

Physicochemical, Microbial, and Microbiome Dynamics in Winery Waste Composting



G. Mitropoulou¹, I. Karapantzou¹, I. Prapa¹, D. Papanikolaou¹, V. Charovas² and Y. Kourkoutas¹

1. Laboratory of Applied Microbiology and Biotechnology, Department of Molecular Biology and Genetics, Democritus University of Thrace, Alexandroupolis, 68100, Greece; 2. Evritika Kellaria S.A., Orestiada, 68200, Greece

INTRODUCTION

Annually, the winemaking industry generates 0.3–0.5 kg of wine byproducts/L, including winter pruning wastes that can be toxic if disposed of without pretreatment, due to high content of organic load and phytotoxic compounds along with high acidity. Composting is the natural process of transforming organic matter to fertilizers, rich in essential nutrients for plant growth by microorganisms under controlled conditions. The organic substrate used in the composting process affects microbial populations, reflecting thus the dynamics of enzymatic activity, decomposition of organic matter, and nitrogen transformations. However, the microbiome associations during the whole process are still unexplored.

Table 1: Concentration of important trace elements and metals during microbial composting of winery wastess

Frace elements/ metals	Day 0	Day 60		
Total Ca	0 10+0 06			
(% dry weight)	0.10±0.00	1.30±0.30		
Active Ca	1 /በ⊥በ 32	<u>በ 13₊</u> በ በን		
(% dry weight)	1.4010.52	0.13±0.02		
CaCO3	1 /IO+O 2 1	1.60±0.5		
(% dry weight)	1.4010.21			
P (ppm)	0.32 ± 0.04	0.25 ± 0.02		
K (ppm)	0.45 ± 0.001	0.40 ± 0.16		
Mg (ppm)	3.92±0.09	2.00 ± 0.64		
Zn (ppm)	580.00 ± 14.10	83.50±23.30		
Mn (ppm)	331.50±2.12	237.50±24.75		
Fe (ppm)	16.50 ± 2.12	12.00 ± 2.83		
Cu (ppm)	32.00±1.41	33.50±2.12		

Table 4: Changes in fungal phylum abundances (%) during microbial composting of winery wastes.

	Fungal phyla	L	Day 0			Day 60		
	Ascomycota			97.43±0.64			90.28±3.05	
	Basidiomycota		2.35 ± 0.64			8.16±2.57		
	Chytridiomyco	ota	0.01 ± 0.00			0.01 ± 0.00		
En	Entomophthoromycota		0.01 ± 0.00			0.01 ± 0.00		
	Glomeromycota		0.03 ± 0.01			0.86 ± 0.40		
	Mucoromycot	a	0.18 ± 0.01			0.68 ± 0.09		
	Tax	onomic	binnir	ng at G	enus le		g_Zygosaccharomyces g_Wickerhamomyces g_Wallemia g_Volutella g_Verticillium g_Trichothecium g_Trichosporon g_Torulaspora g_Torulaspora g_Thielavia g_Terfezia g_Stachybotrys g_Sphaeronaemella g_Spencermartinsia g_Sordaria	
undance (%)	80 -						g_Scedosporium g_Sarocladium g_Saccharomycodes g_Saccharomyces g_Rhodotorula g_Rhizomucor g_Rhizomarasmius g_Reniforma g_Racocetra g_Psora g_Psora g_Pseudaleuria g_Plectosphaerella g_Pichia g_Peziza g_Petriella g_Pesotum g_Penicillium g_Parascedosporium g_Orbilia g_Ophiocordyceps g_Natria	
nulative relative abu	40 –						g_Nectria g_Myrothecium g_Myceliophthora g_Mucor g_Mortierella g_Monacrosporium g_Microdochium g_Microcera g_Metarhizium g_Massaria g_Lophotrichus g_Lophiostoma g_Lophiostoma g_Lophiostoma g_Lomentospora g_Kernia g_Ilyonectria g_Hydnellum g_Humicola g_Holtermanniella g_Heydenia g_Guehomyces g_Graphium g_Gibelluloosis	

OBJECTIVE

The aim of the present study was to investigate the physicochemical, microbial, and microbiome dynamics during the microbial composting of winery waste and assess the suitability of the resulting compost as a biofertilizer for grapevine growth.

MATERIALS AND METHODS

A pilot-scale composting system was utilized and samples were collected over a 60-day period to study the physicochemical parameters, enzymatic activity, microbial counts, and microbiome composition as described previously [1]. Additionally, the potential phytotoxicity of the compost was evaluated using a germination index test with barley seeds and its effectiveness as a substrate for grapevine growth was assessed in a proof-of-concept study [2].

рH Conductivity 9,0 ວ 2,5 8,5 8,0 2,0 표 ^{7,5} 전 7,0 6,5 £ 1,0

RESULTS

Table 3: Changes in bacteria phylum abundances (%) during microbial composting of winery wastes.

Bacterial phyla	Day 0	Day 60
Acidobacteria	0.13±0.05	1.98±0.92
Armatimonadetes	0.03 ± 0.001	0.28 ± 0.12
Bacteroidetes	0.37 ± 0.16	16.62 ± 1.36
Candidatus saccharibacteria	0.01±0.001	0.04±0.001
Chloroflexi	0.01 ± 0.001	0.06 ± 0.02
Cyanobacteria	0.02 ± 0.001	0.02 ± 0.001
Planctomycetes	0.03 ± 0.001	0.03 ± 0.001
Proteobacteria	99.37±0.26	80.97±0.32



Taxonomic binning at Genus level





Figure 3: Taxonomic binning at Genus level of fungi during microbial composting of winery wastes.

CONCLUSION

 Analyzing the microbiome diversity at the beginning and at the end of the composting process of winery waste provided valuable insights about the bioprocess.

- > Germination Index (GI) equal to 133.99 was estimated. Of note, GI values > 80 indicate no phytotoxicity.
- > Effectiveness of the final product was verified by estimating the percentage yield of grapevine leaf dry matter compared to the control samples, leading to values 110 ± 1.3 and $90 \pm$ 1.8 for 25:75 and 50:50 (product: commercial substrate), respectively.



Operational Programme «Eastern Macedonia and Thrace»

Co-funded by Greece and the European Union





artnership Agreement 2014 - 2020

Figure 2: Taxonomic binning at Genus level of bacteria during microbial composting of winery wastes.

ACKNOWLEDGEMENT

The project was co-financed by the European Union and National Resources under the Operational Programme "Eastern Macedonia and Thrace" 2014-2020, AMOP7-0074893, MIS number: 5076618, «AgroWasteCompost»

 Excess enzymatic activities, physicochemical analysis results (low C/N, total N increased, and neutral pH), along with lack of phytotoxicity indicated the suitability of the final product and its effectiveness as a substrate for vine growth was confirmed

✓ More research is still required to fully understand the underlying mechanisms of winery waste biotransformation into efficient biofertilizer and verify its efficiency in real plant

culture conditions.

REFERENCES

[1] Karapantzou et al. (2023). Sustainability, 15(9), 7484. https://doi.org/10.3390/su15097484.

[2] Paradelo et al (2009). Waste Management, 29(2), 579–584. https://doi.org/10.1016/j.wasman.2008.06.019.