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Characterization of New Yeast Strains Isolated from Local Grape Varieties of the Soufli Region in Northern Greece

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Abstract

Over the past few years, consumers' preference for wines with a distinct regional identity has shifted producers to adopt traditional grape varieties and has increased interest in the indigenous grapevine microbiome, due to its potential influence on the wine quality characteristics. In the present study, indigenous *Saccharomyces cerevisiae* strains were isolated from three local grape varieties ("Keratsouda", "Karnachalas", and "Bogialamas") of the Soufli region in Northern Greece. After molecular identification, all strains were evaluated for key oenological traits (ethanol resistance, SO₂ resistance, H₂S production, and amino-acid decarboxylation). Three representative strains (one per grape variety) were further evaluated for their long-term survival after freeze-drying and their fermentation potential at 12 °C and 22 °C. Ethanol content ranged in similar values (11.9–12.5%) for all wines, but kinetic parameters were significantly affected by the strain used and the fermentation temperature. Principal Component Analysis applied on volatile compounds determined by HS-SPME GC/MS indicated that the fermentation temperature rather than the indigenous yeast strains affected significantly the volatile composition in the new wines and resulted in products with distinct sensory profiles.

Keywords: indigenous *S. cerevisiae* strains; regional grape varieties; yeast technological characterization; wine aromatic profile

1. Introduction

Industrial winemaking practices rely on the extensive use of a few selected commercial *Saccharomyces cerevisiae* strains often promoted under different product names and brands [1,2]. The dynamic nature of these starter cultures ensures the effective completion of the alcoholic fermentation by outcompeting the native grape microflora and also provides specific sensory characteristics to the final product [3]. However, as a direct consequence of the decrease of microbial communities involved in fermentations, a series of constraints has risen for the wine sector. Reduction of wine profiles' uniqueness, general flattening of the bouquet and organoleptic uniformity of products, even across different vitivinicultural regions, are the most common malfunctions observed [4,5].

Nevertheless, based on the idea that indigenous yeasts may provide unique flavors that preserve and reflect the regional characteristics of a vineyard, many winemakers have shifted again towards spontaneous fermentations [3,6]. In fact, a vineyard's microflora is an interactive ecosystem, unique for each grape variety, that is highly affected by the geographical location, the climate, as well as the vitivinicultural practices applied. This results in significant species diversity that represent and highlight the regional character of the vineyard, simply called 'terroir' [7–11].

This aspect has led to a growing market and industrial interest in isolating and exploiting indigenous yeasts [12] in an effort to produce unique wines, but also guarantee consistent fermentation outcomes across vintages [7,13]. Various indigenous strains of *S. cerevisiae* have been isolated the past years from various grape varieties (e.g. Cabernet Sauvignon, Malbec, "Gioia del Coll" DOC, Merlot, Garnacha and others) worldwide [9]. These strains have developed

to regional-specific starter cultures showing quick adaptation to their specific environments, enhanced fermentation efficiency and production of unique aroma profiles [11,14,15].

Greek vineyards represent a complex mixture of international and domestic varieties. Specifically, in the Soufli region of Northern Greece “Keratsouda”, a highly productive rose variety with bright pink color, is grown [16]. Similarly, “Karnachalas” and “Bogialamas” are two genetically close varieties that are typically used as a mixture by local winemakers [17]. The indigenous yeast communities of these varieties have not been yet studied, but in an effort to promote the whole region they could be carefully exploited and result in signature wines with high market potential [17,18].

In this context, the aim of the present study was the isolation and of new *S. cerevisiae* strains from the local varieties “Karnachalas”, “Keratsouda”, and “Bogialamas” of the Soufli region in Northern Greece. All strains were selected according to their technological characteristics and evaluated concerning their impact to wine fermentation and quality. Thus, the isolated yeast strains were assessed as potential fermentation starters in order to enhance the expression of the regional characteristics of the produced wines.

2. Materials and Methods

2.1. Isolation of New Yeast Strains from Local Grape Varieties

Grape samples from the 3 local varieties (~2 kg for each), “Karnachalas”, “Keratsouda”, and “Bogialamas” were provided by the company “Winery – Distillery Gr Bellas L.P.” (coordinates: 41.183865, 26.260774) during the 2023 vintage, in late October.

For each variety, stems along with any bruised or rotten grapes were removed by hand and the remaining grapes were crushed in an iMix bag mixer (Interlab, Mourjou, France). The extracted grape must was then transferred to batch bioreactors (0.5 L) and left to perform spontaneous alcoholic fermentation at ~25 °C. Fermentations were monitored daily (by Be° measurements) until full sugar utilization occurred or no further activity was observed. After completion, samples were taken from the mid-zone of each bioreactor, decimally diluted and plated on sterile YPD agar (yeast extract 10 g/L, peptone 20 g/L, dextrose 20 g/L, agar 20 g/L). Agar plates were incubated at 30 °C for 48-72 h and the dilutions on which yeast colonies were clearly distinguishable were chosen for re-streaking. All yeast strains were routinely kept in Malt broth and Malt agar plates (Condalab, Madrid, Spain) at 4 °C until further use. Stock samples were kept at -80 °C by inoculating fully grown cultures in 30% v/v glycerol tubes [6].

The new strains were observed under an optical microscope. After examining their cellular morphology, 10 strains in total, resembling *S. cerevisiae* cells, were selected for molecular identification.

2.2. Molecular Identification of the New Strains

Total DNA of the new isolated strains, was extracted using the Nucleospin® DNA Yeast kit (Macherey-Nagel GmbH & Co. KG, Düren, Germany), following the manufacturer’s instructions. The highly variable Internal Transcribed Spacer (ITS) regions of the ITS1 and ITS2 sequences were amplified with the primers ITS1-4.

Polymerase Chain Reaction (PCR) amplification was performed using the HotStarTaq Plus Master Mix Kit (Qiagen, Germatown, MD, USA), consisting of 35 cycles with the following steps: 94 °C for 3 min, 35 cycles of 94 °C for 30 s, 53 °C for 40 s, 72 °C for 90 s, and the final elongation step at 72 °C for 5 min. PCR products were then subjected to electrophoresis in 2% agarose gel to confirm amplification and to determine the relative intensity of bands. Then, the amplicons were purified using Ampure PB beads (Pacific Biosciences).

PCR products were subjected to electrophoresis on 2% agarose gels to confirm amplification and assess band intensity. Subsequently, a purification step was performed using Ampure XP beads (Beckman Coulter, Brea, California, USA).

Sequences obtained by MR DNA (www.mrdnlab.com, Shallowater, TX, USA) were taxonomically classified using the BLASTn algorithm against the NCBI reference database (www.ncbi.nlm.nih.gov).

2.3. Technological Characterization of the New Strains

- Ethanol resistance: Each yeast strain was spot-inoculated on grape must-agar plates containing increasing concentrations of ethanol (up to 14% vol). Resistance to ethanol was evaluated on the basis of positive growth after 72 h at 26 °C [19].

- SO₂ resistance: Each yeast strain was spot-inoculated on grape must-agar plates containing increasing concentrations of SO₂ in the form of the commercially available potassium metabisulfite. Plates were incubated at 26 °C for 48 h and the colony growth size was assessed (no growth, small growth, medium growth, high growth) [20].
- H₂S production: Each yeast strain was spot-inoculated on Bismuth Sulfite Glucose Glycine Yeast (BIGGY) agar and incubated at 26 °C for 48 h. H₂S production was assessed based on the color formation of the colonies: white (no production); yellow (low production); dark yellow (moderate production); brown or black (high production), as previously described [20].
- Decarboxylation of amino acids: Each yeast strain was spot-inoculated on modified YPG agar plates (5 g/L bacteriological peptone; 5 g/L yeast extract; 1 g/L glucose; 20 g/L agar), as recently described [21]. The medium was supplemented with 10 g/L of each amino acid and 6 mL of a 1% ethanol/water solution of bromocresol purple indicator at a final concentration of 0.5%. Before sterilization, the pH was adjusted to 5.2 in all cases. The plates were incubated at 26 °C for 3-5 days. The formation of a purple ring around the grown colonies indicated positive reaction of the strain.

The strains evaluated for their technological characteristics were grown overnight at Malt broth at 30 °C. Prior to each analysis, they were diluted using sterile ¼ Ringer's solution (VWR International GmbH, Radnor, PA, USA) at a concentration of 10⁵ cfu/mL.

2.4. Freeze-Drying and Monitoring of Yeast Cell Viability During Storage

Three yeast strains (one for each variety) were selected based on their technological characteristics, and examined for cell survival during long-term storage.

Each yeast strain was initially grown overnight at 1 L flask with Malt broth at 30 °C. Yeast cells were collected by centrifugation and subsequently freeze-dried on a BenchTop Pro (Virtis, SP Scientific, Warminster, PA, USA) [22]. The freeze-dried cultures were then stored at 20-25 °C, 4 °C and -18 °C for a period up to 9 months. Cell survival was monitored in frequent intervals (1, 3, 6, and 9 months). 1 g of freeze-dried yeast cells biomass was transferred to 9 mL of sterile ¼ Ringer's solution, serial decimal dilutions were performed, and plate counting on Malt agar plates after incubation at 30 °C for 72 h.

The survival rate % at each time interval was calculated as logcfu/g, divided by the initial yeast cell counts (logcfu/g), and multiplied by 100, as previously described [23].

2.5. Fermentation Efficiency of the New Strains

Fermentation efficiency of the 3 new yeast strains was evaluated in batch fermentations (500 mL) at 22 °C and 12 °C, based on the region's climate during the winemaking season. Varietal grape musts ("Keratsouda" must, and "Karnachalas"/"Bogialamas" must as a mixture) were provided by the company "Winery – Distillery Gr Bellas L.P." immediately after pressing (2024 vintage).

Fermentation inoculums were prepared as previously described [6]. Fully grown yeast strains were firstly inoculated to the grape must of its origin (50 mL) and subsequently re-inoculated to the bioreactors (500 mL) reaching a final concentration of 10⁶ cfu/mL.

All fermentations were performed in triplicates. Wine samples were collected after completion and subjected to chemical analyses.

2.6. Chemical and Sensory Analyses

pH, total acidity, and volatile acidity of wine sampled were determined as previously described [24].

Ethanol, glycerol, residual sugars, and organic acid content were determined by HPLC analysis [Shimadzu chromatography system (Shimadzu Corp., Germany)]. Fermentation parameters were calculated as previously described [24].

Gas Chromatography [MASTER GC Fast Gas Chromatograph (DANI Instruments S.p.a., Italy)] [24] was used for the determination of major volatiles content.

HS-SPME GC/MS analysis [6890N GC, 5973NetworkedMS MSD (Agilent Technologies, USA)] [25] for the determination of minor volatiles content.

Aroma and taste of the produced wines were assessed as previously described [25].

2.7. Statistical Analysis

Statistical significance of oenological parameters was examined by 2-way analysis of variance. Duncan's multiple range test was used to determine significant differences ($P < 0.05$) among results. Statistical significance at $P < 0.05$, coefficients and ANOVA tables were computed by Statistica v.12.0 (Stat Soft Inc., Tulsa, OK, USA).

Principal Component Analysis was computed by XLSTAT 2015.1 (Addinsoft, Paris, France).

3. Results and Discussion

3.1. Yeast Strains Isolation and Characterization

Isolation of indigenous yeast strains was performed after the spontaneous alcoholic fermentations for each grape variety must were completed. Samples were plated on YPD agar and after incubation colonies were selected (from the dish with the greater dilution) based on their characteristics under microscope. Ten colonies, strongly resembling *Saccharomyces* species morphology, proceeded to molecular identification and further evaluation.

The results confirmed that all indigenous strains exhibited $\geq 99\%$ sequence similarity to *Saccharomyces cerevisiae*. The findings confirmed the dominance of *S. cerevisiae* among the yeast microbiota of the studied grape varieties and are consistent with previous studies emphasizing the prevalence of this species during spontaneous alcoholic fermentation [26,27].

All 10 selected indigenous strains showed resistance to SO_2 concentrations ranging 200-300 mg/L (Table 1), as previously documented for indigenous strains [8,28]. All new strains were capable of growing at 14% (v/v) ethanol with the adaptation rate ranging from "low" to "high", although even higher ethanol resistance has been documented in previous studies [7,28,29]. Regarding hydrogen sulfide production, all strains (with the exception of BO-Y1) exhibited high production.

Table 1. Technological characterization of indigenous *S. cerevisiae* strains.

Grape Variety	Strain Code	Ethanol Resistance (14% vol)	*SO ₂ Resistance	H ₂ S Production	Decarboxylation of Amino Acids			
					Arginine	Tyrosine	Tryptophane	Leucine
Karnachalas	KA-Y1	High	200	High	+	+	-	-
	KA-Y3	High	300	High	+	+	+	+
	KA-Y4	Low	300	High	+	+	+	+
	KA-YL2	Medium	300	High	+	+	+	+
	KA-YL4	Medium	300	High	+	+	+	+
Keratsouda	KE-Y1	Low	200	High	+	+	-	-
	KE-Y2	Low	300	High	+	+	-	+
Bogialamas	BO-Y1	Medium	200	Medium	+	+	+	+
	BO-YL1	Medium	300	High	+	+	+	+
	BO-YL2	High	250	High	+	+	+	+

*: Expressed as mg L⁻¹ of potassium metabisulfite; +/-: Positive/negative decarboxylation activity, respectively.

Amino acids are known to contribute to various yeast metabolic pathways. In some cases however, their decarboxylation by certain yeast strains may result to the production of biogenic amines and raise safety concerns for the consumer [30]. In our case, arginine and tyrosine were universally metabolized, as previously shown for indigenous strains [20]. Specifically, arginine is one of the most important sources for yeasts as it is involved in both anabolic and catabolic pathways [31] and tyrosine may be degraded in order to produce important aroma compounds, like fusel alcohols [32]. Tryptophan and leucine were also metabolized with different effectiveness rates among strains, while no response was acquired for histidine, lysine, and phenylalanine.

It is widely known that maintaining fresh microbial cultures for extended periods is a significant challenge for the industry. Thus, the storage of starter cultures between winemaking seasons is highly important and represents a viable and sustainable solution [33]. Microbial preservation mainly relies on application of drying techniques like freeze-drying, thermal-drying and others that offer protection against contamination, maintenance of cell viability through long preservation times, as well as storage and handling convenience [34].

In order to investigate the long-term cell viability of the newly isolated yeast strains, freeze-drying was applied and the freeze-dried cells were stored at ambient (20-25 °C), refrigerator (4 °C), and freezing conditions (-18 °C). In frequent time intervals (1, 3, 6, and 9 months), samples were analyzed and cell viability was determined. The results are summarized in Figure 1. As expected, low storage temperatures (4 °C & -18 °C) resulted in significantly higher ($P < 0.05$) survival rates compared to ambient temperatures, at all cases. Low temperatures are strongly preferred in order to effectively prolong microbial survival [35,36], but still a small drop in yeast counts was inevitable through storage duration, but not significant in all cases. Interestingly, survival rates documented (at 4 °C & -18 °C) ranged in levels similar or better than previously documented for freeze-dried yeast cultures [37].

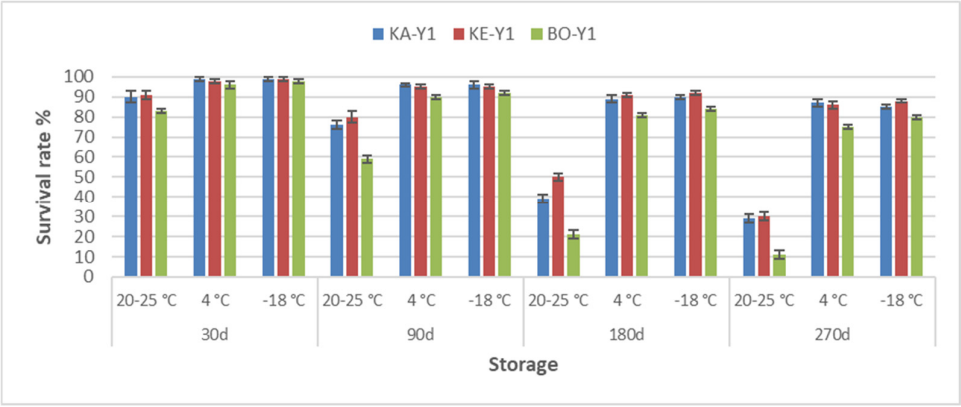


Figure 1. Survival rate % of freeze-dried indigenous *S. cerevisiae* strains during long-term storage at different conditions.

In any case, the survival of dried microorganisms during long-term storage is a dynamic process that depends on various factors such as microbial growth conditions, initial cell concentration, the drying medium, and more [38,39]. Protective agents have shown to further enhance survival of microorganisms [40], but carry-over effect and matters of cost-effectiveness advise against their use [22,41]. However, it should be noted that strain-specific characteristics that may affect the ability of microorganisms to withstand freeze-drying and storage cannot be excluded [42], even amongst different strains of the same species [33].

3.2. Fermentations

All new yeast strains were evaluated regarding their fermentation efficiency in batch fermentations at 22 °C and 12 °C. The temperature selection was based on the region’s climate during the winemaking season (October and November), according to the Hellenic National Meteorological Service [43]. Fermentation kinetic data and other important parameters are presented in Table 2.

Table 2. Kinetic parameters and organic acid profile of wines fermented by indigenous *S. cerevisiae* strains at 12 & 22 °C.

Fermentation Temperature	Yeast Strain Used	Acetaldehyde (mg L ⁻¹)	Ethyl Acetate (mg L ⁻¹)	1-Propanol (mg L ⁻¹)	Isobutanol (mg L ⁻¹)	1-Hexanol (mg L ⁻¹)	Amyl Alcohol (mg L ⁻¹)	Isoamyl Alcohol (mg L ⁻¹)	Methanol (mg L ⁻¹)
12 °C	KA-Y1	206±18a	73±6c	42±3c	81±6c,d	9±1c	71±5c	194±19b	25±3a
	KE-Y1	67±7d	80±6c	7±1d	54±5e	2±1d	49±3d	147±16c	9±1d
	BO-Y1	121±12c	71±8c	75±8a	112±10b	8±1c	102±9a	251±18a	19±2b
22 °C	KA-Y1	168±11b	111±12b	36±2c	158±13a	11±1b	96±7a,b	249±25a	23±3a
	KE-Y1	34±3e	127±11a	61±4b	86±9c	14±2a	84±8b,c	231±16a	15±2c
	BO-Y1	62±4d	31±2d	35±4c	64±4d,e	2±1d	50±5d	104±11d	6±1e

Significant differences ($P < 0.05$) are shown within each column with different letters.

Both the strain used and the fermentation temperature affected significantly ($P < 0.05$) the fermentation duration, the ethanol productivity, the residual sugars content and sugar conversion, the organic acids and glycerol concentration. Total acidity and volatile acidity values were significantly affected ($P < 0.05$) only by the strain used. Strong interactions ($P < 0.05$) were noted in all the above cases. On the contrary, no significant differences were observed for ethanol content, ethanol production yield, and pH values.

As expected, fermentations at low temperatures (12 °C) led to increased fermentation duration [22,24]. Significantly higher residual sugars occurred in all cases at 12 °C, ranging 17.9-38.2 g/L (in mean values), that later resulted in significantly lower conversion values comparing to 22 °C. Notably though, the higher residual sugars content is not considered a fault and may be utilized in order to provide a discreet flavor sweetness to the final product [44–46]. Ethanol content, however, remained unaffected by the low fermentation temperature and ranged in similar values 11.9–12.5% (v/v) in all cases. Ethanol productivities up to 10.5 g/(Ld) were noted in fermentations at 22 °C, but even at 12 °C the observed values were higher or similar to those usually found in traditional winemaking [47,48].

Glycerol is a fermentation-derived secondary metabolite, mainly known for contributing to the wine's flavor by adding sweetness and “fullness”, ultimately providing a “rounder” mouthfeel [49]. In our samples, glycerol content ranged 3.8–11.0 g/L, depending on the fermentation temperature and the strain used. In most cases however, intermediate values (5.7–7.0 g/L), typically encountered in fermentations with indigenous *S. cerevisiae* strains, were observed [8,9].

Regarding organic acids, acetic acid is produced naturally in wine as a result of fermenting microorganisms' metabolism [44]. In low concentrations (up to 0.3 g/L), it may add to the wine's complexity, but certain products (like Canadian ice wines) rely on high concentrations (often exceeding 2.0 g/L) for their special sensory character [50]. In our samples, acetic acid was detected in high quantities only in wines fermented with KE-Y1 strain. Nevertheless, differences (as well as accumulation) of acetic acid is not uncommon in fermentations performed by indigenous *S. cerevisiae* strains and has been previously associated to potentially diminished acetyl-CoA synthetase capabilities [51]. On the other hand, lactic acid was found in levels ranging 0.3-2.3 g/L, but also is widely known to be highly depended on the wine type, the occurrence of malolactic fermentation, the grape cultivar and the viticultural practices applied [52,53].

Total acidity and pH was found in values typical for wines [22,44,45]. On the contrary, elevated volatile acidities were detected in wines fermented with KE-Y1 strain, probably a direct consequence of high acetic acid content. According to OIV, volatile acidity should not exceed values of 1.2 g of acetic acid/L [54]. In special occasions though, high volatile acidity (> 2 g/L) is lawfully permitted and may be utilized as a tool by experienced winemakers in order to develop complex organoleptic attributes [55]. In any case, no fault (like vinegar taint or any other) was detected in our samples during the preliminary sensory evaluation.

3.3. Volatiles

3.3.1. Major Wine Volatiles

Major wine volatiles detected are shown in Table 3. Acetaldehyde, ethyl acetate, isobutanol, 1-hexanol, and methanol concentrations were significantly ($P < 0.05$) affected by both the indigenous strain used and the fermentation temperature. Amyl and isoamyl alcohols, and 1-propanol concentrations were significantly ($P < 0.05$) affected only by the strain used. Strong interactions ($P < 0.05$) were observed between the factors in almost all cases (with the exception of acetaldehyde).

Table 3. Major volatiles of wines fermented by indigenous *S. cerevisiae* strains at 12 & 22 °C.

Fermentation Temperature	Yeast Strain Used	Acetaldehyde (mg L ⁻¹)	Ethyl Acetate (mg L ⁻¹)	1-Propanol (mg L ⁻¹)	Isobutanol (mg L ⁻¹)	1-Hexanol (mg L ⁻¹)	Amyl Alcohol (mg L ⁻¹)	Isoamyl Alcohol (mg L ⁻¹)	Methanol (mg L ⁻¹)
12 °C	KA-Y1	206±18a	73±6c	42±3c	81±6c,d	9±1c	71±5c	194±19b	25±3a
	KE-Y1	67±7d	80±6c	7±1d	54±5e	2±1d	49±3d	147±16c	9±1d
	BO-Y1	121±12c	71±8c	75±8a	112±10b	8±1c	102±9a	251±18a	19±2b
	KA-Y1	168±11b	111±12b	36±2c	158±13a	11±1b	96±7a,b	249±25a	23±3a
22 °C	KE-Y1	34±3e	127±11a	61±4b	86±9c	14±2a	84±8b,c	231±16a	15±2c
	BO-Y1	62±4d	31±2d	35±4c	64±4d,e	2±1d	50±5d	104±11d	6±1e

Significant differences ($P < 0.05$) are shown within each column with different letters.

Acetaldehyde is one of the major wine constituents, which when found in low concentrations is known to provide fruit scents [56]. In wines fermented with the KA-Y1 strain, it was detected in significantly higher ($P < 0.05$) concentrations exceeding 200 mg/L. In general, its presence in concentrations up to 125 mg/L is acceptable in wines [57], while in higher levels it may be considered responsible for the “oxidised” [58] or “flat character” of the product [59]. Nevertheless, concentrations over 220 mg/L are also essential for the special organoleptic character found in products like fortified wines [59]. In any case, no implications were observed in our samples during the preliminary sensory evaluation (subsection 3.5).

Production of ethyl acetate is known to be affected by the fermentation temperature, as well as by the strain of *S. cerevisiae* used, with ‘natural yeast’ fermentations often tending to result in concentrations higher than average [60]. In all our cases however, ethyl acetate levels ranged between 31-127 mg/L (in mean values). Typically, ethyl acetate values up to 150 mg/L are considered to provide a positive fragrance complexity to the final products [44].

Higher alcohols (amyl alcohol, isoamyl alcohol, isobutanol, and 1-propanol) consist a group of compounds that is considered to have the strongest impact on the overall aroma of wine [61]. In low concentrations, they add to the product’s aromatic complexity by adding fruity notes [59,62], but on higher levels, they might confer a masking effect on the character of the wine [63]. It should be noted, however, that whether the overall contribution of higher alcohols is positive or negative is highly variable and depends heavily on the total aromatic context of the product [64]. In our samples, higher alcohols were found in high concentrations for all strains, but never exceeded 550 mg/L (in mean values) [59], and no unpleasant scents were detected.

Methanol and 1-hexanol; on the contrary, were detected in low concentrations < 50 mg/L in all wines.

3.3.2. Minor Volatiles

In total, 41 volatile compounds were identified in the wines fermented with the new *S. cerevisiae* strains. Both the indigenous strain used and the fermentation temperature affected significantly ($P < 0.05$) all the volatile groups of interest (esters, alcohols, organic acids, miscellaneous and total volatile content) and strong interactions ($P < 0.05$) were observed between the two factors in all cases (Table 4). Variations were detected among samples for all volatile compounds detected (Figure 2), potentially determining the final aromatic complexity of each wine.

Table 4. Minor volatiles of wines fermented by indigenous *S. cerevisiae* strains at 12 & 22 °C.

Fermentati on Temperatu re	Yeast Strain Used	Esters Conte nt (mg L ⁻¹)	Alcohols Content (mg L ⁻¹)	Organ ic Acids	Carbonyl Compounds Content (mg L ⁻¹)	Miscellaneous Compounds Content (mg L ⁻¹)	Total Volatiles (mg L ⁻¹)
				Conte nt (mg L ⁻¹)			
12 °C	KA-Y1	63.0c	35.8c	0.7c	19.4c	5.1a,b	124.0b
	KE-Y1	61.1c	42.1b,c	0.6c	11.0d	2.8d	117.6b
	BO-Y1	32.4d	12.8e	2.1a	6.8e	2.4d	56.5c
	KA-Y1	80.3b	43.7b	0.5c	25.7a	4.1c	154.4a
22 °C	KE-Y1	93.2a	26.7d	1.8b	21.7b,c	4.5b,c	147.9a
	BO-Y1	75.7b	54.5a	1.7b	23.2a,b	5.4a	160.6a

Scheme 0. are shown within each column with different letters.

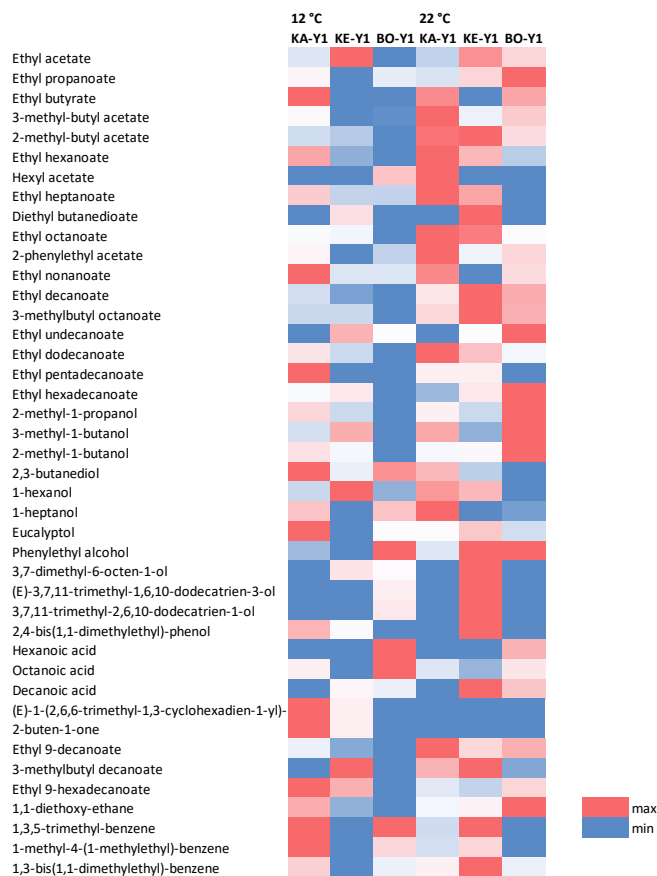


Figure 2. Heatmap of minor volatiles in wines fermented by indigenous *S. cerevisiae* strains at 12 & 22 °C. Differences in volatile concentrations are shown for each compound in blue (min) and red (max) shades. KA-Y1, wine fermented by *S. cerevisiae* strain isolated from “Karnachalas” grapes; KE-Y1, wine fermented by *S. cerevisiae* strain isolated from “Keratsouda” grapes; BO-Y1, wine fermented by *S. cerevisiae* strain isolated from “Bogialamas” grapes.

Esters were the largest group of volatiles detected, with different subcategories like acetate esters and ethyl fatty acid esters present in all samples. In general, esters are fermentation-derived compounds that originate enzymatically from higher alcohols or plain ethanol. Many of them are associated with fruity, floral, sweet, honey and other pleasing scents contributing to the wine aroma profile [65,66]. Fresh notes may be attributed to ethyl propanoate, while ethyl hexanoate, ethyl octanoate, ethyl decanoate, and 2-phenylethyl acetate usually add common fruity notes [67]. Distinct banana-like scents, peer-like scents or apple-peel notes may be attributed to 3-methylbutyl acetate, 2-methylbutyl acetate, and ethyl butyrate, respectively. On the contrary, compounds like ethyl dodecanoate may be responsible for more smokey, toasty and earthy aromas resulting in the dry character of the final product [44,68–70]. In any case, it should be noted that excessive ester amounts may produce pungent odor and foul taste and impact negatively the wine quality.

Alcohols were the second most abundant group of volatile compounds identified. Key compounds like phenylethyl alcohol and 1-hexanol were present in all samples contributing mostly rose aroma and floral-fruity scents, respectively. Similarly, citronellol (adding citrus and sweet notes) was identified in most cases [71–73]. Eucalyptol, characterized by a distinct green aroma, was found in all samples and is believed to be adding “fresh”, and “cool” notes to the product [74]. On the other hand, compounds like 2,3-butanediol, known for adding bitter-sweet notes [44,75] and 1-heptanol (present in most of the samples), may be considered to be of low to none sensory importance for wine, but their presence has been previously associated with the discriminant classification of regional wines [76].

Regarding organic acids, octanoic acid was detected in all samples. Hexanoic and decanoic acid were also detected in low concentrations in most of the wines. Organic acids are mainly known for their low odor threshold limit and their potential impact on wine flavor by contributing to the product's "freshness" [77,78]. Except for their aromatic contributions, in the case of increased concentrations the synthesis of their corresponding esters might also be favored [78].

A few carbonyl compounds were identified in some of the samples. Aldehydes and ketones are produced naturally during fermentation or occur as oxidation byproducts, although in many cases their exact source of origin is difficult to discern [79]. They present low odor thresholds and are known to enhance aroma and flavor complexity [80].

Some miscellaneous compounds were also observed in most of the samples. 1,1-Diethoxyethane, which may indeed contribute to the wine bouquet [44,69], was detected in all cases and is primarily known for its fruity-green notes and refreshing character.

All results derived through Headspace SPME GC/MS were also subjected to Principal Component Analysis. According to the analysis (Figure 3), the fermentation temperature rather than the indigenous yeast strains affected significantly the volatile composition in the new wines. Wines produced at 22 °C were located towards the right half of the plot, while wines produced at 12 °C were located at the left half. Interestingly, wines produced with KA yeast strain were grouped in the middle of the plot and were composed by compounds correlated mostly with PC2.

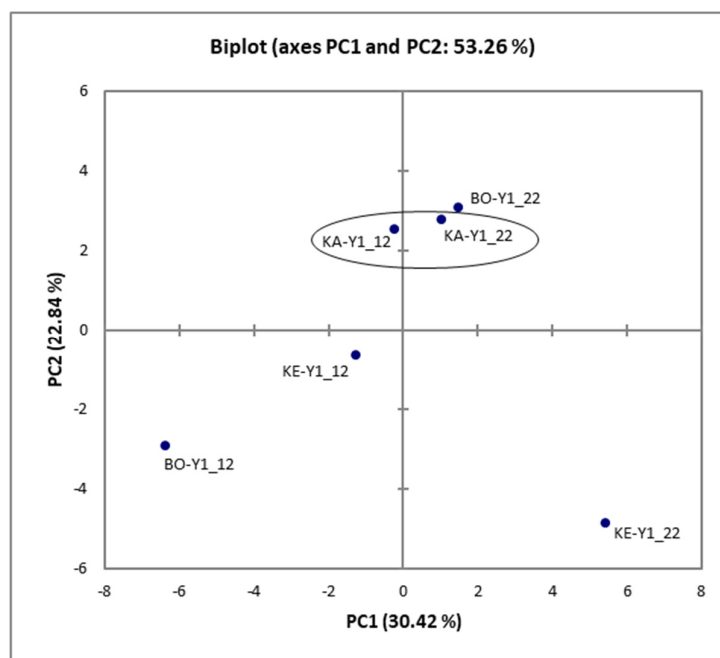


Figure 3. PCA plot of minor volatiles isolated from wines fermented with indigenous yeast strains at 12 & 22 °C. KA-Y1, wine fermented by *S. cerevisiae* strain isolated from "Karnachalas" grapes; KE-Y1, wine fermented by *S. cerevisiae* strain isolated from "Keratsouda" grapes; BO-Y1, wine fermented by *S. cerevisiae* strain isolated from "Bogialamas" grapes. The fermentation temperature is indicated after each sample code.

3.4. Preliminary Sensory Evaluation

All wines fermented with indigenous *S. cerevisiae* strains were evaluated for their sensory aroma and flavor characteristics. According to the results (data not shown), all products were characterized as light-bodied with a predominant wine-like aroma, a mostly sour taste, and a mild aftertaste. A strong aftertaste, however, was noted in wines fermented with KA-Y1 strain, possibly as a result of increased acetaldehyde levels. Nevertheless, the flavor outcome was not unpleasant, but characterized as interesting and piquant-like mostly contributing to the character of the product. Notably, no vinegar or other repulsive scents were detected in any of the wines produced with KE-

Y1 strain (or any other strain), despite the significantly high ($P < 0.05$) content of acetic acid and high volatile acidity values.

3.5. Technological Considerations and Current Limitations

In this work, indigenous *S. cerevisiae* strains were isolated from local grape varieties (“Karnachalas”, “Keratsouda”, and “Bogialamas”) and subsequently used to ferment the grape must (of their origin). Technological advantages derived from this strategy include, among others, the enhanced strain adaptability to each must’s conditions, better regulation of the native microbial flora and improved control of alcoholic fermentation. At the same time, the utilization of lower fermentation temperatures for enhancing the production of aromatic compounds and improving the wine quality is well known between experienced winemakers. These combined actions aim to showcase the unique regional terroir characteristics in the final product [81–83]. Thus, our findings offer local winemakers the opportunity to create and promote ‘signature wines’ by following the example of several Spanish wineries that have already obtained the ‘Pago’ designation. This term constitutes a form of Spanish Geographical Indication [9,84] and is used to describe wines coming from distinct regions with specific soil and microclimate characteristics (like Soufli), produced with traditional vineyard and winemaking techniques and most importantly fermented by the indigenous yeast strains each winery has in possession [85,86].

However, it should be noted that the proposed methodology cannot be adopted uncritically. Future experiments should primarily focus on confirming and replicating the obtained results. Further isolation of new native yeasts able to produce wines with more typical wine characteristics (e.g. lower volatile acidity values) should also be a priority, as a single winemaking season is not sufficiently representative of the region’s terroir. Prioritizing the effective combination of yeast strains for enhanced aroma development, along with the isolation and inoculation of malolactic bacteria, is essential. Innovative technologies such as immobilization on natural supports should also be examined, as they have been shown to boost culture survival during storage, enhance fermentation activity, and contribute positively to the wine’s character [22,23,41,87]. Apart from the above, current findings may be used as a roadmap and provide useful insights for the targeted isolation of native yeasts (from different grape varieties) across other geographical regions. Advanced omics methodologies may be applied in a later stage in order to combine all available data in an effort to predict the functional outcomes of native fermentations. In any case, the detailed technological characterization of the indigenous microbial flora and the aromatic profile of the wines produced (in each grapevine region, like Soufli) is the vital step in order to facilitate positive outcomes not only for the local winemakers, but also for the region, the consumers and wine-industry in general.

4. Conclusions

In the current study, 10 indigenous *S. cerevisiae* strains (ITS sequence similarity $\geq 99\%$) were isolated from local grape varieties in the Soufli region in Northern Greece. All strains exhibited satisfactory ethanol resistance (up to 14% vol), variable SO_2 resistance (200–300 mg/L), predominantly high H_2S production, and distinct amino acid decarboxylation profiles. Long-term survival of 3 freeze-dried selected strains was examined and rates $\geq 75\%$ even after 9 months at low storage temperatures (4 & -18°C) were recorded. Fermentation parameters were significantly influenced by both the strain used and the fermentation temperature (12 or 22°C), but high ethanol content [11.9–12.5% (v/v)] and ethanol productivities higher or similar to those usually found in traditional winemaking were observed. PCA revealed that the fermentation temperature rather than the indigenous yeast strains affected significantly the volatile composition in the new wines resulting in unique sensory profiles. In conclusion, the utilization of indigenous *S. cerevisiae* strains could present a viable strategy for the preservation and promotion of the regional terroir, enabling the production of high quality “signature wines” in various temperatures.

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