



16th Meeting of the Hamburg R-User-Group, 13th Feb 2019

Project "easystats" Making R stats easier!

Daniel Lüdecke d.luedecke@uke.de https://github.com/easystats









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Dominique



Neuropsychologist, psychotherapist, pizza lover 🔊

Postdoc at the Clinical Brain Lab (Singapore) on the neuroscience of deception







https://github.com/easystats

Dominique



Postdoc at the Clinical Brain Lab (Singapore) on the neuroscience of deception





Gerontologist, kind of sociologist, prefers burgers



Postdoc at the Department of Medical Sociology (University Medical Center Hamburg)









- Provide a set of packages that makes it easier to do statistical analysis and reporting with R.
 - □ Low-level (or "core") packages
 - Target group: advanced users and developers
 - Aims (examples): accessor functions to access the internals of models, such as variables, formulas, model frame/data, random effects, their structure and so on...







- Provide a set of packages that makes it easier to do statistical analysis and reporting with R.
 - ☐ Mid-level packages
 - Target group: end-user
 - Aims (examples): computation of model "performance" metrics (R2, ICC, CoD, AIC, BIC and whatnot), model comparison, Bayesian analysis, . . .







- Provide a set of packages that makes it easier to do statistical analysis and reporting with R.
 - ☐ High Level
 - Target groups: non-experts/beginners that want fully-baked solutions to solve their problems and that want to experience the power of R
 - Aims: reporting, plotting







 Provide a set of packages that makes it easier to do statistical analysis and reporting with R.



And most important!

 All packages, especially the low-level packages, should run with minimum dependencies!

```
Package: insight
Type: Package
Title: Easily Access Model Information for Various Model Objects
Description: Although there are generic functions to get information about or
  data from models, many modelling-functions from different packages do not
  provide methods to access these information. 'insight' aims to close this
  gap by providing functions that work for (almost) any model object.
Version: 0.1.0.0001
Date: 2019-01-29
Authors@R: person("Daniel", "Lüdecke", role = c("aut", "cre"), email =
"d.luedecke@uke.de", comment = c(ORCID = "0000-0002-8895-3206"))
Maintainer: Daniel Lüdecke <d.luedecke@uke.de>
License: GPL-3
Depends: R (>= 3.2), stats | and no imports!
Suggests: brms, glmmTMB, lme4, nlme, splines, testthat
Encoding: UTF-8
LazyData: true
```





A first low-level package...

insight





Model objects are terrifying

```
# model frame?
library(nlme)

m <- gls(
  follicles ~ sin(2*pi*Time) +
  cos(2*pi*Time), Ovary,
  correlation = corAR1(form = ~ 1 | Mare)
)

model.frame(m)
#> corStruct parameters:
#> [1] 1.960656
```







Model objects are terrifying

```
# model family?

fm1 <- lme(
   distance ~ age, data = Orthodont
)

family(fm1)
#> Error in UseMethod("family") :
#> no applicable method for 'family'
#> applied to an object of class "lme"
```







Model objects are terrifying

```
# model terms?
library(MCMCglmm)
data(PlodiaP0)
m <- MCMCglmm(
  PO~1, random=~FSfamily, data=PlodiaPO,
  verbose=FALSE, nitt=1300, burnin=300,
  thin=1
all.vars(terms(m))
#> Error in terms.default(m) : no
#> terms component nor attribute
```









Gain insight into your models!





Gain insight into your models!

Thanks to a stunning x-ray-technology, the insight-package allows to easily get insights into your model object!









Gain insight into your models!

- Simple, consistent API:
 - get_*() to retrieve data, find_()* to access model information.









- The goal of this package is to provide tools that make it easy and intuitive to access information contained in various models.
- Although there are generic functions to get information and data from models, many modelling-functions from different packages do not provide methods to access these information.
- insight aims at closing this gap by providing consistent functions that work for (almost) any models.





I'm afraid of no model

```
library(insight)
library(nlme)
m < - gls(
 follicles ~ sin(2*pi*Time) +
 cos(2*pi*Time), Ovary,
 correlation = corAR1(form = ~ 1 | Mare)
get_data(m)
         Time follicles
#> Mare
20
#> 3 1 -0.04545455
                       19
#> ... (truncated)
```







I'm afraid of no model

```
library(MCMCglmm)
data(PlodiaP0)
m <- MCMCglmm(</pre>
  PO~1, random=~FSfamily, data=PlodiaPO,
  verbose=FALSE, nitt=1300, burnin=300,
  thin=1
find_terms(m)
$response
[1] "P0"
$conditional
[1] 1
$random
[1] "FSfamily"
```







I'm afraid of no model

```
library(GLMMadaptive)
m <- mixed_model(</pre>
  count ~ child + camper,
  random = \sim 1 | persons,
  zi_fixed = ~ child + livebait,
  zi_random = ~ 1 | persons,
  data = fish,
  family = zi.poisson()
find_predictors(m, component = "zi")
#> $zero_inflated
#> [1] "child" "livebait"
```





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Dominique & Daniel



