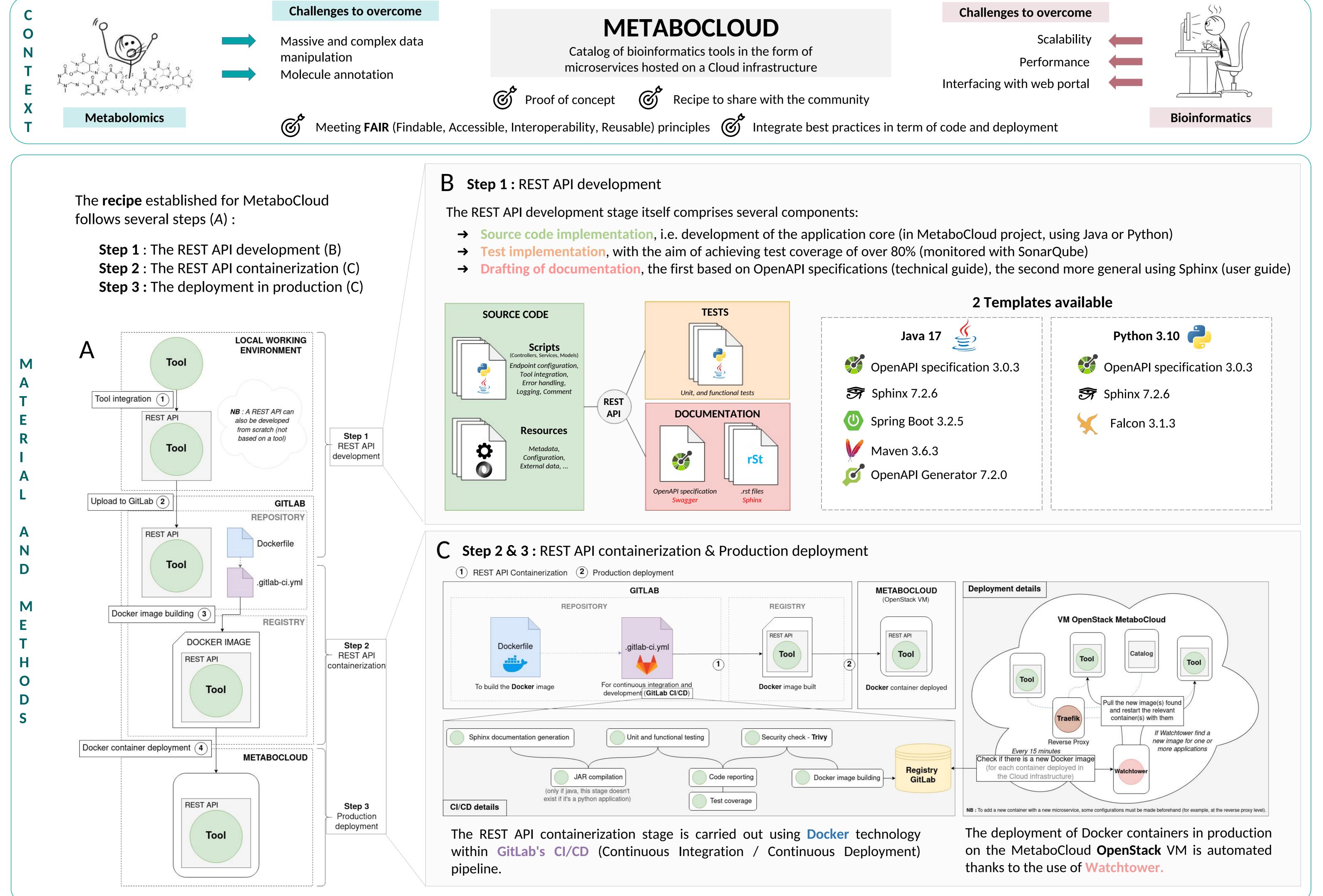
MetaboCloud: A catalog of microservices hosted on a Cloud infrastructure and addressing issues linked to FAIR principles and open science.

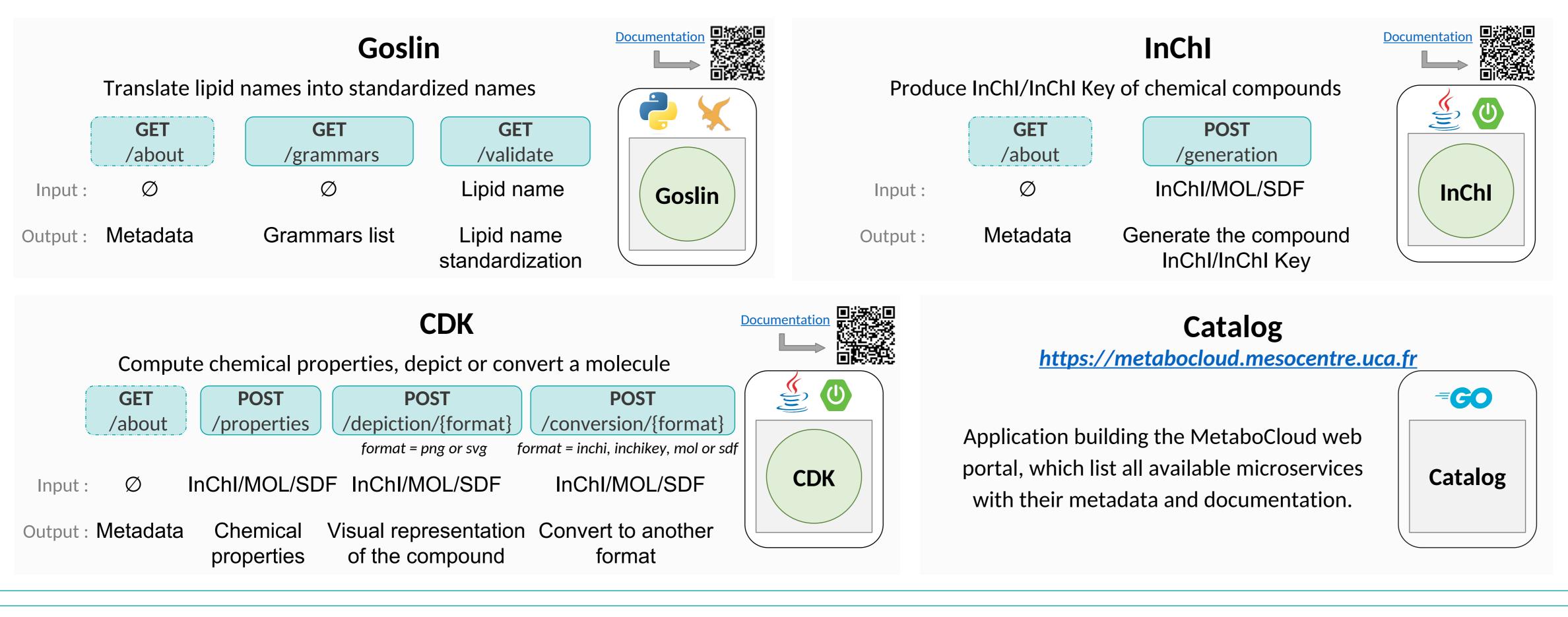
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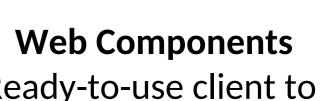
- → Goslin^[1] ("Grammar On Succinct Lipid Nomenclature") microservice developed in python 3.10 from pygoslin 2.1.0
- → CDK^[2] ("Chemistry Development Kit") microservice developed in java 17 from CDK 2.9
- → InChl^[3] microservice developed in java 17 from the binary program InChl 1.06

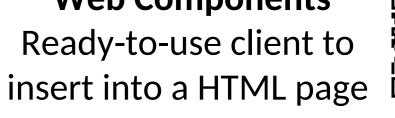


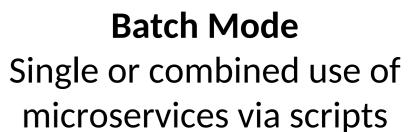
F indable

A ccesible

They can be used in a variety of ways. Either integrated into an application, used in batch mode with scripts, or via their associated web component.







Application

Construction of an application based on microservices architecture

The MetaboCloud microservices infrastructure is based on :

- → Bioinformatic tools
- → From scratch API development if tools are not available
- → Quality testing and documentation
- → An advanced CI/CD work environment managing the construction of a docker image
- → Taken all together in an OpenStack cloud technology environment

Reusable



Web portal associated with metadata (FAIRChecker evaluation)



Can be queried from any programming language, work environment, ...

Version control through containerization

WHAT'S NEXT?

- Development of new microservices
- Maintaining current microservices
- Hackathons directed to developpers/users Current development : a microservice to
- retrieve PNG and MOL stored in a S3 storage

References: [1] Kopczynski D, et al. 2022 Anal Chem. Apr 26;94(16):6097-101. [2] Willighagen EL, et al. 2015 J Cheminformatics. May 30;7:23. [4] Emami Khoonsari P, et al. 2019 Bioinformatics. Oct 1;35(19):3752-3760. **Acknowledgments:** The master internship was supported by Clermont Auvergne University. This work is supported by the French Ministry of faustine.souc@inrae.fr Research and National Research Agency as part of the French metabolomics and fluxomics infrastructure (MetaboHUB-ANR-INBS-0010).











Code source

