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**Table S1.** Analysis of variance (ANOVA) of filamentous fungi detected by NGS in Montepulciano samples at harvest time and at 7<sup>th</sup> and 15<sup>th</sup> day) of microfermentations carried out using aseptically pressed grapes.

Culture-Independent Method (NGS)	Harvest Time			7 <sup>th</sup> Day			15 <sup>th</sup> Day		
	Fungal Species	MO	MC	MNT	MO	MC	MNT	MO	MC
<i>Aspergillus piperis</i>	*	nd	nd	A	nd	A	A	nd	A
<i>Aureobasidium pullulans</i>	B	A	A	A	A	A	A	A	A
<i>Botryosphaeria agaves</i>	A	A	A	A	B	AB	A	B	AB
<i>Botrytis caroliniana</i>	A	A	A	A	A	A	A	A	A
<i>Candida californica</i>	nd	nd	nd	A	A	A	A	A	A
<i>Cladosporium delicatulum</i>	A	A	A	A	A	A	A	A	A
<i>Cladosporium ramotenellum</i>	A	B	B	A	B	AB	A	A	A
<i>Cryptococcus consortionis</i>	B	A	AB	nd	nd	nd	nd	nd	nd
<i>Dissoconium aciculare</i>	nd	nd	*	nd	nd	nd	nd	nd	nd
<i>Epicoccum nigrum</i>	A	A	A	A	A	A	A	A	A
<i>Erysiphe necator</i>	B	AB	A	nd	nd	nd	nd	nd	nd
<i>Filobasidium chernovii</i>	A	A	A	A	A	A	A	A	A
<i>Hanseniaspora uvarum</i>	A	A	A	A	A	A	A	A	A
<i>Hanseniaspora vineae</i>	nd	nd	nd	nd	nd	nd	nd	*	nd
<i>Kodamaea ohmeri</i>	nd	nd	nd	nd	nd	nd	A	nd	A
<i>Lachancea thermotolerans</i>	nd	*	nd	nd	*	nd	nd	*	nd
<i>Lambertella palmeri</i>	nd	nd	*	nd	nd	nd	nd	nd	nd
<i>Metschnikowia chrysoperlae</i>	nd	nd	nd	nd	*	nd	nd	*	nd
<i>Metschnikowia pulcherrima</i>	nd	nd	*	nd	*	nd	nd	*	nd
<i>Meyerozyma guilliermondii</i>	nd	nd	nd	nd	nd	nd	A	nd	A
<i>Mycosphaerella tassiana</i>	A	A	A	A	A	A	A	A	A
<i>Penicillium neocrassum</i>	nd	nd	*	nd	nd	nd	nd	nd	nd
<i>Pichia kluyveri</i>	nd	nd	nd	nd	nd	*	nd	nd	*
<i>Pichia terricola</i>	A	A	nd	A	A	A	A	nd	A
<i>Rhodotorula nothofagi</i>	nd	nd	*	nd	nd	nd	nd	*	nd
<i>Saccharomyopsis vini</i>	*	nd	nd	*	nd	nd	A	nd	A
<i>Sporobolomyces symmetricus</i>	B	B	A	A	nd	A	A	nd	A
<i>Starmerella bacillaris</i>	A	B	AB	A	B	AB	A	B	AB
<i>Stemphylium herbarum</i>	A	A	A	A	A	A	A	A	A

<i>U. m. of Alternaria genus</i>	A	A	A	A	A	A	A	A	A
<i>U. m. of Ascomycota phylum</i>	A	A	A	nd	nd	nd	nd	nd	nd
<i>U. m. of Basidiomycota phylum</i>	C	B	A	nd	nd	nd	A	A	A
<i>U. m. of Fungi kingdom</i>	A	A	A	A	A	A	A	A	A
<i>U. m. of Metschnikowia genus</i>	nd	*	nd	nd	*	nd	nd	*	nd
<i>U. m. of Pleosporales order</i>	B	B	A	nd	nd	nd	nd	nd	nd
<i>U. m. of Strophariaceae family</i>	nd	nd	nd	nd	nd	nd	nd	*	nd
<i>Vishniacozyma carnescens</i>	A	A	A	A	A	A	A	A	A
<i>Zygoascus meyerae</i>	A	nd	A	A	nd	A	*	nd	nd
<i>Zygosaccharomyces bailii</i>	A	nd	A	*	nd	nd	A	A	nd

The significant differences were determined using t-Test ( $P$  value < 0.05). Different letters (A, B, C) within each row indicate significant difference. Only the fungal species > 0.5% of relative abundance were represented. nd = not detected. \* = fungi detected in only one treatment (MO, MT or MNT).

**Table S2.** Analysis of variance (ANOVA) of yeast species detected by culture-dependent method in Montepulciano samples at harvest time and at 7<sup>th</sup> and 15<sup>th</sup> day of microfermentations carried out using aseptically pressed grapes.

Culture-Dependent Method	Harvest Time			7 <sup>th</sup> Day			15 <sup>th</sup> Day		
	MO	MC	MNT	MO	MC	MNT	MO	MC	MNT
<i>Aureobasidium pullulans</i>	A	A	A	nd	nd	nd	nd	nd	nd
<i>Candida californica</i>	*	nd	nd	A	A	A	A	A	A
<i>Cryptococcus carnescens</i>	*	nd	nd	nd	nd	nd	nd	nd	nd
<i>Cryptococcus flavescens</i>	nd	nd	nd	nd	nd	nd	nd	nd	nd
<i>Cryptococcus genus</i>	A	A	nd	nd	nd	nd	nd	nd	nd
<i>Debaryomyces hansenii</i>	nd	*	nd	nd	nd	nd	A	A	nd
<i>Hanseniaspora uvarum</i>	A	A	A	AB	A	B	A	A	nd
<i>Lachancea thermotolerans</i>	nd	*	nd	nd	*	nd	nd	*	nd
<i>Metschnikowia pulcherrima</i>	nd	B	A	nd	*	nd	nd	nd	nd
<i>Pichia kluyveri</i>	nd	nd	nd	nd	nd	nd	*	nd	nd
<i>Pichia sporocuriosa</i>	nd	nd	nd	*	nd	nd	nd	*	nd
<i>Pichia terricola</i>	A	A	A	*	nd	nd	*	nd	nd
<i>Rhodotorula genus</i>	nd	A	nd	nd	nd	nd	nd	nd	nd
<i>Starmerella bacillaris</i>	A	nd	A	A	A	A	A	A	A
<i>Zygoascus meyerae</i>	A	nd	A	nd	nd	nd	nd	nd	nd
<i>Zygosaccharomyces bailii</i>	nd	nd	*	nd	nd	*	A	A	A

The significant differences were determined using t-Test ( $P$  value < 0.05). Different letters (A, B) within each row indicate significant difference. Only the yeast species > 0.5% of relative abundance were represented. nd = not detected. \* = fungi detected in only one treatment (MO, MT or MNT).



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**Table S3.** The main analytical compounds of Montepulciano microfermentations carried out using aseptically pressed grapes.

Samples	Residual Sugars* (g L <sup>-1</sup> )	Ethanol (% v/v)	Volatile Acidity (as Acetic Acid g L <sup>-1</sup> )
MO1	44.7	9.9	6.00
MO2	66.2	8.7	2.76
MO3	58.1	9.1	4.80
MO4	74.7	8.2	13.92
MO5	81.9	7.8	1.75
MO6	79.1	7.9	5.76
MO7	42.3	10.1	16.32
MC1	43.9	9.9	6.72
MC2	22.5	11.1	1.03
MC3	22.4	11.1	1.73
MC4	65.9	8.7	5.04
MC5	58.4	9.1	4.80
MC6	94.6	7.1	3.36
MC7	92.6	7.2	3.36
MNT1	36.4	10.3	14.40
MNT2	4.3	12.1	2.06
MNT3	24.3	11.0	6.00

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\* Total initial sugars 221 g L<sup>-1</sup>.

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**Table S4.** Analytical results of organic (MO), conventional (MC) and not treated (MNT) Montepulciano samples.

Samples	Residual Sugars* (g L <sup>-1</sup> )	Ethanol (% v/v)	Volatile Acidity (as Acetic Acid g L <sup>-1</sup> )
MO	63.86±16.06 <sup>a</sup>	8.80±0.90 <sup>b</sup>	7.33±5.581 <sup>a</sup>
MC	57.20±29.78 <sup>a</sup>	9.17±1.67 <sup>b</sup>	3.72±1.97 <sup>a</sup>
MNT	21.68±16.23 <sup>b</sup>	11.16±0.91 <sup>a</sup>	7.49±6.30 <sup>a</sup>

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\* Total initial sugars 221 g L<sup>-1</sup>. Data are means ± standard deviations. Data with different superscript letters (<sup>a,b</sup>) within each column are significantly different (Duncan test; *p* < 0.05).



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