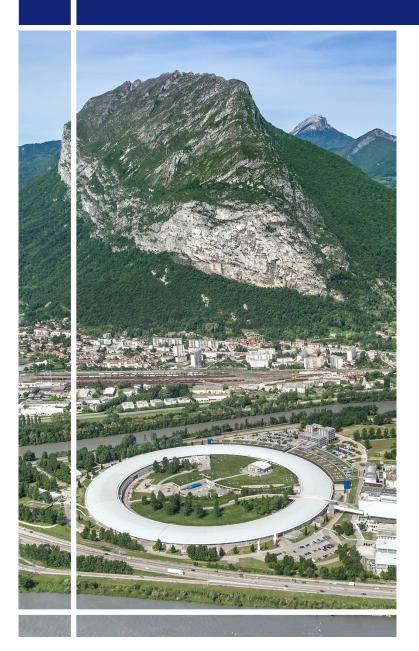


# **ESRF** | The European Synchrotron



# **Ewoks**

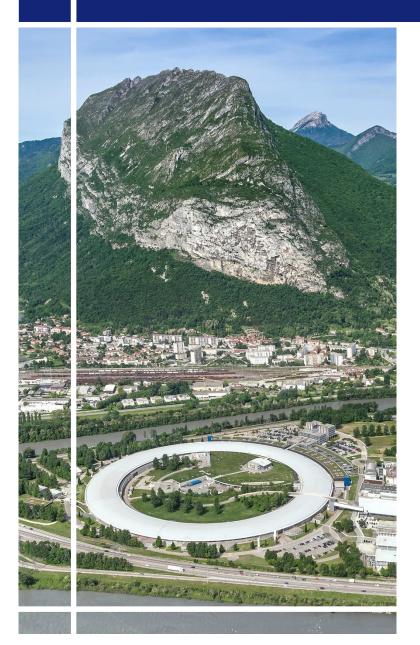
**ESRF Workflow System** 

A meta-workflow system

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## Outline

- 1. What problem does EWOKS solve?
- 2.What does EWOKS provides?
   (how is it "meta"?)
- 3.Getting started with EWOKS
- 4.Examples

#### What problem did we solve by developing EWOKS?

With the ESRF upgrade (**EBS**) and the introduction of the BeamLine Instrumentation Support Software (**BLISS**) more and more beamlines requested support for online data processing.

Solutions were developed independently in the past using various workflow management systems (WMS) like orange, passarelle, scipion, ... or custom tooling.

#### Why workflows?

A workflow is a recipe that fully describes the steps to be taken in a data processing pipeline or the decisions to be made by an automated data acquisition system.

- Reproducibility: results can be regenerated from the recipe
- Reuseability: process many datasets with the same recipe
- Traceability: workflows describe the history of a piece of data
- Human and machine: workflows can be create and executed by both
- Collaborative data processing: experts from different fields can create and execute the same workflow

#### **Design requirements**

We needed a way to share developer efforts across beamlines and techniques while still using the different features provided by existing workflow management systems.

- A workflow created in one system can also be executed in another
- A workflow **task** created in one system can also be executed in another
- **Changes** in technology should not affect users (creating workflows) and developers (implementing workflow tasks).
- Works **light-weight** (PC or laptop) and at **scale** (Beamline or Synchrotron).

EWOKS is **NOT** a workflow management system.

EWOKS is a **meta**-workflow system implemented in python.

It is "meta" in the sense that it **decouples** the workflow definition and task implementation from the WMS.

- EWOKS convention for workflow definition and task implementation
- Bindings for existing workflow management systems



# Many workflow management systems exist





























#### Current EWOKS bindings for existing workflow management systems

	WMS	Features
orange	Orange	Desktop GUI to create and execute workflows
DASK	Dask	Parallelize task execution (locally or on a cluster)
	Pypushflow	Cyclic workflows with conditional links
C Celery	Celery	Schedule workflows (locally or on a cluster)

#### What ewoks provides

- Common workflow definition (NetworkX + mapping for each WMS)
- Common task implementation (+ mapping for each WMS)
- Serialize workflows (NetworkX): json, yaml, ...
- Execution events for monitoring: python logging facility
- Persist/cache task results: parts of a workflow need to execute only once
- Web support: frontend (React) and backend (REST) to create, execute and monitor workflows

#### What ewoks does NOT provide

- Execution engine (except for a basic serial executor for DAGs)
- Data transfer between tasks (although the binding could use the persistence mechanism mentioned above)
- Worker caching
- Desktop GUI



Install EWOKS pip install ewoks

Create a test workflow ("acyclic1" is an test workflow that ships with ewoks) ewoks convert acyclic1 --test test.json -s indent=2

Execute the test workflow and print output ewoks execute test.json --output all

For more information https://ewoks.readthedocs.io

Install EWOKS with the orange binding (desktop GUI) pip install ewoks[orange]

Open workflow in the orange GUI ewoks execute test.json --binding orange

Install EWOKS for web pip install ewoksserver[frontend]

Start the EWOKS web server ewoks-server

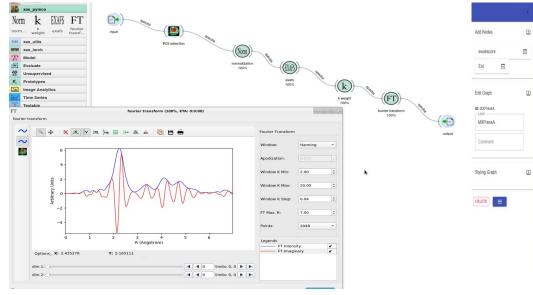
For more information https://ewoks.readthedocs.io

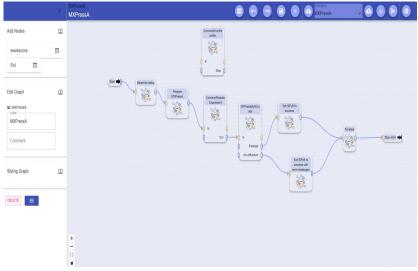


## GUI to create, execute and monitor workflows

Desktop







#### Implement a workflow task

```
from ewokscore import Task
class SumTask(
   Task,
    input_names=["a"],
    optional_input_names=["b"],
    output_names=["result"]
):
    def run(self):
        result = self.inputs.a
        if self.inputs.b:
            result += self.inputs.b
        self.outputs.result = result
```

#### Execute a workflow

```
from ewoks import execute_graph
inputs = [{"id": "task1",
           "name": "a",
           "value": 10}]
# Optionally persist all task outputs
varinfo = {"root_uri": "/tmp/myresults"}
result = execute_graph(workflow,
                       varinfo=varinfo,
                       inputs=inputs)
print(result)
```

#### Define a workflow

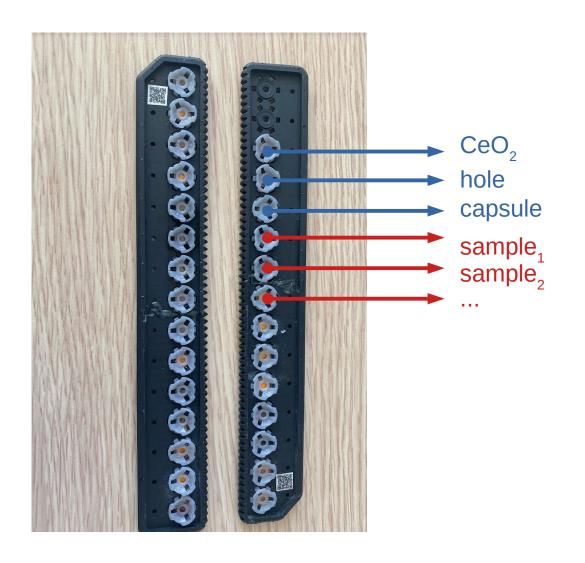
```
nodes = [
        "id": "task1",
        "task_type": "class",
        "task_identifier": "mypackage.SumTask",
        "default_inputs": [{"name": "a",
                             "value": 1}],
    },
{
        "id": "task2",
        "task_type": "class",
        "task_identifier": "mypackage.SumTask",
        "default_inputs": [{"name": "b",
                             "value": 1}],
    },
links = [
        "source": "task1",
        "target": "task2",
        "data_mapping": [{"source_output": "result",
                          "target input": "a"}],
    },
workflow = {"graph": {"id": "testworkflow"},
            "nodes": nodes,
            "links": links}
```

For more information https://ewoks.readthedocs.io



#### 5. Examples: data reduction for X-ray powder diffraction

#### https://twitter.com/ESRF\_Streamline



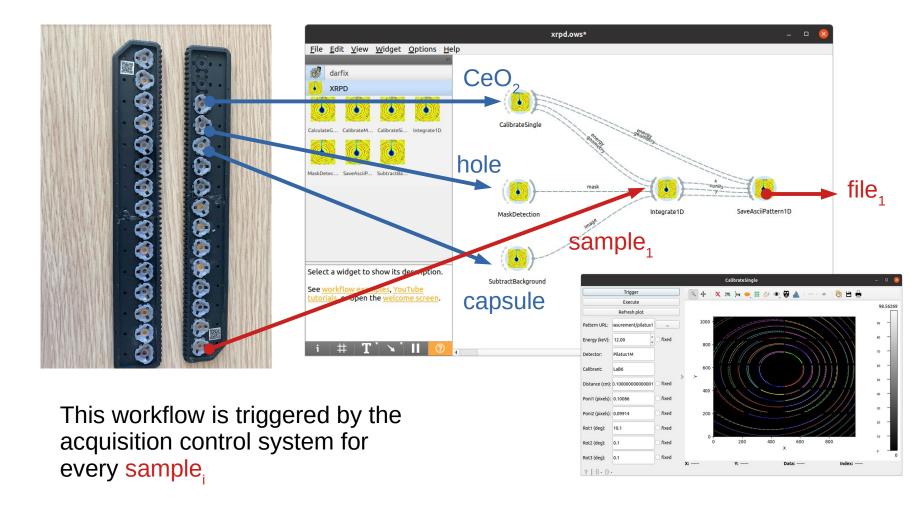
**EWOKS** workflow:

For calibration

For XRPD Data reduction for each sample

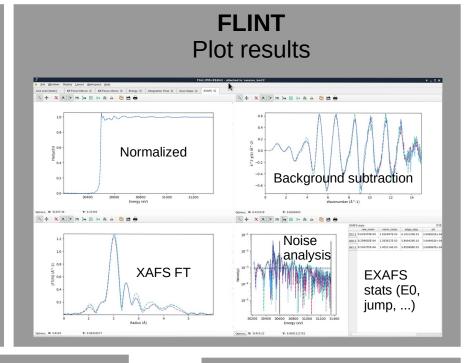
#### 5. Examples: data reduction for X-ray powder diffraction

#### https://twitter.com/ESRF\_Streamline



# 5. Examples: online EXAFS plotting

# 





Save workflow in a file







Every x seconds during scan: send workflow to worker and send result to Flint

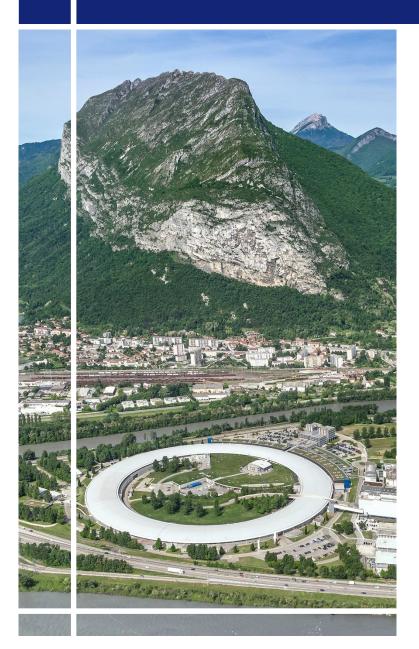


EWOKS (worker)
Execute workflow

redis

Celery





# **Ewoks**

May the force be with you

Documentation https://ewoks.readthedocs.io

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