









Workflow4metabolomics

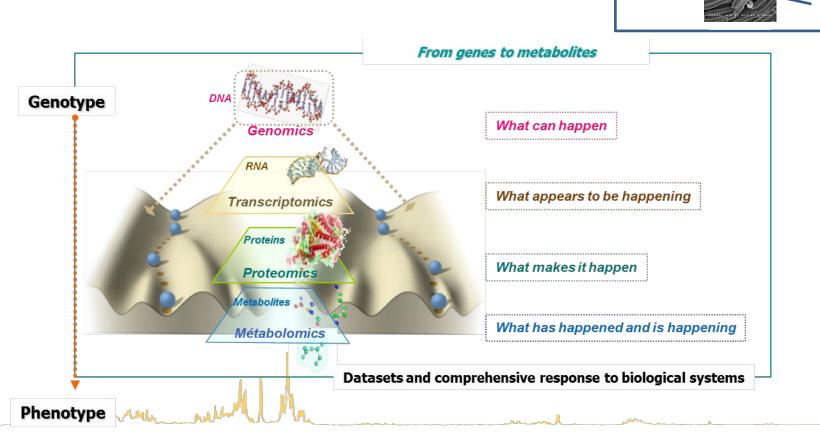
Towards an international computing infrastructure and a tools showcase for Metabolomics

GCC2017 29/06/2017 Gildas Le Corguillé, Franck Giacomoni Christophe Caron

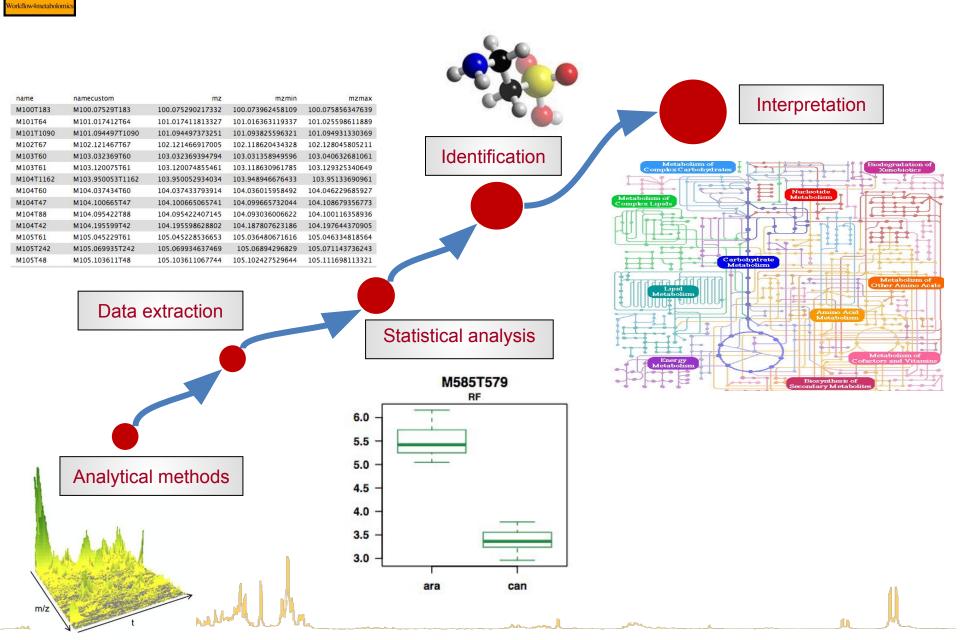


# **Biological context**

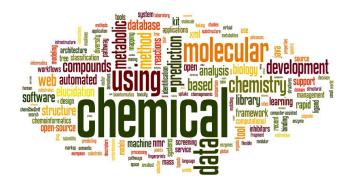
- Identification and quantification of small molecules from the metabolism (metabolites)
- the most recent -omics research domain



## Metabolomics: a data driven approach

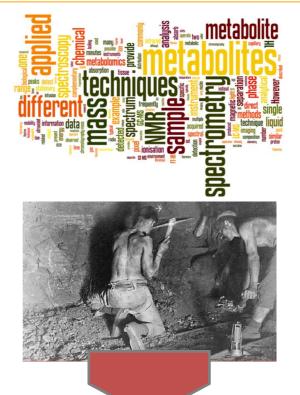


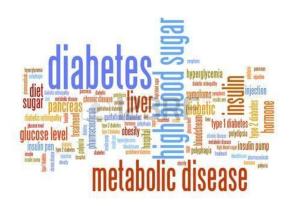
# **Wm** Galaxy, the perfect coach.



maria

orkflow4metabolon

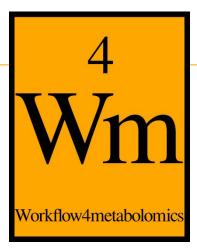






# Mm Once upon a time... W4M





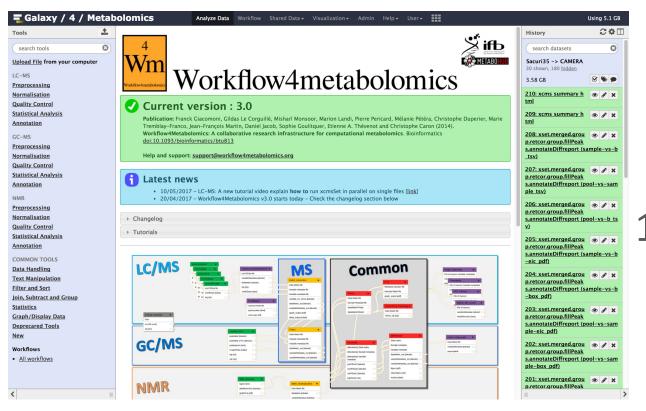
## **The W4M environment**

mr. M

# A main public instance

orkflow4metabolomi

### http://workflow4metabolomics.org

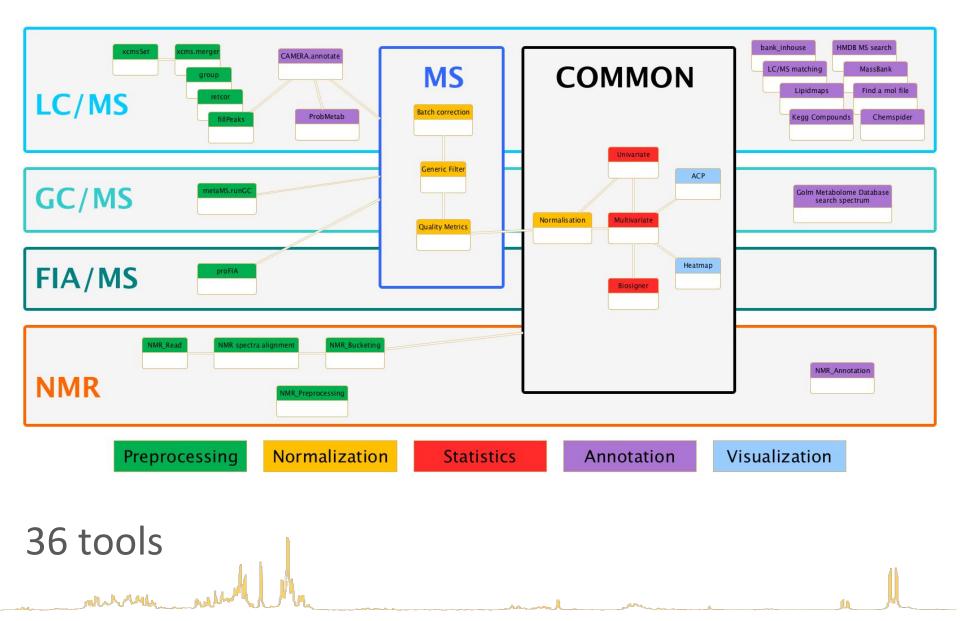


### 800 accounts 25 000 jobs/year

19 related papers

HelpDesk support@workflow4metabolomics.org







Workflow4Experimenters

- 3 sessions since 2014
- 10 Trainers for 20 Trainees
- "Bring your own data" tutoring sessions



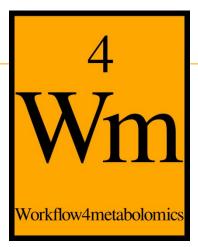




W4M provides DOI to reference histories and workflow usable in papers



http://workflow4metabolomics.org/referenced\_W4M\_histories Paper submitted in *IJBC* 



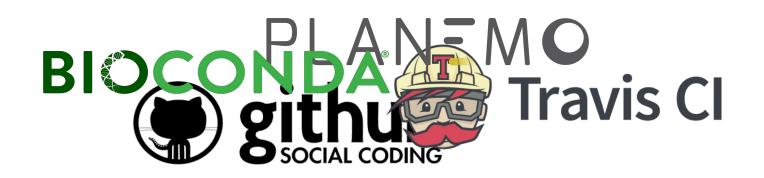
### **BACK OFFICE**

mr. Mah



#### Developments

- -15 developers from 7 laboratories
- -Stick to IUC standards
- -A GitHub: github.com/workflow4metabolomics
- Conda / Planemo / TravisCl



# Model For the second se

# All public W4M tools are distributed:

Dev: github.com/workflow4metabolomics



Wrappers: on main Galaxy ToolShed toolshed.g2.bx.psu.edu/groups#/175812cd7caaf439



VM: with the most common W4M tools workflow4metabolomics.org/virtual-environments



## Docker/Docker: A Galaxy Flavor

docker run -d -p 8080:80 workflow4metabolomics/galaxy-workflow4metabolomics



## **PUSH: Contribution**

The main W4M instance as a Showcase

Hosting on W4M main instance Quality standards (IUC Standard)



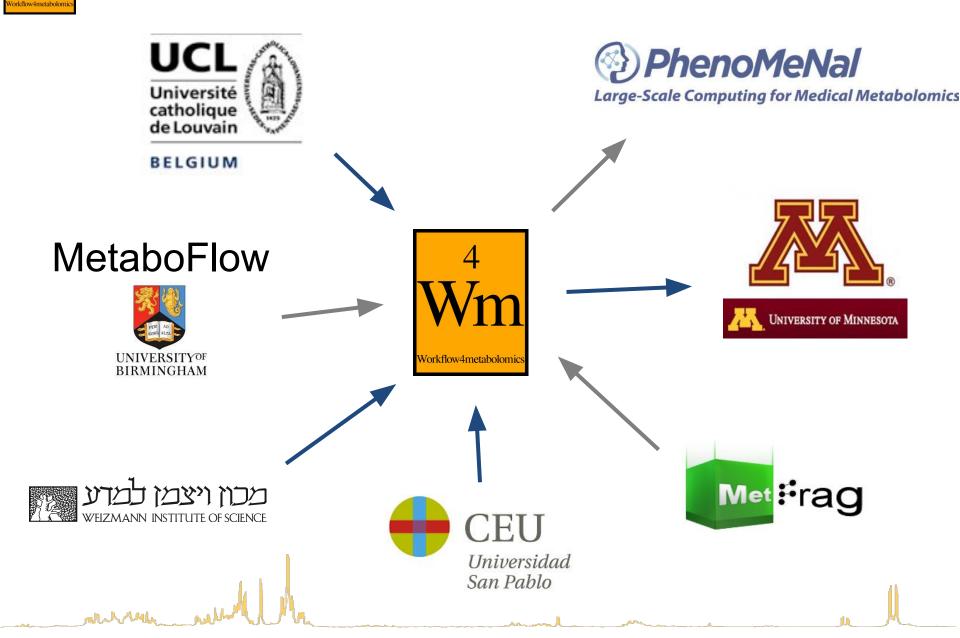
#### Advanced mode

Integration in our reference workflows Follow your exchange formats between tools Collaboration mode if help is needed

However, the support must be done by the developers themselve

https://github.com/workflow4metabolomics/workflow4metabolomics#how-to-contribute

# **Contribution in progress**

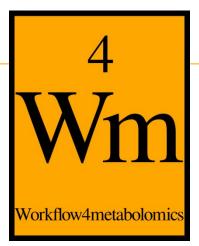


## Galaxy Training Network



#### Metabolomics Icms preprocess × work-in-progress

#355 opened 12 days ago by lecorguille



## PERSPECTIVE

marchal



Visualization



- Interoperability MetaboLights
- Trainings e-Learning (2018)
  - MOOC4Metabolomics
  - Open Course 'e-Bioinformatics4Metabo'





Wm

Vorkflow4metabolomics

## **MERCI** !

