

OSCARS

Open Science Clusters' Action
for Research & Society

Funded Project

FAIRY: FAIRification of YEASTRACT+ to Support Yeast-based Bioeconomy and Health

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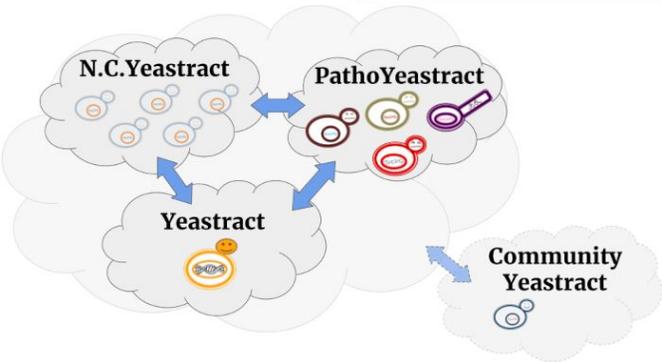
Implemented by



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What problem(s) are you going to solve?

- YEASTRACT+ is a leading resource for yeast transcriptional regulation, but currently **lacks quantitative gene expression data**
 - Many yeast **species remain poorly characterised**, impeding the understanding of their regulatory mechanisms and broader bioeconomic applications
 - YEASTRACT+ currently does not provide a **programmatic API** for data access
 - Current **data** is only **linked** to a few reference databases (e.g. SGD, GO)
 - YEASTRACT+ is not fully compliant with **FAIR principles**
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What are you planning to do to solve the problem?

- Rely on EOSC cloud computing for **raw data curation** (RNA-seq data processing)
 - Enrich regulatory information with **quantitative data**
 - Develop a new **responsive portal** leveraging new curated data
 - With **programmatic access API** exposing data and functionalities
 - Train machine learning models on cross-species regulatory data to:
 - **Assess data quality** (One class classification outlier detection)
 - **Infer** transcriptional **regulatory associations** in poorly-characterised yeasts
 - Update and **Expand the body of data** gathered in the database, expanding it to further yeast species
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What will be the results and how do you plan to make them available to the broader community?

FINAL PRODUCT: An expanded, quantitative and fully FAIR YEASTRACT+ database

Results will be made available through:

- Data and functionalities through our website (<https://yeastract-plus.org/>)
 - Newly curated data available through API + flat files for bulk processing
 - Publication of the upgraded YEASTRACT+ in the NAR Database Issue
 - Participation in bioinformatics, systems biology and yeast biology scientific meetings
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Who is doing it?

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