

Resource Summary Report

Generated by [ASWG](#) on Aug 2, 2026

Phaff Yeast Culture Collection

RRID:SCR_016781

Type: Tool

Proper Citation

Phaff Yeast Culture Collection (RRID:SCR_016781)

Resource Information

URL: <http://phaffcollection.ucdavis.edu/>

Proper Citation: Phaff Yeast Culture Collection (RRID:SCR_016781)

Description: Collection of wild yeast in the world. Academic and industrial researchers utilize this collection for a variety of research and applications, including pigments, enzymes, food ingredients, and hosts for protein expression. Provides contract screening and strain selection services.

Resource Type: data or information resource, portal, database, service resource, organization portal

Keywords: yeast, strain, wild, screening, selection, service

Availability: Commercially available

Resource Name: Phaff Yeast Culture Collection

Resource ID: SCR_016781

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260801T190740+0000

Ratings and Alerts

No rating or validation information has been found for Phaff Yeast Culture Collection.

No alerts have been found for Phaff Yeast Culture Collection.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [ASWG](#) .

Gutierrez-Cano AR, et al. (2024) Characterization of pectinase-producing *Saccharomyces cerevisiae* UCDFST 09-448 and its effects on cull peach fermentations. *Journal of industrial microbiology & biotechnology*, 51.

Tanahashi R, et al. (2023) Isolation of Yeast Strains with Higher Proline Uptake and Their Applications to Beer Fermentation. *Journal of fungi (Basel, Switzerland)*, 9(12).

Wang K, et al. (2021) A novel *Arabidopsis* phyllosphere resident *Protomyces* species and a re-examination of genus *Protomyces* based on genome sequence data. *IMA fungus*, 12(1), 8.

Boundy-Mills K, et al. (2020) Preserving US microbe collections sparks future discoveries. *Journal of applied microbiology*, 129(2), 162.