



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



- 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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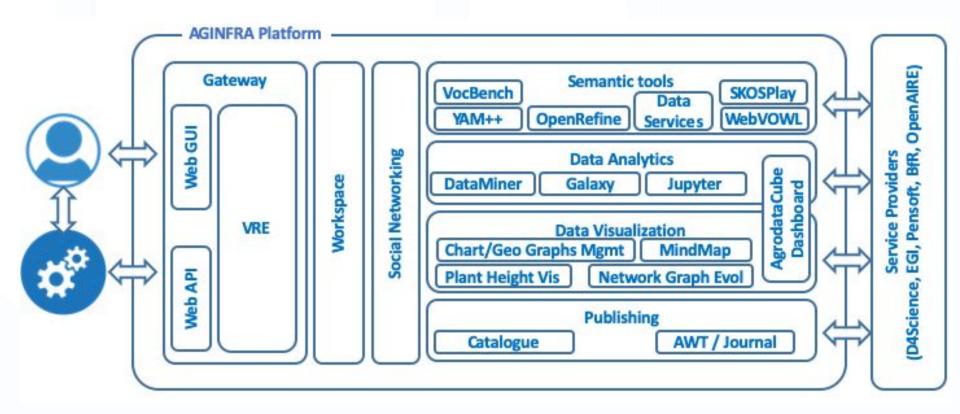


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Background

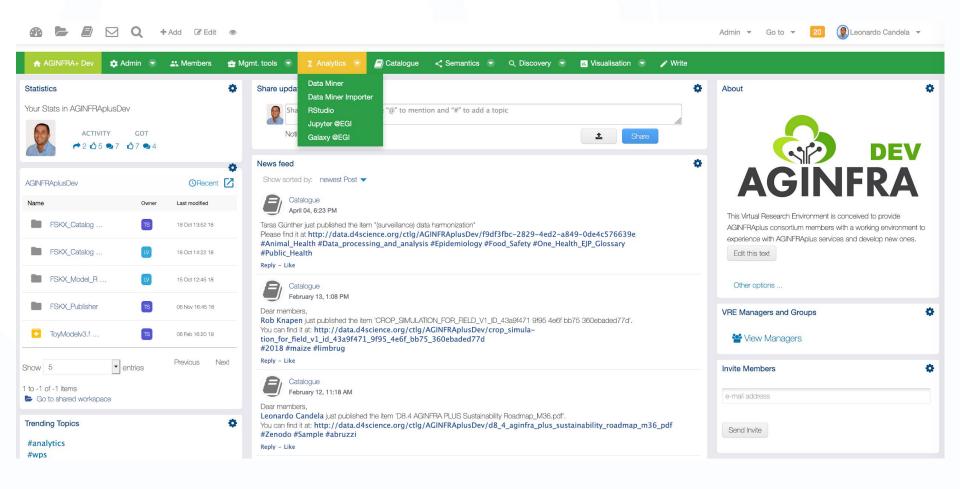
https://aginfra.d4science.org/home





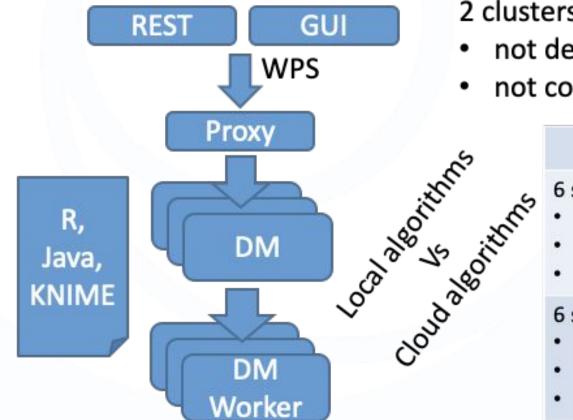
Background

https://aginfra.d4science.org/home



EOSC-hub

Background: DataMiner



2 clusters:

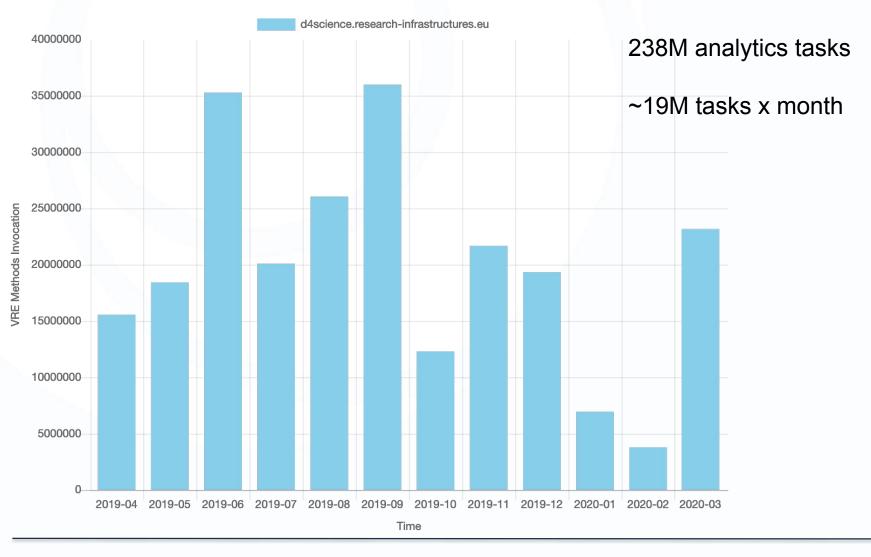
- not dedicated to AGINFRA+
- not co-existing in the same VRE

| Proto | Prod | | |
|-------------------------------|-------------------------------|--|--|
| 6 servers | 15 servers | | |
| 16 vCores | 16 vCores | | |
| • 16 GB RAM | • 32 GB RAM | | |
| • 100 GB Space | • 100 GB Space | | |
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| 16 vCores | 16 vCores | | |
| • 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



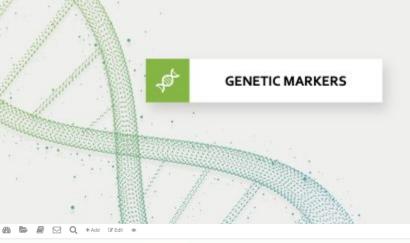
The tool allows a fast identification of speciesspecific genomic loci containing potential genetic markers.

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 | Debug | | | |
| | Start Computation | | | | | |
| | | | | | Samerica | 225 |