



**OpenBioMaps data management service for
biological sciences and biodiversity conservation**

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- The OpenBioMaps (OBM) is a free and open-source database framework for scientific and conservation purposes.
- Science areas:
 - Biology, conservation biology, ecology, biodiversity
- The OBM system is used for data management by nature conservation institutes, biodiversity research and citizen science projects.
- OBM provides a number of services that make day-to-day work with data easier, but it does not yet provide tools for analyzing the data. Particularly for high computing tasks.
- Current use cases:
 - <https://openbiomaps.org/projects/>

- Impressions of some projects



- Develop a background service that supports the interpretation of data from databases on conservation biology and biodiversity.
- A solution that facilitates and generalizes the most common high-computational analysis of data stored in such databases.
- We would like to create a service with EOSC that allows multiple users to run tasks that are above the level of a PC through the same interface. In fact, we would like to develop a “service in service” - specifically for projects that collect nature conservation and biodiversity data.



- The most common computing tasks come from the following areas:
 - Spatial analyses (satellite image processing, large or fine scale spatial queries),
 - ML analyses (satellite image analyses, drone image analyses, distribution models (Random Forest), population dynamics analyses, survival analyses, supervised and non-supervised image classification - photos of individual animals, habitats, survey methods),
 - Conservation genetics analyses.
- To serve these diverse tasks we need a fully configurable VM which let us deploy our service interface (API) which will be available in the OpenBioMaps Network and provide computation capacity access to the involved projects.

- According to our recent experiences in our PC based local computational cluster, the number of processors is the most important in these ecological analyzes. A “typical” analysis is now running at an acceptable (few hours to few days) rate on **16 CORES**.
- The parallel computing requirements of image analysis can be much higher, and **GPU usage can be interesting** there. Some analyzes, for example, genetic analyzes or larger spatial analyzes require **a lot of memory**.

- Minimal Compute and Storage capacity needed for sustaining the Project:
 - 32 CPU cores
 - 96 Gb RAM (3Gb RAM per cores)
 - 2 Tb HD
 - Compute and Storage capacity to fully scale-up the Project after the completion of the pilot:
 - 128 CPU cores
 - 384 Gb RAM (3Gb RAM per cores)
 - 8 Tb HD
 - GPU access will be interesting.
 - **IFCA site** has accepted to provide resources to this EAP case.
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1. Developing a new and flexible API interface together with the OBM user community for supporting conservation science.
 2. Creating a database-based machine learning analysis system that can significantly support the use of data collected in nature conservation and improve the way new data is collected.
 3. Creating a long-term computational layer for OBM
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- Q1:
 - Integration with EGI Cloud Compute.
 - Deploy and Configure OBM node in test environment manually.
- Q2:
 - Create TOSCA Recipes and Ansible roles needed to deploy the application automatically using IM.

- Q3:
 - Deploy OBM node to production environment using the developed recipes.
 - Analyse EOSC data services to be used by the application:
 - EGI DataHub.
 - B2 services (Drop, Find, Handle, Share)
 - EGI Services (Training Infrastructure, Data Transfer)
 - EOSC Marketplace services (GeoDAB, D4Science spatial services, Alien and Invasive Species Virtual Research Environment, Biodiversity, EODC JupyterHub for global Copernicus data)
- Q4:
 - Performance test of all nodes.

- Person Months: 3,5.
- Budget: 21.658 €
 - 3 PMs to IFCA as resource provider.
 - One VM with 48 vCPU, 96GB RAM, 2 Tb HD
 - 0,5 PM to UPV for integration issues.

Thank you for your attention!

Questions?



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