Reproducible environments

Jim Hester RStudio





https://environments.rstudio.com/

No Strategy

Awareness

Shared Baseline

Record & Restore

Validated

scary upgrades
no sharing
old stuff is broken

reprex
sessioninfo()

frozen repo

renv

custom tests

No Strate

Awareness

Baseline

Record & Restore

Validated

scary upgrades no sharing old stuff is broken

reprex sessioninfo() site library frozen repo

renv

internal repo custom tests

devtools::session_info()

Better than utils::sessionInfo() for development packages

Useful for bug reports and colophons

No Strategy

Awarene

Shared Baseline

Restore

Validated

scary upgrades no sharing old stuff is broken

reprex sessioninfo() site library frozen repo

renv

internal repo custom tests

RStudio Package Manager checkpoints

MRAN

Site library

No Strategy

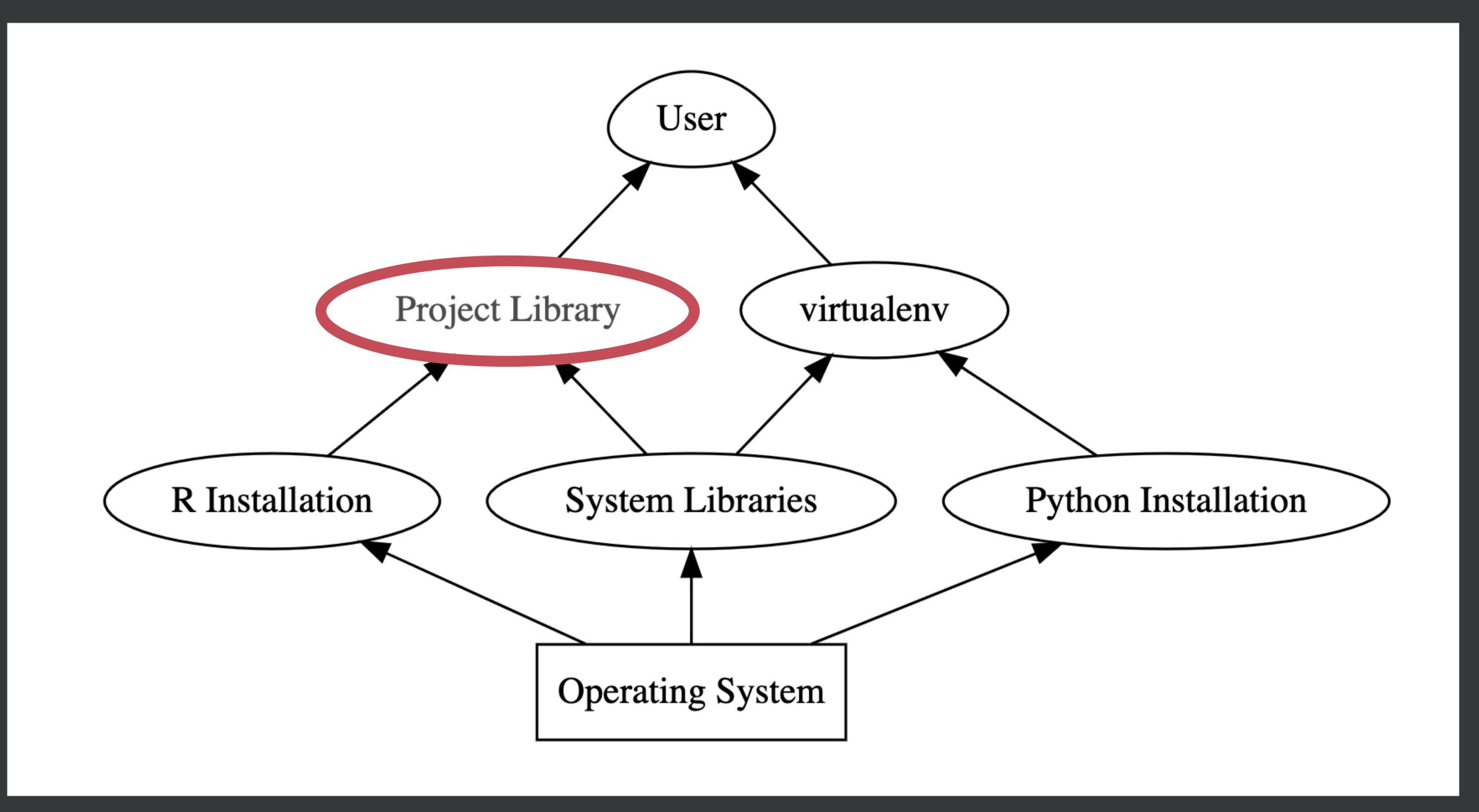
Awareness

Shared Record & Restore

scary upgrades
no sharing
old stuff is broken

reprex
sessioninfo()

site library
frozen repo
renv
internal repo
custom tests



renv (packrat 2.0)

similar to python virtualenv

rstudio.github.io/renv

renv::init()

- 1. Finds existing dependencies (dependencies())
- 2. Creates lock file (snapshot())

renv::snapshot()

```
"renv": {
  "Version": "0.9.2-27"
},
"R": {
  "Version": "3.6.2",
  "Repositories": [
      "Name": "CRAN",
      "URL": "https://cloud.r-project.org"
"Packages": {
  "markdown": {
    "Package": "markdown",
    "Version": "1.0",
    "Source": "Repository",
    "Repository": "CRAN",
    "Hash": "4584a57f565dd7987d59dda3a02cfb41"
  "mime": {
    "Package": "mime",
    "Version": "0.7",
    "Source": "Repository",
    "Repository": "CRAN",
    "Hash": "908d95ccbfd1dd274073ef07a7c93934"
```

renv::restore()

```
> renv::restore()
    The following package(s) will be updated:
63
    # Bioconductor =============
    AnnotationDbi
                             [* -> 1.48.0]

    AnnotationFilter

                             [* -> 1.10.0]
                             [* -> 1.54.0]
    BSgenome
   - Biobase
                             [* -> 2.46.0]
    BiocFileCache
                             [* -> 1.10.2]
                             [* -> 0.32.0]
    BiocGenerics

    BiocParallel

                             [* -> 1.20.0]
    - BiocVersion
                             [* -> 3.10.1]
                             [* -> 2.54.0]
    Biostrings
    DelayedArray
                             [* -> 0.12.0]
                             [* -> 1.22.0]
    GenomeInfoDb
    GenomeInfoDbData
                             [* -> 1.2.2]
                             [* -> 1.22.1]

    GenomicAlignments

    GenomicFeatures
                             [* -> 1.38.0]
    GenomicRanges
                             [* -> 1.38.0]
                             [* -> 2.20.1]
   IRanges
                             [* -> 1.28.0]
    OrganismDbi
   ProtGenerics
                             [* -> 1.18.0]
```


Try renv

install renv and try renv::init() on a real project

Maybe the wtf-packages-report?

restart R

Use devtools::session_info() output

Docker + renv = powerful reproducibility!

