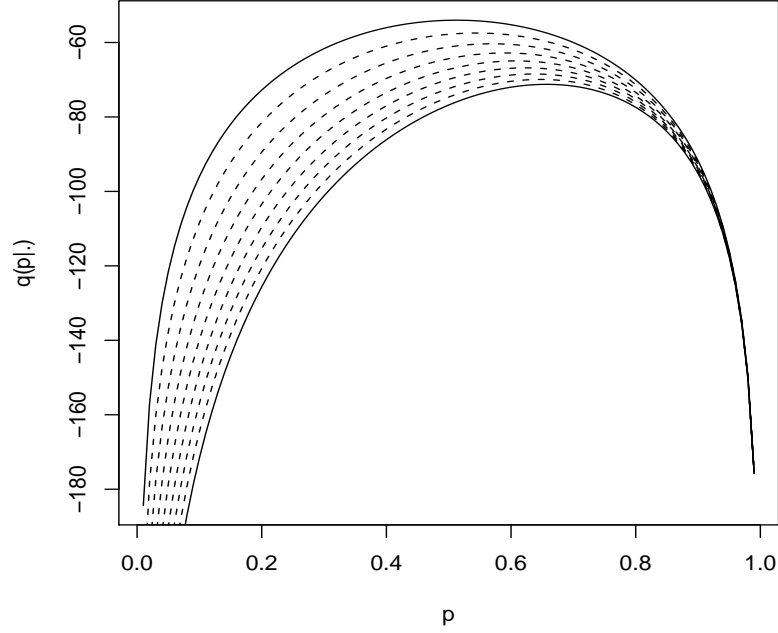


Figure 4.2: Plot of $Q(p|p_0^I)$ versus p for $p_0^I = [0.1, 0.9]$.

$$y_2 = 0.626821497870982_4^5.$$

Any stationary points of the log-likelihood are guaranteed to be contained in the hull of the boxes in the list \mathcal{G} . If a scalar estimate is desired, the midpoint of the hull can be given: $\hat{p} = 0.6268214978709824$.

4.5.2 Univariate t Example

McLachlan and Krishnan (1997) give an example by Arslan et al. (1993) where the EM algorithm can converge to a local *minimum*. A p -dimensional random variable \mathbf{W} is said to have a multivariate t -distribution $t_p(\boldsymbol{\mu}, \Sigma, \nu)$ with location $\boldsymbol{\mu}$, positive definite inner product matrix Σ , and degrees of freedom ν when the density of \mathbf{W} is given by

$$f_p(\mathbf{w}|\boldsymbol{\mu}, \Sigma, \nu) = \frac{\Gamma(\frac{p+\nu}{2})|\Sigma|^{-1/2}}{(\pi\nu)^{p/2}\Gamma(\frac{\nu}{2})\{1 + (\mathbf{w} - \boldsymbol{\mu})^T \Sigma^{-1}(\mathbf{w} - \boldsymbol{\mu})/\nu\}^{(p+\nu)/2}}. \quad (4.2)$$

The example considered is a univariate case of the t -distribution where $\nu = 0.05$, $\Sigma = 1$, and μ is taken as unknown. The observed data is $\mathbf{w} = (-20, 1, 2, 3)$. Ignoring additive and multiplicative constants, the log likelihood is $\log L(\mu) \propto -\sum_i \log\{1 + 20(w_i - \mu)\}$. A plot showing the shape of this log likelihood appears in Figure 4.3.

The function has seven stationary points. The most interesting are the local maxima at $\mu_2 = 1.086$, $\mu_3 = 1.997$, and $\mu_4 = 2.906$. In this complete-data problem it is possible

to graph the log-likelihood and visually choose starting values that will cause a scalar

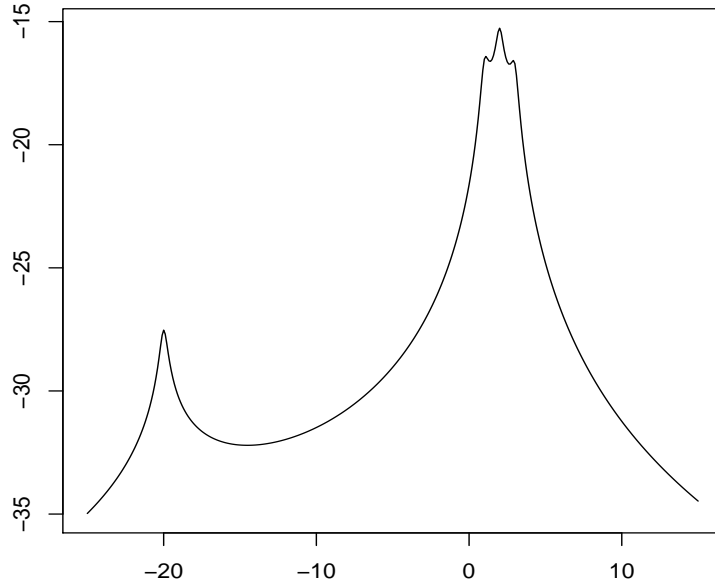


Figure 4.3: Plot of log likelihood function $\log L(\mu)$ versus μ . Local maxima occur at $\mu_1 = -19.993$, $\mu_2 = 1.086$, $\mu_3 = 1.997$, and $\mu_4 = 2.906$.

EM algorithm to converge to each of the local maxima, *and even to a local minimum*. However, the domain of attraction for each stationary point is not necessarily a contiguous region.

Using $\mu_0 = [-1000, 1000]$, the bisection algorithm completes 59 iterations (bisections), during which the length of \mathcal{G} is scarcely longer than the 20 boxes at the final step. These boxes occur in distinct groupings around each of the seven stationary points. While the algorithm actually outputs the list of boxes from \mathcal{G} , for brevity, the hull of each group of boxes and the hull of the associated enclosures of the q functions are given in Table 4.1.

Looking at this table, the nature of each stationary point is not immediately clear. Since this is a univariate case, it would be possible to evaluate the gradient on either side of each ϕ_{S_i} and thereby determine which stationary points are local maxima and which are local minima. However, it is immediately clear from the table that ϕ_{S_5} gives the largest value of $Q(\phi_{S_i}|\phi_{S_i})$ and contains the global maximum of the log-likelihood as displayed by Figure 4.3.

Table 4.1: Enclosures of the stationary points for the univariate t example.

i	ϕ_{S_i}	$Q(\phi_{S_i} \phi_{S_i})$
1	$-19.993164608871_{30}^{29}$	-1.57532666279595_7^4
2	-14.5161774794253_2^0	$-2.098837787645_{302}^{297}$
3	1.08616780631075_0^7	$-1.606093870388_{426}^{397}$
4	$1.3731761015634_{18}^{32}$	$-1.89224275084_{3016}^{2981}$
5	$1.9975126089118_{17}^{24}$	$-1.525009886703_{402}^{386}$
6	$2.6468546770426_{20}^{35}$	$-1.884158362286_{208}^{176}$
7	$2.9056308944679_{75}^{85}$	$-1.617024174245_{707}^{677}$

4.5.3 Binomial-Poisson Mixture Example

This example from [Thisted \(1988\)](#) presents a simple multivariate-parameter example dealing with the number of children per widow in a pension fund.

Children per widow, i	0	1	2	3	4	5	6
Number of widows, n_i	3062	587	284	103	33	4	2

Since the number of widows with no children is larger than would be expected for a Poisson distribution, it is assumed that there are actually two underlying populations. The number of children Y for a widow is modeled as

$$Y \sim \begin{cases} 0 & \text{with probability } \xi \\ \text{Poisson}(\lambda) & \text{with probability } 1 - \xi \end{cases} \quad (4.3)$$

With $\phi = (\lambda, \xi)$, the function to be maximized in the M-step is:

$$q(\phi|\phi_k) = \frac{n_0 \xi_k}{\xi_k + (1 - \xi_k) \exp(-\lambda_k)} \{ \log \xi - \log(1 - \xi) + \lambda \} + N \{ \log(1 - \xi) - \lambda \} + \sum_{i=1}^6 \{ i n_i \log \lambda - n_i \log i! \}. \quad (4.4)$$

Based on a visual examination of the data, the starting values of

$$\phi_0 = (\lambda_0^I, \xi_0^I) = ([0.001, 10], [0.001, 0.999])$$

were chosen as being certain to contain the true parameter values.

Applying the Bisection search, after 52 iterations of bisecting ϕ in both directions, the list \mathcal{G} contains 82 boxes, the first and last of which are

$$y_1 = (1.0378390789897_{57}^{60}, 0.61505669757312_{12}^{14})$$

$$y_{82} = (1.0378390789897_{77}^{80}, 0.61505669757312_{88}^{90}).$$

The hull of the boxes on this list is: $\phi_{S_1} = (1.0378390789897_{57}^{80}, 0.61505669757312_{12}^{90})$.

In this problem, what is important is not the extremely narrow (and hence) high degree of accuracy of ϕ_{S_1} , but the guarantee that considered over the initial parameter space ϕ_0 , the only stationary points of the log-likelihood (if any exist) are guaranteed to be contained in the interval box ϕ_{S_1} . Moreover, if a scalar EM algorithm converges to some stationary point in ϕ_0 , that point will be inside ϕ_{S_1} . Using $\epsilon = 10^{-15}$, the Quick search returns $(1.0378390789897_8^9, 0.61505669757313_2^3)$.

4.5.4 Genetic Example

This example is also taken from [McLachlan and Krishnan \(1997\)](#).

Suppose there are 435 observations from a multinomial distribution as given in Table 4.2 where $r = 1 - p - q$. The observed data is (n_O, n_A, n_B, n_{AB}) and the unknown

Table 4.2: Distribution of data in the genetic example.

Cell	Cell Probability	Observed Frequency
O	r^2	$n_O = 176$
A	$p^2 + 2pr$	$n_A = 182$
B	$q^2 + 2qr$	$n_B = 60$
AB	$2pq$	$n_{AB} = 17$

parameters are $\phi = (p, q)$. As in the multinomial example above, missing data is introduced by splitting the A and B cells across the sum in the cell probability. The q function is given by

$$\begin{aligned}
q(\phi|\phi_k) = & \left(\frac{182}{1 + 2(1 - p_k - q_k)/p_k} + 199 \right) \log(p) + \\
& \left(\frac{60}{1 + 2(1 - p_k - q_k)/q_k} + 77 \right) \log(q) + \\
& \left(594 - \frac{182}{1 + 2(1 - p_k - q_k)/p_k} - \frac{60}{1 + 2(1 - p_k - q_k)/q_k} \right) \log(1 - p - q). \quad (4.5)
\end{aligned}$$

It is not always possible to search the entire portion of the parameter space with one application of the bisection algorithm. In this example, certain combinations of p^I and q^I cause a division by zero error. Specifically, as illustrated in figure 4.4, the gradient does not exist along the lines $p = 0$, $q = 0$, $1 - p - q = 0$, $q = 2 - 2p$, and $2q = 2 - p$. The software can be written to catch division by zero errors and mark a box as containing such until further subdivision occurs. Alternatively, the user can specify a smaller initial region. The only stationary point located inside $\phi_0 = (p_0, q_0) = ([0.00001, 0.45], [0.00001, 0.45])$ is found to be located inside $\phi_S = (0.26444443138466_{694}^{706}; 0.09316881181568_{122}^{200})$.

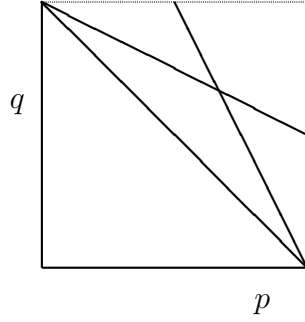


Figure 4.4: The unit square denotes the parameter space for (p, q) . Thick lines denote values for which a division by zero error will occur in calculating the gradient.

4.5.5 Multinomial Example Continued

The multinomial example considered above is presented again in two different ways to illustrate other ways to employ interval analysis to good advantage.

Intervalized Scalar EM Algorithm

For the incomplete-data problem, it is easy to show that

$$(X_2|X_1 + X_2 = 125) \sim \text{Binomial} \left(125, \frac{\frac{1}{4}p}{\frac{1}{2} + \frac{1}{4}p} \right)$$

and the E-Step in the usual scalar EM algorithm becomes $x_{1,k} = 125(\frac{1}{2})/(\frac{1}{2} + \frac{1}{4}p)$ and $x_{2,k} = 125(\frac{1}{4}p)/(\frac{1}{2} + \frac{1}{4}p)$. From the complete-data likelihood of p ,

$$f(p|\mathbf{x}) \propto \left(\frac{1}{2}\right)^{x_{1,k}} \left(\frac{1}{4}p\right)^{x_{2,k}} \left(\frac{1}{4} - \frac{1}{4}p\right)^{x_3+x_4} \left(\frac{1}{4}p\right)^{x_5}$$

the M-Step in the usual scalar EM algorithm is:

$$p_k = \frac{x_{2,k} + x_4}{x_{2,k} + x_3 + x_4 + x_5} = \frac{x_{2,k} + 34}{x_{2,k} + 72}.$$

Using $p_0 = 0.5$ as a starting value and using a convergence tolerance of $\epsilon = 10^{-7}$, the (scalar real) EM algorithm yields:

```
Epsilon:      1e-07
Initial p: 0.5
Iter    p      x2
1      0.608247  25
2      0.624321  29.1502
3      0.626489  29.7373
4      0.626777  29.8159
```

5	0.626816	29.8263
6	0.626821	29.8277
7	0.626821	29.8279
8	0.626821	29.8279

The algorithm converges at the specified tolerance after 8 iterations. In this case, the starting value of 0.5 for p was chosen simply because 0.5 lies exactly halfway between 0 and 1, which define the bounds for possible starting values. A questioning user may well wonder what results would be obtained for different starting values and how the steps of convergence might change. Interval analysis can be used to answer those questions.

This example can easily be programmed in interval arithmetic, though with a slight modification. Because of the dependency problem, narrower interval enclosures of computed values are more likely to be obtained if each variable appears only once in a calculation. The iterates in the EM algorithm for this particular example are therefore written equivalently as:

$$x_{2,k}^I = \frac{125.0}{2/p_k^I + 1} \quad \text{and} \quad p_k^I = 1 - \frac{38}{x_{2,k}^I + 72}$$

The convergence tolerance remains the same as above, but now $p_0^I = [\delta, 1]$ where δ is a small machine number greater than zero. The scalar EM algorithm using interval arithmetic produces the following output:

```

Epsilon: 1e-07
Initial p: [4.94066e-324,1]
i          p          x2
1  [0.472222,0.665689]  [0,41.6667]
2  [0.603656,0.631839]  [23.8764,31.2156]
3  [0.623692,0.627485]  [28.9812,30.0094]
4  [0.626405,0.626910]  [29.7144,29.852]
5  [0.626766,0.626833]  [29.8129,29.8311]
6  [0.626814,0.626823]  [29.8259,29.8284]
7  [0.626821,0.626822]  [29.8277,29.828]
8  [0.626821,0.626822]  [29.8279,29.828]
9  [0.626821,0.626822]  [29.8279,29.8279]

```

It is now easy to see that all scalar starting values of p_0 in the scalar EM algorithm will lead to the same point of convergence, and furthermore the number of iterations to convergence is not highly dependent on the starting value of p . The use of interval arithmetic has allowed the user to consider all possible values of the input parameter at once. This will not always be the case, but is a beneficial feature for the cases where it is possible.

Interval Global Optimization

Since interval analysis is scarcely known in the statistical literature, it will also be useful to mention another optimization method here. In the multinomial example, there is no

missing data and the loglikelihood in this case is given by

$$\log f(p|\mathbf{x}) \propto 125 * \log\left(\frac{2+p}{4}\right) + 38 * \log\left(\frac{1-p}{4}\right) + 34 * \log\left(\frac{p}{4}\right).$$

This can be viewed as an ordinary function to optimize, a task for which interval global optimization is well suited. [Hansen \(1992\)](#) is one of several monographs on this topic. Using the PROFIL software (or a similar package) with an initial interval of $p = [0 + \epsilon, 1 - \epsilon]$, a guaranteed enclosure of a stationary point is returned to the user.

4.6 Conclusions

Interval analysis first gained noticeable development in the 1960s from the work of R. E. Moore. Interval analysis has a fairly extensive literature in some areas, e.g. global optimization, but has seen little development in statistical settings. This paper takes a step at remediating the current state of knowledge by using interval analysis together with ideas from the EM algorithm. The resulting method is capable of finding multiple stationary points of a loglikelihood to a high degree of accuracy. The EM algorithm cannot be relied upon to do this. Unlike other algorithms for optimization, the method retains the ability of the EM algorithm to handle missing-data problems.

Chapter 5

GENERAL CONCLUSIONS

This dissertation has explored various ways in which interval analysis can be utilized in statistical computing. Interval analysis uses intervals instead of real numbers as the basic units of calculation. Interval analysis provides a means to evaluate the range of a function over a given domain. Using this capability, it is possible to obtain numerical answers which are guaranteed to be correct to a certain level of accuracy and to contain the true result. Interval analysis is also able to guarantee finding global optimizers within an initial box.

There are two broad topics covered in this dissertation as related to interval analysis. Several sections of the dissertation are focused on obtaining numerical values which are of a guaranteed accuracy. The application to statistics is in the computation of critical points and tail probabilities of several statistical distributions. For a bivariate chi-square distribution considered in one section, a series expansion is used together with a bound on the truncation error for the series. By evaluating the finite series with intervals and evaluating the truncation error with intervals, an interval enclosure of the true probability is obtained. To find critical points of a distribution, one approach requires solving for the root of an equation. Due to the complicated form of the equation involved, the use of derivatives (as in the Newton-Raphson algorithm) is avoided in favor of a derivative-free root-finding method. An interval secant algorithm (with Illinois modification) is developed and used for finding critical points of the distribution. A bivariate F distribution is also considered using similar techniques with the added complication that each term in the series involves the calculation of an interval Gaussian quadrature rule for evaluating a numerical integral. The methods prove successful for computing guaranteed enclosures of the probabilities to several decimal places (at a minimum). In some cases the guarantee extends to more than a dozen accurate digits. The guaranteed values allowed for the discovery of errors in earlier published tables.

A third section of the dissertation considers the global optimization capabilities of interval analysis. The EM algorithm is widely used in statistics for estimating parameters of a model when data is missing. The original EM paper by [Dempster et al. \(1977\)](#) has been cited in more than 2000 papers since its publication and continues to be an active area of research. Much of the research about the EM algorithm has explored ways to speed up the rate of convergence to a stationary point.

While enjoying enormous popularity, the EM algorithm, like many optimization methods, generally only converges to a stationary point (not necessarily the global optimum). Using special properties of the EM algorithm, an interval enclosure of the gradient of the loglikelihood is derived. The enclosure of the gradient is evaluated over an interval region. Regions where the enclosure of the gradient does not contain zero are therefore known to contain no stationary points. By beginning with an initial region and repeatedly subdividing the region into ever smaller pieces, regions which are known to not contain a stationary point are eliminated from consideration. Any stationary points of the loglikelihood are located in the union of the undiscarded regions. If the initial region is large enough, all stationary points of the loglikelihood will be found. No other known method is capable of achieving the same result.

Interval analysis has been demonstrated to be an effective tool in statistical computing. The unique opportunities and challenges associated with interval analysis promise both opportunities for future research and future rewards.

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