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SACCHAROMYCES CEREVISIAE BIODIVERSITY IN MONFERRATO, NORTH WEST ITALY, AND SELECTION OF INDIGENOUS STARTER CULTURES FOR BARBERA WINE PRODUCTION

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(Article begins on next page)

1 ***Saccharomyces cerevisiae* biodiversity in Monferrato, North West Italy and selection of**
2 **indigenous starter cultures for Barbera wine production**

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ABSTRACT

19 The aim of this study was to examine the biodiversity of *Saccharomyces cerevisiae* isolates from
20 Barbera grapes and musts, from the Monferrato area, in the Piedmont region – North West Italy. An
21 interdelta element PCR analysis was used to identify and discriminate 636 *S. cerevisiae* isolates at a
22 strain level. Ninety-six *S. cerevisiae* that showed different molecular fingerprints were characterized
23 through physiological tests and laboratory scale fermentations. A chemical analysis of experimental
24 wines obtained from inoculated fermentations showed significant differences between the wines.
25 The main variables considered in the strain differentiation were the residual sugars and the
26 production of acetic acid, which ranged from 148.64 to 3.44 g/l and from 0.20 to 0.60 g/l,
27 respectively. As a consequence, strain variability should be considered as a relevant resource to
28 select suitable starter cultures in order to improve or characterize wines with a close bond to the
29 geographic region.

30

31 **Keywords:** *Saccharomyces cerevisiae*, Yeast biodiversity, indigenous starter, Interdelta PCR,
32 Selection.

33

35 INTRODUCTION

36 Wine production is an ancient tradition that has been carried out for centuries through the
37 spontaneous fermentation of grape juice, which takes place due to the presence of indigenous yeasts
38 from different genera and species (FLEET, 2003; PRETORIUS, 2000; ROMANO *et al.*, 2003). The
39 number of species and their presence during fermentation depends on several factors (PRETORIUS
40 *et al.*, 1999), which lead to subsequent wine quality variations from region to region, but also from
41 one year to another, and all this makes the outcome of spontaneous fermentation difficult to predict
42 (PRETORIUS, 2000). In an attempt to address this issue, many winemakers have used pure
43 commercial *Saccharomyces cerevisiae* cultures inoculated into the must (PRETORIUS, 2000).
44 However, it has been suggested that native *S. cerevisiae* strains are better suited to the micro-area
45 climatic conditions of the wine production region (LOPES *et al.*, 2002) and can therefore more
46 easily dominate the natural biota (LA JEUNE *et al.*, 2006).

47 *S. cerevisiae*, the most relevant species in winemaking, is usually chosen as the wine yeast, and the
48 particular strain is chosen according to a set of physiological features that are indicative of their
49 potential usefulness for industrial wine production. In addition to the primary end products of the
50 glycolytic fermentation of glucose and fructose, certain oenological criteria must be considered in
51 order to select yeast strains that show desirable characteristics, including: tolerance and high ethanol
52 production, exhaustion of the sugar in must and high fermentation activity, growth at high sugar
53 concentrations, high glycerol production, resistance and low sulphur dioxide production, good
54 enzymatic profile (high β -glucosidase and proteolytic activities) and low acetic acid formation
55 (ESTEVE-ZARZOSO *et al.*, 2000).

56 At present, there is increasing interest, in the wine community, in the use of indigenous *S.*
57 *cerevisiae* strains that may contribute to the overall sensorial quality of wine and reflect the
58 characteristics of a given region, even in guided fermentations using selected *S. cerevisiae* starter
59 cultures (CAPECE *et al.*, 2010; SUZZI *et al.*, 2012). Recently, in an attempt to respond to these
60 aspects coupled with the current emphasis on the preservation of all forms of genetic biodiversity,
61 some research groups have focused on the selection of yeasts from restricted areas (SETTANNI *et*
62 *al.*, 2012; FRANCESCA *et al.*, 2009; ORLIC *et al.*, 2007; LOPES *et al.*, 2007).

63 We have previously extensively studied the indigenous mycobiota originating from the Barbera
64 grapes from the Monferrato area, Piedmont region, North-West Italy (ALESSANDRIA *et al.*,
65 2015). Barbera grapes produce a ruby-red coloured wine with berry, cherry, plum and spicy
66 flavours, depending on the clone, as well as the location and the age of the plant (BOSSO *et al.*,
67 2011). In this study, *S. cerevisiae* isolates were characterized to establish their genetic and
68 technological variability in order to contribute to the preservation of the *S. cerevisiae* genetic
69 resources of the Barbera of Monferrato *terroir*. Interdelta-PCR was used to help establish the
70 genetic diversity of the isolates. Physiological tests, which focused on the production of
71 extracellular hydrolytic enzymes and on their growth in different ethanol and total SO₂
72 concentrations, were then conducted. Finally, selected genotypes were used to ferment Barbera
73 must, during micro-fermentation trials, in order to evaluate their fermentation potential.

74

75 MATERIAL AND METHODS

76 *Fermentation set up and yeast isolation*

77 Spontaneous fermentations were conducted using *Vitis Vinifera* L. Barbera grapes obtained from
78 fifteen different vineyards (Fig. 1), located in five areas in the Asti and Alessandria districts of the
79 Piedmont region (ALESSANDRIA *et al.*, 2015). The vineyards, which were identified on the basis
80 of their geographical locations, were: 1 (Murisengo), 2 (San Martino Alfieri), 3 (Costigliole d'Asti),
81 4 (Isola d'Asti), 5 (Montegrosso d'Asti), 6 (Agliano Terme), 7 (Vinchio), 8 (Nizza Monferrato), 9

82 (Incisa Scapaccino), 10 (Loazzolo), 11 (Ricaldone), 12 (Alice bel colle), 13 (Acqui -Terme Crocera
83 south west zone), 14 (Acqui Terme - Crocera south est zone) and 15 (Acqui Terme – Dannonna
84 zone).

85 Approximately 25 kg of healthy grapes, without signs of bird damage or *Botryotinia fuckeliana*
86 infection, were harvested. The grapes were crushed in the laboratory and the obtained juice (about
87 15 L volume) was transferred to sterile jugs where it underwent spontaneous fermentation at room
88 temperature (between 22 and 25°C). Yeasts were isolated from each container at the beginning of
89 the fermentation (after 1 day and 3 days), in the middle (after 7 days) and when alcoholic
90 fermentation was completed. The alcoholic fermentation was monitored with a densitometer.
91 Aliquots (0.1 mL each) of several decimal dilutions, in a 0.1% ringer solution (Oxoid, Milan, Italy),
92 were plated on Wallerstein Laboratory Nutrient (WLN) medium (Oxoid) (Pallmann et al., 2001).
93 The plates were incubated at 28°C for 5 days. WLN allows the presumptive identification of the
94 yeast species according to the colony morphology and colour (URSO *et al.*, 2008). At least 10
95 colonies were selected from each sample and at each fermentation stage and were isolated on WLN;
96 priority was given to putative colonies of *Saccharomyces* spp. The isolates were stored at -80 °C in
97 YPD broth (1% (w/v) yeast extract, 2% (w/v) peptone and 2% (w/v) glucose, all obtained from
98 Oxoid) after the addition of glycerol (30%, v/v) (Sigma-Aldrich, Milan, Italy).

99 *Yeast identification*

100 The putative *Saccharomyces* spp. isolates were subsequently identified and simultaneously
101 differentiated at a strain level on the basis of a PCR interdelta element analysis (δ -PCR). In order to
102 conduct the δ -PCR analysis, the total DNA was extracted from 1 millilitre of an overnight culture in
103 YPD broth, according to COCOLIN *et al.* (2000), quantified using a Nanodrop ND-1000
104 spectrophotometer (Celbio, Milan, Italy) and standardized at 100 ng/ μ l. The delta12 (5'-
105 TCAACAATGGAATCCCAAC-3') and delta21 (5'-CATCTTAACACCGTATATGA-3')
106 oligonucleotide primers were used to amplify regions between the repeated interspersed delta
107 sequences (Legras and Karst, 2003). Amplification reactions were performed with a PTC-200 DNA
108 Engine MJ Research thermal cycler (Biorad, Milan, Italy) using the following programme: initial
109 denaturation at 95°C (5 min), 35 cycles of denaturing at 94°C (1 min), annealing at 50°C (1 min),
110 extension at 72°C (1 min) and a final extension at 72°C (10 min). The PCR products were separated
111 in 1.5% agarose gels and stained with ethidium bromide. The resulting fingerprints were analyzed
112 by means of the BioNumerics V4.0 software package (Applied-Maths, Sint-Martens-Latem,
113 Belgium). Similarity among the digitized profiles was calculated using the Pearson correlation, and
114 an average linkage (UPGMA) dendrogram was derived from the profiles. A coefficient of
115 correlation of 85% was arbitrarily selected to distinguish the clusters. The yeasts that were not
116 amplified with δ -PCR, were subsequently identified by using the RFLP of the ribosomal region
117 method as described in ALESSANDRIA *et al.* (2015).

118 *Physiological characterization*

119 *Hydrogen sulphite production*

120 The ability to produce hydrogen sulphite was determined by streaking single colonies onto Biggy
121 agar (Oxoid) and incubating them at 25°C for 48– 72 h. Colony colour was observed and scored as
122 being white, pale hazel, hazel, dark hazel or black.

123 *Enzymatic activities*

124 The esterase, protease and β -glucosidase activities of the isolates were screened as described by
125 ENGLEZOS *et al.* (2015).

126 *Ethanol and SO₂ tolerance assays*

127 Ethanol tolerance and SO₂ tolerance were determined in microplates, according to the method
128 proposed by ARROYO-LOPEZ *et al.* (2010) and TOFALO *et al.* (2012), with some modifications.
129 Yeast Nitrogen Base with amino acids (YNB, 6.7 g/L, [Remel, Lenexa, KS, USA]) and pH 5.5 was
130 supplemented with 20 g/L of glucose and sterilized by filtration using a 0.2 µm membrane filter
131 (VWR, Milan, Italy). The medium was supplemented with different concentrations of ethanol
132 (Sigma) (final concentrations of 0, 12, 14 and 16% v/v) in order to test for ethanol tolerance, while,
133 in order to establish SO₂ resistance, different amounts of total SO₂ were added (after adjustment to
134 pH 3.0) until final concentrations of 0, 50, 100 and 150 mg/L. Cells for the inoculation were
135 prepared from an overnight culture in 1 mL of YPD medium, centrifuged at 9000 rpm for 10 min.
136 The obtained pellet was washed twice in a sterile salt solution (8 g/L NaCl) and then re-suspended
137 in the same solution to obtain a concentration of about 10⁶ cells/mL. The diluted cells (20 µL) were
138 mixed with 180 µL of YNB, prepared as above. The microplates were incubated at 25 °C and the
139 optical density (OD) was measured at 600 nm using a microtiter plate reader (Savatec Instruments,
140 Torino, Italy) at 24 and 48 hours after an orbital shaking of 30 s, in order to re-suspend the cells in
141 the medium before the measurement. YNB without ethanol or SO₂ was used as the control. Cell
142 growth was determined on the basis of the ratio (%) of OD values obtained in medium with and
143 without ethanol or SO₂ for the specific incubation times. Tests were carried out in triplicate. Isolates
144 with a percentage of growth < 10% were considered sensitive.

145 *Microfermentations*

146 The fermentation potential of the different genotypes was evaluated in microfermentation trials.
147 These were carried out in duplicate for 14 days, in 50 mL tubes containing 25 mL of Barbera grape
148 must (120 g/L glucose, 124 g/L fructose, 5.25 titratable acidity as g/L of tartaric acid, pH 3.20 and
149 184 mg/L yeast available nitrogen (YAN)). Before the inoculation, the must was thermally treated
150 at 60 °C for 50 min, and the absence of viable populations was evaluated by plating 100 µL of the
151 must after the treatment on the WLN medium, followed by incubation at 28 °C for 5 days. Pre-
152 cultures were prepared in must at 25 °C for 24 h, and then used to inoculate each fermentation with
153 a cell concentration of 10⁶ per mL, which was determined through a microscopical cell count. The
154 fermentations were carried out under static conditions at 25 °C.

155

156 *Chemical analysis*

157 After 14 days of fermentation, the sugar consumption (glucose and fructose) and the ethanol,
158 glycerol and acetic acid production were evaluated directly by means of HPLC, according to the
159 method proposed by ROLLE *et al.* (2012). YAN was measured using the protocols reported in
160 ENGLEZOS *et al.*, 2016).

161 *Data analysis*

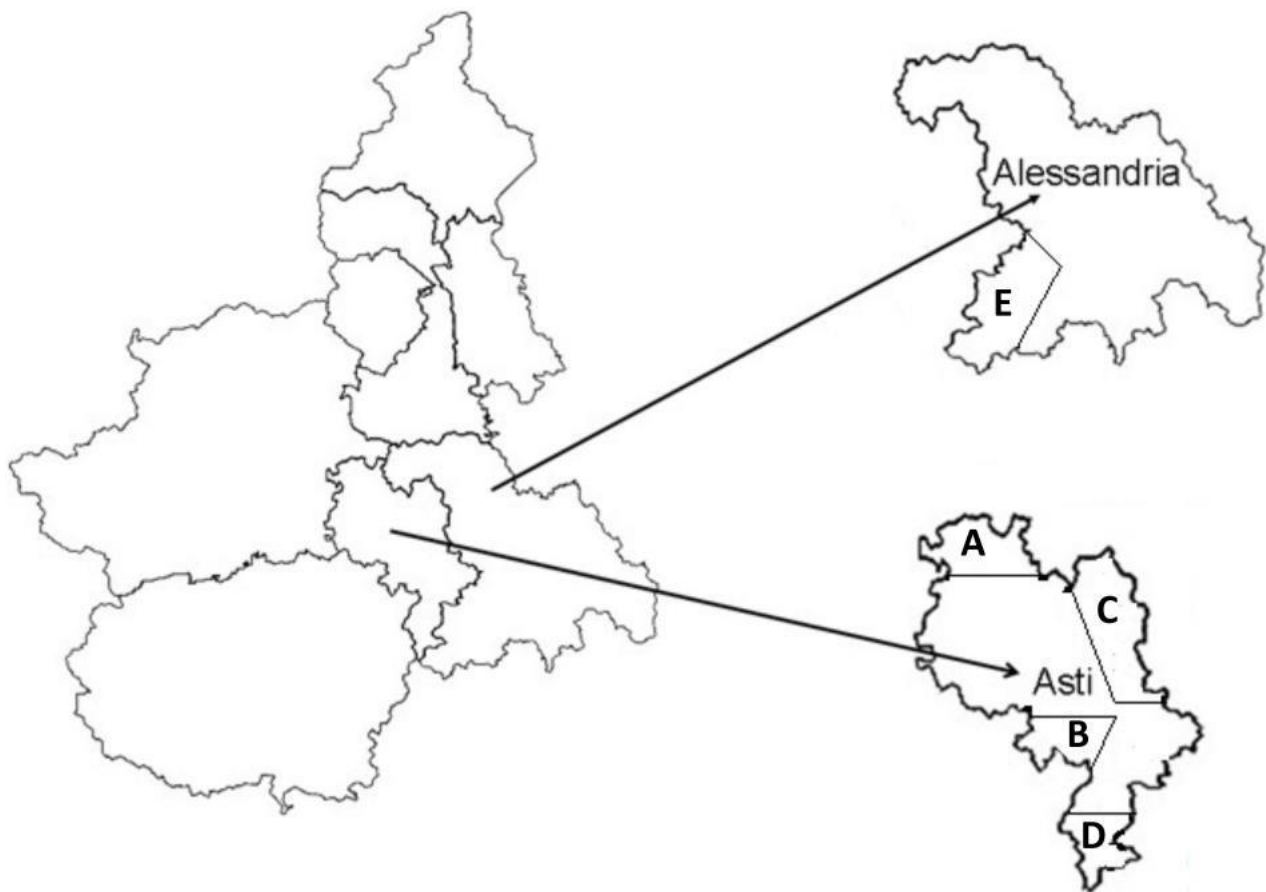
162 The results of the chemical composition of the wines obtained from the micro-fermentation trials
163 were subjected to a Principal Component Analysis (PCA), in order to evaluate the intraspecific
164 biodiversity of *S. cerevisiae* isolates. Statistical analyses were performed using the IBM SPSS
165 Statistics software package (version 19.0, IBM Corp., Armonk, NY, USA).

166

167 **RESULTS AND DISCUSSION**

168 The possible association between territory and yeasts is being actively investigated in recent years
169 and is believed to have a positive impact and influence purchase decision-making by the consumer.
170 The use of indigenous selected yeasts could represent a useful alternative to spontaneous
171 fermentation in order to optimize the typical attributes of the grape variety (CLEMENTE JIMENEZ
172 *et al.*, 2004; REMENTERIA *et al.*, 2003; ROMANO *et al.*, 2008). The goal of this study was to
173 isolate and characterize indigenous *S. cerevisiae* yeasts present on Barbera grapes in the vineyards
174 of the Monferrato area.

175 Fifteen vineyards, located in the Piedmont region (Fig. 1) and cultivated with the Barbera grape
176 variety, were studied during the 2012 harvest season and the collected grapes were crushed to
177 obtain 15 spontaneous alcoholic fermentations. Overall, 943 yeast colonies were isolated during the
178 fermentations, and after molecular identification, most of them (636 isolates) were identified,
179 through the use of δ -PCR, as *S. cerevisiae*. Other species, namely *Hanseniaspora uvarum* (248
180 isolates), *Starmerella bacillaris* (synonym *Candida zemplinina*) (11 isolates), *Pichia anomala* (7
181 isolates) and *Torulasporea delbruecki* (7 isolates) were also isolated, mainly at the beginning and
182 middle of the fermentations.



183

184 **Figure 1.** Geographic location of the fifteen vineyards in the Monferrato region (Asti and
185 Alessandria) with indication of the 5 areas considered for the population genetic analysis reported in
186 Table 2. The distribution of the vineyards in the five areas was as follows: A (Murisengo), B
187 (Montegrosso d'Asti, Costigliole d'Asti, San Martino Alfieri, Agliano Terme, Isola d'Asti), C
188 (Vinchio, Nizza Monferrato, Incisa Scapaccino), D (Loazzolo) and E (Ricaldone, Aquis Terme,
189 Alice bel colle).

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192 The PCR amplification of the δ interspersed sequences was also used to identify the genetic
193 differences among the *S. cerevisiae* isolated during the fermentations. The molecular fingerprinting
194 analysis, using a coefficient of similarity of 85%, allowed numerous strains among the isolates to be
195 distinguished. The dendrogram resulting from the analysis of 636 *S. cerevisiae* isolates highlighted
196 the presence of 62 clusters and 37 strains, which were unique and did not cluster with any other
197 isolate. The most numerous clusters were: XVII and XLI with 60 and 44 isolates, respectively (Tab.
198 1). It is interesting to notice the different level of heterogeneity of *S. cerevisiae* isolated from the
199 different vineyards. For example, most of the isolates from vineyard 1 grouped in one single cluster
200 (XLI), while in other cases (vineyards 12, 13 and 14) a high level of diversity was observed. Only
201 37 *S. cerevisiae* δ -PCR patterns were unique, demonstrating a feeble biodiversity of indigenous *S.*
202 *cerevisiae* strains in Monferrato area. Considering the ratio between the number of *Saccharomyces*
203 isolated and the number of observed patterns, as an approximate biodiversity estimation, our results
204 showed similar values to those found in Portugal (Schuller et al., 2005) and in France (VALERO *et*
205 *al.*, 2007).

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Table 1. Clusters obtained from a comparison of the different fingerprinting profiles of the *S. cerevisiae* isolates examined in this study with/by means of the molecular technique. The arbitrarily selected coefficient of similarity was 85%. The table shows their composition according to the geographical locations (vineyard) from which the isolates were obtained.

Cluster	Number of strains in the cluster	Monferrato's vineyards														
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
I	5	/	/	/	/	/	/	/	/	/	/	5	/	/	/	/
II	15	/	/	/	/	11	/	4	/	/	/	/	/	/	/	/
III	14	/	/	6	/	/	/	4	/	/	/	2	/	2	/	/
IV	6	/	/	6	/	/	/	/	/	/	/	/	/	/	/	/
V	3	/	3	/	/	/	/	/	/	/	/	/	/	/	/	/
VI	6	/	/	/	/	/	/	/	/	/	2	/	/	/	3	1
VII	2	/	/	/	/	/	/	/	/	/	/	/	/	/	/	2
VIII	24	/	14	5	/	/	/	/	2	/	/	/	/	1	/	2
IX	4	/	4	/	/	/	/	/	/	/	/	/	/	/	/	/
X	7	/	/	/	/	3	/	/	/	/	/	4	/	/	/	/
XI	10	/	/	/	/	/	/	/	/	3	/	/	/	7	/	/
XII	8	/	/	/	/	/	/	/	/	4	/	/	/	4	/	/
XIII	7	/	7	/	/	/	/	/	/	/	/	/	/	/	/	/
XIV	18	/	4	/	/	10	/	/	/	/	/	1	/	2	1	/
XV	4	/	/	/	/	/	2	/	/	/	/	/	/	2	/	/
XVI	2	/	/	/	/	/	/	/	/	/	1	/	/	/	1	/
XVII	60	6	4	3	2	1	/	3	8	/	2	/	2	5	9	15
XVIII	6	/	/	/	/	/	/	2	2	/	/	/	/		2	/
XIX	10	/	/	4	/	/	/	/	/	/	/	/	/	4	/	2
XX	15	1	/	/	9	2	/	/	/	/	/	/	/	/	2	1
XXI	13	/	/	/	2	2	/	/	2	/	2	/	/	2	2	1
XXII	28	2	/	1	/	/	/	/	1	/	/	/	1	1	4	18
XXIII	2	/	/	1	/	/	/	/	/	/	/	/	/	1	/	/
XXIV	3	/	/	/	/	/	/	/	/	/	3	/	/	/	/	/
XXV	3	/	/	/	/	/	/	/	/	/	3	/	/	/	/	/
XXVI	4	/	/	/	/	/	/	/	/	/	/	/	/	/	/	4
XXVII	5	/	/	3	/	/	/	/	/	/	/	/	/	/	1	1
XVIII	2	/	/	/	/	/	/	/	/	/	/	/	/	2	/	/
XXIX	10	/	/	/	/	/	/	/	/	/	/	/	/	10	/	/
XXX	6	/	/	/	/	/	/	/	/	/	/	/	/	3	3	/
XXXI	7	/	/	/	/	/	/	/	/	/	3	/	/	3	/	1
XXXII	3	1	/	/	/	/	/	/	/	/	2	/	/	/	/	/
XXXIII	6	/	/	/	/	/	/	/	/	/	/	/	5	1	/	/
XXXIV	5	/	/	/	/	/	/	/	/	/	/	/	5	/	/	/
XXXV	3	/	/	/	1	/	/	/	/	/	/	/	/	2	/	/
XXXVI	5	/	/	/	/	/	5	/	/	/	/	/	/	/	/	/
XXXVII	9	/	/	/	2	/	/	/	2	/	4	/	/	/	/	1
XXXVIII	4	/	/	/	/	/	/	/	/	/	/	/	/	/	/	4

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251 **Table 1. Continued**

Cluster	Number of strains in the cluster	Monferrato's vineyards														
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
XXXIX	2	/	/	/	/	/	/	/	2	/	/	/	/	/	/	/
XL	7	/	/	/	/	/	/	/	/	/	/	/	/	7	/	/
XLI	44	38	/	/	/	/	/	/	/	/	/	/	3	3	/	/
XLII	7	/	/	/	/	/	2	/	1	4	/	/	/	/	/	/
XLIII	9	/	/	5	/	/	/	/	/	1	/	/	2	/	1	/
XLIV	7	/	/	/	/	/	/	/	3	3	1	/	/	/	/	/
XLV	6	/	/	/	/	/	/	/	/	/	/	/	/	6	/	/
XLVI	5	/	/	/	/	/	/	5	/	/	/	/	/	/	/	/
XLVII	10	/	/	/	9	/	/	/	/	/	/	/	/	1	/	/
XLVIII	15	/	/	/	/	/	/	/	/	/	/	13	2	/	/	/
XLIX	6	/	/	/	/	2	4	/	/	/	/	/	/	/	/	/
L	10	/	3	/	/	/	/	/	6	/	1	/	/	/	/	/
LI	4	/	/	/	/	/	/	/	/	3	/	1	/	/	/	/
LII	30	/	/	/	/	2	16	10	/	/	/	/	/	2	/	/
LIII	12	/	/	/	/	2	/	7	/	/	/	/	1	/	2	/
LIV	2	/	/	/	/	/	/	/	/	/	/	/	/	/	2	/
LV	10	/	/	/	/	/	/	/	/	/	/	/	/	/	/	10
LVI	5	/	/	/	/	3	2	/	/	/	/	/	/	/	/	/
LVII	19	2	/	/	/	2	/	/	1	/	/	/	/	6	8	/
LVIII	15	/	/	/	/	3	/	/	/	/	2	/	10	/	/	/
LIX	25	/	/	/	/	/	5	7	/	/	/	/	1	/	12	/
LX	22	/	/	/	/	/	/	/	1	/	7	/	/	12	1	1
LXI	3	/	/	/	/	/	/	/	/	/	1	/	/	2	/	/
LXII	17	/	3	/	/	/	/	/	1	/	/	/	4	8	/	1

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In order to investigate further the *S. cerevisiae* diversity, 96 strains were selected on the basis of the cluster analysis, and screened for desirable oenological characteristics (ESTEVE-ZARZOSO *et al.*, 2000) such as: a low production of hydrogen sulphide and tolerance to a final concentration of 150 mg/L total sulphur dioxide and 16% (v/v) of ethanol (Supplementary Table 1). All the strains exhibited a medium hydrogen sulphide production level; 5% of them appeared to be pale hazel on Biggy agar, while the others were hazel. Concerning the results of the tolerance to SO₂, the selected strains were able to grow in the presence of 50 and 100 mg/L of SO₂ (83% and 60% of the isolates, respectively), while only a few isolates (32%) grew at 150 mg/L of SO₂ within 24 h. Extending the incubation time to 48 h, the number of the isolates that were able to grow at 150 mg/L of SO₂ increased to 63%. Only one strain (ScBa20) was totally inhibited by SO₂. As far as ethanol tolerance is concerned, 60% of the strains grew at 14% v/v within 24 h. Ethanol mainly affected yeast growth by increasing the lag phase, and this evidence explains why after increasing the incubation time to 48 h, 95% of the strains were able to grow in all the ethanol concentrations (Suppl. Tab. 1). β -glucosidase activity was found in only 2.7% of the strains, thus indicating possible production and activity during the fermentation. Protease activity was detected in 37.8% of the tested *S. cerevisiae*, while 21.6% were able to hydrolyse esters (data not shown).

In order to extend the information on the 96 *S. cerevisiae* strains of the Monferrato area, alcoholic fermentations were carried out in Barbera grape must. The experimental wines obtained were analyzed to establish the content of some by-products correlated to the organoleptic quality of the

274 wine and the obtained results are reported in Table 2. The values of the residual sugars ranged from
 275 3.44 to 148.64 g/L. Only seven isolates (ScBa4, ScBa5, ScBa13, ScBa26, ScBa44, ScBa60 and
 276 ScBa63) were able to leave less than 5 g/L of residual sugars after 14 days of fermentation. All the
 277 strains, except ScBa12, ScBa47, ScBa48 and ScBa57, were capable of consuming almost all the
 278 glucose of the must, confirming the glucophylic character of this species. All the yeast strains, that
 279 completed the fermentation, formed low amounts of acetic acid in the wines (less than 0.6 g/L).
 280 Glycerol production was relatively low, ranging from 5.20 to 7.86 g/L. The fermentation purity was
 281 high; most strains had a low ratio between the produced acetic acid and ethanol (range 0.01-0.09)
 282 and only three strains showed values above 0.04. As regards ethanol production, 52% of the strains
 283 produced more than 13%, and 6% of them managed to develop more than 14%.
 284

285 **Table 2.** Chemical analysis of the wines obtained from the fermentation of the pure indigenous *S.*
 286 *cerevisiae* cultures. The data are means \pm standard deviations. With * were reported strains which
 287 were unique and did not cluster with any other isolate.

Strains	Vineyard	Cluster	Residual glucose (g/L)	Residual fructose (g/L)	Glycerol (g/L)	Acetic acid (g/L)	Ethanol (%v/v)	Fermentat ion purity ^a	Ethanol yield ^b	Glycerol yield ^c	Acetic acid yield ^a
ScBa1	11	I	2.18 \pm 0.54	24.23 \pm 2.14	7.00 \pm 0.15	0.45 \pm 0.00	12.57 \pm 0.19	0.036 \pm 0.001	0.058 \pm 0.001	0.032 \pm 0.001	0.0359 \pm 0.0008
ScBa2	13	*	0.88 \pm 0.07	8.68 \pm 0.80	6.86 \pm 0.05	0.51 \pm 0.02	13.77 \pm 0.05	0.037 \pm 0.001	0.059 \pm 0.001	0.029 \pm 0.001	0.0367 \pm 0.0012
ScBa3	5	II	0.65 \pm 0.08	4.43 \pm 1.04	7.01 \pm 0.01	0.56 \pm 0.01	14.11 \pm 0.11	0.039 \pm 0.001	0.059 \pm 0.001	0.029 \pm 0.001	0.0395 \pm 0.0005
ScBa4	7	*	0.68 \pm 0.01	2.77 \pm 1.47	7.42 \pm 0.82	0.29 \pm 0.03	14.14 \pm 0.11	0.021 \pm 0.002	0.059 \pm 0.001	0.031 \pm 0.003	0.0208 \pm 0.0023
ScBa5	7	III	0.38 \pm 0.27	3.08 \pm 1.84	6.89 \pm 0.10	0.30 \pm 0.05	14.27 \pm 0.01	0.021 \pm 0.004	0.059 \pm 0.001	0.029 \pm 0.001	0.0207 \pm 0.0037
ScBa6	7	*	1.14 \pm 0.37	6.57 \pm 2.71	7.08 \pm 0.03	0.31 \pm 0.07	13.89 \pm 0.35	0.022 \pm 0.006	0.059 \pm 0.001	0.026 \pm 0.001	0.0224 \pm 0.0056
ScBa7	3	IV	1.58 \pm 0.73	12.15 \pm 1.93	6.92 \pm 0.16	0.33 \pm 0.00	13.82 \pm 0.03	0.024 \pm 0.001	0.060 \pm 0.001	0.030 \pm 0.001	0.0240 \pm 0.001
ScBa8	2	V	2.98 \pm 3.16	16.78 \pm 18.98	6.54 \pm 0.43	0.47 \pm 0.34	12.60 \pm 0.90	0.037 \pm 0.025	0.056 \pm 0.002	0.029 \pm 0.005	0.0367 \pm 0.0247
ScBa9	5	*	0.69 \pm 0.25	5.68 \pm 4.83	7.07 \pm 0.01	0.29 \pm 0.02	13.73 \pm 0.20	0.021 \pm 0.001	0.058 \pm 0.002	0.030 \pm 0.001	0.0215 \pm 0.0014
ScBa10	10	VI	1.65 \pm 0.49	15.33 \pm 2.45	6.81 \pm 0.14	0.41 \pm 0.04	13.46 \pm 0.13	0.031 \pm 0.002	0.059 \pm 0.001	0.030 \pm 0.001	0.0307 \pm 0.0025
ScBa11	15	VII	1.11 \pm 0.96	11.44 \pm 10.36	7.44 \pm 0.08	0.35 \pm 0.01	13.46 \pm 0.58	0.026 \pm 0.001	0.058 \pm 0.001	0.032 \pm 0.001	0.026 \pm 0.0007
ScBa12	2	VIII	78.13 \pm 0.19	70.52 \pm 0.15	5.20 \pm 0.10	0.43 \pm 0.01	4.90 \pm 0.09	0.088 \pm 0.001	0.051 \pm 0.001	0.054 \pm 0.001	0.0879 \pm 0.0001
ScBa13	2	IX	0.82 \pm 0.25	2.80 \pm 0.68	6.63 \pm 0.08	0.20 \pm 0.02	13.87 \pm 0.28	0.014 \pm 0.002	0.058 \pm 0.001	0.028 \pm 0.001	0.0144 \pm 0.0015
ScBa14	11	X	1.14 \pm 0.72	13.37 \pm 5.12	6.99 \pm 0.07	0.29 \pm 0.03	13.59 \pm 0.45	0.021 \pm 0.003	0.059 \pm 0.001	0.030 \pm 0.001	0.0213 \pm 0.0033
ScBa15	14	XI	2.88 \pm 0.61	28.44 \pm 1.79	7.10 \pm 0.03	0.44 \pm 0.01	12.38 \pm 0.05	0.035 \pm 0.001	0.058 \pm 0.001	0.033 \pm 0.001	0.0352 \pm 0.0008
ScBa16	14	XII	3.71 \pm 0.58	28.71 \pm 1.54	6.82 \pm 0.22	0.38 \pm 0.04	12.04 \pm 0.14	0.032 \pm 0.003	0.057 \pm 0.001	0.032 \pm 0.001	0.0319 \pm 0.0028
ScBa17	2	XIII	1.09 \pm 0.07	6.93 \pm 0.34	7.37 \pm 0.01	0.39 \pm 0.01	14.03 \pm 0.03	0.028 \pm 0.001	0.059 \pm 0.001	0.031 \pm 0.001	0.0275 \pm 0.0004
ScBa18	2	*	0.89 \pm 0.37	11.64 \pm 5.35	6.46 \pm 0.65	0.41 \pm 0.18	13.00 \pm 0.32	0.031 \pm 0.013	0.056 \pm 0.001	0.028 \pm 0.004	0.0314 \pm 0.0129
ScBa19	5	XIV	0.83 \pm 0.23	8.09 \pm 5.27	6.85 \pm 0.09	0.36 \pm 0.05	13.34 \pm 0.71	0.028 \pm 0.006	0.056 \pm 0.004	0.029 \pm 0.001	0.0275 \pm 0.0055
ScBa20	14	XV	0.63 \pm 0.24	7.80 \pm 2.79	7.26 \pm 0.02	0.41 \pm 0.00	13.89 \pm 0.07	0.029 \pm 0.001	0.060 \pm 0.001	0.031 \pm 0.001	0.0292 \pm 0.0004
ScBa21	10	XVI	1.58 \pm 0.21	10.27 \pm 2.32	6.72 \pm 0.04	0.44 \pm 0.06	13.40 \pm 0.15	0.030 \pm 0.001	0.0059 \pm \pm 0.001	0.030 \pm \pm 0.001	0.0234 \pm 0.003
ScBa22	15	XVII	1.54 \pm 0.62	17.88 \pm 4.13	7.88 \pm 0.12	0.38 \pm 0.04	12.88 \pm 0.14	0.030 \pm 0.004	0.057 \pm 0.001	0.035 \pm 0.001	0.0297 \pm 0.0037
ScBa23	13	*	1.04 \pm 0.65	13.68 \pm 5.92	7.35 \pm 0.04	0.34 \pm 0.01	13.21 \pm 0.26	0.026 \pm 0.001	0.058 \pm 0.001	0.032 \pm 0.001	0.026 \pm 0.0012
ScBa24	14	*	1.18 \pm	15.75 \pm	7.13 \pm	0.34 \pm	13.14 \pm	0.026 \pm	0.058 \pm	0.031 \pm	0.0261 \pm

4			0.04	0.21	0.20	0.01	0.19	0.001	0.001	0.001	0.0007
ScBa2	7	*	3.86 ± 1.76	22.9 ± 5.06	6.81 ± 0.17	0.26 ± 0.01	12.82 ± 0.71	0.021 ± 0.001	0.059 ± 0.001	0.031 ± 0.001	0.0206 ± 0.0006
ScBa2	7	XVIII	0.69 ± 0.08	4.20 ± 1.12	7.13 ± 0.21	0.25 ± 0.04	13.96 ± 0.22	0.018 ± 0.003	0.058 ± 0.001	0.030 ± 0.001	0.0181 ± 0.0035
ScBa2	14	XIX	0.73 ± 0.48	9.18 ± 6.55	6.88 ± 0.13	0.24 ± 0.04	13.72 ± 0.64	0.017 ± 0.004	0.059 ± 0.001	0.029 ± 0.001	0.0174 ± 0.0037
ScBa2	4	XX	0.75 ± 0.10	7.31 ± 2.50	6.65 ± 0.12	0.32 ± 0.07	13.68 ± 0.30	0.024 ± 0.006	0.058 ± 0.001	0.028 ± 0.001	0.0237 ± 0.0058
ScBa2	5	XXI	1.84 ± 0.13	16.38 ± 0.98	6.80 ± 0.11	0.41 ± 0.05	13.32 ± 0.20	0.031 ± 0.004	0.059 ± 0.001	0.030 ± 0.001	0.0307 ± 0.0043
ScBa3	15	XXII	5.12 ± 2.03	31.28 ± 5.30	6.65 ± 0.03	0.44 ± 0.01	11.8 ± 0.54	0.037 ± 0.001	0.057 ± 0.001	0.032 ± 0.001	0.037 ± 0.0008
ScBa3	2	XXIII	1.15 ± 0.44	11.07 ± 4.09	6.59 ± 0.01	0.40 ± 0.06	13.56 ± 0.41	0.029 ± 0.005	0.058 ± 0.001	0.028 ± 0.001	0.0293 ± 0.0051
ScBa3	10	XXIV	0.96 ± 0.47	6.09 ± 0.38	7.03 ± 0.84	0.37 ± 0.02	13.67 ± 0.17	0.027 ± 0.002	0.058 ± 0.001	0.03 ± 0.004	0.0269 ± 0.0015
ScBa3	10	XXV	0.65 ± 0.17	5.69 ± 1.43	6.42 ± 0.17	0.44 ± 0.03	13.97 ± 0.27	0.031 ± 0.003	0.059 ± 0.002	0.027 ± 0.001	0.0313 ± 0.003
ScBa3	13	*	1.13 ± 0.54	18.34 ± 1.89	7.78 ± 0.05	0.32 ± 0.00	12.82 ± 0.01	0.025 ± 0.001	0.057 ± 0.001	0.035 ± 0.001	0.0249 ± 0.0001
ScBa3	7	*	0.62 ± 0.20	5.13 ± 1.56	7.44 ± 0.03	0.52 ± 0.02	13.93 ± 0.20	0.037 ± 0.002	0.058 ± 0.001	0.031 ± 0.001	0.0372 ± 0.0019
ScBa3	15	XXVI	1.14 ± 0.42	15.81 ± 2.76	6.69 ± 0.08	0.40 ± 0.06	12.99 ± 0.12	0.031 ± 0.004	0.057 ± 0.001	0.029 ± 0.001	0.0306 ± 0.004
ScBa3	3	XXVII	0.57 ± 0.13	5.55 ± 2.26	6.59 ± 0.07	0.33 ± 0.01	13.87 ± 0.12	0.024 ± 0.001	0.058 ± 0.001	0.028 ± 0.001	0.0239 ± 0.0012
ScBa3	10	*	0.79 ± 0.02	7.81 ± 1.69	6.99 ± 0.32	0.41 ± 0.02	13.23 ± 0.50	0.031 ± 0.002	0.056 ± 0.003	0.030 ± 0.001	0.0310 ± 0.0002
ScBa3	14	XXVII I	1.58 ± 0.35	17.36 ± 1.76	7.86 ± 0.10	0.31 ± 0.01	12.99 ± 0.23	0.024 ± 0.001	0.058 ± 0.001	0.035 ± 0.001	0.0240 ± 0.0010
ScBa4	14	XXIX	2.68 ± 0.47	24.44 ± 0.38	7.49 ± 0.68	0.42 ± 0.05	12.24 ± 0.90	0.036 ± 0.007	0.054 ± 0.004	0.034 ± 0.003	0.0356 ± 0.0068
ScBa4	12	*	1.40 ± 0.86	19.33 ± 4.79	7.12 ± 0.16	0.39 ± 0.02	13.01 ± 0.28	0.03 ± 0.002	0.058 ± 0.001	0.032 ± 0.001	0.0297 ± 0.0025
ScBa4	13	XXX	2.52 ± 0.62	25.2 ± 1.99	6.98 ± 0.23	0.36 ± 0.00	12.6 ± 0.17	0.029 ± 0.001	0.058 ± 0.001	0.032 ± 0.001	0.0287 ± 0.0003
ScBa4	15	XXXI	3.07 ± 1.42	26.09 ± 4.50	6.91 ± 0.20	0.36 ± 0.02	12.50 ± 0.68	0.029 ± 0.001	0.058 ± 0.002	0.032 ± 0.001	0.0289 ± 0.0001
ScBa4	1	XXXII	0.62 ± 0.01	3.63 ± 0.59	6.99 ± 0.08	0.27 ± 0.01	14.05 ± 0.04	0.019 ± 0.001	0.058 ± 0.001	0.029 ± 0.001	0.0189 ± 0.0003
ScBa4	12	XXXII I	3.88 ± 0.02	30.01 ± 0.48	6.86 ± 0.11	0.41 ± 0.03	12.11 ± 0.02	0.034 ± 0.002	0.058 ± 0.001	0.033 ± 0.001	0.0338 ± 0.0020
ScBa4	12	XXXI V	3.32 ± 0.14	24.06 ± 0.06	7.69 ± 0.27	0.39 ± 0.00	12.43 ± 0.22	0.031 ± 0.001	0.057 ± 0.001	0.035 ± 0.001	0.0313 ± 0.0003
ScBa4	4	XXXV	61.26 ± 8.62	48.21 ± 7.22	5.65 ± 0.44	0.52 ± 0.02	7.42 ± 0.89	0.071 ± 0.006	0.055 ± 0.001	0.042 ± 0.002	0.0708 ± 0.0058
ScBa4	6	XXXV I	68.47 ± 0.49	53.25 ± 1.42	5.51 ± 0.29	0.54 ± 0.02	6.86 ± 0.28	0.079 ± 0.001	0.056 ± 0.002	0.045 ± 0.002	0.0786 ± 0.0003
ScBa4	15	XXXV II	1.39 ± 0.90	17.01 ± 3.35	6.94 ± 0.05	0.37 ± 0.00	12.88 ± 0.40	0.029 ± 0.001	0.057 ± 0.001	0.031 ± 0.001	0.0288 ± 0.0012
ScBa5	9	*	2.78 ± 0.19	25.25 ± 0.23	6.75 ± 0.15	0.39 ± 0.01	12.6 ± 0.08	0.031 ± 0.001	0.058 ± 0.001	0.031 ± 0.001	0.0311 ± 0.0003
ScBa5	15	XXXV III	2.48 ± 0.03	24.83 ± 0.26	6.81 ± 0.18	0.36 ± 0.01	12.49 ± 0.22	0.029 ± 0.001	0.058 ± 0.001	0.031 ± 0.001	0.0291 ± 0.0001
ScBa5	8	XXXI X	0.71 ± 0.09	8.32 ± 0.62	6.67 ± 0.11	0.43 ± 0.02	13.73 ± 0.08	0.031 ± 0.001	0.058 ± 0.001	0.028 ± 0.001	0.0312 ± 0.0010
ScBa5	11	*	1.78 ± 1.01	21.99 ± 5.78	6.80 ± 0.04	0.42 ± 0.03	12.60 ± 0.14	0.033 ± 0.002	0.057 ± 0.001	0.031 ± 0.001	0.0333 ± 0.0019
ScBa5	14	XL	1.72 ± 1.39	15.94 ± 7.53	7.06 ± 0.01	0.31 ± 0.00	13.24 ± 0.42	0.023 ± 0.001	0.058 ± 0.001	0.031 ± 0.001	0.0233 ± 0.0011
ScBa5	1	XLI	2.11 ± 1.08	17.06 ± 5.14	6.70 ± 0.06	0.24 ± 0.01	13.18 ± 0.48	0.018 ± 0.001	0.059 ± 0.001	0.030 ± 0.001	0.0181 ± 0.0015
ScBa5	6	XLII	1.85 ± 0.39	15.86 ± 1.64	6.78 ± 0.01	0.32 ± 0.01	13.35 ± 0.16	0.024 ± 0.001	0.059 ± 0.001	0.030 ± 0.001	0.0238 ± 0.0007
ScBa5	3	XLIII	68.19 ± 0.5	45.76 ± 2.71	5.21 ± 0.13	0.60 ± 0.00	7.39 ± 0.17	0.081 ± 0.002	0.057 ± 0.001	0.040 ± 0.002	0.0811 ± 0.0024
ScBa5	8	XLIV	4.84 ±	30.43 ±	6.53 ±	0.43 ±	11.73 ±	0.038 ±	0.055 ±	0.031 ±	0.0375 ±

8			3.73	10.18	0.20	0.01	1.22	0.004	0.002	0.003	0.0045
ScBa5	1	*	3.60 ±	28.12 ±	6.84 ±	0.42 ±	12.20 ±	0.034 ±	0.057 ±	0.032 ±	0.0341 ±
9			1.45	4.22	0.01	0.00	0.46	0.001	0.001	0.001	0.0011
ScBa6	10	XLIV	0.69 ±	4.12 ±	7.81 ±	0.30 ±	14.08 ±	0.022 ±	0.059 ±	0.033 ±	0.0216 ±
0			0.20	2.80	0.20	0.02	0.03	0.001	0.001	0.001	0.0013
ScBa6	14	XLV	0.56 ±	7.10 ±	6.96 ±	0.42 ±	13.79 ±	0.030 ±	0.058 ±	0.029 ±	0.0305 ±
1			0.05	0.59	0.19	0.00	0.18	0.001	0.001	0.001	0.0007
ScBa6	13	*	0.98 ±	13.73 ±	6.43 ±	0.32 ±	13.21 ±	0.024 ±	0.058 ±	0.028 ±	0.0239 ±
2			0.08	0.37	0.17	0.01	0.28	0.001	0.001	0.001	0.0001
ScBa6	14	*	0.39 ±	3.85 ±	7.20 ±	0.50 ±	13.58 ±	0.038 ±	0.056 ±	0.030 ±	0.0377 ±
3			0.15	1.71	0.07	0.14	0.48	0.012	0.002	0.001	0.0116
ScBa6	14	*	0.98 ±	7.81 ±	6.87 ±	0.32 ±	13.33 ±	0.025 ±	0.056 ±	0.029 ±	0.0248 ±
4			0.54	1.64	0.14	0.09	0.61	0.008	0.002	0.001	0.0079
ScBa6	7	XLVI	1.07 ±	9.91 ±	6.46 ±	0.23 ±	13.42 ±	0.017 ±	0.057 ±	0.028 ±	0.0173 ±
5			0.51	4.72	0.09	0.01	0.50	0.001	0.001	0.001	0.0013
ScBa6	4	XLVII	3.00 ±	25.73 ±	6.76 ±	0.39 ±	12.15 ±	0.032 ±	0.056 ±	0.031 ±	0.0319 ±
6			0.12	0.44	0.01	0.01	0.07	0.001	0.001	0.001	0.0005
ScBa6	4	*	3.17 ±	25.26 ±	7.72 ±	0.37 ±	12.71 ±	0.029 ±	0.059 ±	0.036 ±	0.0292 ±
7			1.05	2.74	1.37	0.01	0.56	0.001	0.004	0.007	0.0002
ScBa6	4	*	0.89 ±	9.31 ±	6.60 ±	0.22 ±	13.46 ±	0.016 ±	0.057 ±	0.028 ±	0.0162 ±
8			0.22	2.35	0.22	0.03	0.31	0.002	0.001	0.001	0.0017
ScBa6	4	*	0.85 ±	8.44 ±	6.54 ±	0.24 ±	13.41 ±	0.018 ±	0.057 ±	0.028 ±	0.0182 ±
9			0.42	4.20	0.03	0.02	0.27	0.002	0.001	0.001	0.0021
ScBa7	12	XLVII	4.27 ±	31.23 ±	6.88 ±	0.40 ±	12.27 ±	0.033 ±	0.059 ±	0.033 ±	0.0330 ±
0		I	1.33	3.89	0.02	0.03	0.27	0.001	0.003	0.001	0.0014
ScBa7	11	*	0.72 ±	4.54 ±	6.72 ±	0.30 ±	13.88 ±	0.021 ±	0.058 ±	0.028 ±	0.0214 ±
1			0.33	2.92	0.09	0.01	0.39	0.001	0.001	0.001	0.0001
ScBa7	6	XLIX	0.66 ±	7.05 ±	6.84 ±	0.40 ±	13.78 ±	0.029 ±	0.058 ±	0.029 ±	0.0291 ±
2			0.09	0.76	0.01	0.01	0.08	0.001	0.001	0.001	0.0009
ScBa7	14	*	0.78 ±	7.81 ±	7.10 ±	0.42 ±	13.82 ±	0.031 ±	0.059 ±	0.030 ±	0.0307 ±
3			0.04	0.64	0.04	0.00	0.02	0.001	0.001	0.001	0.0002
ScBa7	8	L	0.93 ±	9.96 ±	6.64 ±	0.40 ±	13.49 ±	0.030 ±	0.058 ±	0.028 ±	0.0300 ±
4			0.28	2.72	0.01	0.01	0.11	0.001	0.001	0.001	0.0007
ScBa7	9	LI	3.57 ±	27.49 ±	6.79 ±	0.38 ±	12.35 ±	0.031 ±	0.058 ±	0.032 ±	0.0309 ±
5			2.41	8.01	0.13	0.01	0.66	0.002	0.001	0.001	0.0022
ScBa7	9	*	3.63 ±	30.74 ±	6.58 ±	0.37 ±	12.01 ±	0.031 ±	0.057 ±	0.031 ±	0.0308 ±
6			1.97	4.86	0.05	0.00	0.29	0.001	0.001	0.001	0.0008
ScBa7	9	*	3.96 ±	30.12 ±	6.61 ±	0.37 ±	11.93 ±	0.031 ±	0.057 ±	0.031 ±	0.0314 ±
7			1.15	3.88	0.04	0.03	0.41	0.001	0.001	0.001	0.0011
ScBa7	7	*	1.02 ±	9.71 ±	6.15 ±	0.31 ±	13.42 ±	0.023 ±	0.057 ±	0.026 ±	0.0228 ±
8			0.22	2.42	0.11	0.04	0.11	0.003	0.001	0.001	0.0033
ScBa7	11	*	1.23 ±	12.73 ±	6.52 ±	0.23 ±	13.37 ±	0.017 ±	0.058 ±	0.028 ±	0.0169 ±
9			0.37	2.91	0.13	0.01	0.06	0.001	0.001	0.001	0.0007
ScBa8	8	*	6.79 ±	36.88 ±	6.57 ±	0.41 ±	11.22 ±	0.037 ±	0.056 ±	0.033 ±	0.0366 ±
0			2.11	3.98	0.00	0.02	0.36	0.003	0.001	0.001	0.0032
ScBa8	7	LII	1.96 ±	15.41 ±	6.96 ±	0.40 ±	13.30 ±	0.030 ±	0.059 ±	0.031 ±	0.0299 ±
1			1.25	6.55	0.03	0.01	0.38	0.002	0.001	0.001	0.0017
ScBa8	6	LIII	1.00 ±	10.89 ±	6.65 ±	0.21 ±	13.36 ±	0.016 ±	0.057 ±	0.029 ±	0.0157 ±
2			0.06	0.92	0.04	0.00	0.07	0.001	0.001	0.001	0.0001
ScBa8	5	LIV	2.43 ±	17.35 ±	6.66 ±	0.38 ±	12.82 ±	0.030 ±	0.057 ±	0.030 ±	0.0298 ±
3			1.29	5.92	0.15	0.01	0.45	0.002	0.001	0.001	0.0017
ScBa8	15	LV	2.14 ±	23.56 ±	6.68 ±	0.40 ±	12.47 ±	0.032 ±	0.057 ±	0.031 ±	0.0318 ±
4			0.86	3.33	0.05	0.02	0.29	0.002	0.001	0.001	0.0022
ScBa8	14	*	3.92 ±	29.25 ±	6.70 ±	0.38 ±	12.10 ±	0.031 ±	0.057 ±	0.032 ±	0.0314 ±
5			1.20	3.40	0.02	0.00	0.16	0.001	0.001	0.001	0.0008
ScBa8	6	LVI	1.06 ±	10.25 ±	6.73 ±	0.42 ±	13.60 ±	0.031 ±	0.058 ±	0.029 ±	0.0307 ±
6			0.15	1.61	0.13	0.01	0.20	0.001	0.002	0.001	0.0011
ScBa8	1	LVII	1.74 ±	21.21 ±	6.85 ±	0.39 ±	12.54 ±	0.031 ±	0.057 ±	0.031 ±	0.0311 ±
7			0.96	4.94	0.07	0.02	0.20	0.002	0.001	0.001	0.0021
ScBa8	12	LVIII	2.85 ±	25.68 ±	6.64 ±	0.37 ±	12.39 ±	0.029 ±	0.058 ±	0.031 ±	0.0295 ±
8			2.21	9.50	0.24	0.02	0.02	0.002	0.003	0.003	0.0016
ScBa8	13	*	2.24 ±	19.4 ±	7.12 ±	0.33 ±	13.07 ±	0.025 ±	0.059 ±	0.032 ±	0.0253 ±
9			0.24	0.70	0.07	0.02	0.18	0.001	0.001	0.001	0.0009
ScBa9	13	LIX	2.55 ±	24.87 ±	7.43 ±	0.32 ±	12.29 ±	0.026 ±	0.057 ±	0.034 ±	0.0258 ±
0			0.28	0.73	0.20	0.01	0.13	0.001	0.002	0.001	0.0011
ScBa9	10	LX	2.64 ±	19.53 ±	6.36 ±	0.37 ±	12.64 ±	0.029 ±	0.057 ±	0.029 ±	0.0292 ±
1			0.41	1.64	0.17	0.01	0.11	0.001	0.001	0.001	0.0007
ScBa9	14	LXI	3.43 ±	26.05 ±	6.49 ±	0.36 ±	12.06 ±	0.030 ±	0.056 ±	0.030 ±	0.0301 ±

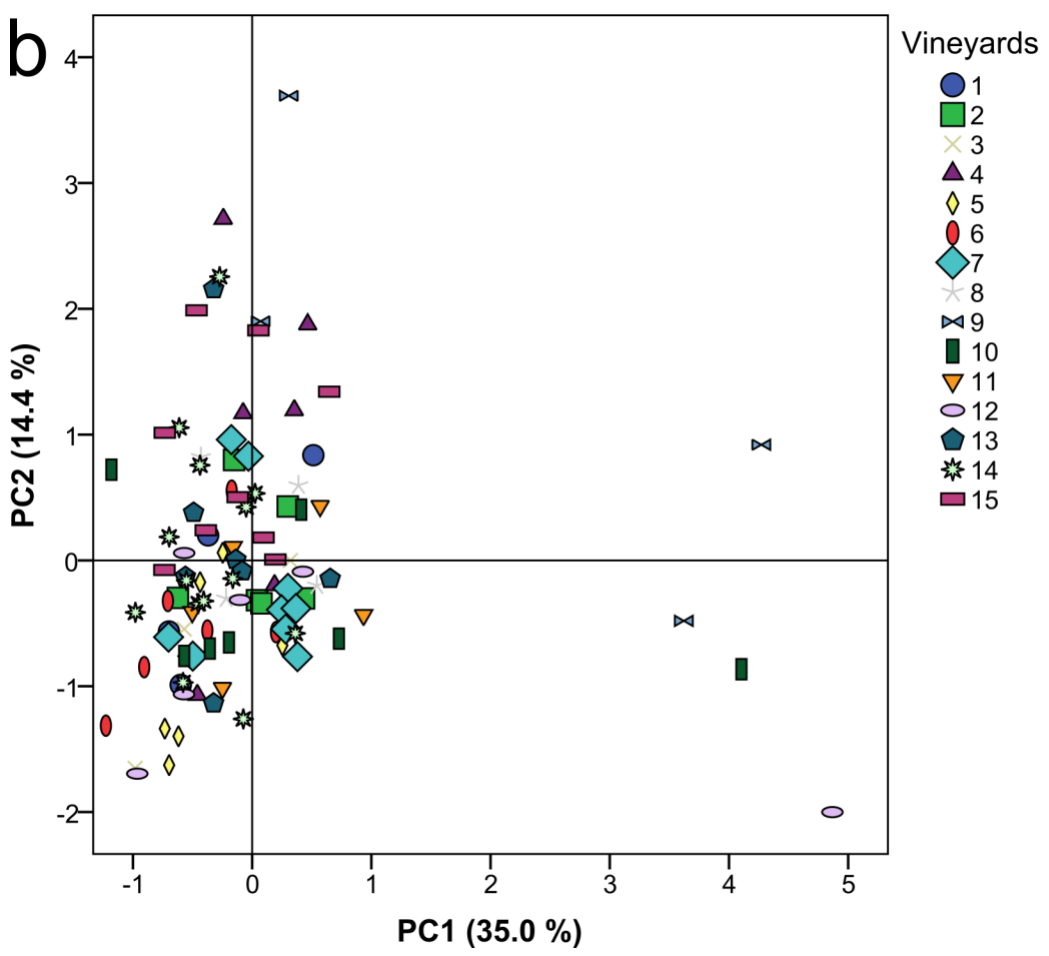
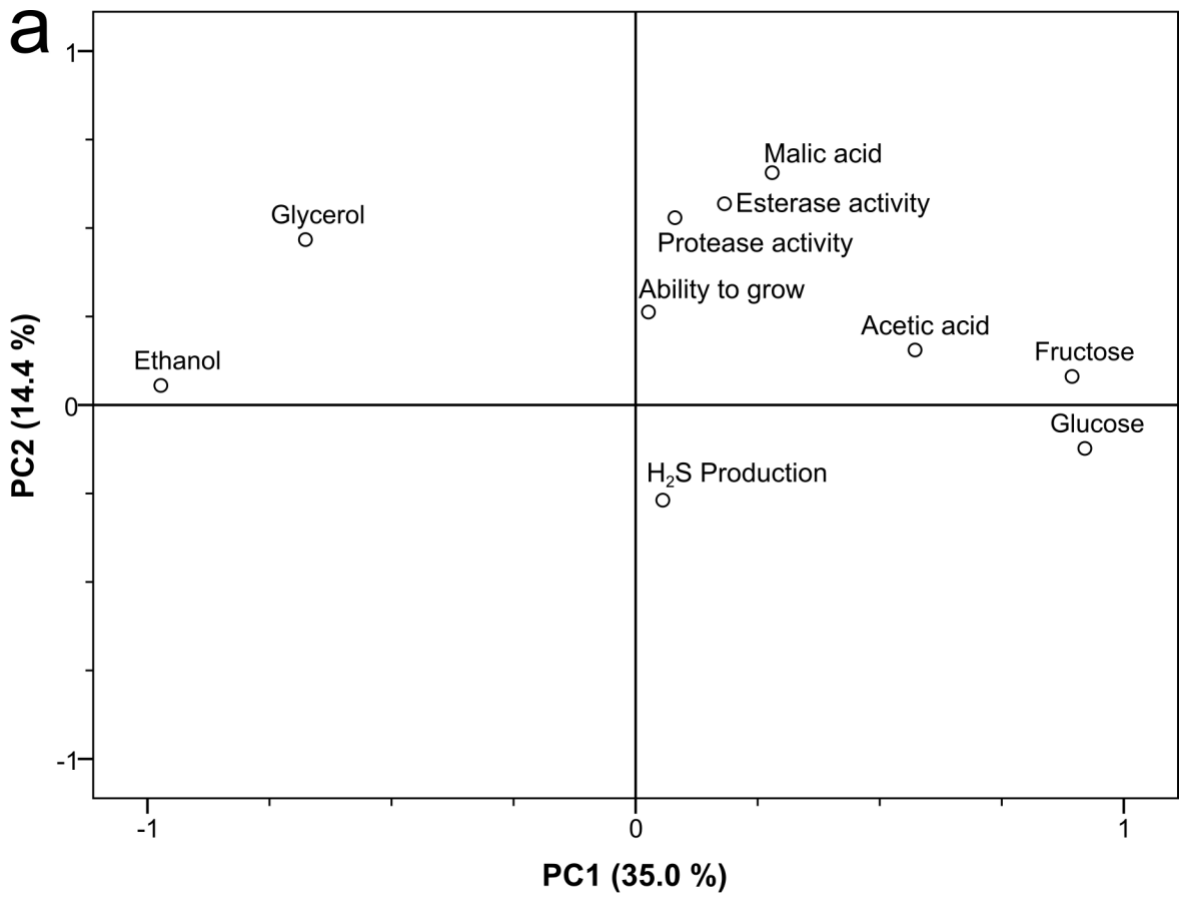
2			2.51	8.16	0.02	0.01	0.33	0.001	0.001	0.002	0.0001
ScBa9	6	LXII	0.67 ± 0.31	6.69 ± 5.00	6.46 ± 0.18	0.21 ± 0.01	13.55 ± 0.34	0.016 ± 0.001	0.057 ± 0.002	0.027 ± 0.001	0.0156 ± 0.0001
ScBa9	5	XX	0.83 ± 0.05	10.16 ± 0.53	6.59 ± 0.05	0.28 ± 0.00	13.46 ± 0.05	0.021 ± 0.001	0.058 ± 0.001	0.028 ± 0.001	0.0205 ± 0.0003
ScBa9	15	LXII	2.42 ± 0.73	24.31 ± 2.48	6.55 ± 0.17	0.37 ± 0.01	12.31 ± 0.24	0.030 ± 0.002	0.057 ± 0.002	0.030 ± 0.001	0.0301 ± 0.0017
ScBa9	12	LIX	3.18 ± 1.87	25.13 ± 5.26	6.67 ± 0.06	0.40 ± 0.02	12.12 ± 0.28	0.033 ± 0.001	0.056 ± 0.001	0.031 ± 0.001	0.0329 ± 0.0007

288 a Fermentation purity: acetic acid (g/L)/ethanol % (v/v), b Ethanol yield: ethanol % (v/v)/sugar
289 consumption (g/L), c Glycerol yield: glycerol (g/L)/sugar consumption (g/L), d Acetic acid: acetic
290 acid (g/L)/sugar consumption(g/L).

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293 The physiological data from the growth tests at 50 mg/L of SO₂ after 24 h (enological conditions,
294 variable: “Ability to grow”), the presence or absence of enzymatic activities (esterase and protease
295 activity), H₂S production and the chemical composition (glucose, fructose, malic and acetic acids,
296 glycerol, and ethanol) of the wines obtained after 14 days of fermentation were used to evaluate the
297 technological diversity of the strains. A Principal Component Analysis (PCA) was carried out and
298 the outcome is presented in Figure 2, including the loadings plot (Figure 2a) and the scatter plot
299 (Figure 2b). PC1 (35.0 % of variance explained) was better correlated with the strains that left high
300 level of residual sugars present in the wine and that produced high amount of acetic acid, while PC2
301 (14.4 % of variance explained) was mainly correlated to the high production of glycerol and low
302 degradation of malic acid (Figure 2a). Two groups may be differentiated according to the scatter
303 plot (PC1 or PC2 with values close or higher than 2, respectively). Some strains from vineyards 4,
304 13, 14, and 15 produced high glycerol amounts and preserved malic acid contents, and two isolates
305 from vineyards 10 and 12 were unsatisfactory due to their sugars degradation. In addition, all the
306 four isolates from vineyard 9 were present either in the first or in the second group (Figure 2b).

307



309 **Figure 2.** Principal component analysis of the *S. cerevisiae* strains from the fifteen vineyards (as
310 reported Figure 1), according to the wine chemical composition: loadings plot (a) and scatter plot
311 (b). The fifteen vineyard considered were: 1 (Murisengo), 2 (San Martino Alfieri), 3 (Costigliole
312 d'Asti), 4 (Isola d'Asti), 5 (Montegrosso d'Asti), 6 (Agliano Terme), 7 (Vinchio), 8 (Nizza
313 Monferrato), 9 (Incisa Scapaccino), 10 (Loazzolo), 11 (Ricaldone), 12 (Alice bel colle), 13 (Acqui
314 Terme Crocera south west), 14 (Acqui Terme Crocera south est) and 15 (Acqui Terme Dannonna).

315

316 **CONCLUSIONS**

317 The present study investigated the genetic and technological diversity of autochthonous *S.*
318 *cerevisiae* in the North-West of Italy, in the Monferrato area. It was possible to genetically and
319 phenotypically differentiate the strains. In order to investigate the ability of the isolated strains to
320 properly ferment Barbera must, at pilot scale level first and in industrial settings after, relevant
321 technological characteristics, such as sugar consumption and acetic acid production, should be taken
322 into consideration. All the data presented here were obtained from pasteurized natural must, and the
323 ability of the selected strains to dominate the natural grape and must mycobiota should therefore be
324 determined throughout the fermentation process. In parallel, the ability to complete the fermentation
325 in the competitive environment of a natural grape must should also be confirmed. Finally, the
326 production of compounds, such as alcohols, esters, carbonyl compounds and fatty acids that have an
327 impact on the sensory characteristics of wine should also be evaluated.

328

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331 www.wildwine.eu/). The information in this document only reflects the authors' views and the
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471 **Supplementary table 1.** Results of the resistance to ethanol and SO₂ of the tested strains. The
472 values presented are the ratio between the OD of the isolates in broth with and without ethanol or
473 SO₂ times 100 at the specific incubation times. The values are the means of triplicate experiments.

Strains	SO ₂ growth (mg/L)						Ethanol growth (% vol.)					
	24 hours of incubation			48 hours of incubation			24 hours of incubation			48 hours of incubation		
	50	100	150	50	100	150	12	14	16	12	14	16
ScBa1	36	6	14	81	74	81	0	0	0	64	0	0
ScBa2	82	87	71	93	99	89	100	93	100	100	100	100
ScBa3	81	46	36	100	91	100	36	6	7	97	47	0
ScBa4	34	3	4	100	72	36	2	1	1	17	0	0
ScBa5	81	46	36	98	99	96	36	6	7	98	73	4
ScBa6	44	46	16	87	67	52	11	1	1	13	1	1
ScBa7	58	48	42	95	88	61	9	4	4	90	6	5
ScBa8	76	49	42	98	94	93	8	4	5	85	7	5
ScBa9	70	42	43	96	92	94	5	4	5	72	4	5
ScBa10	26	15	9	70	69	60	19	1	0	70	0	0
ScBa11	22	6	6	87	83	10	2	1	1	5	0	0
ScBa12	9	6	2	44	7	7	4	2	2	3	1	1
ScBa13	60	32	32	99	93	83	9	3	3	97	3	3
ScBa14	38	2	1	85	55	16	51	1	0	61	0	0
ScBa15	26	3	3	72	30	14	49	0	0	58	0	0
ScBa16	2	4	23	6	40	77	51	0	0	61	0	0
ScBa17	12	0	0	91	54	17	37	0	0	76	0	1
ScBa18	47	0	0	55	9	0	1	1	0	38	0	0
ScBa19	87	58	52	85	84	75	40	0	0	77	0	0
ScBa20	63	0	0	64	2	0	0	0	0	13	0	0
ScBa21	52	31	46	60	59	50	55	0	0	83	0	0
ScBa22	96	91	83	100	97	96	100	95	97	98	93	99
ScBa23	67	63	54	96	66	40	96	70	71	100	92	100

Strains	SO ₂ growth (mg/L)						Ethanol growth (% vol.)					
	24 hours of incubation			48 hours of incubation			24 hours of incubation			48 hours of incubation		
	50	100	150	50	100	150	12	14	16	12	14	16
ScBa1	36	6	14	81	74	81	0	0	0	64	0	0
ScBa2	82	87	71	93	99	89	100	93	100	100	100	100
ScBa3	81	46	36	100	91	100	36	6	7	97	47	0
ScBa4	34	3	4	100	72	36	2	1	1	17	0	0
ScBa5	81	46	36	98	99	96	36	6	7	98	73	4
ScBa6	44	46	16	87	67	52	11	1	1	13	1	1
ScBa7	58	48	42	95	88	61	9	4	4	90	6	5
ScBa8	76	49	42	98	94	93	8	4	5	85	7	5
ScBa9	70	42	43	96	92	94	5	4	5	72	4	5
ScBa10	26	15	9	70	69	60	19	1	0	70	0	0
ScBa11	22	6	6	87	83	10	2	1	1	5	0	0
ScBa12	9	6	2	44	7	7	4	2	2	3	1	1
ScBa13	60	32	32	99	93	83	9	3	3	97	3	3
ScBa14	38	2	1	85	55	16	51	1	0	61	0	0
ScBa15	26	3	3	72	30	14	49	0	0	58	0	0
ScBa16	2	4	23	6	40	77	51	0	0	61	0	0
ScBa17	12	0	0	91	54	17	37	0	0	76	0	1
ScBa18	47	0	0	55	9	0	1	1	0	38	0	0
ScBa19	87	58	52	85	84	75	40	0	0	77	0	0
ScBa20	63	0	0	64	2	0	0	0	0	13	0	0
ScBa21	52	31	46	60	59	50	55	0	0	83	0	0
ScBa22	96	91	83	100	97	96	100	95	97	98	93	99
ScBa23	67	63	54	96	66	40	96	70	71	100	92	100