R Packages

Statistical Programming

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What are R packages?

R packages are just a collection of files (R code, compiled code, data, documentation, etc.) that live in your library path.

When you run library(pkg) the functions (and objects) in the package's namespace are attached to the global search path.

```
dir(.libPaths())
```

```
"airports"
                                                    "archive"
                                                                          "arrayhelpers"
                                                                                                "AsioHeaders"
##
     [1] "abind"
                               "assert.that"
                                                     "av"
                                                                          "available"
                                                                                                "backports"
     [6] "askpass"
                               "base64enc"
                                                     "BayesFactor"
                                                                          "bayesplot"
                                                                                                "beeswarm"
         "base"
                               "benchmarkme"
                                                    "benchmarkmeData"
                                                                          "BH"
                                                                                                "bit"
    [16] "bench"
    [21] "bit64"
                               "blob"
                                                     "bookdown"
                                                                          "boot"
                                                                                                "branchMover"
    [26] "brew"
                               "brio"
                                                    "broom"
                                                                          "broom.mixed"
                                                                                                "bslib"
                               "callr"
                                                     "cellranger"
    [31] "cachem"
                                                                          "checklist"
                                                                                                "checkmate"
                               "chromote"
                                                    "chron"
                                                                          "class"
                                                                                               "classInt"
    [36] "cherryblossom"
    [41] "cli"
                               "clipr"
                                                    "clisymbols"
                                                                          "cluster"
                                                                                                "cmdstanr"
                                                     "collections"
    [46] "coda"
                               "codetools"
                                                                          "colorblindr"
                                                                                                "colorspace"
```

Search path

```
search()
## [1] ".GlobalEnv"
                           "package:stats"
                                                "package:graphics"
                                                                    "package:grDevices" "package:utils"
## [6] "package:datasets"
                           "package:methods"
                                                "Autoloads"
                                                                    "package:base"
 library(diffmatchpatch)
 search()
   [1] ".GlobalEnv"
                                 "package:diffmatchpatch" "package:stats"
                                                                                    "package:graphics"
                                 "package:utils"
                                                           "package:datasets"
                                                                                    "package:methods"
    [5] "package:grDevices"
    [9] "Autoloads"
                                 "package:base"
```

Loading vs attaching

"package:base"

[9] "Autoloads"

If you do not want to attach a package you can directly use functions via :: or load it with requireNamespace().

```
loadedNamespaces()
   [1] "Rcpp"
                          "grDevices"
                                            "digest"
                                                              "diffmatchpatch" "R6"
                                                                                                 "isonlite"
   [7] "magrittr"
                          "evaluate"
                                            "datasets"
                                                              "xaringan"
                                                                               "stringi"
                                                                                                 "rlang"
  [13] "utils"
                          "iquervlib"
                                            "bslib"
                                                              "graphics"
                                                                               "rmarkdown"
                                                                                                 "base"
                          "stringr"
                                            "xfun"
                                                              "vaml"
                                                                               "fastmap"
                                                                                                 "compiler"
  [19] "tools"
## [25] "stats"
                          "htmltools"
                                            "knitr"
                                                              "methods"
                                                                               "sass"
requireNamespace("forcats")
## Loading required namespace: forcats
loadedNamespaces()
                                                              "diffmatchpatch" "R6"
   [17] "Rcpp"
                          "grDevices"
                                            "digest"
                                                                                                 "isonlite"
   [7] "magrittr"
                          "evaluate"
                                            "datasets"
                                                              "xaringan"
                                                                               "stringi"
                                                                                                 "rlang"
                                            "bslib"
                                                                               "rmarkdown"
                                                                                                 "base"
  [13] "utils"
                          "iquervlib"
                                                              "graphics"
                          "tools"
                                                                               "vaml"
  [19] "forcats"
                                            "stringr"
                                                              "xfun"
                                                                                                 "fastmap"
## [25] "compiler"
                                            "htmltools"
                                                              "knitr"
                                                                               "methods"
                                                                                                 "sass"
                          "stats"
search()
    [1] ".GlobalEnv"
                                  "package:diffmatchpatch"
                                                             "package:stats"
                                                                                       "package:graphics"
    [5] "package:grDevices"
                                  "package:utils"
                                                             "package:datasets"
                                                                                       "package:methods"
```

Where to R packages come from

We've already seen the two primary sources of R packages:

CRAN:

```
install.packages("diffmatchpatch")
```

GitHub:

```
remotes::install_github("rundel/diffmatchpatch")
```

there is one other method that comes up (particularly around package development), which is to install a package from local files.

Local install:

```
R CMD install diffmatchpatch_0.1.0.tar.gz

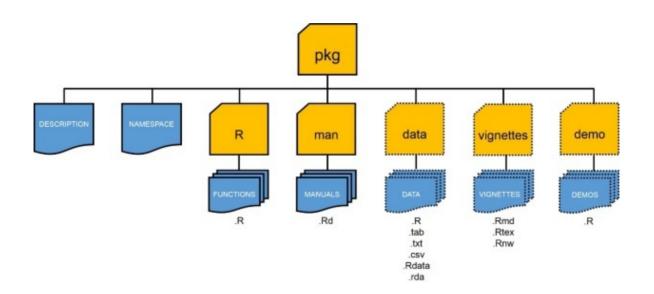
devtools::install("diffmatchpatch_0.1.0.tar.gz")
```

What is CRAN

It is the Comprehensive R Archive Network which is the central repository of R packages.

- Maintained by the R Foundation and run by a team of volunteers, ~22k packages
- Retains all current versions of released packages as well as archives of previous versions
- Similar in spirit to Perl's CPAN, TeX's CTAN, and Python's PyPI
- Some important features:
 - All submissions are reviewed by humans + automated checks
 - Strictly enforced submission policies and package requiements
 - All packages must be actively maintained and support upstream and downstream changes

Structure of an R Package



Core components

- DESCRIPTION file containing package metadata (e.g. package name, description, version, license, and author details). Also specifies package dependencies,
- NAMESPACE details which functions and objects are exported by your package
- R/ folder containing R script files (.R)
- man/ folder containing R documentation files (.Rd)

The following components are optional, but quite common:

- tests/ folder contain unit tests
- src/ folder containing code to be compiled (usually C / C++)
- data/ folder containing example data sets (exported as .Rdata via save())
- inst/ files that will be copied to the package's top-level directory when it is installed (e.g. examples or data files that don't belong in data/)

Package contents

Source Package

```
fs::dir_tree("~/Desktop/Projects/diffmatchpatch/")
```

```
~/Desktop/Projects/diffmatchpatch/
##
       DESCRIPTION
##
       LICENSE.md
##
       NAMESPACE
       NFWS.md
##
##
##
            RcppExports.R
            diff.R
##
           diffmatchpatch-package.R
##
##
            match R
##
            options.R
##
            patch.R
            print.R
##
##
       README, Rmd
       README.md
##
##
       cran-comments.md
##
       diffmatchpatch.Rproj
##
       inst
##
            include
##
            diff match patch.h
##
       man
            diff.Rd
##
##
            dmp_options.Rd
##
            match.Rd
##
            patch.Rd
```

Installed Package

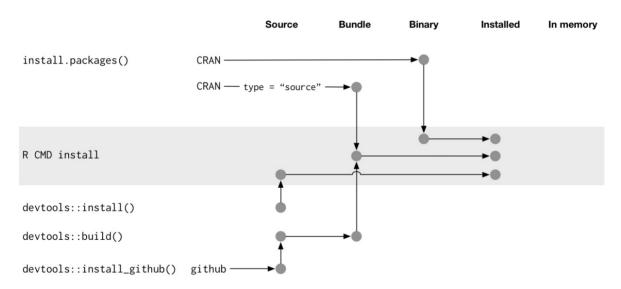
```
fs::dir_tree(system.file(package="diffmatchpatch"))
## /usr/local/lib/R/4.1/site-library/diffmatchpatch
       DESCRIPTION
##
       INDEX
##
       Meta
           Rd.rds
##
           features rds
##
            hsearch.rds
##
##
           - links.rds
##
           nsInfo.rds
##
          - package.rds
##
       NAMESPACE
##
       NFWS.md
##
##
            diffmatchpatch
##
           diffmatchpatch.rdb

    diffmatchpatch.rdx

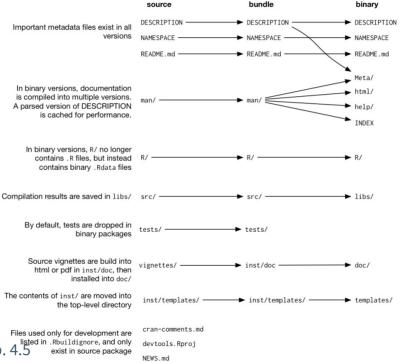
##
##
       help
##
            AnIndex
##
            aliases rds
##
           diffmatchpatch.rdb
##
            diffmatchpatch.rdx
            paths.rds
##
##
       html
##
            00Index.html
##
           R.css
```

A deeper dive on diffmatchpatch

Package Installation



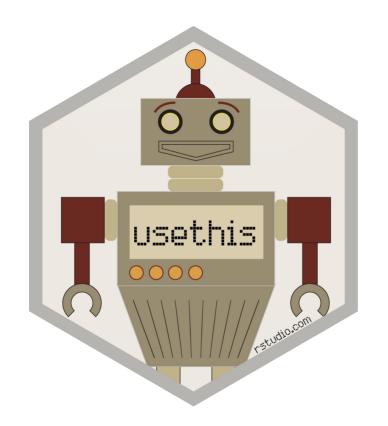
Package Installion - Files



Package development

What follows is an opinionated introduction to package development,

- this is not the only way to do thing (none of the following are required)
- I would strongly recommend using:
 - RStudio
 - RStudio projects
 - GitHub
 - usethis
 - o roxygen2



usethis

This is an immensely useful package for automating all kinds of routine (and tedious) tasks within R

- Tools for managing git and GitHub configuration
- Tools for managing collaboration on GitHub via pull requests (see pr_*())
- Tools for creating and configuring packages
- Tools for configuring your R environment (e.g. .Rprofile and .Renviron)
- and much much more

Live demo - Building a Package