# PBS Modelling 0.60: User's Guide

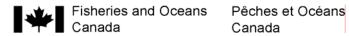
(Draft Report)

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2006

# **Canadian Technical Report of Fisheries and Aquatic Sciences xxxx**



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# Canadian Technical Report of Fisheries and Aquatic Sciences

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Fisheries and Aquatic Sciences xxxx

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by

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#### **ABSTRACT**

Schnute, J.T., Couture-Beil, A., and Haigh, R. 2006. PBS Modelling 1: user's guide v.0.60 (draft version). Can. Tech. Rep. Fish. Aquat. Sci. xxxx: xxx + xxx p.

This draft report describes the R package *PBS Modelling*, which contains software to facilitate the design, testing, and operation of computer models. The initials *PBS* refer to the Pacific Biological Station, a major fisheries laboratory on Canada's Pacific coast in Nanaimo, British Columbia. Initially designed for fisheries scientists, this package has broad potential application in many scientific fields. *PBS Modelling* focuses particularly on tools that make it easy to construct and edit a customized graphical user interface (GUI) appropriate for a particular problem. Although our package depends heavily on the R interface to Tcl/Tk, a user does not need to know Tcl/Tk. In addition to GUI design tools, *PBS Modelling* provides utilities to support data exchange among model components, conduct specialized statistical analyses, and produce graphs useful in fisheries modelling and data analysis. Examples implement classical ideas from fishery literature, as well as our own published papers. The examples also provide templates for designing customized analyses using other R libraries, such as *PBS Mapping*, *odesolve*, and *BRugs*. Users interested in building new packages can use *PBS Modelling* and a simpler enclosed package *PBS Try* as prototypes. An appendix describes this process completely, including the use of C code for efficient calculation.

#### **RÉSUMÉ**

Schnute, J.T., Couture-Beil, A. et Haigh, R. 2006. PBS Modelling 1: Guide de l'utilisateur v.0.60 (version provisoire). Can. Tech. Rep. Fish. Aquat. Sci. xxx: xxx + xxx p.

Google translation: Ce projet de rapport décrit le paquet PBS de R modelant, qui contient le logiciel pour faciliter la conception, l'essai, et l'opération des modèles d'ordinateur. Les initiales PBS se rapportent à la station biologique Pacifique, un laboratoire important de pêche sur la côte Pacifique du Canada dans Nanaimo, Colombie britannique. Au commencement conçu pour des scientifiques de pêche, ce paquet a la large application potentielle dans beaucoup de domaines scientifiques. PBS modelant des foyers en particulier sur les outils qui le rendent facile à construire et éditer une interface utilisateur graphique adaptée aux besoins du client (GUI) s'approprient pour un problème particulier. Bien que notre paquet dépende fortement de l'interface de R à Tcl/Tk, un utilisateur n'a pas besoin de savoir Tcl/Tk. En plus des outils de conception de GUI, modeler de PBS fournit des utilités à l'échange d'informations supplémentaires parmi les composants modèles, les analyses statistiques spécialisées par conduite, et les graphiques de produit utiles dans la pêche modelant et l'analyse de données. Les exemples mettent en application des idées classiques de la littérature de pêche, aussi bien que nos propres papiers édités. Les exemples fournissent également des calibres pour concevoir des analyses adaptées aux besoins du client en utilisant d'autres bibliothèques de R, telles que tracer de PBS, odesolve, et BRugs. Les utilisateurs intéressés à de nouveaux paquets de bâtiment peuvent employer modeler de PBS et un essai inclus plus simple du paquet PBS comme prototypes. Une annexe décrit ce processus complètement, y compris l'utilisation du code de C pour le calcul efficace..

#### **Preface**

After working with fishery models for more than 30 years, I've used a great variety of computer software and hardware. Currently, the free distribution of R (R Development Core Team 2006a) provides an excellent platform for software development. Furthermore, the associated network of contributed libraries on CRAN (Comprehensive R Archive Network, CRAN: <a href="http://cran.r-project.org/">http://cran.r-project.org/</a>) gives access to a wealth of algorithms from many users in various fields. This disciplined system allows users, like the authors of this package, to distribute software that extends the utility of R in new directions.

At various times, I've used software in Basic, Fortran (Mittertreiner and Schnute 1985), Pascal, C, and C++ to implement ideas in published papers. Usually this software goes stale in time, due to minimal documentation, changing operating systems, the lack of portable libraries, and many other factors. Because R includes a rich library of statistical software that operates on multiple platforms, my colleagues and I can now distribute software that actually works when other people try it. The user community includes us, because we often find that we can't remember how to operate our own software after a few weeks or months, let alone years. Although writing a good R package requires considerable effort, the result often pays off in portability, communication, and long term usage.

*PBS Modelling* tries to accomplish several goals. First, it anticipates the need for model exploration with a graphical user interface, a so-called GUI (pronounced gooey). We make this easy by encapsulating key features of the Tcl/Tk library into convenient tools fully documented here. A user need not learn Tcl/Tk to use this package. Everything required appears in Appendix B. You might want to start by running the function testWidgets(). Co-author Rowan Haigh likes the subtitle: "modelling the world with gooey substances."

Second, we want to demonstrate interesting analyses related to our work in fishery management and other fields. The function runExamples() illustrates some of these, as described further in Section 5. The code for all of them appears in the R library directory PBSmodelling\Examples. We demonstrate the power of other R libraries, such as BRugs (to perform Bayesian posterior sample with the application WinBUGS), odesolve (to solve differential equations numerically), and PBSmapping (to draw maps and perform spatial analyses).

Third, *PBS Modelling* serves as a prototype for building a new R package, as summarized in Appendix A. We illustrate two methods of calling C code (.C and .Call), and discuss many other technical issues encountered while building this library.

Finally, to use R effectively, we've found it convenient to devise a number of "helper" functions that facilitate data exchange, graphics, function minimization, and other analyses. We include these here for the benefit of our users, who may choose to ignore them. We hope that *PBS Modelling* inspires interest in interactive models that demonstrate applications in many fields. Enjoy!

Jon Schnute, August 2006

#### 1. Introduction

This draft report describes software to facilitate the design, testing, and operation of computer models. The package *PBS Modelling* is distributed as a freely available library for the popular statistical program R (R Development Core Team 2006a). The initials *PBS* refer to the Pacific Biological Station, a major fisheries laboratory on Canada's Pacific coast in Nanaimo, British Columbia. Previously, we produced the R library *PBS Mapping* (Schnute et al. 2004), which draws maps and performs various spatial operations. Although both packages (which can run separately or together) include examples relevant to fishery models and data analysis, they have broad potential application in many scientific fields.

Computer models allow us to speculate about reality, based on mathematical assumptions and available data. The full implications of a model usually require numerous runs with varying parameter values, data sets, and hypotheses. A customized graphical user interface (or GUI, often pronounced "gooey") facilitates this exploratory process. *PBS Modelling* focuses particularly on tools that make it easy to construct and edit a GUI appropriate for a particular problem. Some users may wish to use this package only for that purpose. Other users may want to explore the examples included, which demonstrate applications of likelihood inference, Bayesian analysis, differential equations, computational geometry, and other modern technologies. In constructing these examples, we take advantage of the diversity of algorithms available in other R libraries.

In addition to GUI design tools, *PBS Modelling* provides utilities to support data exchange among model components, conduct specialized statistical analyses, and produce graphs useful in fisheries modelling and data analysis. Examples implement classical ideas from fishery literature, as well as our own published papers. The examples also provide templates for designing customized analyses using the R libraries discussed here. In part, *PBS Modelling* provides a (very incomplete) guide to the variety of analyses possible with the R framework. We anticipate many revisions of our library, as we find time to include more examples.

*PBS Modelling* depends heavily on Peter Dalgaard's (2001, 2002) R interface to the Tcl/Tk package (Ousterhout 1994). This combines a scripting language (Tcl) with an associated GUI toolkit (Tk). In our library, we simplify GUI design with the aid of a "window description file" that specifies the layout of all GUI components and their relationship with variables in R. We support only a subset of the possibilities available in Tcl/Tk, but we customize them in ways intended specifically for model design and exploration (Appendix B). A user of *PBS Modelling* does not need to know Tcl/Tk.

Computer models typically involve a variety of components, such as code, data, a user interface, and documentation. Figure 1 illustrates the tangled relationships that sometimes accompany computer model design. *PBS Modelling* allows the GUI to become a device for organizing components, as well as running and testing software (Figure 2). The project might involve other applications, as well as R itself.

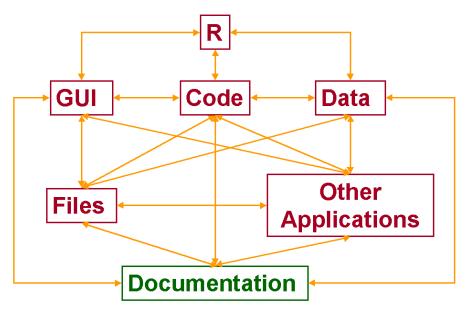


Figure 1. Tangled relationships among computer model components.

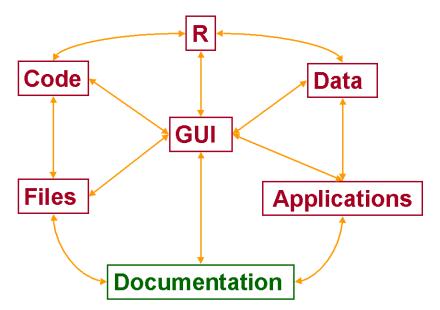


Figure 2. Computer model components organized with a graphical user interface (GUI).

In *PBS Modelling*, project design normally begins with a text file that describes the GUI. Additional files may contain code for R and other applications, which sometimes require code written in languages other than R. For example, the R *BRugs* library (to perform Bayesian inference using Gibbs sampling) requires a file with the intended statistical model, written in the language of a separate program *WinBUGS*. In other contexts, a user might write C code to get acceptable performance from model components that require extensive computer calculations. This code might be compiled as a separate program or linked directly into a customized R package.

Section 2 of this report describes the process of designing a GUI to operate a computer model. Components can share data through text files in a specialized "PBS format" presented in Section 3. These correspond naturally to list objects within R. Section 4 describes additional tools for customized graphics and data analysis. In Section 5, we highlight briefly some of the examples in our initial release, although we expect the list to expand in future versions.

Appendix A describes the process of building *PBS Modelling* in a Windows environment. A simple enclosed package *PBS Try* gives a prototype for building any R package, including the use of C code to speed calculations. Appendix B gives the complete syntax for all visual components (called *widgets*) available for a writing a window description file to specify a customized GUI. Appendix C shows the help files included with the library.

To use *PBS Modelling*, run R and install the package from the R GUI (click "Packages", "Install package(s)..., select a mirror, and choose PBSmodelling from the list of packages). Windows users can also obtain an appropriate zip file from the authors of this report or directly from the CRAN web site <a href="http://cran.r-project.org/">http://cran.r-project.org/</a>.

## 2. GUI tools for model exploration

The practical task of writing appropriate code for the R Tcl/Tk package can sometimes become a bit daunting, particularly if the GUI window requires extensive design and change. For a restricted set of Tk components (called *widgets*), *PBS Modelling* makes it much easier to design and use GUIs for exploring models in R. A user needs to supply two key parts of a GUI-driven analysis:

- a window description file (an ordinary text file) that completely specifies the desired layout of widgets and their relationship with functions and variables in R, and
- R code that defines relevant functions, variables, and data.

This section begins with an example to illustrate the main ideas, and then gives complete details for constructing window description files that can be used to generate GUIs.

#### 2.1. Example: Lissajous curves

A Lissajous curve (<a href="http://mathworld.wolfram.com/LissajousCurve.html">http://mathworld.wolfram.com/LissajousCurve.html</a>), named after one of its inventors Jules-Antoine Lissajous, represents the dynamics of the system

$$x = \sin(2\pi mt), \quad y = \sin[2\pi (nt + \phi)],\tag{1}$$

where time t varies from 0 to 1. During this time interval, the variables x and y go through m and n sinusoidal oscillations, respectively. The constant  $\phi$ , which lies between 0 and 1, represents a cycle fraction of phase shift in y relative to x. We want to design a GUI that allows us to explore this model by plotting Lissajous curves (y vs. x) for various choices of the parameters  $(m, n, \phi)$ . We also want to vary the number of time steps k and choose a plot that is either lines or points.

**Table 1.** Two text files associated with the "Lissajous Curve" project. The first gives a description of the GUI window used to manage the graphics. The second contains R code to draw a Lissajous curve.

# File 1: LissajousCurve.txt window title="Lissajous Curve"

```
vector length=4 names="m n phi k"
    labels="'x cycles' 'y cycles' 'y phase' points" \
    values="2 3 0 1000"
radio name=ptype text=lines value="l" mode=character
radio name=ptype text=points value="p" mode=character
button text=Plot function=drawLiss
```

#### File 2: LissajousCurve.r

```
drawLiss <- function() {
  getWinVal(scope="L");
  tt <- 2*pi*(0:k)/k;
  x <- sin(2*pi*m*tt); y <- sin(2*pi*(n*tt+phi));
  plot(x,y,type=ptype);
  invisible(NULL); }</pre>
```

This analysis can be accomplished with the R code and window description file shown in Table 1. Assume that these two files reside in the current working directory and that *PBS Modelling* has been installed in R. Start an R session from this directory, and type the following three lines of code in the R command window:

```
> require(PBSmodelling)
> source("LissajousCurve.r")
> createWin("LissajousCurve.txt")
```

The first line assures that *PBS Modelling* is loaded, the second defines the function drawLiss for drawing Lissajous curves, and the third creates a window that can be used to draw curves corresponding to any choice of parameters. Figure 3 shows the resulting GUI window interface. When the <Plot> button is clicked, the curve in Figure 4 appears in the R graphics window. This corresponds to the default parameter values:

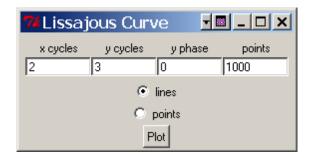
$$m = 2, n = 3, \phi = 0, k = 1000.$$
 (2)

The GUI allows different Lissajous figures to be drawn easily. Simply change parameter values in the four entry boxes, and click on <Plot>.

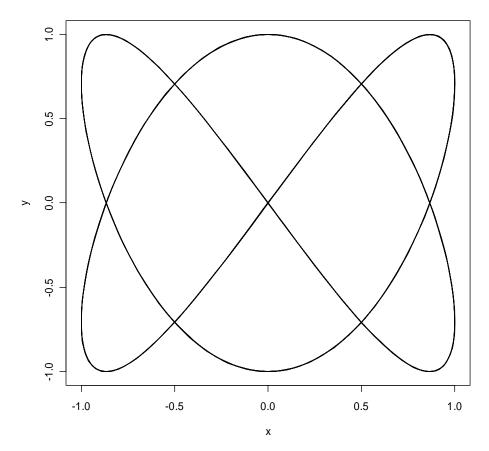
The description file (Table 1) specifies a window titled "Lissajous Curve" with a vector of four entries. These correspond to quantities with the R variable names m, n, phi, and k. The corresponding window (Figure 3) will contain four entry boxes that allow these quantities to be changed. A label for each quantity emphasizes its conceptual role: the number of cycles for x or y, the phase shift for y, and the number of points plotted. Initial values correspond to those listed in (2). The backslash ( $\setminus$ ) character indicates that a widget description (in this case,

a vector) continues on the next line. A pair of radio buttons, both corresponding to an R variable named ptype, allow selection between "lines" and "points" when drawing the plot. The graph is actually drawn (i.e., the R function drawLiss is called) when the user presses a button that contains the text "Plot". In general, we use the symbols <...> to designate a button or keystroke, such as the <Plot> button or the radio buttons <lines> and <points>.

The file of R code (Table 1) implements the algorithm (1) for computing k points on a Lissajous curve. The function drawLiss has no arguments, but gets values of the R variables m, n, phi, k, and ptype from the GUI window via a call to the *PBS Modelling* function getWinVal. The argument scope="L" implies that these variables have local scope within this function only. (Another choice scope="G" would give the variables global scope.)



**Figure 3.** GUI generated by the description file LissajousCurve.txt in Table 1. It contains five widgets: the window titled "Lissajous Curve", a vector of four entries, two linked radio buttons (and <points>), and a <Plot> button.



**Figure 4.** Default graph for the "Lissajous Curve" project, obtained by clicking the <Plot> button in Figure 3. The *x* variable goes through two cycles while the *y* variable goes through 3 cycles. A line graph is drawn through 1000 points generated by the algorithm (1).

#### 2.2. Window description file

A window description file currently supports the following 17 widgets:

- 1. window an entire new window;
- 2. menu a menu grouping;
- 3. menuitem an item in a menu;
- 4. grid a rectangular block of widgets;
- 5. label a text label;
- 6. button a button linked to an R function that runs a particular analysis and generates a desired output, perhaps including graphics;
- 7. check a check box used to turn a variable off or on, with corresponding values 0 or 1;
- 8. radio one of a set of mutually exclusive radio buttons for making a particular choice;
- 9. null –a blank widget that can occupy an empty space in a grid;
- 10. entry a field in which a scalar variable (number or string) can be altered;
- 11. text an entry box that supports multiple lines of text;
- 12. vector an aligned set of entry fields for all components of a vector;

- 13. matrix an aligned set of entry fields for all components of a matrix;
- 14. data an aligned set of entry fields for all components of a data frame, where columns can have different modes;
- 15. slide a slide bar that sets the value of a variable;
- 16. slideplus an extended slide bar that also displays a minimum, maximum, and current value:
- 17. history a device for archiving parameter values corresponding to different model choices, so that a "slide show" of interesting choices can be preserved.

The description file is an ordinary text file that specifies each widget on a separate line. However, any one widget description can span multiple lines by using a backslash character (\) to indicate the end of an incomplete line. For example, the single line:

```
label text="Hello World!"
is equivalent to:
label \
  text="Hello World!"
```

Meaningful indentation is highly recommended, but not compulsory. The three-line description of a vector widget in Table 1 illustrates a readable style.

Each widget has named arguments that control its behaviour, analogous to the named arguments of a function in R. Some (required) arguments must be specified in the widget description. Others (not required) can take default values. All widgets have a type argument equal to one of the 17 names above, although the word type can be omitted in the description file. Appendix A gives an alphabetic list of all these widgets, along with detailed descriptions of all arguments. As in calls to R functions, argument names can be omitted as long as they conform to the order specified in the detailed widget descriptions given below. Nevertheless, we recommend that all argument names be specified, except possibly the name type, which is always the first argument for each widget. Unlike R functions, where commas separate arguments, the arguments in a widget description are separated by white space.

In a description file, all argument values are treated initially as strings. In addition to specifying a line break, the backslash can be used to indicate five special characters: single quote \', double quote \", tab \t, newline \n, and backslash \\. If an argument value does not include spaces or special characters, then quotes around the string are not required. Otherwise, double quotes must be used to delineate the value of an argument. Single quotes indicate strings nested within strings. For example, the vector in Table 1 has four labels specified by the string argument

```
labels="'x cycles' 'y cycles' 'y phase' points"
```

A hash mark (#) that is not within a string begins a comment, where everything on a line after the hash mark is ignored. As mentioned above, an isolated backslash (not part of a special character) indicates continuation onto the next line. A break can even occur in the middle of a string, such as the long label

```
label text="This long label with spaces \
  spans two lines in the description file"
```

In this case, leading spaces in the second line are ignored, to allow meaningful formatting in the description file. Intentional spaces in a long string should appear prior to the backslash on the first line.

Although the type argument (like vector) for a widget can never be abbreviated, other arguments follow the convention used with named arguments in R function calls. For a given widget type, the available arguments can be abbreviated, as long as the abbreviations uniquely identify each argument. For example, the vector in Table 1 could be specified as:

Unlike variable names in R, widget names and their arguments are not case sensitive. Some users may prefer to write all type variables in upper case or with an initial capital letter. For example, the names WINDOW, VECTOR, RADIO, and BUTTON could be used to emphasize the widgets in Table 1.

#### 2.3. Window support functions

PBS Modelling includes functions designed to connect R code with GUI windows. Every window has a name argument (with default name=window), and windows with different names can coexist. When running a program with multiple windows, only one window will be current (i.e., selected by the user) at any particular time. The function createWin uses a description file to generate one or more windows, where each window has a name (perhaps the default) taken from the file. If a window with the specified name already exists, it will be closed before the new window is opened. When designing and testing a GUI, this feature ensures that a new version automatically replaces the previous one. The function closeWin, which takes a vector of window names, closes all windows named in the vector. With no arguments, closes all windows that are currently open.

Although createWin normally builds a GUI from a description file, it will also accept a vector of strings equivalent to such a file. Thus, a file of R source code can define a GUI directly, without the need for a separate description file. Table 2 illustrates how this can be done in a simple case. To see the character vectors equivalent to a given description file (say, winDesc.txt), type the R command:

```
scan("winDesc.txt", what=character(), sep="\n")
```

In particular, if the description file includes a backslash or double quote character, the corresponding R string must represent it as  $\ \ '$ , respectively. Despite this alternative of embedding window descriptions in R source files, we recommend writing separate files to define GUIs, except perhaps for very simple models.

**Table 2.** A simple file of R source code with character strings that define a GUI. No separate window description file is required.

#### File: Simple.r

```
# window description strings
winStr=c(
    "window",
    "entry name=n value=5",
    "button function=myPlot text=\"Plot sinusoid\"");

# function to plot a sinusoid
myPlot <- function() {
    getWinVal(scope="L");
    x <- seq(0,500)*2*n*pi/500;
    plot(x,sin(x),type="l"); };

# commands to create the window
require(PBSmodelling); createWin(winStr,astext=TRUE)</pre>
```

Internally, *PBS Modelling* uses a list object in the process of generating a GUI from a description file. The functions compileDescription and parseWinFile give lists that correspond to description files. Just as createWin can act directly on a character vector, it can also act on a suitably defined list, rather than a file. This feature makes it possible to replace a description file with R code that defines the corresponding list, although we recommend against this practice in most cases.

R programs need to share data with a GUI window. *PBS Modelling* provides three functions that deal with values of R variables named in a description file:

- getWinVal returns values from the current window;
- setWinVal sets values in the current window;
- clearWinVal clears global values associated with the current window.

Some models associate a particular action with a single parameter vector. In such cases the function <code>createVector</code> generates a GUI directly, without the need for a corresponding description file.

Several functions support file display and manipulation from a GUI:

- openFile opens a file using the default program for the file extension;
- promptOpenFile shows the current directory for choosing a file to open;
- promptSaveFile shows the current directory for naming a file to save.

*PBS Modelling* includes a history widget designed to collect interesting choices of GUI variables so that they can be redisplayed later, rather like a slide show. This widget has

buttons to add and remove GUI settings from the current collection, to scroll backward and forward, and to clear all entries from the collection. Other buttons allow entire history files to be saved or loaded.

Normally, a user would invoke a history widget simply by including a reference to it in the description file. However, *PBS Modelling* includes some support functions for customized applications:

- initPBShistory initializes data structures for holding a collection of history data;
- addPBShistory saves the current window settings to the current history record;
- rmPBShistory removes the current record from the history;
- backPBShistory and forwPBShistory move backward and forward among the history records;
- jumpPBShistory moves to a specified record in the history;
- exportPBShistory and importPBShistory save and load histories from files;
- clearPBShistory removes all records from the current collection.

The help file for initPBShistory shows an example that uses these functions directly.

#### 2.4. Internal data for windows

PBS Modelling uses two list variables PBS.win and PBS.options in the global environment to store information relevant to its current settings. In particular, PBS.win has four components that contain data about the current GUI window, where

- \$vars is a vector with the names of all R variables declared in the description file;
- \$funs is a vector with the names of all R functions declared in the description file;
- \$actions is a vector with action names (optionally declared in the description file) that indicate recent actions taken by the user in the current GUI;
- \$windowname is the name of the currently active window.

The functions showVars, showFuns, showActions, and showWin can also be used to return these character vectors. If multiple windows are present, PBS. win automatically gets updated to the data for the window currently selected.

After using createWin to produce a GUI, the vectors PBS.win\$vars and PBS.win\$funs provide useful summaries of names declared in the current project. Furthermore, the vector PBS.win\$actions provides a record of GUI actions taken by the user, starting with the most recent and working backwards. By default, the action associated with a widget is its type; for example a button has default action=button. In general, however, the description file could give a unique action name to each potential action, so that the vector would give an unambiguous record of user actions.

If a widget invokes the function openFile, the associated action should be the file name. By definition, openFile has the default argument PBS.win\$actions[1].

Currently, PBS.options acts primarily to store default program names associated with file extensions. On a Windows platform, the native R function shell.exec (called by

openFile) automatically chooses a default from the registry. For this reason, our distribution specifies an empty list:

```
PBS.options$openFile=list().
```

The default can, however, be overwritten by specifying explicit list components, such as: html='"c:/Program Files/Mozilla Firefox/firefox.exe" %f'

where %f denotes the file name in the string passed to the operating system. On Unix platforms, it may be essential to specify default this way. Future versions of our library may include other options, such as default width for a data entry field or the maximum number of actions.

## 3. Functions for data exchange

Computer models usually require data exchange between model components. For example, as described above, the functions <code>getWinVal</code> and <code>setWinVal</code> move data between an R program and the GUI. Other applications, such as those written separately in C, may have the ability to write data to files that R can read. In cases like this, it would be convenient to have variable names in the C code correspond to variables with the same names in R. <code>PBS Modelling</code> can facilitate this process with the functions <code>readList</code> and <code>writeList</code>, which convert a text file to an R <code>list</code> and <code>vice-versa</code>. Another function <code>unpackList</code> creates local or global variables with names that match the list components.

Table 3 illustrates a data file in PBS format, legible by readList. The file contains lines with an initial dollar sign (like \$x in Table 3) that specify a list component name in R, followed by one or more lines of data. Data items are separated by white space. A single item of data corresponds to a scalar in R, multiple items on a single line correspond to a vector, and multiple lines of data correspond to a matrix with the number of columns determined by the first line of data. Thus, in Table 3, \$x is a scalar, \$y is a vector of length 4, and \$z is a 2×4 matrix. The format also supports four possible data type definitions on a line preceded by \$\$:

- \$\$ vector mode=numeric names=""
- \$\$ matrix mode=numeric ncol colnames="" byrow=TRUE
- \$\$ data modes=numeric ncol colnames byrow=TRUE
- \$\$ array mode=numeric dim fromright=TRUE

Table 3 illustrates their use in specifying \$a, \$b1, and \$b2. Matrices and data frames can be read by row or column. This choice determines the order of reading the data, and white space (including line breaks) merely signifies breaks between data items. Array objects with three or more dimensions can be read in two ways, with indices varying first from the right or from the left. For example, data for an array indexed by [i,j,k] are read by varying i first with fixed j and k if fromright=TRUE. Similarly, k varies first if fromright=FALSE.

**Table 3.** Sample data file for *PBS Modelling*. The function readList converts this file to a list object with six components: a scalar \$x, a logical vector \$y, two matrices (\$z,\$a), and two data frames (\$b1,\$b2). The matrix \$a is read by column, and \$b1=\$b2.

```
$x
0
$у
T F TRUE FALSE
ŜΖ
11.1 12.2 13.3 14.4
15.5 16.6 17.7 1.88e+01
$a
$$matrix ncol=2 byrow=FALSE colnames="a b"
5 1 2 3
$b1
$$data ncol=3 modes="numeric logical character" \
  byrow=TRUE colnames="N L C"
5 T aa
3 F bb
8 T cc
10.5 F dd
$b2
$$data ncol=3 modes="numeric logical character" \
 byrow=FALSE colnames="a b c"
5 3 8 10.5
TFTF
aa bb cc dd
```

As in widget descriptions, arguments may be omitted in favour of their defaults, and the \$\$ line may be continued across multiple lines by using a backslash character \. For a matrix, the argument ncol is required. Similarly, a data object (i.e., a data frame) must specify ncol and a vector colnames of length ncol. Also, modes must have length 1 (so that all entries in the data frame have the same mode) or length ncol. An array must have a complete dim argument, a vector giving the number of dimensions for each index.

As indicated earlier, *PBS Modelling* can use this specialized data format as a convenient means of capturing data from other programs. For example, to export data from an external C program, write C code that generates a data file in PBS format, where component names in the file match the C variable names. Then read the resulting file into an R session with the function readList, and use unpackList to produce local or global R variables. At this point, both R and C share data with the same variable names.

To considerable extent, R has native support for reading and writing a variety of text files, including the functions scan, cat, source, dump, dget, dput, read, write, read.table, and write.table. External programs sometimes utilize R formats for their input data. For example, the program WinBUGS (Speigelhalter et al., 2004), which implements Bayesian inference using Gibbs sampling, uses data files written in a list format closely related to the R syntax produced by the dput function. If the file myData.txt has this format, then either of the two R commands

```
myData <- dget("myData.txt");
myData <- eval(parse("myData.txt"));</pre>
```

produces a corresponding R list object named myData.

We should, however, add a word of caution here. When R saves array data in dput format, it converts the array to a vector by varying the indices from left to right. For example, a matrix with indices [i,j] is saved as a vector in which i varies for each fixed j. In effect, the data are stored by column. This sometimes gives an unnatural visual appearance. In English, the eye reads naturally from left to right, then down. Matrices are normally displayed by row, with column index j varying for each fixed i. WinBUGS, supported by the R package BRugs (Thomas 2004), requires input data formatted in this visually meaningful way. More generally, WinBUGS reads arrays by varying the indices from right to left. The BRugs function bugsData writes data in this format, but users must take special care in reading WinBUGS data with the dget function.

# 4. Support functions for graphics and analysis

As mentioned in the preface, we have devised a number of functions that make it easier for us to work in R. Some of them, such as plotBubbles, relate to techniques discussed in our published work (e.g., Richards et al. 1997; Schnute and Haigh 2006). Others just provide convenient utilities. For example, testCol("red") shows all colours in the palette colors() that contain the string "red".

#### 4.1. Graphics utilities

resetGraph	Resets various graphics parameters to defaults, with mfrow=c(1,1)
expandGraph	Sets various graphics parameters to make graphs fill out available space
drawBars	Draw a linear bar plot on the current graph
plotCsum	Plots cumulative sum of a vector, with value added

#### 4.2. Data management

clearAll .......Function to clear all data in the global environment

pad0 .......Pads numbers with leading zeroes (string)

show0 ......Shows decimal places including zeroes (string)

view ......Views first n rows of a data frame or matrix

comparePars......Find the difference between two par vectors \*\*\*\*\* not implemented compareLists......Find the difference between two lists

#### 4.3. Function minimization and maximum likelihood

\*\*\*\*\* These concepts have not been implemented in *PBS Modelling 0.60*.

#### New data type:

parVec - a data frame with columns val, min, max, active (logical), and possible row names:

scalePar scales parameters to [0,1] restorePar get actual parameters from scaled values

calcMin(pvec, func, tol) calculates the minimum of func, starting at pvec

#### 4.4. Handy utilities

calcGM	Calculates the geometric mean of a vector of numbers
findPat	Find all strings that include any string in a vector of patterns
pause	Pause, typically between graphics displays
showArgs	Show the arguments for a specified widget in Appendix B
testWidgets	GUI to test all widgets listed in Appendix B

view......View of first few lines of a (potentially large) matrix or data frame

## 5. Examples

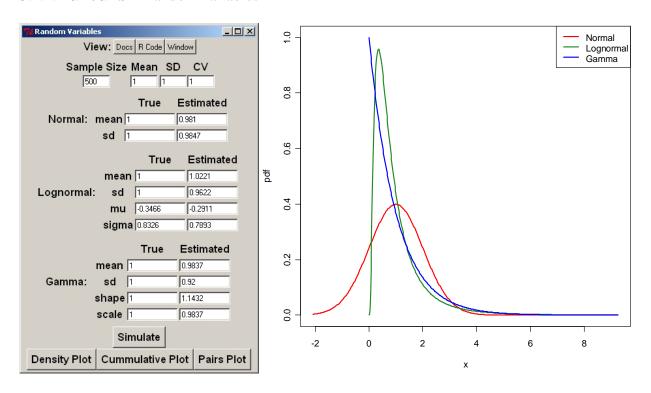
As mentioned in the Preface, *PBS Modelling* includes a variety of examples that illustrate applications based on this and other libraries. Generally, each example comprises R-code, a window description file, documentation, and other supporting files. All relevant code appears in the R library directory PBSmodelling\Examples, where example xxx typically has corresponding files xxx.r, xxxWin.txt, and xxxDoc.txt or xxxDoc.pdf. The function runExamples() brings up a window that runs the examples in a temporary directory located on the path defined by the environment variable Temp.

Alternatively, you can copy all the files from PBSmodelling\Examples to a directory of your choice and open R in this working directory. To run example xxx, type source("xxx.r") on the R command line. For instance, source("LissFig.r") creates Lissajous figures (described earlier) and includes the history widget for collecting settings that the user wishes to retain. Sourcing LissFig.r invokes the windows description file LissFigWin.txt, which produces the GUI.

These examples work correctly only if a user's computer has been set to associate ".txt" and "r" with suitable text editors. Similarly, the Acrobat Reader must be installed, so that "\*.pdf" is associated with that program. In the descriptions below, we often refer to GUI elements in quotation marks. For example, "Model" usually refers to the button labelled "Model". In some cases, this becomes shorthand for "Press the button labelled Model".

#### 5.1. Random variables

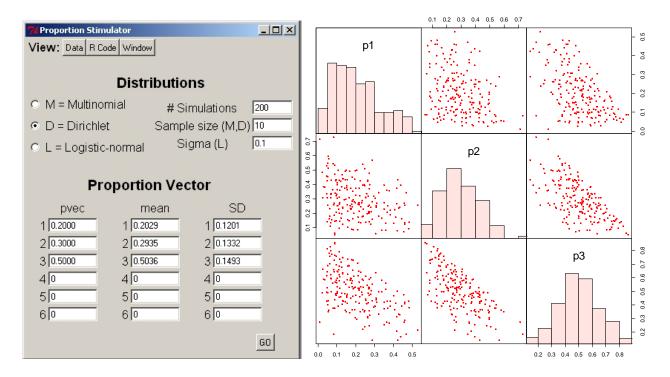
#### 5.1.1. RanVars – Random variables



**Figure 5.** RanVars GUI (left) and density plot (right). Simulations are based on 500 random draws with mean =1 and SD=1.

The RanVars example draws samples from three continuous random distributions (normal, lognormal, and gamma) with a common mean  $\mu$  and standard deviation  $\sigma$ . The documentation ("Docs" button) shows relevant formulas that connect distribution parameters with the moments  $\mu$  and  $\sigma$  Estimated parameter values from a simulation (invoked by "Simulate") are displayed in the GUI alongside the true values (Figure 5). We use only the straightforward formulas in the documentation, without bias correction formulas like those described by Aitchison and Brown (1969). Three buttons at the bottom of the GUI portray the data visually as density curves, cumulative proportions, and paired scatter plots.

#### 5.1.2. RanProp – Random proportions

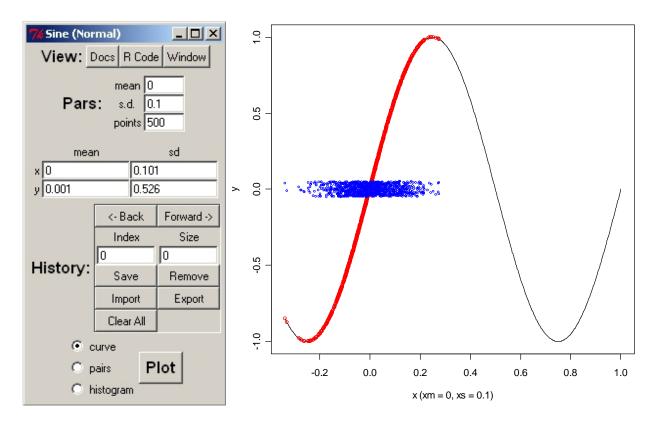


**Figure 6.** RanProp GUI (left) and pairs plot (right). Simulations are based on 200 random draws where n = 10 for the multinomial and Dirichlet distributions and  $\sigma = 0.1$  for the logistic-normal distribution. The pairs plot portrays results for the Dirichlet.

The RanProp example simulates up to five random proportions drawn from one of three distributions – multinomial, Dirichlet, and logistic-normal. The observed proportion means and standard deviations are reported in the GUI (Figure 6), and a graphical display renders the points as a paired scatter plot. After defining options in the GUI, including the vector "pvec" of true underlying proportions, press "Go". Schnute and Haigh (2006) provide further technical details about these three distributions.

\*\*\*\* This example still needs a documentation file.

#### 5.1.3. SineNorm - Sine normal

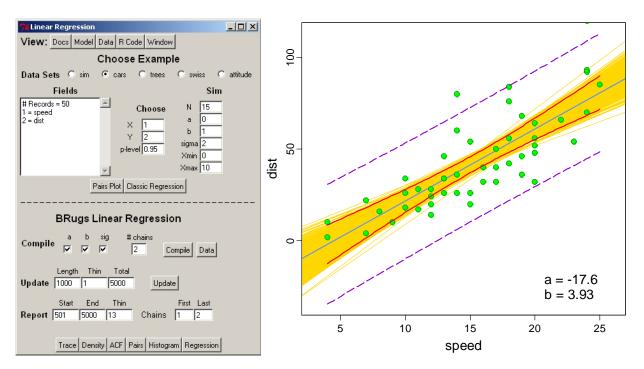


**Figure 7.** SineNorm GUI (left) and plot (right). Simulations are based on 500 random draws of  $y = \sin(2\pi x)$ , where x is normal with mean  $\mu = 0$  and standard deviation  $\sigma = 0.1$ . Blue points portray jittered values of x, and red points show corresponding values of y.

The SineNorm example illustrates a somewhat unconventional random variable  $y = \sin(2\pi x)$ , where x is normal. The GUI allows you to specify the mean  $\mu$  and standard deviation  $\sigma$  of x. If  $\mu = 0$  and  $\sigma$  is small, the transformation is nearly linear, so that y is approximately normal. If  $\sigma$  is large, the transformation concentrates y near -1 and 1. Figure 7 illustrates the transformation when  $\sigma$  has the moderate value 0.1. Try  $\sigma = 10$  to see how values y tend to occur near the peaks and troughs of the sine function, where the slope is relatively flat.

#### 5.2. Statistical analyses

## 5.2.1. LinReg – Linear regression

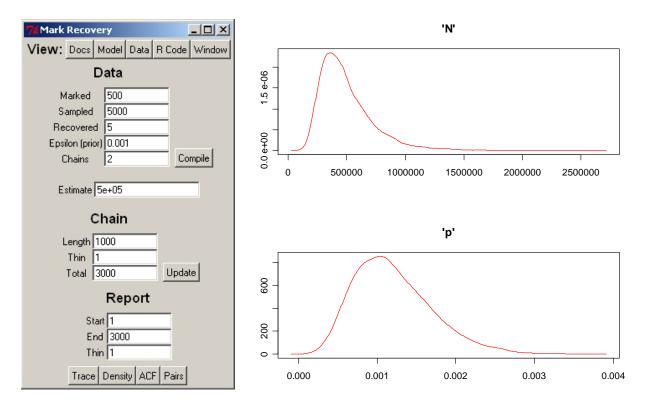


**Figure 8.** LinReg GUI (left) and regression plot (right). The linear regression uses the cars dataset (n=50) to predict dist vs. speed. The plot shows observations (green circles), fitted line (solid blue line), the 95% confidence limits of the fitted model (solid red lines), the 95% CL of the data (dashed purple lines), and the fits using the Bayes posterior estimates of (a,b) (gold lines).

The example LinReg estimate parameters in a linear regression y = a + bx using either simulated data or data objects that come with the R-package. We compare classical frequentist regression with results from Bayesian analysis, using the BRugs library to interface with the program WinBUGS. After selecting various data options, "Pairs Plot" shows a pairs plot (x, y) and "Classic Regression" adds confidence limits (at "p-level") from regression theory. Red and violet curves show bounds for a prediction or a new observation, respectively, each conditional on x. If the data came from simulation, a blue line portrays the truth, with specified values a and b, that must be estimated from the data.

A corresponding Bayesian analysis uses the WinBUGS model shown by pressing "Model". Choose parameters to monitor (normally all of them): the intercept a, the slope b, and the predictive standard deviation  $\sigma$ . After specifying a number of sample chains for the MCMC sample, press "Compile" to compile the model with these settings. "Update" generates samples in "Length" increments. Additional buttons at the bottom of the GUI allow you to explore the MCMC output. Posterior samples of (a,b) correspond to sample lines. The "Regression" button illustrates these in relationship to confidence limits from a frequentist analysis (Figure 8).

#### 5.2.2. MarkRec - Mark-recovery



**Figure 9.** MarkRec GUI (left) and density plots (right). A low recovery of marked fish can lead to fat tails in *N* due to occasional large spikes in the population estimate.

The example MarkRec performs a Bayesian analysis of a mark-recovery experiment in which M fish are marked and allowed to disperse randomly in the population. Later, a sample of size S is removed from the population and R marks are recovered. Both the total population N and the marked proportion P are unknown, where

$$p = \frac{M}{N} \cong \frac{R}{S}.$$

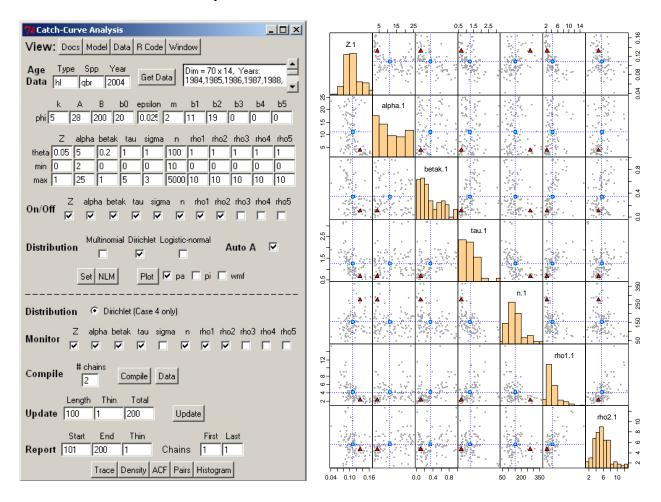
In one version of the theory, R is binomially distributed with probability p in a sample of size S, and the above approximation suggests the estimate

$$\hat{N} = \frac{S}{R}M = \frac{M}{R}S.$$

When recoveries are low ( $R \approx 0$ ), the posterior distribution of N exhibits a fat tail (Figure 9).

As in LinReg, "Model" shows the MarkRec model for WinBUGS, which (deliberately) includes an illegitimate prior that depends on the data. By increasing an initially small quantity  $\varepsilon$ , this fake prior allows the tail of N values to be arbitrarily clipped. Schnute (2006) gives some historical perspective to this analysis, in the context of work by W.E. Ricker.

#### 5.2.3. CCA – Catch-curve analysis



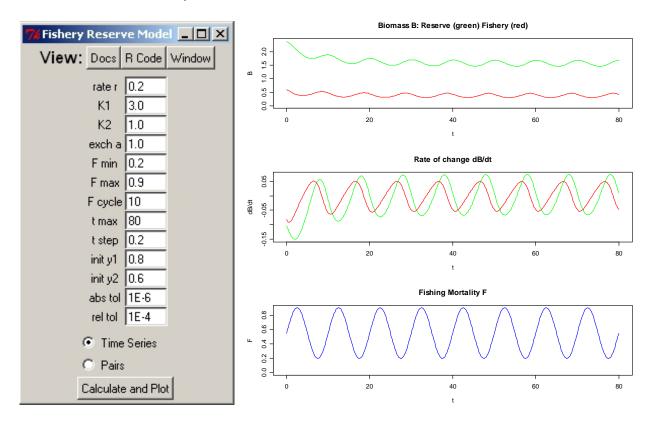
**Figure 10.** CCA GUI (left) and parameter pairs plot (right). Comparison of Bayes posterior distribution of CCA model parameter estimates from chain 1 (*N*=100). Symbols indicate means (blue squares) and modes (red triangles). Diagonal shows parameter estimate distributions.

The example CCA illustrates a catch-curve model proposed by Schnute and Haigh (2006). It incorporates effects of survival, selectivity, and recruitment anomalies on age structure data from a single year. After making various model choices, press "Set", "NLM" (which may take several seconds), and "Plot" to view the maximum likelihood estimates and their relationship with the data. A WinBUGS model ("Model") allows us to calculate posterior distributions. (See the last few lines of "Model".) As in MarkRec, select parameters to monitor, specify a number of chains, and "Compile" the model. "Update"s may be slow, but eventually they produce interesting posterior samples (Figure 10). "Docs" gives details of the deterministic model, and the Dirichlet distribution is used to describe error in the observed proportion.

We include this example to illustrate a somewhat realistic WinBUGS model that can be used to estimate parameters for a population dynamics model. We will provide further information when the paper (Schnute and Haigh 2006) is published. *PBS Modelling* includes the data for this example as the matrix CCA.qbr.hl.

#### 5.3. Other applications

## 5.3.1. FishRes – Fishery reserve



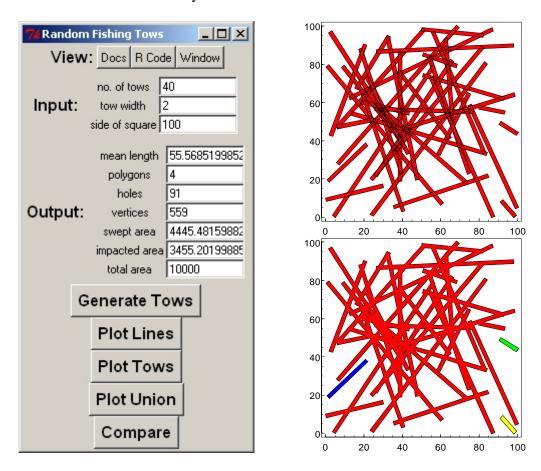
**Figure 11.** FishRes GUI (left) and time series (right) of population biomass, rates of biomass change, and fishing mortality.

The example FishRes models a fish population associated with a marine reserve using differential equations, which are solved numerically with the odesolve library. The dynamic equations:

$$\begin{split} \frac{dB_{1}}{dt} &= rB_{1} \left( 1 - \frac{B_{1}}{K_{1}} \right) + a \left( \frac{B_{2}}{K_{2}} - \frac{B_{1}}{K_{1}} \right) \\ \frac{dB_{2}}{dt} &= rB_{2} \left( 1 - \frac{B_{2}}{K_{2}} \right) - a \left( \frac{B_{2}}{K_{2}} - \frac{B_{1}}{K_{1}} \right) - F(t)B_{2} \\ F(t) &= F_{\min} + \frac{F_{\max} - F_{\min}}{2} \left( 1 + \sin \frac{2\pi t}{n} \right) \end{split}$$

describe the biomass  $B_i$  in region i=1,2, where region 1 is a reserve and region 2 experiences a periodic fishing mortality rate F, with minimum and maximum values  $F_{\min}$  and  $F_{\max}$ . The two regions have a common growth rate F, but different carrying capacities F, a parameter F determines the movement rate from the region of higher density to the other region (Figure 11).

#### 5.3.2. FishTows – Fishery tows



**Figure 12.** FishTows GUI (left) and simulated tow track (right). Tow track plots show 40 random tows in a square with side length 100. Each tow has width 2, and the rectangle encompasses 10,000 square units. *Top*: The individual rectangles, with 160 vertices, have areas that sum to 4,445 square units. *Bottom*: The union includes a complex polygon (red) and three isolated rectangles (blue, green, yellow) that cover only 3,455 square units. The complex polygon (red) has 547 vertices and 91 holes.

The example FishTows provides a simulator of fishery tow tracks using the PBSmapping library. The example demonstrates the difference between swept area and area impacted by trawls that often cover the same ground repeatedly. This application can be regarded an exotic random number generator, where tows initially join two points picked from a uniform random distribution within a square of a given side length. Three parameters (the number of tows, the tow width, the side length) determine several random variables, including the mean tow length, the areas swept and impacted, the numbers of polygons and holes in the union set of tows, and the number of vertices in the union. Each of these would also have a variance and an overall distribution generated by many runs of this example.

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# Appendix A. Building PBSmodelling and other packages

The R project defines a standard for creating a package of functions, data, and documentation. You can obtain a comprehensive guide to "Writing R Extensions" (R Development Core Team 2006b, R-exts.pdf) from the CRAN web site or the R GUI (see the References above). Ligges (2003) and Ligges and Murdoch (2005) provide useful introductions. We have designed PBSmodelling and a very simple enclosed package PBStry as prototypes for package development. This Appendix summarizes the steps needed to:

- A.1. install the required software;
- A.2. build PBS Modelling from source materials;
- A.3. write source materials for a new package and compile them;
- A.4. include C code in a package.

Our discussion applies only to package development on a computer running Microsoft Windows 2000, XP, or (maybe) later. We particularly highlight issues that have proved troublesome for us. The R library directory PBSmodelling\PBStools contains batch files that can assist the process. For example, you might locate this directory as C:\Utils\R\R-2.3.1\library\PBSmodelling\PBStools.

#### A.1. Installing required software

Building R packages requires six pieces of free software. Even if some of this software is already installed, it may be helpful (or even essential) to update to the latest versions. We recommend installing each package on a path that does *not* include spaces. For example, avoid using C:\Program Files, even if that happens to be part of a package's default path. In this appendix, we use C:\Utils as a root directory for all required software. The list below shows versions available at the time of writing this report, along with suggested paths.

- 1. **R** itself, currently version 2.3.1 (C:\Utils\R\R-2.3.1). We assume that R is already installed from the CRAN web site <a href="http://cran.r-project.org/">http://cran.r-project.org/</a> and that it runs correctly on your computer. We also assume that the package PBSmodelling is installed in R.
- 2. A set of **UNIX tools** for building R packages in Windows (no version number shown, C:\Utils\Rtools). Obtain the file tools.zip from the web site <a href="http://www.murdochsutherland.com/Rtools/">http://www.murdochsutherland.com/Rtools/</a>. Uncompress this file in C:\Utils, and (if you wish) rename the base directory from \tools to \Rtools. These tools are *essential*. DO NOT plan to use

- programs with the same name in an installation of Cygwin or any other UNIX emulator that happens to be installed on your computer.
- 3. **Perl**, currently version 5.8.8.817 (C:\Utils\Perl). Obtain this from the Active State web site: <a href="http://www.activestate.com/Products/ActivePerl/Download.html">http://www.activestate.com/Products/ActivePerl/Download.html</a>.
- 4. **MinGW**, currently version 5.0.3 (C:\Utils\MinGW), the minimalist GNU C compiler for Windows. Obtain a small installation file from <a href="http://www.mingw.org/">http://www.mingw.org/</a>, and use this as a package manager to download and install at least the basic components.
  - Alternatively, you may wish to obtain **Dev-C++**, currently version 4.9.9.2 (C:\Utils\DevCpp). Obtain a complete installation file from <a href="http://www.bloodshed.net/">http://www.bloodshed.net/</a>. This package includes a distribution of MinGW and a convenient integrated development environment (IDE) for compiling and testing C programs. Based on our current experience, Dev-C++ works as an adequate substitute for MinGW in building R packages.
- 5. **MiKTeX**, currently version 2.5 (C:\Utils\MiKTeX), available from <a href="http://www.miktex.org/">http://www.miktex.org/</a>. This processor for TeX and LaTeX files helps typeset help files within a package. If you don't have version 2.5 or later, take this opportunity to upgrade. Download the "basic" installation file, and install these components only. You can add more LaTeX packages from the Internet later, as required. (MiKTeX often does this automatically.) Take some time to investigate the MiKTeX package manager (mpm.exe or go to the "Programs" menu and select "MiKTeX 2.5", "Browse Packages").
  - The text editor **WinEdt** (available from <a href="http://www.winedt.com/">http://www.winedt.com/</a>) provides a convenient GUI for editing LaTeX files and operating MiKTeX. Combined with the R package RWinEdt, it can also serve as an editor and interface for R. However, it is available only as shareware that requires a fee for long-term use, unlike any other software mentioned here.
- 6. The Microsoft **HTML Help Workshop** (no version shown, C:\Utils\HHW). You need the installation file HtmlHelp.exe, currently at <a href="http://www.microsoft.com/office/ork/xp/appndx/appa06.htm">http://www.microsoft.com/office/ork/xp/appndx/appa06.htm</a>. If Microsoft no longer makes it available there, you can obtain the file from <a href="http://www.murdoch-sutherland.com/Rtools/">http://www.murdoch-sutherland.com/Rtools/</a>. After installation, we think you can safely ignore a message that "This computer already has a newer version of HTML Help". (If anyone has different information, please let us know.)

After these six pieces of software are installed, you're ready to start building R packages. For this purpose, create a new directory that can contain your packages; such as D:\Rdevel\. Within the R library directory, which could be C:\Utils\R\R-2.3.1\library\, find the subdirectory PBSmodelling\PBStools. Copy all the batch files there into your new packages directory. You should have these ten files:

- definePaths.bat, checkPaths.bat related to the installation;
- checkPBS.bat, buildPBS.bat, packPBS.bat, unpackPBS.bat related to PBS Modelling;
- check.bat, build.bat, pack.bat, unpack.bat related to the construction of new packages.

You need to change definePaths.bat so that it reflects the paths you chose in the above six installations. For example, your version of this batch file might contain the lines

```
set R_PATH=C:\Utils\R\R-2.3.1\bin
set TOOLS_PATH=C:\Utils\Rtools\bin
set PERL_PATH=C:\Utils\Perl\bin
set MINGW_PATH=C:\Utils\MinGW\bin
set TEX_PATH=C:\Utils\MiKTeX\miktex\bin
set HTMLHELP_PATH=C:\Utils\HHW
```

Notice that each path, except the last, ends in a bin subdirectory.

Hopefully, your installation is now complete. In you new packages directory, run checkPaths.bat from a command line or double-click the icon. This script verifies that a few essential files lie on the indicated paths. If everything is correct, you should see the message "All program paths look good". Otherwise, you'll see a warning about software that doesn't appear on your specified paths.

You may wish to inspect all the batch files with a text editor. They don't use your system PATH environment variable. Instead, each one defines a new local path appropriate for building R packages (via checkPaths.bat). A SETLOCAL command ensures that this change doesn't alter your system's permanent environment.

#### A.2. Building PBSmodelling

Once all the required software is installed, the batch files discussed above make it fairly easy to build PBSmodelling. We assume that you have already created the directory discussed in Appendix A.1, say D:\Rdevel, for building R packages and that it contains the relevant eight batch files. In particular, definePaths.bat should reflect your installation paths and checkPaths.bat should report the message that "All program paths look good". Then follow these steps:

- 1. On the CRAN web site <a href="http://cran.r-project.org/">http://cran.r-project.org/</a>, go to "Packages" on the left and find PBSmodelling. Download the file PBSmodelling\_x.xx.tar.gz into D:\Rdevel. Then rename this file (or copy it and rename the copy) so that the version number is removed. You should now have the file PBSmodelling.tar.gz in D:\Rdevel.
- 2. In the development directory D:\Rdevel, double-click the icon for unpackPBS.bat or type the command unpackPBS in a corresponding command window. This should extract the contents of PBSmodelling.tar.gz, preserving directory structure, into a subdirectory \PBSmodelling with five sudirectories: \data, \inst, \man, \R, and \src.

Our batch file uses the command tar -xzvf PBSmodelling.tar.gz, where tar.exe appears in the \Rtools directory (section A.1, step 2). The command line parameters specify a verbose (v) extraction (x) of the given file (f), after filtering with gzip (z).

If you use other software for this extraction, please ensure that it is configured to handle UNIX files correctly. For example, "WinZip" has an option to extract a "TAR file with smart CR/LF conversion". This must be turned off.

3. In the base directory D:\Rdevel, double-click the icon for checkPBS.bat or type the command checkPBS in a corresponding command window. If all software is installed correctly and D:\Rdevel\PBSmodelling correctly represents the contents of the .tar.gz file, you should see a series of DOS messages reporting "OK" to various tests. A distinct pause might accompany the message: "checking whether package 'PBSmodelling' can be installed ...".

You might also encounter a delay as MiKTeX downloads the LaTeX package lmodern, part of a larger package lm. If this is really slow, you can abort the process and install lm with the MiKTeX package manager, as discussed in step 5 of section A.1. Choose a remote server near you. You only need to do this once. When it's finished, run checkPBS.bat again.

- 4. Examine the new directory E:\Rdevel\PBSmodelling.Rcheck created by the check process in step 2. The text files 00check.log and 00install.out show detailed results.
- 5. In the base directory D:\Rdevel, double-click the icon for buildPBS.bat or type the command buildPBS in a corresponding command window. This creates the file D:\Rdevel\PBSmodelling.zip, which could be used to install PBSmodelling from a local zip file.
- 6. Again in the base directory D:\Rdevel, double-click the icon for packagePBS.bat or type the command packagePBS in a corresponding command window. This creates a new package distribution file PBSmodelling\_x.xx.tar.gz that replaces the one downloaded from CRAN in step 1.

If these steps all work without problems, you can be pretty sure that the requisite software is installed correctly and that you understand the basic steps needed to build R packages.

#### A.3. Creating a new R package

R packages require a special directory structure. The R function package.skeleton automatically creates this structure, but (without further work) it does not produce a package that can be compiled. Although PBSmodelling has the requisite structure, it is perhaps too complicated to serve as a convenient prototype. For this reason, we include a small subset PBStry that illustrates the key details. You can make a new package simply by editing the files in PBStry. You need a suitable editor (e.g., WinEdt or the Notepad) to view and change various text files.

1. Start by locating the file PBStry\_x.xx.tar.gz in the R library directory \PBSmodelling\PBStools. Copy this file into your development directory, such as D:\Rdevel, and rename it (or copy and rename the copy) to obtain the file PBStry.tar.gz.

2. Follow steps similar to those in section A.2 to unpack, check, build, and re-package PBStry. You must now use a DOS command window in D:\Rdevel to issue the four commands

unpack PBStry check PBStry build PBStry pack PBStry

which invoke the batch files unpack.bat, check.bat, build.bat, and package.bat. The first command should give you a new subdirectory \PBStry, along with its five sudirectories: \data, \inst, \man, \R, and \src.

3. Use your editor to open the file DESCRIPTION in the root directory \PBStry. This file, essential in every R package, contains key information in a special format (RDCT 2006b, section 1.1.1). The following example illustrates a minimal set of required fields.

Package: MyPack Version: 1.00 Date: 2006-08-31 Title: My R Package

Author: User of PBS Modelling Maintainer: User of PBS Modelling

Depends: R (>= 2.3.1)

Description: My customized R functions

License: GPL version 2 or newer (recommended)

The package name in DESCRIPTION must agree with the directory name in which this file lies. For example, if you change PBStry to MyPack in DESCRIPTION and rename the directory from \PBStry to \MyPack, you have effectively changed the package name. Similarly, if you change the version to 1.01, you have effectively changed the version number that appears in the file names for distributing your package.

4. The subdirectory \PBStry\R contains all R code used by the package. For example, PBStry includes six R functions (calcFib, calcGM, calcSum, findPat, pause, and We view). The six files could be combined into a single file (such as PBStry.R), but we use separate files here for clarity. The functions all have relatively simple code, hopefully comprehensible to users with limited R experience. Five of them come from PBSmodelling. Two of them (calcFib, calcSum) call compiled C code, as we discuss more completely in section A.4 below.

By convention, the distinct file zzz.R defines code for initializing the package. In this case the function .First.lib, calls library.dynam to load a dynamic link library (PBStry.dll) created from compiled C code during the build process.

When a version number changes, the DESCRIPTION file must be changed accordingly. We also like to make a corresponding change in zzz.R, so that the version number appears on the R console when the library is loaded. PBStry illustrates this possibility for zzz.R.

5. The subdirectory \PBStry\data contains all data objects that come with the package. Here, the binary file QBR.rda holds a matrix of quillback rockfish (*Sebastes maliger*)

sample data used in the CCA example above (section 5.2.3). The same data matrix is called CCA.gbr.hl in PBSmodelling.

If you want to add data to a new package, first create the object (e.g., myData) in R and then execute the command:

```
save(myData,file="myData.rda")
```

The object name must match the prefix in the file name, and the suffix must be .rda. Include the resulting file in your package's \data subdirectory.

- 6. The subdirectory \PBStry\man contains a documentation file for every object in the package. PBStry has six functions and one data set, so the \man subdirectory has seven corresponding R documentation files (\*.Rd). An additional file PBStry.Rd documents the package as a whole. Rd files use a rather complex scripting language (RDCT 2006b, section 2) that can be converted to help files in several formats (PDF, HTML, text). For many packages, the examples in PBStry may provide adequate prototypes. They represent three distinct cases: functions (e.g., calcGM.Rd, findPat.Rd), data sets (QBR.Rd), and complete packages (PBStry.Rd).
- 7. The subdirectory \PBStry\src contains source code for C code to be compiled into the dynamic link library PBStry.dll. We include sample files to calculate Fibonacci numbers iteratively (fib.c) and to add the components of a numeric vector (sum.c). In section A.4, we discuss the linkage between R code and compiled C functions.
- 8. Finally, the subdirectory \PBStry\inst contains files that are to be included directly in the R library tree for PBStry when the package is installed. The file PBStry-Info.txt briefly describes the context and purpose of the trial package.

If you have successfully followed the steps above, you have actually built two R packages, PBSmodelling and PBStry. Furthermore, you're reasonably familiar with the contents of PBStry. You can use the files in that small package as prototypes for writing your own R package, which might contain R code in the subdirectory \R. data in \data, C source code in \src, and R documentation in \man.

The larger package PBSmodelling offers more prototypes and uses a somewhat different style. The main directory includes the required DESCRIPTION file, plus a second file NAMESPACE that lists all objects available to a user of the package. Effectively, the namespace mechanism distinguishes between objects provided by the package and other (hidden) objects required for the implementation, but not intended for public use. Our NAMESPACE file contains the rather cryptic instruction: exportPattern("^[^\\.]"). The R string "^[^\\.]" translates to the regular expression ^[^\.] that designates any pattern not starting with a period (.). We don't export "dot" objects, whose names in R start with a period. (For more complete information, see the file PBSMfunctions.txt in the subdirectory \inst or the R library directory for PBSmodelling.) The namespace file must also import functions required from other packages. Because PBSmodelling relies heavily on tcltk, the file includes the command: import(tcltk).

In PBStry, without a namespace, the file zzz.R defines the initializing function .First.lib, as mentioned in step 4 above. By contrast, the namespace protocol in PBSmodelling requires a different name for the initializing function: .onLoad in zzz.R.

In summary, we recommend building a new package by editing, adding, and deleting prototype files in PBStry. Our batch files can facilitate tests and debugging. For more advanced work, particularly packages with a namespace protocol, look at PBSmodelling. Have a current version of RDCT (2006b) available, and consult that manual when necessary. We find it useful to keep the PDF file open and to use Acrobat's search feature (Ctrl-F) to find topics of interest.

### A.4. Embedding C code

R provides two functions, .C() and .Call(), for invoking compiled C code. PBStry includes two simple examples that use .C(), probably the method of choice for simple packages. The .Call() function uses a more complex interface that offers better support for R objects. PBSmodelling includes a simple example to illustrate this calling convention.

# Calling C functions from R using .C()

**Table A1.** C representations of R data types.

R Object	С Туре
logical	int *
integer	int *
double	double *
complex	Rcomplex * 1
character	char **

<sup>1</sup> Rcomplex is defined in Complex.h.

The .C() calling convention uses the following key concepts:

- R must allocate the appropriate length and type of variables before calling a C function.
- R objects are transformed into an equivalent C type (Table A.1), and a pointer to the value is passed into the C function. All values are returned by modifying the original values passed in.
- A C function called by .C() must have return type void, because values are returned only by accessing the predefined R function arguments.
- C code written for the shared DLL must not contain a main function.
- Within a C function, dynamically allocated memory must be de-allocated by the programmer before the function returns. Otherwise a memory leak will likely occur.
- .C() returns a list similar to the '...' list of arguments passed in, but reflecting any changes made by the C code. (See the help file for .C)

**Table A2.** Two text files associated with a .C() call in PBStry. R code in the first file calls C code in the second.

```
File 1: calcFib.R
calcFib <- function(n, len=1) {</pre>
  if (n<0) return(NA);
  if (len>n) len <- n;
  retArr <- numeric(len);</pre>
  out <- .C("fibonacci", as.integer(n), as.integer(len),</pre>
             as.numeric(retArr), PACKAGE="PBStry")
  x \leftarrow out[[3]]
  return(x) }
                         File 2: fib.c
void fibonacci(int *n, int *len, double *retArr) {
  double xa=0, xb=1, xn=-1; int i, j;
  /* iterative loop */
  for(i=0;i<=*n;i++) {
    /* initial conditions: fib(0)=0, fib(1)=1 */
    if (i \le 1) \{ xn = i; \}
    /* fib(n)=fib(n-1)+fib(n-2) */
    else {xn = xa+xb; xa=xb; xb=xn; }
    /* save results if iteration i is within the
       range from n-len to n */
    j=i - *n + *len - 1;
    if (j>=0) retArr[j] = xn;
  } /* end loop */
} /* end function */
```

The function calcFib in PBStry illustrates an application of these concepts (Table A2). The R function uses C code to calculate the first n Fibonacci numbers iteratively, where a vector holds the last len numbers calculated. After ensuring that n and len satisfy obvious constraints, the R code creates a return array retArr of the appropriate length. The .C call passes n, len, and retArr by reference to the C function fibonacci. On exit, the vector out contains a list corresponding to the input variables n, len, and retArr, so that the third component out[[3]] holds the modified vector of values calculated by fibonacci. We encourage you also to examine the second example in PBStry, associated the files calcSum.R and sum.c.

**Table A3.** .Call() example adapted from PBSmodelling, with two associated text files. R code in the first file calls C code in the second.

```
File 1: callFib.R
callFib <- function(n, len=1) {</pre>
  out <- .Call("fibonacci2", as.integer(n),</pre>
               as.integer(len), PACKAGE="PBSmodelling")
  return(out) }
                            File 2: fib2.c
#include <R.h>
#include <Rdefines.h>
SEXP fibonacci2(SEXP sexp_n, SEXP sexp_len) {
  /* ptr to output vector that we will create */
  SEXP retVals;
  double *p_retVals, xa=0, xb=1, xn; int n, len, i, j;
  /* convert R variables into C 'int's */
  len = INTEGER VALUE(sexp len);
  n = INTEGER_VALUE(sexp_n);
  /* Allocate space for the output vector */
  PROTECT(retVals = NEW_NUMERIC(len));
  p_retVals = NUMERIC_POINTER(retVals);
  /* iterative loop */
  for(i=0;i<=n;i++) {
    /* initial conditions: fib(0)=0, fib(1)=1 */
    if (i \le 1) \{ xn = i; \}
    /* fib(n)=fib(n-1)+fib(n-2) */
    else { xn = xa+xb; xa=xb; xb=xn; }
    /* save results if iteration i is within the
       range from n-len to n */
    j=i - n + len - 1;
    if (j>=0) p_retVals[j] = xn;
  } /* end loop */
  UNPROTECT(1);
  return retVals;
} /* end fibonacci2 */
```

### Calling C functions from R using .Call()

The .C() convention requires a fairly simple conversion of R objects into C types (Table A.1). By contrast, .Call() provides extra structure that enables C to handle R objects directly (RDCT 2006b, section 4.7). This function uses "S-expression" SEXP types defined in rinternals.h., a file in the \include directory of the R installation. An SEXP pointer can reference any type of R object. The .Call() convention uses the following key concepts:

- C functions called by R must accept only SEXP typed arguments. These arguments should be treated as read only.
- Similarly, C functions called by R must have SEXP return types.
- The Programmer must protect R objects from the R garbage collector, and must release protected objects before the function terminates. R provides macros for this task.
- C code written for the shared DLL must not contain a main function.
- Within a C function, dynamically allocated memory must be de-allocated by the programmer before the function returns. Otherwise a memory leak will likely occur.

The function callFib in Table A3 (adapted from .fibCall in PBSmodelling) illustrates an application of these concepts. As before, the R function uses C code to calculate the first n Fibonacci numbers iteratively, where a vector holds the last len numbers calculated. The code in callFib assumes that n and len already satisfy the necessary constraints. The simple .Call to fibonacci2 looks very natural. Input values n and len produce the output vector out, where the C code must somehow determine what out should be. Not surprisingly, it requires more complicated C code to make this happen.

The C function fibonacci2 (Table A3) first loads header files that include the required definitions from R. All input and output variables belong to type SEXP. Other internal variables have the standard C types double and int. Functions like INTEGER\_VALUE() convert R types into C types. The SEXP vector retVals of return values is created by the R constructor NEW\_NUMERIC() and then protected from garbage collection by PROTECT(). After all required variables are defined and type cast correctly, the iterative loop of calculations follows the earlier example in Table A2. Finally, the only protected vector retVals is released by UNPROTECT(1), and the standard closing command return retVals returns the output vector from fibonacci2.

Obviously, it takes some time and effort to become familiar with the specialized R types, constructors, and conversion functions. For this reason, it's probably easier at first to use .C(), rather than .Call().

# **Appendix B. Widget descriptions**

This appendix lists PBS Modelling widgets in alphabetical order. Details for each widget include a description, usage, arguments, and an illustrated example. In specifying a widget, the user can arrange named arguments in any order. If arguments are not named, they must appear in the order specified by the argument list, similar to named arguments in an R function.

# **Button**

### Description

A button linked to an R function that runs a particular analysis and generates a desired output, perhaps including graphics.

# Usage

```
type=button text="Calculate" font="" width=0 function=""
   action="button" sticky="" padx=0 pady=0
```

# Arguments

# Example

```
window title="Widget = button"
button text="Push Me"
```



## Check

# Description

A check box used to turn a variable off or on, with corresponding values FALSE or TRUE.

### Usage

```
type=check name checked=FALSE text="" font="" function=""
   action="check" sticky="" padx=0 pady=0
```

#### Arguments

namename of R variable altered by this check box (required)	
checkedif TRUE, the box is checked initially and the variable is set to 1	
textidentifying text placed to the right of this check box	
fontfont for check text - specify family (Times, Helvetica, or Courier), size (as	
point size), and style (bold, italic, underline, overstrike), in any order	
functionR function to call when the check box is changed	
actionvalue for PBS.win\$action when altering this check box	
stickyoption for placing the widget in available space; valid choices are:	
N, NE, E, SE, S, SW, W, NW	
padxspace used to pad the widget on the left and right	
padyspace used to pad the widget on the top and bottom	

## Example

```
window title="Widget = check"
check name=junk checked=T text="Check Me"
```



#### Data

## Description

An aligned set of entry fields for all components of a data frame. The data widget can accept a variety of modes. The user must keep in mind that rowlabels and collabels should conform to R naming conventions (no spaces, no special characters, etc.). If mode is logical, fields appear as a set of check boxes that can be turned on or off using mouse clicks.

#### Usage

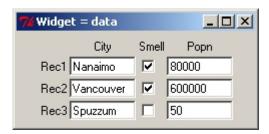
```
type=data nrow ncol names modes="numeric" rowlabels=""
  collabels="" rownames=X colnames= font="" values=
  byrow=TRUE function="" enter=TRUE action="data" width=6
  sticky="" padx=0 pady=0
```

nrownumber of rows (required)
ncolnumber of columns(required)
nameseither one name or a set of nrow*ncol names used to store the data
frame in R (required)
modesR modes for the data frame, where valid modes are:
numeric, integer, complex, logical, character
rowlabelseither one label or a vector of nrow labels used to label rows of this data
frame in the display

collabelseither one label or a vector of ncol labels used to label columns of this
data frame in the display
fontfont for labels - specify family (Times, Helvetica, or Courier), size (as
point size), and style (bold, italic, underline, overstrike), in any order
valuesdefault values (either one value for all data frame components or a set of
nrow*ncol values)
byrowif TRUE and nrow*ncol names are used, interpret the names by row;
otherwise by column. Similarly, interpret nrow*ncol initial values.
functionR function to call when any entry in the data frame is changed
enterif TRUE, call the function only after the <enter> key is pressed</enter>
actionvalue for PBS. win action when changing any component of this data
frame
widthcharacter width to reserve for the each entry in the data frame
stickyoption for placing the widget in available space; valid choices are:
N, NE, E, SE, S, SW, W, NW
padxspace used to pad the widget on the left and right
padyspace used to pad the widget on the top and bottom

### Example

```
window title="Widget = data"
data nrow=3 ncol=3 names=Census byrow=FALSE \
    modes="character logical numeric" width=10 \
    rowlabels="Rec1 Rec2 Rec3" collabels="City Smell Popn" \
    values="Nanaimo Vancouver Spuzzum T T F 80000 600000 50"
```



# **Entry**

### Description

A field in which a scalar variable (number or string) can be altered.

### Usage

```
type=entry name value="" width=20 label="" font="" function=""
  enter=TRUE action="entry" mode="numeric" sticky="" padx=0
  pady=0
```

### Arguments

name.....name of R variable corresponding to this entry (required) value.....default value to display in the entry

widthcharacter width to reserve for the entry
labeltext to display above the entry box
fontfont for label - specify family (Times, Helvetica, or Courier), size (as point
size), and style (bold, italic, underline, overstrike), in any order
functionR function to call when the entry is changed
enterif TRUE, call the function only after the <enter> key is pressed</enter>
actionvalue for PBS.win\$action when making this entry
modeR mode for the value entered, where valid modes are:
numeric, integer, complex, logical, character
stickyoption for placing the widget in available space; valid choices are:
N, NE, E, SE, S, SW, W, NW
padxspace used to pad the widget on the left and right
padyspace used to pad the widget on the top and bottom

# Example

```
window title="Widget = entry"
entry name=junk value="Enter something here" width=20
    mode=character
```



### Grid

# Description

Creates space for a rectangular block of widgets. Spaces must be filled. Widgets can be any combination of available widgets, including grid.

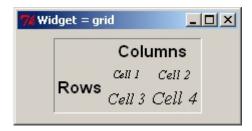
### Usage

```
type=grid nrow=1 ncol=1 toptitle="" sidetitle="" topfont=""
    sidefont="" byrow=TRUE borderwidth=1 relief="flat"
    sticky="" padx=0 pady=0
```

nrow	number of rows in the grid
ncol	number of columns in the grid
toptitle	.title to place above grid
sidetitle	title to place on the left side of the grid.
topfont	font for top of grid - specify family (Times, Helvetica, or Courier), size (as
	point size), and style (bold, italic, underline, overstrike), in any order
sidefont	font for left of grid - specify family (Times, Helvetica, or Courier), size (as
	point size), and style (bold, italic, underline, overstrike), in any order
byrow	if TRUE, create widgets across rows, otherwise down columns

#### Example

```
grid 2 2 relief=groove toptitle=Columns sidetitle=Rows
    topfont="Helvetica 12 bold" sidefont="Helvetica 12 bold"
    label text="Cell 1" font="times 8 italic"
    label text="Cell 2" font="times 10 italic"
    label text="Cell 3" font="times 12 italic"
    label text="Cell 4" font="times 14 italic"
```



# History

#### Description

Allows the user to save and recover previous combinations of widget settings. A desired realization can be saved to a hidden stack using the Add button. Each saved realization has an Index within the stack of total saves Size. The user can scroll back and forward through the stack, invoking saved widget settings. Realizations no longer desired can be deleted using the Remove button.

#### Usage

```
type=history name="default" archive=TRUE sticky="" padx=0
   pady=0
```

namename of history archive
archiveif TRUE, user can import previous sessions or export the current session
stickyoption for placing the widget in available space; valid choices are:
N, NE, E, SE, S, SW, W, NW
padxspace used to pad the widget on the left and right
padyspace used to pad the widget on the top and bottom

# Example

window title="Widget = history"
history archive=TRUE



# Label

### Description

Creates a text label. If the text argument is left blank, label emulates the null widget.

#### Usage

```
type=label text="" font="" sticky="" padx=0 pady=0
```

# Arguments

text ......text to display in the label

font ......font for label - specify family (Times, Helvetica, or Courier), size (as point size), and style (bold, italic, underline, overstrike), in any order

sticky ......option for placing the widget in available space; valid choices are:

N, NE, E, SE, S, SW, W, NW

padx ......space used to pad the widget on the left and right

pady .....space used to pad the widget on the top and bottom

### Example

```
window title="Widget = label"
label text="Information Label"
```



#### **Matrix**

#### Description

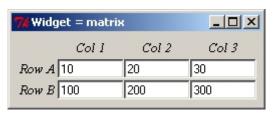
An aligned set of entry fields for all components of a matrix. If the mode is logical, the matrix appears as a set of check boxes that can be turned on or off using mouse clicks.

### Usage

```
type=matrix nrow ncol names rowlabels= collabels= rownames=""
  colnames="" font="" values="" byrow=TRUE function=""
  enter=TRUE action="matrix" mode="numeric" width=6 sticky=""
  padx=0 pady=0
```

# Arguments

nrownumber of rows (required)
ncolnumber of columns(required)
nameseither one name or a set of nrow*ncol names used to store the matrix in
R (required)
rowlabelseither one label or a vector of nrow labels used to label rows of this
matrix in the display
collabelseither one label or a vector of ncol labels used to label columns of this
matrix in the display
fontfont for labels - specify family (Times, Helvetica, or Courier), size (as
point size), and style (bold, italic, underline, overstrike), in any order
valuesdefault values (either one value for all matrix components or a set of
nrow*ncol values)
byrowif TRUE and nrow*ncol names are used, interpret the names by row;
otherwise by column. Similarly, interpret nrow*ncol initial values.
functionR function to call when any entry in the matrix is changed
enterif TRUE, call the function only after the <enter> key is pressed</enter>
actionvalue for PBS.win\$action when changing any component of this
matrix
modeR mode for the matrix, where valid modes are:
numeric, integer, complex, logical, character
widthcharacter width to reserve for the each entry in the matrix
stickyoption for placing the widget in available space; valid choices are:
N, NE, E, SE, S, SW, W, NW
padxspace used to pad the widget on the left and right
padyspace used to pad the widget on the top and bottom



#### Menu

### Description

A menu grouping. Submenus can either be menu or menuitem.

### Usage

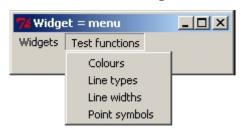
```
type=menu nitems=1 label font=""
```

#### Arguments

```
nitems ......number of items or submenus to include in the menu
label......text to display as the menu label (required)
font ......font for menus - specify family (Times, Helvetica, or Courier), size (as
point size), and style (bold, italic, underline, overstrike), in any order
```

### Example (assuming that the R functions have been defined)

```
window title="Widget = menu"
menu nitems=1 label="Widgets"
    menuitem label="Show arguments" func=showArgs
menu nitems=4 label="Test functions"
    menuitem label="Colours" func=testCol
    menuitem label="Line types" func=testLty
    menuitem label="Line widths" func=testLwd
    menuitem label="Point symbols" func=testPch
```



### MenuItem

#### Description

One of nitems following a menu command.

#### Usage

```
type=menuitem label font="" function action="menuitem"
```

```
label......text to display as the menu item label (required)

font......font for submenus - specify family (Times, Helvetica, or Courier), size (as point size), and style (bold, italic, underline, overstrike), in any order function......R function to call when the menu item is clicked (required) action.....value for PBS.win$action when selecting this menu item
```

#### Null

### Description

Creates a null widget, useful for padding a grid with blank cells that appear as empty space.

### Usage

```
type=null padx=0 pady=0
```

### Arguments

```
padx.....space used to pad the label on the left and right pady.....space used to pad the label on the top and bottom
```

#### Example

```
grid 2 2 relief=raised toptitle=Top sidetitle=Side
    topfont="Courier 10 bold" sidefont="courier 10 bold"
    label text="Here" font="courier 8"
    null
    null
    label text="There" font="courier 8"
```



# Radio

## Description

One of a set of mutually exclusive radio buttons for making a particular choice. Buttons with the same value for name act collectively to define a single choice among the alternatives.

#### Usage

```
type=radio name value text="" font="" function=""
   action="radio" mode="numeric" sticky="" padx=0 pady=0
```

name	name of R variable altered by this radio button, where radio buttons with
	the same name define a mutually exclusive set (required)
value	value of the variable when this radio button is selected (required)
text	identifying text placed to the right of this radio button
font	font for radio buttons - specify family (Times, Helvetica, or Courier), size
	(as point size), and style (bold, italic, underline, overstrike), in any order
function	R function to call when this radio button is selected
action	value for PBS, winSaction when this radio button is selected

# Example

```
window title="Widget = radio"
grid 1 4
    radio name=junk value=0 text="None"
    radio name=junk value=1 text="Option A"
    radio name=junk value=2 text="Option B"
    radio name=junk value=3 text="Option C"
```



# Slide

# Description

A slide bar that sets the value of a variable. This widget only accepts integer values.

### Usage

```
type=slide name from=0 to=100 value=NA showvalue=FALSE
  orientation="horizontal" function="" action="slide"
  sticky="" padx=0 pady=0
```

# Example

```
window title="Widget = slide"
slide name=junk from=1 to=1000 value=225 showvalue=T
```



### **SlidePlus**

### Description

An extended slide bar that also displays a minimum, maximum, and current value. This widget accepts real numbers.

#### Usage

```
type=slideplus name from=0 to=1 by=0.01 value=NA function=""
enter=FALSE action="slideplus" sticky="" padx=0 pady=0
```

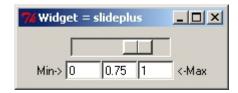
### Arguments

namename of the numeric R variable corresponding to this slide bar (required)
fromminimum value of the variable
tomaximum value of the variable
byminimum amount for changing the variable's value
valueinitial slide value, where the default is the specified from value
functionR function to call when the slide value is changed
enterif TRUE and the slide value is changed via the entry box, call the function
only after the <enter> key is pressed</enter>
actionvalue for PBS.win\$action when changing the slide value, either by
moving the slide bar or changing the value in the entry box
stickyoption for placing the widget in available space; valid choices are:
N, NE, E, SE, S, SW, W, NW
padxspace used to pad the widget on the left and right
padyspace used to pad the widget on the top and bottom

### Note

To facilitate retrieving and setting the minimum and maximum values, two additional variables are created by suffixing ".max" and ".min" to the given name.

```
window title="Widget = slideplus"
slideplus name=junk from=0 to=1 by=0.01 value=0.75
```



#### **Text**

### Description

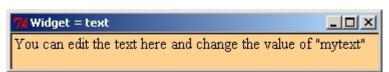
An information text box that can display messages, results, or whatever the user desires. The displayed information can be either fixed or editable.

#### Usage

```
type=text name height=8 width=30 bg="white" mode="character"
font="" value="" borderwidth=1 relief="sunken" edit=TRUE
```

# Arguments

```
window title="Widget = text"
text name=mytext height=2 width=55 bg="#FFD18F" \
    font="times 11" borderwidth=1 relief="sunken" edit=TRUE \
    value="You can edit the text here and change the value of \"mytext\""
```



### Vector

# Description

An aligned set of entry fields for all components of a vector. If the mode is logical, the vector appears as a set of check boxes that can be turned on or off using mouse clicks.

### Usage

```
type=vector names length=0 labels="" values="" font=""
  vertical=FALSE function="" enter=TRUE action="vector"
  mode="numeric" width=6 sticky="" padx=0 pady=0
```

# Arguments

1	nameseither one name (for a whole vector) or a vector of names for individual
	variables used to store the values in R (required)
	lengthrequired only if a single name is given for a vector of length greater than 1
	labelseither one label of a vector of length labels used to label the vector in the display
	± •
,	valuesdefault values (either one value for all vector components or a vector of
	length values)
1	fontfont for vector labels - specify family (Times, Helvetica, or Courier), size
	(as point size), and style (bold, italic, underline, overstrike), in any order
7	verticalif TRUE, display the vector as a vertical column with labels on the left;
	otherwise display it as a horizontal row with labels above
1	functionR function to call when any entry in the vector is changed
(	enterif TRUE, call the function only after the <enter> key is pressed</enter>
ć	actionvalue for PBS.win\$action when changing any component of this vector
7	modeR mode for the vector, where valid modes are:
	numeric, integer, complex, logical, character
7	widthcharacter width to reserve for the each entry in the vector
Ş	stickyoption for placing the widget in available space; valid choices are:
	N, NE, E, SE, S, SW, W, NW
1	padxspace used to pad the widget on the left and right
1	padyspace used to pad the widget on the top and bottom



#### Window

# Description

Create a new window. Windows are used as a palette upon which widgets are placed. Each open window has a unique name. The function closeWin closes all windows unless a specific name (or vector of names) is provided by the user. Also, if createWin opens a window with a name already in use, the older window is closed before the new window is opened.

#### Usage

```
type=window name="window" title="" vertical=TRUE
```

#### Arguments

#### Example

window title="Widget = window (upon which all other widgets
 are placed)"



# Appendix C. PBS Modelling functions and data

This appendix lists the functions currently made available by *PBS Modelling*, along with a list of their dependencies on hidden "dot" functions. For more information about the hidden functions, see the file PBSMfunctions.txt in the R library directory for PBSmodelling.

#### C.1. Objects in *PBS Modelling*

CCA.qbr.hl Dataset: sampled counts of quillback rockfish (*Sebastes maliger*)

addArrows

addLabel

Add a label to a plot using relative (0:1) coordinates

Add a label to a plot using relative (0:1) coordinates

Add a legend to a plot using relative (0:1) coordinates

Calculate Fibonacci numbers by several methods

Calculate the geometric mean, allowing for zeros

Calculate Tobjects from the global environment

clearWinVal Remove all current widget variables

closeWin Close GUI window(s)

compareLists Compare two non-recursive lists

compileDescription Convert and save a window description as a list

create Vector Create a GUI with a vector widget

createWin Create a GUI window

drawBars Draw a linear barplot on the current plot expandGraph Expand the plot area by adjusting margins

exportPBShistory Export a saved history

findPat Search a character vector to find multiple patterns

genMatrix Generate test matrices for plotBubbles getWinVal Retreive widget values for use in R code

importPBShistory Import a history list from a file

initPBShistory Create structures of a new history widget openFile Open a file with the associated program

pad0 Pad numbers with leading zeroes

parseWinFile Convert a window description file into a list
pause Pause between graphics displays or other calculations

pickCol Display an interactive colour selection palette

plotBubbles Construct a bubble plot from a matrix

plotCsum Plot cumulative sum of data promptOpenFile Display an "Open File" dialog promptSaveFile Display a "Save File" dialog

readList Read a list from a file in PBS Modelling format

resetGraph Reset par values for a plot

runExamples Run GUI examples included with *PBS Modelling* 

setWinVal Update widget values

show0 Convert numbers into text with specified decimal places

showArgs Display expected widget arguments

testCol Display colour palette
testLty Display line types
testLwd Display line widths
testPch Display plotting symbols

testWidgets Display sample GUIs and their source code unpackList Unpack list elements into variables view Display first n rows of an object

writeList Write a list to a file in PBS Modelling format

# C.2. Function dependencies

This appendix documents function dependencies within *PBS Modelling*. All functions appear as underlined entries in alphabetic order. If a function depends on others, the list of dependencies appears below the underlined name. Following a standard in UNIX and R, functions whose name begins with a period (*dot functions*) are considered hidden from the user. *PBS Modelling* enforces this standard through the NAMESPACE protocol discussed above in section A.3.

- d-d1b	
<u>.addslashes</u>	.createWidget.grid
and a Commont Marila	.createWidget.history
<pre>.autoConvertMode</pre>	.createWidget.label
	.createWidget.matrix
.buildgrid	.createWidget.null
.createTkFont	.createWidget.radio
.createWidget	.createWidget.slide
	.createWidget.slideplus
<u>.catError</u>	$. {\tt createWidget.text}$
	.createWidget.vector
.convertMatrixListToDataFrame	
.getMatrixListSize	
.setMatrixElement	.createWidget.button
	.createTkFont
<pre>.convertMatrixListToMatrix</pre>	.extractData
.getMatrixListSize	
.setMatrixElement	.createWidget.check
	.createTkFont
.convertMode	.extractData
.convertMode	
	.createWidget.data
.convertPararmStrToList	.createWidget.grid
.catError	.stopWidget
.trimWhiteSpace	
<del>-</del>	.createWidget.entry
.convertPararmStrToVector	.createWidget.grid
.catError	.extractData
.trimWhiteSpace	
	.createWidget.grid
.convertVecToArray	.buildgrid
.getArrayPts	
.mapArrayToVec	.createWidget.history
· · · · · · · · · · · · · · · · · · ·	.createWidget.grid
.createTkFont	initPBShistory
.convertPararmStrToVector	
. conver er draf inder 10 vec eor	.createWidget.label
.createWidget	.createTkFont
.createWidget.button	. 01 04001111 0110
.createWidget.check	.createWidget.matrix
.createWidget.data	.createWidget.matrix
<del>-</del>	.stopWidget
.createWidget.entry	.scopwiaget

	.trimWhiteSpace
.createWidget.null	.getParamOrder
.createWidget.radio	
.createTkFont	.getPBS.win
.extractData	.extractVar
.createWidget.slide	.getReadListParamOrder
.extractData	
	.hash
<pre>.createWidget.slideplus .extractData</pre>	.isReallyNull
.extractbata	.inCollection
.createWidget.text	<u>:::::::::::::::::::::::::::::::::::::</u>
.createTkFont	.initPBSoptions
.createWidget.vector	.isReallyNull
.createWidget.grid	
.stopWidget	.mapArrayToVec
.extractData	.matrixHelp
.getPBS.win	.matrixHelp
.extractFuns	.parsegrid
.extractVar	.parsegrid
.convertMatrixListToDataFrame	.parsemenu
.convertMatrixListToMatrix	.parsemenu
.matrixHelp	·parsemena
.PBSdimnameHelper	.PBSdimnameHelper
_	
.fibC	<u>.readList.P</u>
	.catError
.fibCall	.readList.P.convertData
	.stripComments
.fibClosedForm	.trimWhiteSpace
.fibR	.readList.P.convertData
	.autoConvertMode
.getArrayPts	.catError
	.convertMode
<pre>.getMatrixListSize</pre>	$. \verb convertPararmStrToVector  \\$
.getMatrixListSize	<pre>.convertVecToArray</pre>
	.getParamFromStr
<pre>.getParamFromStr</pre>	.getReadListParamOrder
.catError	
.convertPararmStrToList	<pre>.searchCollection</pre>
.isReallyNull	
.searchCollection	.setMatrixElement
.stripSlashes	$.\mathtt{setMatrixElement}$
.stripSlashesVec	

.setWinValHelper calcGM .getPBS.win .setWinValHelper clearAll clearPBShistory .stopWidget .updatePBShistory .stripComments rmPBShistory .stripComments .trimWhiteSpace clearWinVal .stripSlashes closeWin .catError .isReallyNull .stripSlashesVec compareLists .catError .isReallyNull compileDescription .trimWhiteSpace parseWinFile writeList .updatePBShistory setWinVal createVector .validateWindowDescList createWin .getParamOrder createWin .validateWindowDescWidgets .createWidget .validateWindowDescWidgets .getPBS.win .hash .getParamOrder .isReallyNull .writeList.P .validateWindowDescList .addslashes parseWinFile addArrows drawBars addLabel expandGraph addLegend exportPBShistory promptSaveFile writeList addPBShistory .updatePBShistory getWinVal findPat backPBShistory forwPBShistory .updatePBShistory .updatePBShistory setWinVal setWinVal calcFib genMatrix .fibC .fibCall getWinVal .fibClosedForm .getPBS.win .fibR .isReallyNull

importPBShistory
.updatePBShistory
promptOpenFile

readList

initPBShistory

jumpPBShistory
.updatePBShistory
getWinVal
setWinVal

openFile

.initPBSoptions .isReallyNull openFile

pad0

parseWinFile

.getParamFromStr

.parsegrid
.parsemenu

 $. \verb|stripComments| \\$ 

.trimWhiteSpace

<u>pause</u>

pickCol

plotBubbles

plotCsum
addLabel
resetGraph

promptOpenFile
.trimWhiteSpace

promptSaveFile
promptOpenFile

readList
.readList.P

resetGraph

rmPBShistory

.updatePBShistory

setWinVal

runExamples closeWin createWin getWinVal setWinVal

setWinVal
.isReallyNull
.setWinValHelper

show0

showArgs

.getParamOrder

testCol

testLty

testLwd resetGraph

testPch resetGraph

testWidgets closeWin createWin getWinVal setWinVal

unpackList

view

writeList
.writeList.P

addArrows

Add Arrows to a Plot Using Relative (0:1) Coordinates

# Description

Calls arrows function using relative (0:1) coordinates.

# Usage

```
addArrows(x1, y1, x2, y2, ...)
```

# Arguments

x1	x-coordinate at base of arrow
y1	y-coordinate at base of arrow
x2	x-coordinate at tip of arrow
y2	y-coordinate at tip of arrow
	additional paramaters for arrows

# Details

```
Lines will be drawn from (x1[i],y1[i]) to (x2[i],y2[i])
```

### See Also

```
addLabel
addLegend
```

```
tt=seq(from=-5,to=5,by=0.01)
plot(sin(tt), cos(tt)*(1-sin(tt)), type="1")
addArrows(0.2,0.5,0.8,0.5)
addArrows(0.8,0.95,0.95,0.55, col="#FF0066")
```

addLabel

Add Label to a Plot Using Relative (0:1) Coordinates

# Description

Places a label in a plot using relative (0:1) coordinates

# Usage

```
addLabel(x, y, txt, ...)
```

# Arguments

x x-axis coordinate in the range (0,1); can step outside y y-axis coordinate in the range (0,1); can step outside x txt desired label at (x,y)

... additional arguments passed to text function

#### See Also

addArrows addLegend

# Examples

```
resetGraph()
addLabel(0.75,seq(from=0.9,to=0.1,by=-0.10),c('a','b','c'), col="#0033AA")
```

addLegend

Add Legend to a Plot Using Relative (0:1) Coordinates

# Description

Places a legend in a plot using relative (0:1) coordinates.

### Usage

```
addLegend(x, y, ...)
```

### **Arguments**

x x-axis coordinate in the range (0,1); can step outside y y-axis coordinate in the range (0,1); can step outside

... arguments used by legend, such as "lines", "text", or "rectangle"

#### See Also

addArrows

addLabel

calcFib

Several Methods to Calculate Fibonacci Numbers

### Description

Computes Fibonacci numbers using four different methods: 1) iteratively using R code, 2) via the closed function in R code, 3) iteratively in C using the .C function, and 4) iteratively in C using the .Call function.

# Usage

```
calcFib(n, len=1, method="C")
```

### **Arguments**

n nth fibonacci number to calculate

len A vector of length len showing previous fibonacci numbers

method Select method to use: C, Call, R, closed

#### Value

Vector of the last len Fibonacci numbers calculated.

calcGM

Calculate the Geometric Mean

# Description

Calculate the geometric mean of a numeric vector, possibly excluding zeros and/or adding an offset to compensate for zero values.

# Usage

```
calcGM(x, offset = 0, exzero = TRUE)
```

# Arguments

x vector of numbers

offset value to add to all components, including zeroes

exzero if T, exclude zeroes (but still add the offset)

#### Value

geometric mean of the modified vector  $\mathbf{x} + \mathbf{offset}$ 

#### Note

NA values are automatically removed from x

# Examples

```
calcGM(c(0,1,100))
calcGM(c(0,1,100),offset=0.01,exzero=FALSE)
```

CCA.qbr.hl

Dataset of Sampled Counts of Quillback Rockfish (Sebastes maliger)

# Description

Count of sampled fish-at-age for quillback rockfish (Sebastes maliger) in Johnstone Strait, British Columbia, from 1984 to 2004.

# Usage

```
data(CCA.qbr.hl)
```

#### **Format**

A matrix with 70 rows (ages) and 14 columns (years). Attributes "syrs" and "cyrs" specify years of survey and commercial data, respectively.

```
[,c(3:5,9,13,14)] Counts-at-age from research survey samples [,c(1,2,6:8,10:12)] Counts-at-age from commercial fishery samples
```

All elements represent sampled counts-at-age in year. Zero-value entries indicate no observations.

#### **Details**

Handline surveys for rockfish have been conducted in Johnstone Strait (British Columbia) and adjacent waterways (126°37'W to 126°53'W, 50°32'N to 50°39'N) since 1986. Yamanaka and Richards (1993) describe surveys conducted in 1986, 1987, 1988, and 1992. In 2001, the Rockfish Selective Fishery Study (Berry 2001) targeted quillback rockfish Sebastes maliger for experiments on improving survival after capture by hook and line gear. The resulting data subsequently have been incorporated into the survey data series. The most recent survey in 2004 essentially repeated the 1992 survey design. Fish samples from surveys have been supplemented by commercial handline fishery samples taken from a larger region (126°35'W to 127°39'W, 50°32'N to 50°59'N) in the years 1984-1985, 1989-1991, 1993, 1996, and 2000 (Schnute and Haigh 2006).

#### Note

Years 1994, 1997-1999, and 2002-2003 do not have data.

#### Source

```
Fisheries and Oceans Canada - GFBio database: http://www-sci.pac.dfo-mpo.gc.ca/sa-mfpd/statsamp_GFBio.htm
```

#### References

Berry, M.D. 2001. Area 12 (Inside) Rockfish Selective Fishery Study. Science Council of British Columbia, Project Number FS00- 05.

Schnute, J.T., and Haigh, R. 2006. Compositional analysis of catch curve data with an application to *Sebastes maliger*. ICES Journal of Marine Science (in revision).

Yamanaka, K.L. and Richards, L.J. 1993. 1992 Research catch and effort data on nearshore reef-fishes in British Columbia Statistical Area 12. Canadian Manuscript Report of Fisheries and Aquatic Sciences 2184, 77 pp.

```
# Plot age proportions (blue bubbles = survey data, red = commercial)
data("CCA.qbr.hl", package="PBSmodelling")
```

clearAll

 $Remove\ All\ R\ Objects$ 

# Description

Generic function to clear .RData in R

# Usage

clearAll()

clearWinVal

Remove All Widget Variables

# Description

Removes all global variables that share a name in common with any widget variable name as defined in PBS.win\$vars. Use this with caution.

### Usage

```
clearWinVal()
```

#### See Also

getWinVal

closeWin

Close GUI windows

# Description

The closeWin function closes (destroys) one or more windows made with createWin.

# Usage

```
closeWin(name=names(.PBS.tclHash))
```

# Arguments

name

a vector of window names to close, as defined in the window description file's WINDOW widget.

### See Also

createWin

compareLists

Compares two non-recursive lists

# Description

Displays any differences between two non-recursive lists.

# Usage

```
compareLists(a,b,verbose=FALSE)
```

# Arguments

a list ab list b

verbose be verbose

#### compileDescription

Converts and Saves a Window Description Into a List

# Description

compileDescription converts a Window Description File into an equivalent Window Description List. The list is complete - meaning all default values have been added to the list.

# Usage

```
compileDescription(descFile, outFile)
```

# Arguments

descFile filename of markup file.

outFile filename indicating where to save outputed R source code.

#### **Details**

The Window Description File is converted into a list, which is then converted into R code and saved to the outfile.

#### See Also

parseWinFile
createWin

createVector

Create a GUI with a Vector Widget

# Description

createVector creates a basic window containing a vector and a submit button. This provides a quick way to create a window without the need for a window description file.

# Usage

```
createVector(vec, vectorLabels=NULL, func="", windowname="vectorwindow")
```

# **Arguments**

vec

a vector of strings representing widget variables. If it is named, the names are used as the variable names, and the values are used as the default value

of the widget

vectorLabels

option vector of strings to be used as labels above each widget.

func

string value of function name to be called when new data is entered or

when "GO" is pressed.

windowname

window name required if multiple vector windows are created.

#### See Also

createWin

# Examples

createWin

Create a GUI Window

### Description

The createWin function takes a window markup file, and creates a window based on the markup file.

# Usage

```
createWin(fname, astext=FALSE)
```

#### Arguments

fname filename of markup file or list returned from parseWinFile.

astext if true, fname is intrupted as a vector of strings. with each element repre-

senting a line of the source of a window description file

#### **Details**

The markup file contains a single widget per line. Widgets can span multiple lines by including a backslash ('\') as the last character of the line, which then ignores the newline.

For more details about widget types, and the markup file see the pdf located in the installation directory.

It is possible to use a Window Description List produced by compileDescription rather than a filename for the fname argument.

Another alternative is to set astext=TRUE and pass in a vector of characters for fname. This vector of characters represents the file contents. Each element of the vector is equivalent to a new line in the window description file.

#### Value

PBS.win contains window information such as: present values, present variable names, present variable values, and triggered action values. This information is encapsulated in a list which is set as the global PBS.win variable, which is also returned.

PBS.win

vars Current widget values

funs Functions required by Window

names Variable names: names(PBS.win\$vars)
action Action that triggered a function call

### See Also

parseWinFile
getWinVal
setWinVal

closeWin

compileDescription

createVector

initPBShistory for an example of using astext=TRUE

# Examples

```
#see file testWidgets\LissWin.txt in PBSmodelling package directory
# window title="Lissajous Curve"
# grid 1 2
    label text=Pars: font=bold
   vector length=4 names="m n phi k" \
      labels="'x cycles' 'y cycles' 'y phase' points" \
#
      values="2 3 0 1000" vertical=T
# grid 1 2
    label text=History: font=bold
   history
# grid 1 2
   grid 2 1
#
      radio name=ptype text=lines value="1" mode=character
      radio name=ptype text=points value="p" mode=character
# button text=Plot func=drawLiss font=bold
# Calculate and draw the Lissajous figure
drawLiss <- function()</pre>
 getWinVal(scope="L");
 ti <- 2*pi*(0:k)/k;
 x <- sin(2*pi*m*ti);</pre>
 y <- sin(2*pi*(n*ti+phi));</pre>
 plot(x,y,type=ptype);
  invisible(NULL);
}
## Not run:
require(PBSmodelling);
createWin(system.file("testWidgets/LissWin.txt",package="PBSmodelling"))
## End(Not run)
```

drawBars

Draw a Linear Barplot on the Current Plot

# Description

Draws a linear barplot on the current graph.

## Usage

```
drawBars(x, y, width, base = 0, ...)
```

x x-coordinatesy y-coordinates

... further graphical parameters (see 'par') may also be supplied as arguments

### Examples

```
plot(0:10,0:10,type="n")
drawBars(x=1:9,y=9:1,col="deepskyblue4",lwd=3)
```

expandGraph Expand Plot Area by Adjusting Margins

# Description

Tries to maximize the area of multiple plots by minimizing margins.

# Usage

```
expandGraph(mar=c(4,3,1.2,0.5), mgp=c(1.6,.5,0),...)
```

### **Arguments**

numerical vector of the form 'c(bottom, left, top, right)' specifying the

margins of the plot

mgp numerical vector of the form 'c(axis title, axis labels, axis line)' specifying

the margins for axis title, axis labels, and axis line

... Additional graphical parameters to be passed to par()

### See Also

 ${\tt resetGraph}$ 

par

# Examples

exportPBShistory Export Saved History

# Description

Exports the current history List.

### Usage

```
exportPBShistory(hisname="", fname="")
```

### Arguments

hisname Name of the history list to export. If it is set to "", the value from

PBS.win\$action[1] will be used instead.

fname Where to save the history to. If it is set to "", a save file window will be

displayed.

#### See Also

```
importPBShistory
initPBShistory
promptSaveFile
```

findPat

Search a Vector With Multiple Patterns

# Description

Searches all patterns in pat from vec, and returns the matched elements in vec.

# Usage

```
findPat(pat, vec)
```

### **Arguments**

pat character vector of patterns to match in vec vec character vector where matches are sought

#### Value

A character vector of all matched strings.

# Examples

```
#find all strings with a vowel, or that start with a number
findPat(c("[aeoiy]", "^[0-9]"), c("hello", "WRLD", "11b"))
```

genMatrix

Generate Test Matrices for plotBubbles

### Description

Generates a test matrix of random numbers (mean = mu and standard deviation = sigma), primarily for plotBubbles.

### Usage

```
genMatrix(m,n,mu=0,sigma=1)
```

# **Arguments**

m number of rows

n number of columns

mu mean of normal distribution

sigma standard deviation of normal distribution

#### Value

An m by n matrix with normally distributed random values.

#### See Also

plotBubbles

### Examples

plotBubbles(genMatrix(20,6))

getWinVal

Retreive Widget Values

# Description

An optional scope argument allows the function to create local or global variables based on the list that is returned.

#### Usage

```
getWinVal(v=NULL, scope="", asvector=FALSE, windowname="")
```

### Arguments

v vector of variable names to be retrieved. If NULL, it is set to every widget.

scope If "", do not set any variables. If "L" create variables local to the parent

frame that called the function. If "G" create global variables.

asvector return a vector instead of a list. WARNING: if a widget variable is a true

vector or matrix, this will not work.

windowname Which window to select values from. If "" is given, it will use the most

recently active window determined from PBS.win\$windowname.

#### Value

A list (or vector) with named components, based on the variable name as the key, and the value of the associated widget as the value.

#### See Also

parseWinFile
setWinVal
clearWinVal

### Description

Imports a history list from a file, and places it as the history list identified by hisname.

### Usage

```
importPBShistory(hisname="", fname="")
```

#### Arguments

hisname The imported list is placed into the history list identified by hisname. If it

is set to "", the value from PBS.win\$action[1] will be used instead.

fname Which file to import. If it is set to "", an open file window will be displayed.

#### See Also

```
exportPBShistory
initPBShistory
promptOpenFile
```

initPBShistory

Customized History Widget Functions

### Description

The functions: initPBShistory, rmPBShistory, addPBShistory, forwPBShistory, backPB-Shistory, jumpPBShistory are made available to those who would like to use history, without using the history widget. In other words, these functions allow users to create a custom history widget.

```
initPBShistory(hisname, indexname=NULL, sizename=NULL, overwrite=TRUE)
rmPBShistory(hisname="")
addPBShistory(hisname="")
forwPBShistory(hisname="")
backPBShistory(hisname="")
jumpPBShistory(hisname="", index="")
clearPBShistory(hisname="")
```

The name of the history "list" to manipulate. If it is omitted, the function hisname will use the value of PBS.win\$actions[1] as the history name. This allows functions to be called directly from the window description file. With the exception of initPBShistory which must be called before createWin(). indexname The name of the index entry widget in the window description file. If it is NULL, then the current index feature will be disabled. The name of the current size entry widget. If it is NULL, then the current sizename

size feature will be disabled.

index Index to the history item. if it is left as "", then value will be extracted

from the widget identified by indexname

If set to true, history (matching hisname) will be cleared. Otherwise, the overwrite

two will be merged.

#### **Details**

PBS Modelling includes a pre-built history widget designed to collect interesting choices of GUI variables so that they can be redisplayed later, rather like a slide show.

Normally, a user would invoke a history widget simply by including a reference to it in the description file. However, PBS Modelling includes some support functions for customized applications.

To create a customized history, each button must be described separately in the window description file rather than making reference to the history widget.

The history "List" must be initialized before any other functions may be called. The use of a unique history name (hisname) is used to associate a unique history session with the supporting functions.

The indexname and sizename arguments correspond to the given names of entry widgets in the window description file which will be used to display the current index, and total size of the list. the indexname entry widget can also be used by jumpPBShistory to retrieve an index to jump to.

#### See Also

importPBShistory exportPBShistory

#### **Examples**

#Example of creating a custom history widget that saves values #whenever the "plot" button is pressed, and updates the plot when #back or next is pushed. A custom history is needed to achieve this #functionality since the normal history widget does not update plots require(PBSmodelling)

```
#create a window Description to be used with createWin using astext=TRUE
#PS: watch out for escaping special characters which
     must be done twice (first for R, then PBSmodelling)
winDesc <- '
window title="Custom History"
vector names="a b k" labels="a b points" font="bold" values="1 1 1000" function=myPlot
grid 1 3
 button function=myHistoryBack text="<- Back"</pre>
 button function=myPlot text="Plot"
 button function=myHistoryForw text="Next ->"
grid 2 2
 label "Index"
  entry name="myHistoryIndex" width=5
 label "Size"
  entry name="myHistorySize" width=5
#convert text into vector with each line represented as a new element
winDesc <- strsplit(winDesc, "\n")[[1]]</pre>
#custome functions required to update plots after restoring history values
myHistoryBack <- function() {</pre>
 backPBShistory("myHistory")
 myPlot(saveVal=FALSE) #show the plot with saved values
myHistoryForw <- function() {</pre>
 forwPBShistory("myHistory")
 myPlot(saveVal=FALSE) #show the plot with saved values
}
myPlot <- function(saveVal=TRUE) {</pre>
  #save all data whenever plot is called (directly)
  if (saveVal)
    addPBShistory("myHistory")
  getWinVal(scope="L")
  tt <- 2*pi*(0:k)/k;
  x \leftarrow (1+\sin(a*tt)); y \leftarrow \cos(tt)*(1+\sin(b*tt));
  plot(x, y)
}
initPBShistory("myHistory", "myHistoryIndex", "myHistorySize")
createWin(winDesc, astext=TRUE)
```

openFile

Open Files With Associated Program

### Description

openFile attempts to open a file, based off the command set in the PBS.options\$openfile list. If PBS.options\$openfile[[extension]] is set openFile will replace all occurrences of "%f" with the absolute path of the filename, and then execute the command. Otherwise, if no command is set, shell.exec() will be used.

# Usage

```
openFile(fname)
```

### **Arguments**

fname

filename to be opened.

# Examples

```
## Not run:
#setup firefox to open .html files
PBS.options$openfile$html='"c:/Program Files/Mozilla Firefox/firefox.exe" %f'
openFile("foo.html")
## End(Not run)
```

pad0

Pads Numbers with leading zeroes

### Description

Takes numbers, converts them to integers then text, and pads them with leading zeroes.

#### Usage

```
pad0(x, n, f = 0)
```

#### Arguments

x Vector of numbers

n Length of padded integer

f Factor of 10 to expand x by

#### Value

A character vector representing **x** with leading zeros.

parseWinFile

Convert Window Description File into a List

# Description

Parses a Window Markup file into the list format expected by createWin()

### Usage

```
parseWinFile(fname, astext=FALSE)
```

### **Arguments**

fname file name of window markup file.

astext if true, fname is interpreted as a vector of strings. with each element

representing a line of the source of a window description file

#### Value

A list representing a parsed window description file that can be directly passed to createWin.

### Note

All widgets are encapsulated into a 1 column by N row grid.

#### See Also

```
createWin
compileDescription
```

# Examples

```
## Not run:
x<-parseWinFile(system.file("examples/LissFigWin.txt",package="PBSmodelling"))
createWin(x)
## End(Not run)</pre>
```

pause

Pause

### Description

Pause, typically between graphics displays

### Usage

```
pause(s = "Press <Enter> to continue")
```

### Arguments

S

prompt text

**PBSmodelling** 

PBS Modelling

# Description

PBS Modelling provides software to facilitate the design, testing, and operation of computer models. It focuses particularly on tools that make it easy to construct and edit a customized graphical user interface (GUI). Although it depends heavily on the R interface to the Tcl/Tk package, a user does not need to know Tcl/Tk.

The package contains examples that illustrate models built with other R packages, including PBS Mapping, odesolve, and BRugs. It also serves as a convenient prototype for building new R packages, along with instructions and batch files to facilitate that process.

The root library directory of PBSmodelling includes a complete user guide PBSmodelling-UG.pdf. To use this package effectively, please consult the guide.

PBS Modelling comes packaged with several examples which can be accessed with the runExamples() function.

pickCol

Display Interactive Colour Selection Palette

# Description

Display an interactive colour chooser. If returnValue is true, then the equivalent hex colour value is returned, otherwise if returnValue is faluse, an intermediate GUI is used to display the hex value and no value is returned to R.

### Usage

```
pickCol(returnValue=TRUE)
```

# Arguments

returnValue If true only display the colour chooser and return the hex value. Otherwise use an intermediate GUI to display the hex value

#### Value

A hexidecimal colour value.

### See Also

testCol

plotBubbles

Construct a Bubble Plot from a Matrix

# Description

Constructs a bubble plot for a matrix z.

```
plotBubbles(z, xval = FALSE, yval = FALSE, rpro = FALSE,
cpro = FALSE, rres = FALSE, cres = FALSE, powr = 1,
clrs = c("black", "red"), size = 0.2, lwd = 2, debug = FALSE, ...)
```

Z	input matrix
xval	x-values for the columns of z. if xval=TRUE, first row contains x-values for the columns
yval	y-values for the rows of z. if yval=TRUE, first column contains y-values for the rows
rpro	if rpro=TRUE, convert rows to proportions
cpro	if cpro=TRUE, convert columns to proportions
rres	if rres=TRUE, use row residuals (subtract row means)
cres	if cres=TRUE, use column residuals (subtract column means)
powr	power transform. radii proportional to $z^{powr}$ . powr=0.5 gives bubble areas proportional to z
clrs	colours used for positive and negative values
size	size (inches) of the largest bubble
lwd	line width for drawing circles
debug	display debug information if true
	additional arguments for symbols function

# See Also

 ${\tt genMatrix}$ 

# Examples

```
plotBubbles(genMatrix(20,6))
```

plotCsum

Plots Cumulative Frequency of Data

# Description

Plots cumulative frequency of data

```
plotCsum(x, add = FALSE, ylim = c(0, 1), xlab = "Measure",
ylab = "Cumulative Proportion", ...)
```

x vector of values
add if TRUE, add cumulative frequency curve to current plot
ylim limits for y-axis
xlab label for x-axis
ylab label for y-axis
... additional arguments for plot.

### Examples

```
x <- rgamma(n=1000,shape=2)
plotCsum(x)</pre>
```

### Description

Displays the default open file prompt provided by the Operating System.

### Usage

```
promptOpenFile(initialfile="", filetype=list(c("*", "All Files")), open=TRUE)
```

### **Arguments**

initialfile file name of text file containing the list.

filetype a list of character vectors indicating what filetypes to look for. Each vector

is of length one or two, and specifies the file extension, or "\*" (for all file-types). The second element is an optional name for the file type describing

the file type.

open If True display Open file prompt, if False display Save File prompt.

#### Value

The filename and path of file selected by user.

#### See Also

promptSaveFile

### Examples

promptSaveFile

Display Save File Dialog

### Description

Displays the default save file prompt provided by the Operating System.

### Usage

```
promptSaveFile(initialfile="", filetype=list(c("*", "All Files")), save=TRUE)
```

# Arguments

initialfile file name of text file containing the list.

filetype a list of character vectors indicating what filetypes to look for. Each vector

is of length one or two, and specifies the file extension, or "\*" (for all file-types). The second element is an optional name for the file type describing

the file type.

save If True display Save file prompt, if False display Open File prompt.

#### Value

The filename and path of file selected by user.

#### See Also

```
promptOpenFile
```

#### Examples

readList

Read a List From a File

# Description

Reads in a list which was saved to a file by writeList. It can support a native R list, or PBSmodelling P format. It is detected automatically.

For information about "P" format, see writeList.

## Usage

```
readList(fname)
```

# Arguments

fname

file name of text file containing the list.

#### See Also

writeList

unpackList

resetGraph

Reset plot par Values

# Description

Reset the plot par() to default values to ensure the plot takes up the whole canvas.

# Usage

resetGraph()

### See Also

resetGraph

runExamples

Display PBS Modelling Examples

### Description

Displays an interactive demo GUI to display PBS Modelling examples.

The example source files can be found in the PBSmodelling/examples directory located in the R library.

#### Usage

runExamples()

#### **Details**

Some examples make use of external packages which must be installed in order to work correctly.

LinReg, MarkRec, and CCA Require the Brugs package.

FishRes requires the odesolve package.

FishTows requires the PBSmapping package.

setWinVal

Update Widget Values

# Description

setWinVal updates a widget with a new value. The vars argument expects a list or vector with named elements. Every element name corresponds to the widget name which will be updated with the supplied element value.

The vector and matrix widgets can be updated in several ways. If more than one name is given for the names argument then each element is treated as if it simply an entry widget. If however, a single name is given, and the value returned by getWinVal is a vector or matrix, the whole widget can be updated by passing an appropriately sized vector or widget. Alternatively each element can be updated by appending the index in square braces to the end of the name.

```
setWinVal(vars, windowname)
```

vars a list or vector with named components.

windowname Which window to select values from. If "" is given, it will use the most

recently active window determined from PBS.win\$windowname.

#### **Details**

The data widget can also be updated in the same fashion as the matrix; however, when updating a single element, a "d" must be added after the brackets. This is due to the internal coding of PBS Modeling. Example: "foo[1,1]d"

#### See Also

```
getWinVal
createWin
```

### Examples

show0

Convert Numbers Into Text With Specified Decimal Places

### Description

Return character representation of number with specified decimal places.

# Usage

```
show0(x, n, add2int = FALSE)
```

#### Arguments

x Number as scalar or vector

n Number of decimal places to show, include zeroes

add2int If TRUE, add zeroes on the end of integers

### Examples

```
frame()

#do not show decimals on integers
addLabel(0.25,0.75,show0(15.2,4))
addLabel(0.25,0.7,show0(15.1,4))
addLabel(0.25,0.65,show0(15,4))

#show decimals on integers
addLabel(0.25,0.55,show0(15.2,4,TRUE))
addLabel(0.25,0.5,show0(15.1,4,TRUE))
addLabel(0.25,0.45,show0(15,4,TRUE))
```

showArgs

Display Expected Widget Arguments

# Description

Displays the order and default values of widget arguments. The list can be shortened by specifying a single widget name. Otherwise all widgets are displayed.

### Usage

```
showArgs(widget="")
```

### **Arguments**

widget

Only displays information about this one widget

testCol

Display Colour Palette

# Description

Provides a testing bed for displaying colours on a graph. Colours can be specified in any of 3 different ways: 1) by a color name, 2) by a hexidecimal color code created by rgb() or 3) by an index into the color palette.

```
testCol(cnam=colors()[1:20])
```

cnam

vector of colour names to display

#### See Also

pickCol

# Examples

```
testCol(c("sky","fire","sea","wood"))

testCol(c("plum","tomato","olive","peach","honeydew"))

testCol(rainbow(63))

#display all colours set in the colour palette
testCol(1:length(palette()))

#they can even be mixed
testCol(c("#9e7ad3", "purple", 6))
```

testLty

 $Display\ Line\ Types$ 

# Description

Displays line types available

# Usage

```
testLty(newframe = TRUE)
```

### Arguments

newframe

if true, create a new blank frame, otherwise overlay current frame

testLwd Display Line Widths

# Description

User can specify particular ranges for lwd. Colours can also be specified and are internally repeated as necessary.

#### Usage

```
testLwd(lwd=1:20, col=c("black","blue"), newframe=TRUE)
```

### **Arguments**

lwd line widths to test colours to test

newframe if true, create a new blank frame, otherwise overlay current frame

# Examples

```
testLwd(3:15,col=c("salmon","aquamarine","gold"))
```

testPch

Display Plotting Symbols

### Description

Allows the user to specify particular ranges (increasing continuous integer) for pch.

#### Usage

```
testPch(pch=1:100, ncol=10, grid=TRUE, newframe=TRUE, bs=FALSE)
```

### Arguments

pch symboles to view

ncol number of columns (can only be 2, 5, or 10). Most sensibly this is set to

10

grid if T, grid is plotted for visual aid

newframe if T, reset the graph, otherwise overlay on top of the current graph

bs if T, shows backslash characters used in text statements. (e.g.  $30\272C =$ 

30°C)

#### Examples

testPch(123:255)
testPch(1:25,ncol=5)
testPch(41:277,bs=TRUE)

testWidgets

Displays Sample GUIs and Source Code

### Description

Displays an interactive demo GUI to provide several sample GUIs along with window description source code. It is possible to modify the sample source code in the provided text box which can then be recreated with the button below.

The window description source files can be found in the PBSmodelling/testWidgets directory located in the R library.

#### Usage

testWidgets()

#### **Details**

The following are the widgets and default values supported by PBS Modelling. See Appendix B for a detailed description.

```
button text="Calculate" font="" width=0 function="" action="button" sticky="" padx=0 pady=0

check name checked=FALSE text="" font="" function="" action="check" sticky="" padx=0 pady=0

data nrow ncol names modes="numeric" rowlabels="" collabels="" rownames="X" colnames="Y" font="" values="" byrow=TRUE function="" enter=TRUE action="data" width=6 sticky="" padx=0 pady=0

entry name value="" width=20 label="" font="" function="" enter=TRUE action="entry" mode="numeric" sticky="" padx=0 pady=0

grid nrow=1 ncol=1 toptitle="" sidetitle="" topfont="" sidefont="" byrow=TRUE borderwidth=1 relief="flat" sticky="" padx=0 pady=0

history name="default" archive=TRUE sticky="" padx=0 pady=0
```

label text="" font="" sticky="" padx=0 pady=0

```
matrix nrow ncol names rowlabels="" collabels="" rownames="" colnames=""
font="" values="" byrow=TRUE function="" enter=TRUE action="matrix"
mode="numeric" width=6 sticky="" padx=0 pady=0
menu nitems=1 label font=""
menuitem label font="" function action="menuitem"
null padx=0 pady=0
radio name value text="" font="" function="" action="radio"
mode="numeric" sticky="" padx=0 pady=0
slide name from=0 to=100 value=NA showvalue=FALSE orientation="horizontal"
function="" action="slide" sticky="" padx=0 pady=0
slideplus name from=0 to=1 by=0.01 value=NA function="" enter=FALSE
action="slideplus" sticky="" padx=0 pady=0
text name height=8 width=30 edit=FALSE bg="white" mode="character"
font="" value="" borderwidth=1 relief="sunken" edit=TRUE padx=0 pady=0
vector names length=0 labels="" values="" font="" vertical=FALSE
function="" enter=TRUE action="vector" mode="numeric" width=6 sticky=""
padx=0 pady=0 window name="window" title="" vertical=TRUE
```

#### See Also

createWin, showArgs

unpackList

Unpack List Elements Into Variables

### Description

Function to make local or global variables (depending on the scope) from the named components of a list.

```
unpackList(x, scope="L")
```

x list to unpack.

scope If "L" create variables local to the parent frame that called the function. If

"G" create global variables.

### Value

A character vector of unpacked variable names.

### See Also

readList

# Examples

```
x<-list(a=21,b=23)
unpackList(x)
print(a)</pre>
```

view

Display First n Rows of an Object

# Description

Views first n rows of a data frame or matrix or first n elements of a vector or list. All other objects are simply reflected.

# Usage

```
view(obj, n = 5)
```

# Arguments

obj Object to view

n First n elements of the obj to view

writeList

Write a List to a File

### Description

Writes an ASCII text representation in either "D" format, or "P" format. The "D" format makes use of dput and dget and produces an R representation of the list. The "P" format attempts to represents a simple list in an easy to read format.

### Usage

```
writeList(x, fname, format="D", comments="")
```

## Arguments

fname file name of text file containing the list

x list to write out

format of file to create. "P" or "D"

comments vector of character strings to be used as initial comments in the file

#### Details

The "D" format is equivalent to using dput and dget, which supports all R objects.

The "P" format only supports named lists of vectors, matrices, and data-frames. Nested lists are not supported. In the simplest form, the "P" format consists of a named list element prefixed with a dollar sign (\$) on a single line with data on the following line(s). All following data will be until a new variable name is found (denoted by \$), or the end of a file is reached.

If multiple lines of data are used, then the data is treated as a matrix, and all rows must have the same amount of values (separated by whitespace).

It is possible to specify more advanced options by including a line with two dollar signs (\$\$) that follows on the next immediate line after the variable name declaration. These options allow names for vectors, and colnames, as well as data.frames objects. For complete details see the PBS modelling PDF.

#### Note

"P" format only supports a list of vectors, or matrices, but cannot support sub-lists. However "D" format supports all R objects

#### See Also

readList

dput