# From Momocs to MomX

Design principles

Vincent Bonhomme

SMEF 2021 - online ed

Julien Claude

Morphometrics
with R





2014

mostly

wrapping

mostly

wrapping



Journal of Statistical Software

February 2014, Volume 56, Issue 13. http://www.jstatsoft.org/

Momocs: Outline Analysis Using R

Vincent Bonhomme French Institute of Pondicherry

Cédric Gaucherel
French Institute of Pondicherry

Sandrine Picq UMR CBAE, Montpellier

Julien Claude University of Montpellier II



complete refactoring



2021 (?)



+ A Evin

+ V Bonhomme

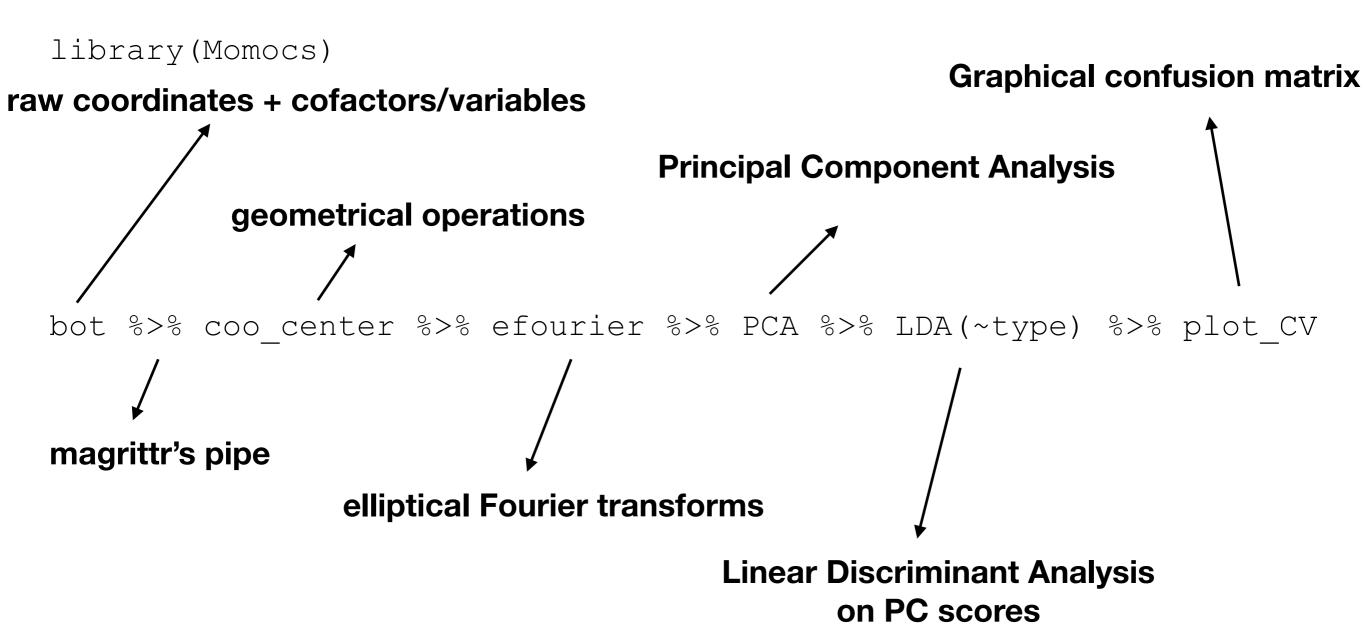
From the very beginning, Momocs aimed to be an unified and complete environment for morphometrics in R

# What's fine with Momocs?

library(Momocs)

bot %>% coo\_center %>% efourier %>% PCA %>% LDA(~type) %>% plot\_CV

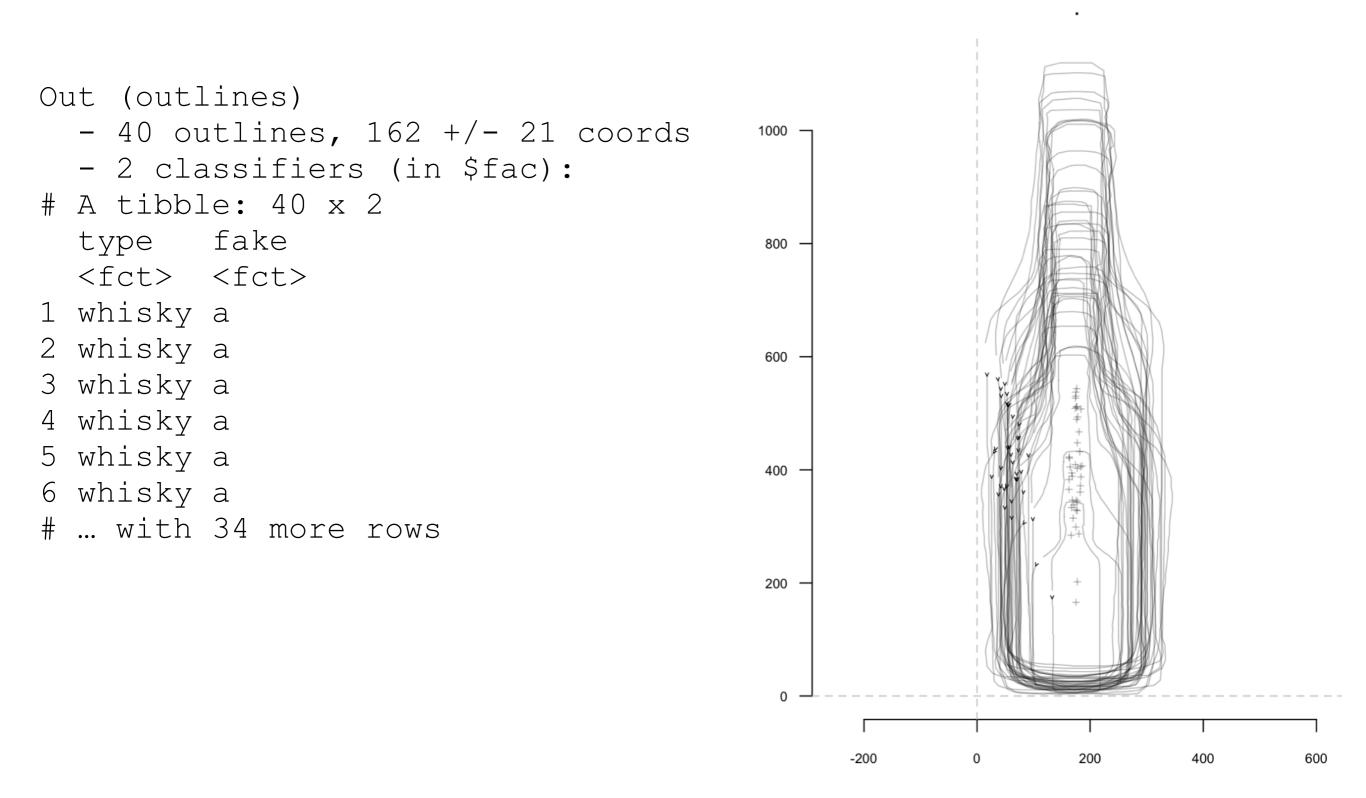
Consistant, compact, pipe-friendly grammar

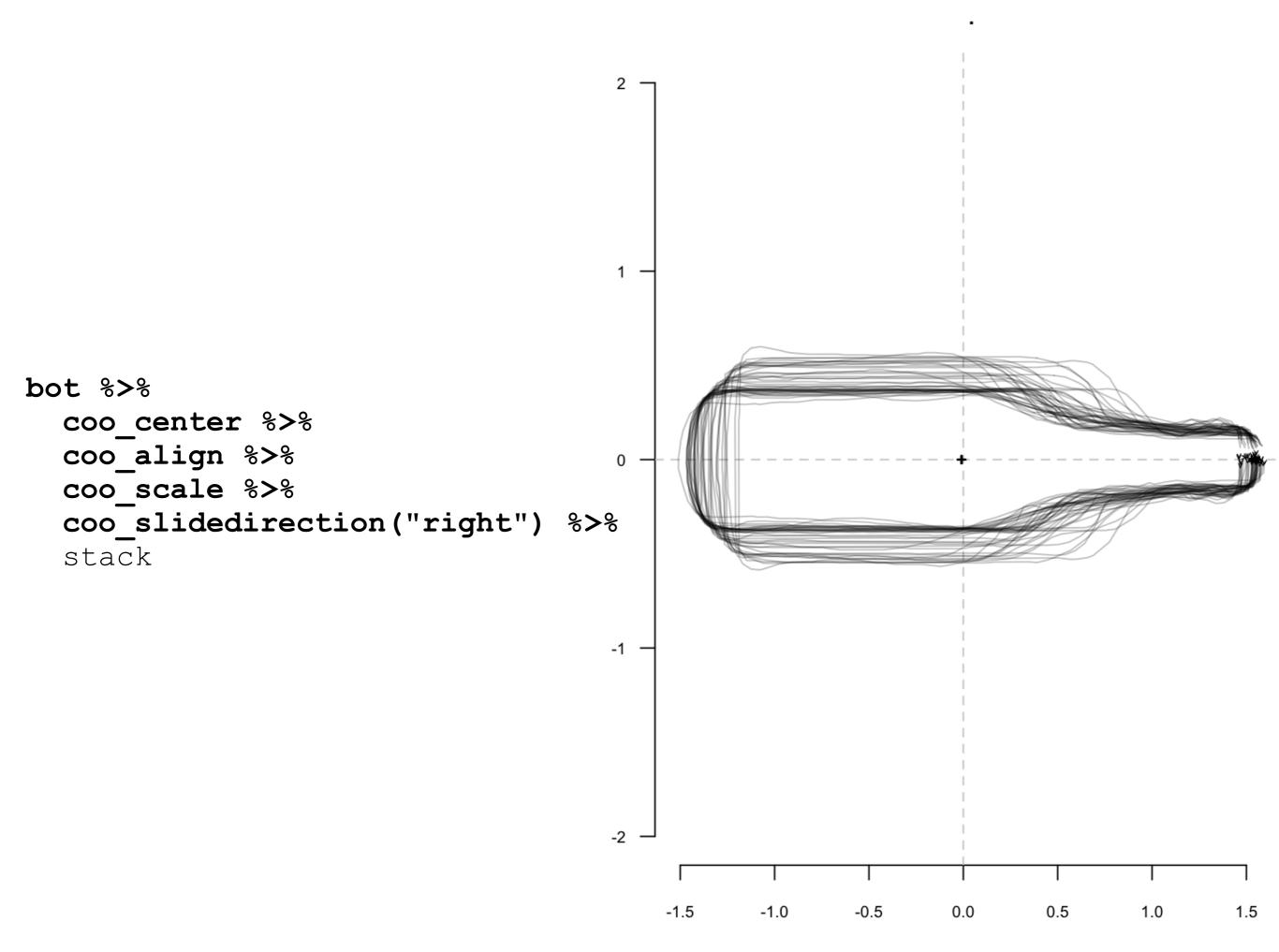


Consistant, compact, pipe-friendly grammar

#### > bot

#### bot %>% stack



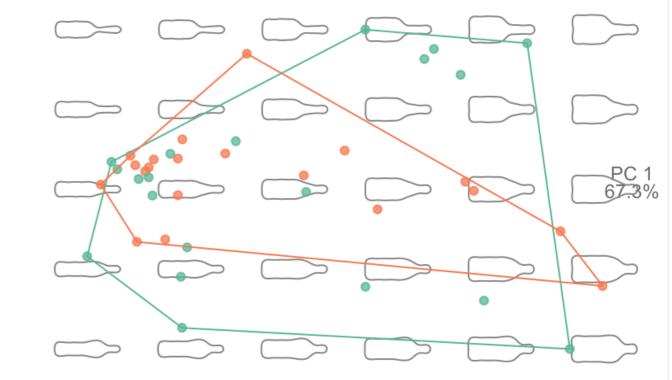


```
bot %>%
  coo center %>%
  coo align %>%
  coo scale %>%
  coo slidedirection("right") %>%
  efourier(norm=FALSE)
'nb.h' set to 10 (99% harmonic power)
An OutCoe object [ elliptical Fourier analysis ]
 - $coe: 40 outlines described, 10 harmonics
# A tibble: 40 x 2
 type fake
 <fct> <fct>
1 whisky a
2 whisky a
3 whisky a
4 whisky a
5 whisky a
6 whisky a
# ... with 34 more rows
```

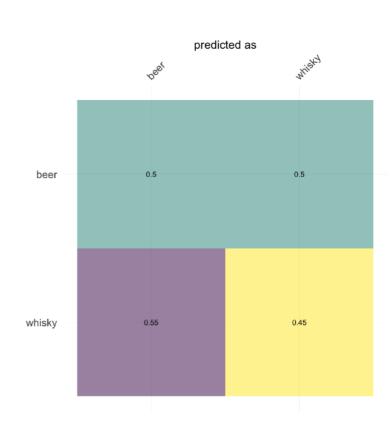
```
PC 2
16.5%
```

```
beerwhisky
```

```
bot %>%
  coo_center %>%
  coo_align %>%
  coo_scale %>%
  coo_slidedirection("right") %>%
  efourier(norm=FALSE)%>%
  PCA %>%
  plot_PCA(~type)
```



```
bot %>%
  coo center %>%
  coo align %>%
  coo scale %>%
  coo slidedirection("right") %>%
  efourier (norm=FALSE) %>%
  PCA %>%
  LDA (~type) %>%
 plot CV()
 * Cross-validation table ($CV.tab):
        classified
actual beer whisky
           10
 beer
                  10
  whisky 11
 * Class accuracy ($CV.ce):
 beer whisky
  0.50 0.45
```



<sup>\*</sup> Leave-one-out cross-validation (\$CV.correct): (47.5% - 19/40):

#### in brief

Not that bad for the regular front-user, Quite boring for serious coders, A nightmare for developers (ie myself)

## Continuously written for 10 years

- > R (and my R) have improved a LOT in the meantime
- > poorly integrate with the tidyverse so far because objects are not native data.frames
- > For complex projects (ie my post-docs) the time patching >>> time exploring :(
- > I spend more time handling the email hotline than improving the doc or the code :(

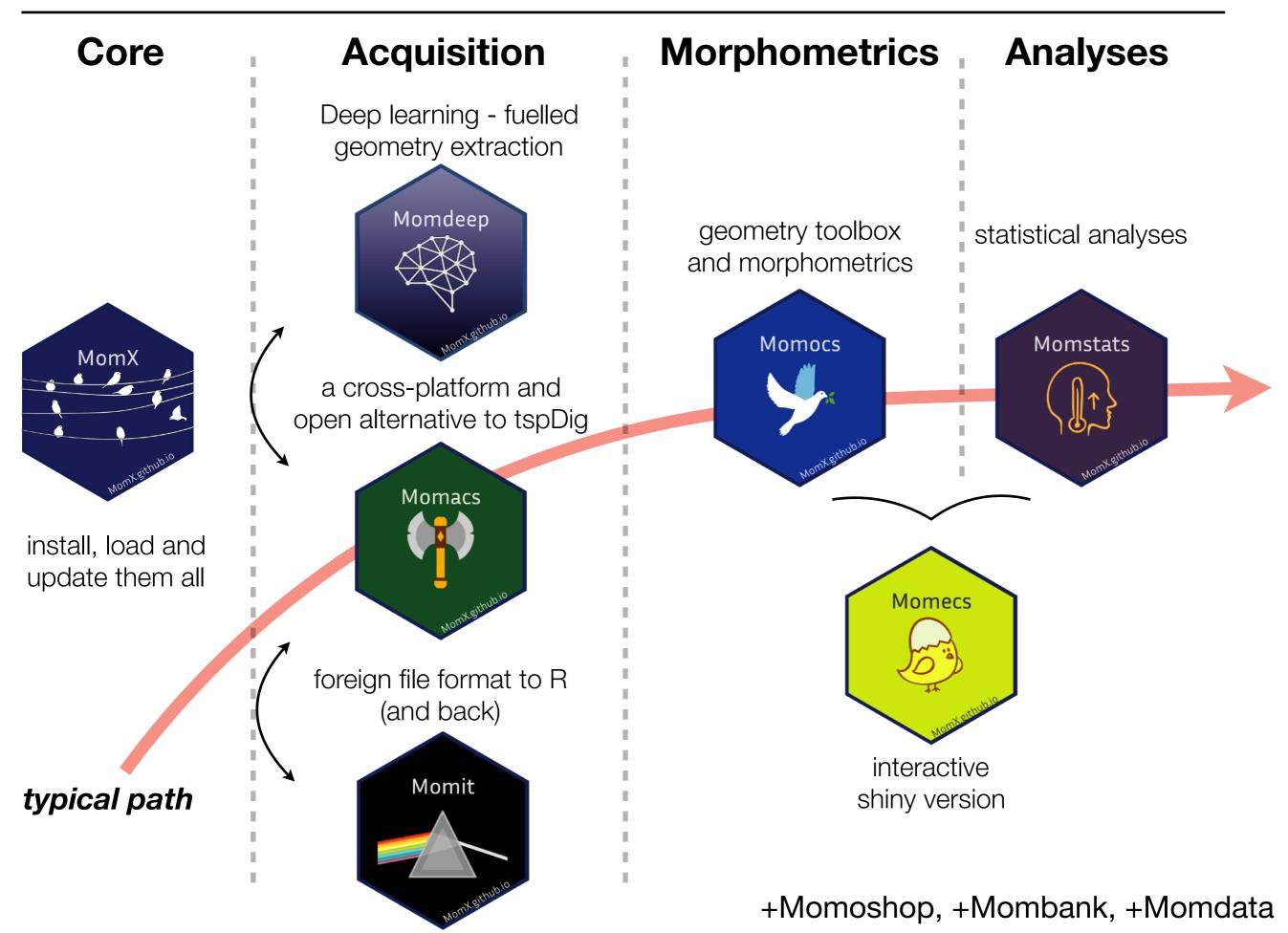
## Too big (~20k lines of code + doc)

- > Hard to extend, to test, to improve
- > API, syntax, ..., became quite inconsistent

### Inherent philosophy was outdated

- > data.frame is the natural object for data science (incl. morphometrics)
- > tidyverse is the natural ecosystem for data science (incl. morphometrics)
- > small is beautiful

Hence the motivation for a complete refactoring of Momocs



1. Everything is a tibble (which is a data.frame, and also a list)
magrittr, ggplot2, dplyr, tidyr, stringr, purrr, tidyeval

2. Every package (and function) does a single task and does it well small is beautiful a single bug does not turn down everything

3. Everything is archived, continuously integrated, tested and documented coding locally -> github -> travis-ci -> code coverage -> pkgdown

fully automated

momx.github.io

#### Complete

- > data acquisition and post-processing (**Momacs**)
- > data sharing/import/export, checking, handling (**Momit**)
- > geometric operations on shapes and morphometrics (Momocs2)
- > statistics (**Momstats**)
- > interactive version of MomX (**Momecs**)
- > automated reporting (**Momscribe**)
- > share datasets and manage collections (Mombank)

#### Convenient and scalable

- > Arrive and leave whenever you want
- > Interoperable packages (tidyverse ready)
- > Easier to optimise (C++)
- > Packages can be dockerised, deployed
- > Foundations to handle millions of shapes

### Open-source and gratis

- > Free plan: written in pure R + GPL3
- > Academic plan: citations + collaborations
- > Training plan: to R and MomX

to fund myself and MomX development

## Towards release

## 2021

v 0.1 (Momit, Momocs2, Momstats) release on CRAN companion paper

#### State of the union

pkg	website	github	travis	last commit	code coverage
MomX	go	go	build error	last commit april	
Momit	go	go	build passing	last commit may	
Momacs	go	go	build passing	last commit march	-
Momocs2	go	go	build passing	last commit may	
Momstats	go	go	build passing	last commit april	
Momdata	go	go	build passing	last commit april	

## 2022

fund myself and MomX Momacs, Momecs, Momdeep more tutorials fill the gaps

many details and working things there:

momx.github.io

feel free to join whatever your skills are!
bonhomme.vincent@gmail.com