

Making an R package

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Why develop your own packages?

R packages are an ideal way to package and distribute R code and data for re-use by others.

- **Community:** CRAN / packages part of R success
- **cross-platform / cross-OS:** packages are portable
- **getting R code to collaborators:** distribution
- **reproducibility:** aided greatly by identifiable package versions
- **version control:** learn about git (or svn)

Part I

Package structure

Software Prerequisites

There are two main prerequisites for building R packages:

1. GNU software development tools including a C/C++ compiler; and
2. LaTeX for building R manuals and vignettes.
3. Rtools (Windows)

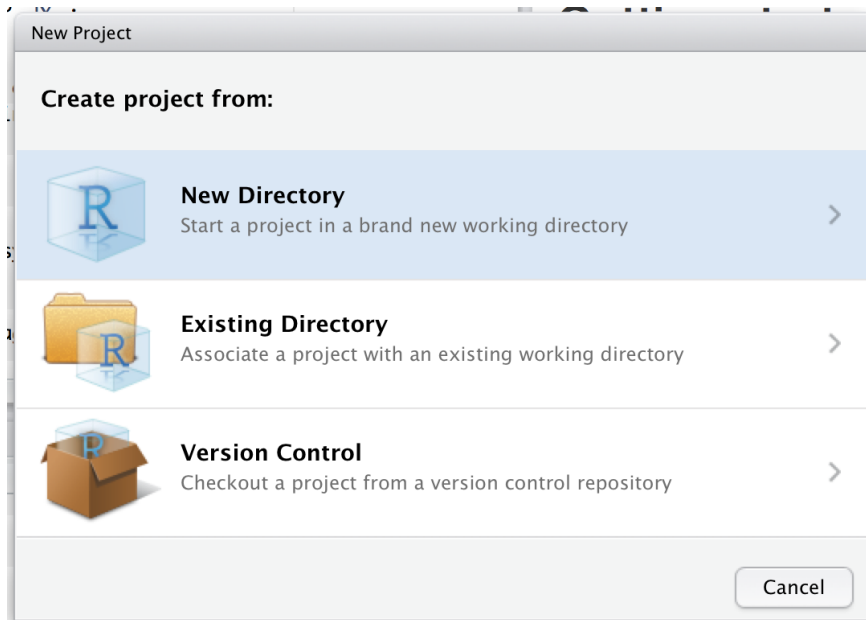
Reference: 1. R packages by Hadket wickham. See also at <http://r-pkgs.had.co.nz/>

Requirements for a name

There are three formal requirements: - the name can only consist of letters, numbers and periods, i.e., .; - it must start with a letter; - it cannot end with a period.

Strategies for creating a name

- Pick a unique name you can easily Google.
- Avoid using both upper and lower case letters.
- Find a word that evokes the problem and modify it so that it's unique:
 - plyr is generalisation of the apply family, and evokes pliers.
 - knitr (knit + r) is “neater” than sweave (s + weave).
 - testdat tests that data has the correct format.
- Use abbreviations:
 - Rcpp = R + C++ (plus plus)
 - lvplot = letter value plots.
- Add an extra R:
 - stringr provides string tools.
 - tourr implements grand tours (a visualisation method).
 - gistr lets you programmatically create and modify GitHub gists.



Creating a New Package

1. Click File | New Project.
2. Choose “New Directory”:

3. Then “R Package”:

4. Then give your package a name and click “Create Project”:

The package you just created

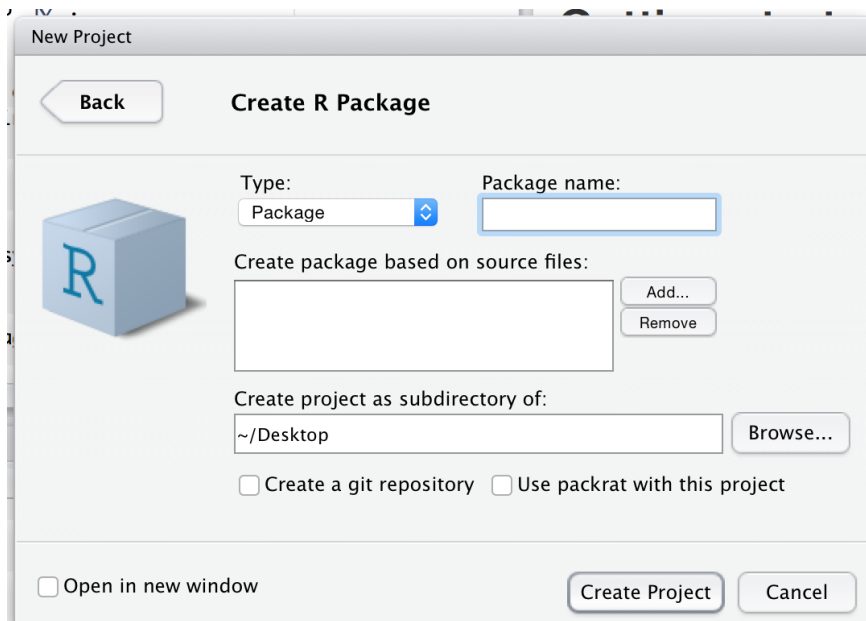
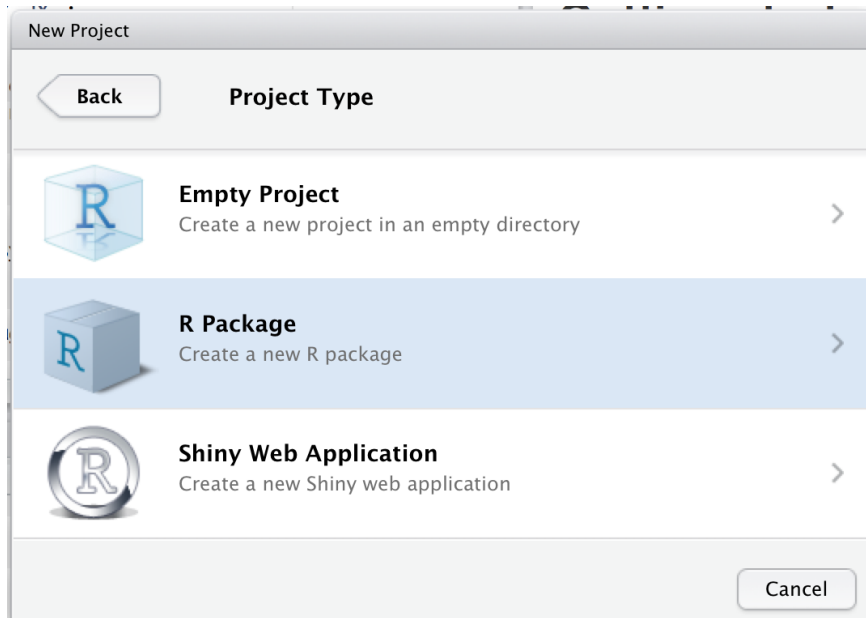
The smallest usable package, one with three components:

1. An R/ directory.
2. A basic DESCRIPTION file.
3. A basic NAMESPACE file.

It will also include an RStudio project file, pkgname.Rproj, that makes your package easy to use with RStudio.

Five states a package across its lifecycle:

1. Source
2. Bundled
3. Binary
4. Installed
5. In-memory



Source packages

- The development version of a package that lives on your computer.
- A source package is just a directory with components like R/, DESCRIPTION

Bundled packages

- A **bundled** package is a package that has been compressed into a single file.
- Package bundles in R use the extension .tar.gz.
- The main differences between an uncompressed bundle and a source package are:
 - Vignettes are built so that you get HTML and PDF output instead of Markdown or LaTeX input.
 - Your source package might contain temporary files used to save time during development, like compilation artefacts in src/. These are never found in a bundle.
 - Any files listed in .Rbuildignore are not included in the bundle.

- **.Rbuildignore** prevents files in the source package from appearing in the bundled package.
- It allows you to have additional directories in your source package that will not be included in the package bundle.
- A typical example

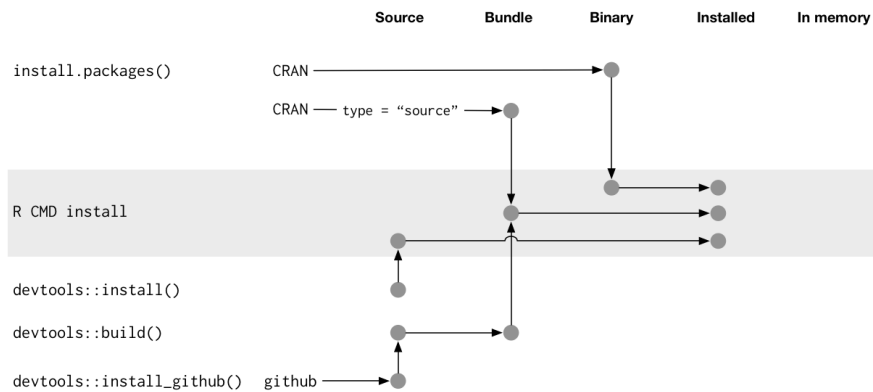
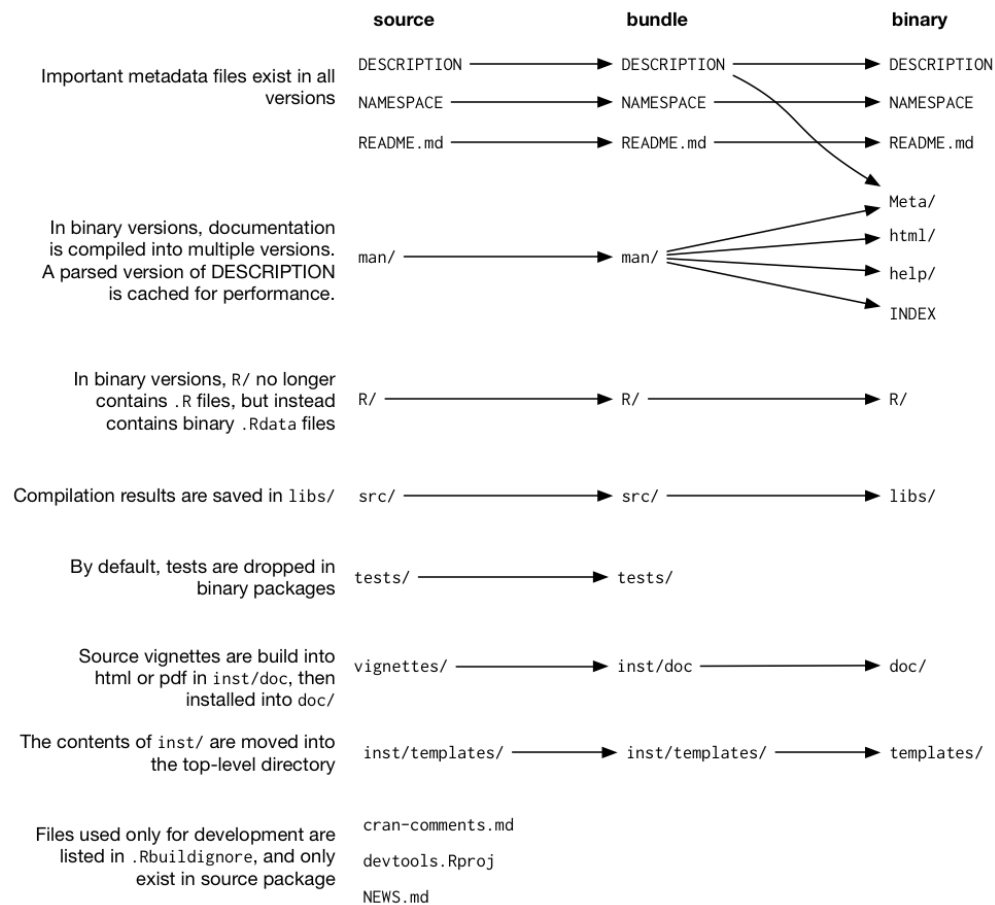
```
^.*\.Rproj$      # Automatically added by RStudio,  
^\.Rproj\.user$  # used for temporary files.  
^README\.Rmd$    # An Rmarkdown file used to generate README.md  
^cran-comments\.md$ # Comments for CRAN submission  
^NEWS\.md$       # A news file written in Markdown  
^\.travis\.yaml$ # Used for continuous integration testing with travis
```

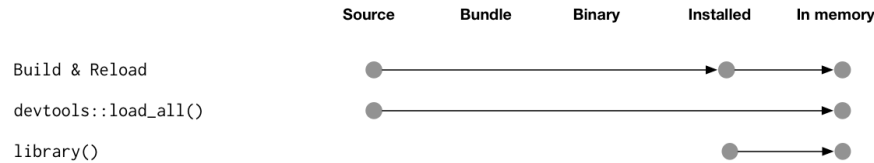
Binary packages

- If you want to distribute your package to an R user who doesn't have package development tools, you will need to make a **binary** package.
- Binary packages are platform specific:
 - Mac binary packages end in .tgz
 - Windows binary packages end in .zip

Installed packages

- An **installed** package is just a binary package that has been decompressed into a package library.
- The following diagram illustrates the many ways a package can be installed.
- An easier way: using **RStudio**





In memory packages

- To use a package, you must load it into memory.
- To use it without providing the package name, you need to attach it to the search path.

```
# Automatically loads devtools
devtools::install()

# Loads and _attaches_ devtools to the search path
library(devtools) # or
require(devtools)
install()
```

- `library()` is not useful when you're developing a package because you have to install the package first.

What is a library?

- A library is simply a directory containing installed packages.
- Almost every one has at least two libraries:
 - one for packages you're installed
 - one for the packages that come with every R installation (like base, stats, etc).

```
.libPaths()
```

```
## [1] "C:/Users/wench/Documents/R/win-library/3.6"
## [2] "C:/Program Files/R/R-3.6.1/library"
```

```
lapply(.libPaths(), dir)[[2]]
```

```
## [1] "base"      "boot"      "class"     "cluster"
## [5] "codetools" "compiler"  "datasets"  "foreign"
## [9] "graphics"  "grDevices" "grid"      "KernSmooth"
## [13] "lattice"   "MASS"      "Matrix"    "methods"
## [17] "mgcv"      "nlme"      "nnet"      "parallel"
## [21] "rpart"     "spatial"   "splines"   "stats"
## [25] "stats4"    "survival"  "tcltk"     "tools"
## [29] "translations" "utils"
```

```
lapply(.libPaths(), dir)[[1]]
```

```
## [1] "abind"      "AMIAS"      "arules"
## [4] "arulesViz"  "askpass"    "assertthat"
## [7] "backports"  "base64enc"  "BeSS"
## [10] "BH"         "BiocManager" "BiocVersion"
## [13] "bitops"     "bootstrap"  "brew"
## [16] "broom"      "callr"      "car"
```

## [19]	"carData"	"caTools"	"cdcsis"
## [22]	"cellranger"	"cghAMIAS"	"cghFLasso"
## [25]	"cli"	"clipr"	"clisymbols"
## [28]	"cmprsk"	"coin"	"colorspace"
## [31]	"commonmark"	"conquer"	"corpcor"
## [34]	"corrplot"	"covr"	"cowplot"
## [37]	"crayon"	"crosstalk"	"curl"
## [40]	"data.table"	"dendextend"	"DEoptimR"
## [43]	"desc"	"devtools"	"digest"
## [46]	"diptest"	"dlstats"	"DNAcopy"
## [49]	"doParallel"	"dplyr"	"DT"
## [52]	"ellipsis"	"Epi"	"etm"
## [55]	"evaluate"	"fansci"	"farver"
## [58]	"fastmap"	"fdrtool"	"flexmix"
## [61]	"FNN"	"forcats"	"foreach"
## [64]	"formatR"	"fpc"	"freeknotsplines"
## [67]	"fs"	"fusedBeSS"	"futile.logger"
## [70]	"futile.options"	"gclus"	"gdata"
## [73]	"GeneNet"	"generics"	"genlasso"
## [76]	"GGally"	"ggdendro"	"ggforce"
## [79]	"ggformula"	"ggplot2"	"ggpubr"
## [82]	"ggrepel"	"ggsci"	"ggsignif"
## [85]	"ggstance"	"gh"	"git2r"
## [88]	"glmnet"	"glmulti"	"glue"
## [91]	"gplots"	"gridExtra"	"gtable"
## [94]	"gtools"	"gurobi"	"haven"
## [97]	"hdi"	"hexbin"	"highr"
## [100]	"hms"	"htmltools"	"htmlwidgets"
## [103]	"httpuv"	"httr"	"igraph"
## [106]	"ini"	"ISLR"	"iterators"
## [109]	"jiebaR"	"jiebaRD"	"jpeg"
## [112]	"jsonlite"	"kernlab"	"knitr"
## [115]	"ks"	"l0tf"	"labeling"
## [118]	"lambda.r"	"lars"	"lassoshooting"
## [121]	"later"	"latticeExtra"	"lazyeval"
## [124]	"leaflet"	"leaflet.providers"	"libcoin"
## [127]	"lifecycle"	"limSolve"	"linprog"
## [130]	"lme4"	"lmtest"	"longitudinal"
## [133]	"lpSolve"	"magick"	"magrittr"
## [136]	"maptools"	"markdown"	"MatrixModels"
## [139]	"matrixStats"	"mclust"	"memoise"
## [142]	"mime"	"minqa"	"mixtools"
## [145]	"modeltools"	"modSaRa"	"mosaic"
## [148]	"mosaicCore"	"mosaicData"	"MrBeSS"
## [151]	"MTE"	"multcomp"	"multicool"
## [154]	"munSELL"	"mvtnorm"	"ncvreg"
## [157]	"nlme"	"nloptr"	"numDeriv"
## [160]	"openssl"	"openxlsx"	"ordinal"
## [163]	"parcor"	"pbkrtest"	"pense"
## [166]	"perry"	"PGEE"	"pillar"
## [169]	"pkgA"	"pkgbuild"	"pkgconfig"
## [172]	"pkgload"	"plogr"	"plotly"
## [175]	"plyr"	"PMA"	"png"
## [178]	"polyclip"	"polynom"	"ppls"

## [181] "prabclus"	"praise"	"prettyunits"
## [184] "processx"	"progress"	"promises"
## [187] "pryr"	"ps"	"purrr"
## [190] "qap"	"quadprog"	"quantmod"
## [193] "quantreg"	"R.matlab"	"R.methodsS3"
## [196] "R.oo"	"R.utils"	"R6"
## [199] "randomForest"	"raster"	"rcmdcheck"
## [202] "RColorBrewer"	"Rcpp"	"RcppArmadillo"
## [205] "RcppEigen"	"readr"	"readxl"
## [208] "registry"	"rematch"	"rematch2"
## [211] "remotes"	"reshape"	"reshape2"
## [214] "rex"	"rio"	"rlang"
## [217] "rlmDataDriven"	"rmarkdown"	"robustbase"
## [220] "robustHD"	"robustsubsets"	"robustsubsets-master"
## [223] "roxygen2"	"rprojroot"	"rrpack"
## [226] "rrr"	"RSAVS"	"rstatix"
## [229] "rstudioapi"	"rversions"	"sandwich"
## [232] "scales"	"scalreg"	"scatterplot3d"
## [235] "secure"	"segmented"	"SemiPar"
## [238] "seriation"	"sessioninfo"	"shiny"
## [241] "SIS"	"slam"	"snow"
## [244] "snowfall"	"sourcetools"	"sp"
## [247] "SparseM"	"spikeslab"	"stringi"
## [250] "stringr"	"sys"	"testthat"
## [253] "TH.data"	"tibble"	"tidyr"
## [256] "tidyselect"	"tinytex"	"tseries"
## [259] "TSP"	"tsvd"	"TTR"
## [262] "tweenr"	"ucminf"	"usethis"
## [265] "utf8"	"vcd"	"vctrs"
## [268] "venn"	"VennDiagram"	"viridis"
## [271] "viridisLite"	"visNetwork"	"wbs"
## [274] "whisker"	"withr"	"wordcloud"
## [277] "xfun"	"XML"	"xml2"
## [280] "xopen"	"xtable"	"xts"
## [283] "yaml"	"zeallot"	"zip"
## [286] "zoo"		

library() and require()

- The main difference between `library()` and `require()` is what happens if a package isn't found.
- While `library()` throws an error, `require()` prints a warning message and returns `FALSE`

```
library(blah)
```

```
require(blah)
```

```
## Loading required package: blah
```

```
## Warning in library(package, lib.loc = lib.loc, character.only = TRUE,
## logical.return = TRUE, : there is no package called 'blah'
```