Rserve, renv, flask, Vue.js dans un docker pour intégrer des données omiques avec ASTERICS Élise Maigné, Nathalie Vialaneix

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- project (2020/2023) funded by
- with a case study contribution from



Tool designed for interactive and integrative data analysis, oriented toward omics data.







The most dangerous part of the talk: the live demo...



https://asterics.miat.inrae.fr







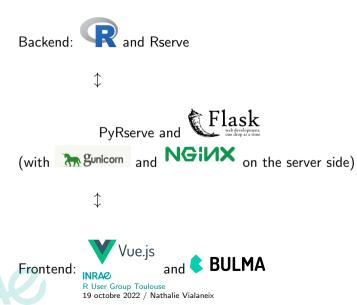




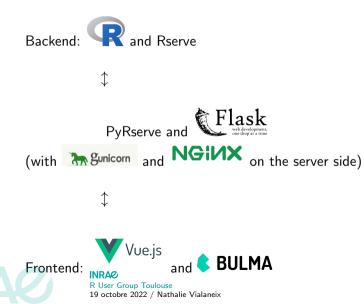








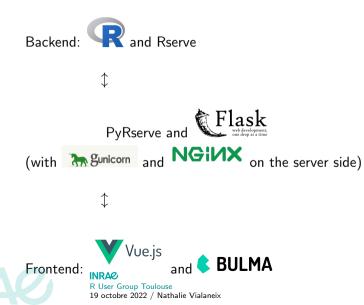




Why this choice?

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- Vuejs: flexible and modular programming





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Compared to shiny?

- could have been a bit more flexible
- shiny deployment is hard (free shiny server / ShinyProxy)
- But: slower, more complicated to program, Rsessions are frequently hijacking the application. 4



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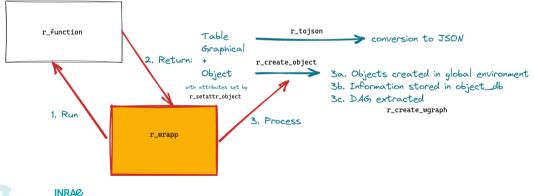
Input arguments: dataset names (and not the actual object!) \Rightarrow need for an object database + use of get and assign.



lacksim Main technical bottleneck from the old R side

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How does it work?



(stabilized with **renv**), Python, and nginx

docker-compose used to combine them

From a user point of view:

- Just pull the images from our registry
- ▶ ... and use our docker-compose file (just one command line)!

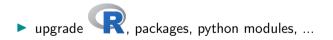




a few bugs to fix

a few other features to implement (metagenomics analyses are among them)

test the deployment on other servers (with docker)









Online application: https://asterics.miat.inrae.fr

User documentation:

https://asterics.pages.mia.inra.fr/user_documentation/ Code repository (with installation instructions): https://forgemia.inra.fr/asterics/asterics Suggest an issue: https://forgemia.inra.fr/asterics/asterics-issues/-/issues or asterics-tlse@inrae.fr Want to contribute? Technical documentation:

https://asterics.pages.mia.inra.fr/doc_asterics/

