

Creating R Packages

A very short introduction

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Why Create Your Own R Packages?

Share to increase efficiency

- Share commonly used code between projects (of yours)
- Improve reproducibility
- Make sure code keeps working
- Add documentation to help yourself
- Control what gets used when

What is in an R Package?

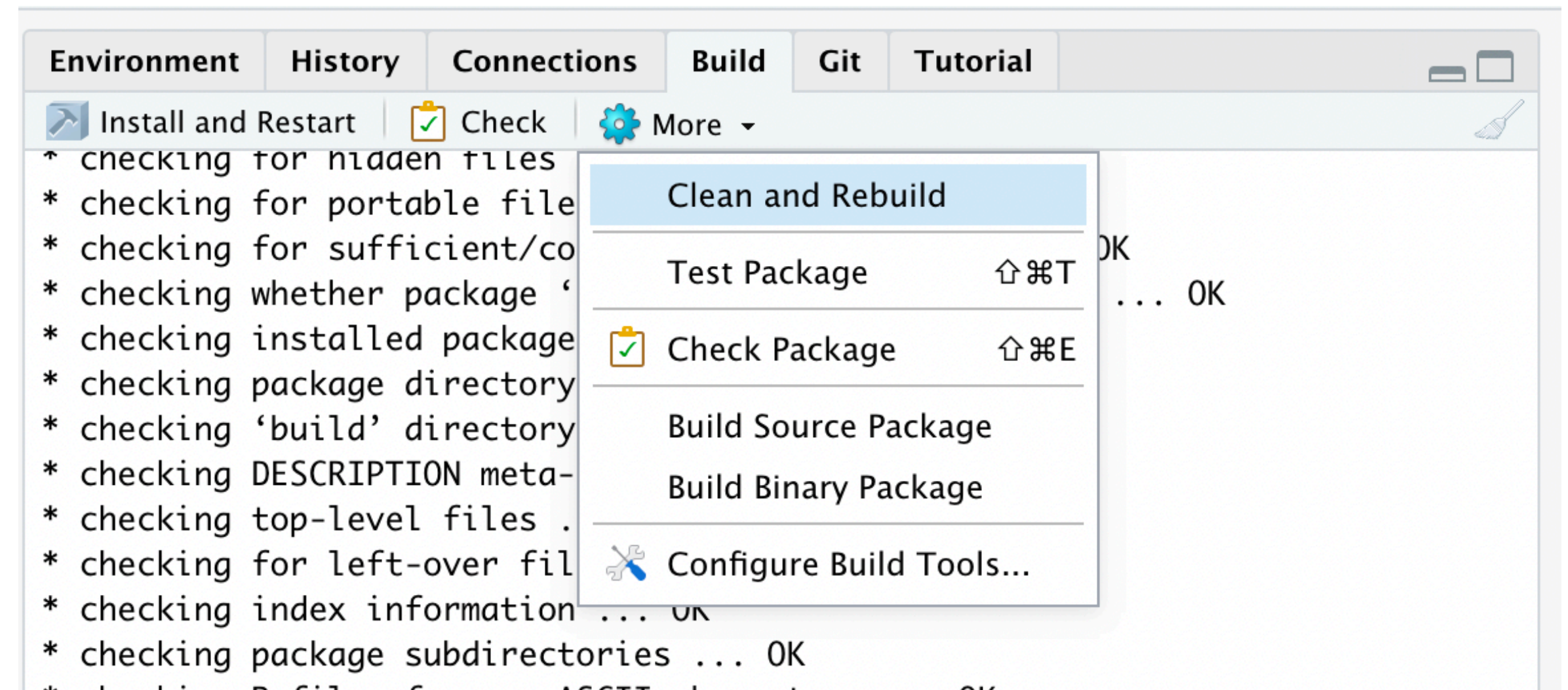
More than just R functions

- R functions
- Datasets
- Documentation: help, examples and vignettes
- Tests
- Namespace information (import & export)
- To be compiled code (C, C++, Fortran)
- Author information

Convert R Scripts to R Package

Let R do it for you

- The old / default way: `package.skeleton` gets you started and automatically converts functions and creates help files.
- Or: Use the `devtools` R package (Cheatsheet: <https://devtools.r-lib.org>)
- R Studio has lots of tools to help



Documentation

Keep code & help close

- Pure R package structures keeps R code in .R files and documentation in .Rd files.
- Keeping things in sync is a hassle to do “by hand”.
- Roxygen2 lets you add documentation in your R files and generates lots automatically.
- Also deals with imports and exports.

```
401 ##' Use \code{compare} to quickly compare the estimated parameters of
402 ##' the fits of multiple lmerMod or rlmerMod objects.
403 ##'
404 ##' @title Create comparison charts for multiple fits
405 ##' @param ... objects to compare, or, for the \code{\link{xtable}}
406 ##' functions: passed to the respective \code{\link{xtable}} function.
407 ##' @param digits number of digits to show in output
408 ##' @param dnames names of objects given as arguments (optional)
409 ##' @param show.rho.functions whether to show rho functions in output.
410 ##' @keywords models utilities
411 ##' @examples
412 ##' \dontrun{
413 ##'   fm1 <- lmer(Yield ~ (1|Batch), Dyestuff)
414 ##'   fm2 <- rlmer(Yield ~ (1|Batch), Dyestuff)
415 ##'   compare(fm1, fm2)
416 ##'   require(xtable)
417 ##'   xtable(compare(fm1, fm2))
418 ##'   str(getInfo(fm1))
419 ##' }
420 ##' @export
421 compare <- function(..., digits = 3, dnames = NULL,
422                      show.rho.functions = TRUE) {
423   linfos <- list(...)
424   if (!missing(dnames) && !is.null(dnames)) names(linfos) <- dnames
425   linfos <- lapply(linfos, getInfo)
426   ...
```

Share with Others

CRAN is not the only way

- R CMD build creates an .tar.gz archive you can install elsewhere
- R CMD INSTALL --build <your-package>.tar.gz if you need to compile code for someone on Windows (produces a .zip archive that everybody can install).
- Put your package on github, etc, and install via `devtools::install_github("kollerma/robustlmm")`
- But CRAN is definitely the easiest way to share your package

Publish your Package on CRAN

The CRAN police will be watching you

- If you follow the rules, it's easy. Make sure you know them: <https://cran.r-project.org/web/packages/policies.html>
- Check your package:
- Run R CMD `check --as-cran <your-package>.tar.gz`
- with several versions of R (previous, current and development)
- Submit via web-form: <https://xmpalantir.wu.ac.at/cransubmit/>

Tests

What isn't tested will break

- Write examples for every function you export / expose to the user.
- All examples are run when running R CMD check
- Add test scripts to tests/ subdirectory.
- Simplest way: accompany .R files in tests/ with .Rout files. (Hint: Copy them over from previous R CMD check runs.) Any changes will be highlighted.
- Write unit tests via testthat R package (<https://testthat.r-lib.org>).

R Documentation

The R Packages “Bible”

- Writing R Extensions
<https://cran.r-project.org/doc/manuals/r-release/R-exts.html>
- Whatever your problem, this usually has the solution.