

Unraveling metabolically active fungal-bacterial diversity in commercial organic vineyard soils

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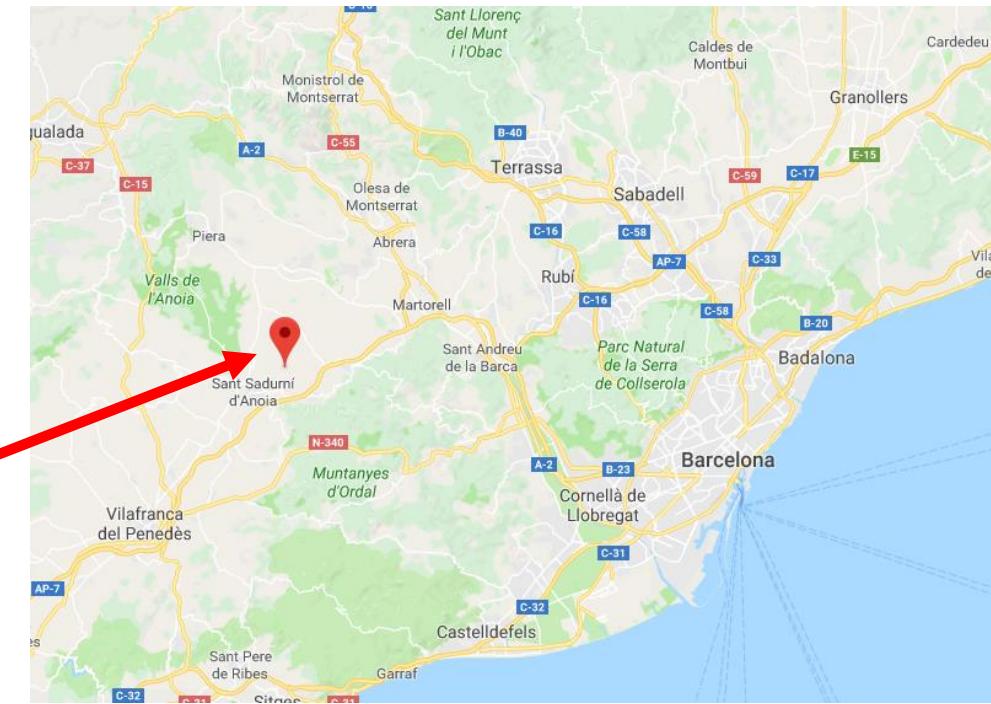
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SPAIN**

OBJECTIVES

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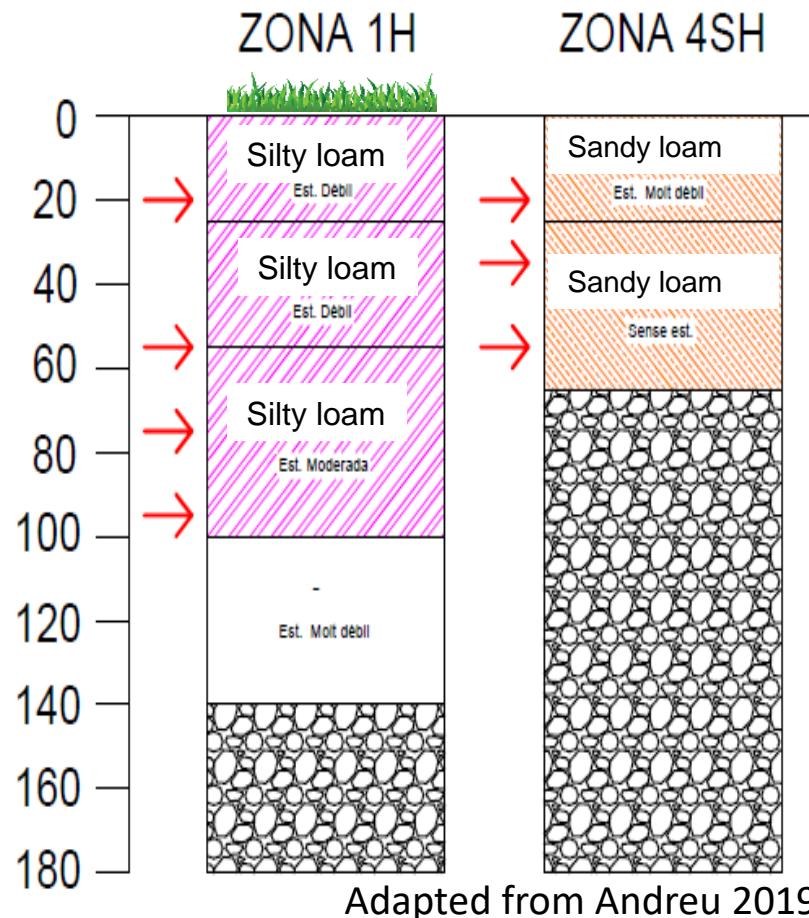
To assess the impact of phenology (pre-bloom and post-harvest periods) on the diversity of metabolically active soil-rhizosphere microbiota in a commercial vineyard in Sant Sadurní d'Anoia, a typical cava and wine producing region (Penedès DO, Catalonia, Spain).



Location of Catalonia (dark green)

Plot coordinates: [41°49'N 1°28'E](#)

- *Vitis vinifera* variety 'Macabeu', grafted onto 41-B (*V. berlandieri* x *V. vinifera*), 20 years old.
- Total Surface 6 ha, vine spacing: 1,0 x 2,6 m.
- Organic and rainfed farming.
- Fertilizer: 20 tones (composted cow manure)/ha every 4 years.
- Permanent cover crop and **silt loam** soil in Zone 1 and tillage and **sandy loam** soil in Zone 4.



Mediterranean climate (SMC)

Temperature
Summer: 23–24 °C
Winter: 6–8 °C

Rainfall
- Annual 550 mm
- Maximum rate in fall and minimum rate in summer

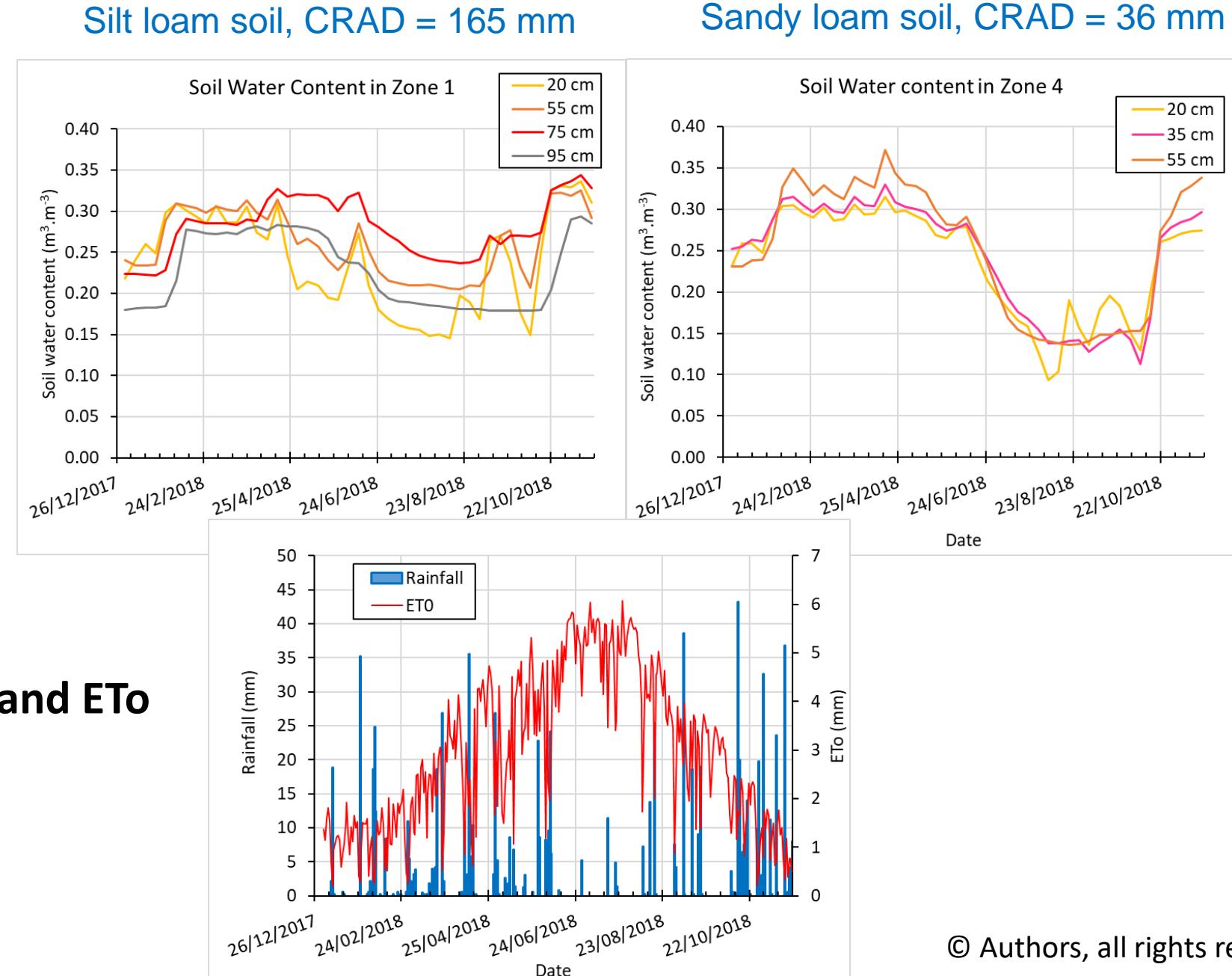
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RESULTS

Soil water content

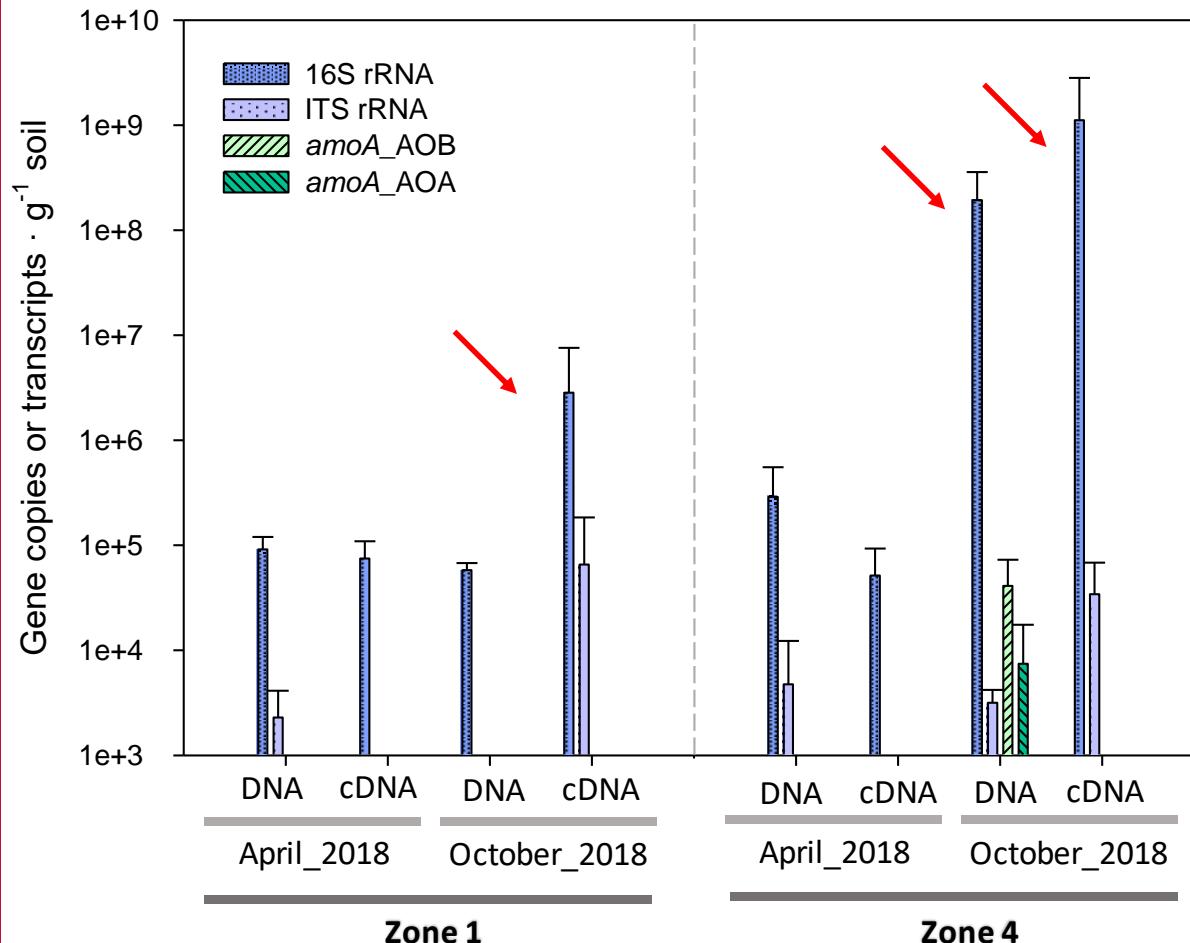
Measured with FDR probes (model 10HS, (Decagon) at different soil depths

Rainfall and ETo



RESULTS

Microbial assessment: Quantitative PCR



April 2018: Pre-harvest

October 2018: Post-harvest

In post-harvest period, bacterial and fungal population were more enriched and active, specially in zone 4.

In zone 1, active bacterial population increased 2 orders of magnitude.

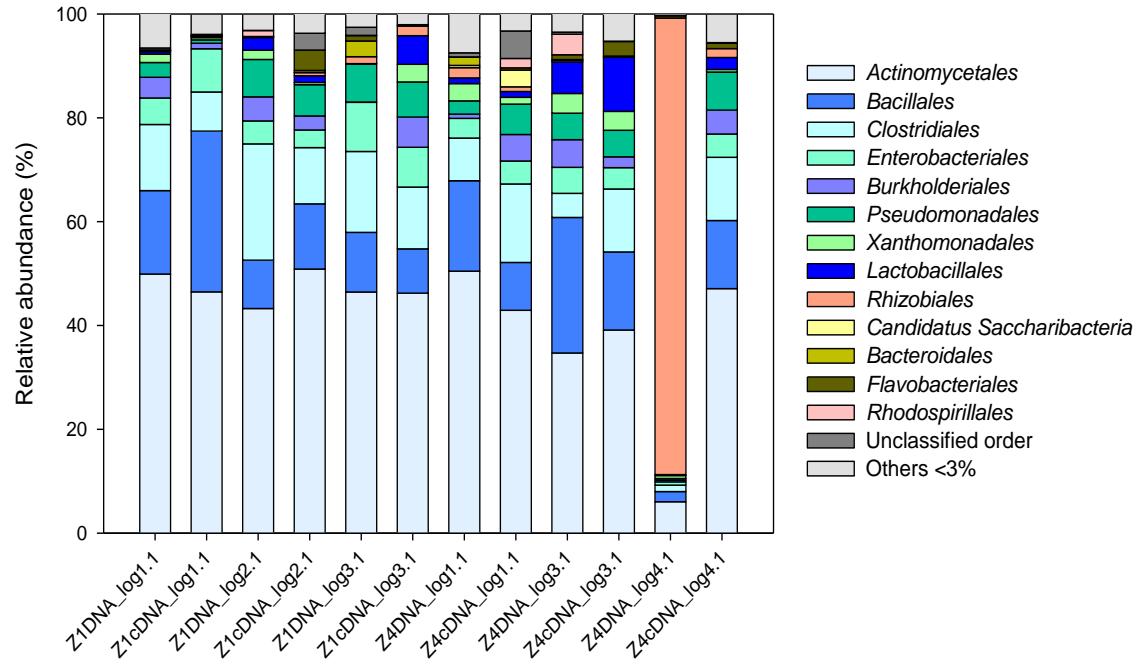
In zone 4, metabolically active bacteria increased 5 orders of magnitude.

The harvest and climate conditions promoted this activity.

Bacteria and Archaea Ammonia Oxidizers (AOB and AOA), were detected in, post-harvest period.

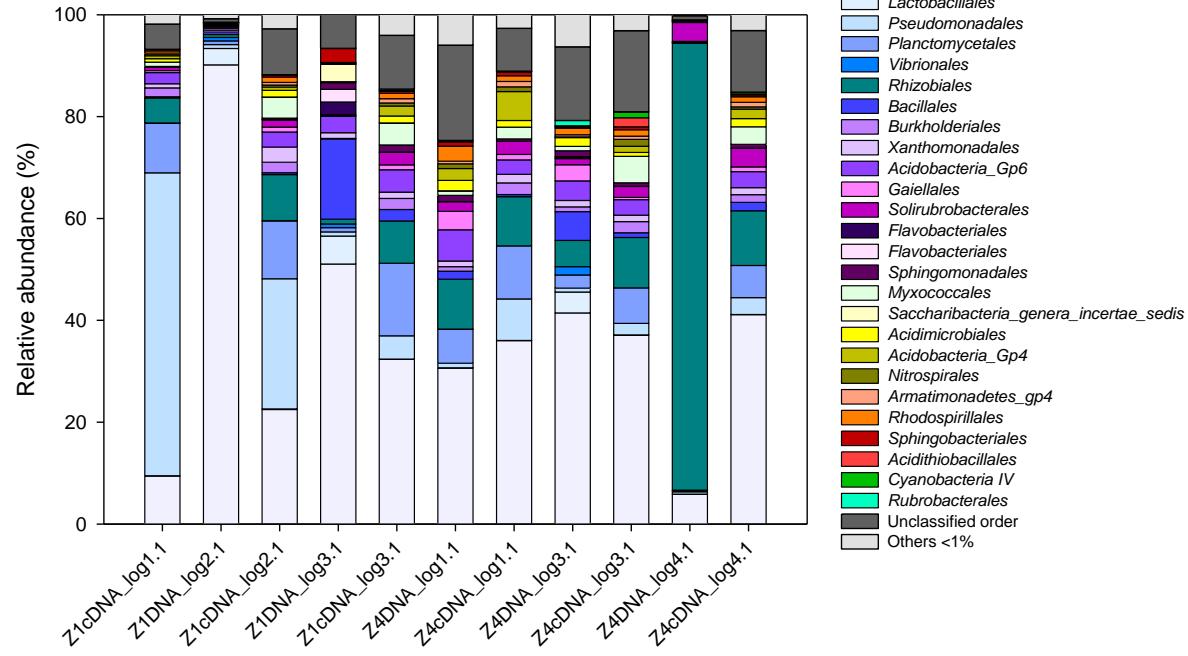
Nitrospirales presence and activity (MiSeq data), could hamper ammonia-oxidizers activity.

April 2018: Pre-harvest



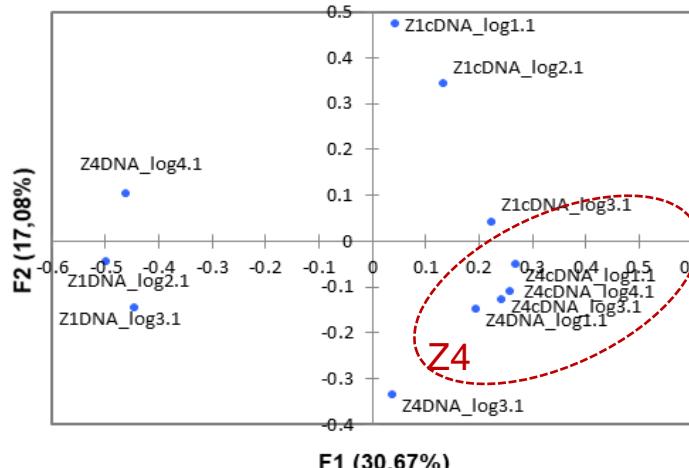
Bacterial Diversity

October 2018: Post-harvest

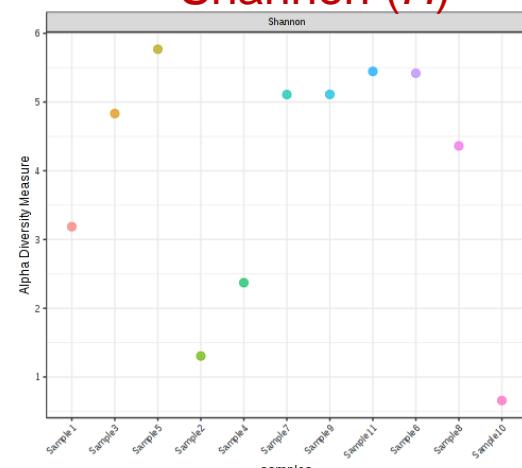


Bacterial Diversity: POST-HARVEST

PCoA



Shannon (H)

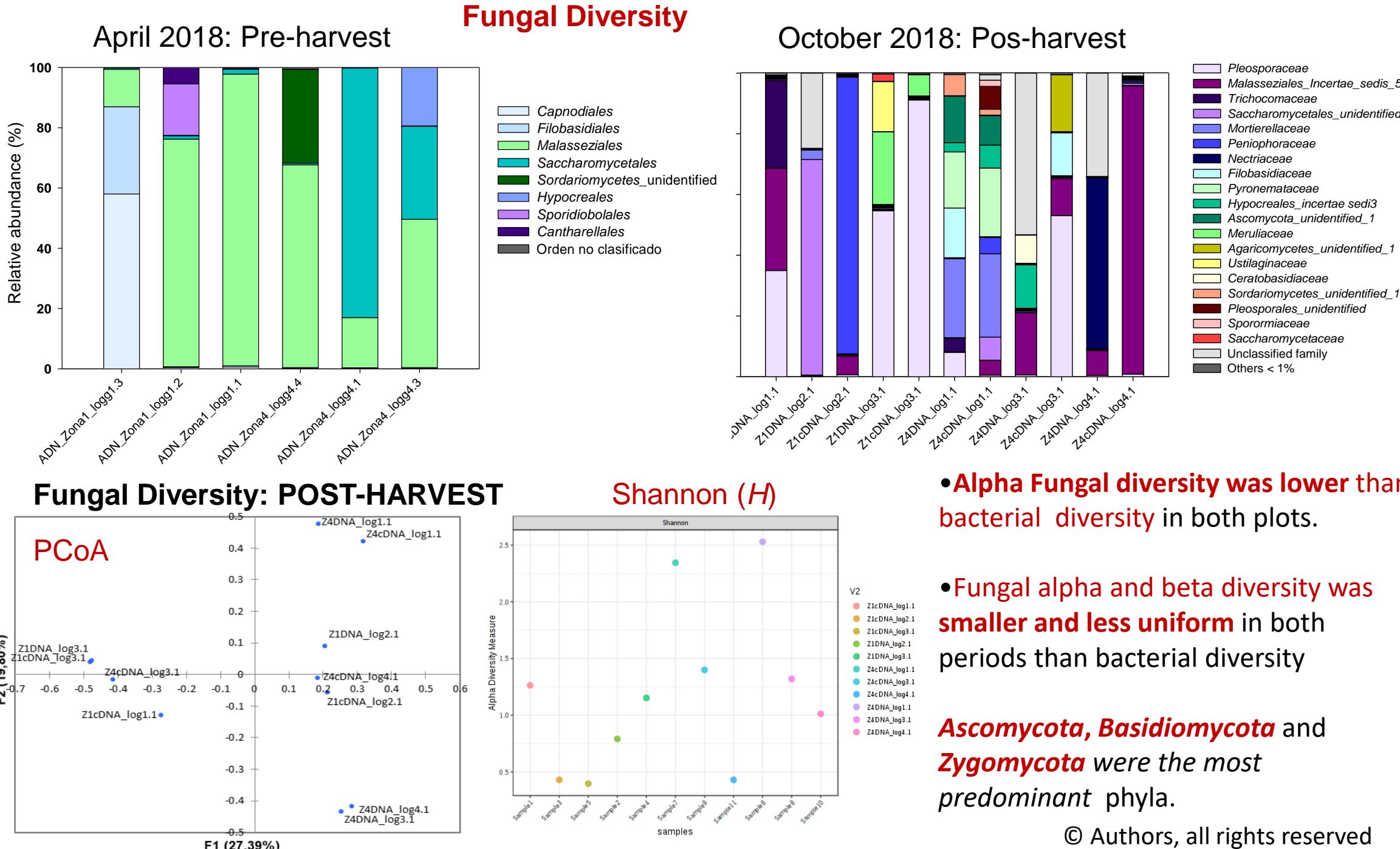


- Total and functional bacterial diversity in Z4 is more homogeneous in postharvest period than Z1 zone.

- Actinobacteria** (mainly by *Actinomycetales* order), **Proteobacteria** (mainly by *Rhizobiales* and *Pseudomonadales* orders) were the most predominant phylotypes.

- Clostridiales** (*Firmicutes* phylum) phylotypes were completely replaced during post-harvest season.

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CONCLUSIONS

- Soil bacterial and fungal communities were more metabolically active during post-harvest than pre-harvest season in both zones.
- No metabolically-active fungal community was detectable in pre-harvest period. Fungal populations were less diverse than bacterial diversity.
- Both environmental conditions and the mechanical harvest may promote microbial growth due to sugar availability in soil (-30 cm) linked to rainy periods. The vineyard, in post-harvest period, could be more exposed to phytopathogens, so this could be an interesting period to control these communities and consequently the potential infection for the next productivity period.
- High throughput sequencing analysis (16S/ITS MiSeq) revealed that the microbial diversity was specific both for each plot each plant and time period.
- The diversity of bacterial and fungal populations increased during post-harvest season.
- Simultaneous RNA/DNA-based molecular biology tools could improve the knowledge of metabolically active microbial populations in soils at different seasons and phenological stages of rainfed vineyard.
- These can be important in order to study and evaluate the potential and real emissions of greenhouse gases from vines in Mediterranean conditions under climate change.
- This information must be used to accomplish the compromises developed in COP 21 and COP 22, related to mitigation strategies.