

The openCARP cardiac electrophysiology simulator

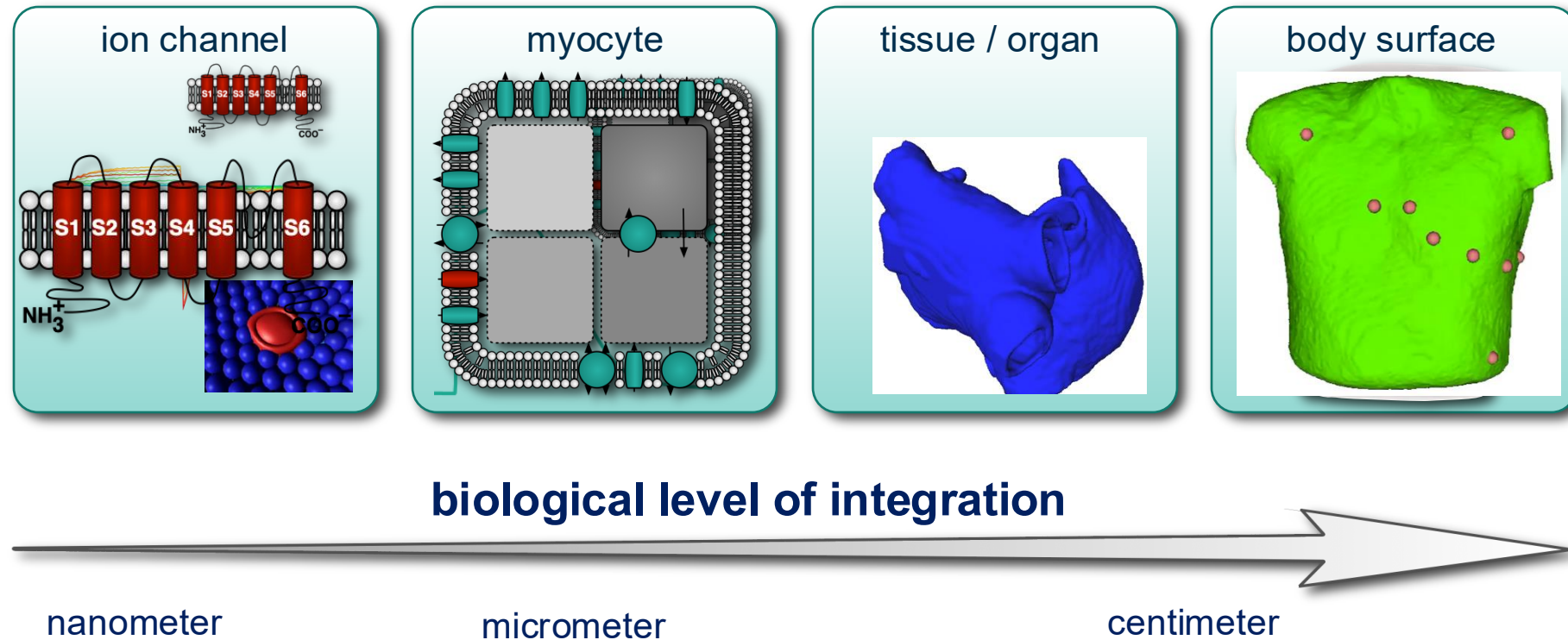
www.openCARP.org
helmholtz.software/software/opencarp

Axel Loewe
Institute of Biomedical Engineering (IBT)

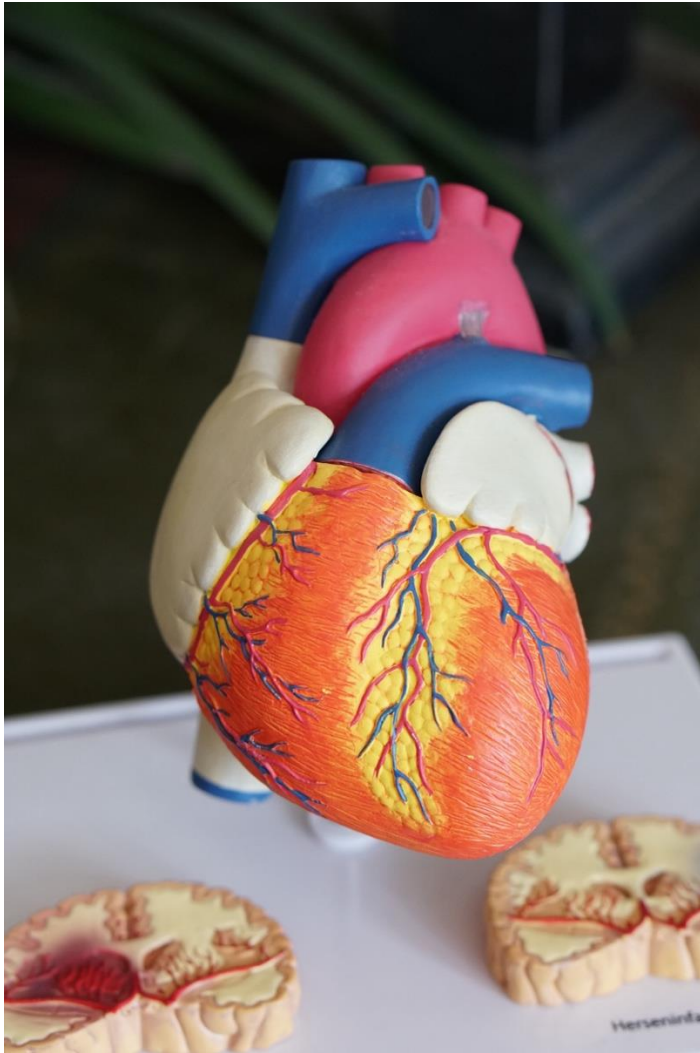


1. How can mathematics and software help address heart diseases?
2. The openCARP ecosystem
3. Research Software Engineering infrastructure

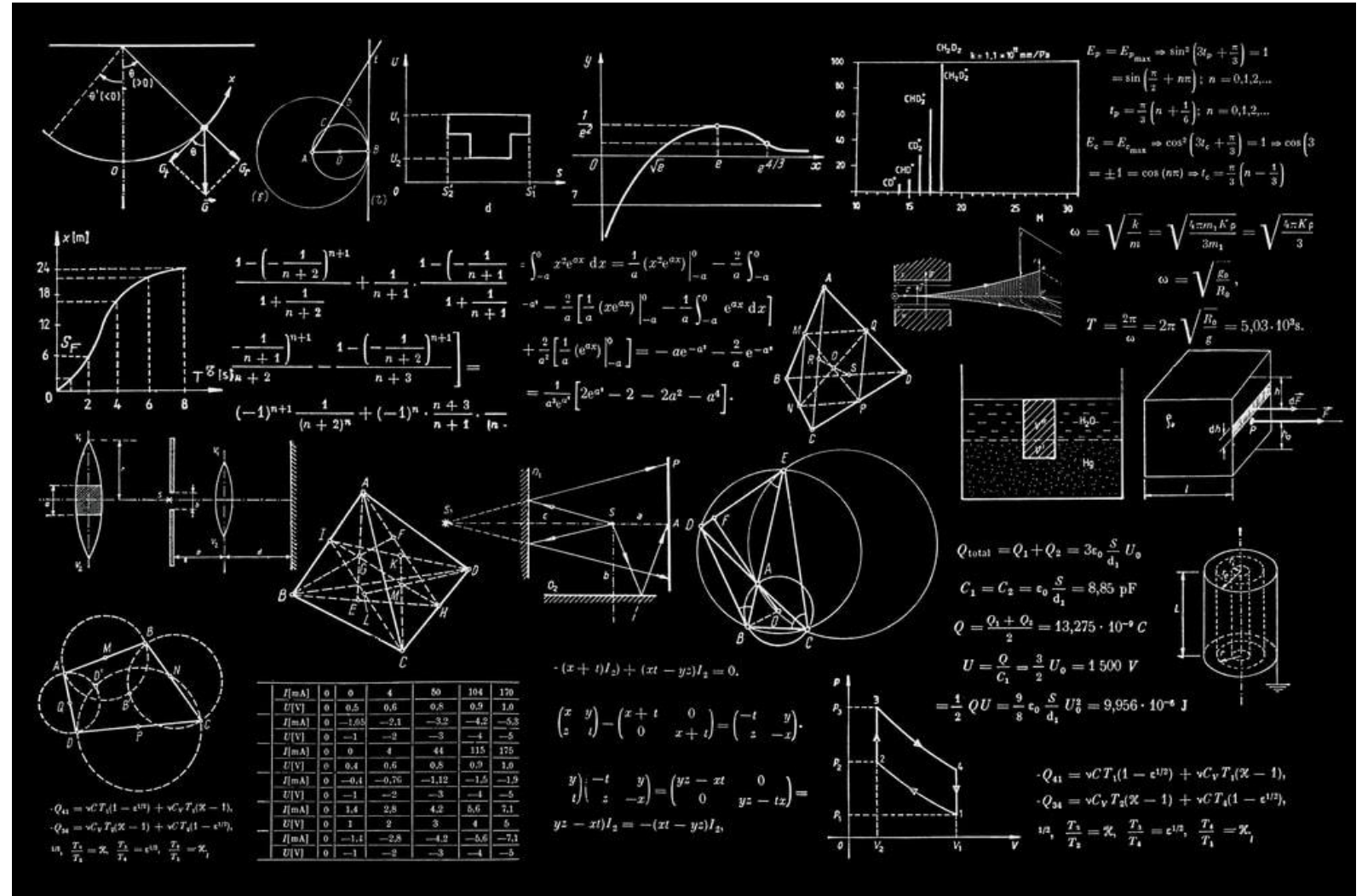
The heart as a multi-scale system



How can math help?



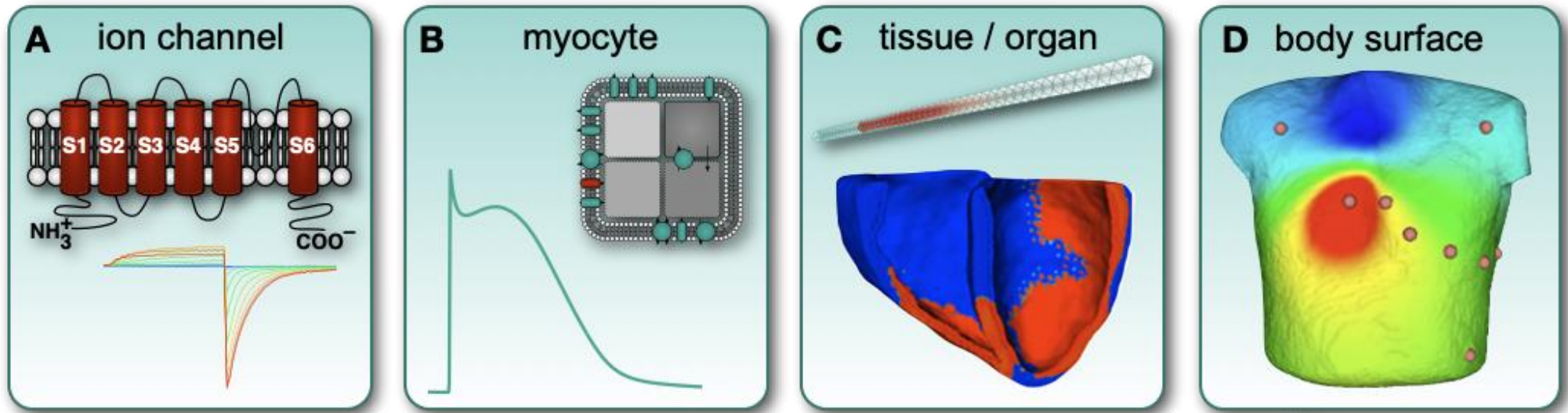
Robina Weermeijer, unsplash.com



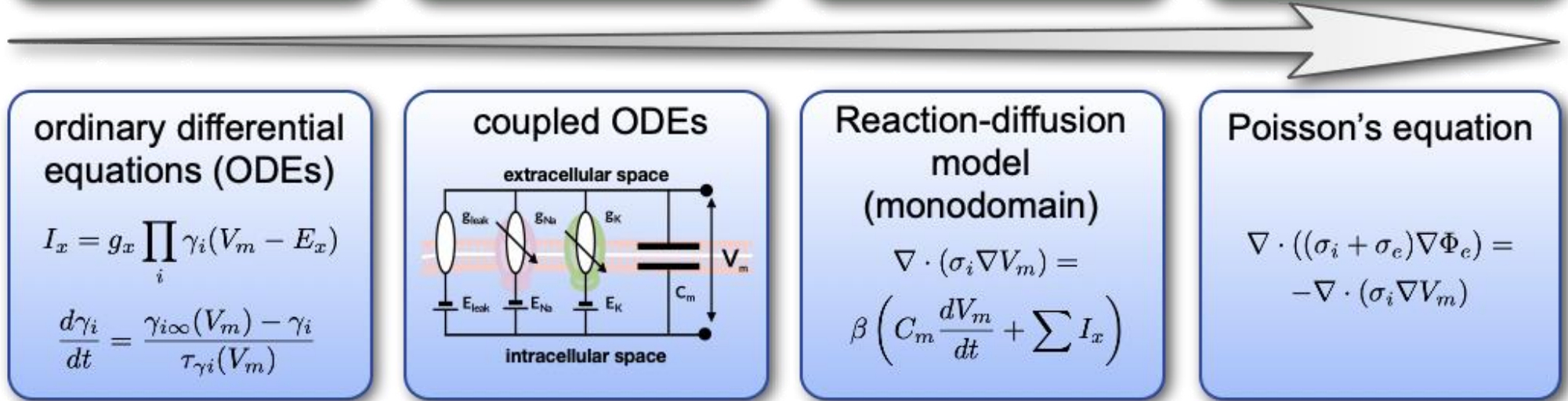
Dan-Cristian Pădureț, unsplash.com

Multi-scale system modeling of cardiac electrophysiology

Biological Level
of Integration

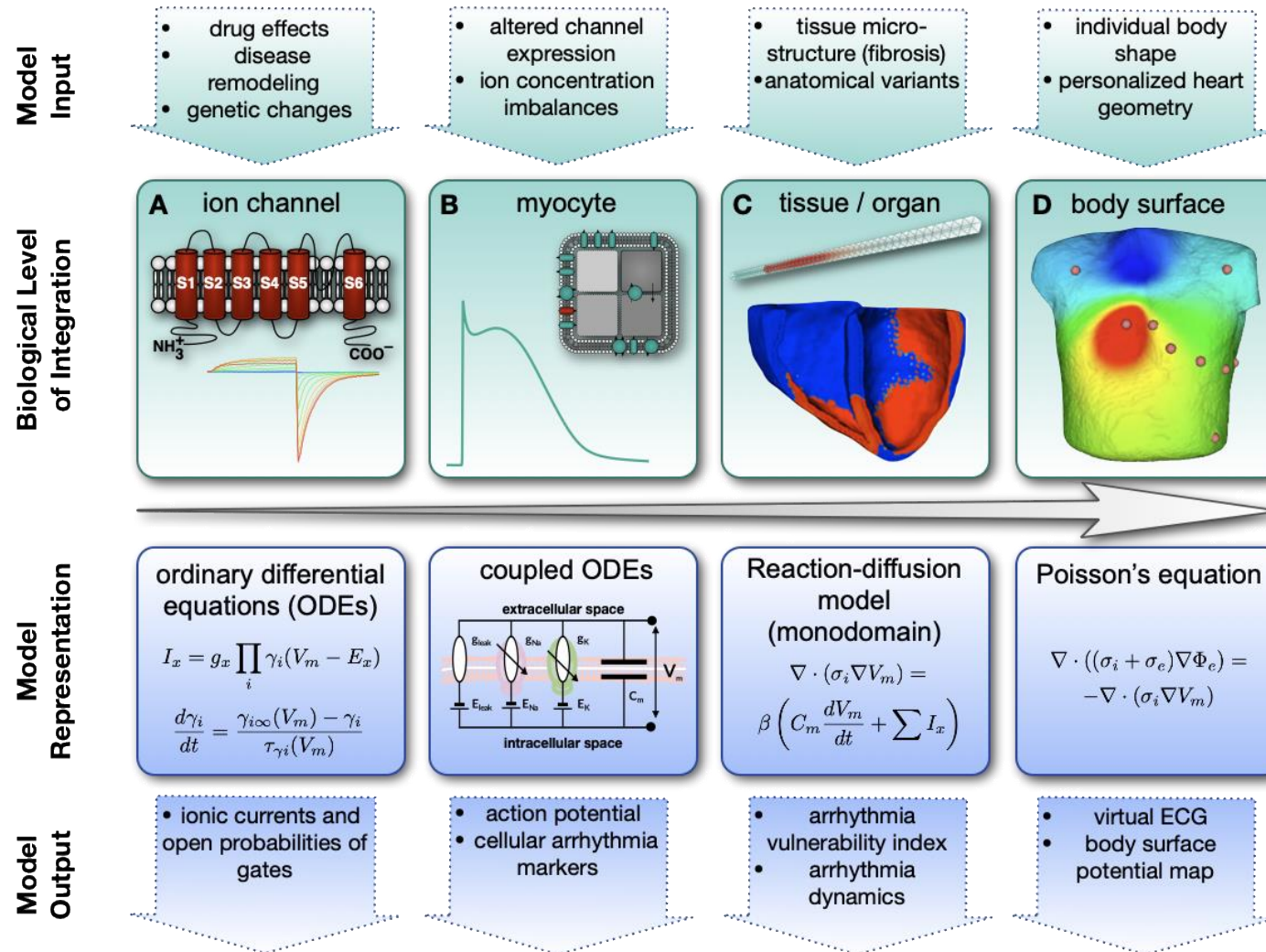


Model
Representation



Loewe et al. "Computational modelling of biological systems now and then: revisiting tools and visions from the beginning of the century" PTRSA 2025;383:20230384

Multi-scale system modeling of cardiac electrophysiology



Computer modeling & simulation

Controlled research environment

- to understand fundamental physiology and pathomechanisms
- to evaluate diagnostic and therapeutic approaches (in silico studies)
- to optimize device design

Personalised medicine through digital twins



Kate Miller



DOC RABE Meda / stock.adobe.com

DSL Vorhofflimmern

Deutsche Herzstiftung

Azzolin et al., „Personalized ablation vs. conventional ablation strategies to terminate atrial fibrillation and prevent recurrence“, *Europace* 2023;25:211-22

Computer modeling & simulation

Controlled research environment

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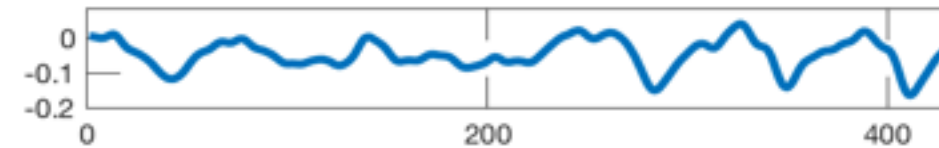
Personalised medicine through digital twins

Scalable research environment to generate big, quality-controlled dataset for machine learning

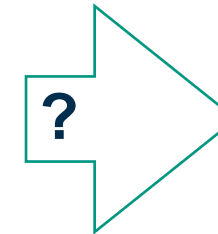
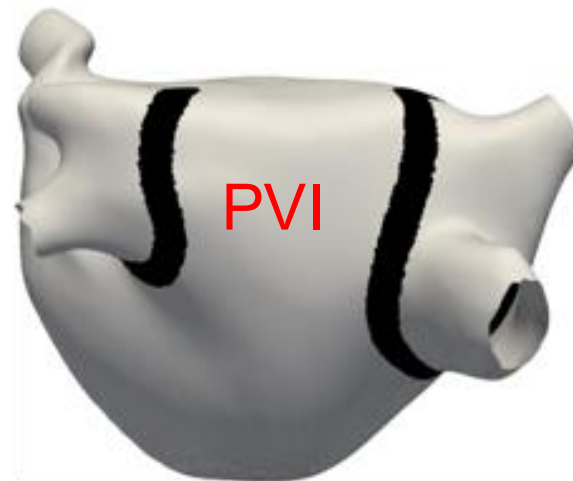
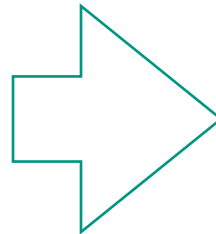


Will pulmonary vein isolation be successful?

ECG



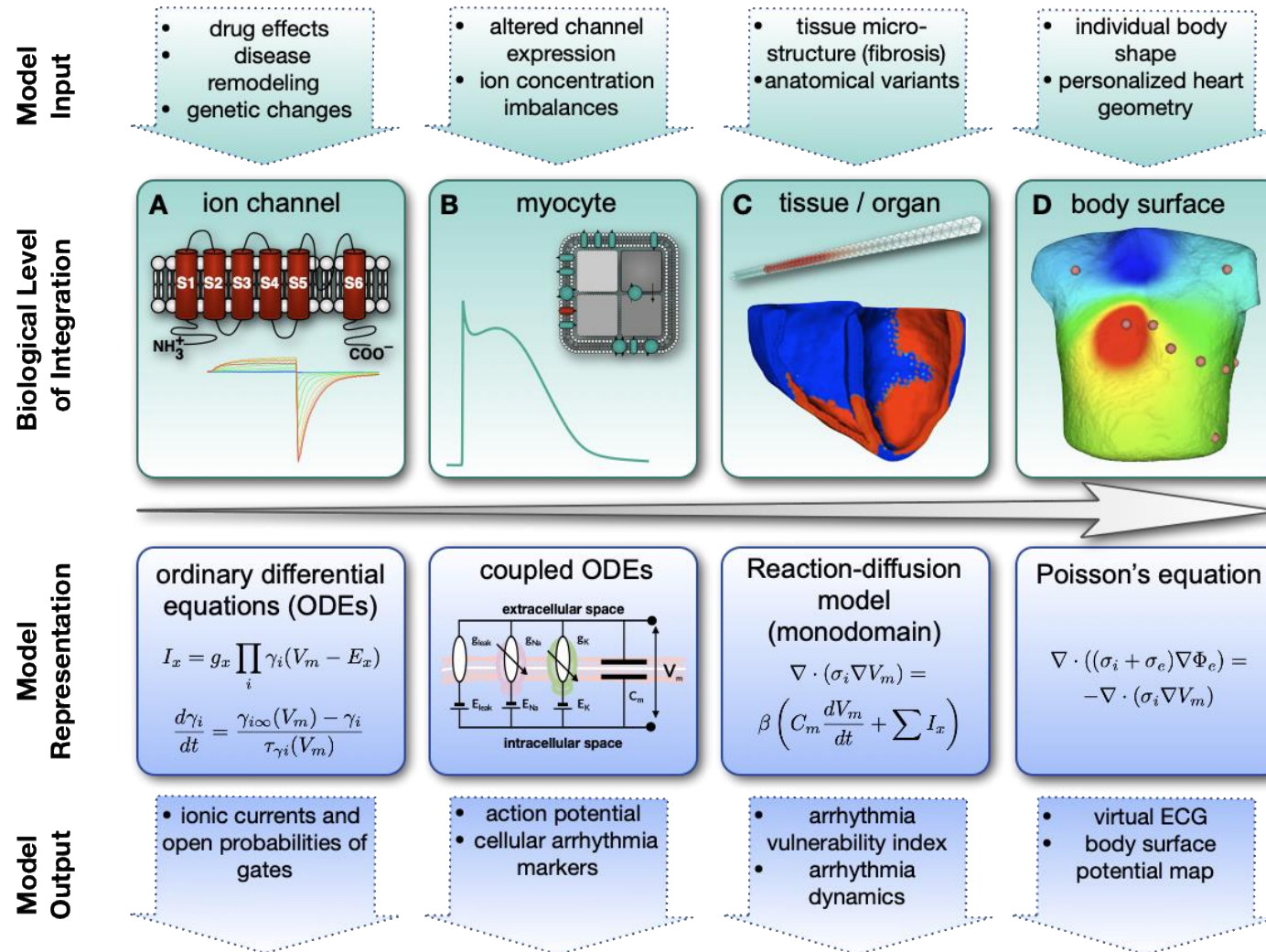
**Atrial
fibrillation**



**Atrial
fibrillation**

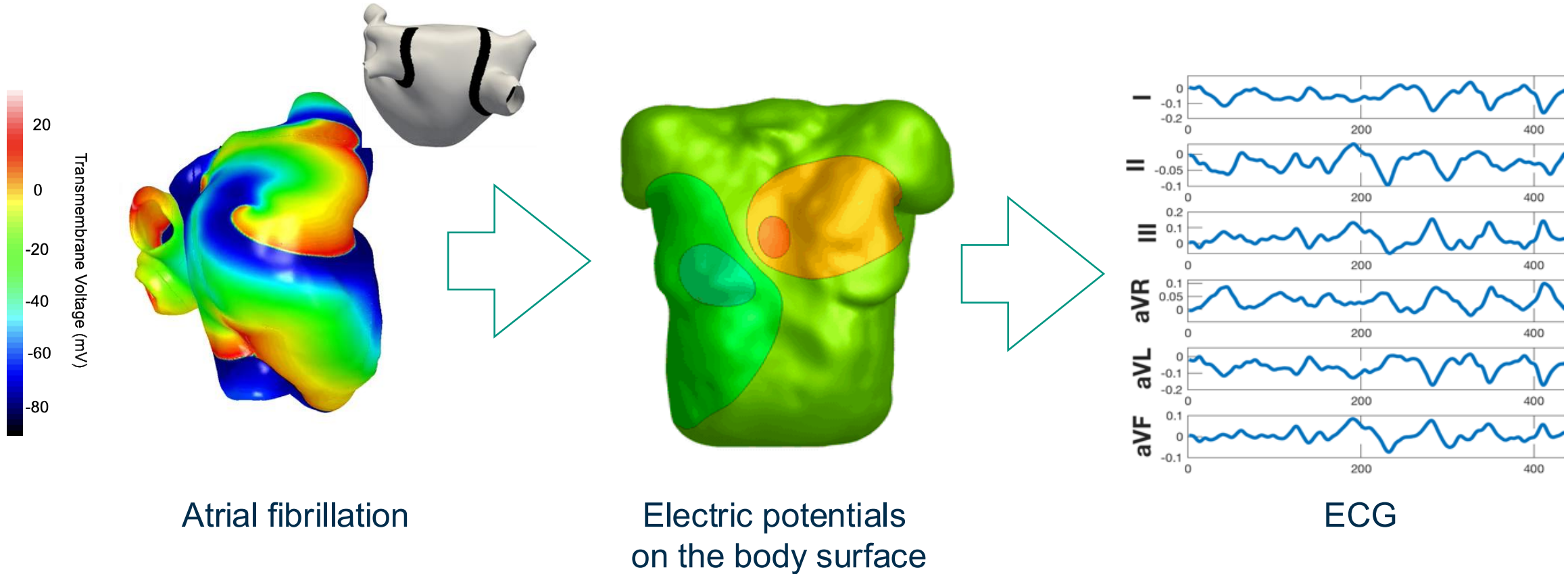


Multi-scale system modeling of cardiac electrophysiology



Loewe et al. "Computational modelling of biological systems now and then: revisiting tools and visions from the beginning of the century" PTRSA 2025;383:20230384

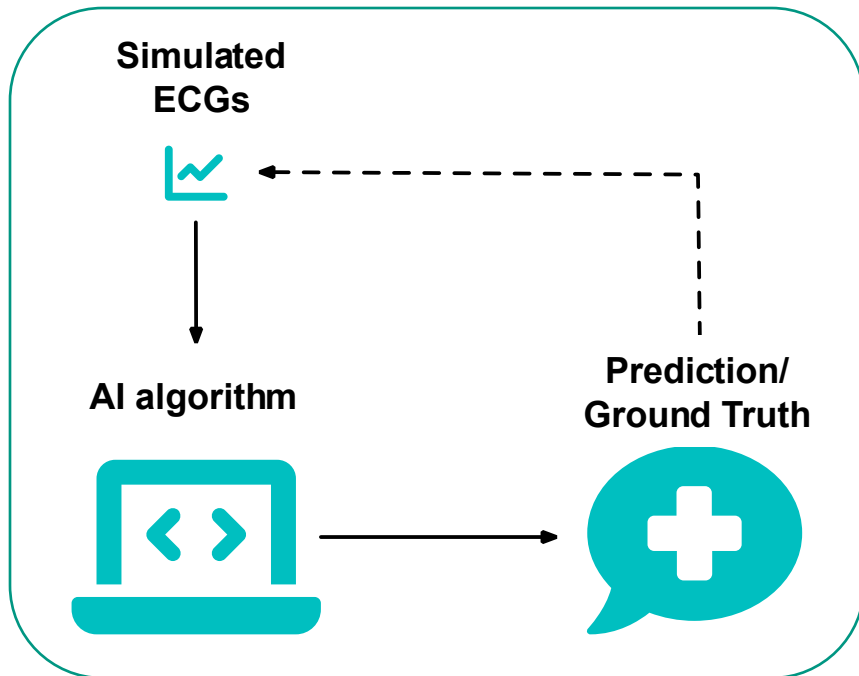
ECG simulations



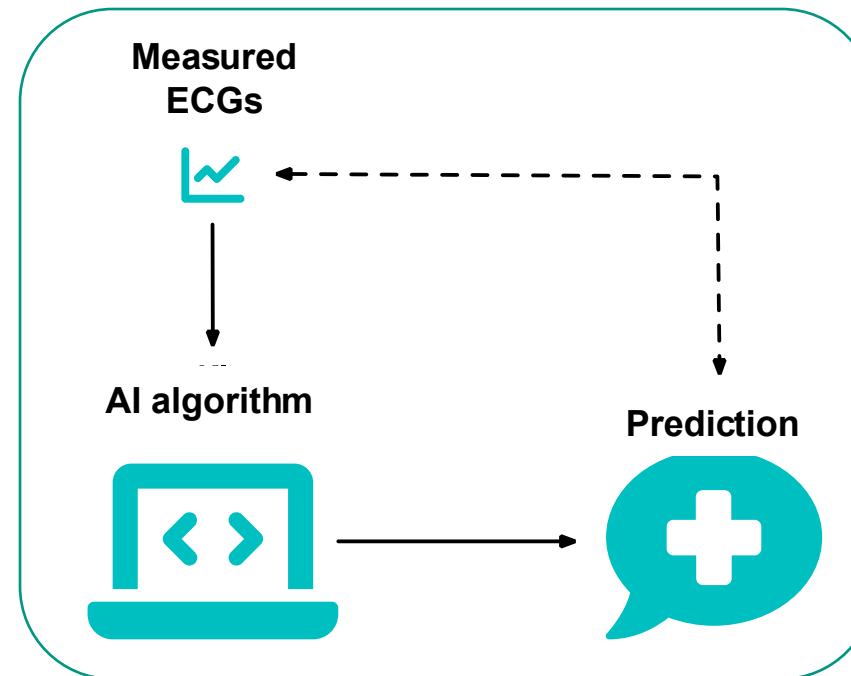
> 1000 simulated atrial fibrillation episodes

Machine learning

Training phase



Test phase



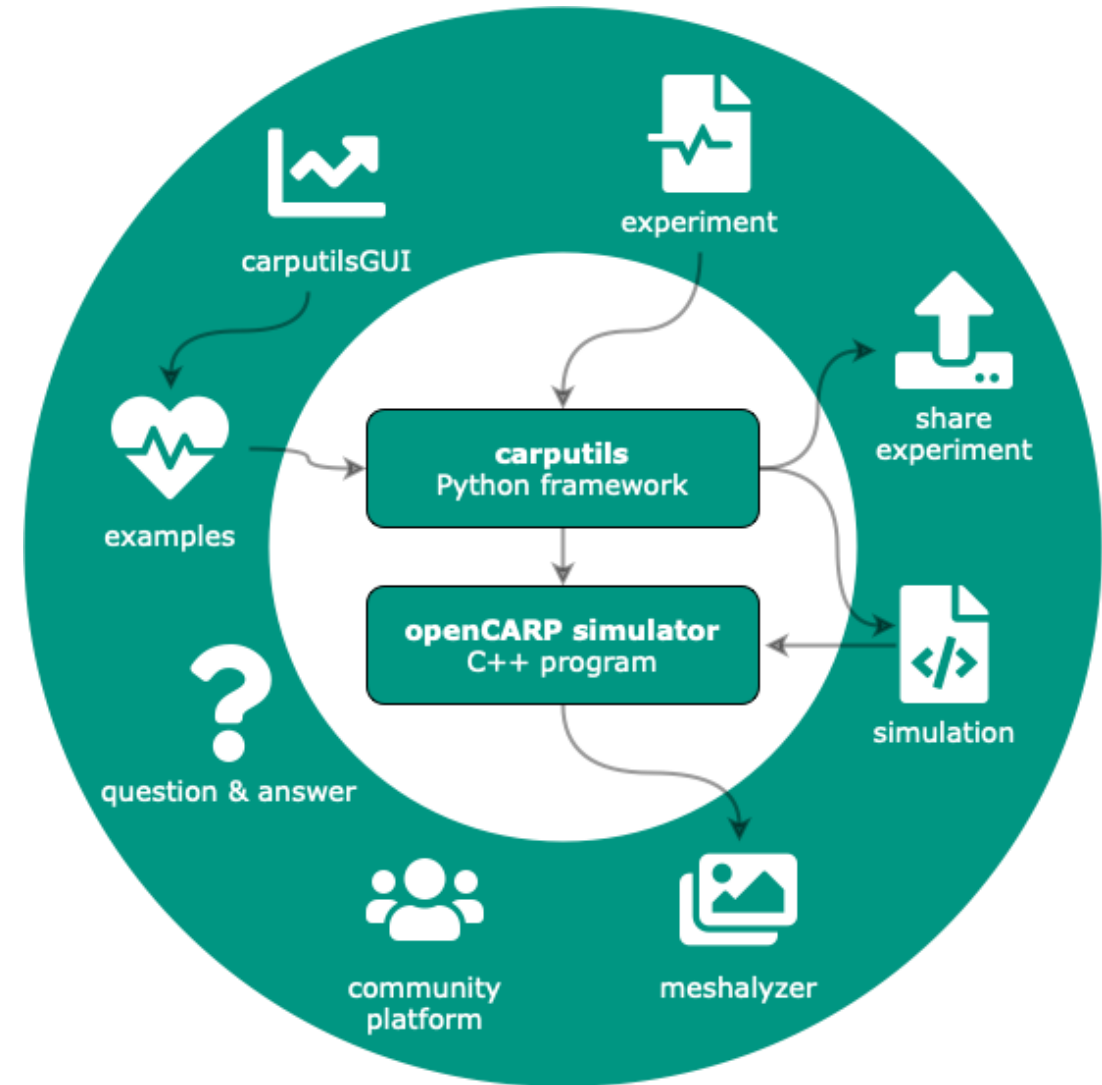
46 patients:
82.6% specificity
73.9% sensitivity
93.5% consistency

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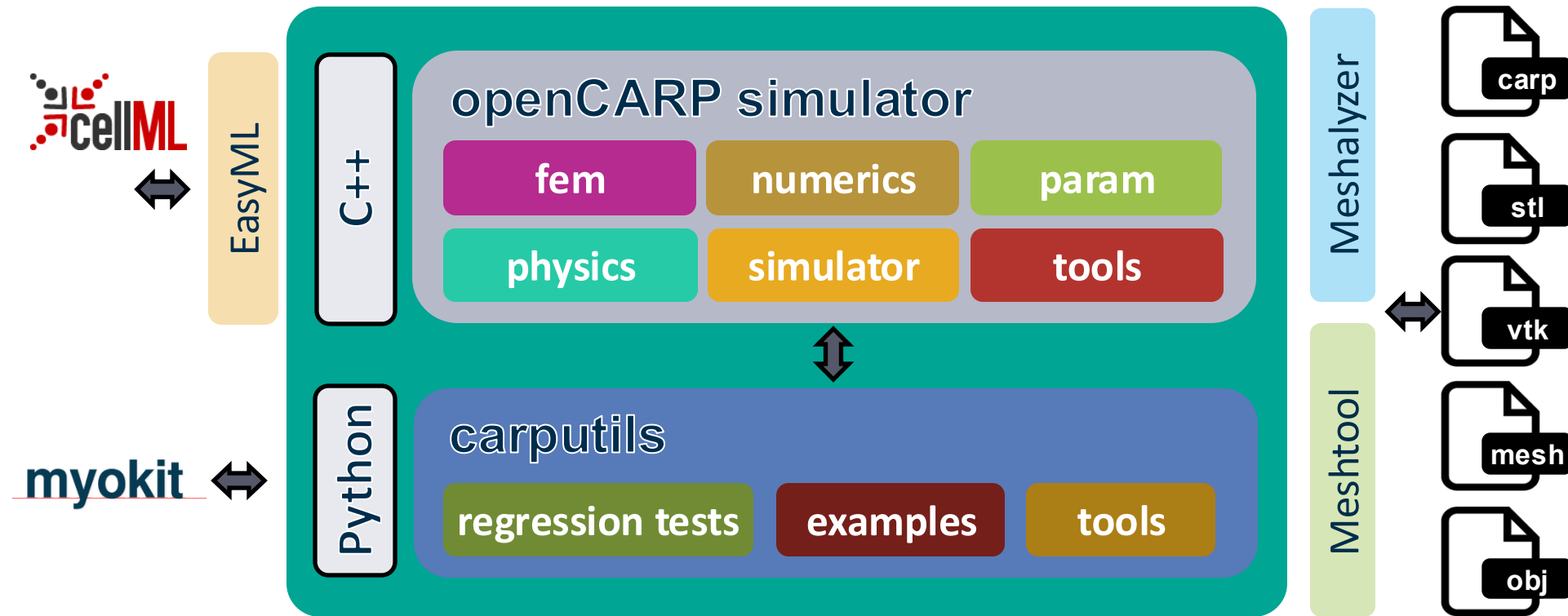
The openCARP ecosystem

openCARP: C++ cardiac electrophysiology simulator, free for academic, non-commercial use. Simulations from ion channel to organ level.

carputils: Python framework to develop complex simulation pipelines, i.e. to automate in silico experiments including all modeling and simulation steps.



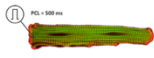
Global architecture of the openCARP simulation platform



Examples

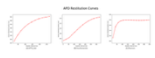
Mini-experiments coded up in carputils

www.openCARP.org > Documentation > Examples



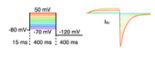
Basic single cell EP

This example introduces the basic steps of running EP simulations in an isolated myocytes




APD restitution

Action potential duration (APD) restitution example in single cell. As pacing frequency is increased, APD shortens to maintain a one to one stimulus to responses




Voltage clamp

This example explains the basic usage of bench for performing voltage clamp experiments.




EasyML to C code

This tutorial explains the basics of using the code generation tool limpet_fe.py for generating ODE solver code for models of cellular dynamics.



EasyML basics


Here you will get more information about the cell model math format EasyML.



Import CellML

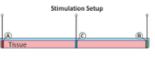
Here you learn how to import CellML data into openCARP and what problems might occur during this process

single cell simulations



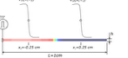
Basic tissue EP

This example introduces to the basics of using the openCARP executable for simulating EP at the tissue and organ scale



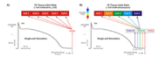
Extracellular stimulation

In this example you learn how to stimulate a tissue from the extracellular space




Tuning Conduction Velocity

This tutorial introduces the background for the relationship between conduction velocity and tissue conductivity




Init tissue from cell

This tutorial demonstrates how to initialize a cardiac tissue with state variables obtained from a single-cell stimulation



Adjust parameters

This tutorial demonstrates how to adjust parameters in tissue simulations to match experimental data for conduction velocity, APD, and wavelength



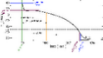
CV restitution

This example demonstrates how to compute conduction velocity restitution in cardiac tissue

tissue simulations


visualization

pre- and postprocessing



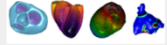
IGB utils

This example demonstrates several utilities to manipulate IGB files.



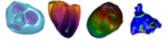
limpetGUI

limpetGUI is a simple tool for visualizing traces output generated by bench



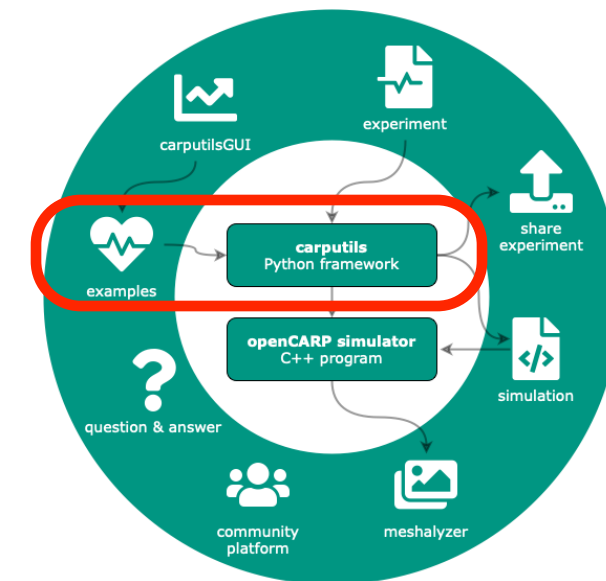
Meshalyzer

meshalyzer is a 4D visualization tool for openCARP allowing interactions to investigate data



Meshalyzer II

A few advanced features of meshalyzer are explained here to make life simpler

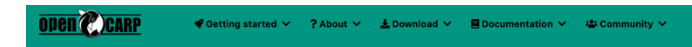
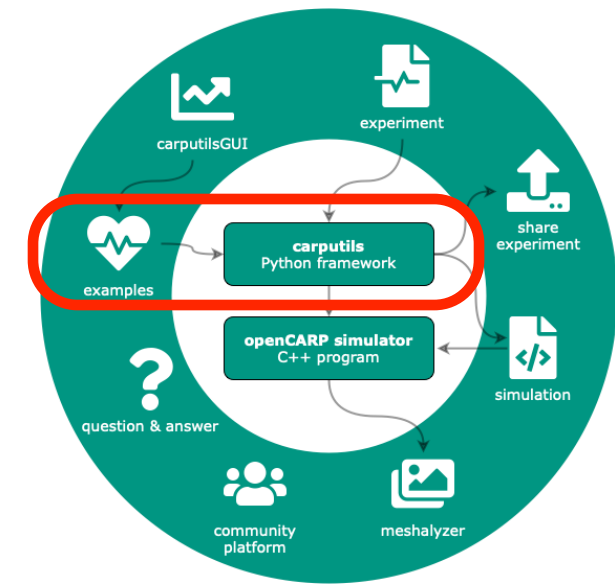


Onboarding Tutorials

Interactive Jupyter notebooks

www.openCARP.org > Getting Started > Onboarding Tutorials

The screenshot shows a JupyterLab interface for the 'openCARP onboarding' tutorial. The title bar indicates 'tutorial.ipynb'. The left sidebar shows a file explorer with a folder icon and a search icon. The main content area displays the tutorial title 'Single Cell Tutorial' and the subtitle 'Basic functionality I'. Below the title, the author 'Gunnar Seemann' and the date 'January 29, 2024' are listed. The 'Introduction' section contains three informational boxes: 'New to Jupyter Lab?' with a link to a video, 'Don't know how to proceed in a tutorial?' with instructions on using the 'display_quiz' command, and 'Important' with a note about pink background output. The interface includes a top menu bar with 'File', 'Edit', 'View', 'Run', 'Kernel', 'Tabs', 'Settings', and 'Help'.



Onboarding tutorials in openCARP JupyterLab

You will have the best experience with the tutorials, if you run openCARP JupyterLab in a local Docker installation. Follow the [instructions](#) to set up Docker locally. After the openCARP Docker is running, you should reload this page. If you try this on a Mac, please don't use Safari, as you wouldn't get a connection to localhost. Firefox or Chrome should work.

If you don't want to use a local Docker version, you can continue with just clicking on the cards. mybinder will be opened, but you have limited resources, it will take some time to start the front page and all data and input will be lost if you restart after some time.

Single Cell Tutorials

Aim of the single cell tutorials is to make you familiar with the command line interface of the single cell electrophysiology model program bench to develop single cell experiments. You also will learn basics about carputils. We strongly suggest to go through them in sequence.

All onboarding tutorials will be opened on [mybinder](#).

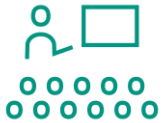
Three cards representing the single cell tutorials. The first card, 'Overview page of onboarding tutorials', features a lightbulb icon and a link to the overview page. The second card, 'Basic functionality I', features a line graph icon and a link to the first tutorial. The third card, 'Basic functionality II', features a line graph icon and a link to the second tutorial. Each card includes a brief description of the tutorial's content.

Community platform

www.openCARP.org > Community



Newsletter



User & contributor meetings



Q&A forum



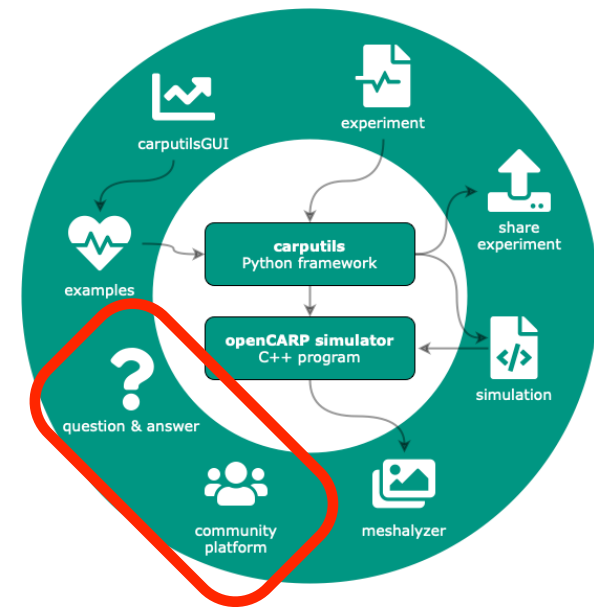
Issue tracker



Share experiments



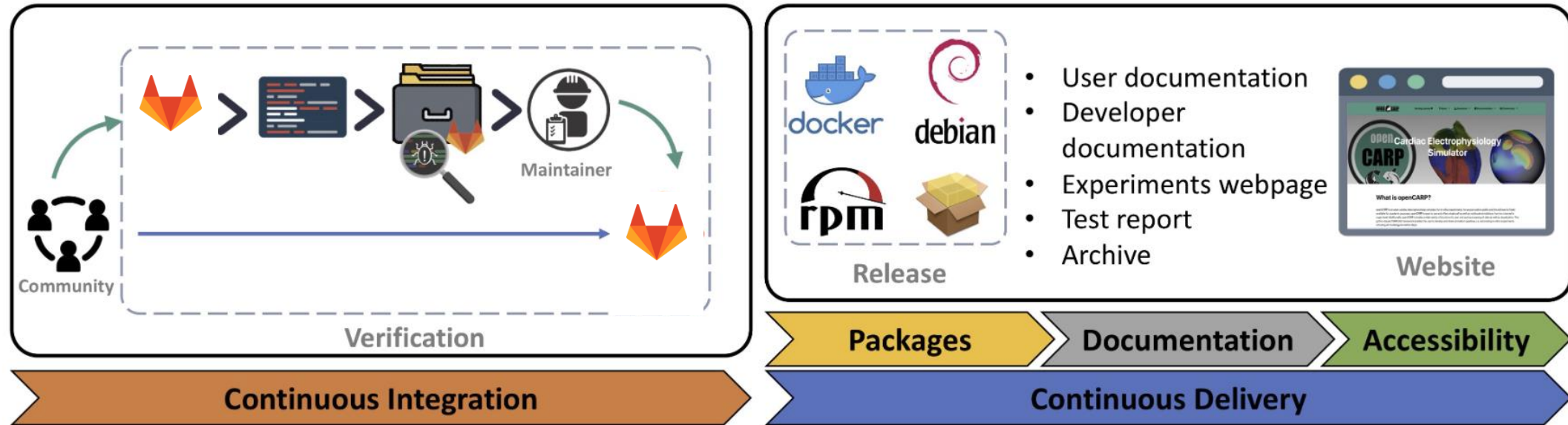
Contributing



Bach et al. "The openCARP CDE – Concept for and implementation of a sustainable collaborative development environment for research software".
Bausteine Forschungsdatenmanagement 2022;2022(1):64–84. DOI: 10.17192/bfdm.2022.1.8368.

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Infrastructure to support maintainability and sustainability



Collaborative development environment based on Gitlab, automated integration and delivery.

Maintainability: version control, code review, automated testing, automated benchmarking, automated generation of packages, documentation & website content.

Sustainability: automated long-term preservation and citable publishing of software releases along with relevant metadata on persistent research data repository.

Bach et al. "The openCARP CDE – Concept for and implementation of a sustainable collaborative development environment for research software".
Bausteine Forschungsdatenmanagement 2022;2022(1):64–84. DOI: 10.17192/bfdm.2022.1.8368.

The FAIR4RS Principles

Findable: Software, and its associated metadata, is easy for both humans and machines to find.

Accessible: Software, and its metadata, is retrievable via standardised protocols.

Interoperable: Software interoperates with other software by exchanging data and/or metadata, and/or through interaction via application programming interfaces (APIs), described through standards.

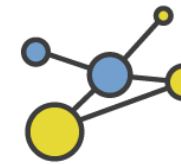
Reusable: Software is both usable (can be executed) and reusable (can be understood, modified, built upon, or incorporated into other software).



Findable



Accessible



Interoperable



Reusable

FACILE-RS: Automated Metadata Conversion and Software Publication Based on CodeMeta



Publishing software according to the **FAIR Principles for Research Software (FAIR4RS)** increases transparency, reproducibility, and reusability of research.

Adopting the FAIR4RS principles requires substantial effort from developers, including:

- maintaining software metadata in several standard formats (DataCite, Citation File Format (CFF), CodeMeta, ...),
- assigning each software version a unique persistent identifier.

FACILE-RS allows to **generate various metadata formats** from CodeMeta metadata and to **publish software on reputable research data repositories** in an automated way.

FACILE-RS: archival and long-term preservation of research software repositories made easy

Marie Houllon¹, Jochen Klar², Ziad Boutanios¹, Tomas Stary¹, Terry Cojean^{1,3}, Hartwig Anzt^{1,3}, and Axel Loewe¹

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DOI: 10.21105/joss.07330

Software

- Review
- Repository
- Archive

Editor: Ana Trisovic

Reviewers:

- @scaera
- @NicolettaNinkovic1

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Published: 26 June 2025

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Summary

The Python package FACILE-RS (Findability and Accessibility through Continuous Integration with Less Effort for Research Software) facilitates the archival and long-term preservation of research software repositories. It consists of a set of scripts that simplify the maintenance of software metadata by automating its generation and synchronization in various formats from a single manually maintained metadata file. FACILE-RS also makes it easier to publish and archive software releases according to the Open Science paradigm and the FAIR (Findable, Accessible, Interoperable, Reusable) principles for Research Software by offering tools to automate the creation of releases and the upload to persistent research data repositories.

In particular, FACILE-RS automates:

- Creating a DataCite record (DataCite Metadata Working Group, 2021) based on CodeMeta files (Boettiger, 2017) present in repositories
- Creating a CFF (Citation File Format) file (Druskat et al., 2021) from CodeMeta files
- Creating archive packages in the BagIt (Kunze et al., 2018) or the BagPack (RDA Research Data Repository Interoperability WG, 2019) formats
- Creating a release on the GitLab development platform using the GitLab API
- Archiving software releases persistently on Zenodo
- Archiving software releases persistently using the RADAR service (Kraft et al., 2016)
- Using content from Markdown files, BibTeX files, or Python docstrings to create web pages within the Grav CMS

The scripts can be run manually, but they have been designed to be used within workflow automation systems such as GitLab CI/CD or GitHub Actions, in order to reduce the need for manual intervention when maintaining metadata and creating persistent software releases.

Statement of need

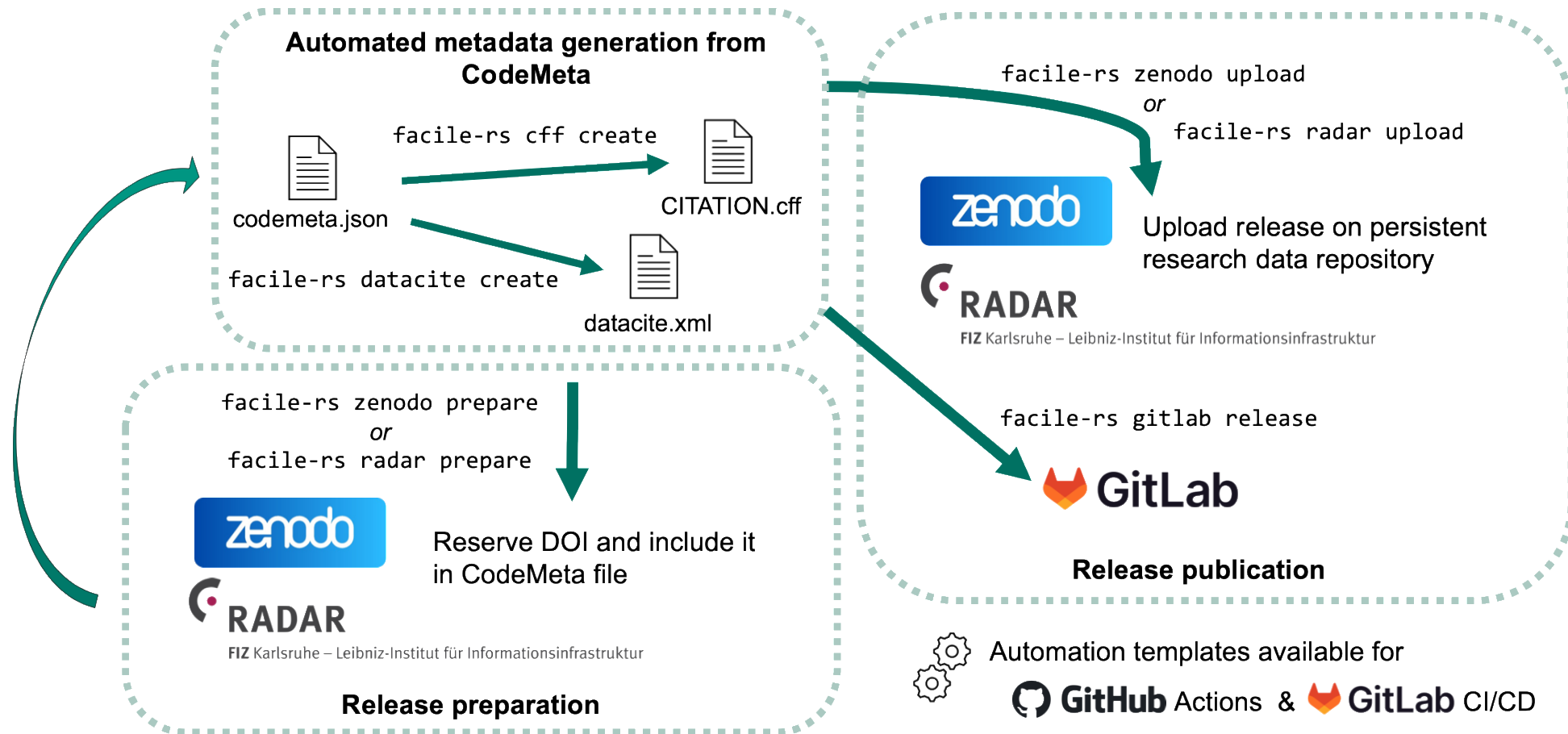
Research software development is a fundamental aspect of academic research (Anzt et al., 2021), and it is now widely acknowledged that the FAIR principles (Wilkinson et al., 2016), established to improve the reusability of research data, should also be applied to research software. However, specific aspects of research software like executability or evolution over time require these guidelines to be adapted. Therefore, the FAIR principles for Research Software (FAIR4RS) have been introduced (Chue Hong et al., 2021).

In particular, reproducible research requires software and associated metadata to be easily findable by both machines and humans, and retrievable via standardised communication protocols. In this context, several metadata standards are widely used across the scientific community:

Houllon et al. (2025). FACILE-RS: archival and long-term preservation of research software repositories made easy. *Journal of Open Source Software*, 12(110), 7330. <https://doi.org/10.21105/joss.07330>

Houllon et al. “FACILE-RS: archival and long-term preservation of research software repositories made easy”. *JOSS* 2025; accepted

FACILE-RS: Findability and Accessibility through Continuous Integration with Less Effort for Research Software



<https://git.opencarp.org/openCARP/FACILE-RS>

Houillon et al. "FACILE-RS: archival and long-term preservation of research software repositories made easy". JOSS 2025; accepted

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Summary



Lessons learned

- **Collaboration** makes many things easier but some also more complex
- Fostering a **user community** is effort worthwhile
- **Automating** routine tasks makes life easier but also these need to be maintained
- Positive: increasing awareness for **software as research infrastructure**

Open challenges

- Uptake of best practices can be **sluggish**
- Maintaining **know-how** in teams with very few permanent members
- Legal and fiscal **sponsorship**
- **Scaling** support to a growing user base
- Tracking **usage**
- **Maintaining “smaller” software**



EuroHPC
Joint Undertaking



Co-funded by
the European Union



Research Software Engineering (RSE) at KIT