# The 1st International Online Conference on Fermentation

MDPI

12-13 November 2025 | Online

# Indigenous yeast biodiversity revealed by bioprospecting methods in South West Switzerland

Yannick Barth <sup>1</sup>, Corentin Descombes <sup>1</sup>, Benoît Bach <sup>2</sup> and François Lefort <sup>1</sup>

types (left to right)

Vine bark

Whole berries

Flowers Insects

- <sup>1</sup> Plants and Pathogens Group, Research Institute INTNP, HES-SO University of Applied Sciences and Arts Western Switzerland, Genève, Switzerland
- <sup>2</sup> Changins, HES-SO University of Applied Sciences and Arts Western Switzerland, Nyon, Switzerland;

#### **INTRODUCTION & AIM**

Yeasts colonize various natural and anthropogenic habitats, notably vineyards and wineries. There is a growing interest in indigenous yeast starters, reflective of the local yeast biodiversity and potentially adapted to cultivated grape varieties. Native yeasts participate to the microorganism's diversity of distinct niches in the vineyards and wineries and could be linked to the notion of "terroir". They could be a unique microbial signature of the "terroir" and therefore an invaluable resource for winemakers.

The present bioprospection reported here covered **8 vineyards and their associated wineries** across 4 Swiss cantons (Geneve, Vaud, Neuchâtel, and Valais) in South-Western Switzerland.



Figure 1: Sampled vineyards and wineries Geneva (red) Vaud (blue) Neuchâtel (green) Valais (violet)

#### **METHOD**

#### **Collection of samples**

Saccharomyces yeasts were collected from winery equipment and surfaces using sterile swabs, with five samples per area. In the vineyard, non-Saccharomyces yeasts were sampled from vine bark, flowers, insects, and berries, with five samples per type in each parcel.

Figure 2: Samples

## Selective liquid medias

Samples were inoculated into YPD medium containing chloramphenicol to inhibit bacteria and ethanol to select for yeasts (3% ethanol for vineyard, 6% for winery) and incubated at 22°C for 48 hours before plating.

#### Isolation of colonies

Cultures were diluted and plated onto WLN agar. Colonies were selected according to color, shape, and growth characteristics. Morphologically distinct colonies were subcultured to obtain pure cultures.

#### Genetic identification by PCR

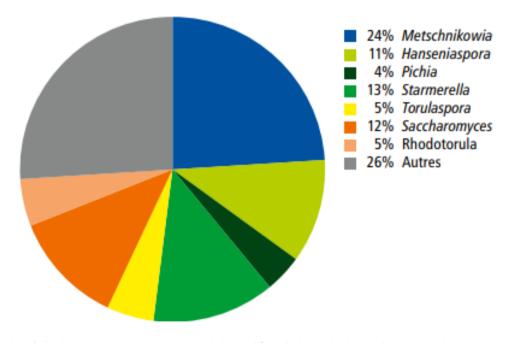
After DNA extraction, the D1/D2 domain of 26S rDNA was amplified by PCR, purified and Sanger sequenced. Species identity was obtained by sequence comparison to the GenBank database.

#### **RESULTS & DISCUSSION**

A total of 221 yeast isolates were genetically identified, encompassing 25 genera and 45 species.

**Table 1:** Yeast occurrence by viticultural domain

		Wine estate								
Genus	Yeast species	République (GE)	Chambet (GE)	Gros (GE)	Bonvillars (VD)	Brunner (NE)	Auvernier (NE)	Cornulus (VS)	Grand Brûlé (VS)	Total
Aureobasidium	Aureobasidium pullulans	0	0	2	0	0	0	0	0	2
Candida	Candida boidinii	0	0	0	0	0	1	0	0	1
	Candida friedrichii	0	0	0	1	0	0	0	0	1
	Candida norvegica	0	0	0	0	0	1	0	0	1
	Candida pseudofarinosa	0	0	0	0	1	1	0	0	2
	Candida tsuchiyae	0	0	0	1	0	0	0	0	1
Citeromyces	Citeromyces matritensis	0	0	0	1	0	0	0	0	1
Curvibasidium	Curvibasidium cygneicollum	0	0	2	0	4	0	0	0	6
Cystobasidium	Cystobasidium pinicola	0	0	0	0	0	0	1	2	3
Debaryomyces	Debaryomyces hansenii	1	0	0	0	0	0	0	0	1
Filobasidium	Filobasidium magnum	0	1	0	0	0	0	0	0	1
Hanseniaspora	Hanseniaspora osmophila	0	0	0	0	0	1	0	0	1
	Hanseniaspora uvarum	4	3	8	5	0	6	0	0	26
Hyphopichia	Hyphopichia burtonii	0	0	1	0	0	0	0	0	1
Kazachstania	Kazachstania servazii	3	0	0	0	0	0	0	0	3
Kwoniella	Kwoniella europaea	0	0	0	0	1	0	0	0	1
Lachancea	Lachancea thermotolerans	0	0	2	0	0	0	1	0	3
Metschnikowia	Metschnikowia fructicola	2	0	1	2	0	0	2	0	7
	Metschnikowia gruessii	1	0	1	0	0	0	0	0	2
	Metschnikowia koreensis	0	0	1	0	0	0	0	0	1
	Metschnikowia pulcherrima	9	0	0	3	0	0	4	1	17
	Metschnikowia reukaufii	7	5	3	2	1	0	2	1 1	21
	Metschnikowia viticola	0	4	0	0	0	1	0	0	5
Meyerozyma	Meyerozyma caribbica	Ö	0	2	0	1	0	0	0	3
	Meyerozyma guilliermondii	0	1	1	0	0	0	0	0	2
Papiliotrema	Papiliotrema terrestris	ő	0	0	0	0	0	0	1	1
Pichia	Pichia kluyveri	ő	1	0	0	0	0	0	0	1
	Pichia manshurica	0	0	0	0	0	1	0	0	1
	Pichia membanifaciens	ő	1	0	3	0	2	3	0	9
Rhodosporidium	Rhodosporidium babjevae	ő	1	0	0	0	0	0	0	1
· ·	Rhodotorula graminis	2	0	1	0	0	1	0	0	4
Rhodotorula	Rhodotorula mucilaginosa	1	0	0	0	0	0	0	0	1
Saccharomyces	Saccharomyces cerevisiae	4	2	0	11	1	13	1	0	32
Solicoccozyma	Solicoccozyma aeria	1	0	0	0	0	0	0	0	1
Sporidiobolus	Sporidiobolus pararoseus	0	0	5	0	0	0	0	0	5
Starmerella	Starmerella apicola	o o	0	0	1	0	0	0	1	2
	Starmerella bombi	2	1	0	0	0	0	1	1	5
	Starmerella bombicola	4	4	0	1	1	2	2	4	18
	Starmerella orientalis	1 7	0	0	0	0	0	1	0	2
	Starmerella sorbosivorans	0	0	0	0	0	0	2	0	2
	Starmerella vitae	1	0	0	0	0	0	0	0	1
Torulaspora	Torulaspora delbrueckii	- 2	0	0	1	0	2	1	4	10
Vishniacozyma	Vishniacozyma carnescens	0	1	0	0	0	0	0	0	1
	Vishniacozyma carnescens Vishniacozyma victoriae	0	1	1	0	0	0	0	0	2
<i>Vickerhamomyces</i>		0	4	0	0	0	3	2	0	9
, , , , , , , , , , , , , , , , , , , ,	Wickerhamomyces anomalus	45	4		32				-	9
25 genera	45 species	45	30	31	32	10	35	23	15	1



**Figure 3:** Main yeast genera identified in eight vineyards and wineries of Western Switzerland

#### **CONCLUSION**

This approach revealed a rich diversity of indigenous yeasts with a potential to enhance and diversify wine flavors. Some of the non-Saccharomyces species found here are already used commercially, highlighting the potential of these yeasts as valuable resources for winemaking.

### FUTURE WORK / REFERENCES

The *Metschnikowia* subset of this yeast collection has undergone characterization in co-fermentations with *S. cerevisiae* (doi:10.3390/foods12244485). Characterization studies, including micro-fermentations alongside chemical and sensory analyses, will be extended to additional species. Another subset of has been evaluated as biocontrol agents against *Botrytis cinerea*, some yeasts showed a very strong effect against.