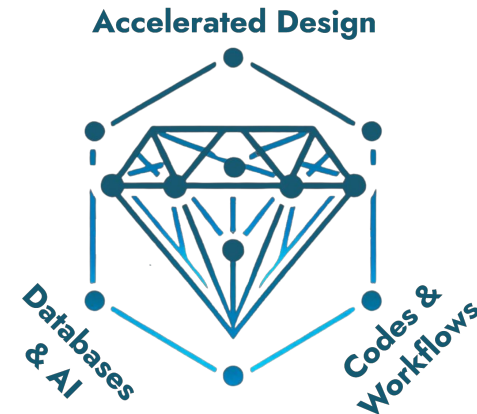


Journée utilisateurs Gricad
2025

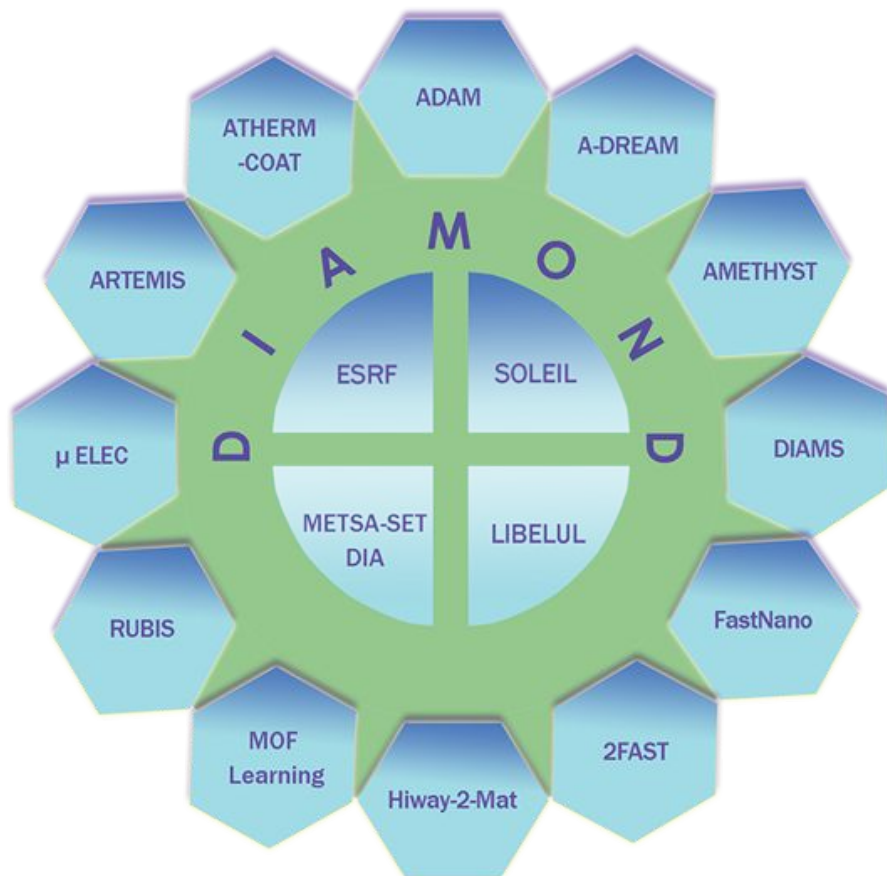
***Diamond
platform***
*For accelerated
materials design*



DIAMOND : ANR-22-PEXD-0015

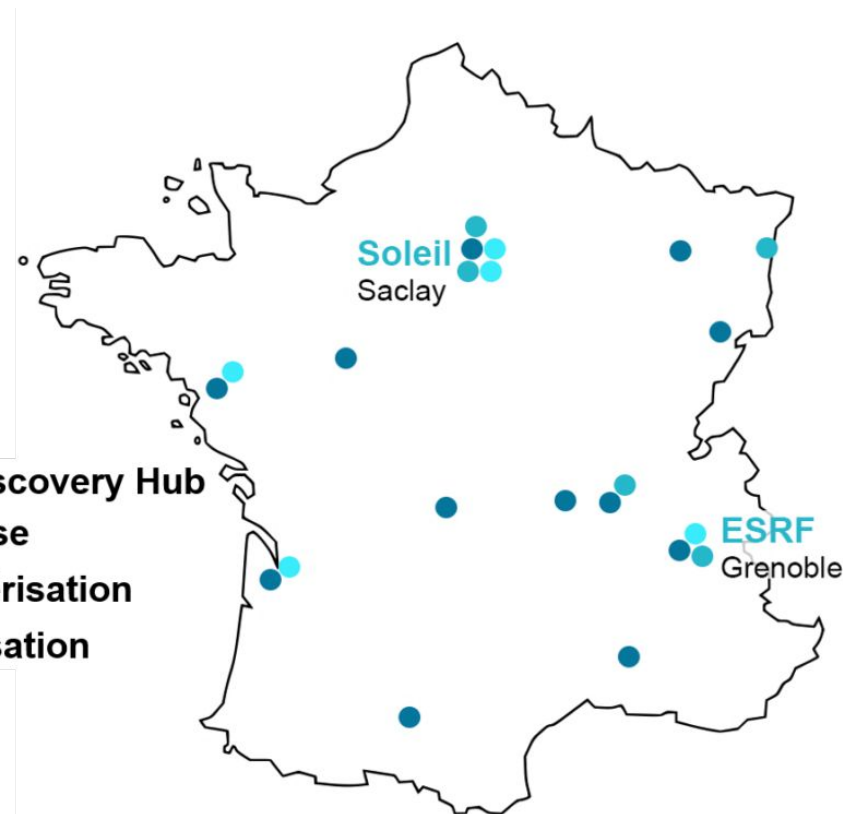


DIADEM Projects & Platforms



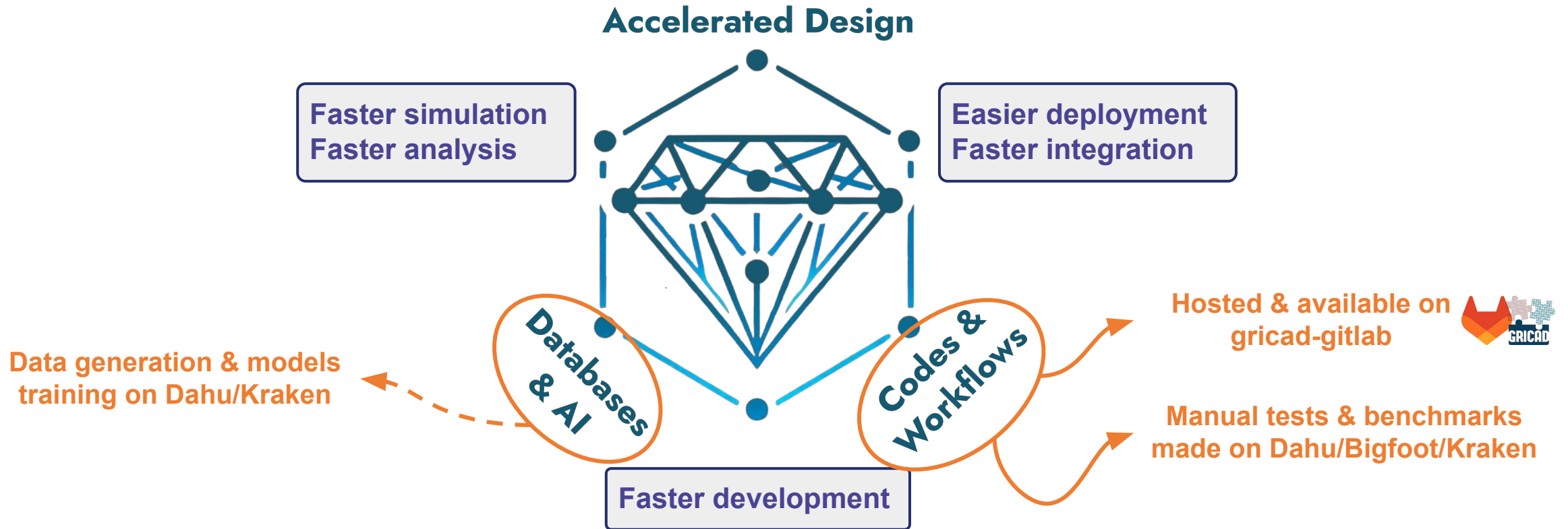
DIADEM Discovery Hub

- Synthèse
- Caractérisation
- Modélisation



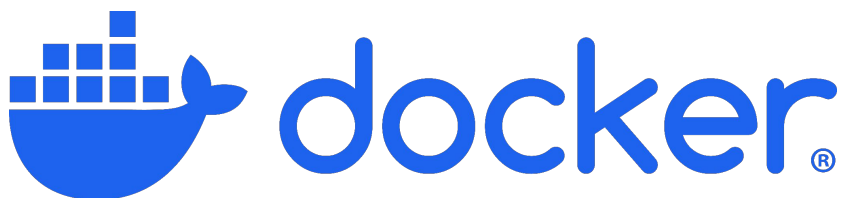
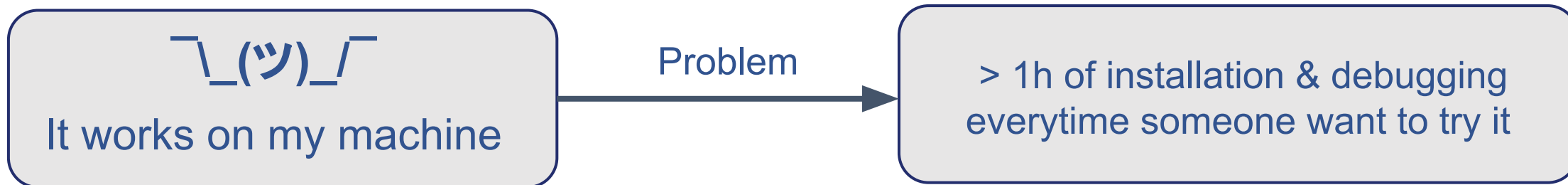


Diamond, the numerical platform





Containerization, a software need



Micro-services / Personal use



High Performance Computing



Containerization issues



build : Aug, 8th



build : Nov, 6th



Package differences
can cause **diverging
results** and/or **errors**!



Reproducibility, a scientific need

The ability to reproduce the environment of software that accompanies research work should not be considered a mundanity or an exercise that's "overkill". The ability to rerun, inspect, and modify software are the natural extension of the scientific method. Without a companion reproducible software environment, research papers are *merely the advertisement of scholarship*, to paraphrase Jon Claerbout.

- **Ludovic Courtès**, *Maintainer of Guix*



Back to
this commit



Commit History	
5azdr8k	Feb, 5th
df77a4m	Feb, 6th
...	
2abzh5f	May, 20th

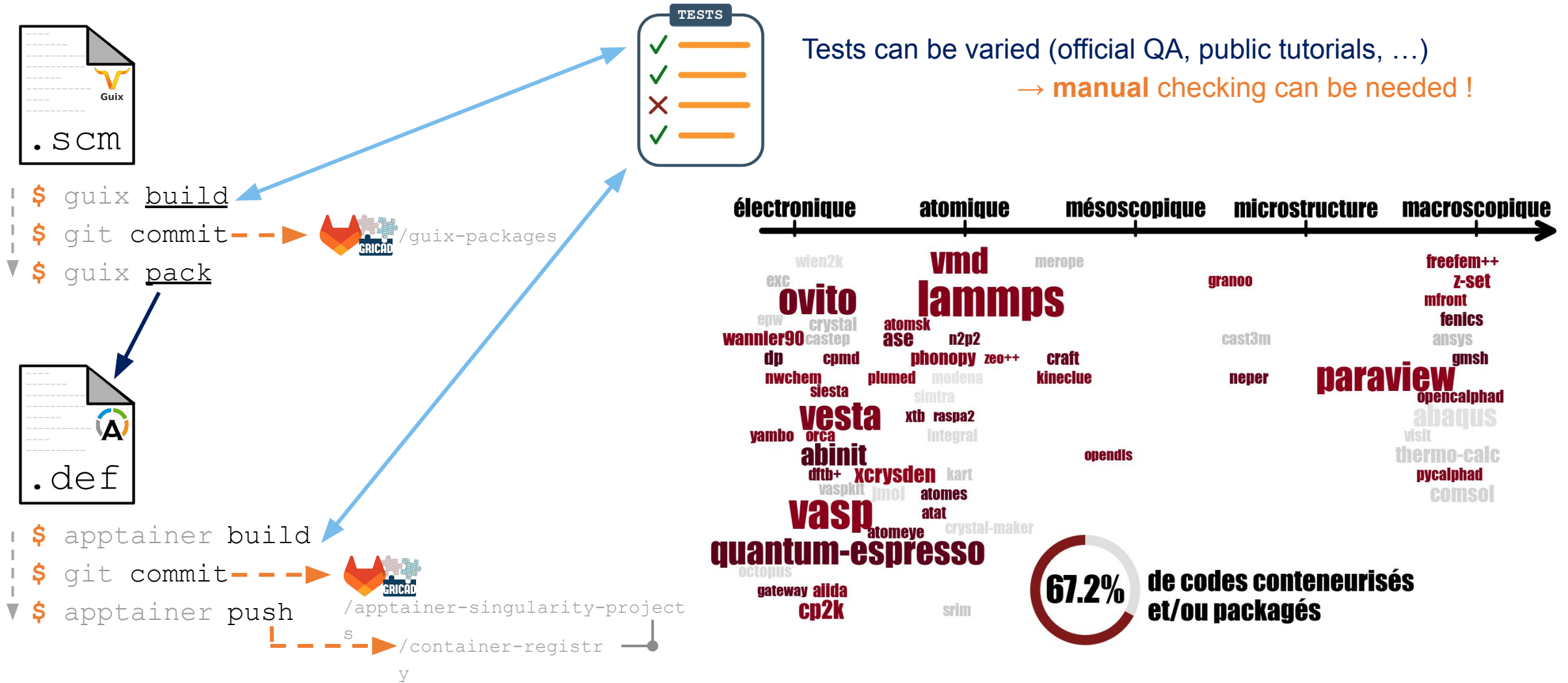
```
$ guix install python-numpy
```

```
$ guix time-machine --commit=df77a4m -- install python-numpy
```

Every dependency is frozen as it was at a given commit time



Containerization-packaging process





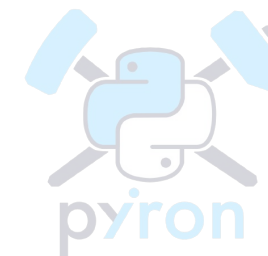
Access DIAMOND codes & workflows

- A dedicated space in the website was set to host the workflows and codes documentations.
- The **first demonstrator workflow** of MOFLearning (a Python workflow for gas absorption simulations with RASPA and Zeo++) is **already public**, with tutorials and benchmark accessible via our website.
- As soon as the workflows give rise to **scientific contributions** and **be published**, they will be made available via our **GRICAD Gitlab**, at the same address as all our other tools and plugins.
- Open source code are available **directly** after we manage to package them

Why use AiiDA ?



Covalent



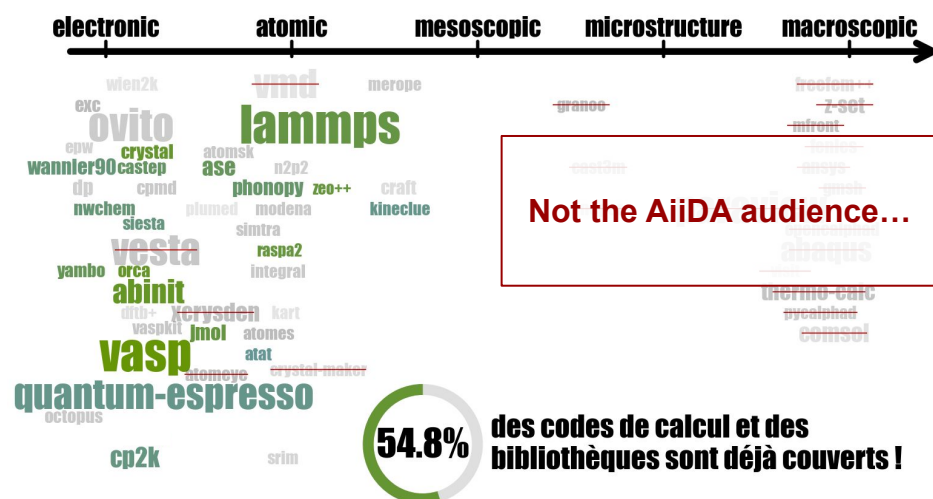
and many more ...

Choice motivated by :

- Proximity to AiiDA developers at SIMaP
- The list of existing plugins
- The list of existing schedulers

In practice, AiiDA relies in plugins

- **for codes**, to communicate with simulation software
- **for schedulers**, to deploy calculations in HPC facilities
- **for communication protocols**, to talk between different machines performing a task



Here is where we did most of the technical work! We did: *aiida-oar*, *fifo-transport*, *aiida-n2p2*, *aiida-numodis*... Also forks of *aiida-lammps* and *aiida-vasp* with customizations to better serve our workflows.

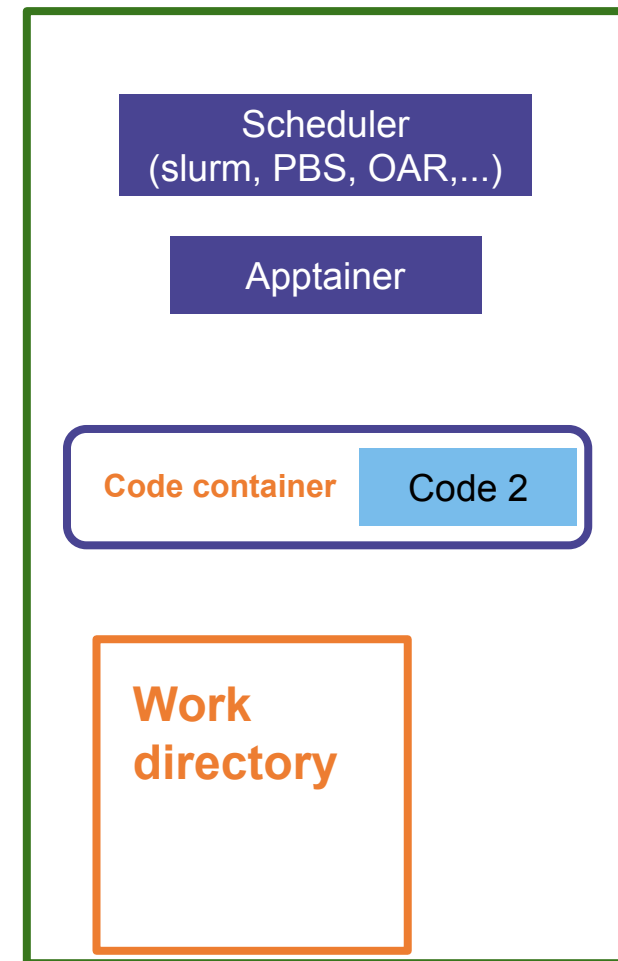
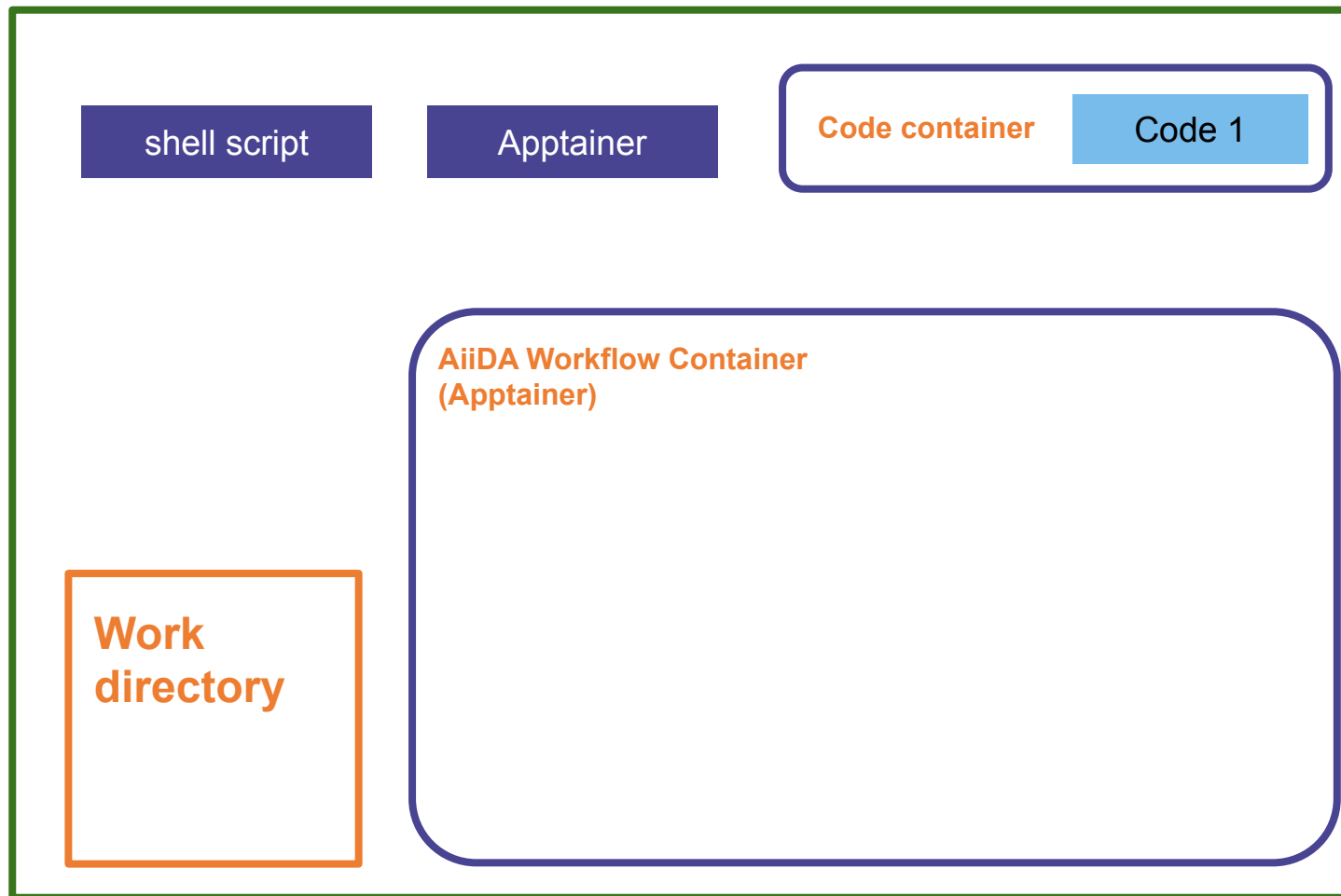


AiiDA workflow

Local machine (workflow server)

Remote machine (HPC Facility)

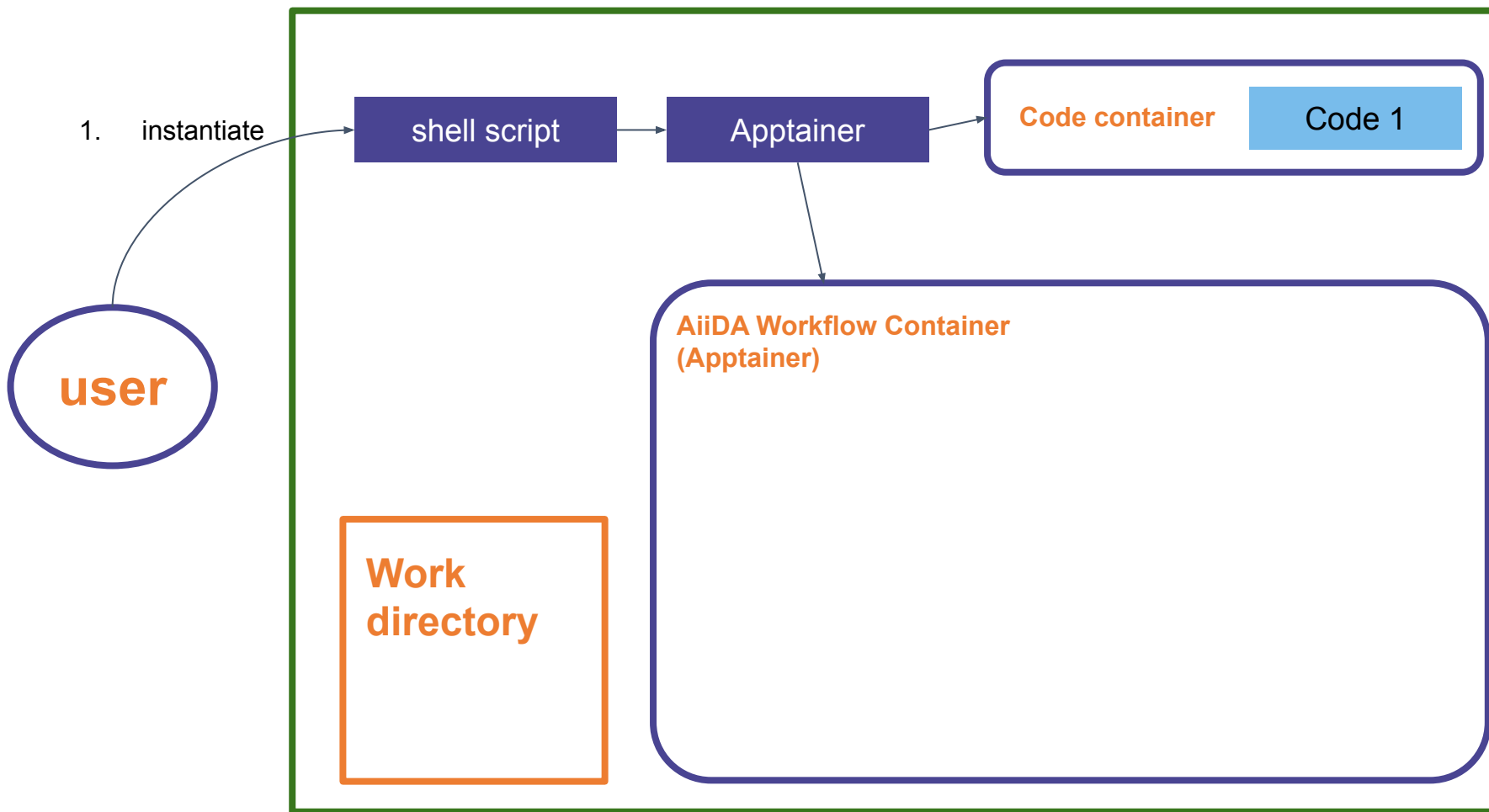
user



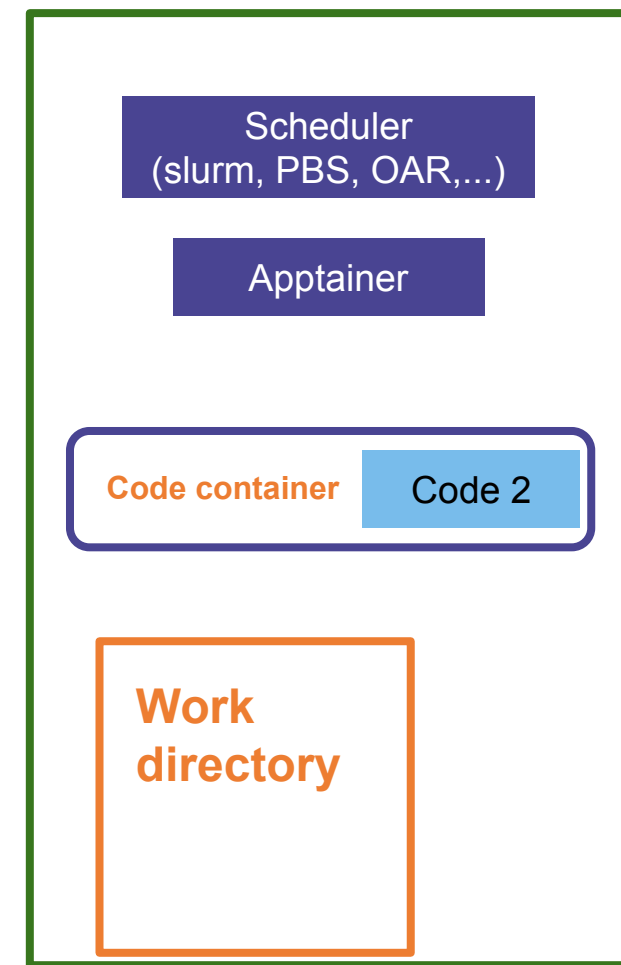


AiiDA workflow

Local machine (workflow server)



Remote machine (HPC Facility)

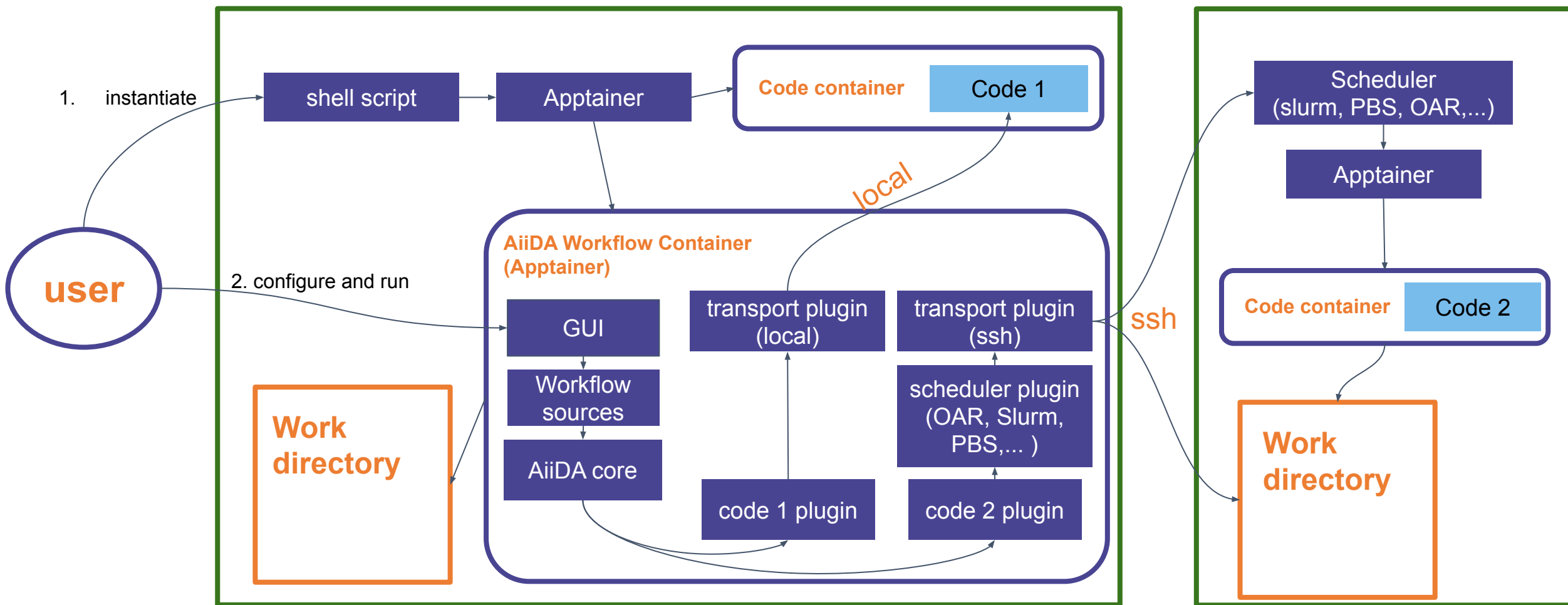




AiiDA workflow

Local machine (workflow server)

Remote machine (HPC Facility)

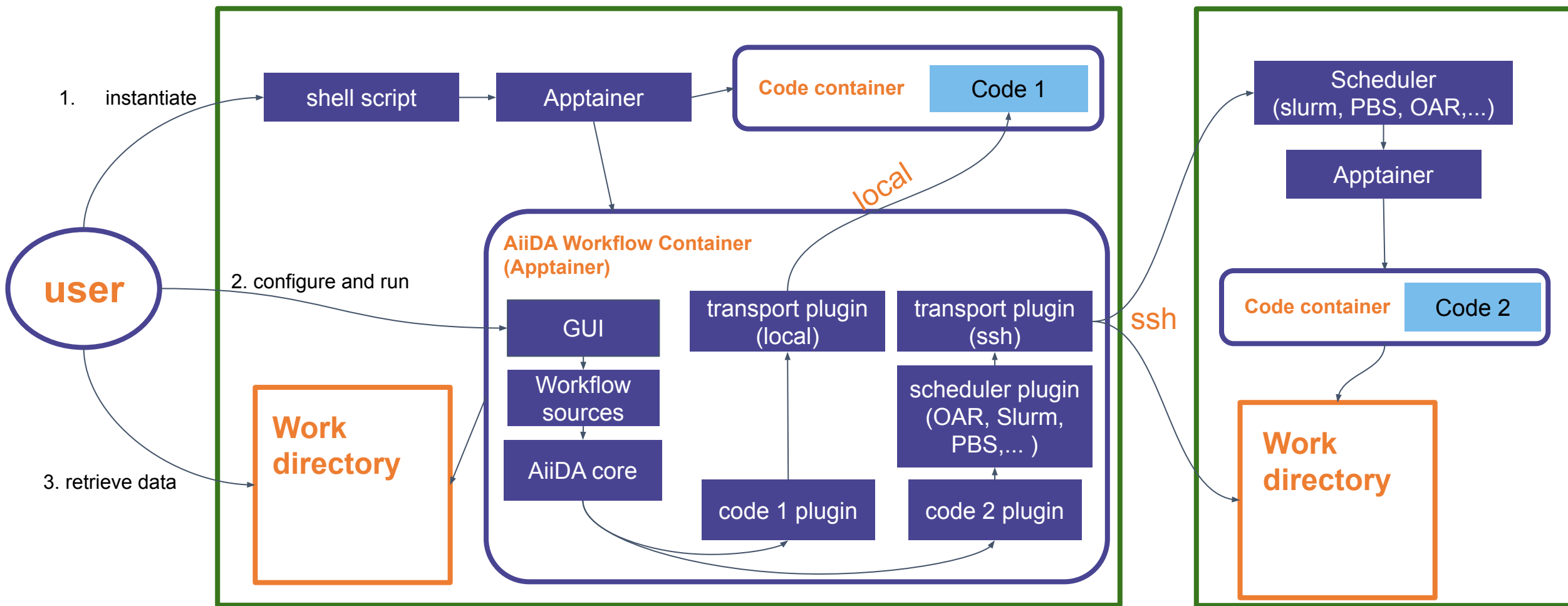




AiiDA workflow

Local machine (workflow server)

Remote machine (HPC Facility)

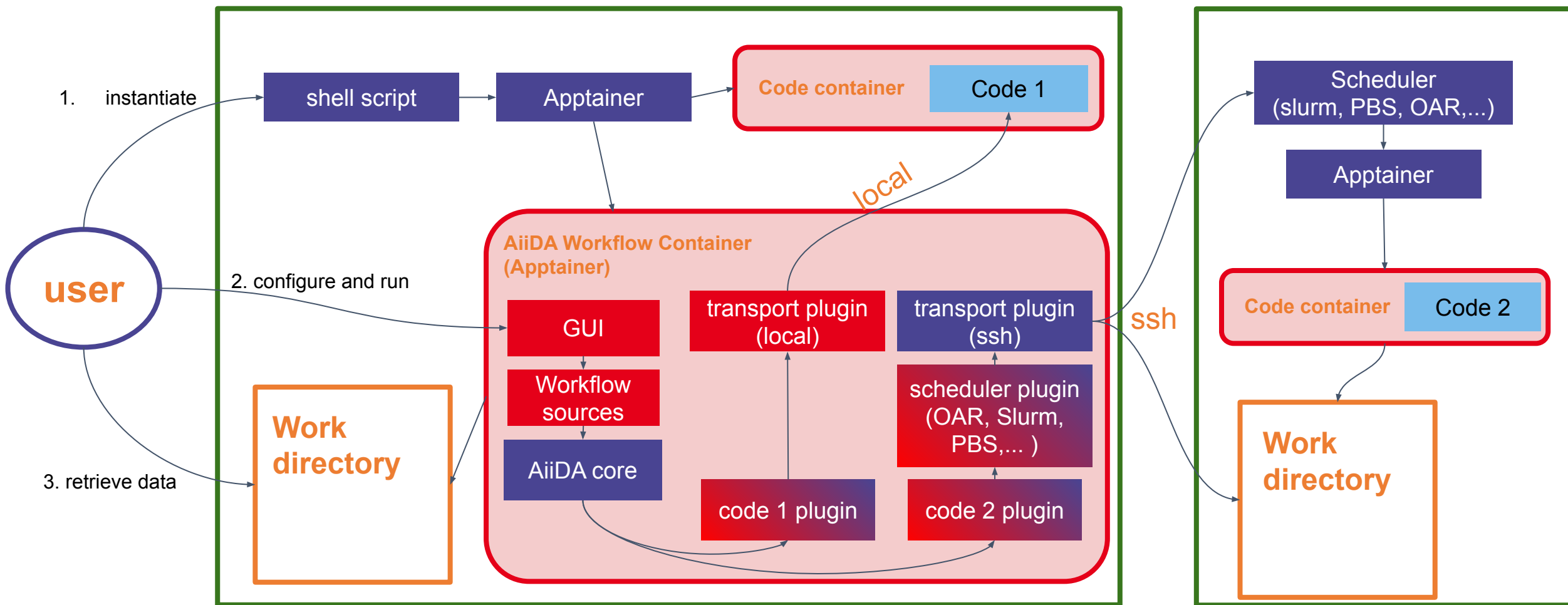




AiiDA workflow

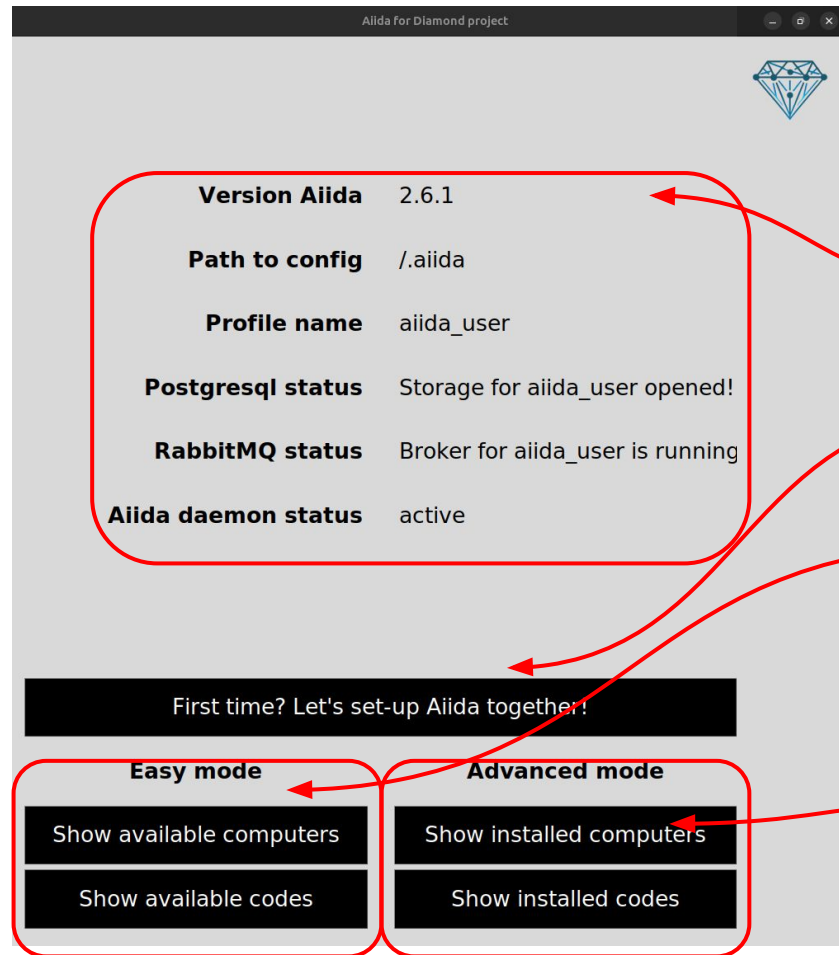
Local machine (workflow server)

Remote machine (HPC Facility)





Development of GUI for AiiDA Workflow Manager



The main objective is to make using AiiDA as easy as possible!

Multiple views for all users :

- **Information Panel**, displaying current AiiDA setup
- A **Beginner Mode**, building fast setup, ready in a click!
- An easy click-and-play menu to install codes or computers from **pre-made configuration files**:
[VASP, LAMMPS, QE,...] and [DAHU, JEAN-ZAY,...]
- An **Advanced Mode** to those who want to edit details of even delete codes or computers already installed



Our Workflows:

- **NUMODIS+LAMMPS workflow**

Atomistically informed dislocation dynamics simulations



Scientific specialists:

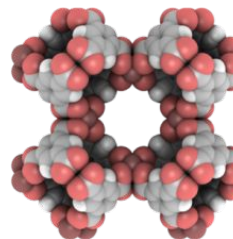
Laurent Dupuy

Status of development:

Stable, Containerized, Private

- **MOFLearning workflows**

Evaluate gas- absorption properties of MOFs, screening large databases



Arthur Hardiagon
François-Xavier Coudert

Stable, Containerized, Private, Production

- **Quantum Espresso workflow**

Screening of materials for electronic structure related properties



Ashna Jose
Mattia Ragni
Roberta Poloni
Martin Uhrin

WIP

- **VASP workflow**

Procedural AIMD trajectories aiming the training MLIPs for metals



Jonathan Chapignac
Noël Jakse

WIP, Containerized, Private, Production

- **MLIP workflow - n2p2 [WP3]**

Automate the training and validation of MLIPs with the n2p2 package



Arthur France-Lanord
Marco Saitta

WIP, Private





Workflows VASP

A Workflow for liquid alloys MD via VASP

- **Implementation and analysis:** J P A Mendonca, N Jakse
- **Technical support / containerization:** B Arrondeau, D Bissuel, P-A Bouttier
- Initially proposed by Noel and P-A as a **test case** for the platform, to evaluate the automation limits and guide our development efforts (martyr WF)
- Some more tangible interests in this workflow:
 - There is **MD data produced by Noels group** that still needs time expensive (human time) exploitation. E.g., doing structural evolution analysis in MDs with about 20k atoms and more the 1k non correlated frames.
 - The automated workflow can be extremely **useful for WP3**, creating datasets for future MLIPs.
 - There are at least two people in Noels group that can be very interested and active **beta testers**
- We are starting with AiiDA, but one of the goals is also to **reproduce it with other WF managers**, to test their limits and capabilities

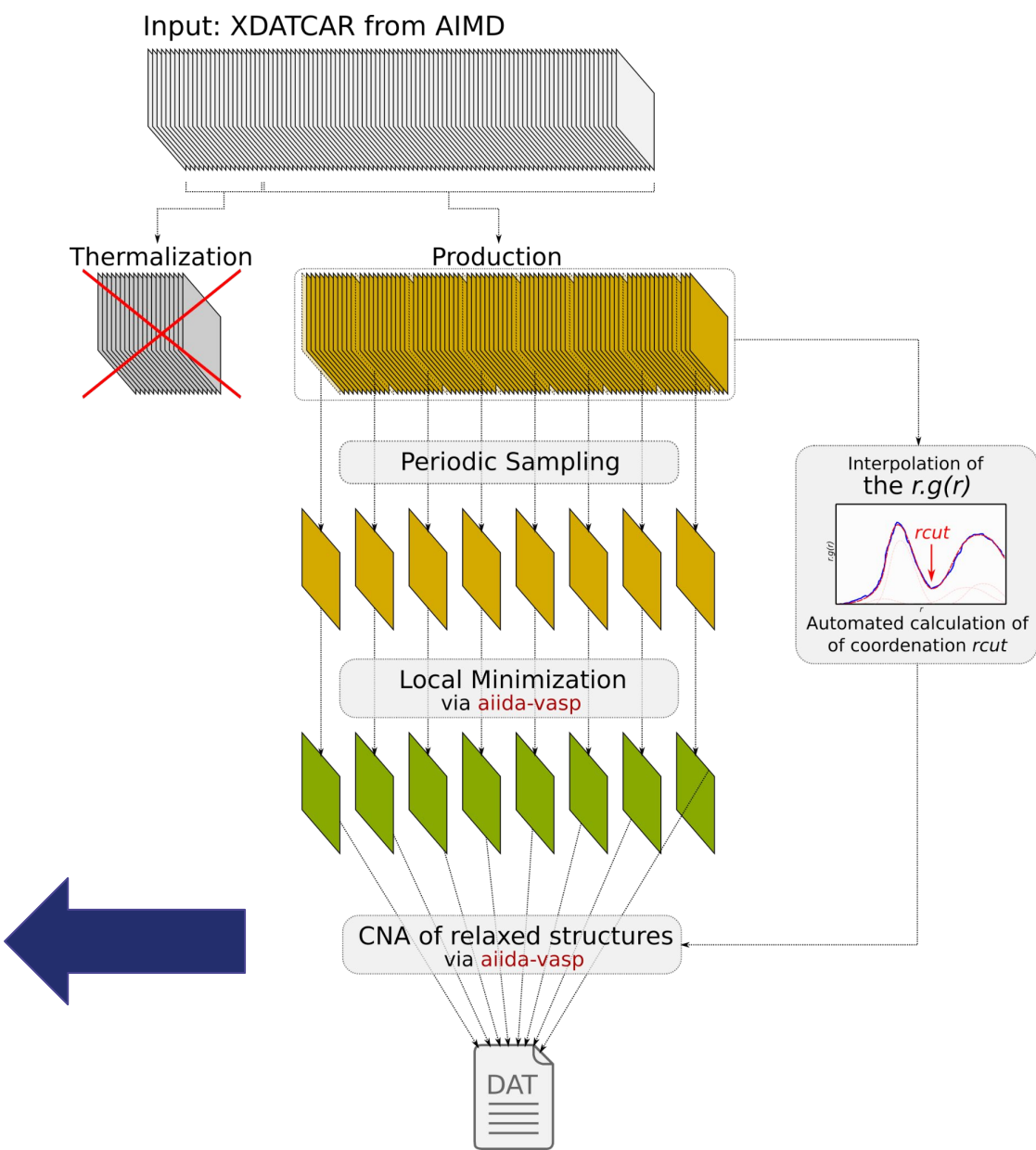
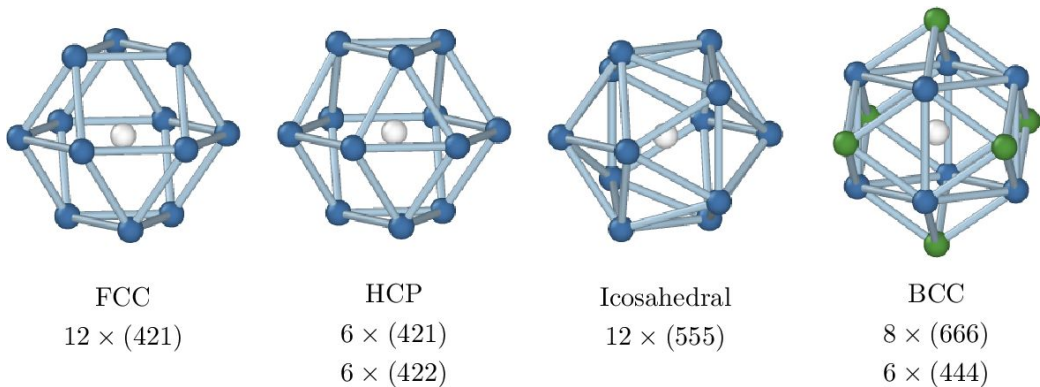


Workflows VASP - Module 1

Workflows organization:

WF1 - MD structural analysis

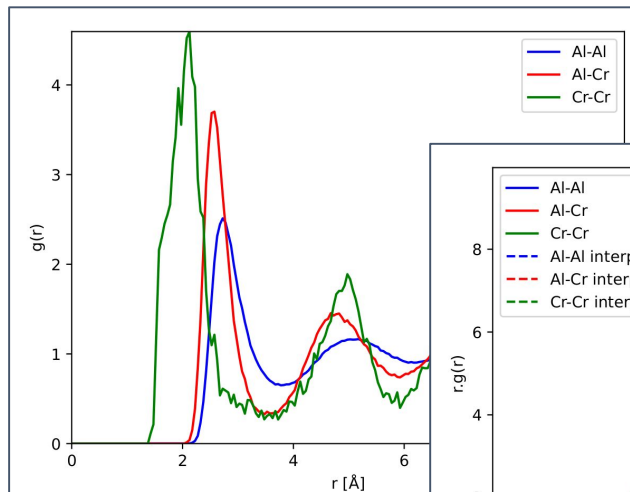
- Input: MD trajectory from VASP
- Process: Sampling > Structure Relaxation > Structural Analysis
- Output: Inherent Structures and CNA analysis (per atom / global)



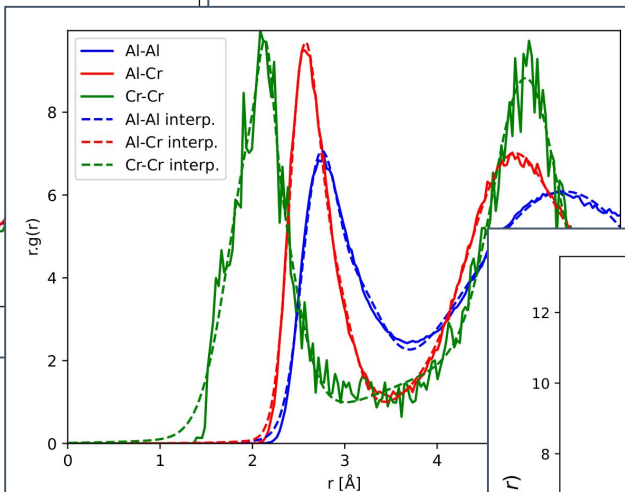


Workflows VASP - Module 1

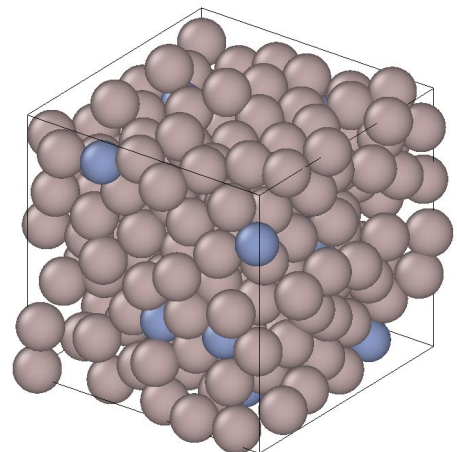
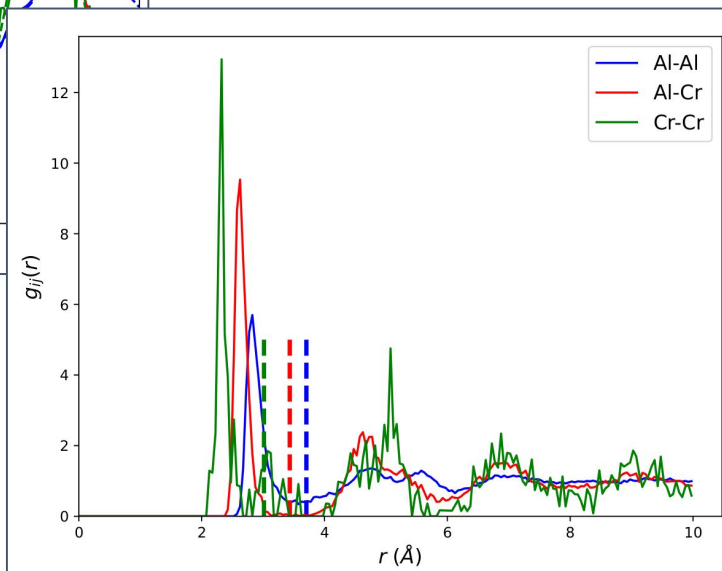
$g(r)$ from MD



$r.g(r)$ from MD



$g(r)$ inherent structures



Each structure to optimise
= 64 cores for 24h

CNA Analysis

```
-----FILE POSCAR_8000 -----
POSCAR_8000
1 1 13 [3 0 0]:1 [4 2 1]:1 [4 2 2]:3 [4 3 3]:3 [5 4 4]:3 [5 5 5]:2
2 1 10 [2 0 0]:3 [3 1 1]:2 [3 2 2]:1 [4 2 2]:2 [4 3 3]:1 [5 4 4]:1
3 1 12 [3 0 0]:2 [3 1 1]:1 [3 2 2]:1 [4 2 2]:3 [4 3 3]:3 [5 4 4]:1 [5 5 5]:1
...
254 2 10 [4 3 3]:3 [4 4 4]:1 [5 4 4]:1 [5 5 5]:5
255 2 12 [4 3 3]:2 [5 4 4]:2 [5 5 5]:8
256 2 10 [3 1 1]:2 [4 3 3]:6 [5 5 5]:2

Total number of pairs for atom type 1: 2632
Total number of pairs for atom type 2: 196
Total number of pairs for atom type 3: 0
Total number of pairs for atom type 4: 0

Number of atoms type 1: 238
Number of atoms type 2: 18
Number of atoms type 3: 0
Number of atoms type 4: 0

Coordination number type 1: 11.058823529411764
Coordination number type 2: 10.888888888888889
Coordination number type 3: 0.0
Coordination number type 4: 0.0
Coordination number total : 11.046875

CNA particle type 1
[1 0 0]:18 0.006839
[2 0 0]:206 0.078267
[2 1 1]:34 0.012918
[3 0 0]:228 0.086626
[3 1 1]:484 0.183891
[3 2 2]:144 0.054711
[4 1 1]:42 0.015957
[4 2 1]:132 0.050152
[4 2 2]:418 0.158815
[4 3 3]:485 0.184271
[4 4 4]:4 0.001520
[5 3 2]:4 0.001520
[5 3 3]:4 0.001520
[5 4 4]:238 0.090426
[5 5 5]:180 0.068389
[6 6 6]:11 0.004179

CNA particle type 2
[3 0 0]:2 0.010204
[3 1 1]:6 0.030612
[4 2 1]:6 0.030612
[4 2 2]:14 0.071429
[4 3 3]:57 0.290816
[4 4 4]:4 0.020408
[5 4 4]:22 0.112245
[5 5 5]:84 0.428571
[6 6 6]:1 0.005102

CNA particle type 3
CNA particle type 4
```

Chemical bound close
to crystallization:

6.8% Al

42.9% Cr



Workflows VASP - Module 2

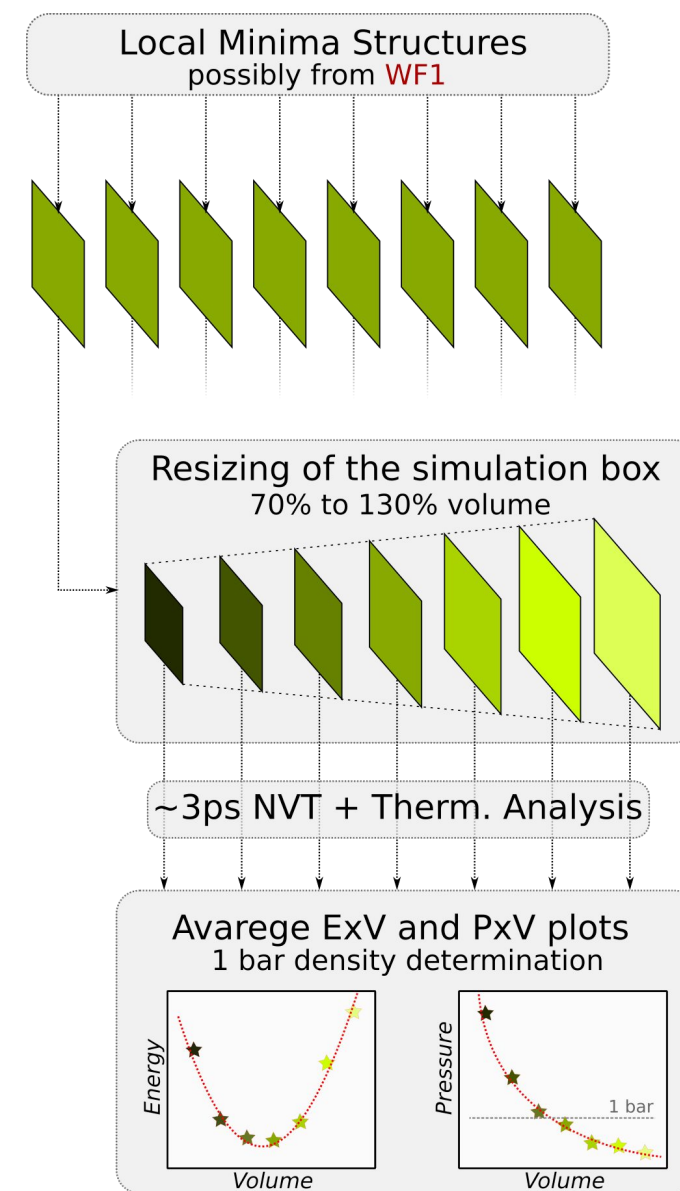
Workflows organization:

WF1 - MD structural analysis

- Input: MD trajectory from VASP
- Process: Sampling > Structure Relaxation > Structural Analysis
- Output: Inherent Structures and CNA analysis (per atom / global)

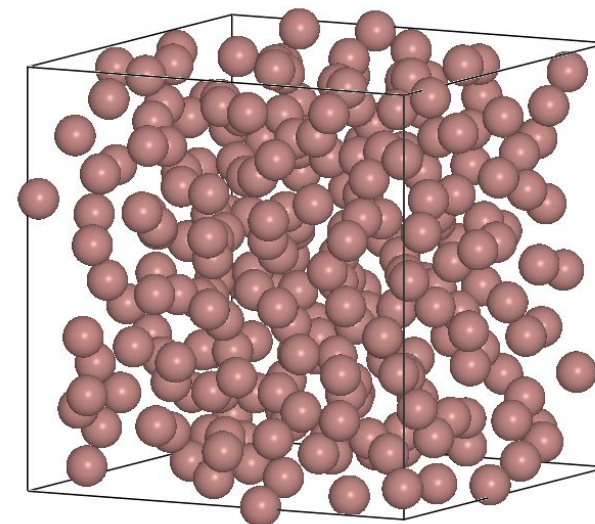
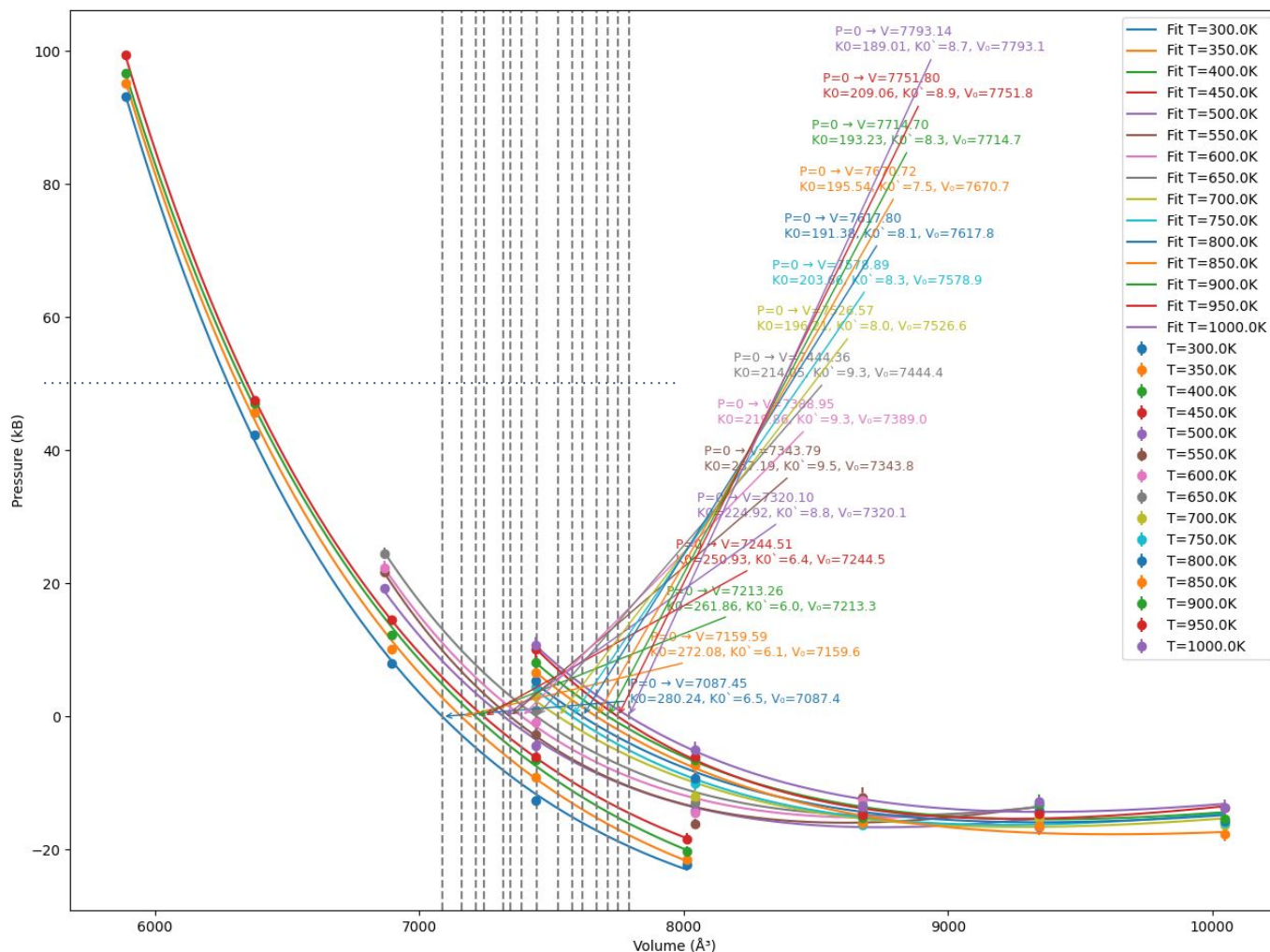
WF2 - Volume optimization

- Input: Collection of structures (like WF1 outputs) and temp. [list?]
- Process: Resize volume in range(70%,130%) > ~3ps NVTs > Thermodynamic
- Output: ExV and VxP plots, 1 bar predicted V





Workflows VASP - Module 2



In₂₅₀

Each temperature studied:

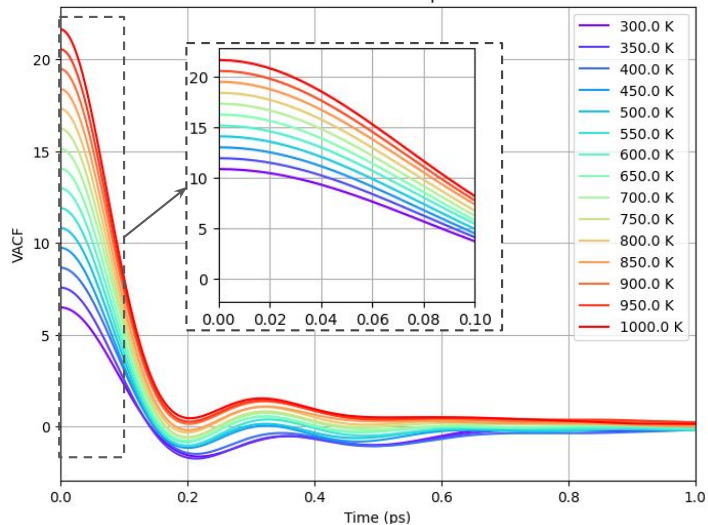
5 *ab initio* MDs
64 cores/48h each

Volumes of equilibrium allow the study to be done at ideal thermodynamics conditions with standard NVT Molecular Dynamics!

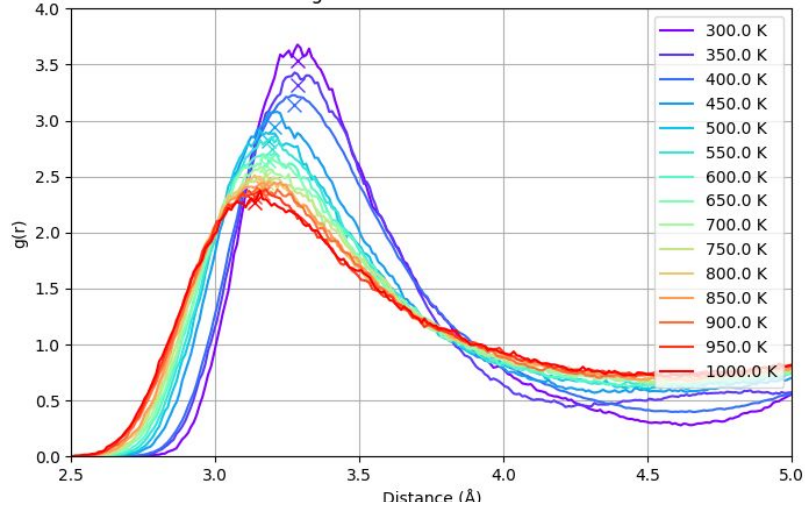


Workflows VASP - Module 2

VACF at Different Temperatures



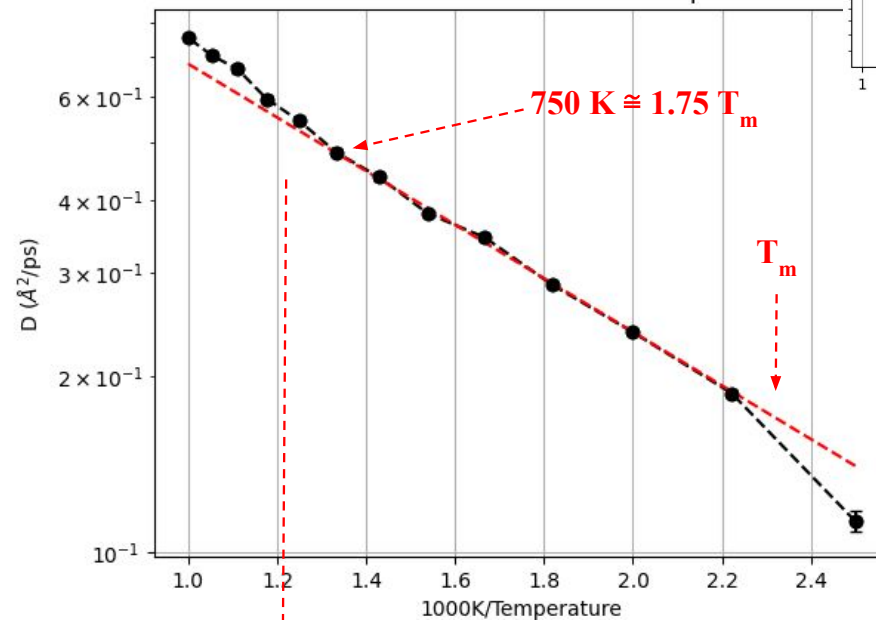
Average Radial Distribution Function



Indium melting point:

$$T_m = 429.8 \text{ K}$$

Diffusion Coefficient vs Inverse Temperature



Arrhenius fit:

$$D(x) = D_0 * \exp(s * x)$$

$$D_0 = 1.949077 \text{ \AA}^2/\text{ps}$$

$$s = -1.051576 \text{ (dimensionless per 'x' where } x=1000/T)$$

$$\Rightarrow \ln(D) = \ln(D_0) - 1.051576 * (1000/T)$$

Derived activation energy:

$$E_a = 0.090618 \text{ eV} = 8.743 \text{ kJ/mol}$$



Workflows VASP - Module 3

Workflows organization:

WF1 - MD structural analysis

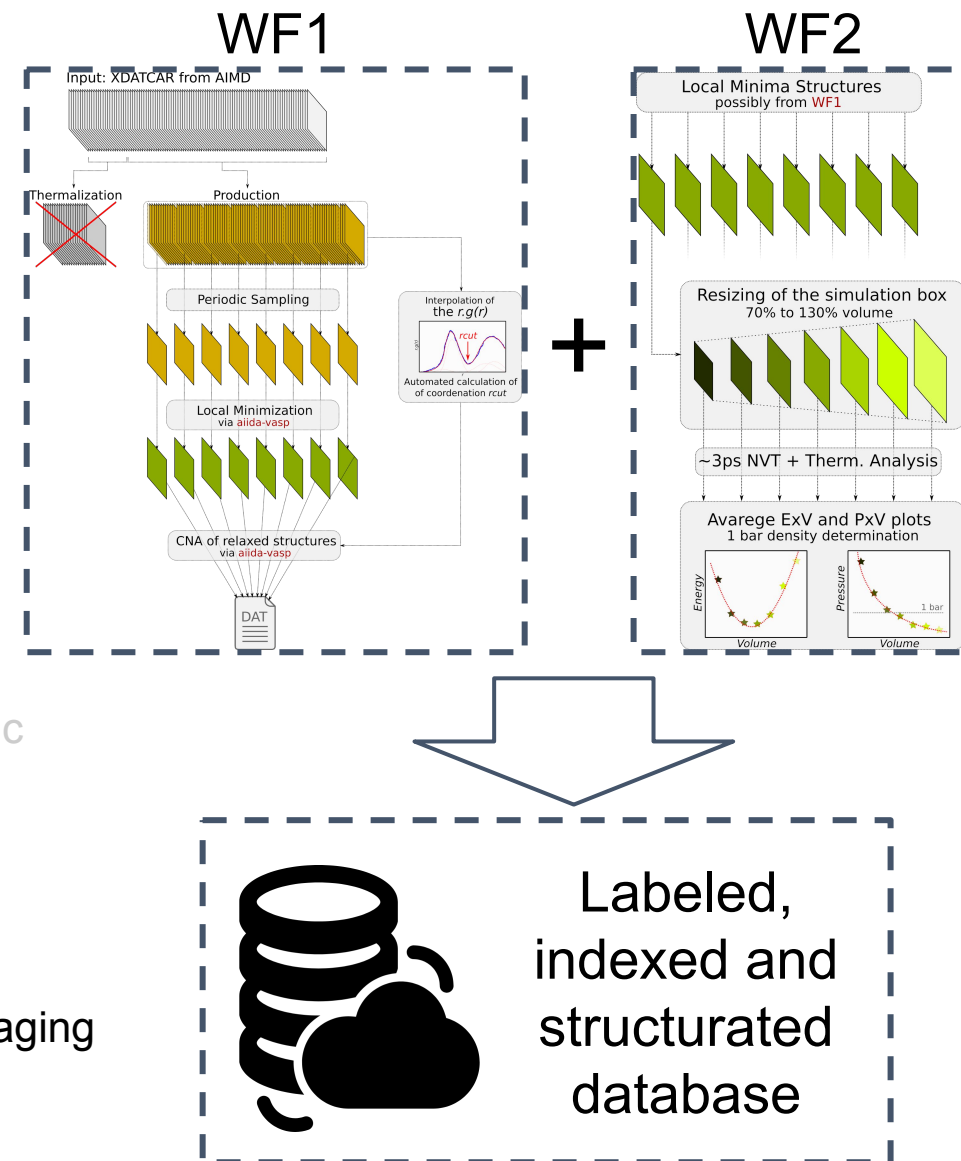
- Input: MD trajectory from VASP
- Process: Sampling > Structure Relaxation > Structural Analysis
- Output: Inherent Structures and CNA analysis (per atom / global)

WF2 - Volume optimization

- Input: Collection of structures (like WF1 outputs) and temp. [list?]
- Process: Resize volume in range(70%,130%) > ~3ps NVTs > Thermodynamic
- Output: ExV and VxP plots, 1 bar predicted V

WF3 - MLIP DataBase builder

- Input: Single structure, range of volumes and temperatures
- Process: Long NPT 1 bar > WF1 > WF2 > Single points / descriptors > Packaging
- Output: Database (structures, descriptors, energies, forces)

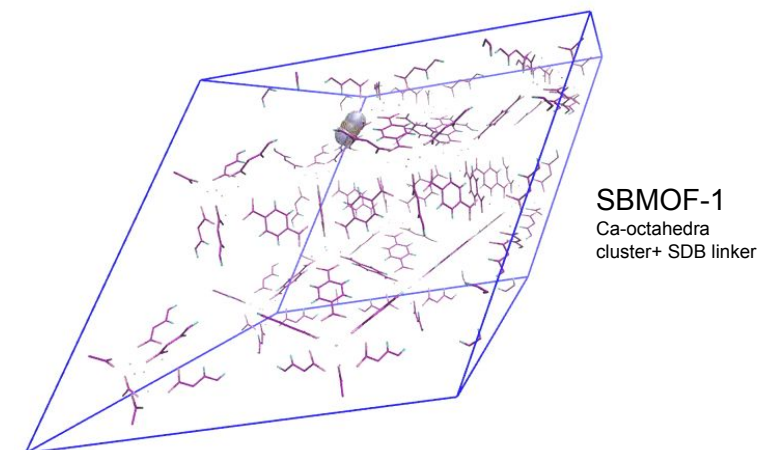
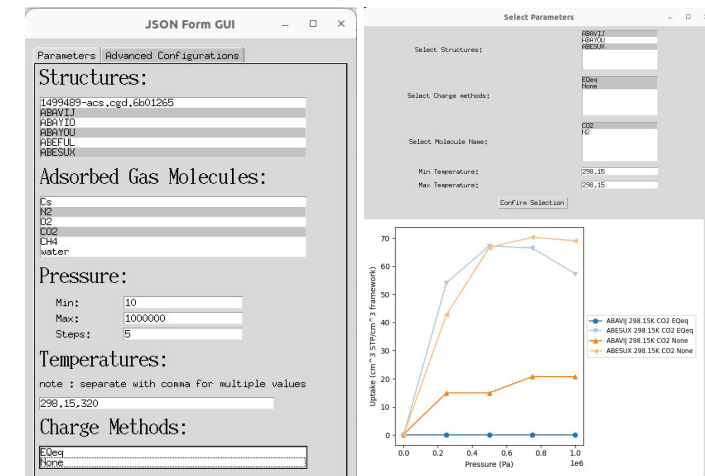
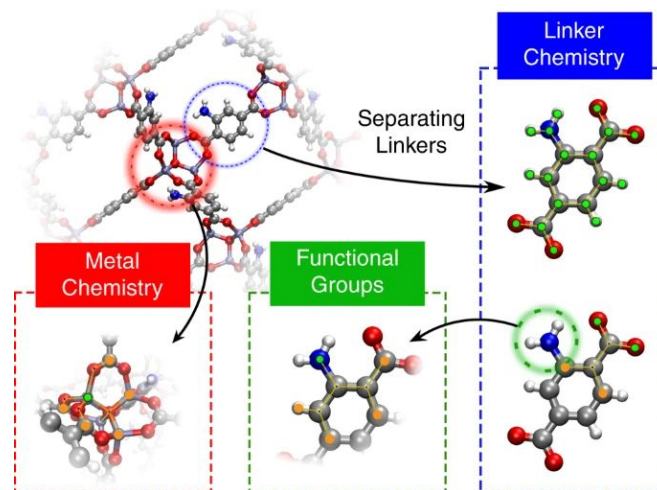
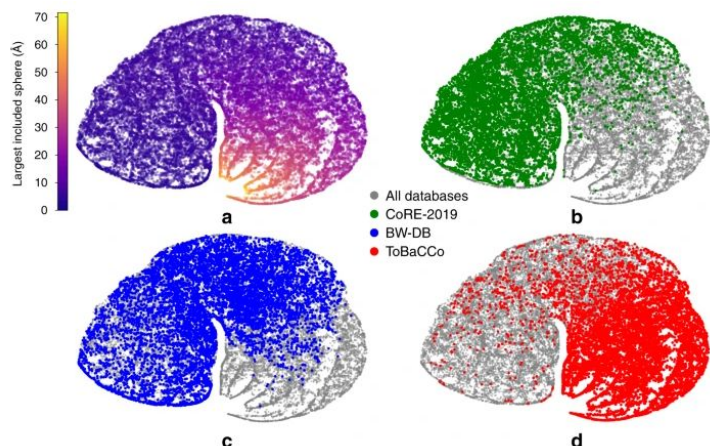




Use Case : MOF Learning

Workflow Python

- **Implementation and analysis:** A Hardiagon, FX Coudert
- **Technical support / containerization:** JPA de Mendonca, D Bissuel
- Compute **adsorption properties** (Monte-Carlo)
- Materials : Metal-organic Frameworks from **CoRE-MOF 2019 database**
- **single-CPU** calculations



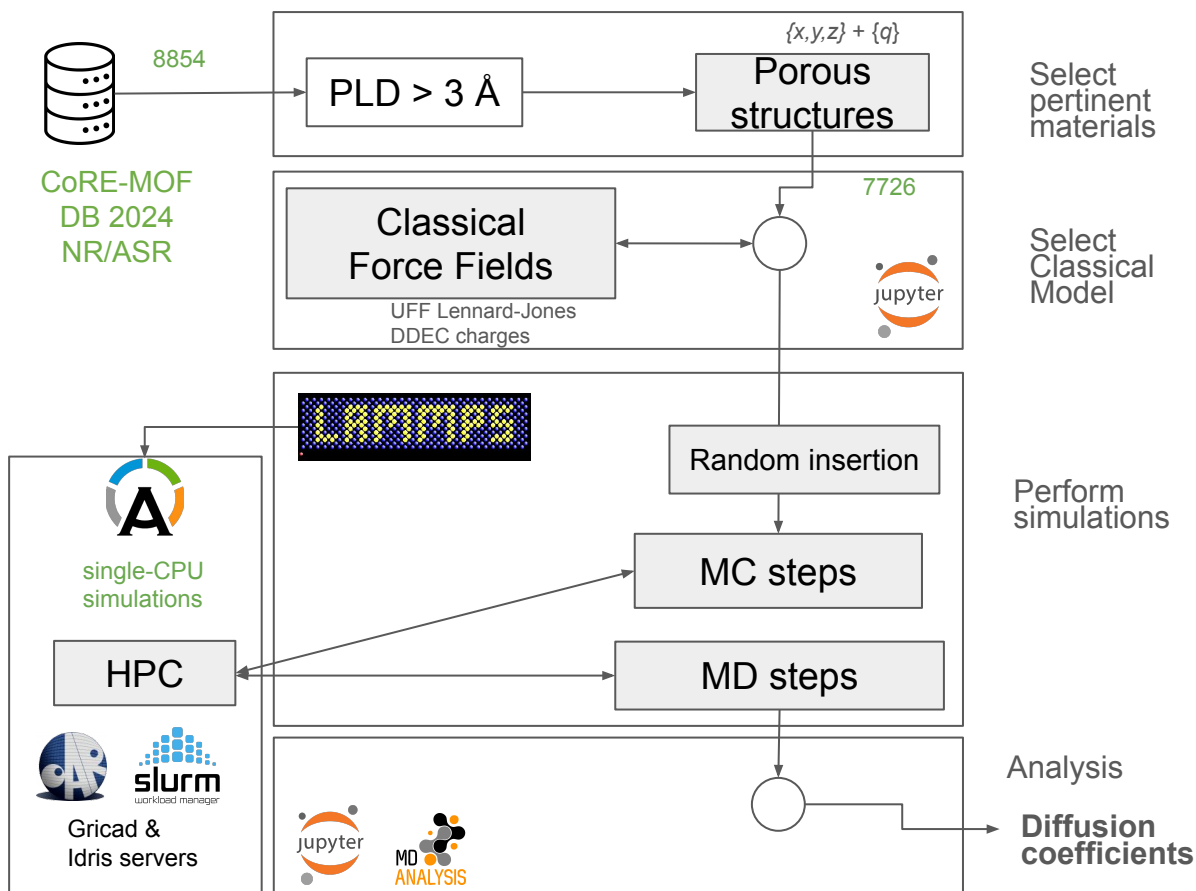
Functionalities

- JSON input/outputs
- GUI for input/output analysis
- containers
- Benchmarks (Dahu, Gricad)



Use Case : MOF Learning

Workflow Diffusion



Current State

Operational bash workflow

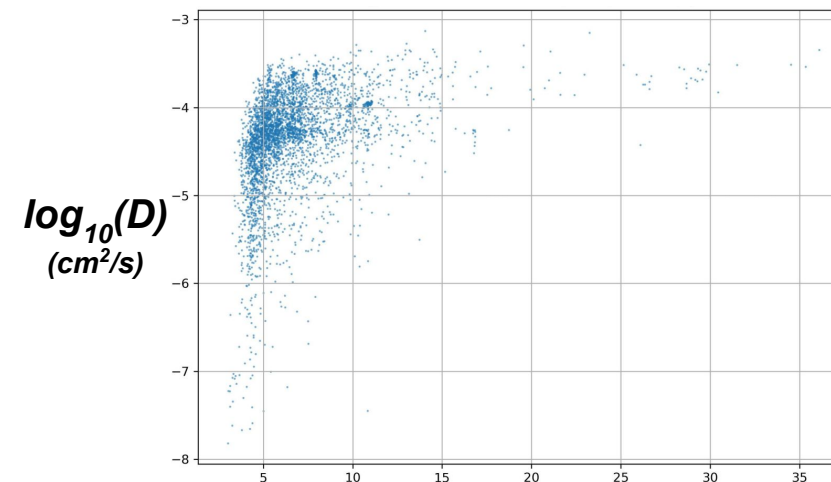
- use container with LAMMPS
- screening with **Krypton** gas

Aiida workflow tools

- simulations in 2 HPCs
- store database
- Template for Krypton
- Template for CO₂

Intern student:
William Xu

2 weeks for 7726 1-cpu
calculations (each ~48h)

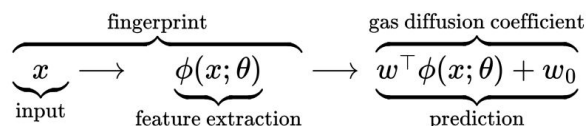




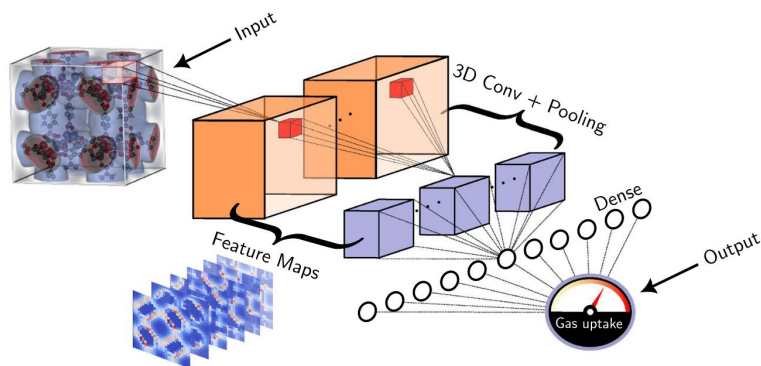
Use Case : MOF Learning

- Deep Learning prediction

- **Small model** : Convolutional Neural Networks
- **Fast predictions** 48h → 30 s
- **Informative descriptor = energy grids (PES)**



- **Accuracy** : level of theory for PES evaluation

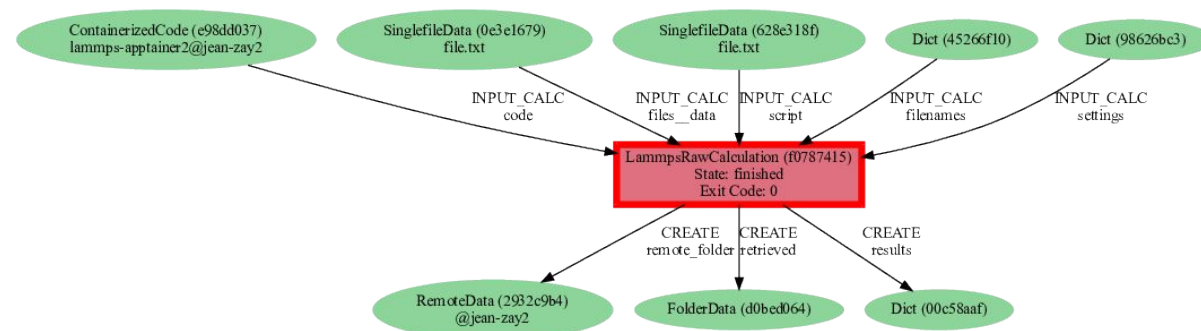


Sarikas, Scientific Reports, 2024

[moxel](#)

- Perspectives

- **CNN training** on 3D maps
 - hyperparameters optimization
 - grid resolution
 - comparison with other methods (SOAP + RF)
- **High-throughput simulations** with **CO₂** in MOFs
 - charges and electrostatics maps
 - Workflow manager : Aiiida



Thanks!
Any questions?



<https://diamond-diadem.github.io/en/>