# BLSS - THE BERKELEY INTERACTIVE STATISTICAL SYSTEM: AN OVERVIEW 

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## 1. Introduction

BLSS (pronounced bliss) is a highly interactive statistics software system which runs on UNIX-based computers and workstations. It has grown out of the instructional needs of our department: We needed a single system which is both easy enough for students in freshman statistics courses (and first-year TA's) to learn and to use, but flexible and powerful enough for use in advanced undergraduate and beginning graduate classes. This paper is a brief overview of the software itself and its use in the Berkeley instructional program. For an extensive description of the software see Abrahams and Rizzardi [1988].

## 2. Instructional Philosophy

Why use statistical software in instruction? To some people the answer is obvious; but to some it may seem to be a passing fad. We see two main purposes, and they apply to all statistics courses at all levels - even our large introductory one-semester courses.
1). To help develop statistical intuition. Students should use real statistical techniques on real data - including large datasets (hundreds of observations). This develops their intuition about the art and science of analyzing data in a way which paper-and-pencil textbook problems cannot.
2). To help develop intuition for the probability and mathematics behind the statistics. Students should try out and experiment with statistical techniques, probability simulations, etc., both as a whole and - especially at the intermediate and advanced levels in terms of their individual mathematical building blocks. This is the antidote to thinking of the techniques as 'black boxes'.
Developing intuition is crucial to understanding.
Of course, students should be able do all this without getting bogged down in the details of computer use!

## 3. User Interface and Datasets

BLSS provides a simple, command-oriented user interface which also recognizes assignments and algebraic expressions. For example, the command:

```
- regress x y
```

forms the regression of $y$ on $x$, and the command:

- help regress
shows the on-line documentation for the regress command. The command:

```
. a = sqrt (b^2 + c^2)
```

assigns to $a$ the value of the expression $\sqrt{b^{2}+c^{2}}$, and the command:

```
. }s=t[4,5
```

assigns to $s$ the value of element $[4,5]$ of the matrix $t$. (The ' $\because$ ' in these examples is the prompt character which BLSS types when it is ready to accept commands.)

Datasets (the objects $a, b, c, s, t, x$, and $y$ in the preceding examples) are stored as familiar objects: they can be scalars, row vectors, column vectors, matrices, or three-way arrays. The work area can contain as many datasets as necessary, each with its own name, shape, and dimensions. The list command makes a list of datasets in the work area and describes their shape. For example, it might tell us:

| Contents of your work area: |  |  |
| :---: | :---: | :---: |
| a | dataset, dims= $(1,7)$ | (row vector) |
| b | dataset, dims= $(1,7)$ | (row vector) |
| c | dataset, dims= 11,7$)$ | (row vector) |
| ozone | dataset, dims=(366,13) | (matrix) |
| plover | dataset, dims=(42,5) | (matrix) |
| s | dataset, dims= $(1,1)$ | (scalar) |
| t | dataset, dims= $(10,10)$ | (matrix) |
| x | dataset, dims=(24,1) | (column vector) |
| Y | dataset, dims= 24,1$)$ | (column vector) |

In this example, and throughout this paper, we use slanted monospace font to show what the user types and regular monospace font to show what BLSS types.

## 4. Elementary Statistics

Elementary courses use BLSS for basic statistics commands such as descriptive statistics, stem-and-leaf diagrams, scatterplots, regression, etc. The following three commands, for example, fetch a dataset (of final examination scores) from the BLSS data library, show descriptive statistics for it, and make a stem-and-leaf diagram:

```
. load finalexam
Loaded "finalexam" from the BLSS system data area.
. stat finalexam
Statistics: finalexam
. stemleaf finalexam
N = 62, min = 6, 25% = 45, 50% = 60, 75% = 73, max = 92
Leaf digit unit (ldu) = 1 (1|2 represents 12.)
    0|69
    1|29
    ||9
    3|235699
    4|01455566
    5102245678889
    6|11234556789
    710113356789
    8।00223346
    9|12
```

For these simple univariate statistics, the command forms are no more complex than:

Subscripts can be used to specify columns, rows, and elements within a dataset. As a special case, single subscripts can be used to denote entire columns. For example, the command:

```
. scat plover[1] plover[2]
```

makes a scatterplot of the first column of the plover dataset (as the $X$ variable) against the second (as the $Y$ variable). Alternatively, options (which are enclosed in curly brackets \{\}) can be used to specify columns. Results of commands can be placed in output datasets. For example, the command:

```
. regress plover {x=1,2} {y=3} > y.hat resids coefs
```

forms the regression of column 3 of plover on columns 1 and 2 . The command displays the results:

| Dependent variable: |  | plover[3] |  |  |
| :---: | :---: | :---: | :---: | :---: |
| Independent variables: |  | plover[1 2] |  |  |
| Observations | 42 | Parameters | s 3 |  |
| Parameter | Estimate | SE | t-Ratio | P-Value |
| intercept | -14.264 | 0.91567 | -15.5779 | 0.0000 |
| coef 1 | 0.23824 | 0.017456 | 13.6480 | 0.0000 |
| coef 2 | 0.67424 | 0.043260 | 15.5859 | 0.0000 |
| Residual SD | 0.11289 | Residual V | Variance | 0.012744 |
| Multiple R | 0.97397 | Multiple R | R-squared | 0.94861 |

and it creates output datasets named y.hat, resids, and coefs which contain the fitted values, regression residuals, and estimated intercept and coefficients. (Of course, many other options and outputs are available for this command, but we do not discuss them here.)

The ' $>$ ' symbol is called the output dataset separator; everything to its right is an output dataset. Think of it as an arrow. The general form is:

$$
\text { command input-datasets } \rightarrow \text { output-datasets }
$$

In general, every number which is printed in an output display (such as in the regress display above) is also available as an output dataset should the user want it.

## 5. Elementary Probability and Sampling

BLSS provides several commands which illustrate concepts of probability and sampling in an elementary context. For example, students sometimes find the definition of a confidence interval confusing. The confid command illustrates the definition via simulation. Unless the information is provided (via input datasets and options on the command line), confid asks the student for the number and frequency of tickets to put in a box (using the vocabulary of Freedman, Pisani, and Purves [1978]), the number of draws per sample, and the number of samples:

```
. confid
Either: enter the numbers on the tickets in the box;
    Or: on separate lines, enter each ticket's number and frequency.
Finish with RETURN and CTRL-D.
5 4 8 6
(Control-D)
Box saved in: box00
Number of draws per sample? 100
Number of samples? 10
```

```
10 samples of 100 draws each are made from box00.
Population mean is 5. Population SD is 2.
```



BLSS provides several other such simulations. A coin-tossing simulation asks the student whether to use a fair coin or not, how many tosses to make, etc.; it keeps track of the cumulative number and proportion of heads, the difference from the expected value, etc. Another box-model simulation illustrates the sampling variation of the estimated average and standard deviation: it is similar to the confid simulation shown above, but instead displays summary statistics for each sample. (A separate sample command allows samples to be drawn - with or without replacement - from a real dataset.)

The demoiv command (named after De Moivre) provides an elementary illustration of the central limit theorem: it displays the scaled probability histogram for one observation from the box, and for the sum of $2,4,8,16, \ldots$ draws. Students find demoiv particularly entertaining - as they try (unsuccessfully) to design boxes which will defeat the central limit theorem.

## 6. Intermediate Use

BLSS provides many commands for specific statistical techniques, including: frequency counts, cross-tabulation, confidence intervals, hypothesis tests ( t -tests, z -tests, $\chi^{2}$-tests), probability functions, random number generation, matrix decompositions, time series analysis, etc.

However, its real power for instruction at intermediate and advanced levels comes from being able to assemble individual commands into larger sequences, combined with a number of useful matrix and array operators. We illustrate with a standard example from an intermediate level course - generating a random vector $\mathbf{x}$ from a multivariate Normal distribution with mean vector $\mu$ and covariance matrix $\Sigma$ using the Cholesky decomposition: $\mathbf{x}=\mathbf{z T}+\mu$, where $\mathbf{T}^{\prime} \mathbf{T}=\Sigma$ and $\mathbf{x}, \mathbf{z}$, and $\mu$ are row vectors, $\mathbf{z}$ of independent standard Normals. To create the vector $m u$, give the command:

$$
. m u=5,-2,0
$$

The comma operator ',' used here catenates the numbers together into a row vector (it can also catenate together column vectors and matrices which have the same number of rows); this is one way to create a new dataset. Another way to create a dataset is to read it in with the read command; we use this to enter the covariance matrix sigma:

```
. read > sigma
```

```
Type your data one row per line; finish with RETURN and CTRL-D.
    9 -6 3
-6 29 -2
    3 -2 5
(Control-D)
Read 9 values into "sigma"; dims=(3,3) (matrix).
```

(The read command automatically infers the dimensions of the new dataset from how it was typed in.) To compute the Cholesky square root of sigma and place it in $t$ we use the chol command:

```
. chol sigma > t
```

To show $t$, simply type its name:
. $t$

| 3.000 | -2.000 | 1.000 |
| ---: | ---: | ---: |
| 0.000 | 5.000 | 0.000 |
| 0.000 | 0.000 | 2.000 |

To check that the Cholesky decomposition is correct we can look at $\Sigma-T^{\prime} \mathbf{T}$ :

| • sigma $-t^{\prime \prime \# * t}$ |  |  |
| ---: | ---: | ---: |
| 0.000 | 0.000 | 0.000 |
| 0.000 | 0.000 | 0.000 |
| 0.000 | 0.000 | 0.000 |

(The ' operator is matrix transposition, and the \#* operator is matrix multiplication.)
Instead of generating a single random vector, we generate 100 simultaneously. We start by using the rgau command ('r' for random and 'gau' for Gaussian) to generate a matrix $z$ of dimensions $(100,3)$ containing random standard Normals:

```
. rgau {dims=100,3} > z
```

To obtain our 100 random 3-vectors as a matrix $x$ :

```
. x = z#*t + mu
```

Note that the addition operation here added a 100-by-3 matrix to a length 3-row vector, and yielded a 100-by-3 matrix. The row vector $m u$ was conceptually 'expanded' into a matrix of the appropriate dimension; that is, it set $x_{i j}=[\mathbf{z T}]_{i j}+\mu_{j}$ for all $i$ and $j$. BLSS provides this 'dimension-expansion' feature for all elementwise binary operators. It is useful for many array operations, such as centering and scaling columns or rows of a matrix, etc.

Because intermediate results can be examined and variations can be experimented with, assembling sequences of commands to perform more elaborate operations is an excellent way for students to build intuition.

## 7. Convenience Features

BLSS provides a number of convenience features which make life easier.
Aliases allow you to create abbreviations for commands and command-and-option combinations. As a trivial example, if the word 'regress' is too long to type, the command:

```
. alias reg regress
```

creates an alias name 'reg' for it. Thereafter, whenever you type the command reg, it is just as if you had typed the command regress; the new command reg can accept any of the inputs, options, or outputs of regress. More usefully, aliases can include inputs and options. For example, the stat command (which shows descriptive statistics) normally computes variances and standard deviations using the divide-by-( $N-1$ ) formula. The option $\{\mathbf{d n}=0\}$ makes it use the divide-by- $N$ formula instead. To always use this option, give the command:

```
. alias stat stat {dn=0}
```

(This alias value contains two words, 'stat' and ' $\{\mathrm{dn}=0\}$ '. In general, alias values can contain any number of words.) Most BLSS commands provide options which change their default behavior or output displays. Thus, aliases can be used to customize BLSS commands and displays to one's own taste.

Strings are similar to aliases, except that they allow for abbreviations to be placed in the middle of the command line as well as the beginning of it. Thus, they can provide abbreviations for long sets of options, long dataset names, etc.

BLSS keeps a record of commands typed in the current session; this is known as the command history. Commands from the history can be displayed, repeated, and edited (for example, to change an option or parameter value without re-typing the entire command).
Command files (or macros) can be written which combine existing commands into new commands; the new commands are invoked in the same way (using the same syntax, etc.) as existing commands. A convenient way to construct command files is to edit the record of commands in your history after you have given some set of commands that you want to repeat.
Aliases, strings, and command files - as well as datasets, on-line help files, and a startup message - can be installed on a class-wide basis. Thus, the instructor can provide a customized environment for all students in the class. For example, if the instructor is using a textbook which defines variances and standard deviations using the divide-by- $N$ formula, he can install the alias shown above, so that the students automatically get the same results as in the textbook.

## 8. Software Interfacing

BLSS provides software interfaces to the operating system (UNIX) at several levels.
For common tasks - such as re-routing text output displays from the terminal screen to a disk file, sending files to a lineprinter (or laserprinter), and editing files (using the editor of one's choice) - BLSS has builtin interfaces to the appropriate UNIX software; to perform these tasks it is unnecessary to leave BLSS and enter UNIX. (This also means that students need not learn any UNIX commands; they can 'live' within BLSS.)

In addition, BLSS and UNIX can be interfaced at the command level: UNIX commands can be called directly from within BLSS, and vice versa. This allows for rapid integration of BLSS with other application software.
Finally, BLSS can be interfaced at the subroutine level - with UNIX routines, or with
the user's own subroutines. In fact, BLSS commands are nothing other than UNIX commands (written in either C or Fortran) which call the BLSS subroutine library. The result is that - although we have been emphasizing its instructional uses - BLSS provides a ready-made frontend for running one's own Fortran (or C) subroutines.

Here is a simple example of a complete Fortran program which does so. It takes one input dataset and creates one output dataset.

```
c Very simple Fortran calling program
    to a user-supplied Fortran subroutine.
One input dataset (x) and one output dataset (y).
Minimal BLSS library support:
    No option handling. No dynamic memory allocation.
    real x(10000), y(10000)
    ... Open the input dataset ...
    ix = iopeni("x", "-ERR-")
    ... Find out its number of rows and columns ...
    nr = inr(ix)
    nc = inc(ix)
    ... Check dataset size against allocated memory ...
        if (nr*nc .gt. 10000)
+ call error("Dimensions cannot exceed nr*nc > 10000.")
... Open the output dataset ...
iy = iopeno("y", "-ERR-", 1, nr, nc)
... Read the input data matrix ...
call mread(ix, x, nr, nc, "", 0)
... Check for no NA's (missing values) ...
    call mchk (ix, x, nr, nc, "-ERR-")
    ... User-supplied subroutine to obtain y from x ...
    call mysubr(x, y, nr, nc)
    ... Write the output data matrix ...
    call mwrite(iy, y, nr, nc)
    ... Clean up and exit ...
    call iexit(0)
    end
    ... This could be any user-supplied subroutine ...
    ... For the purpose of illustration, we supply a trivial one ...
    subroutine mysubr(x, y, nr, nc)
    real x(nr,nc), y(nr,nc)
    do 10 j = 1, nc
        do 10 i = 1, nr
                        y(i,j) = -x(i,j)
    continue
    return
    end
```

The input and output datasets are called $x$ and $y$ inside the program, but can have any name as far as the user is concerned. The functions iopeni and iopeno take care of opening the input and output datasets and discovering their names from the command line; the subroutines mread and mwrite take care of reading and writing data one matrix-worth at a time. Other subroutines and functions (inr, inc, mchk) perform utility tasks such as returning the number of rows and columns in the input dataset, checking to see whether it contains NA's (missing values), etc. The BLSS library provides many more capabilities such as: checking for options, performing dynamic memory allocation for Fortran, a variety of error-condition handling, etc. Thus the calling program can be as simple or as complex as desired, in terms of options, defaults, and so forth.
The calling program can be invoked directly from UNIX as well as from within BLSS this is helpful if you are in the process of writing your subroutine and want to make a lot of use of the editor, the compiler, and the debugger.

## 9. Current Status

At U.C. Berkeley, BLSS is used in statistics courses at all levels, from freshman through graduate, as well as in quantitative courses taught by other departments. The following table summarizes its use in courses at Berkeley in the 1987-88 academic year.

|  | Number of <br> One-Semester <br> Courses | Total <br> Enrollment |
| :--- | :---: | :---: |
| Statistics Department | 4 | 912 |
| Freshman/Sophomore level | 13 | 342 |
| Junior/Senior level | 3 | 134 |
| $\quad$ Graduate level |  |  |
| Other Departments | 2 | 44 |
| Anthropology | 3 | 164 |
| Chemistry | 1 | 61 |
| Electrical Engineering | 1 | 5 |
| Forestry | 27 | 1662 |

## 10. Future Directions

BLSS is under continuing development by the Department; the design and programming effort involves several people. For the immediate future, we are concentrating on:

- Improved graphics. Our plan is to add high-quality, flexible, command-driven graphics capabilities which work with a variety of output devices including Tektronixcompatible graphics terminals, X-windows, and laserprinters, as well as character devices such as lineprinters and regular terminals.
- A full-screen, menu-driven user interface is being written: Some of our instructors believe that menus are preferable to commands for use in elementary level classes.
- More statistics capabilities. Obviously we cannot add everything; so, in order to help set priorities, we made a widespread survey of universities earlier this year. Three new capabilities currently being worked on are: elementary (rank-based) nonparametric methods, analysis of variance for factorial designs, and ARMA and ARIMA routines.


## 11. Why UNIX?

Why did we choose UNIX as the operating system on which to run BLSS? By limiting ourselves to a single operating system, we can concentrate on developing better software rather than on satisfying the idiosyncrasies of many different operating systems; and from our standpoint, UNIX is the most important operating system for the foreseeable future. It is the only vendor-independent operating system which runs on a wide variety of hardware (from desktop machines to mainframes and even supercomputers), and its networking and resource-sharing capabilities far exceed those of any other we know of. Our department has found that UNIX provides an excellent environment for both research
and teaching. In particular, running classes is much easier on UNIX than on PC's, because UNIX eliminates the need for multiple copies of software, datasets, help files, and other resources on each PC, and because it provides easy communication between instructor, teaching assistants, and students.

Three years ago, our department considered switching over from UNIX to a network of PC's or Macintoshes for instruction - but we decided against it based on the strengths of UNIX.

## 12. Summary

BLSS meets a wide range of instructional needs, from elementary courses for freshman nonmajors through beginning courses for graduate students.
In our view, the purpose of BLSS (or the computer, in general) in statistical instruction is to help students develop intuition: for both statistics itself and for the probability and mathematics behind the statistics.

BLSS provides a command-oriented user interface which also recognizes assignments and algebraic expressions.

BLSS stores datasets intuitively (as scalars, row vectors, column vectors, matrices, etc.). The work area can contain as many datasets as needed, each with its own name and shape.

For elementary courses, BLSS provides standard statistics commands and also elementary probability demonstrations and simulations.

For intermediate and advanced courses, BLSS provides many specific statistical techniques - but its real power comes from being able to assemble sequences of commands and low-level matrix and array operations to accomplish more elaborate tasks.

Convenience features allow the user to customize the BLSS environment to his own liking. Course instructors can apply such customizations to an entire class.

The lineprinter, the editor, and UNIX command-level capabilities can be used directly from within BLSS. Commands can be written which interface BLSS and UNIX at the subroutine level.

BLSS enjoys widespread use at U.C. Berkeley. Over 1600 students used it here last year in courses at all levels, both within the Statistics Department and outside it.

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