Building an R Package in Windows

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1 Introduction

Building a package in R is intimidating at first. R is a very programmable language, but many users write scripts and source them each time they load R due to the hurdles required to build a package. Also, though many people have gone through the process, in many cases one has to experience it for themselves before it becomes easy. In this document, we attempt to make the transition a bit easier.

The first natural question to ask is: why build a package in the first place? Depending on the situation, there are a few reasonable answers. If you have multiple functions, then a package allows you to load and use all of them in one step rather than multiple steps. You also have convenient documentation ready and can use it with the conventional R help function "?". Perhaps most importantly, other people can download and install your functions and use them.

Throughout this document we will assume that you are using a Windows machine. The sequence on a Mac should not be too different.

2 Building the Package for the First Time

The first time you build your package is different from all of the others. This time you will set up the help files and the DESCRIPTION and NAMESPACE files. To initiate this process, you need to have all of the functions for your initial build sourced in your current session of R. In addition, if you plan to include a data set in your package, have this loaded as well. Make sure that you only have functions and data loaded in the current session that you want to have included in the package. For the purposes of this document, we will call our package test. It will include a function called descrip(), which produces descriptive statistics for a variable or data set, and a data set called mri, from Scott S. Emerson M.D., Ph.D., from http://www.emersonstatistics.com/datasets/mri.txt.

2.1 package.skeleton()

Making help files ...

Once you have everything loaded, run the function

```
package.skeleton("test")

Which will print the following statements:

Creating directories ...

Creating DESCRIPTION ...

Creating NAMESPACE ...

Creating Read-and-delete-me ...

Saving functions and data ...
```

Done.

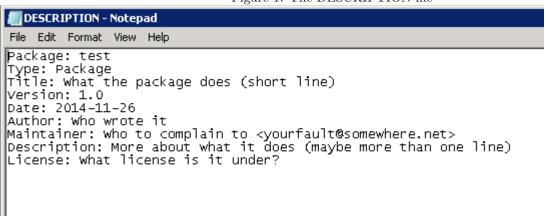
Further steps are described in './test/Read-and-delete-me'.

The file will be created in the same directory where your R folder is. This is usually in My Documents by default. You can specify a directory to place the package files in with the path argument.

2.2 The DESCRIPTION file

The first file mentioned in the package.skeleton() output is the DESCRIPTION file. This file must be edited (can be edited using Notepad or some other text editor) before the package can be compiled. If you pull it up in Notepad, it will look like this: The lines that need to be edited are the Title, Author, Maintainer, Description, and

Figure 1: The DESCRIPTION file

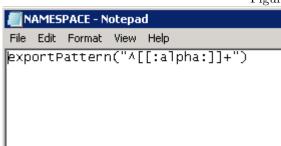


License. If your package depends on another package (say you write a function that can handle Surv variables, for survival analysis, and thus depends on the survival package) the convention is to add a line titled Imports after the Description line. In the case where we needed the survival package, we would write "Imports: survival". This line ensures that if the required packages are not loaded, R will throw an error when the user attempts to load test.

2.3 The NAMESPACE file

The second file created is the NAMESPACE file. This file controls which functions are imported or exported by the package. Initially, it looks like Figure ??. This initial line tells the package to export all of the files that have

Figure 2: The NAMESPACE file



a corresponding help file. If you need to export additional functions, write "export(functionname)". If you need to import functions (for example, from the survival package as above) write "import(functionname)" to import specific functions or "import(packagename)" to import all functions from a package. This file and the DESCRIPTION file MUST be in the package folder, or else it cannot be built.

2.4 The Read-and-delete-me file

The last file created in the main package folder is the Read-and-delete-me file (Figure ??). It tells you the most basic items to complete before your package can be built. This file MUST be deleted before the package can be

Figure 3: The Read-and-delete-me file

```
File Edit Format View Help

* Edit the help file skeletons in 'man', possibly combining help files for multiple functions.

* Edit the exports in 'NAMESPACE', and add necessary imports.

* Put any C/C++/Fortran code in 'src'.

* If you have compiled code, add a useDynLib() directive to 'NAMESPACE'.

* Run R CMD build to build the package tarball.

* Run R CMD check to check the package tarball.

Read "Writing R Extensions" for more information.
```

built.

2.5 The man files

Navigate to the package directory. Notice that there is a subfolder titled "man". This contains the skeleton of the help manual files for each function in your package. In our case, this folder contains three files: descrip, mri, and test-package. All three of these files must be edited before we can build. If you open up the descrip file, the first few lines will look like Figure ??. The second item in the file is titled alias. For those who have many functions

Figure 4: Sample Manual File

```
\name{descrip}
     \alias{descrip}
    %- Also NEED an '\alias' for EACH other topic documented here.
 3
 4
    \title{
        ~~function to do ... ~~
 5
    %%
 6
 7
    \description{
        --- A concise (1-5 lines) description of what the function does. ---
 8
10
    \usage{
    descrip(..., strata = NULL, subset = NULL, probs = c(0.25, 0.5, 0.75), geomInclude
    = FALSE, replaceZeroes = FALSE, restriction = Inf, above = NULL, below = NULL,
    labove = NULL, rbelow = NULL, lbetween = NULL, rbetween = NULL, interval = NULL,
    linterval = NULL, rinterval = NULL, lrinterval = NULL, version = FALSE)
12
    %- maybe also 'usage' for other objects documented here.
13
14
    \arguments{
15
       \item{\dots}{
16
    %%
            ---Describe \code{\dots} here---
17
18
       \item{strata}{
19
    %%
            ~~Describe \code{strata} here~~
20
       \item{subset}{
21
22
    %%
             ~Describe \code{subset} here~~
23
    }
       \item{probs}{
24
25
             ~Describe \code{probs} here~~
26
27
      \item{geomInclude}{
28
             ~Describe \code{geomInclude} here~~
29
    }
30
       \item{replacezeroes}{
31
    9696
            ---Describe \code{replaceZeroes} here---
32
33
      \item{restriction}{
```

in their package, but only want the user to have access to a few of them, this line allows the package to pass the check command. If you only want the user to use a few functions, you can delete the man files for your hidden functions and add them as aliases to the man file that they are associated with. For example, if we had a function testDescrip that performed a necessary part of the descrip function but was located in a separate file, we could delete the man file for testDescrip and put textDescrip in the alias line of the man file for descrip.

The usage line shows all of the default values in your R file for descrip. This usually does not need to be edited, but all of the other lines must be.

3 Building the Package

Once you have edited all of the man files, you are almost ready to build the package. First, however, you must install Rtools. This is a package for building R packages, and is available at http://cran.r-project.org/bin/windows/Rtools/. Once you download and run the installer, a wizard window will pop up and guide you through the process. Accepting all of the defaults is fine (unless you want to build R itself, which we generally don't). After Rtools is installed, however, you are not done. For this to work correctly, your PATH variable must be set correctly. To find the PATH variable, go to Control Panel -> System -> Advanced System Settings -> Advanced Tab. Then go to Environment Variables and find PATH, and click Edit. Check to make sure that there are three entries for Rtools:

c:\Rtools\bin;c:\Rtools\perl\bin;c:\Rtools\MinGW\bin;

Make sure that there are no spaces in-between entries. If R is not already in the path, make sure to add an entry for it. In this test case, R is located in

C:\Users\brianw26\R\R-3.1.1\bin\x64

so we add this to the PATH variable. Also, if you do not already have MikTex installed, it is recommended to do so and is located at http://miktex.org/download (this helps create nice pdfs for the help files).

Now we can build the package. First, open a command prompt session. Navigate to the directory containing the package. Then type

R CMD build packagename

This will create a tar.gz file. Next, type

R CMD check packagename_versionnumber.tar.gz

Where the text after check is the name of the tar.gz file created in the first step. This is where most of the problems come in. The check function makes sure that your package will run and that all of the code works properly (especially if you have examples in your man files). Considerable amounts of time may be spent making sure all of the errors go away. However, you may see other flags as well. These do not need to all be taken care of, even if you are submitting to CRAN. Though it is ideal that all of the flags are taken care of, if you have certain functions or man files which raise flags and can justify them to CRAN, you are okay. The flags which are fine are WARNING flags, but only in certain parts (check CRAN documents for which are okay to have). If there are no errors, last run

R CMD INSTALL --build packagename

which will create a .zip file which can be downloaded and installed by Windows users. A Mac user must build the package on their machine for the proper tar.gz file to be built.

4 Updating the Package

After the initial build of the package, you may add functions or edit existing functions. Then it is necessary to update the package! First, you must edit the DESCRIPTION file to reflect a new version number - while this is not strictly necessary, it is good practice. Second, make sure that you have a man file for each new function you create, or that you update the man file for an updated existing function. Now that you have a folder for the package, simply place the ".R" files in the R subfolder of the package and they will be included when the package is updated. Last, run the three commands from the command prompt in the previous section. This will build the new version of the package!