## Building Interoperability in Existing Software Ecosystems with S3 Classes

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#### Lots of data wrangling is required to:

- Make successive pieces fit well with one another
- Swap one piece for another equivalent one





Precious time wasted reformatting inputs and outputs.

Very costly in times of emergency.





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# Efforts invested now in interoperability will pay important dividends later down the line.



An international multi-stakeholder project to

harmonise the ecosystem of epidemiology tools in R



Make existing tools interoperable

- V
- Support existing tools to adopt global standards



Develop a sustainable community



Epiverse

TRACE

powered by data.org









#### But we also care about preserving the ecosystem!



More about this in my "CRAN Task View Analysis" poster tomorrow



#### **Conversion functions do not scale**

XKCD 1406: Universal Converter Box, by Randall Munroe, CC BY-NC



Related article: https://voltrondata.com/codex/open-standards



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#### **Conversion functions do not scale**



## Standards are the only viable solution

Related article: <u>https://voltrondata.com/codex/open-standards</u>



#### Standards are hard...





#### Standards are hard... but not impossible!





#### How to fix both ends of the pipe?









1. Community engagement





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2. Technical strategy







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2. Technical strategy

Keep an eye out for links to blog posts with more details!





What is S3?

"S3 is informal and ad hoc, but there is a certain elegance in its minimalism: you can't take away any part of it and still have a useful OO system." Hadley Wickham



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- Mostly an advanced dispatch system based on the presence of the specific attribute
- Used by all R users, even if they don't realize it!



#### •••

fn <- ecdf(rnorm(12))
plot(fn)</pre>

#### •••

m <- lm(bill\_length\_mm ~ bill\_depth\_mm, data = penguins)
plot(m)</pre>

Im(bill\_length\_mm ~ bill\_depth\_mm)





ecdf(x)









#### How to fix both ends of the pipe the R way?





More details at <a href="https://epiverse-trace.github.io/posts/parent-class">https://epiverse-trace.github.io/posts/parent-class</a>



What makes a good S3 class for interoperability?

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Inherit from data.frame where possible

More details at https://epiverse-trace.github.io/posts/parent-class



Go the extra mile to provide support for the tidyverse:

- Support for tibbles as well as data.frames
- Support for dplyr verbs

More details at https://hugogruson.fr/posts/compa-tibble/ & https://epiverse-trace.github.io/posts/extend-dataframes/



Support for tibbles as well as data.frames:

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Support for tibbles as well as data.frames:

- Do not rely on implicit drop value



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#### Support for tibbles as well as data.frames:

- Do not rely on implicit drop value



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Support for tibbles as well as data.frames:

- Do not rely on implicit drop value
- Do not rely on partial matching (df\$c instead of df\$col)



Support for dplyr verbs:

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 Methods for names()<-, [<-, will provide automatic support for most dplyr verbs

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Support for dplyr verbs:

- Methods for names()<-, [<-, will provide automatic support for most dplyr verbs
- If full compatibility is required, you need extra methods for dplyr\_row\_slice(), dplyr\_col\_modify(), dplyr\_reconstruct()

More details at https://epiverse-trace.github.io/posts/extend-dataframes/



How to fix both ends of the pipe the R way?

S3 class







Transposition of "progressive enhancement" web dev concept.



Transposition of "progressive enhancement" web dev concept.

Ensure good experience for all users/packages, even those who don't support the standards, while allowing enhanced experience and extra features when using the standards.



Transposition of "progressive enhancement" web dev concept.

Two key ideas:

- Adding a new method is invisible
- Adding a new attribute is invisible

More details at <a href="https://epiverse-trace.github.io/posts/progressive-enhancement/">https://epiverse-trace.github.io/posts/progressive-enhancement/</a>



How to add S3 support "invisibly", without breaking changes?

More details & caveats at <u>https://epiverse-trace.github.io/posts/s3-generic/</u>



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How to add S3 support "invisibly", without breaking changes?



	#' @export centroid <- function(coords, weights) {
	UseMethod("centroid")
	}
>	<pre>#' @rdname centroid #' #' @export centroid.default &lt;- function(coords, weights) {</pre>
	}

More details & caveats at <a href="https://epiverse-trace.github.io/posts/s3-generic/">https://epiverse-trace.github.io/posts/s3-generic/</a>



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```
• • •
centroid <- function(coords, weights) {</pre>
  UseMethod("centroid")
centroid.default <- function(coords, weights) {</pre>
centroid.pointset <- function(coords, weights = NULL) {</pre>
  centroid(coords$coords, coords$weights)
```



How to fix both ends of the pipe the R way?

S3 class

R package



More details at <a href="https://epiverse-trace.github.io/posts/progressive-enhancement/">https://epiverse-trace.github.io/posts/progressive-enhancement/</a>



- If already returning parent from standard, update to return standard





- If already returning parent from standard, update to return standard
- If not possible to update to standard, return a classed output to allow custom dispatch or conversion functions







### For forward-compatibility, class inheritance should never be tested with ==!

More details at <a href="https://developer.r-project.org/Blog/public/2019/11/09/when-you-think-class.-think-again/index.html">https://developer.r-project.org/Blog/public/2019/11/09/when-you-think-class.-think-again/index.html</a>



#### How to fix both ends of the pipe the R way?







#### Conclusion

Three steps to add interoperability in an existing ecosystem:

- 1.Develop standards inheriting from well-established classes (e.g., data.frame)
- 2.Add support for these standards in function inputs by adding new methods
- 3.Add support for these standards in function outputs





This is not necessarily the ideal way to design and implement S3 support in general.

This approach is specifically thought to add S3 support in an existing ecosystem with minimal disruption.



#### Thanks to collaborators and for your attention!

