Package management

Creating and maintaining project-relative package libraries with renv

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Resources

- to read more on today's topic, check out:
- Ch. 10 (Basic reprodubility: freezing packages) from Rodrigues (2023)
- the renv website
 - or CRAN documentation and vignettes therein (e.g.,: Introduction to renv)

Packages

- most open source software (like R) has a range of libraries available
 - created by other users/developers and shared for free
- the benefit of open software (besides being free) is that we don't have to wait for an updated version to be released by a company
 - and anybody can create an R package to facilitate certain tasks or fix some problem
- this is part of the reason for the success and popularity of R
 - someone else has likely created a package for some problem or need you have

CRAN packages

- the Comprehensive R Archive Network: R's central software repository
 - currently 20,888 available!
- an archive of the most recent package versions
- for a package to be included in the CRAN, it must go through a lot of tests and checks
 - any updates or changes must again be reviewed before being added to CRAN
- CRAN packages can be installed using install.packages(), as we've been doing

Developer packages

- often hosted on GitHub or GitLab, where packages are typically developed before being reviewed and added to the CRAN
 - benefit: you can make whatever you changes to your package that you like without having to pass a review on the CRAN
- since CRAN packages are often developed on GH or GL, pre-release (beta) versions will often be available on a GH repo
- packages/package versions on GH cannot be installed via install.packages()
 - we'll see later how to do this

Dependencies

- some packages are dependent on specific versions of other packages
 - if so, you will be prompted during installation to install these dependencies
 - but beware: sometimes this overwrites an existing package version you already have,
 which can break code that was written with this older version
- this is especially true because, as our projects are currently set up, we have one global package version on our computer
 - so analyses we ran 3 years ago would've used older versions of packages
 - when we update these packages for current analyses, this might disrupt the code from 3 years ago
- we'll see one (partial) solution for this problem soon
- pacman package (optional)
 - a package management tool
 - we'll use the p_load() function to replace install.packages() and library() in our worksflow
 - takes a list of packages, and checks if each package is installed already
 - if yes, the package is loaded (as with library())
 - if no, the package is installed (as with install.packages()) and then loaded (as with library())
 - only works with CRAN packages (which is all we have for now anyway)

Package versions

- packages can be updated at any time
 - if hosted on the CRAN, they newer versions are first reviewed/rigorously tested
 - if hosted on GitHub/Lab, nobody needs to check the update before publication
- if you want to check which version of a package you're using, you can run packageVersion("package")

```
packageVersion("ggplot2")
```

```
[1] '3.5.1'
```

Updating packages

- to check if a package needs updating, you can:
 - go to Tools > Check for package updates, or
 - run update.packages()
- each will tell you which packages can be updated to which versions
 - and give you the option of updating these packages

Package library

- where do all these installed packages go?
 - a folder that contains all the packages, called a library
- to find out where this (global) package library is, run .libPaths()

.libPaths()

• the output should currently produce a single file path, something like:

```
> .libPaths()
[1] "/Library/Frameworks/R.framework/Versions/4.4-arm64/Resources/library"
```

• this is the location of your global package library

Package versions and reproducibility

- we've seen that package versions and dependencies can easily break our existing code
- this means that older projects that were built using previous package versions won't be able to run if we update these packages in our global package library
 - also a problem in the future: our current code will depend on the package versions we're using today
- we need a project-relative package library that is independent of the global library
 - we'll use the renv package to do this

renv

- renv aids in maintaining reproducible environments in R projects
- available on the CRAN

Listing 1 Run in the Console

install.packages("renv")

- main benefit: creates a self-contained, independent library per R Project
 - avoids cross-library package contamination
- renv freezes and stores package versions used in a project
- but does not make a project reproducible across R versions and machines
 - that's because older package versions are not always compatible with newer computational environments

Limits of renv

renv...

...can

- keep track of packages and their versions
- create a project-specific library per R version
- automatically load/install these package versions

...cannot

- make a project reproducible across all computational environments
- load/install package versions that are incompatible with current R versions or computational environments
- guarantee full long-term reproduciblity

reny workflow

- Figure 1 visualises a project workflow with renv
- next we'll see how we use these functions to set-up and maintain a project-specific package library

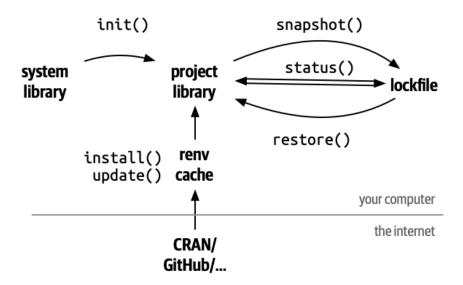


Figure 1: Source: CRAN vignette 'Introduction to renv' (all rights reserved)

Initialise project library

- run the following in the Console or in a code chunk but with #| eval: false
 - we only want to run this *once* per R Project
 - when working in an actual project, I would just run this in the console
 - for learning/documenting how to use renv, I would keep this in a code chunk with
 #| eval: false
- you should see something like this in the Console:

Listing 2 In the Console or with eval: false

```
renv::init()
```

New files

- renv::init() creates three new files or directories
 - renv.lock
 - renv/
 - .Rprofile
- explore these files/folders and see if you can figure out what they contain

renv.lock

- contains metadata about the packages and their versions that you have installed
 - this is enough metadata to re-install these package versions on a new machine
- two main components:
 - R: info on R version and list of repositories where packages were installed from
 - Packages: a record per package with necessary info for re-installation

renv/

- importantly, contains your project-relative library/
 - this is instead of using the one library on your computer
- provides us with "isolation": the package versions used in an R Project is independent of the global library
 - in other words, different R Projects can use different package versions
 - updating packages globally, or in one project, will not affect other project libraries

.RProfile

- runs whenver you (re-)start your R Project
- at this point, should contain a single line:

source("renv/activate.R")

- if you go to this R script, you'll send a lot of code
 - this essentially loads in your project library

Project library

• now if we re-run .libPaths(), we should see our project library

Listing 3 Run in the Console

.libPaths()

- > .libPaths()
- [1] "/Users/danielapalleschi/Documents/IdSL/Teaching/SoSe24/M.A./r4repro_SoSe2024/renv/librations/
- [2] "/Users/danielapalleschi/Library/Caches/org.R-project.R/R/renv/sandbox/macos/R-4.4/aarch
 - [1] is the local project library path
 - [2] is the path to a global package cache that **renv** maintains so that you don't repeatedly download packages to your machine for each project library
 - e.g., if we already have ggplot2 installed globally on our machine, whenever we
 want to add it to a project library we don't need to re-install it entirely from the
 CRAN (unless we want a different package version)

Installing more packages

- which packages are stored in renv.lock?
 - only those that are used within your project
- packages not used in your project but installed in your global library aren't included
 - to add these packages, or any other packages you want, you need to (re-)install them locally within your project
- let's install a package that we'll use later on: 1me4

```
# as usual
install.packages("lme4")
# or with the renv package
renv::install("lme4")
```

- if you already have a package on your machine (in your global library), renv will just grab it from the global cache
- if not, it will be downloaded from CRAN

Installing a new package

- let's also install a package I'm confident you don't already have on your machine (as you might've already worked with lme4 in other classes)
 - [brms] for Bayesian regression models using Stan

```
install.packages("brms")
renv::install("brms")
```

• and if we want a specific package version:

```
renv::install("brms@2.19.0")
```

Installing developer packages

- not all packages are available on the CRAN
 - we can install developer packages from GitHub or GitLab using, e.g., the install_github() function from either the remotes or devtools package (both are very common)

```
remotes::install_github("paul-buerkner/brms")
devtools::install_github("paul-buerkner/brms")
```

• or we can use renv::install()

```
# most recent version
renv::install("paul-buerkner/brms")
```

```
# a specific previous version, for which you'll need the commit ID
renv::install("paul-buerkner/brms@db6ddde90ba533cb3942bc5a62b03803773b9844")
```

Lockfile status

- you should make a habit of checking the status of your lockfile
 - you can do this by running the following:

```
renv::status()
```

• ideally, you'll usually get the following message:

```
> renv::status()
No issues found -- the project is in a consistent state.
```

• but if you've installed or updated some packages, you will get a list of any packages that are out-of-sync or haven't been stored in the lockfile (as should be our case)

Updating renv.lock file

• to update the lockfile and library, simply run:

```
renv::snapshot()
```

- you'll be given a list of changes to be made and asked if you want to proceed
 - if not problems are mentioned, then you can go ahead

Updating packages

• to update packages using renv, we can use:

```
renv::update()
# or
renv::update.packages()
```

- this will not automatically store the updated versions in the lockfile
 - to do this, include the argument lock = TRUE
- you can also use these functions to only check by including check = T

Restoring lockfile

```
renv::restore()
```

- this will restore the current project's package versions to be those stored in the lockfile
 - but only if the library was built in the same R version
 - otherwise, all packages need to be installed, and might not function the same
- useful if you
 - want to revert to the stored package versions
 - want to run your project on another computer (e.g., a collaborator)

Additional packages

- some other packages that can be useful for package management or reproducibility
- groundhog: version control for CRAN, GitHub, and GitLab packages
 - uses groundhog.library() instead of library() to load packages
 - can take a list of libraries (or an object which contains such a list) and a date as arguments
 - will then install the package versions that were available at the given date
- issues can arise when package versions were built on a previous version of R, and are no longer supported
 - this can cause the installation to fail (just like with renv)

Session Info

- whether you're using renv or not, always end a script with sessionInfo()
 - with dynamic reports: this will print out the package versions used to produce the output
 - in R: you can save the info as an object and save it as an RDS file
 - * or run it, copy-and-paste the output in the script, and comment it all out

sessionInfo()

```
R version 4.4.0 (2024-04-24)
Platform: aarch64-apple-darwin20
Running under: macOS Ventura 13.2.1
Matrix products: default
BLAS:
        /Library/Frameworks/R.framework/Versions/4.4-arm64/Resources/lib/libRblas.0.dylib
LAPACK: /Library/Frameworks/R.framework/Versions/4.4-arm64/Resources/lib/libRlapack.dylib;
locale:
[1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8
time zone: Europe/Berlin
tzcode source: internal
attached base packages:
[1] stats
              graphics grDevices datasets utils
                                                      methods
                                                                base
loaded via a namespace (and not attached):
 [1] digest_0.6.35
                       fastmap_1.2.0
                                         xfun_0.44
                                                           magrittr_2.0.3
 [5] knitr_1.47
                       htmltools_0.5.8.1 rmarkdown_2.27
                                                           cli_3.6.2
 [9] renv_1.0.7
                       compiler_4.4.0
                                       rprojroot_2.0.4
                                                           here_1.0.1
[13] rstudioapi_0.16.0 tools_4.4.0
                                         evaluate_0.23
                                                           Rcpp_1.0.12
[17] yaml_2.3.8
                       magick_2.8.3
                                         rlang_1.1.4
                                                           jsonlite_1.8.8
```

References

Rodrigues, B. (2023). Building reproducible analytical pipelines with R.