

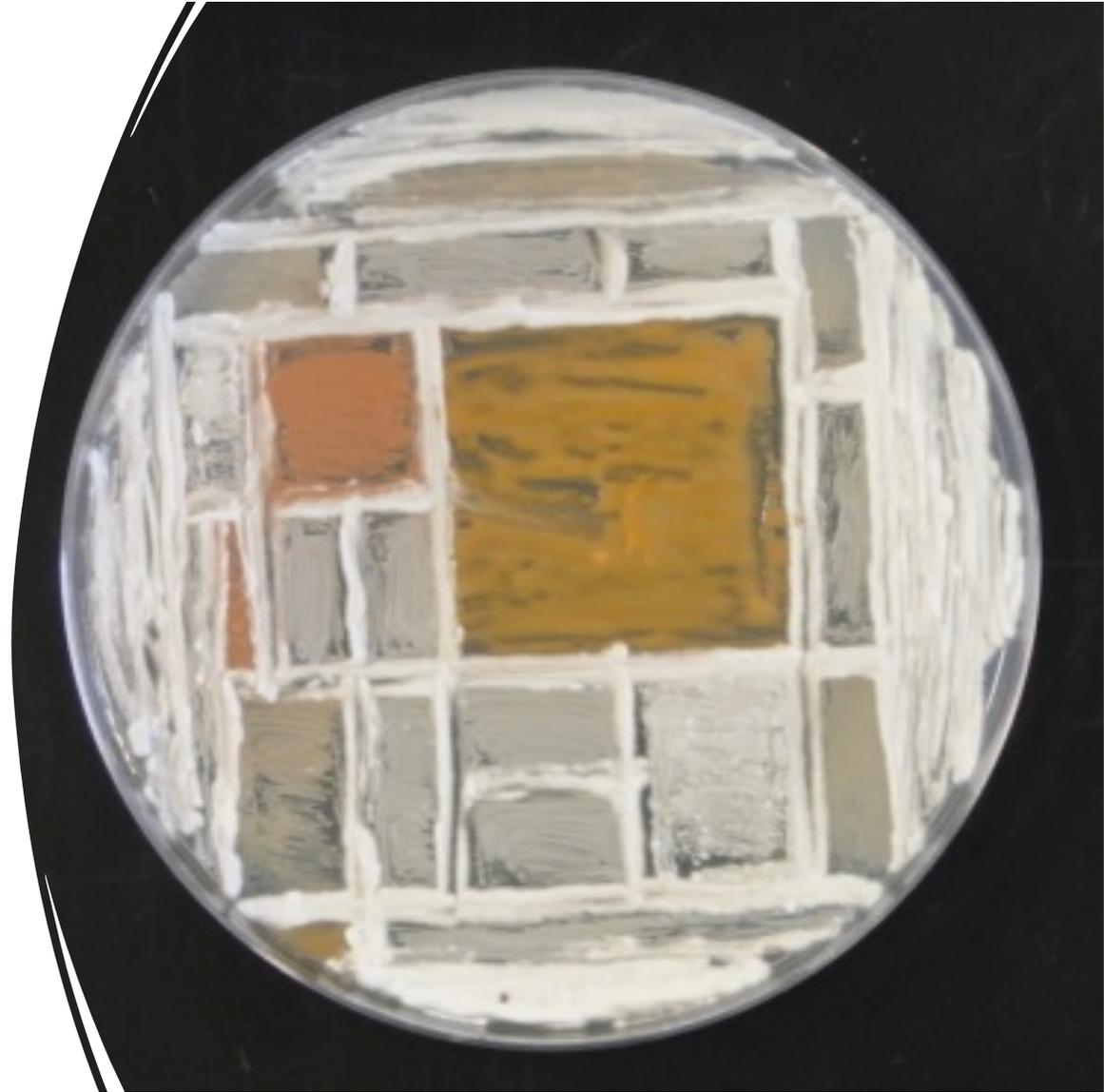
Three essential collection elements: Microbes, data, and expertise

Kyria Boundy-Mills

Curator, Phaff Yeast Culture Collection

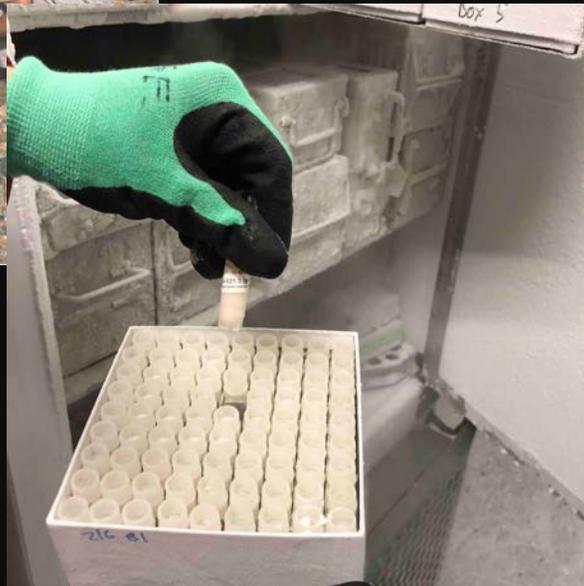
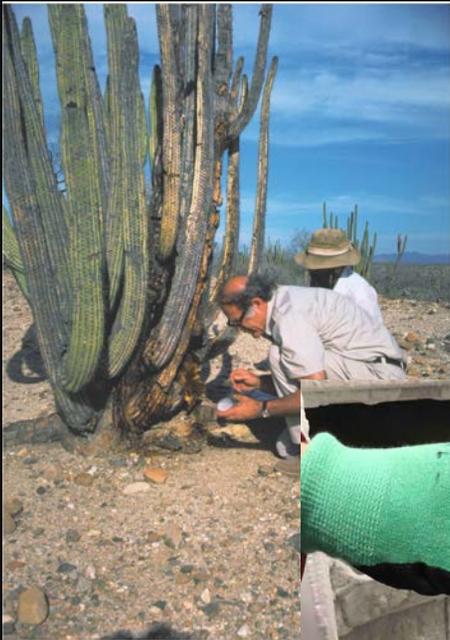
University of California
Davis

WDCM 2 Nov 2023 Shenzhen



Phaff Yeast Culture Collection

- 9,000 strains; over 1,500 species
- Ecology research
 - Food, plants, animals, other
- 1 to 800 strains per species (independent isolates)
- Wild, non-GMO
- Phenotype database
- In GCM, and our own catalog:
<https://phaffcollection.ucdavis.edu>





1. MICROBES



**Currently adding 5,000 yeast strains from
Dr. Marc-André Lachance**

Criteria for selecting:

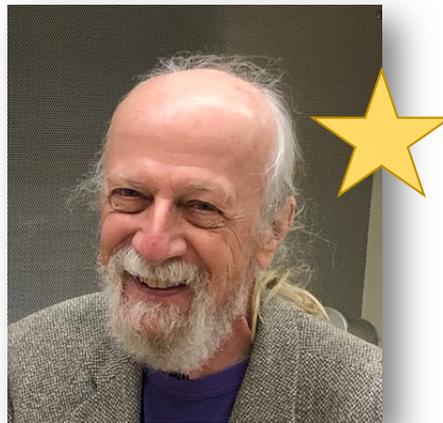
- Important discoveries
 - Cited in publications
 - Type strain of new species
- Investment in data
 - Genome sequenced; Phenotype data
- Immediate access needed
- Costly or impossible to replace
 - “Moon rocks”
 - Classical mutations, habitat destroyed
- Complements holdings
 - Genetic, geographic, temporal, phenotype diversity
- Few or no limitations on use

Adapted from: Flattau, P. E., M. Boeckmann, R. d. I. Cruz, P. Lagasse, N. Mitchell, M. Patterson, and D. Singpurwalla. 2007. Scientific collections: Mission-critical infrastructure for federal scientific agencies. Science and Technology Policy Institute.

Tips for researchers

2019 publication by USCCN

- Organize specimens
 - Cull duplicates; assign unique ID
- Database
 - Source, genotype, phenotype, documentation
 - Share database
- Preserve
- Long-term plan
 - Future home; funding



Boundy-Mills, K., K. McCluskey, P. Elia, J. A. Glaeser, D. L. Lindner, D. R. Nobles Jr, J. Normanly, F. M. Ochoa-Corona, J. A. Scott, T. J. Ward, K. M. Webb, K. Webster and J. E. Wertz (2020). "Preserving US microbe collections sparks future discoveries." *Journal of Applied Microbiology* 129(2): 162-174.



2. DATA: Lachance data sheets

The image displays a series of handwritten data sheets, likely from a laboratory notebook or a specialized data collection form. The sheets are densely packed with text and tables, covering a wide range of topics related to microbial strains. The data is organized into columns and rows, with various headings and sub-headings. The handwriting is clear and legible, indicating a systematic approach to data collection. The sheets are numbered, and the overall layout suggests a comprehensive record of experimental results and observations.

Over 5,000 strains

- Assimilation of C, N compounds
- Fermentation of C compounds
- Killer activity
- Enzymes
- Morphology

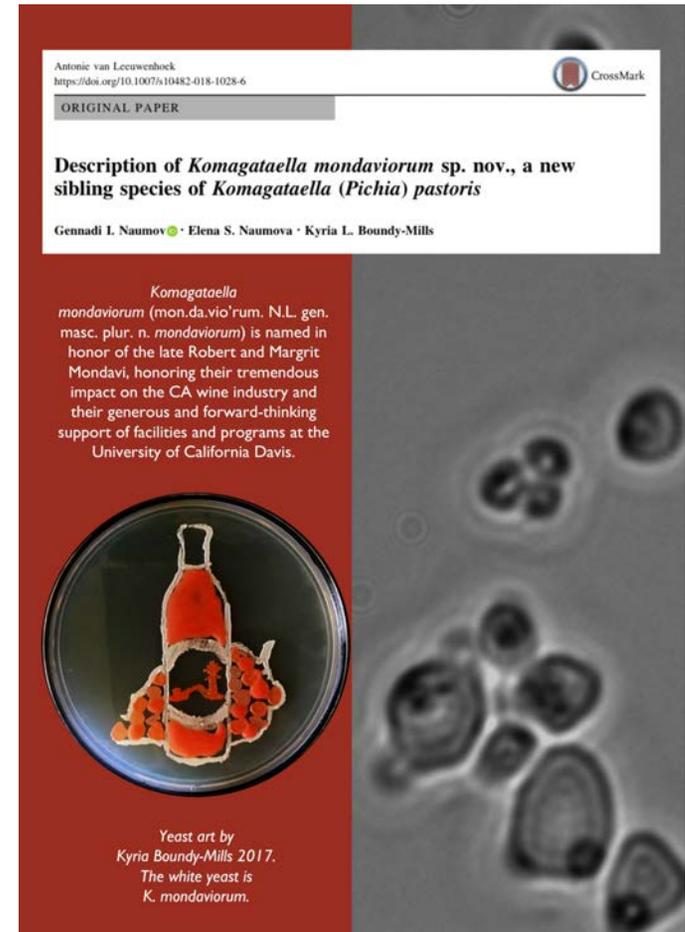
Stress tolerance:

- High, low temperatures
- High, low pH
- High sugar, salt; ethanol tolerance
- Growth inhibitors



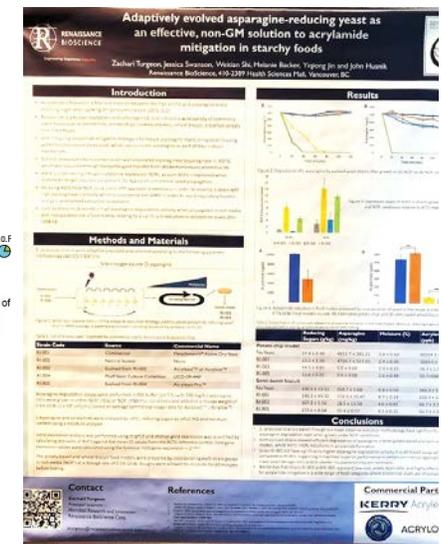
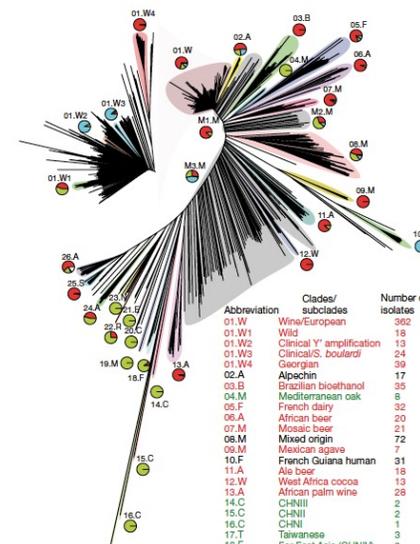
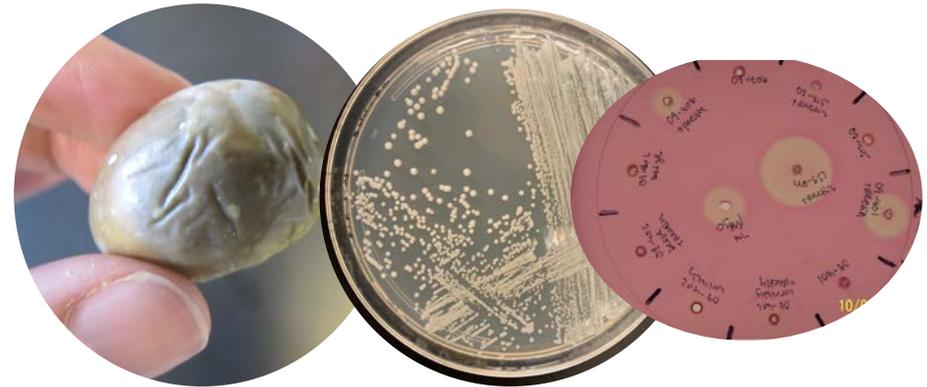
3. EXPERTISE of collection personnel

- SOURCE:
 - Acid tolerance study → isolated from low pH substrates
- TRADITIONAL PHENOTYPE DATA:
 - Ethanol tolerance study → ethanol tolerance data
- RECENT DATA:
 - New species descriptions → ribosomal sequences
 - *Komagataella mondaviorum*
 - *Rasporella diana*
- CONNECTING RESEARCHERS:
 - Asparaginase, acrylamide story



Many uses of one yeast strain: *Saccharomyces cerevisiae* UCDFST 09-448

- 2009: Caused spoilage of olives – excessive pectinase activity
- 2011: Sent to researchers in France who sequenced genomes of 1,011 *S. cerevisiae* strains
 - Published in Nature in 2018
- 2020: Renaissance Biosciences found interesting asparaginase gene in a few *S. cerevisiae* strains including 09-448
 - Asparagine in foods can become acrylamide; Asparaginase before cooking reduces acrylamide in product
 - Renaissance used adaptive evolution to improve the strain
 - In 2022, Renaissance produced 2,200 kg of dried yeast for food companies to put into baked and fried foods to reduce acrylamide



REQUEST: Can I see your collection's Disaster Plan?

MICROBES, DATA:

- Fire, flood, earthquake
- War
- Pandemic
- Freezer malfunction
- Computer crash

PERSONNEL:

- Illness, death
- Retirement
- Funding cuts

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