Proper identification of individuals is crucial for acknowledging and  
studying their scientific work, be it journal articles or pieces of  
software. In this tech note, one year after CRAN started supporting  
ORCIDs, we shall explain why and how to use unique author identifiers in  
DESCRIPTION files.

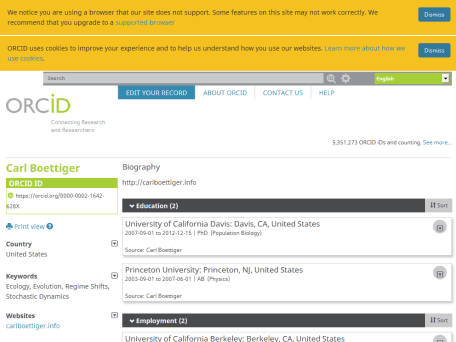
**Why use ORCIDs on CRAN?**

When analyzing the authorship of CRAN packages, one can look at authors’  
names and email addresses. Names can be written with and without quotes,  
email addresses change.

So all in all, using an unique ID per author makes sense. That’s  
something offered for the academic world and beyond by  
[ORCID](https://www.crossref.org/community/orcid/), ORCID meaning “Open  
Researcher and Contributor ID”. Anyone can set up an ORCID account for  
free and link it up to the rest of their online persona, as well as to  
their employment and works. When the authors of a scientific paper give  
their ORCID ID to the publisher, [these can be included in the html and  
PDF versions of a  
paper](https://orcid.org/content/journal-article-display-guidelines).

The advantages of using ORCID IDs in DESCRIPTION are:

* **Unique identifiers for free!** Sure, there’s no verification so  
  someone could use your identity, but at least, it should help  
  assigning your own work to you.
* **Clickable name on CRAN pages, METACRAN pages and pkgdown  
  websites!** E.g. when clicking on Carl’s green bubble on codemetar  
  CRAN page, one gets to [his ORCID  
  profile](https://orcid.org/0000-0002-1642-628X) including a link to  
  his personal website.

" />

Even for non academics, making one’s online persona more accessible from  
a CRAN DESCRIPTION can’t hurt.

**How to manipulate ORCIDs in DESCRIPTION?**

So, how does one add add and interact with ORCID in DESCRIPTION?*!*

Library("desc")

**Search authors by ORCID**

desc offers many functions for manipulating authors: adding or  
deleting them or their roles. Now you can perform these operations using  
ORCID to identify authors to be modified. E.g. if you want to add a role  
to an existing author, say Carl who worked hard enough to earn the “aut”  
role, you’d run

desc::desc\_add\_role(role = "aut",

orcid = "0000-0002-1642-628X")

Obviously this very example above has probably little real-life  
applicability, but you could imagine manipulating DESCRIPTION files *en  
masse* using a table of ORCIDs and new roles.

**Add ORCID to authors of a package**

The new add\_orcid() method and desc\_add\_orcid() functions make it  
possible to add ORCID IDs to authors directly instead of via the  
comment argument. So you could write some script

desc::desc\_add\_orcid("0000-0002-2815-0399",

given = "Maëlle",

family = "Salmon")

and run it inside all your local package folders to add your ORCID. It  
will add this ORCID ID to the author whose given name is “Maëlle” and  
whose family name is “Salmon”. If you only write

desc::desc\_add\_orcid("0000-0002-2815-0399",

given = "Maëlle")

and there are two Maëlle’s, the function will error and you’ll need to  
run it with more specific arguments.

**Add all of *your* identity at once!**

The coolest feature of desc ORCID support might be that  
desc::desc\_add\_me() will now use the ORCID of the environment variable  
ORCID\_ID if you created it. Hence, once per machine it’d now be  
recommended to

* install the whoami package necessary for desc::desc\_add\_me() to  
  work. install.packages("whoami").
* create an environment variable ORCID\_ID, editing .Renviron by  
  calling e.g. usethis::edit\_r\_environ(). Here’s how the  
  corresponding .Renviron line looks for me:

ORCID\_ID="0000-0002-2815-0399"

desc::desc\_add\_me() looks for your name and email addresses in your  
existing configuration .So say you’ve  
just been told to add yourself as contributor of a package you’ve  
improved: inside the package directory, you’d just need to run  
desc::desc\_add\_me() and voilà ! No risk to wrongly copy-paste your  
ORCID.

As a conclusion of this section, not only are ORCIDs useful, but they’re  
getting citizen status in the world of as-automatic-as-possible package  
development. Now, we shall see how one could use ORCIDs inside R to  
gather information about package authors.

**How to study CRAN authors via their ORCID**

The API requires authentication so the first step after installing the  
package is to run:

rorcid::orcid\_auth()

and to save the resulting token as an ORCID\_TOKEN environment variable  
in .Renviron (again, usethis::edit\_r\_environ() can help with that).  
Authentication warrants your having an ORCID account, so if you haven’t  
one yet, [register via this link](https://orcid.org/register).

Then, we’ll query all comment’s of Authors@R of CRAN packages. Some  
CRAN packages use the Author field instead, which is not recommended  
and doesn’t allow using comment.

*Side-note, in general, how would one analyze stuff over the whole CRAN  
collection?*

* *Some information can be found in tools::CRAN\_package\_db() as seen  
  in* [*this recent tech  
  note*](https://ropensci.org/technotes/2018/09/10/github-badges/)*.*
* *There’s* [*an official read-only mirror of each all CRAN packages on  
  GitHub*](https://github.com/cran) *thanks to METACRAN.*
* *METACRAN serves a* [*CRAN package  
  database*](https://r-pkg.org/services#crandb)*, wrapped by an R  
  package of course.*
* *One could try setting up a local CRAN mirror.*

Thanks to Gábor, who maintains METACRAN, for patiently answering my  
questions about meta-CRAN things. If we were after the comments section  
of just a few packages, we could query METACRAN database like so:

url <- "https://crandb.r-pkg.org/-/latest?keys=[%22desc%22,%22codemetar%22]"

obj <- jsonlite::fromJSON(url, simplifyVector = FALSE)

lapply(obj, function(x) eval(parse(text = x$`Authors@R`))$comment)

but in our case, when wanting to get all the current CRAN authors, it’s  
easier to use tools::CRAN\_package\_db(). As a side-note, if you find a  
better solution than the functions defined below parse\_authors() and  
parse\_author() (some purrr magic?) to coerce all authors information  
to a data.frame, feel free to comment under the post!

library("magrittr")

all\_info <- tools::CRAN\_package\_db()

all\_info <- all\_info[!is.na(all\_info$`Authors@R`),]

all\_info <- all\_info[, c("Package", "Authors@R")]

names(all\_info)[2] <- "authors\_at\_r"

null\_to\_na <- function(x){

if(is.null(x)){

NA

}else{

x

}

}

parse\_author <- function(author){

author <- as.person(author)

tibble::tibble(given = null\_to\_na(author$given),

family = null\_to\_na(author$family),

email = null\_to\_na(author$email),

comment = as.character(null\_to\_na(author$comment)),

orcid = as.character(null\_to\_na(author$comment["ORCID"])))

}

parse\_authors <- function(authors\_at\_r){

authors <- try(eval(parse(text = authors\_at\_r)),

silent = TRUE)

if(inherits(authors, "try-error")){

return(list(NA))

}

else{

list(purrr::map\_dfr(authors, parse\_author))

}

}

cran\_authors <- all\_info %>%

as.data.frame() %>%

dplyr::select(Package, authors\_at\_r) %>%

dplyr::rowwise() %>%

dplyr::mutate(authors = parse\_authors(authors\_at\_r)) %>%

tidyr::unnest(authors)

This is how the resulting table looks like. Out of 13147 packages on  
CRAN we are looking at 5421, i.e. packages using the Authors@R field.  
There is one line per author in a package.

str(cran\_authors)

## Classes 'tbl\_df', 'tbl' and 'data.frame': 16272 obs. of 7 variables:

## $ Package : chr "abbyyR" "abc" "abc" "abc" ...

## $ authors\_at\_r: chr "person(\"Gaurav\", \"Sood\", email = \"gsood07@gmail.com\", role = c(\"aut\", \"cre\"))" "c( \n person(\"Csillery\", \"Katalin\", role = \"aut\", email=\"kati.csillery@gmail.com\"),\n person(\"Le"| \_\_truncated\_\_ "c( \n person(\"Csillery\", \"Katalin\", role = \"aut\", email=\"kati.csillery@gmail.com\"),\n person(\"Le"| \_\_truncated\_\_ "c( \n person(\"Csillery\", \"Katalin\", role = \"aut\", email=\"kati.csillery@gmail.com\"),\n person(\"Le"| \_\_truncated\_\_ ...

## $ given : chr "Gaurav" "Csillery" "Lemaire" "Francois" ...

## $ family : chr "Sood" "Katalin" "Louisiane" "Olivier" ...

## $ email : chr "gsood07@gmail.com" "kati.csillery@gmail.com" NA NA ...

## $ comment : chr NA NA NA NA ...

## $ orcid : chr NA NA NA NA ...

We can count the number of occurrences of ORCIDs.

(cran\_authors %>%

dplyr::filter(!is.na(orcid)) %>%

dplyr::pull(Package) %>%

unique() %>%

length() -> orcid\_pkg\_no)

## [1] 613

Only 613 packages have at least an ORCID ID out of 13147 CRAN packages  
(you can’t have an ORCID ID without using the Authors@R field).

There are 512 unique ORCID IDs. Let’s look at the most prolific authors  
with an ORCID ID.

cran\_authors %>%

dplyr::filter(!is.na(orcid)) %>%

dplyr::count(orcid, sort = TRUE) %>%

head(n = 10) -> most\_prolific

knitr::kable(most\_prolific)

| **orcid** | **n** |
| --- | --- |
| 0000-0003-1444-9135 | 29 |
| 0000-0002-4035-0289 | 21 |
| 0000-0003-0918-3766 | 21 |
| 0000-0003-4198-9911 | 19 |
| 0000-0003-4097-6326 | 16 |
| 0000-0001-8301-0471 | 13 |
| 0000-0003-0645-5666 | 12 |
| 0000-0001-5243-233X | 11 |
| 0000-0001-5670-2640 | 11 |
| 0000-0002-8584-459X | 10 |

Then we can use rorcid to get a glimpse of their profile.

rorcid::as.orcid(most\_prolific$orcid)

## [[1]]

## 0000-0003-1444-9135

## Name: Chamberlain, Scott

## URL (first): http://ropensci.org/

## Country: US

## Keywords: ecology, open access, bioinformatics, evolution, R

##

## [[2]]

## 0000-0002-4035-0289

## Name: Ooms, Jeroen

## URL (first): https://github.com/jeroen

## Country: NL

## Country: US

## Keywords:

##

## [[3]]

## 0000-0003-0918-3766

## Name: Zeileis, Achim

## URL (first): https://eeecon.uibk.ac.at/~zeileis/

## Country: AT

## Keywords:

##

## [[4]]

## 0000-0003-4198-9911

## Name: Hornik, Kurt

## URL (first):

## Country:

## Keywords:

##

## [[5]]

## 0000-0003-4097-6326

## Name: Leeper, Thomas

## URL (first): http://www.lse.ac.uk/government/whosWho/Academic%20profiles/ThomasLeeper.aspx

## Country: GB

## Keywords: Public Opinion, Survey Experiments, Political Behaviour, R, Research Design

##

## [[6]]

## 0000-0001-8301-0471

## Name: Hothorn, Torsten

## URL (first):

## Country:

## Keywords:

##

## [[7]]

## 0000-0003-0645-5666

## Name: Xie, Yihui

## URL (first): https://yihui.name

## Country:

## Keywords:

##

## [[8]]

## 0000-0001-5243-233X

## Name: Tang, Yuan

## URL (first): https://terrytangyuan.github.io/about/

## Country: US

## Keywords: Machine Learning, Data Visualization, Open Source, Software Engineering

##

## [[9]]

## 0000-0001-5670-2640

## Name: Rudis, Bob

## URL (first): http://rud.is/b

## Country: US

## Keywords: cybersecurity, r, data visualization, web crawling/scraping

##

## [[10]]

## 0000-0002-8584-459X

## Name: You, Kisung

## URL (first): https://kisungyou.github.io/

## Country: KR

## Keywords:

Over all authors with an ORCID, using rorcid one could extract their  
location and much more, but this is beyond the scope of this tech note.

Inversely, if one wanted to find CRAN packages by ORCID ID, instead of  
ORCID IDs by CRAN package, one could make the most of METACRAN. There is  
no search by ORCID, but since everything in DESCRIPTION is indexed by  
METACRAN, one can use the ORCID ID as search term.

* .

**Conclusion**

In this tech note we made the case for using ORCID IDs as identifiers  
for the authors of CRAN packages, for academics and non-academics as  
well.

For academics, another aspect of making their software contributions  
valuable is getting the software they write as work inside their ORCID  
profile.

But this is probably a topic for another time, such as would be the  
topic of CRAN potentially adopting other IDs like GitHub username,  
ideally via Keybase.

let’s hope ORCID IDs will be more adopted by academics and non-academics  
alike for:

* Making CRAN package authors easier to identify and their online  
  persona easier to find.
* Making academic CRAN package authors’ scientific contributions  
  easier to access when studying packages.

Have *you* started desc::add\_orcid()-ing? Don’t hesitate to comment  
below!