Spark Meeting Galaxy-Server



Dr. Kristin Reiche, Alexander Scholz Fraunhofer IZI, Leipzig

Prof. Dr. Ezio Bonifacio, Dr. Juliana Roscito CRTD / TU Dresden

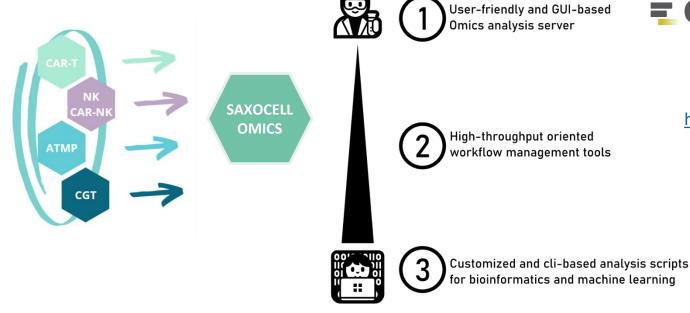
Dr. Jan Ewald, Max Joas ScaDS.AI / Uni Leipzig

Dr. David Langenberger, Dr. Mario Fasold ecSeq Bioinformatics, Leipzig



OMICS — Support for data analyses







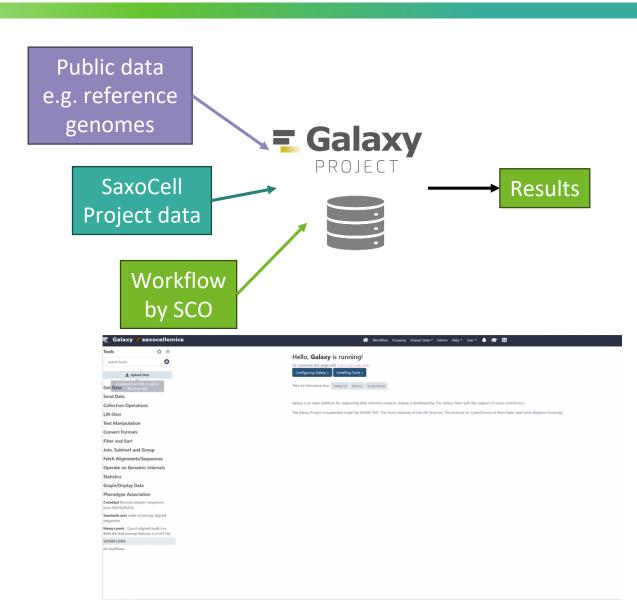
https://service.scadsai.uni-leipzig.de/saxocellomics/galaxy-dev

Galaxy project



Galaxy is an **open-source** and **community-driven** platform for **FAIR** data analysis that enables users to:

- Use tools from various domains that can be plugged into workflows
- ... via a graphical web interface.
- **Share** data, workflows and results
- Ensure reproducibility by standardized and documented workflows.
- Use either public or self-hosted Galaxy analysis server

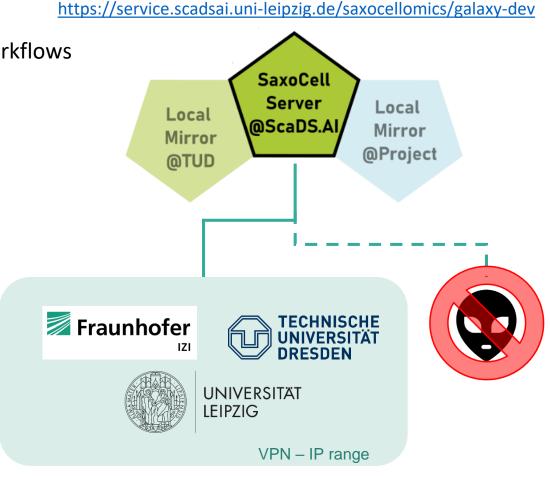


Advantages for SaxoCell



Galaxy server @ScaDS.AI (Leipzig University)

- Full control and security over your (sensitive) data and workflows
- **Scalability** (currently reduced resources in first phase)
 - Depending on demand: more resources at ScaDS.Al server
 - Easy roll-out on your infrastructure
- SCO will establish standardized workflows.
- You can decide:
 - Perform your own analysis or by SCO team
 - Create and tweak your own workflow or use existing workflows
 - Share or not share data, workflows and results with other users



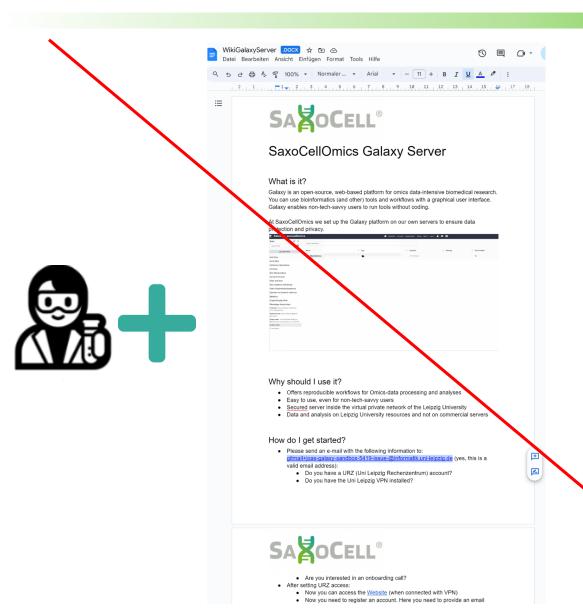
FAQ: Modus Operandi





FAQ: Introduction









FAQ: Getting Started



- Do you have access to:
 - IZI VPN 📀
 - Leipzig University VPN
 - TU Dresden VPN
- Contact us
 - gitmail+joas-galaxy-sandbox-5419-issue-@informatik.uni-leipzig.de

FAQ: Sign up



- Use your name:
 - Max_Joas
 - World_Best_Scientist777
- No annoying mail verification
- Let's see for yourself:
 - https://service.scadsai.uni-leipzig.de/saxocellomics/galaxy-dev/

FAQ: Running Analysis



- What can I do?
 - Almost everything



- RNA-seq: raw -> pathways
- Genome annotation
- Chip seq analysis
-
- Tools: https://toolshed.g2.bx.psu.edu
- Workflows: https://usegalaxy.org/workflows/list-published (#1320)
- What are the limitations?



- 80 GB user quote
- RAM
- Concurrent use
- Can I use Galaxy as a team?



share data, workflows and history

FAQ: Privacy & Security



- Server hosted by Leipzig University
- Secured by VPN (only accessible within)
- Analysis server:
 - no data backup
 - user is master of data (delete)
 - workflows backed-up
- Downtime possible
 - users notified
- Roll-out on your servers



FAQ: Getting Help

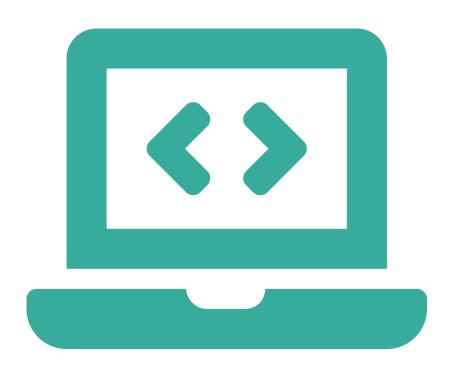


There are multiple channels where you can get help:

- contact us
- galaxy training https://training.galaxyproject.org
- galaxy community help: https://help.galaxyproject.org
- ask a question on https://www.biostars.org
- ask a question on https://stackoverflow.com
- check out our FAQ <u>https://docs.google.com/document/d/1IsHne_fxngYHVRDYRYxrE_e96ZrJcss8/edit?usp=sharing&ouid=115296162071832875425&rtpof=true&sd=true</u>

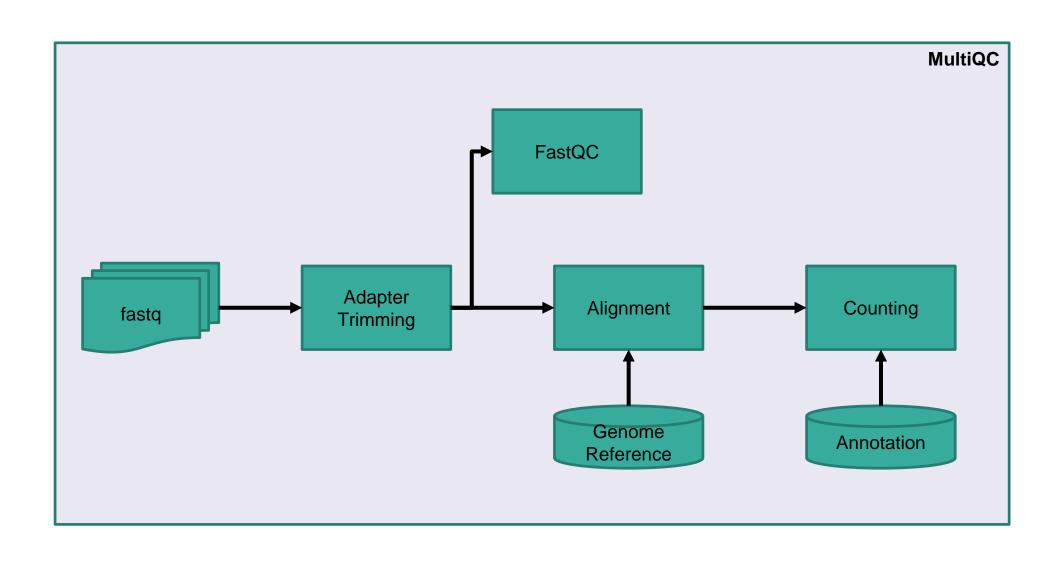
Hands-on





Hands-on





CONTACT US





ezio.bonifacio@tu-dresden.de juliana.roscito@tu-dresden.de





jan.ewald@uni-leipzig.de joas@informatik.uni-leipzig.de



kristin.reiche@izi.fraunhofer.de ulrike.weirauch@izi.fraunhofer.de alexander.scholz@izi.fraunhofer.de



david.langenberger@ecseq.com mario.fasold@ecseq.com