



LifeWatch ERIC e-Science for NIS research workshop | 20-21 May 2021

An event organised within the framework of the LifeWatch ERIC Internal Joint Initiative and with the support of the ENVRI-FAIR project



Session 1: Long-term monitoring of hard-bottom marine communities ARMS Workflow - development, current status, functionalities



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1. Forewords about ICT presentations

Two sections in each ICT presentation

Associated Workflow

Explain technical aspects related to the current deployment

Tesseract Architecture

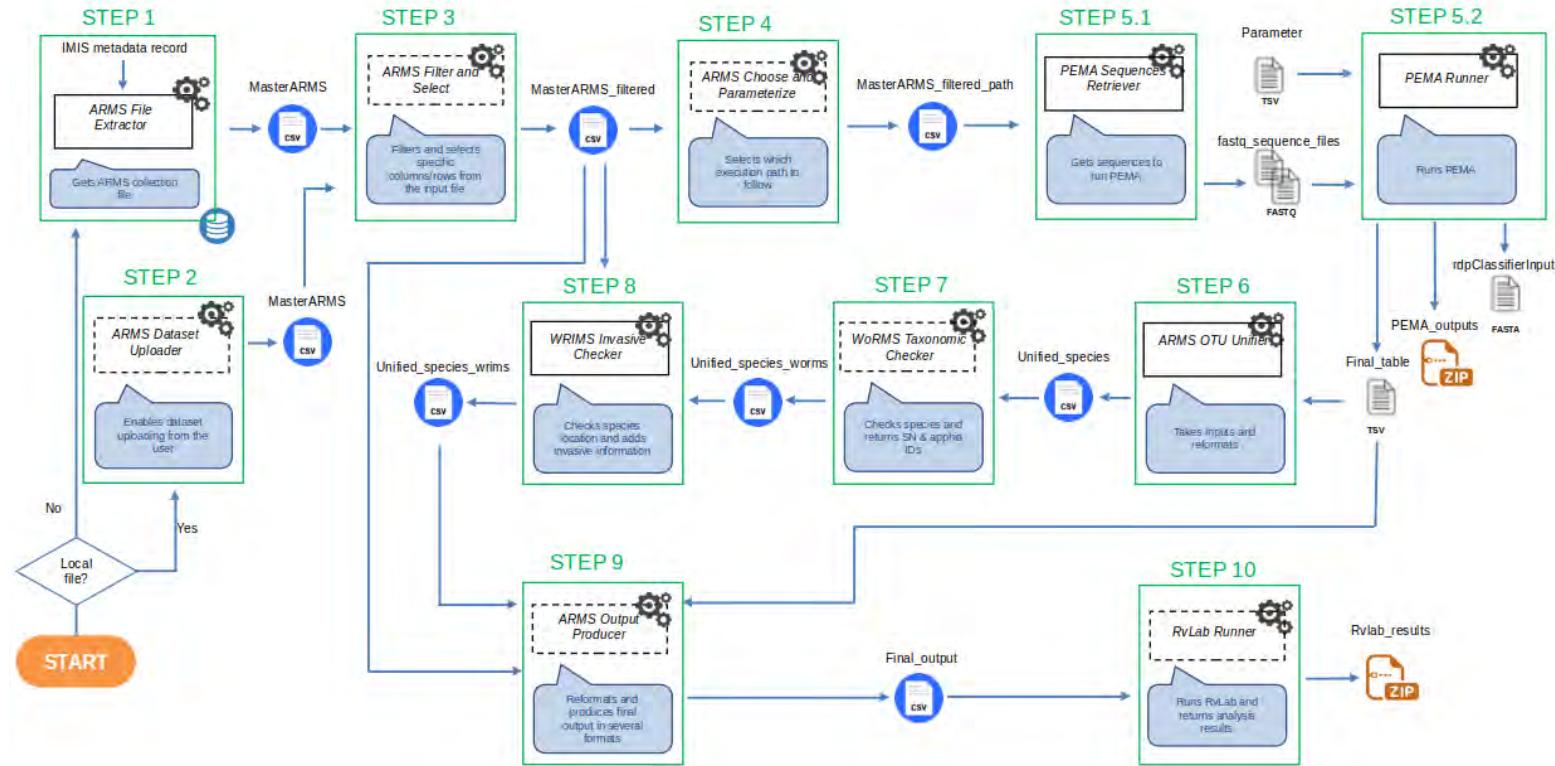
Explain technical aspects of the architecture **tested** in the development of the workflows

2. Current deployment of ARMS workflow

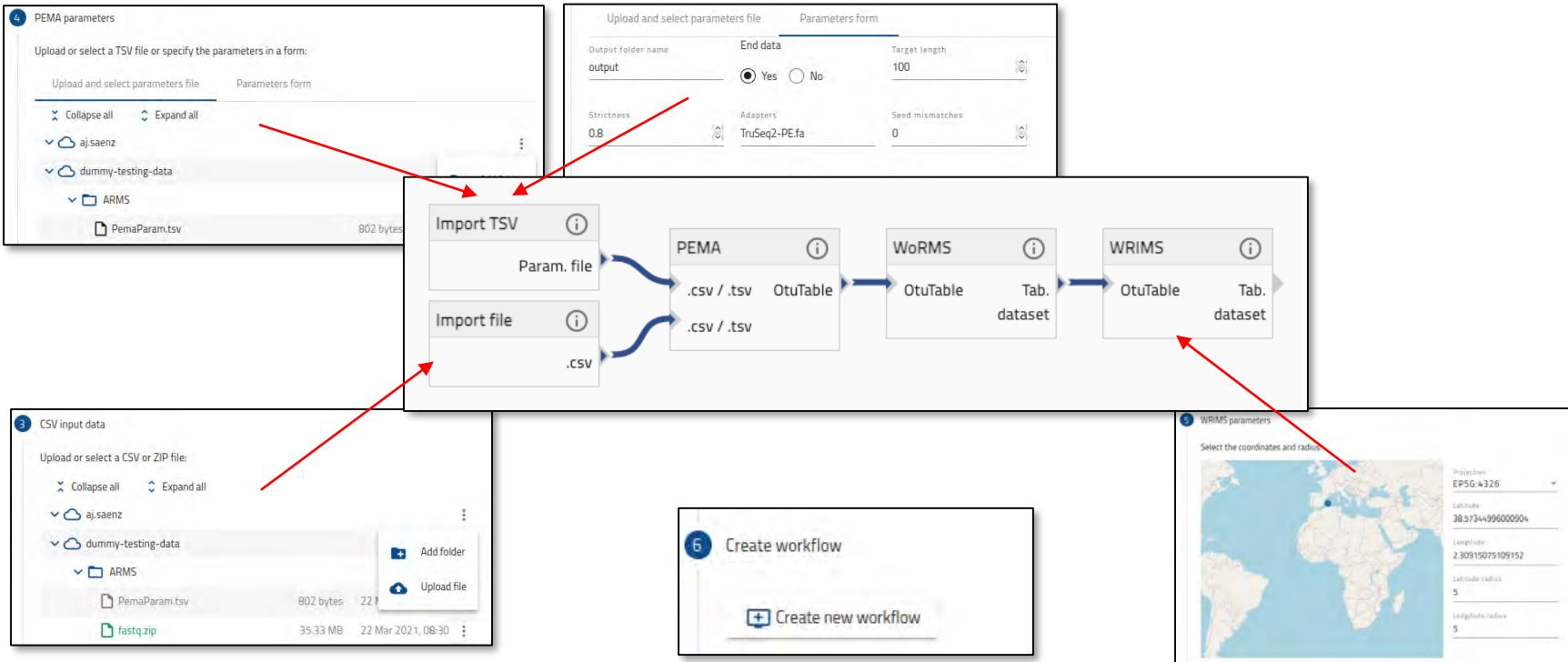
Associated Workflow

Explain technical aspects
related to the current
deployment

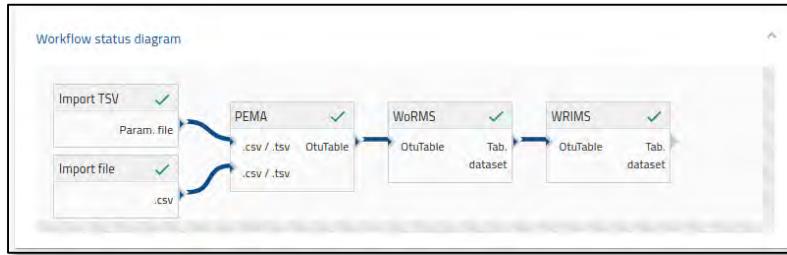
ARMS Validation Case



ARMS UI Current Implementation



ARMS UI Current Implementation



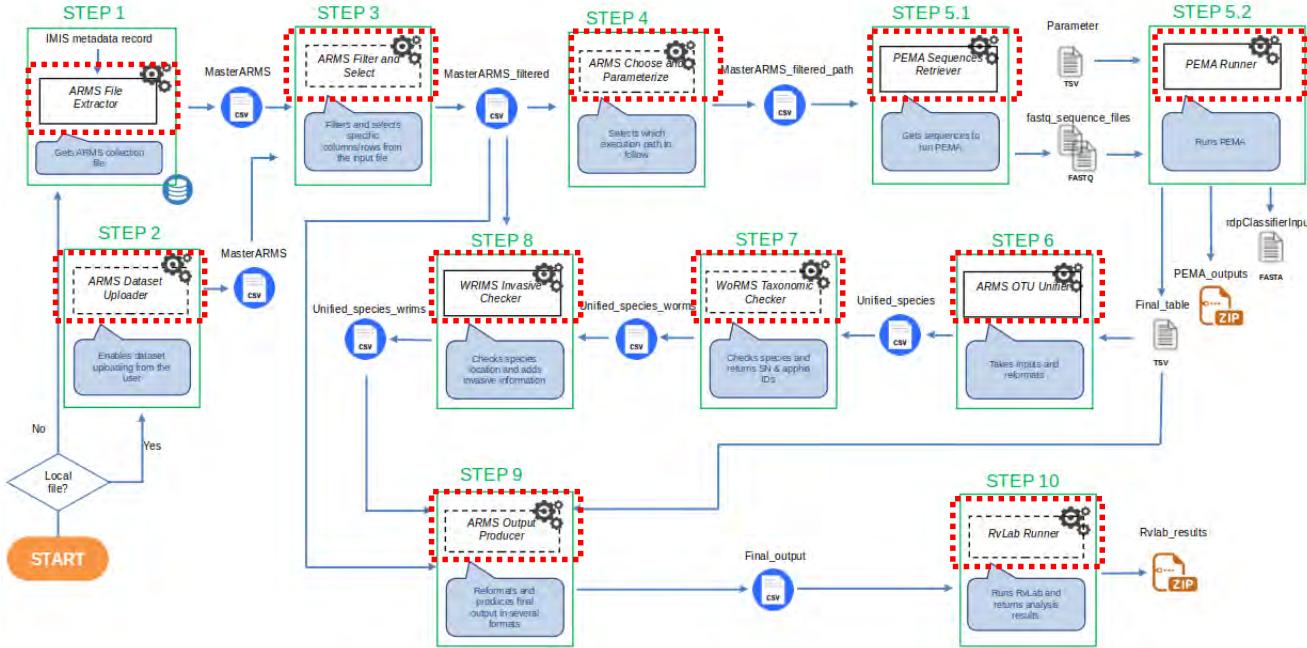
```

{
  "version": 0,
  "name": "ComponentImportFile1",
  "createdAt": "2021-04-23T16:44:00Z",
  "updatedAt": null,
  "params": null,
  "inputs": null,
  "result": {
    "message": "Success",
    "files": [
      {
        "path": "ComponentImportFile1/fastq.zip",
        "size": 35.33,
        "type": "zip"
      },
      {
        "path": "ComponentImportFile1/log.txt",
        "size": 363,
        "type": "text"
      }
    ],
    "log": [
      {
        "path": "ComponentImportFile1/log.txt",
        "size": 372,
        "type": "text"
      }
    ]
  },
  "status": "DONE"
}
  
```

Workflow output files

Workflow output files			
		File	Date
<input type="checkbox"/> Collapse all		<input type="checkbox"/> Expand all	
<input type="checkbox"/> 971683ea8b7c475999df4a88989bc7b4			
<input type="checkbox"/> ComponentImportFile1			
<input type="checkbox"/>	fastq.zip	35.33 MB	23 Apr 2021, 16:44
<input type="checkbox"/>	log.txt	363 bytes	23 Apr 2021, 16:44
<input type="checkbox"/> ComponentImportTSV1			
<input type="checkbox"/>	log.txt	372 bytes	23 Apr 2021, 16:44
<input type="checkbox"/>	out.tsv	800 bytes	23 Apr 2021, 16:44
<input type="checkbox"/> ComponentPEMA1			
<input type="checkbox"/>	final_table.tsv	9.11 KB	23 Apr 2021, 16:46
<input type="checkbox"/>	log.txt	7.68 MB	23 Apr 2021, 16:46
<input type="checkbox"/> ComponentWRIMS1			
<input type="checkbox"/>	classification.csv	43 bytes	23 Apr 2021, 16:47
<input type="checkbox"/>	log.txt	1.04 KB	23 Apr 2021, 16:47
<input type="checkbox"/> ComponentWoRMS1			
<input type="checkbox"/>	log.txt	9.8 KB	23 Apr 2021, 16:47
<input type="checkbox"/>	otu_table.csv	1.06 KB	23 Apr 2021, 16:47

ARMS Wrappers Implementation and Generalization



A “wrapper” provides two aspects:

- An interface to the workflow engine
- An execution environment and implementation

One of our goals is to generalize them so that we can reuse

ARMS File Extractor

Label: ARMS File Extractor

Description: Retrieves MasterARMS.csv

Input: ARMS URL

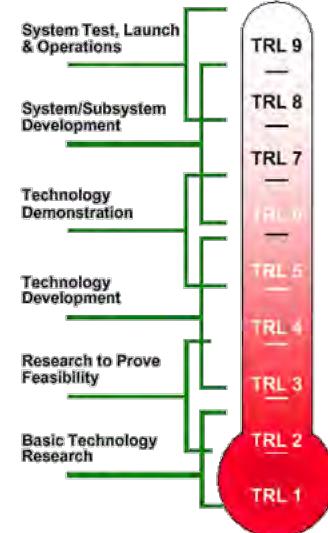
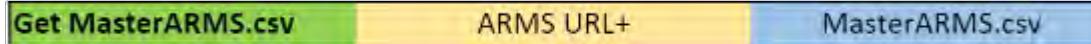
Output: MasterARMS (csv)

Metadata Harmonized: YES

TRL: 5

GUI Integration: YES

Technology: R / Python



Technology
Readiness
Level

PEMA Sequences Retriever

Label: PEMA Sequences Retriever

Description: Retrieves Sequences fasta files for
PEMA

Input: MasterARMS (csv)

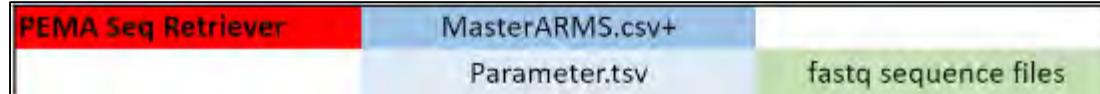
Output: fastq_sequence_files (fastq)

Metadata Harmonized: YES

TRL: 4

GUI Integration: YES

Technology: R / Python



PEMA Runner

Label: PEMA Runner

Description: Executes PEMA code and produces results

Input: fastq_sequence_files (fastq)

Output: Final_table (tsv),
rdpClassifierInput (fasta),
PEMA_outputs (zip)

Metadata Harmonized: YES

TRL: 5

GUI Integration: YES

Technology: Docker / R / Python

PEMA Runner	fastq sequence files	
	Parameter.tsv	final_table.tsv
	rdpClassifierInput.fasta	PEMA_outputs.zip

ARMS OTU Unifier

Label: ARMS File Extractor

Description: Takes input CSV, TSV and XLS and
unifies them

Input: Final_table (tsv)

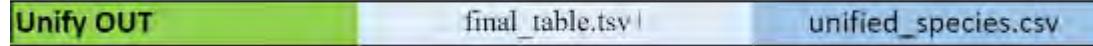
Output: Unified_species (csv)

Metadata Harmonized: YES

TRL: 2

GUI Integration: NO

Technology: R / Python



WoRMS Taxon Check

Label: WoRMS Taxon Check

Description: Species names checked by WoRMS

Input: Unified_species (csv)

Output: Unified_species_worms (csv)

Metadata Harmonized: YES

TRL: 3

GUI Integration: YES

Technology: R / Python

WoRMS taxon check

unified_species.csv+

unified_species.csv

WRIMS Invasive Check

Label: WRIMS Invasive Check

Description: Checks Species location and returns invasive

Input: Unified_species_worms (csv), MasterARMS (csv)

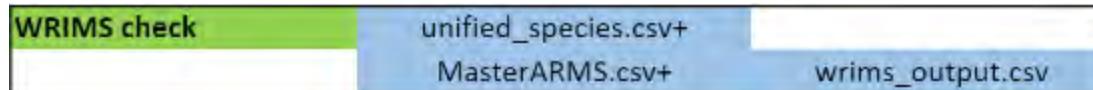
Output: MasterARMS (csv)

Metadata Harmonized: YES

TRL: 2

GUI Integration: NO

Technology: R / Python



Produce output

Label: Produce output

Description: Produces final output in various formats

Input: MasterARMS_filtered (csv),
Unified_species_wrims (csv),
Final_table (tsv)

Output: Final_output (csv)

Metadata Harmonized: YES

TRL: 3

GUI Integration: YES

Technology: R / Python

Produce output	MasterARMS.csv	
	unified_species.csv	
	final_table.tsv	final_output

Run RvLab

Label: Run RvLab

Description: Runs RvLab and returns analysis results

Input: Final_output (csv)

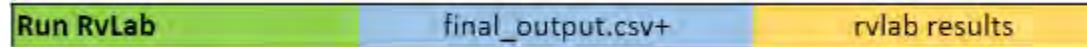
Output: Rvlab_results.zip

Metadata Harmonized: YES

TRL: 3

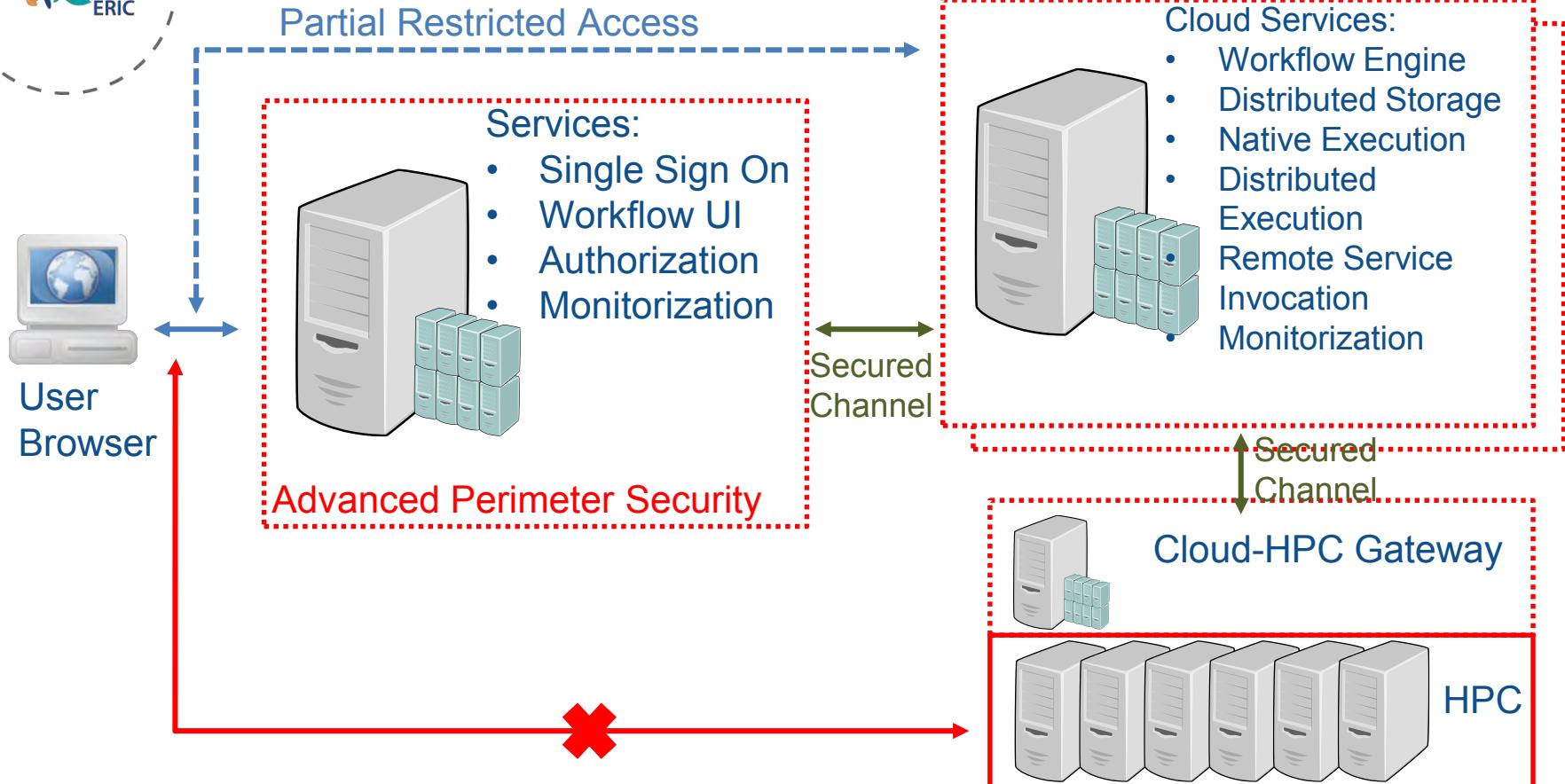
GUI Integration: NO

Technology: Docker / R / Python



3. Security and Distributed Heterogeneous Executions

Tesseract Architecture
Explain technical aspects of
the architecture **tested** in
the development of the
workflows





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Thanks!

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