



# AGINFRA+: Virtual Research Environments to Support Agriculture and Food Research Communities

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Dissemination level: Public



o eosc-hub.eu





### **Overview**

- <u>Aim</u>: **Deploy a DataMiner cluster and make it** available to D4Science communities
- <u>Maturity</u> (TRL) level: TRL9
  - EGI FedCloud
- EOSC services:
  - EOSC Federated Authentication
  - EOSC Accounting Service (through EGI FedCloud)
  - EOSC Monitoring Service



# **Summary - Integration**

- Cloud Compute service
  - IFCA-LCG2 cloud site since April 17th
  - SLA (01/04 -> 31/12)
    - https://documents.egi.eu/public/ShowDocument?docid=2875

### EOSC Accounting Service

- Sum Elapsed \* Num Processors: 67980 hours
- May: 36 tasks
  - 15K hours, 241Gbytes
- June: 25 tasks
  - 25K hours, 170Gbytes
- July: 1 task
  - 14K hours, 65Gbytes



# **Summary - Integration**

### • EOSC Monitoring Service

- Availability/Uptime
  - JIRA issue: <u>https://jira.eosc-hub.eu/browse/EOSCWP10-117</u>
- Specific Nagios probes will be implemented to check the performance of the DM cluster
  - Currently on discussion if this is feasible through DM cluster API
  - JIRA issue: <u>https://jira.eosc-hub.eu/browse/EOSCWP10-116</u>



### Summary - Roadmap status

Quarter 1	<ul> <li>Identify the initial resource capacity [DONE]</li> <li>Engage with EOSC-hub providers [DONE]</li> <li>Deploy DM cluster in a single cloud provider [IN PROGRESS][DONE]</li> <li>Enabling EOSC-hub monitoring and accounting [DONE]</li> </ul>
Quarter 2	<ul> <li>Quarterly review of performance thresholds         <ul> <li>Analytics tasks increased 10% [May: 36, June: 3, July: 1]</li> <li>Availability (uptime) &gt; 95% [DONE]</li> </ul> </li> <li>Availability of monitoring probes for checking the status of the DM cluster (through API) [IN PROGRESS, JIRA's EOSCWP10-117]</li> </ul>
Quarter 3	<ul> <li>Quarterly review of performance thresholds (see Q2)</li> <li>Integration of monitoring probes in EOSC Monitoring service</li> </ul>
Quarter 4	<ul> <li>Quarterly review of performance thresholds (see Q2)</li> <li>Review of EAP experience and assessment of</li> </ul>
	<ul> <li>operational continuity of the application</li> <li>Registration of AGINFRA+ VREs in the EOSC</li> </ul>



# - Summary Technical Coordination

#### **Community Requirements DB**

#### <u>AGINFRA+ entry</u> created:

- Capacity Reqs
- Technical Reqs: JIRA issues
  - EAP services not listed in JIRA, how can they be tracked?

### **EOSC Portal**

Not yet there, should this be done by the EAP (through provider portal)?

Requirement number	Requirement title	Link to Requirement JIRA ticket	Source Use Case
Example	EOSC-hub to provide an FTS data transfer service	EOSCWP10-21 - EOSC-hub to provide an FTS data transfer service IN PROGRESS	UC1
RQ1	DM cluster deployment in the EGI Cloud Compute service	DONE	UC1
RQ2	Tracking compute consumption		UC2
RQ3	Get performance metrics: number of analytics tasks	i n° analytics tasks May: 36 June: 3	UC3
RQ4	Get performance metrics: availability / uptime (percentage)	EOSCWP10-117 - Get performance metrics: availability / uptime for AGINFRA+ service TODO	UC3
RQ5	Display of performance metrics in Nagios	EOSCWP10-116 - Display of performance metrics of DM cluster in EOSC monitoring service T000	UC3

### Thank you for your attention!

Questions?





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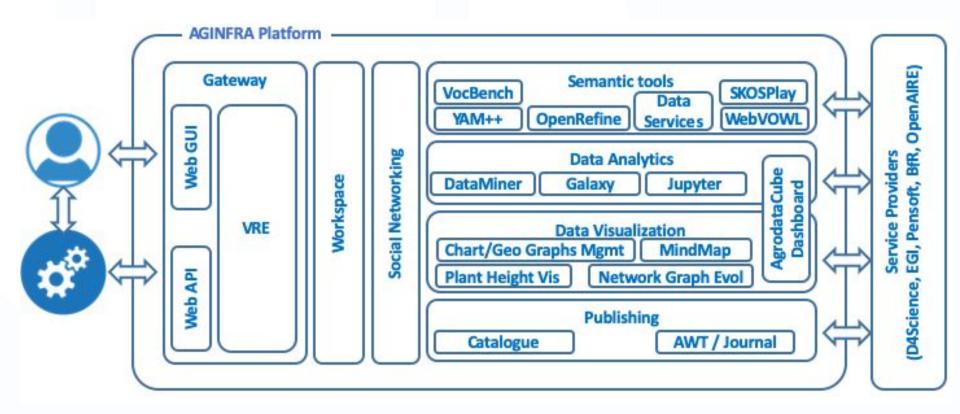


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## Background

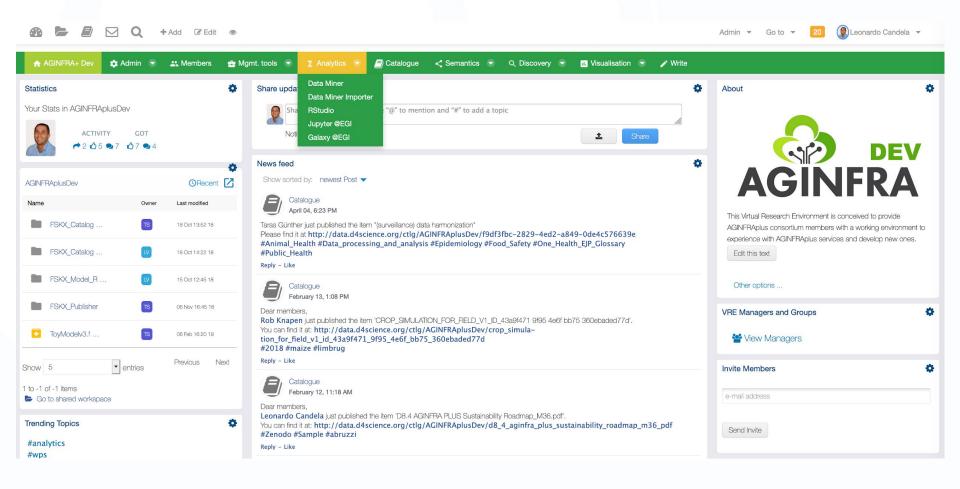
#### https://aginfra.d4science.org/home





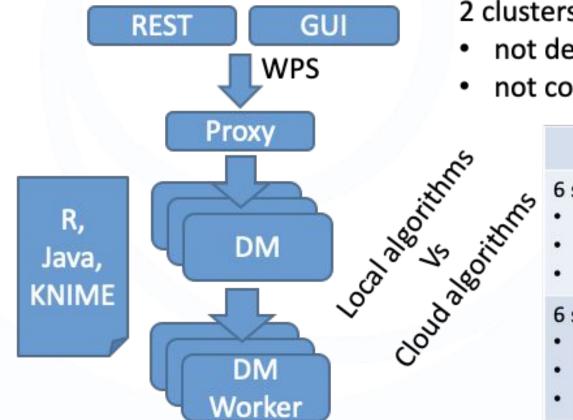
### Background

#### https://aginfra.d4science.org/home



**EOSC-hub** 

# **Background: DataMiner**



2 clusters:

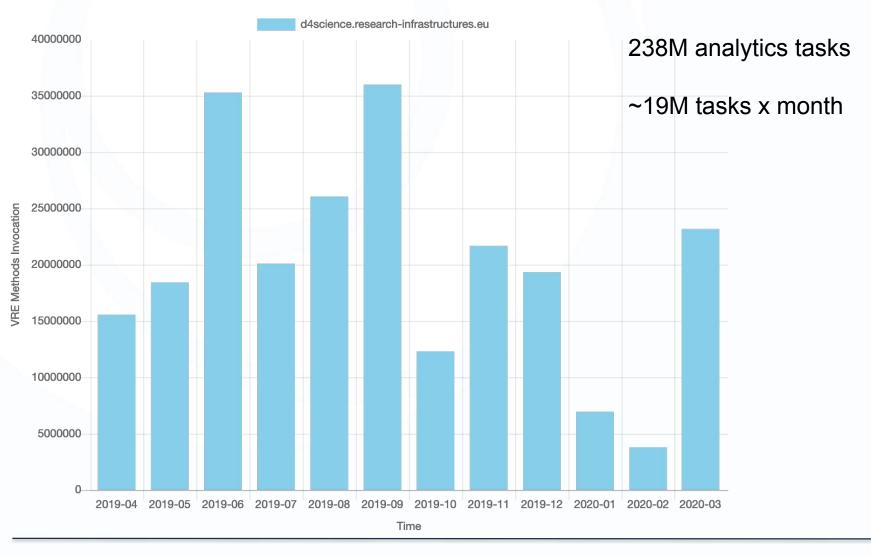
- not dedicated to AGINFRA+
- not co-existing in the same VRE

Proto	Prod		
6 servers	15 servers		
<ul> <li>16 vCores</li> </ul>	<ul> <li>16 vCores</li> </ul>		
• 16 GB RAM	• 32 GB RAM		
• 100 GB Space	• 100 GB Space		
6 servers	15 servers		
<ul> <li>16 vCores</li> </ul>	<ul> <li>16 vCores</li> </ul>		
• 16 GB RAM	• 32 GB RAM		
• 100 GB Space	• 100 GB Space		



### **Background: DataMiner**

d4science.research-infrastructures.eu VRE Methods Invocation [Method Invocation]





### The BioCoS use case



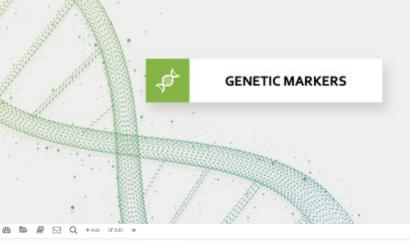
The tool allows a fast identification of speciesspecific genomic loci containing potential genetic markers.

The functional principle is based on the identification of sequences non-tolerated in a predefined set of genomes (Query) but evolutionary conserved in the genome of interest (Target).

It can be applied for the identification of genetic markers for inter- and intra-species analyses – in base of the biological need.

Advantages:

- 1) Overcomes whole genome alignments
- 2) Reduces sequencing needs



🕈 AGINFRA+ for BioCoS 🛛 🏚 Admin 🕤	Members ∑ Analytics	Catalogue	< Semantics 💿	Q Discovery 🕤 🛛	🖪 Visualisation 🕤	🖍 Write
DataMiner	go back		Access to the Data Space	8	Execute an Experiment	Check th
Operators	Computations Execution	Call Service Profile				
🛸 🗄 🛃 🗡 🗡 🕹 User 🕶	Tools: 🙁 Remove All Operators					
BIOCOS (1) V					•	
Inter Species Cloud Inter Species Cloud (Published by Glancarlo Panichi (giancarlo panichi) on 2020/02/10 15:51 GMT)	Inter Species Cloud (Publisher	ed by Giancarlo Panich	i (giancarlo.panichi) on		Inter Species Cloud	ł
BLACK BOX (1)	genomeDir:	Select Item Genome I	Directory			
CHARTS (1)	targetGenomeDir:	Select Item Target Ge	nome Directory			
PIPELINE (1)	minAwLen: 1		Min AW Len			
		nteger Value	Max AW Len			
		U teger Value	Place Party Learn			
		600	MAW Memory in MB			
		iteger Value				
			Verbose			
	debug: fa	alse	<ul> <li>Debug</li> </ul>			
	Start Computation					
					Samerica	225