

## Understanding Robustness of *Yarrowia lipolytica* for Undetoxified Biomass Hydrolysate Utilization

Seunghyun Ryu<sup>1\*</sup> ([sryu2@utk.edu](mailto:sryu2@utk.edu)), Caleb Walker,<sup>1</sup> Stephanie Thompson<sup>2</sup>, Richard J. Giannone<sup>3</sup>, Patricia J. Slininger<sup>2</sup>, Bruce S. Dien<sup>2\*</sup>, ([bruce.dien@usda.gov](mailto:bruce.dien@usda.gov)) and Cong T. Trinh<sup>1\*</sup> ([ctrinh@utk.edu](mailto:ctrinh@utk.edu))

<sup>1</sup>University of Tennessee, Knoxville; <sup>2</sup>National Center for Agricultural Utilization Research, USDA-ARS, Peoria, IL; <sup>3</sup>Oak Ridge National Laboratory, Oak Ridge, TN;

**Project Goals:** To elucidate and harness the exceptional robustness of novel, undomesticated *Y. lipolytica* isolates from genetic diversity screening as a bioenergy-relevant microbial platform for efficient conversion of undetoxified biomass hydrolysates into designer bioesters with continuous recovery using solvent extraction.

**Abstract text.** Robustness is an important phenotype for bioenergy microbes to acquire but is difficult to engineer. The oleaginous yeast, *Yarrowia lipolytica*, is an exceptionally robust microbe that can naturally tolerate stressful environments, assimilate a wide range of substrates, and produce high-value chemicals. In this study, we aim to understand and harness the robust characteristics of *Y. lipolytica* for utilizing mixed C5 and C6 sugars in undetoxified switchgrass biomass hydrolysates. From a screen of 57 undomesticated *Y. lipolytica* isolates, we selected top-performing strains exhibiting robust growth and lipid accumulation in biomass hydrolysates and subjected them to comprehensive growth, lipid production, and multi-omics characterization. We then engineered these top performing strains through a combination of targeted pathway engineering followed by adaptive laboratory evolution to further enhance their inherent robustness. Multi-omics profiling of these robust *Yarrowia* strains revealed novel genotype, regulation and cellular processes that positively impact their utilization of biomass hydrolysate sugars. These foundational studies provide key insight into the genetic mechanisms responsible for robustness in *Yarrowia* strains, enabling targeted engineering strategies to be deployed for enhanced production of biofuels and bioproducts from lignocellulosic biomass.

### References

- 1 Walker, C., Dien, B., Giannone, R.J., Slininger, P., Thompson, S.R., **Trinh, C.T.\***, 2021. Exploring Proteomes of Robust *Yarrowia lipolytica* Isolates Cultivated in Biomass Hydrolysate Reveal Key Processes Impacting Mixed Sugar Utilization, Lipid Accumulation, and Degradation, *mSystems*, **6**(4): e00443-21.
2. Walker, C., Ryu, S. & Trinh, C. T. Exceptional solvent tolerance in *Yarrowia lipolytica* is enhanced by sterols. *Metabolic Engineering* **54**, 83-95, doi:<https://doi.org/10.1016/j.ymben.2019.03.003> (2019).

3. Ryu, S. & Trinh, C. T. Understanding Functional Roles of Native Pentose-Specific Transporters for Activating Dormant Pentose Metabolism in *Yarrowia lipolytica*. *Applied and Environmental Microbiology* **84**, doi:10.1128/aem.02146-17 (2018).
4. Walker, C. *et al.* Draft Genome Assemblies of Five Robust *Yarrowia lipolytica* Strains Exhibiting High Lipid Production, Pentose Sugar Utilization, and Sugar Alcohol Secretion from Undetoxified Lignocellulosic Biomass Hydrolysates. *Microbiol Res Announc* **7**, e01040-01018 (2018).
5. Quarterman, J., Slininger, P. J., Kurtzman, C. P., Thompson, S. R. & Dien, B. S. A survey of yeast from the *Yarrowia* clade for lipid production in dilute acid pretreated lignocellulosic biomass hydrolysate. *Applied microbiology and biotechnology* **101**, 3319-3334, doi:10.1007/s00253-016-8062-y (2017).
6. Ryu, S., Labbé, N. & Trinh, C. T. Simultaneous saccharification and fermentation of cellulose in ionic liquid for efficient production of  $\alpha$ -ketoglutaric acid by *Yarrowia lipolytica*. *Applied microbiology and biotechnology* **99**, 4237-4244, doi:10.1007/s00253-015-6521-5 (2015).
7. Ryu, S., Hipp, J. & Trinh, C. T. Activating and Elucidating Metabolism of Complex Sugars in *Yarrowia lipolytica*. *Applied and Environmental Microbiology* **82**, 1334-1345, doi:10.1128/aem.03582-15 (2016).

*Funding statement.* This work is supported by the U.S. Department of Energy, Office of Science, Office of Biological and Environmental Research, Genomic Science Program under Award Number DESC0019412.