# OP01: Galaxy Framework Exordium &

# HS03: Visualization of RNA-Seq with Galaxy framework

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## Overview

- Oral presentation
  - Introduction to Galaxy
  - Galaxy Training (training.galaxyproject.org)

- Hands-on presentation
  - An example <u>Workflow</u> which shows about visualization of RNA-Seq results

## Contribute to community

- Global community
  - Public instances
    - usegalaxy.org
    - usegalaxy.eu
    - Usegalaxy.org.au
- Domain specific
  - Cheminformatics
- Gitter channels
- Github Project
- Create an account on usegalaxy.eu or usegalaxy.org

- Regional Galaxy groups
  - Africa: <a href="https://africa.usegalaxy.eu/">https://africa.usegalaxy.eu/</a>
  - o India:
    - usegalaxy.in
    - TMS Foundation
      - Become Member
        - https://forms.gle/rZkmk5tWP4XG36fg8
- Gmentee/Gmentor program
  - https://galaxy-mentor-network.ne tlify.app/

# Galaxy global training

# **GTN Tapas**

bit.ly/gtntapas





Free Global Galaxy Training!
March 14-18, 2022
Covid, RNA-Seq, Single Cell,
Proteomics, ML, Annotation,
Climate, Ecology & More!



## Thanks to Fidele & Prof.Ludwig-Müller

Thanks for attention & Happy to take questions?

# Setting up your galaxy account

### Setting up Galaxy account

Go to the "Login or Register" link at the top of Galaxy interface and choose "Register" (unless of course you already have an account):



[19/10/2021] EBI is currently facing some issues with their FTP server and we are experiencing network failures. They are looking into it and will fix it asap. Please retry later if possible.

Welcome to Galaxy, please log in	
Public Name or Email Address	
Password	
Forgot password? Click here to reset your password.  Login	
elXi* LDGIN	
Don't have an account? Register here.	

#### COVID-19 Research!

Want to learn the best practices for the analysis of SARS-CoV-2 data using Galaxy? Visit the Galaxy SARS-CoV-2 portal. We mirror all public SARS-CoV-2 data from ENA in a Galaxy data library for your convenience. The Galaxy community has created COVID-19 dedicated training materials. Please check our recent activities for more details.

If you need help submitting your data to public archives, like ENA, please get in touch. We will support you in sharing your data.

#### This is the Indian galaxy server, Welcome Home!

Namaste All! Welcome to the India Galaxy workbench; a comprehensive set of tools and workflows dedicated to accelerate your bioinformatics analyses focusing Bharat. This workbench is built on the Galaxy framework thanks to Galaxy Europe we are hosting the seeds of Indian Galaxy to facilitate more wider user base.

To know more about the events, training webinars focused on Galaxy for Indian data analysis community visit wwww.galaxyproject.in. If you want to talk about Galaxy and any required help about it join the gitter adda (अड्डा).

1. Training

Please register only one account - we provide this service free of charge and have limited computational resources. Multi-accounts are tracked and will be subjected to account termination and data deletion.

Create a Galaxy account
Email Address
Password
Confirm password
Public name
Your public name is an identifier that will be used to generate addresses for information you share publicly. Public names must be at least three characters in length and contain only lower-case letters, numbers, dots, underscores, and dashes (``, `_', `-').  Create
Already have an account? Log in here.

How to visualize the expression of the genes across the samples?

## Heatmaps

- Commonly used to visualize differentially expressed genes
- Heatmap2 tool in Galaxy
- Uses the heatmap.2 function from the R gplots package
- Study from <u>Fu et al. (2015)</u>

## Dataset

- Normalized Counts file
- Differentially expressed results file
- Genes of interest

# Let's go for demonstration!

# Thank you

Any auestion?

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# Extra Slides