

# Submitting to CRAN and Continuous Integration

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[http://johnmuschelli.com/smi\\_hackathon\\_2018.html](http://johnmuschelli.com/smi_hackathon_2018.html)

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# Unit Testing

- Runs specific code for test cases.
- `testthat` and `RUnit` packages
- `usethis::use_testthat()`

```
context("All my data tests")
test_that("Data is installed", {
  expect_is(sri24_image_df(), "data.frame")
})
```

- **see** `expect_equal`, `expect_identical`, etc.

Unit Testing: Any issue/error/edge case deserves a unit test

# Code Coverage: `covr` package

- Calculates the code coverage: % of lines tested when package is tested
  - excludes white space
  - `covr::package_coverage()` - defaults to tests unit tests only
  - `covr::package_coverage(type = "all")` - runs tests, vignettes, examples.

# Code Coverage: `covr` package

Online services that allow you to view the coverage online:

- Coveralls: <https://coveralls.io/>
- Codecov: <https://codecov.io/>

# Checking Packages Locally

- Check the Package: Build → Check Package.
  - Make sure Build → Configure Build Tools. Put `--as-cran` in Check Package area.

```
devtools::check(args = c('--as-cran'))
```

- Runs examples, tests, and check imports/suggests/etc.

# Checking Packages Locally

```
devtools::check(args = c('--as-cran'))
```

Before submitting to CRAN:

- Run `available::available("PACKAGE_NAME")`: checks package name
- Shows `notes`: non-global visible things are a by-product of `tidyverse` non-standard evaluation

# Continuous Integration: Travis and Appveyor

- Builds and checks R packages on Windows (Appveyor) and Linux/OS X (Travis CI)
- Sign up with GitHub, works well with GitHub
- Installs R, runs R CMD check --as-cran, and provides result



Travis CI



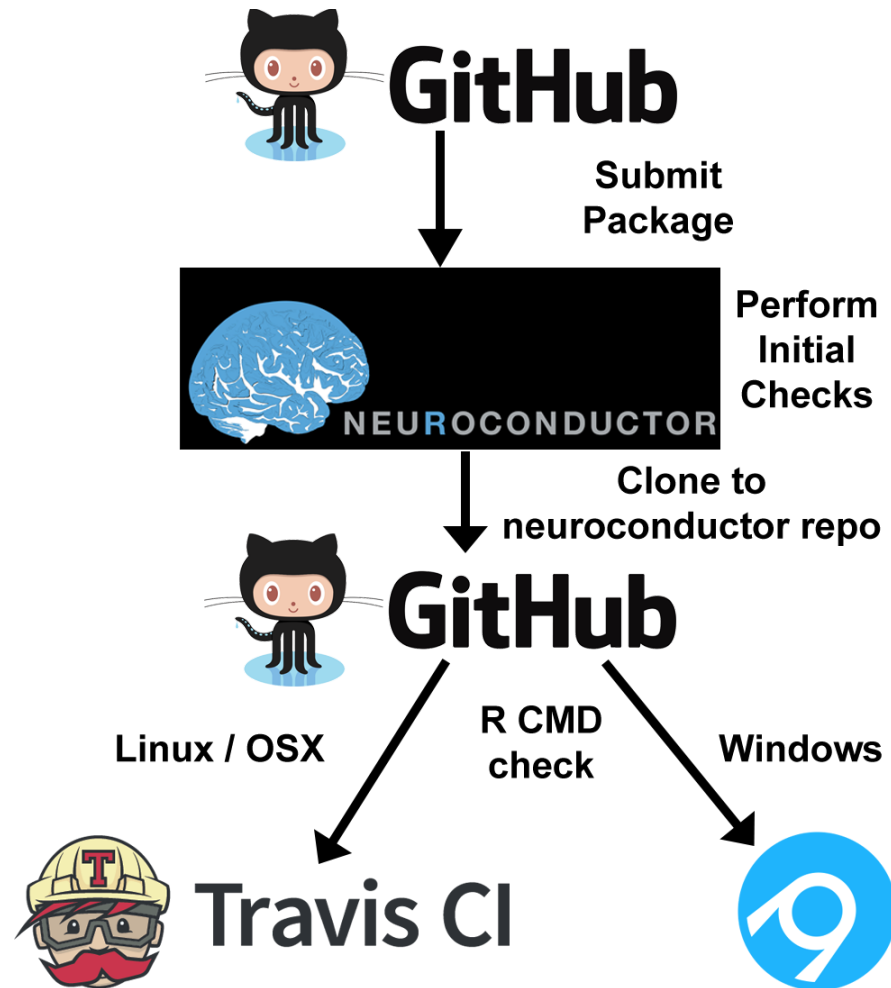


# Development

## Pipeline:

Check the package for stability

- uses current Neuroconductor Packages



# Using CI and Coverage: Badges

- `usethis::use_travis()` - creates `.travis.yml`
  - add `warnings_are_errors: true`

Add coverage: `usethis::use_coverage()` - pick  
`coveralls/codecov`:

```
after_success:
```

- `Rscript -e 'covr::coveralls(type = "all")'`
- `Rscript -e 'covr::codecov(type = "all")'`

- `usethis::use_appveyor()` - creates  
`appveyor.yml`

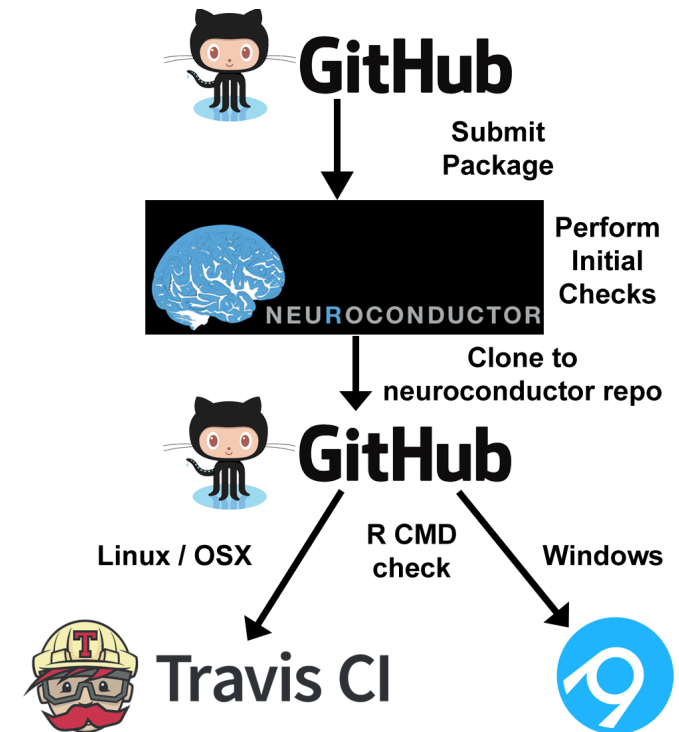
# Submitting to CRAN

- [Writing R Extensions](#) - all things for a package
- CRAN policy (long): <https://cran.r-project.org/web/packages/policies.html>
- `devtools::release()` (needs `NEWS.md` and `cran-comments.md`)
- RStudio → Build → Build Source Package
  - builds vignettes, creates a `.tar.gz`, upload to <https://cran.r-project.org/submit.html>

# Submitting to Neuroconductor

[https://neuroconductor.org/su  
bmit-package](https://neuroconductor.org/su<br/>bmit-package)

- Must be a GitHub repo
- Verify via maintainer email
- See [https://neuroconductor.org/l  
ist-packages/all](https://neuroconductor.org/l<br/>ist-packages/all)



# Solutions to Common Problems

- Clear out `man` folder
- Replace `NAMESPACE` with blank file with `#`  
Generated by `roxygen2`: do not edit by hand at the top
- Check your `Imports/Suggests/Depends`

# Recommendations

- Use `Imports`. Only in very rare cases, use `Depends`, unless depending on R version.
- Use `@importFrom` directives instead of `@import`
  - helps you identify problems (moving functions)
- Use the “Go to file/function” search in RStudio
- Separate files for each function (personal preference)