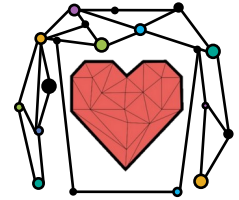


Web Interface for Cardiac Simulations

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From Proposal for WP1:

6. Integrate the components delivered by the other work packages into a production-ready HPC simulation platform for cardiac electrophysiology.

Results of a Survey

- 28 people (20 users) participated in the survey
- The majority runs bench and openCARP directly
- People have difficulties with installing dependencies and compiling openCARP (9 out of 15 that install openCARP from sources)
- Only a few (3) use docker

Desired features:

- Allow to retrieve results of the simulation
- Easy run on HPC resources
- Connection with published experiments
- Visualization in the web-based system
- Templates for carputils experiments
- Submission of carputils experiments from the interface
- Interactive control of the parameters of carputils script
- Access to carputils tutorials
- Simplified sharing of experiments with the community

Target groups

- Beginner users – example tutorials to facilitate learning
- Experienced users – implementation of own experiments

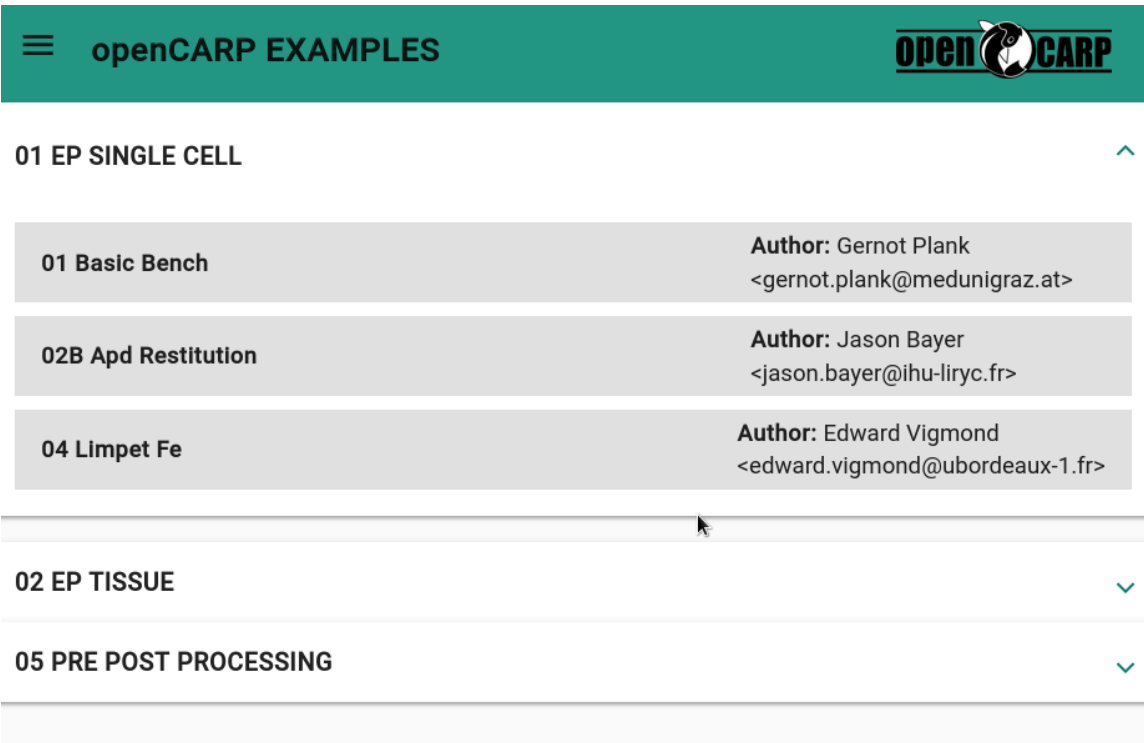
Technology

- Carputils GUI
- Streamlit or Trame Python frameworks
- JupyterLab

Carputils GUI

- Python backend using Flask API
- JavaScript frontend using Angular framework
- ParaViewWeb for visualizations
- Postgress database
- Nginx web server
- Deployed as docker-compose services

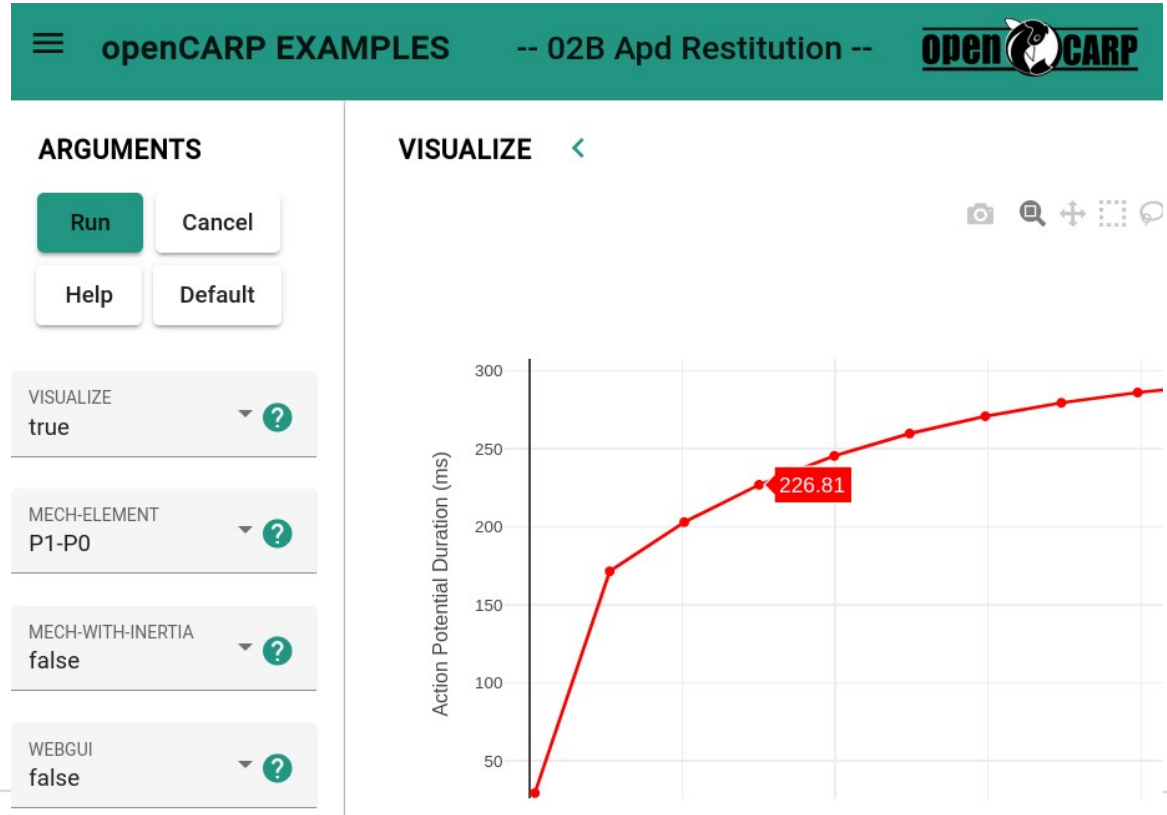
- Automatic conversion from selected tutorials
- Both single cell and tissue simulations
- Self-hosted or provided as service



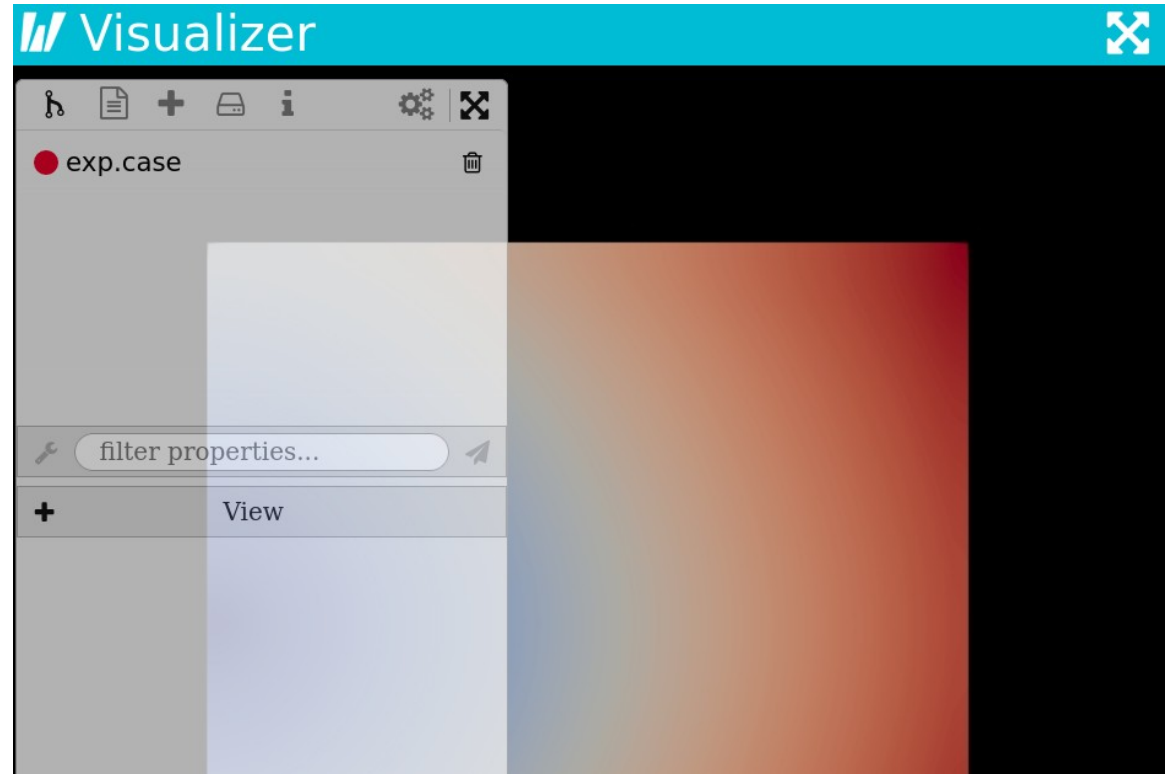
The screenshot shows the 'openCARP EXAMPLES' section of the Carputils GUI. It features a green header with a hamburger menu icon and the text 'openCARP EXAMPLES'. Below the header, there are three main categories: '01 EP SINGLE CELL', '02 EP TISSUE', and '05 PRE POST PROCESSING'. Each category is followed by a list of specific examples with their authors and contact information.

| Example ID | Author | Contact |
|---------------------|----------------|---------------------------------|
| 01 Basic Bench | Gernot Plank | <gernot.plank@medunigraz.at> |
| 02B Apd Restitution | Jason Bayer | <jason.bayer@ihu-liryc.fr> |
| 04 Limpet Fe | Edward Vigmond | <edward.vigmond@ubordeaux-1.fr> |

- Single cell experiment visualized using plotly graphics library



- Tissue level simulations visualized through ParaViewWeb interface



Streamlit and Trame

- Python frameworks for data science
- Simple implementation
- Python libraries for processing and visualization

```
# Streamlit code
st.title("Simulating the effects of channel mutations")

form = st.form(key='input parameters')
col1, col2, col3 = form.columns(3)
mutation = col1.selectbox("Channel mutation", mutations)

iks_choice = ["Loewe", "Verkerk", "Wilders"]
iks_channel = col2.radio("IKs channel", iks_choice)

duration = col3.radio("Duration of the simulation in milliseconds", (1000, 5000, 10000))

run = form.form_submit_button(label="Submit")
```

Simulating the effects of channel mutations

Channel mutation

CTRL

IKs channel

- Loewe
 Verkerk, Wilders

Duration of the simulation in milliseconds

- 1000
 5000
 10000

Submit

Streamlit Framework

- Video demonstration

- Documents combining **text**, formatted mathematical **formulas**, and **code** as well as **data**, **figures** and **tables** generated by that code
- Used to present linear data analysis pipelines
- Intuitive user interaction through selection widgets
- Large and supporting community of users
- Freely available and highly customizable

- Deployed on many HPC system such as:
 - ◆ BW HPC Center
 - ◆ Jülich Supercomputing Center
 - ◆ Cineca HPC Center

- JupyterLab interface on BWUniCluster
- Container mode allows arbitrary image, e.g. docker.opencarp.org
- JupyterLab components automatically installed

Select your resources

The grayed out fields contain a reasonable preselection of resources.
Other values can be selected in advanced mode.

Number of CPU-cores: 1

Good availability

Number of GPUs: 0

Runtime: 0.5 hour

Partition: single

Amount of memory: 4GB

JupyterLab-Basemodule: Container Mode

Auto-Reservation:

Advanced Mode:

Container Mode:

--container-image

--container-name

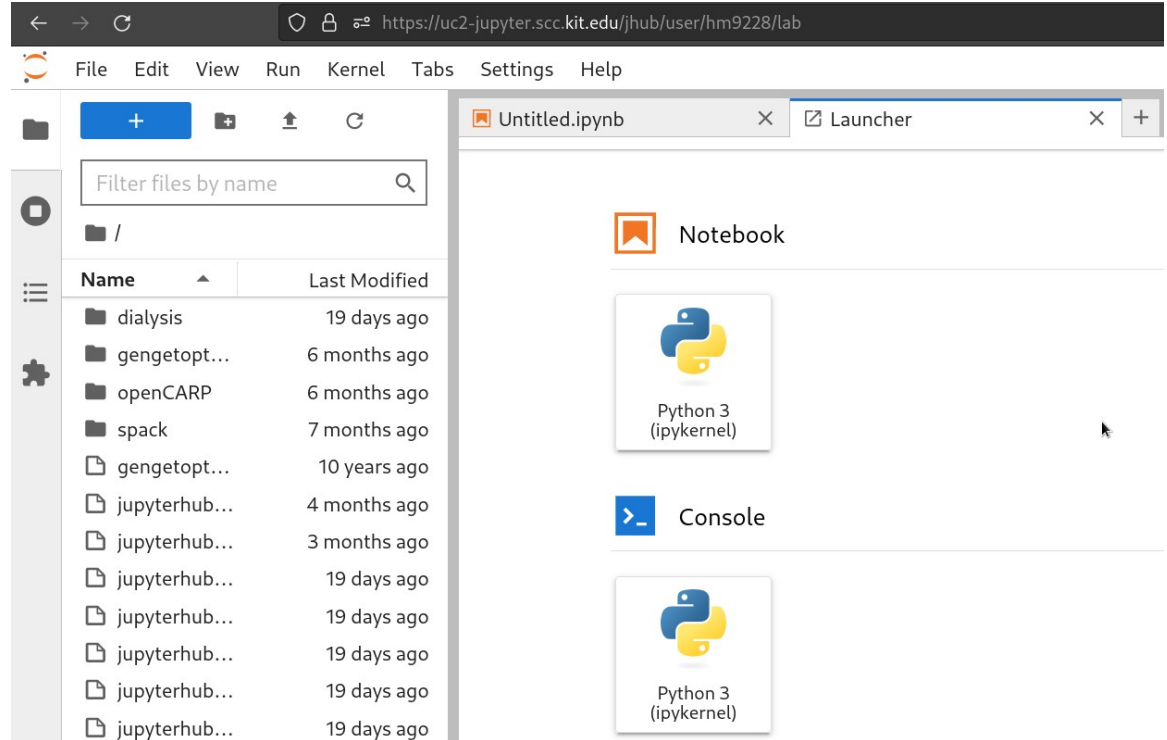
--container-mount-home

--container-mounts=<default mounts>

--no-container-remap-root

Spawn

- Mounted \$HOME directory provides persistent access to notebook files



The screenshot displays the JupyterLab web interface in a browser window. The address bar shows the URL `https://uc2-jupyter.scc.kit.edu/jhub/user/hm9228/lab`. The interface includes a top menu bar with options: File, Edit, View, Run, Kernel, Tabs, Settings, and Help. On the left, a file browser sidebar shows a search bar labeled "Filter files by name" and a list of files and folders. The files are organized into a table with columns for "Name" and "Last Modified".

| Name | Last Modified |
|---------------|---------------|
| / | |
| diagnosis | 19 days ago |
| gengetopt... | 6 months ago |
| openCARP | 6 months ago |
| spack | 7 months ago |
| gengetopt... | 10 years ago |
| jupyterhub... | 4 months ago |
| jupyterhub... | 3 months ago |
| jupyterhub... | 19 days ago |
| jupyterhub... | 19 days ago |
| jupyterhub... | 19 days ago |
| jupyterhub... | 19 days ago |
| jupyterhub... | 19 days ago |
| jupyterhub... | 19 days ago |

On the right, the main workspace area shows a "Notebook" section with a Python 3 (ipykernel) icon and a "Console" section with a Python 3 (ipykernel) icon. The browser tabs show "Untitled.ipynb" and "Launcher".



Simulation using openCARP backend

First, we define the simulation. This uses the `subprocess` module that calls openCARP and return pandas dataframe with the results.

```
In [6]: import subprocess
import pandas as pd
from random import randint

def simulation(imp_par,duration):
    n = randint(0,1e6)
    command = ["bench",
               "--numstim=0",
               "--duration={}".format(duration),
               "--imp=Loewe",
               "--dt=0.01",
               "--dt-out=0.1",
               "--fout=tmp",
               "--imp-par={}".format(imp_par),
               "-v",
               "--trace-no={}".format(n)]

    call = subprocess.run(command, capture_output=True)
    data = pd.read_csv("Trace_{}.dat".format(n), header=None, sep="\t")

    return data
```

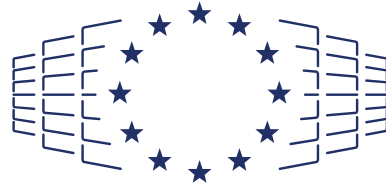
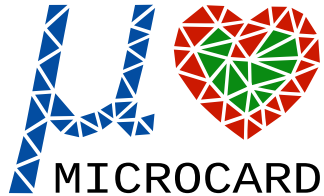
Now we prepare the input parameters

- Video demonstration

- Document usage of JupyterLab with openCARP
- Convert some tutorial into JupyterLab format
 - Extract docstrings from ``run.py`` and convert from `*.rst` to markdown
 - Add Jupyter cells in markdown
 - ``mystnb-to-jupyter`` to convert markdown to `*.ipynb` format
- Explore possible usage of meshalyzer and ParaView for 3d visualization within JupyterLab

- Write a tutorial on Streamlit or Trame framework with openCARP

Thank you for your attention!



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