

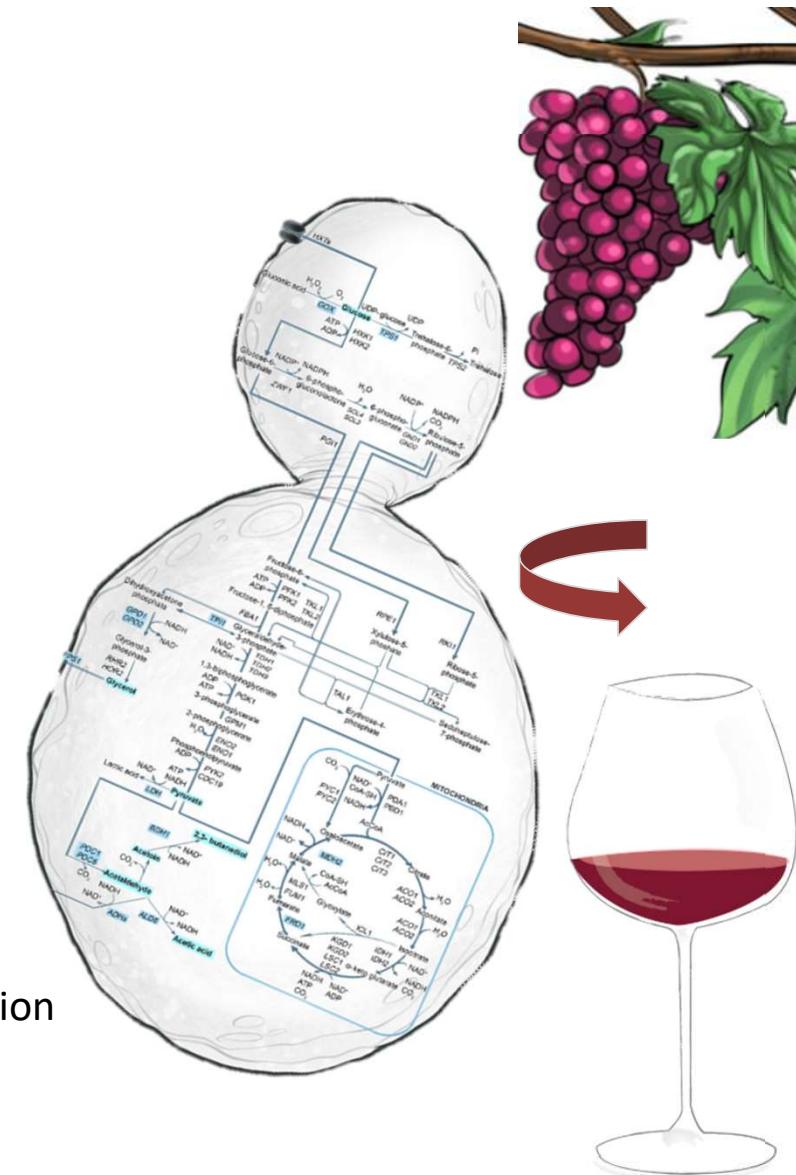


Yeasts from (organic) grapes and wines: Biodiversity and how to manage un-inoculated fermentations

Ana Hranilovic, Marina Bely, Vladimir Jiranek, Warren Albertin,
Isabelle Masneuf-Pomarede

Yeasts in winemaking

- Role:
 - alcoholic fermentation (AF)
 - conversion of sugars to ethanol and CO₂
 - production/modulation of flavour-active compounds
- Species:
 - *Saccharomyces cerevisiae*
 - ‘wine yeast’ capable of completing AF
 - rare in the vineyards (grape 1/1000)
 - *Non-Saccharomyces*
 - different yeasts that occur on grapes/in cellars
 - metabolic contribution to wine profiles
 - e.g.(de)acidification and enhanced aroma production



Examples of yeasts from grapes and wines

Wine Server, UC Davis

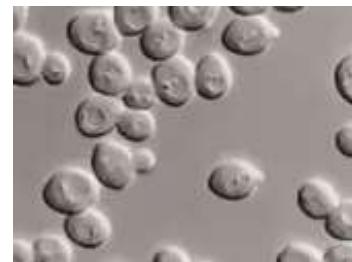
Hanseniaspora uvarum

- common on grapes



Torulaspora delbrueckii

- aroma/flavour enhancement



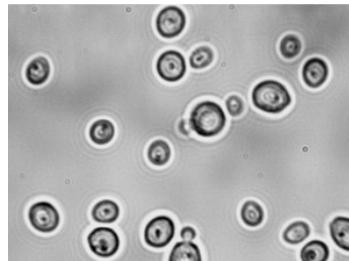
Candida zemplinina

- fructophilic character



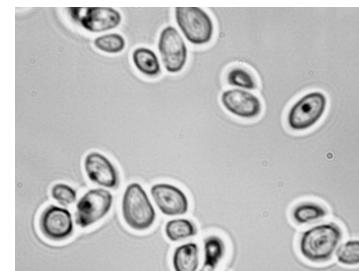
Lachancea thermotolerans

- lactic acid production
- lower ethanol production



Metschnikowia pulcherrima

- lower ethanol production
- bioprotection



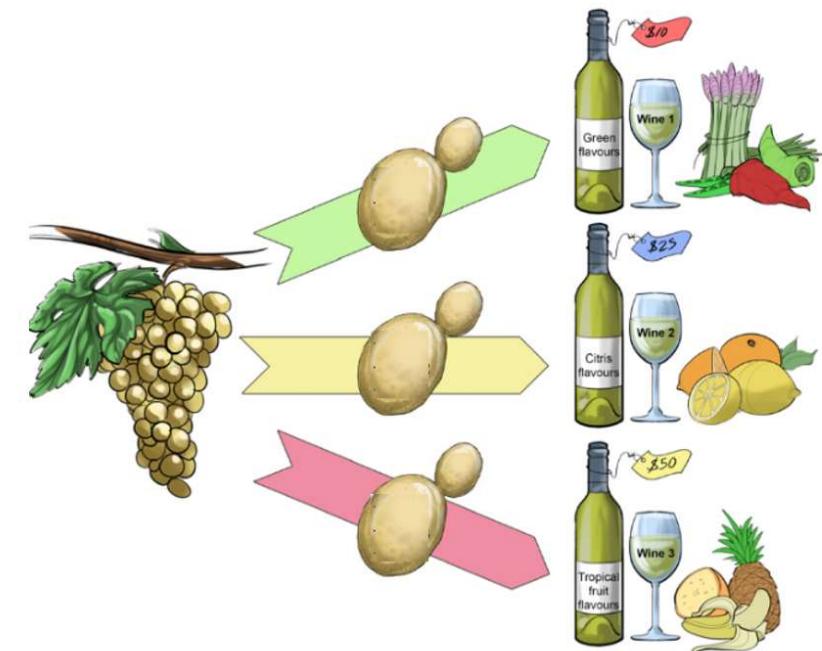
Schizosaccharomyces pombe

- malic acid degradation



Different fermentation modalities – pros and cons

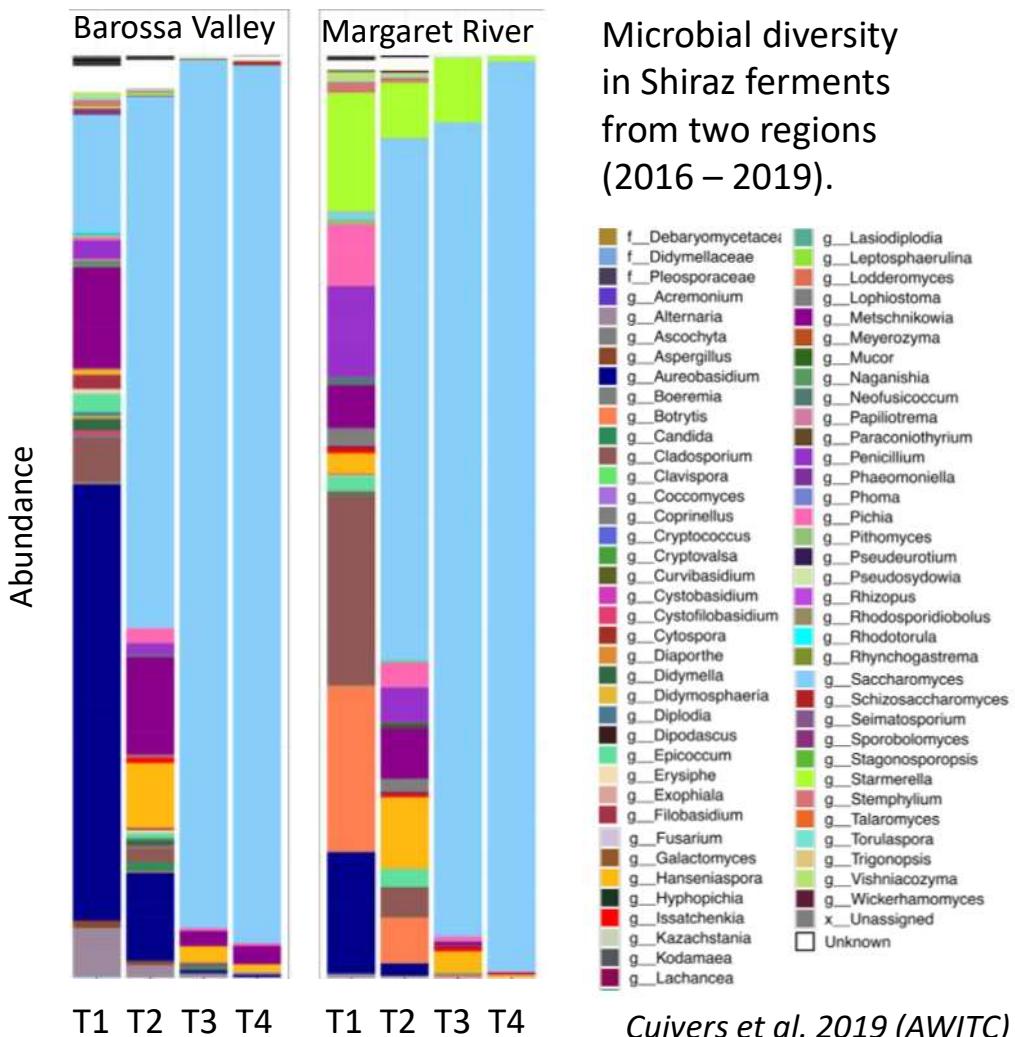
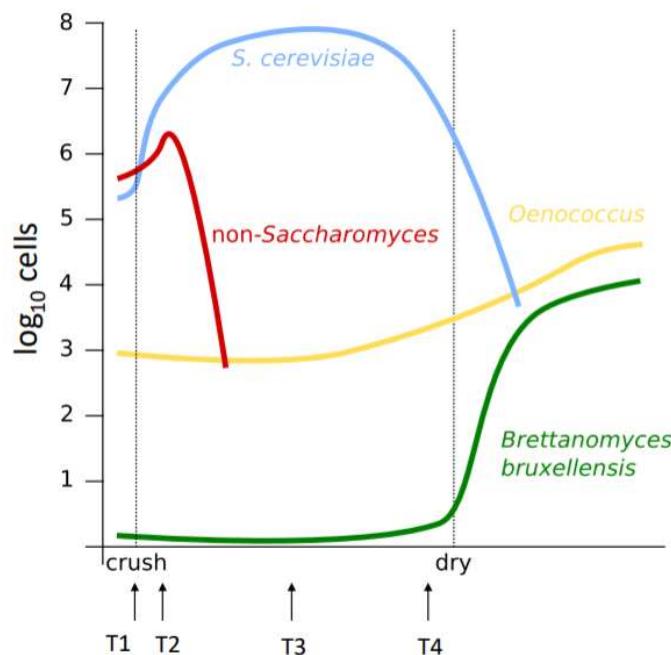
- **Inoculated fermentations with *S. cerevisiae***
 - timely and reliable fermentation
 - costs and wine ‘uniformity’
- **Multistarter fermentations with non-*Saccharomyces* yeasts**
 - modulation of wine chemical and sensory profiles
 - costs and management
- **Un-inoculated fermentations**
 - ‘microbial terroir’ and stylistic distinctness
 - lower inputs (ADY cost, refrigeration, nutrients)
 - marketing opportunities (organic/natural/... wines)
 - risks of stuck/sluggish fermentations and erratic results



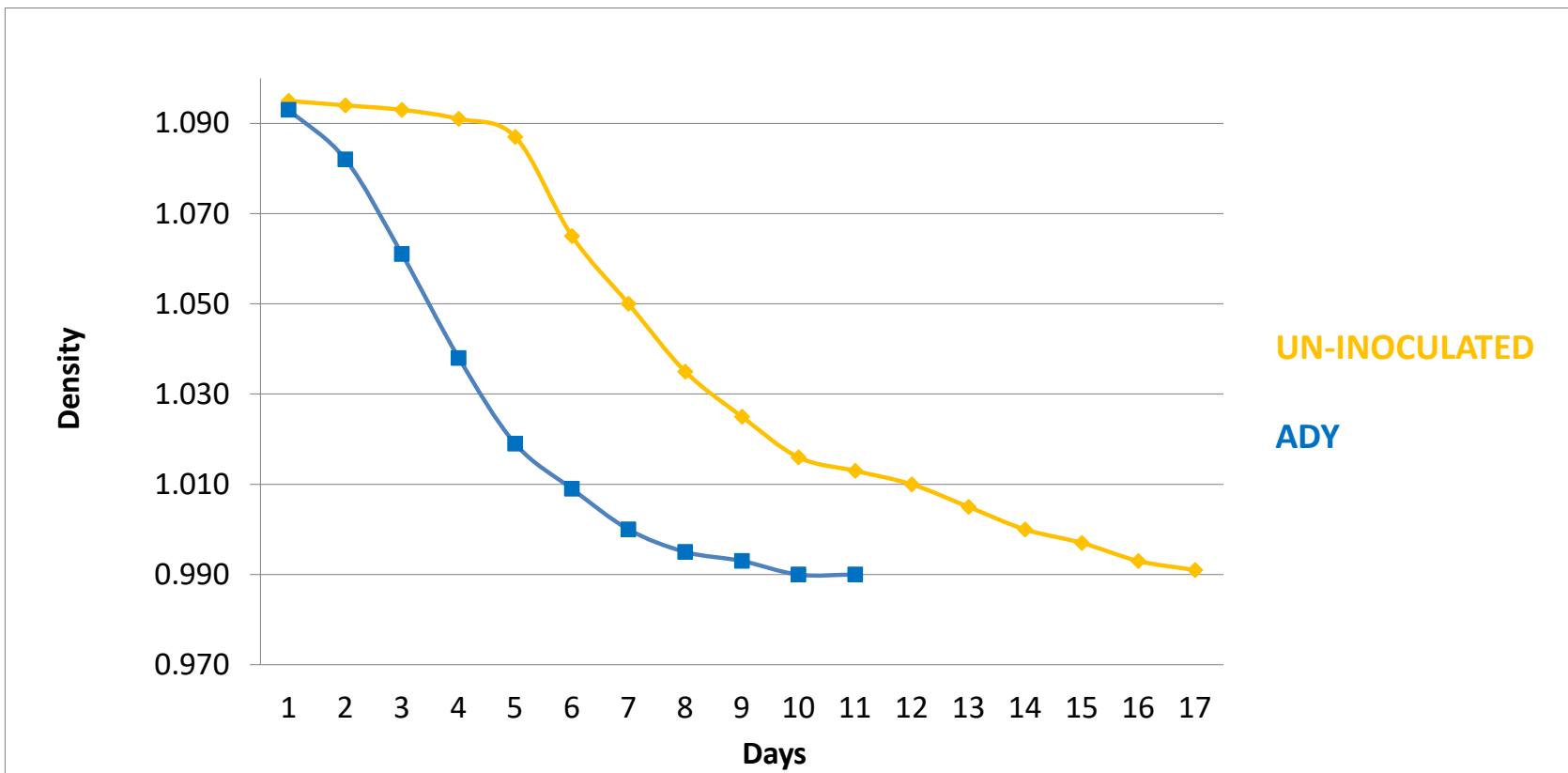
Population dynamics in un-inoculated fermentations

- succession of species:

- non-*Saccharomyces* at early stages taken over by *S. cerevisiae*
- variable composition and dynamics



Managing fermentations with indigenous yeasts

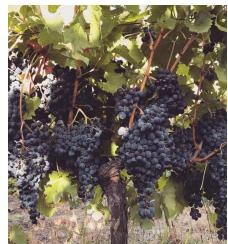


Comparison of fermentation dynamics with indigenous yeasts and ADY.

Pied de cuve technology

1) at the beginning of harvest: from grapes

- collecting healthy ripe grapes 8-10 days before harvest (~150 kg for 50 hL)
- pressing the grapes
- 30 ppm SO₂,
- monitoring AF; 20-25 ° C
- tasting at mid-fermentation
- use pied de cuve; 3-5% of the tank



2) during harvest time: from fermented juice/must



3-5%

Tank A



3-5%

Tank B



Tank C

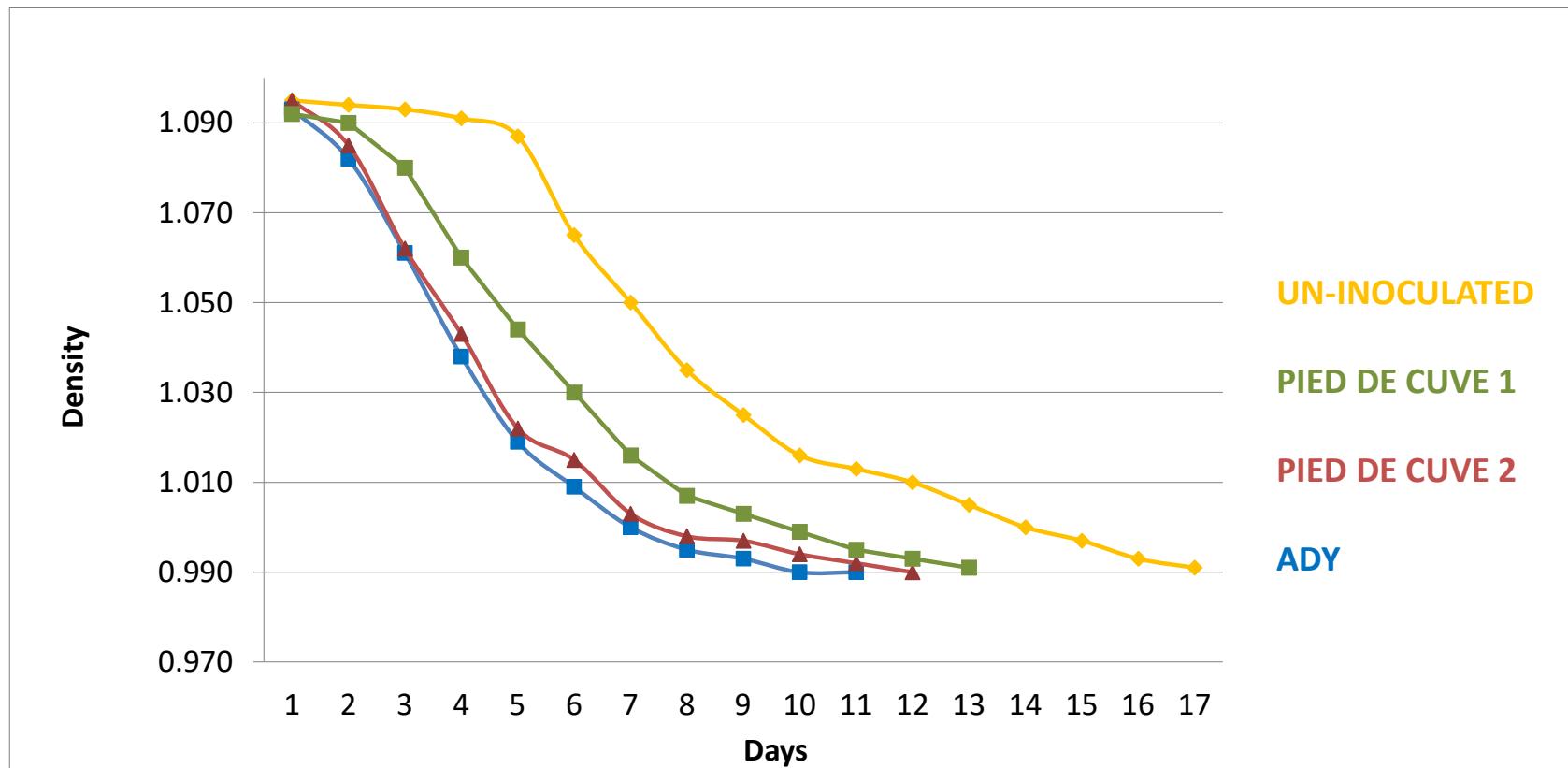


3-5%



Tank D

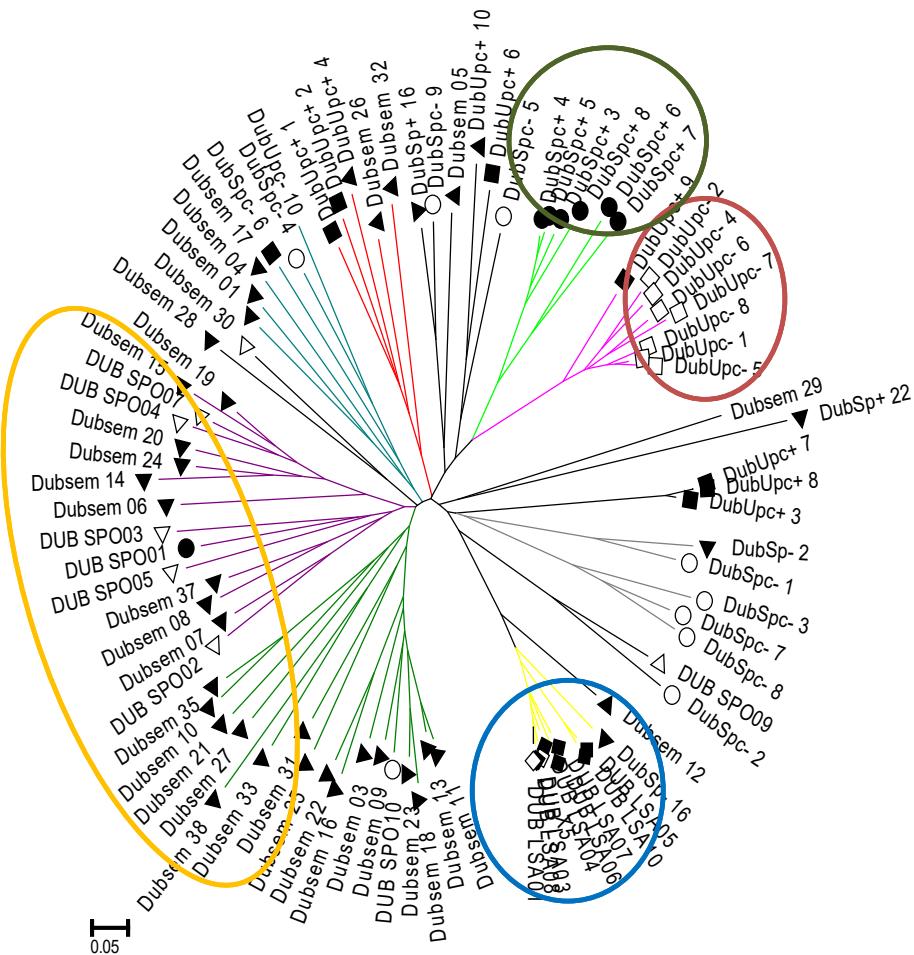
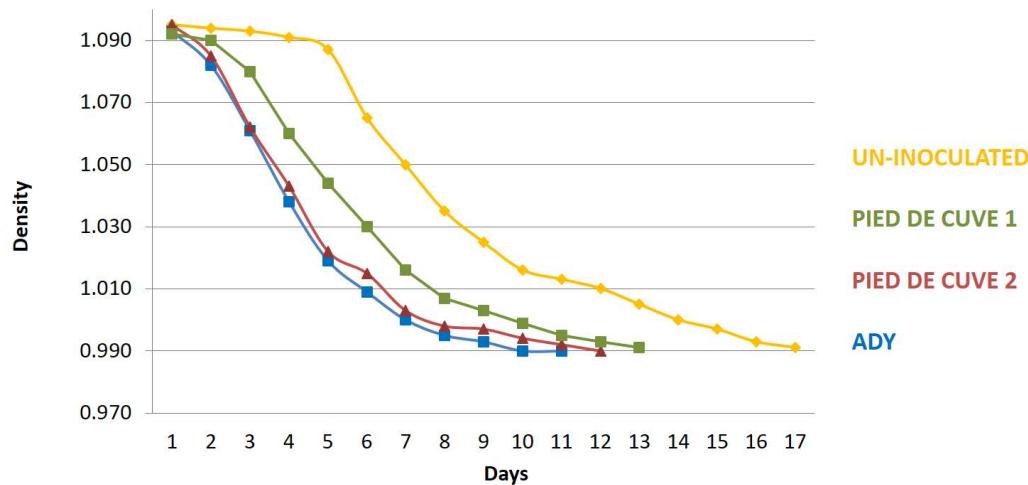
Pied de cuve – ensuring timely and reliable fermentation



Comparison of fermentation dynamics with indigenous yeasts, ADY and two *pied de cuve* treatments.

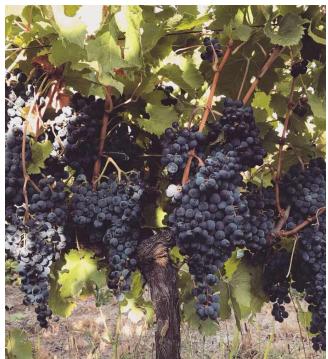
Pied de cuve and *S. cerevisiae* diversity

- higher initial *S. cerevisiae* rates
 - selection of a dominant population

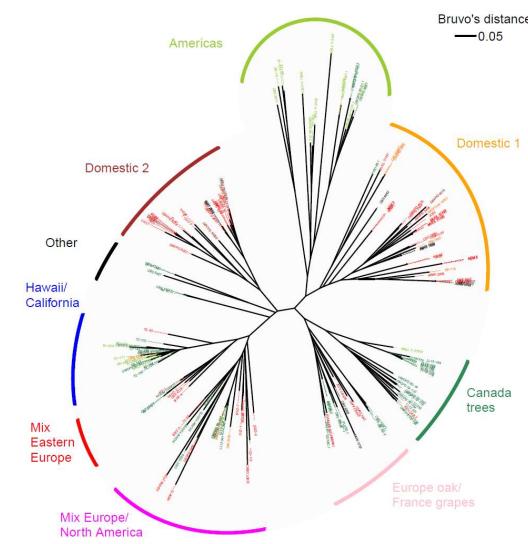


Genetic diversity of *S. cerevisiae* isolates at 75% of AF; dendrogram constructed with genotyping data on 16 microsatellite loci (NJ clustering)

Selection of indigenous yeasts

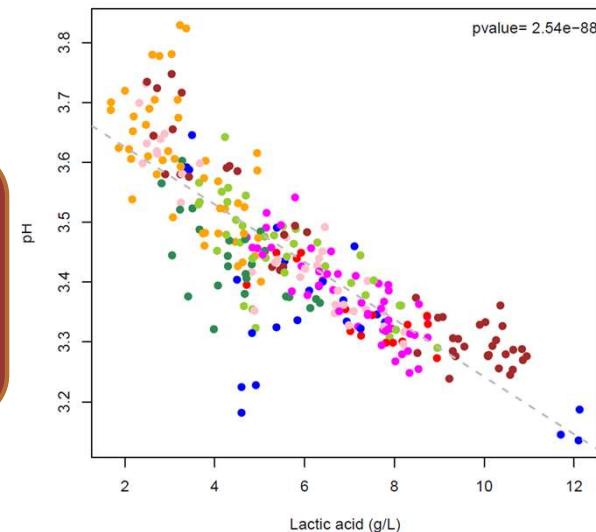


Population-wide selection of *Lachancea thermotolerans* for wine starters



Genotyping of
~200 isolates

Oenological
performance of 94
strains



RESEARCH ARTICLE

The evolution of *Lachancea thermotolerans* is driven by geographical determination, anthropisation and flux between different ecosystems

Ana Hranilovic^{1,2}, Marina Bely³, Isabelle Masneuf-Pomareda^{3,4}, Vladimir Jiranek^{1,2}, Warren Albertin^{3,5*}

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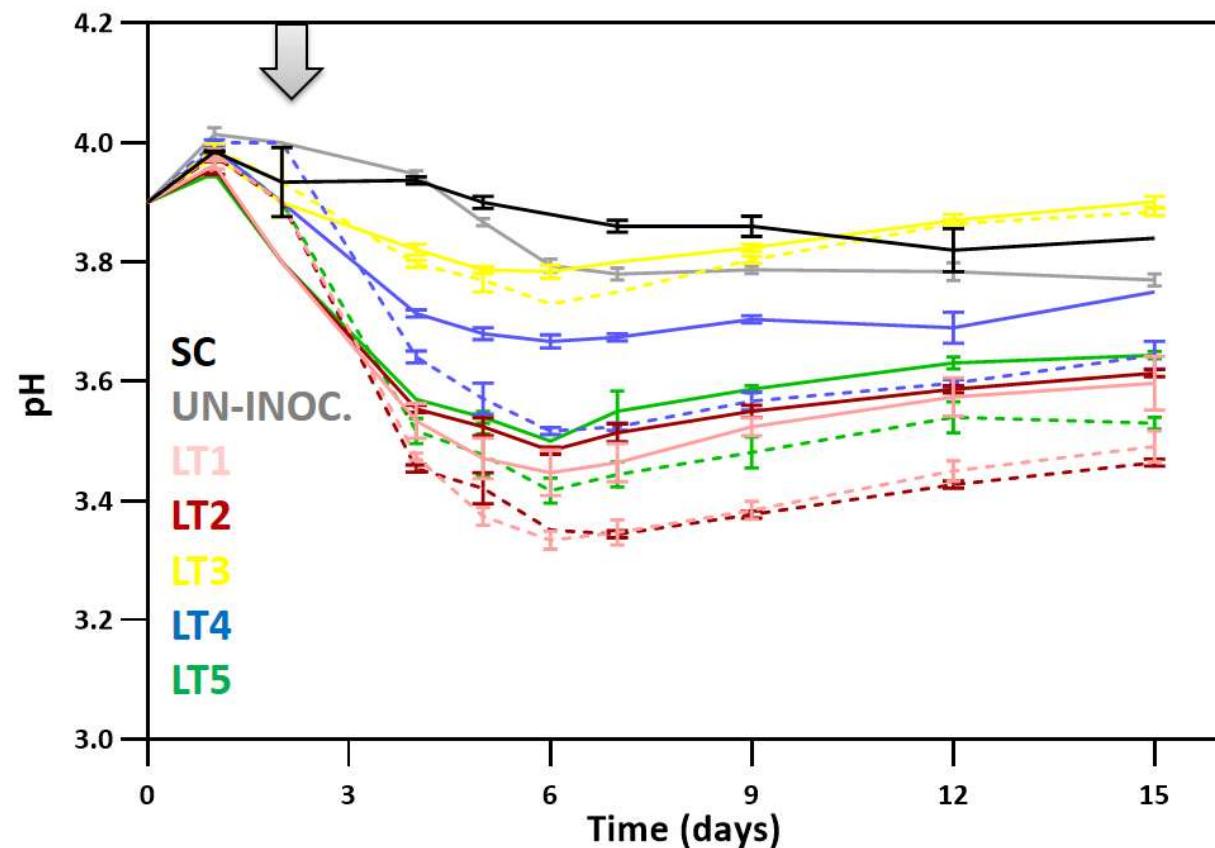
OPEN

Oenological traits of *Lachancea thermotolerans* show signs of domestication and allopatric differentiation

Ana Hranilovic^{1,2}, Joanna M. Gambetta³, Leigh Schmidtke^{1,3}, Paul K. Boss^{1,4}, Paul R. Grbin^{1,2}, Isabelle Masneuf-Pomareda^{5,6}, Marina Bely⁵, Warren Albertin^{5,7} & Vladimir Jiranek^{1,2}

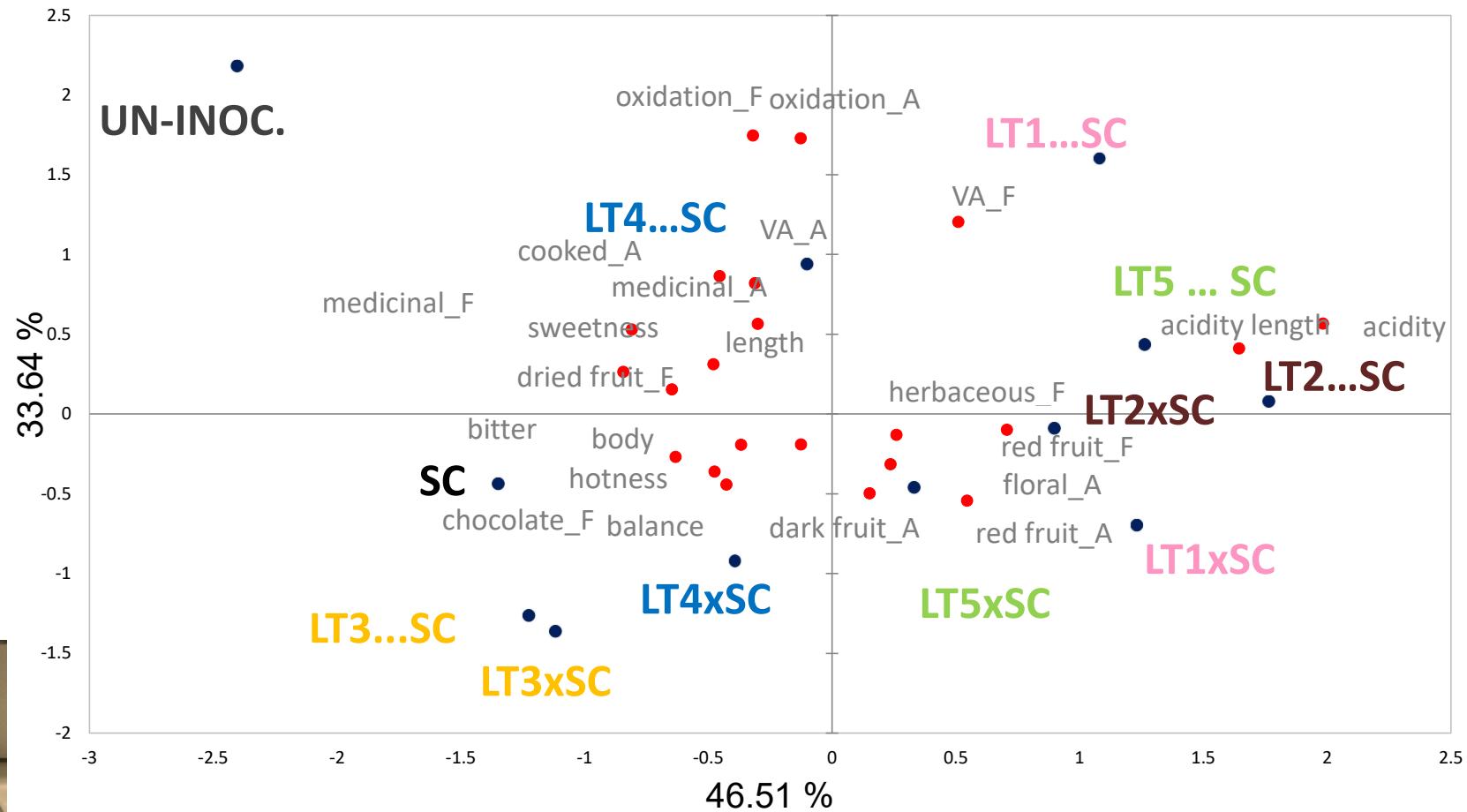


Pilot-scale winemaking trials in warm climate Merlot



Fermentation and acidification kinetics in Merlot. Full lines represent co-inoculations and dotted lines sequential inoculations of *L. thermotolerans* and *S. cerevisiae* (arrow).

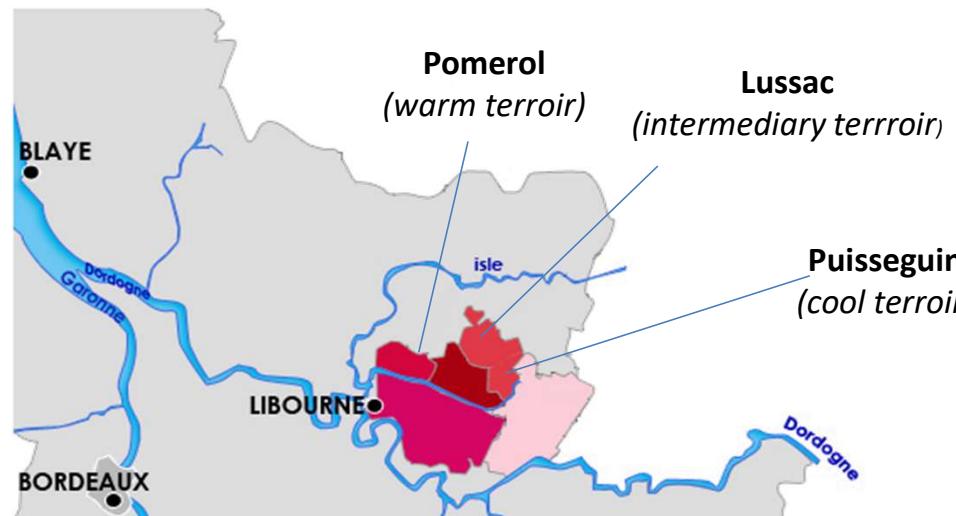
Sensory analysis of Merlot wines



PCA built with attributes significantly ($p < 0.05$) affected by the yeast treatment.

Rate-all-that-apply method; 47 tasters; sequential and co-inoculations are represented with x and ... symbols, respectively.

Organic viticulture and grape berry microbiota



PhD thesis Guilherme Martins, 2012

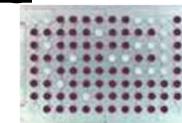


cell
suspension

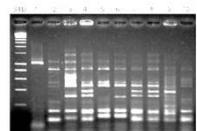
Isolation and
enumeration



Metabolic profiling
(Biolog™)



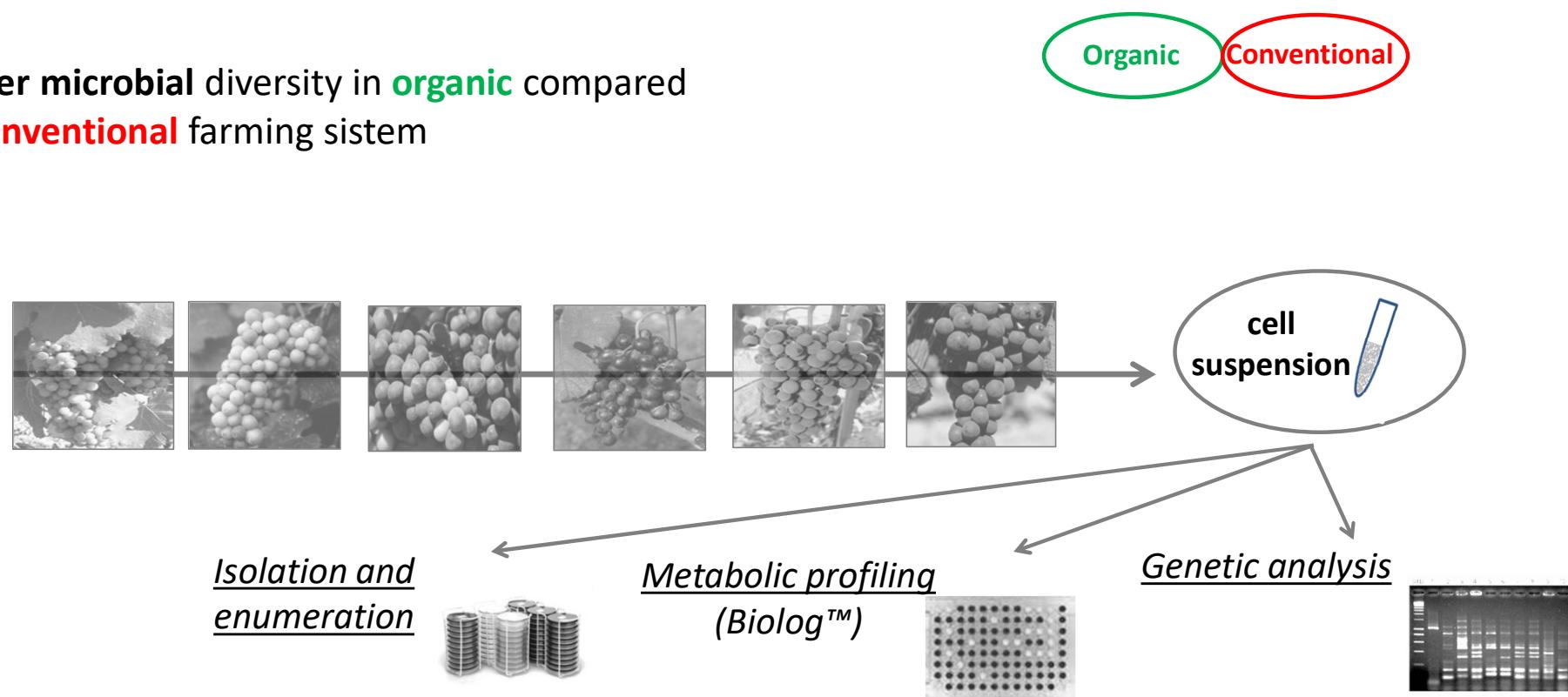
Genetic analysis



Organic viticulture and grape berry microbiota

PhD thesis Guilherme Martins, 2012

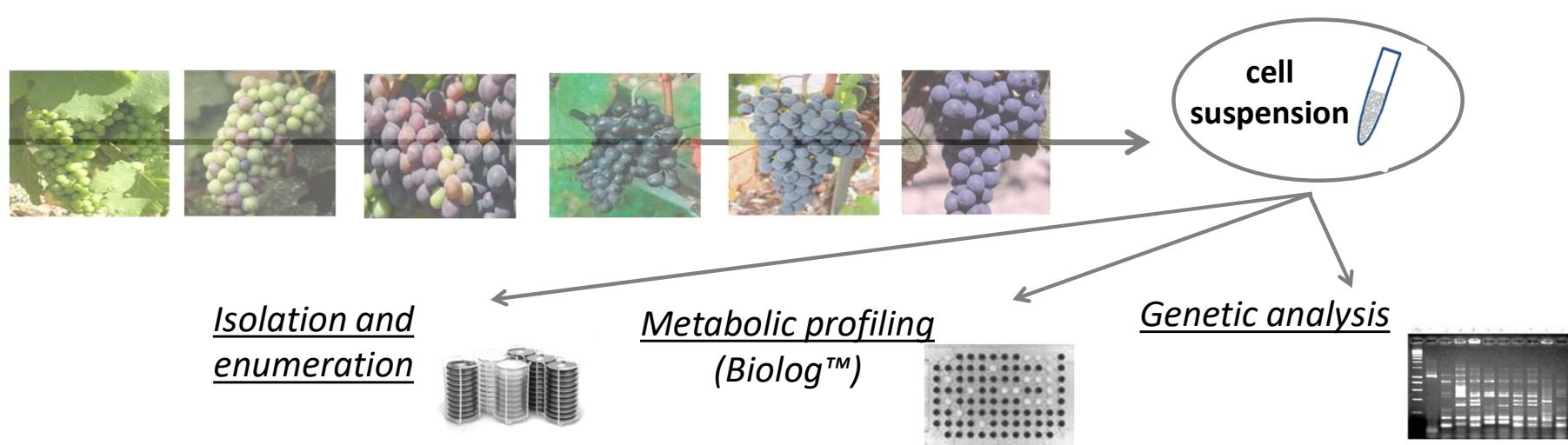
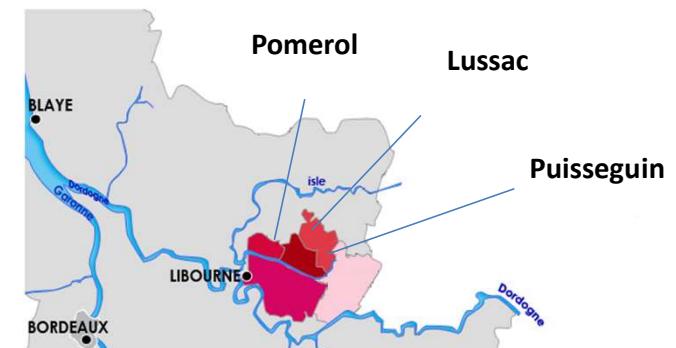
- **Significant effect** of the farming system on population size and diversity
- **Higher microbial diversity** in **organic** compared to **conventional** farming sistem



Organic viticulture and grape berry microbiota

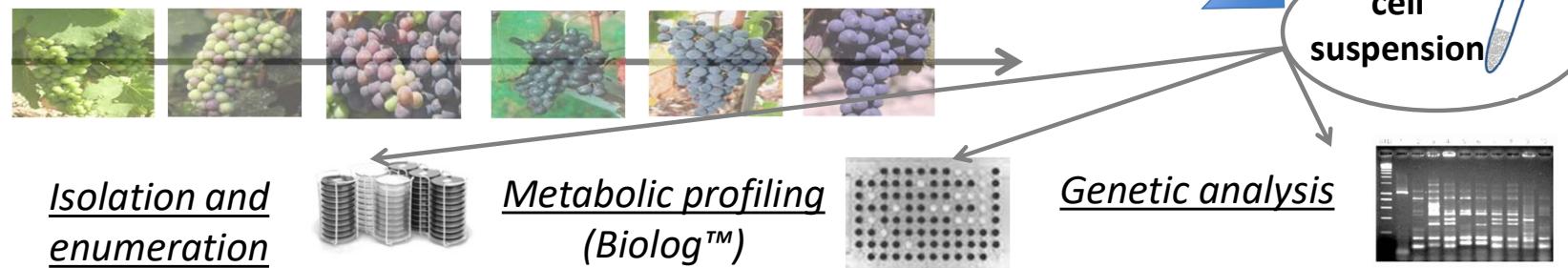
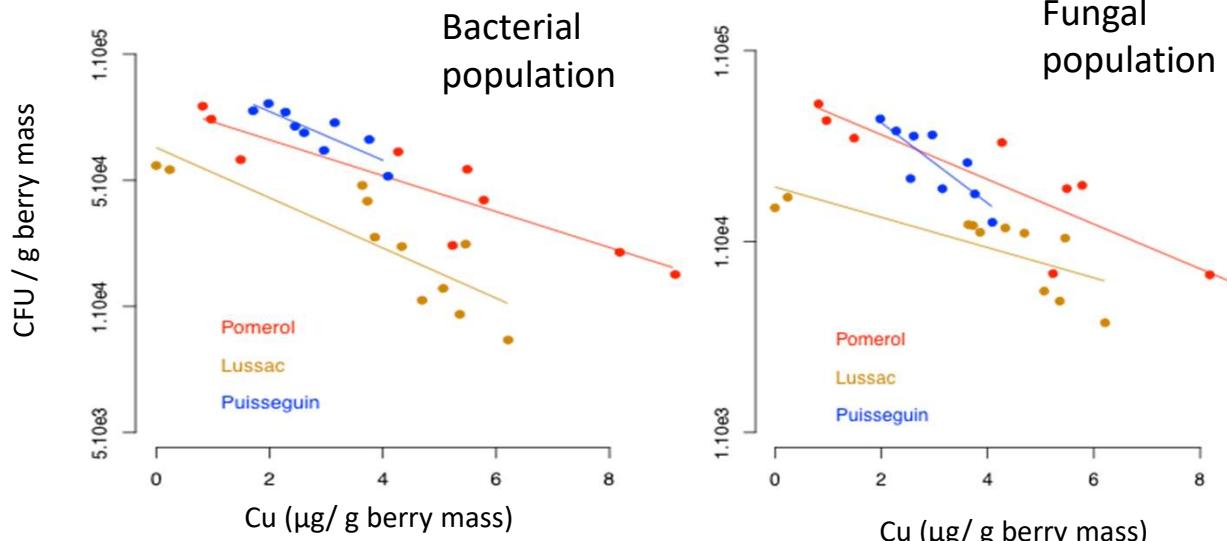
PhD thesis Guilherme Martins, 2012

- Significant effect of the farming system on microbial population size and diversity
- Higher microbial diversity in **organic** compared to conventional farming systems
- Preponderant effects of **microclimate** and **phenological stage**



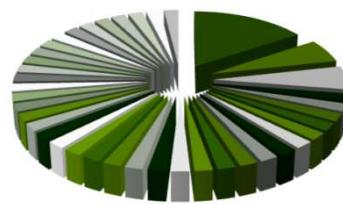
Copper additions negatively affect :

- **Population size**
($p < 0,01$; $\rho = 0.87 - 0.95$)
- **Population diversity:**
 - **metabolic profiles**
(e.g. Shannon index based on Biolog® metabolic profiles;
 $p < 0,0001$; $\rho = 0,90$)
 - **genetic profiles**
(e.g. bacterial diversity
 $p < 0,012$; $\rho = 0,58$)



Link between microbial diversity and ‘complexity’?

- Not clear answer from the literature
- Microbial diversity ≠ wine ‘complexity’
- Chemical complexity ≠ sensory complexity



The complexity of wine: clarifying the role of microorganisms

Sophie Tempère¹ · Axel Marchal¹ · Jean-Christophe Barbe^{1,2} · Marina Bely¹ · Isabelle Masneuf-Pomarede^{1,2} · Philippe Marullo^{1,3} · Warren Albertin^{1,4} 

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Summary of different fermentation modalities

mix of indigenous *S. cerevisiae* and non-*Saccharomyces* yeasts

pre-selection of multiple species/strains

inoculation of *S. cerevisiae* strain (active dry yeast; ADY)



Pretorius, 2016



Thank you for your attention!

