Construção de Pacotes em R: da Teoria à Prática Utilizando o RStudio

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Construção de Pacote

Por que criar um pacote no R?

- Organização de funções, exemplos e conjuntos de dados associados a um mesmo tema.
- Divulgação:
 - Fácil distribuição;
 - Comunidade familiarizada com a instalação e utilização de pacotes.

Construção de Pacote

Pré-Requisitos

- Programar em R
- R, RStudio e RTools atualizados
- Pacotes:
 - "devtools" atualizar o pacote, simular instalação
 - "roxygen2" criar documentação
 - "testthat" testes de funções, validações
 - "usethis" substituto do "devtools" (mais intuitivo)
 - "knitr" Rmarkdown
- git instalado versionamento
- compilador de LaTeX instalado "vignettes" (manuais)

Construção de Pacote

Estrutura de um pacote no R

- Arquivo "DESCRIPTION" metadados do pacote (nome, autor, versão, licença etc)
- Arquivo "NAMESPACE" lista funções que estarão disponíveis para os usuários
- Diretório "R" funções (arquivos .R)
- Diretório "man" manuais (arquivos .Rd).
- Opcionais:
 - Diretório "data" arquivos de dados para exemplos ou testes
 - Diretório "tests" testes das funções do pacote
 - Diretório "vignettes" explicações mais didáticas sobre as funções e o pacote

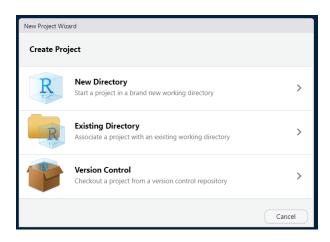


Figura: RStudio: File - New Project - New Directory

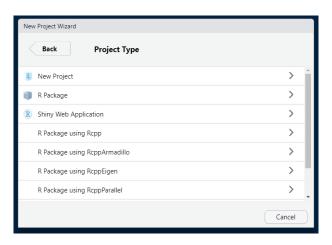


Figura: R package

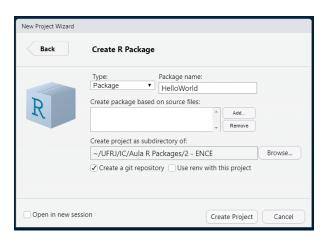


Figura: Pacote "HelloWorld"

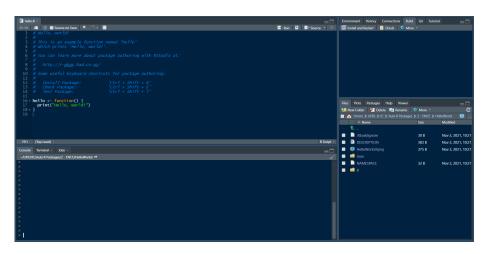


Figura: Tela Inicial

```
hello.R
           Source on Save
 16 • hello <- function() {
      print("Hello, world!")
 18 4 }
```

Figura: Função "hello"

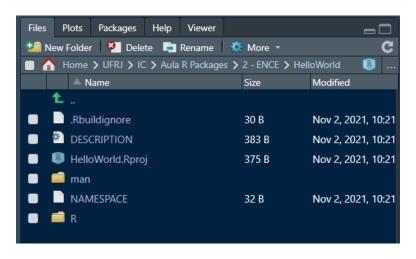


Figura: Arquivos Padrão

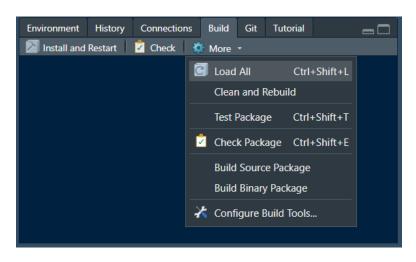


Figura: Aba "Build"



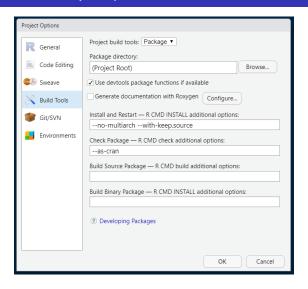


Figura: "Project Options"

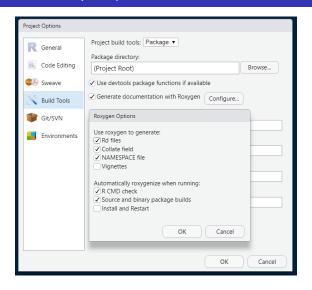


Figura: Geração de arquivos com o "Roxygen"



Figura: Documentação com o "Roxygen" - função ("Insert Roxygen Skeleton")

```
18  #' Title
19  #'
20  #' @return
21  #' @export
22  #'
23  #' @examples
24  hello <- function() {
25  print("Hello, world!")
26  }
27
```

Figura: Documentação com o "Roxygen" - estrutura padrão

```
@param nome nome desejado
23
       @return frase com o nome escolhido
    #'@examples
    #' hello("Bruno")
29
    #' @export
   hello <- function(nome) {
      print(paste("Hello, ",nome, "!", sep=""))
```

Figura: Documentação com o "Roxygen" - estrutura preenchida

```
==> devtools::document(roclets = c('rd', 'collate', 'namespace'))

Updating Helloworld documentation
First time using roxygen2. Upgrading automatically...

Updating roxygen version in C:\Users\pedro\OneDrive\Documentos\UFRJ\I
C\aula R Packages\2 - ENCE\Helloworld/DESCRIPTION

Loading Helloworld
Warning: The existing 'hello.Rd' file was not generated by roxygen2,
and will not be overwritten.

Warning: The existing 'NAMESPACE' file was not generated by roxygen2,
and will not be overwritten.

Documentation completed
```

Figura: Documentação com o "Roxygen" - comando "Document"

```
=>> devtools::document(roclets = c('rd', 'collate', 'namespace'))
Updating Helloworld documentation
First time using roxygen2. Upgrading automatically...
Loading Helloworld
Writing NAMESPACE
Writing hello.Rd
Writing NAMESPACE
Documentation completed
```

Figura: Documentação com o "Roxygen" - comando "Document"

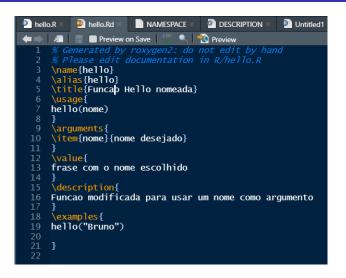


Figura: Documentação com o "Roxygen" - Diretório "man" - hello.Rd

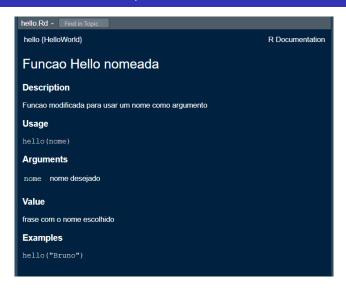


Figura: Documentação com o "Roxygen" - ajuda da função hello()

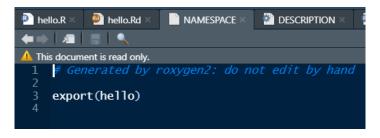


Figura: Documentação com o "Roxygen" - arquivo "Namespace"

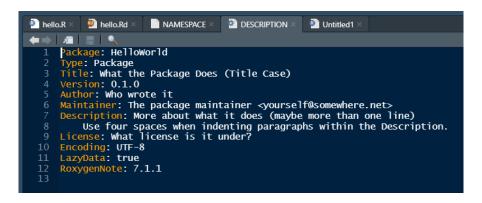


Figura: Arquivo "Description" - padrão

```
nello.R
          DESCRIPTION
                          nello.Rd
                                      NAMESPACE
                                                     DESCRIPTION >
                                                                     Untitled1
     Package: BayesSampling
      Type: Package
     Title: Bayes Linear Estimators for Finite Population
     Version: 1.1.0
     Date: 2021-04-24
     Authors@R: c(person("Pedro", "Soares Figueiredo",
                            role = c("aut", "cre"),
                            email = "pedrosfig@hotmail.com",
comment = c(ORCID = "0000-0003-2279-2881")),
                    person("Kelly C.", "M. Gonçalves",
                            role = c("aut", "ths"),
                            email = "kelly@dme.ufrj.br"
                            comment = c(ORCID = "0000-0002-4524-547X")))
     Maintainer: Pedro Soares Figueiredo <pedrosfig@hotmail.com>
     Description: Allows the user to apply the Bayes Linear approach to finite population with the Simple Random Sampling - BLE_SRS() - and
          the Stratified Simple Random Sampling design - BLE_SSRS() - (both without replacement), to the Ratio estimator (using auxiliary
          information) - BLE_Ratio() - and to categorical data - BLE_Categorical().
          The Bayes linear estimation approach is applied to a general linear regression model for finite population prediction in BLE_Reg()
          and it is also possible to achieve the design based estimators using vague prior distributions.
          Based on Goncalves, K.C.M. Moura, F.A.S and Migon, H.S. (2014) <a href="https://www150.statcan.gc.ca/n1/en/catalogue/12-001-X201400111886">https://www150.statcan.gc.ca/n1/en/catalogue/12-001-X201400111886</a>.
     URL: https://www150.statcan.gc.ca/n1/en/catalogue/12-001-X201400111886, https://github.com/pedrosfig/BayesSampling
     License: GPL-3
     Encoding: UTF-8
    LazyData: true
     RoxygenNote: 7.1.1
     Depends: R (>= 3.5)
     Imports:
          MASS.
          Matrix.
          stats.
          matrixcalc
     Suggests:
          knitr.
          rmarkdown.
          TeachingSampling
     VignetteBuilder: knitr
     Language: en-US
```

Figura: Arquivo "Description" - pacote "BayesSampling"

```
ainv <- MASS::ainv
      bdiag <- Matrix::bdiag
      var <- stats::var
      sample <- base::sample
      colnames <- base : colnames
      is.symmetric.matrix <- matrixcalc::is.symmetric.matrix
      is.positive.definite <- matrixcalc::is.positive.definite
 17:1
        (Top Level)
                                                    use_package(package, type = "Imports", min_version =
                                                       NULL)
Console
         Terminal
                    lohs
                                                    use package() adds a CRAN package dependency to DESCRIPTION
~/UFRJ/IC/Aula R Packages/2 - ENCE/HelloWorld/
                                                    and offers a little advice about how to best use it, use dev package ()
                                                    adds a versioned dependency on an in-development GitHub package.
                                                    adding the repo to Remotes so it will be automatically installed from the
                                                    correct location.

    use package

             use_package_doc
                                                    Press F1 for additional help
  usethis::use_package
```

Figura: Funções importadas de outros pacotes - "usethis"

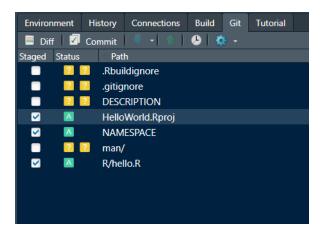


Figura: Git - versionamento

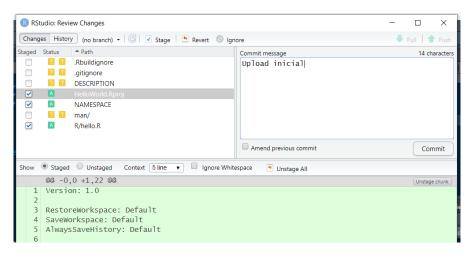


Figura: Git - preparando "Commit"

```
Julo, Neview Changes
  Git Commit
                                                                                  Close
   >>> C:/Program Files/Git/bin/git.exe commit -F C:/Users/pedro/AppData/Local/Temp/
   [master (root-commit) 7388aa7] Upload inicial
    3 files changed, 58 insertions(+)
    create mode 100644 HelloWorld.Rproj
    create mode 100644 NAMESPACE
    create mode 100644 R/hello.R
```

Figura: Git - "Commit" feito

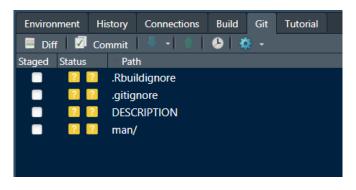


Figura: Git - arquivos atualizados não aparecem mais

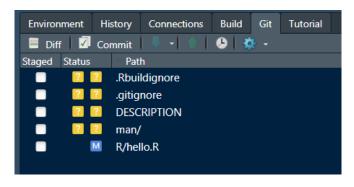


Figura: Git - arquivo hello.R modificado

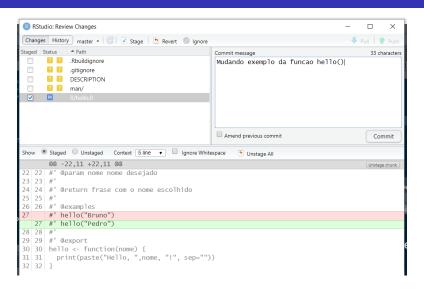


Figura: Git - mudanças evidenciadas

```
Git Commit
                                                                              Close
>>> C:/Program Files/Git/bin/git.exe commit -F C:/Users/pedro/AppData/Local/Temp/
 [master baf88f3] Mudando exemplo da funcao hello()
 1 file changed, 1 insertion(+), 1 deletion(-)
```

Figura: Git - arquivo alterado

```
🔯 Install and Restart 💆 Check 🏻 🔅 More 🔻
==> devtools::check(args = c('--as-cran'))
Updating HelloWorld documentation
Loading HelloWorld
Writing NAMESPACE
Writing NAMESPACE
```

Figura: Check - "devtools"

```
See
     'C:/Users/pedro/OneDrive/Documentos/UFRJ/IC/Aula_R Packages/2 - ENCE/HelloWorl
d.Rcheck/00check.log'
   for details
Duration: 26.2s
  Non-standard license specification:
    What license is it under?
  Standardizable: FALSE
Erro: R CMD check found WARNINGS
Execu@@o interrompida
Exited with status 1.
```

Figura: Check - warning

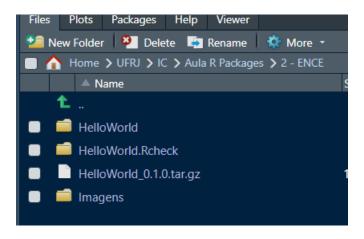


Figura: Check - diretório criado



Figura: Check - pacote "rhub"

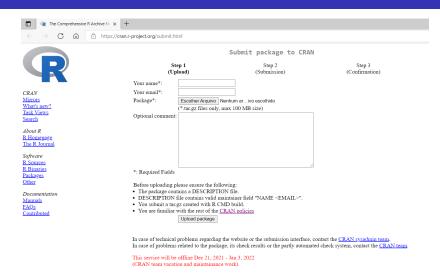


Figura: Submissão ao CRAN

BayesSampling: Bayes Linear Estimators for Finite Population

Allows the user to apply the Bayes Linear approach to finite population with the Simple Random Sampling - BLE_SRS() - and the Stratified Simple Random Sampling design - BLE_SSRS() - (both without replacement), to the Ratio estimator (using auxiliary information) - BLE_Ratio() - and to categorical data - BLE_Categorical(). The Bayes linear estimation approach is applied to a general linear regression model for finite population prediction in BLE_Reg() and it is also possible to achieve the design based estimators using vague prior distributions. Based on Gonçalves, K. CM, Moura, F.A.S and Migon, H.S.(2014) < https://www.150.statean.gc.ca/n/len/catalogue/12-001-X201400111886>.

Version: 1.1.0 Depends: R (> 3.5)

Imports: MASS, Matrix, stats, matrixcale
Suggests: knitr. rmarkdown, TeachingSampling

Published: 2021-05-01

Author: Pedro Soares Figueiredo 🙃 [aut. cre], Kelly C. M. Goncalves 🙃 [aut. ths]

Maintainer: Pedro Soares Figueiredo <pedrosfig at hotmail.com>

License: GPL-3

URL: https://www150.statcan.gc.ca/n1/en/catalogue/12-001-X201400111886,

https://github.com/pedrosfig/BayesSampling

NeedsCompilation: no

Language: en-US
Materials: README

CRAN checks: BayesSampling results

Figura: CRAN - BayesSampling

```
Materials:
                   README
CRAN checks:
                   BayesSampling results
Documentation:
Reference manual: BayesSampling.pdf
Vignettes:
                  BLE Categorical
                  BLE Ratio
                  BLE Reg
                  BLE SRS
                  BLE SSRS
                  BayesSampling
Downloads:
                  BayesSampling 1.1.0.tar.gz
Package source:
Windows binaries: r-devel: BayesSampling 1.1.0.zip, r-release: BayesSampling 1.1.0.zip, r-oldrel:
                  BayesSampling 1.1.0.zip
macOS binaries: r-release (arm64): BayesSampling 1.1.0.tgz, r-release (x86 64): BayesSampling 1.1.0.tgz, r-oldrel:
                  BayesSampling 1.1.0.tgz
Old sources:
                  BayesSampling archive
Linking:
Please use the canonical form https://CRAN.R-project.org/package=BayesSampling to link to this page.
```

Figura: CRAN - BayesSampling



Figura: CRAN - manual padrão: BayesSampling.pdf

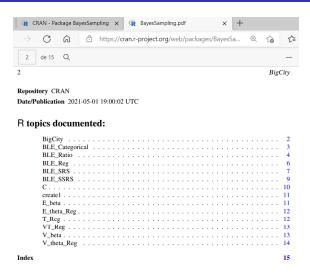
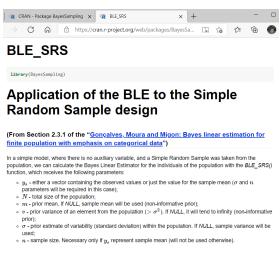


Figura: CRAN - manual padrão: BayesSampling.pdf



Vague Prior Distribution

Figura: "vignette" - função BLE_SRS()

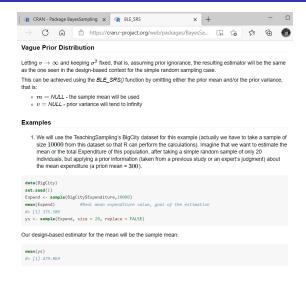
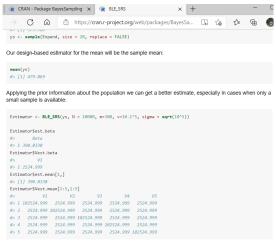


Figura: "vignette" - função BLE_SRS()



2. Example from the help page

Figura: "vignette" - função BLE_SRS()

Referências



Hartigan, J. A. (1969). Linear Bayesian methods. J. Roy. Statist. Soc. Ser. B, 31, 440–454.

Roger D. Peng, Sean Kross, and Brooke Anderson (2017). Mastering Software Development in R: https://bookdown.org/rdpeng/RProgDA/

Zacks, S. (2002). In the footsteps of Basu: The predictive modelling approach to sampling from finite population. Sankhyā: The Indian Journal of Statistics, Series A, 64, 532-544.

Muito Obrigado!

Agradecimentos:



- Contato: pedrosfig@hotmail.com
- Pacote: https://cran.r-project.org/package=BayesSampling
- Links: https://github.com/pedrosfig/Sem_ENCE_Pacotes_R

