CeSGO

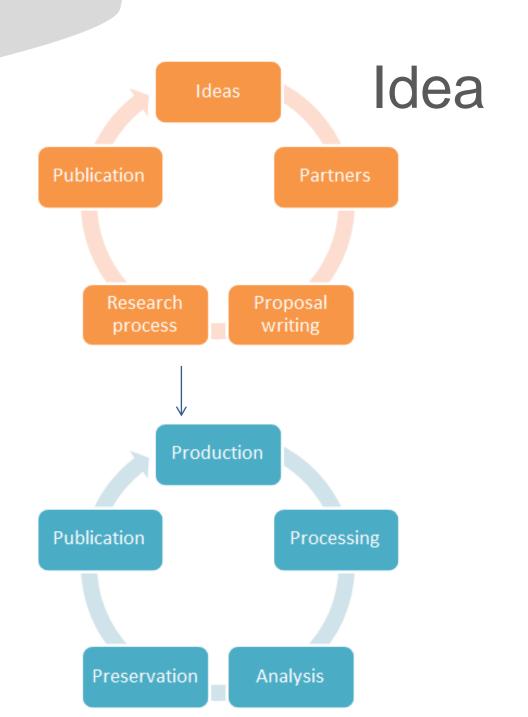
An environment for scientific collaboration

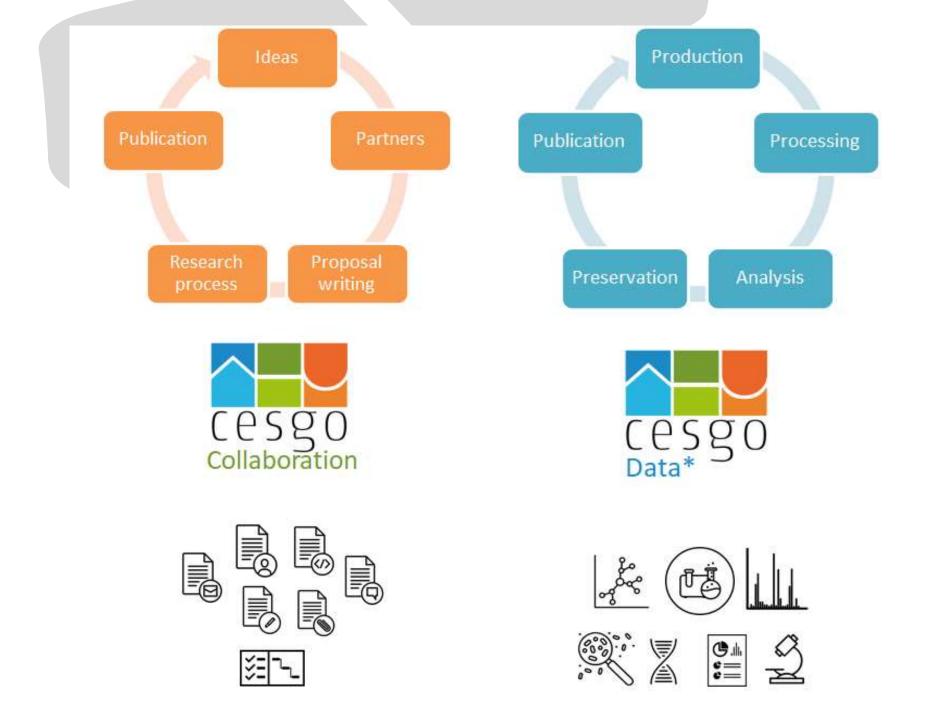
INTRODUCTION

Background

- 2010: IBiSA funding
 - EMME project
 - First steps on experimental metadata
- 2012 : Région Bretagne et Pays de la Loire funding
 - e-Biogenouest and CeSGO
 - Based on HubZero solution
- 2015 : European funding (CPER)
 - CeSGO v2
 - Five services for projects and data life cycle

To provide an integrated environment to help scientists to work from project ideas to publication through data production and management





What can be done with CeSGO



Create groups



Share ideas



Exchange



Edit and share documents



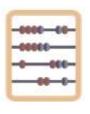
Create websites



Manage and share files



Manage scientific data



Launch compute jobs



Manage projects

CeSGO services



Collaboration

Exchange with you community



Data access

Sync and share research data



Research sharing

Store and publish research data



Projects

Manage your projects efficiently with kanban



Instant

Live chat with your partners



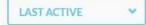
Powered by Wordpress

COLLABORATION

Main features

- Work alone or inside a group
- Documents: create, edit and manage authorizations
- Files: upload files (pdf, images, etc.)
- Forums: create topics inside a group
- Websites: create a website with Wordpress

Groups



ALL TYPES Search Groups...

WHO'S ONLINE



ALL GROUPS



BIOGENOUEST.ORG

active 5 days, 23 hours ago

Refonte du site web du GIS Biogenouest : suivi, avis, et tests.

Private Group / 10 members



Communication-Group created to support the communication of a research project or team Internship-Group created to support an intership



CESGO

active 6 days, 21 hours ago

Group to share informations about CeSGO internal project

Project tasks

Report

Private Group / (3) members

GenOuest-Group dedicated to the GenOuest core facility



GENODOCK

active 1 week, 3 days ago

GenoDock est un nouvel environnement dédié à la gestion de données génomiques. Cet outil offre la possibilité de mettre à disposition de la communauté des génomes ainsi que leurs annotations (structurales et fonct [...]

Private Group / 5 members



GenOuest-Group dedicated to the GenOuest core facility

GenOuest R&D-Research & Developement group dedicated to GenOuest core facility

MEMBERS

Newest Active Popular



Cyril MONJEAUD a few seconds ago



Grégoire Siekaniec an hour ago



Léa Cabioch an hour ago



Olivier Collin 3 hours ago

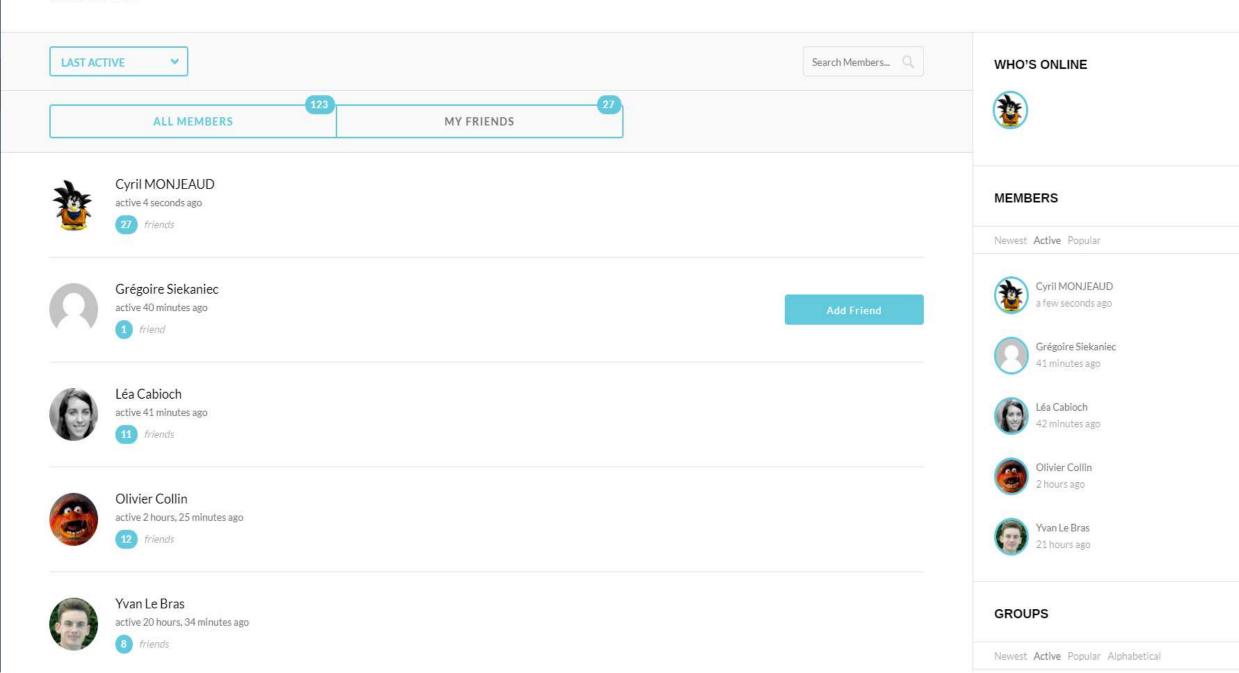


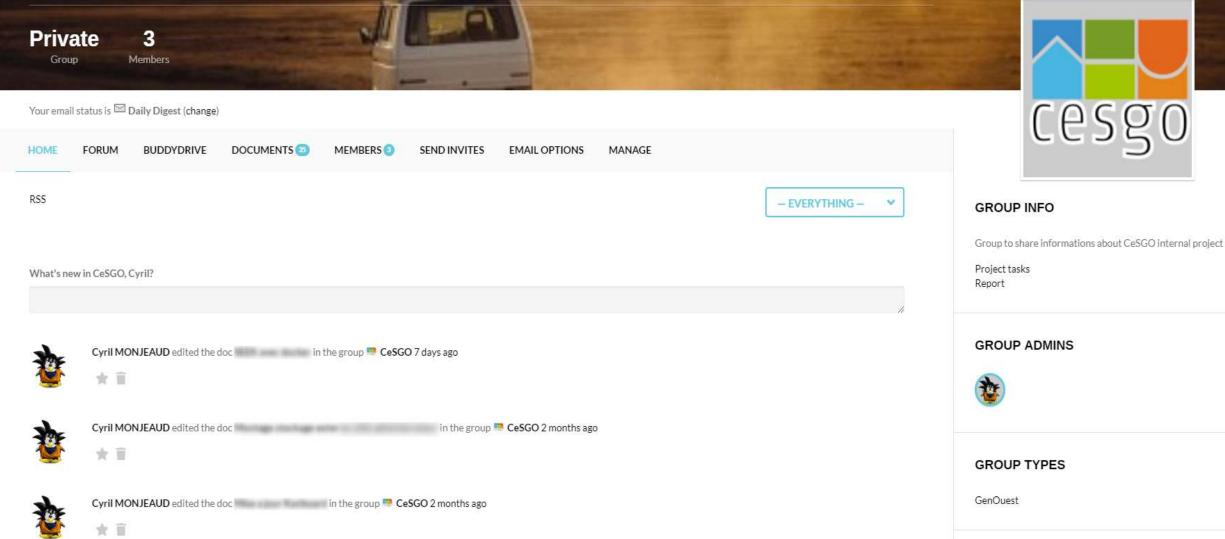
Yvan Le Bras 21 hours ago

GROUPS

Newest Active Popular Alphabetical

Members



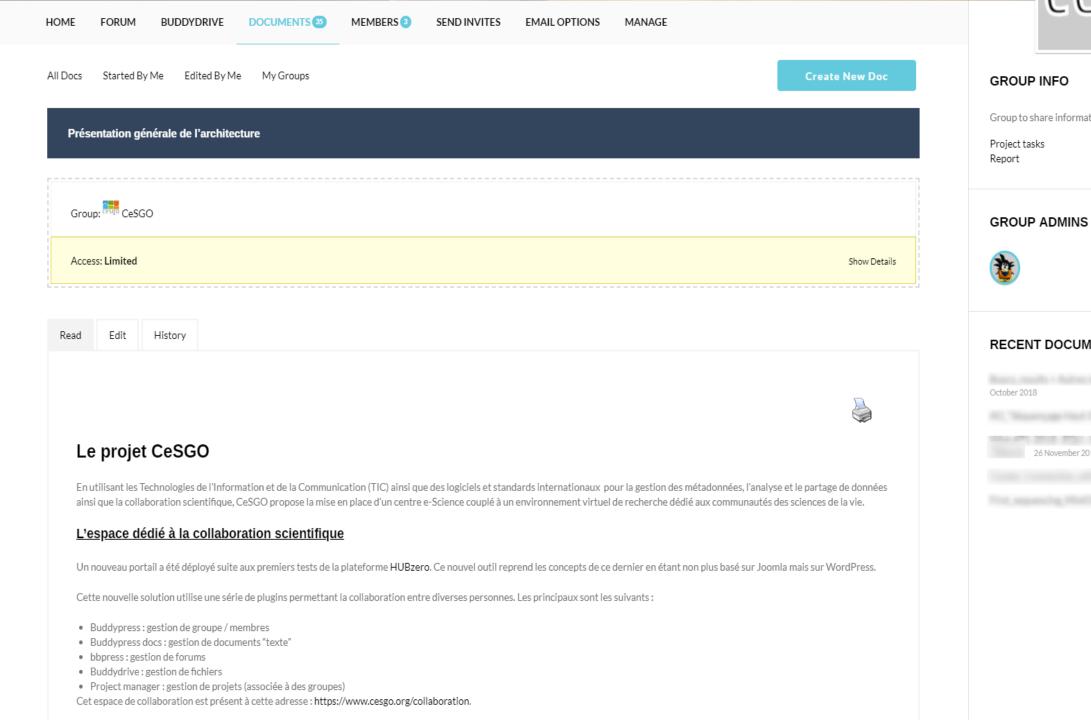


Cyril MONJEAUD created the doc in the group Residue CeSGO 2 months ago

Anthony Bretaudeau joined the group # CeSGO 2 months ago

RECENT DOCUMENTS







GROUP INFO

Group to share informations about CeSGO internal project

Project tasks Report



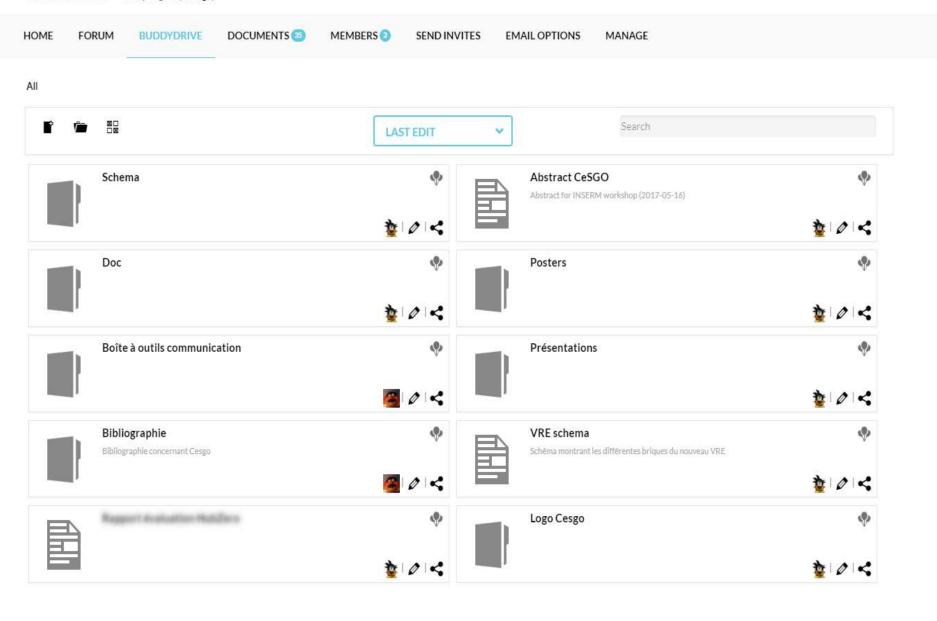
RECENT DOCUMENTS

October 2018

26 November 2018

26 November 2018

15 November 2018





GROUP INFO

Group to share informations about CeSGO internal project

Project tasks

Report

GROUP ADMINS



GROUP TYPES

GenOuest

RECENT DOCUMENTS

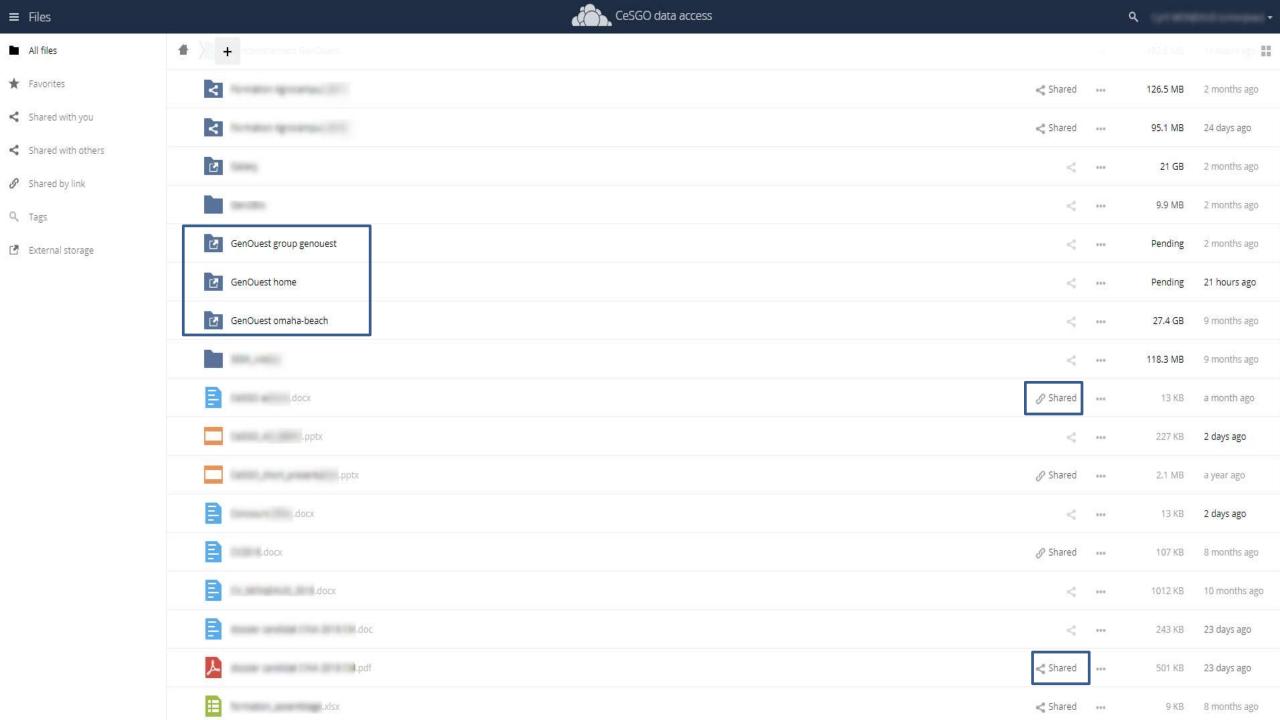


Powered by Owncloud

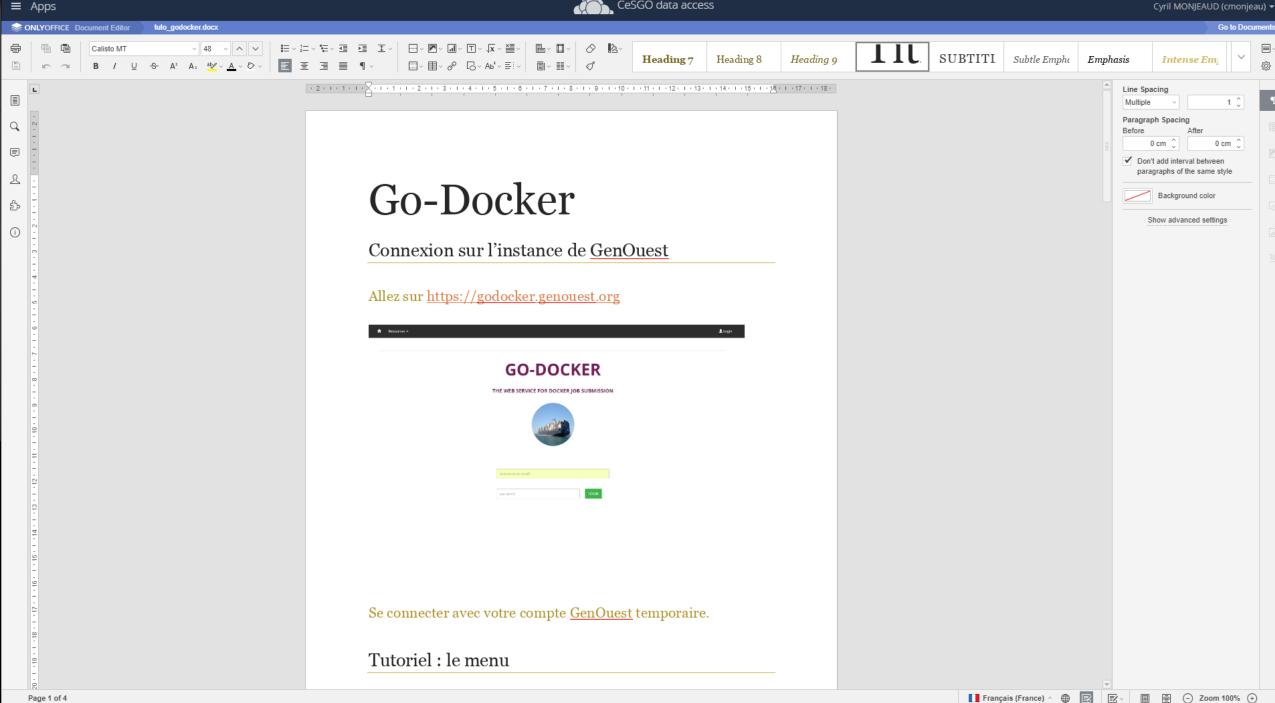
DATA ACCESS

Main features

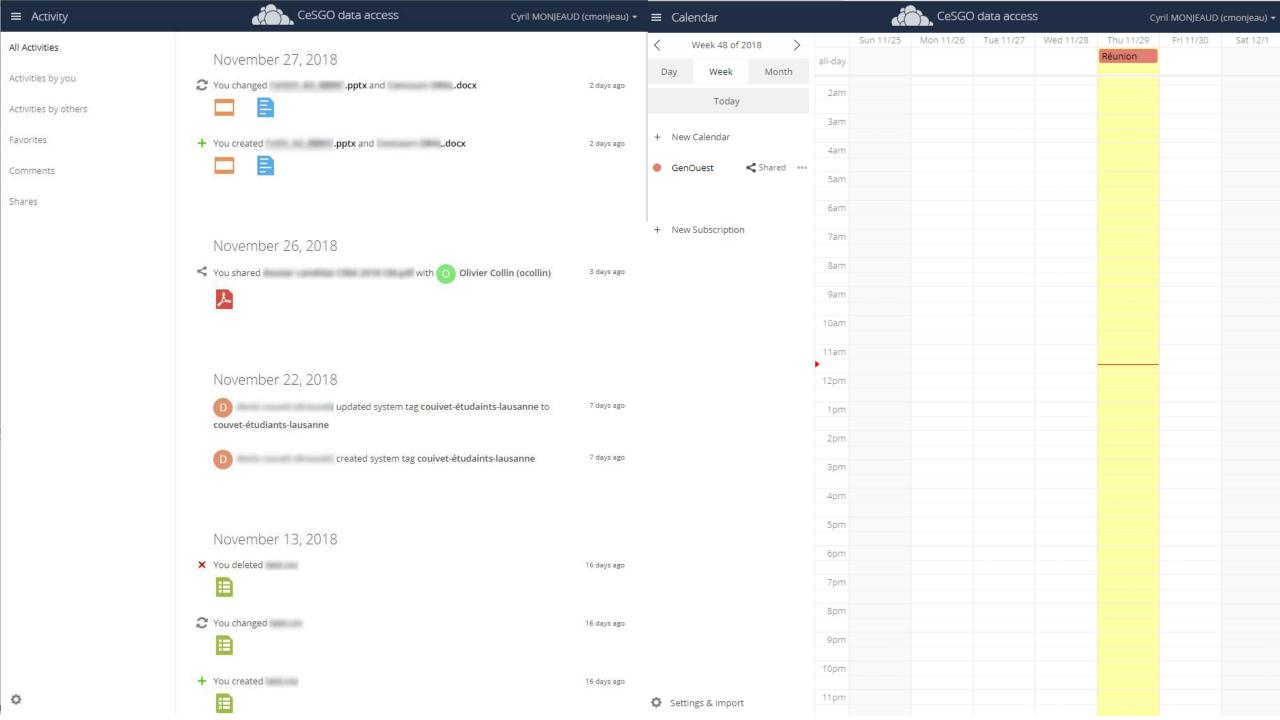
- Storage: limited to 50Go
- Accessibility: GenOuest folders (home, group, omaha)
- Sharing: generate a public link or share with user / group
- Online collaborative documents
- Synchronization: desktop clients and applications are available
- External: add external storage (FTP / SFTP/ etc.)







Page 1 of 4

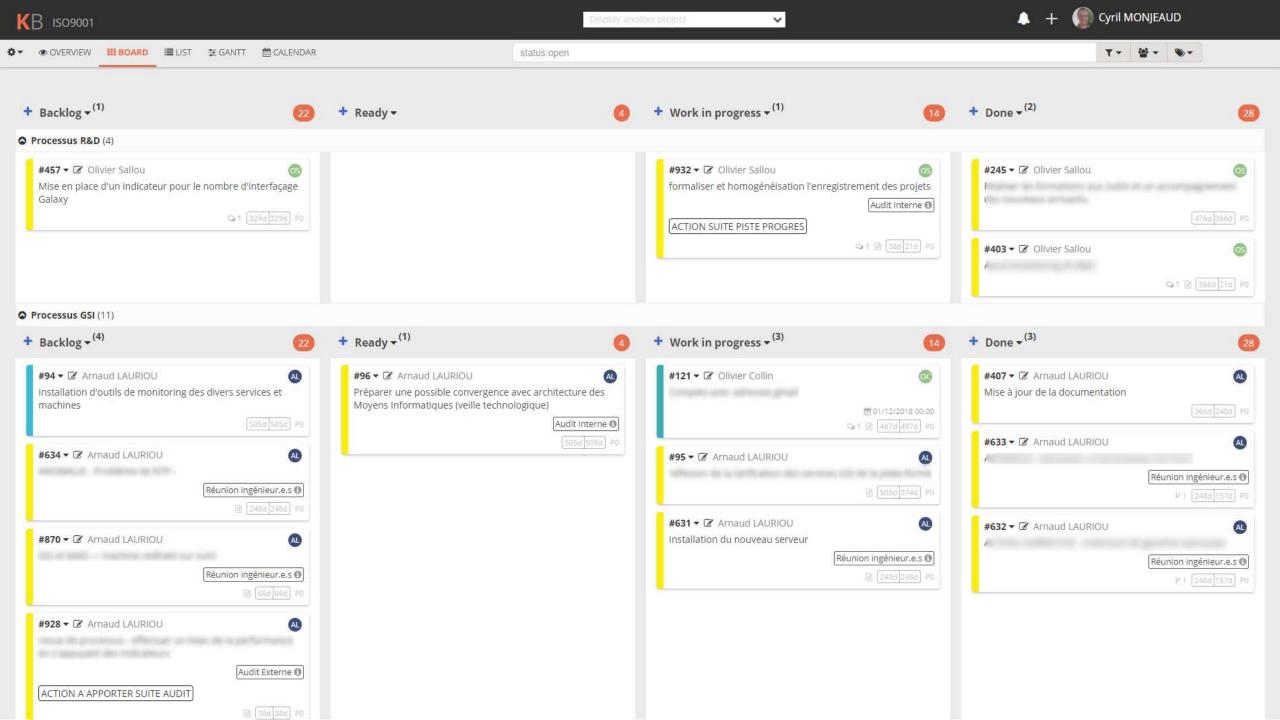


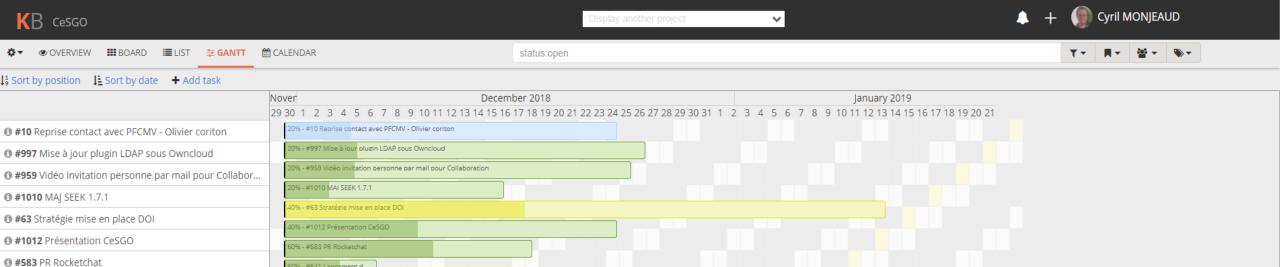
Powered by Kanboard

PROJECTS

Main features

- Projects: private or collaborative based on Kanban
- Gantt chart
- Permissions: share a project with users and manage roles
- Notifications: email and web notifications.
- Activities: Reports and graphs of each project

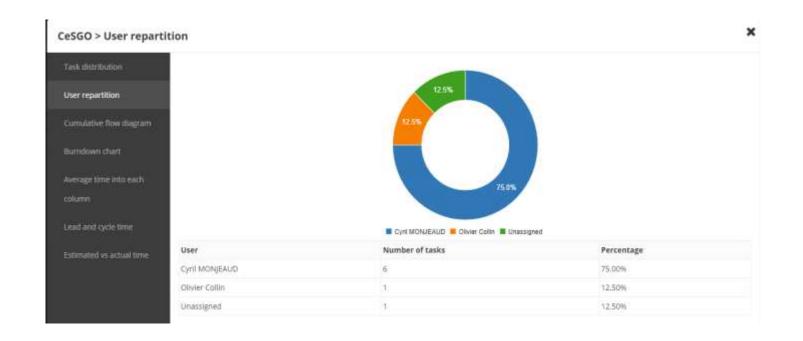




Moving or resizing a task will change the start and due date of the task.

1 #641 Lancement des dockers auto

60% - #641 Lancement d...



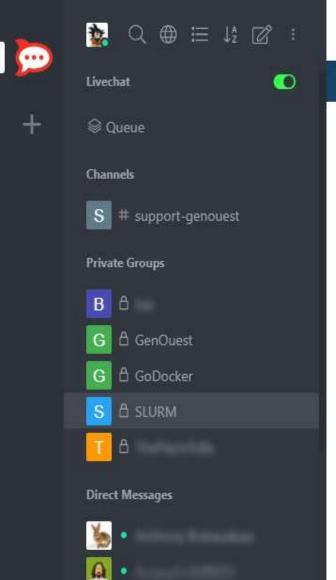


Powered by Rocket.Chat

INSTANT

Main features

- Chat rooms: communicate and collaborate using team chat
- Direct chat: private conversation between two users
- Video conference: create a video conference inside a room
- Anywhere: desktop & mobile apps







Parlons SGE en fait

```
Exception in thread "main" java.awt.AWTError: Can't connect to X11 window server using 'localhost: 27.0' as the value of
the DISPLAY variable.
       at sun.awt.X11GraphicsEnvironment.initDisplay(Native Method)
       at sun.awt.X11GraphicsEnvironment.access$200(X11GraphicsEnvironment.java:65)
       at sun.awt.X11GraphicsEnvironment$1.run(X11GraphicsEnvironment.java:115)
       at java.security.AccessController.doPrivileged(Native Method)
       at sun.awt.X11GraphicsEnvironment.<clinit>(X11GraphicsEnvironment.java:74)
       at java.lang.Class.forName@(Native Method)
       at java.lang.Class.forName(Class.java:264)
       at java.awt.GraphicsEnvironment.createGE(GraphicsEnvironment.java:103)
       at java.awt.GraphicsEnvironment.getLocalGraphicsEnvironment(GraphicsEnvironment.java:82)
       at sun.awt.X11.XToolkit.<clinit>(XToolkit.java:126)
       at java.lang.Class.forNameO(Native Method)
       at java.lang.Class.forName(Class.java:264)
       at java.awt.Toolkit$2.run(Toolkit.java:860)
       at java.awt.Toolkit$2.run(Toolkit.java:855)
       at java.security.AccessController.doPrivileged(Native Method)
       at java.awt.Toolkit.getDefaultToolkit(Toolkit.java:854)
       at javax.swing.UIManager.getSystemLookAndFeelClassName(UIManager.java:611)
       at uk.ac.babraham.FastQC.FastQCApplication.main(FastQCApplication.java:322)
```



5:21 PM

autant pour moi, c'est bon pour le x11, j'ai mis à jour la doc







Powered by SEEK

RESEARCH SHARING

FAIR data

- FAIR = Findable, Accessible, Interoperable, and Re-usable.
- This service aims to improve the integration of the FAIR data model, by helping <u>linking data with its experimental information</u>
- Data can be imported from a local storage, or through a direct link to the data (such as the Data access service links)

Experimental information

ISA formalism is used to describe biological activities.

Investigation

Project context

e.g. Growth control of the eukaryote cell: a systems biology study in yeast

Study

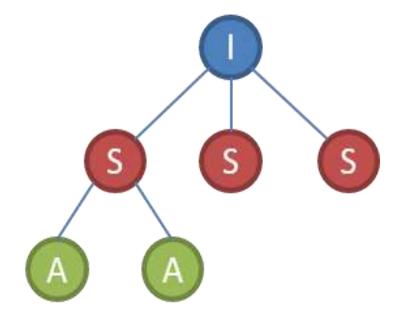
A unit of research

e.g. Study of the impact of changes in flux on the transcriptome under different nutrient limitations

Assay

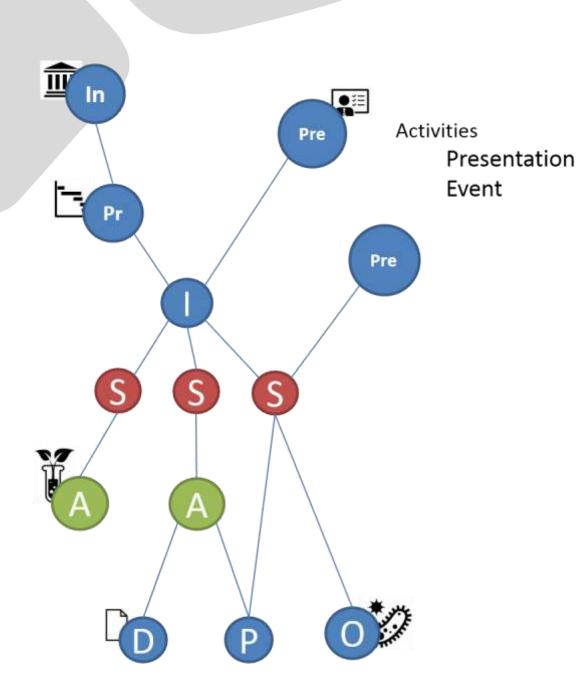
Analytical measurement

e.g. Protein expression profiling using mass spectrometry



Assets

- Yellow pages
 - Institution
 - Project
- Biological Experiments
 - Investigation
 - Study
 - Assay
- Assets
 - Data file (upload and remote)
 - Model (sbml)
 - SOP (protocoles)
 - Publication



Home / Projects Index / CeSGO project



En utilisant les Technologies de l'Information et de la Communication (TIC) ainsi que des logiciels et standards internationaux pour la gestion des métadonnées, l'analyse et le partage de données ainsi que la collaboration scientifique, CeSGO propose la mise en place d'une plate-forme e-Science couplée à un environnement virtuel de recherche centré sur les besoins des communautés des sciences de la vie.



SEEK ID: https://research-sharing.cesgo.org/projects/2 CesGO PALs: No PALs for this Project

Public web page: https://www.cesgo.org

Organisms: Escherichia coli, Lactococcus lactis, Oncorhynchus mykiss

Related items

People (3)

Institutions (3)

Investigations (2)

Studies (2+1) Assays (2+1) Data files (7+2) Models (1) SOPs (1) Publications (4)

CeSGO Admin 🦞 🔮



Projects: Training, CeSGO project Institutions: Default Institution

No description specified

Disciplines: Not specified Roles: Not specified Expertise: Not specified

Tools: Not specified

Cyril Monjeaud 🍄



Projects: CeSGO project, Rapsodyn, EnginesOn project, FEED-A-GENE, Training, CORAIL MAD

Institutions: GenOuest, Biogenouest

Disciplines: Bioinformatician Roles: Not specified

Expertise: Data Management, Software Engineering Tools: Web services, cloud, Workflows, docker, Python, galaxy

No description specified

Olivier Collin 🍳



Projects: CeSGO project, Rapsodyn, Training

Institutions: GenOuest

(b) orcid.org/0000-0002-8959-8402

No description specified

Disciplines: Bioinformatician Roles: Project Coordinator

Expertise: Data Management, Bioinformatics

Tools: Not specified

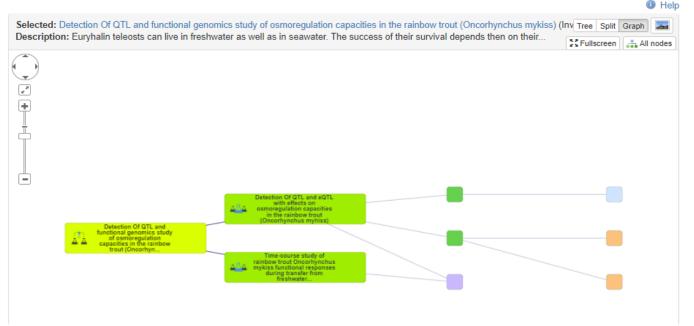
Home / Investigations Index / Detection Of QTL and functional genomics study of osmoregulation capacities in the rainbow trout (Oncorhynchus mykiss)

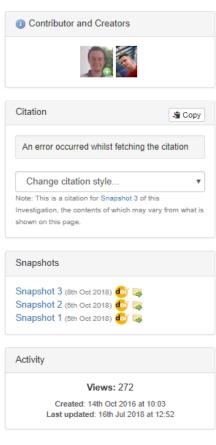
Detection Of QTL and functional genomics study of osmoregulation capacities in the rainbow trout (Oncorhynchus mykiss)

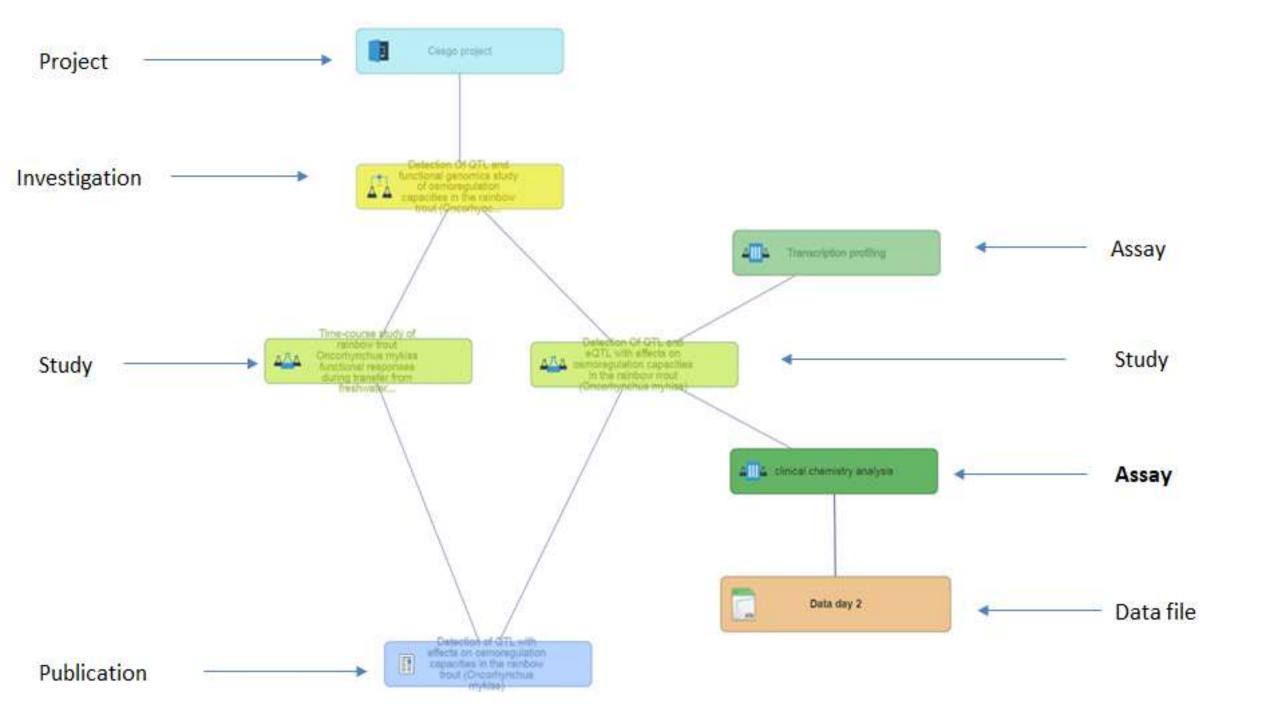
Euryhalin teleosts can live in freshwater as well as in seawater. The success of their survival depends then on their osmotic acclimation capacities. The objective of my work is to describe acclimation processes in the salted water at the rainbow trout by a study coupling functional genomic and genetic approaches. From a first differential gill transcriptomic study, a list of candidates genes was established. This study also allowed to investigate the physiological answer to a hyperosmotic kinetics challenge. Main results reveal good euryhalinity capacities of the tested trouts and a maximum transcriptomic answer 24 h after the seawater transfer. Biological processes involved in the acclimation mechanisms are also proposed. A second part of this work consisted of the characterisation of the genetic control of processes linked to seawater acclimation in rainbow trout. Using as characters, a two times repeated 24h post seawater transfer plasma sodium and chloride levels, as well as gill weight, unitrait and multitrait analyses allowed to reveal 18 QTL among whom 9 are qualified as robust. A last approach of eQTL detection then allowed, based on a gill transcriptomic analysis and on the results of the first two approaches, to offer 69 exclusive candidates genes. If the majority of these genes are offered as acting at functional level, some are suggested as positional candidates. It is the first time that a blending transcriptomic differential approach coupled to a QTL / eQTL study is led to a non-sequenced genome aquacole interest teleost for the acclimation capacity to different osmotic environments. The existence of a genetic element having an influence on the capacities of osmoregulation in teleost is shown here. These results lay the road with cobblestones for a definite investigation of the genetic bases of seawater acclimation processes in teleosts.

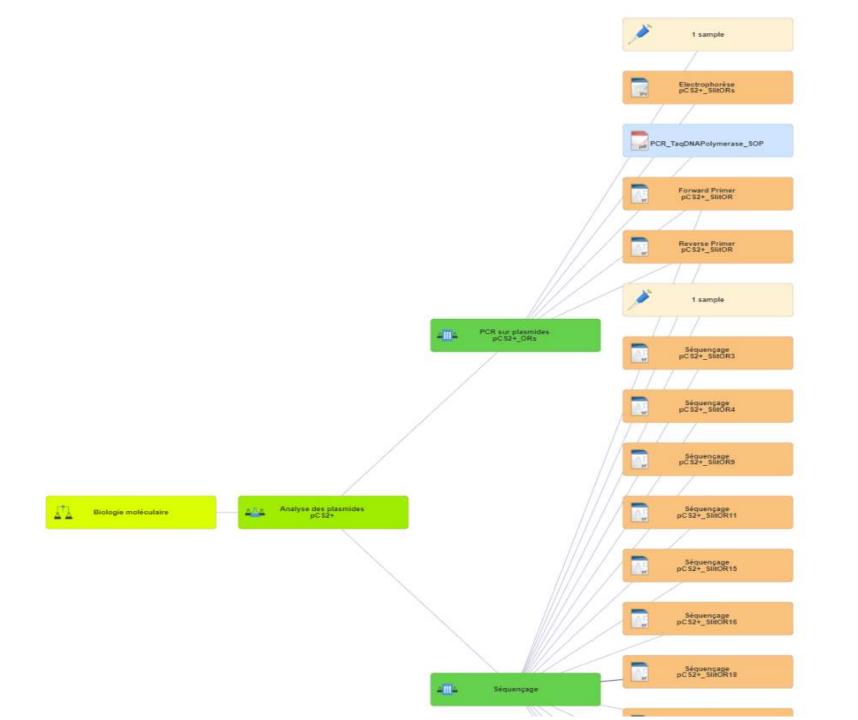
SEEK ID: https://research-sharing.cesgo.org/investigations/1

Projects: CeSGO project









CONCLUSION

People and link

People

- Yvan Le Bras CeSGO v1
- Cyril Monjeaud CeSGO v2
- Olivier Collin CeSGO v1-v2
- With all the GenOuest team

CeSGO portal

https://www.cesgo.org









QUESTIONS?