

Comparative metabolic profiling to investigate the contribution of *O. oeni* MLF starter cultures to red wine composition

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Abstract In this research work we investigated changes in volatile aroma composition associated with four commercial *Oenococcus oeni* malolactic fermentation (MLF) starter cultures in South African Shiraz and Pinotage red wines. A control wine in which MLF was suppressed was included. The MLF progress was monitored by use of infra-red spectroscopy. Gas chromatographic analysis and capillary electrophoresis were used to evaluate the volatile aroma composition and organic acid profiles, respectively. Significant strain-specific variations were observed in the degradation of citric acid and production of lactic acid during MLF. Subsequently, compounds directly and indirectly resulting from citric acid metabolism, namely diacetyl, acetic acid, acetoin, and ethyl lactate, were also affected depending on the bacterial strain used for MLF. Bacterial metabolic activity increased concentrations of the higher alcohols, fatty acids, and total esters, with a larger increase in ethyl esters than in acetate esters. Ethyl lactate, diethyl succinate, ethyl octanoate, ethyl 2-methylpropanoate, and ethyl propionate concentrations were increased by MLF. In contrast, levels of hexyl acetate, isoamyl acetate, 2-phenylethyl acetate, and ethyl acetate were reduced or remained unchanged, depending on the strain and cultivar evaluated. Formation of ethyl butyrate, ethyl propionate, ethyl 2-methylbutyrate, and ethyl isovalerate was related to specific bacterial strains used, indicating possible differences in

esterase activity. A strain-specific tendency to reduce total aldehyde concentrations was found at the completion of MLF, although further investigation is needed in this regard. This study provided insight into metabolism in *O. oeni* starter cultures during MLF in red wine.

Keywords Malolactic fermentation (MLF) · Volatile aroma composition · MLF starter cultures · *Oenococcus oeni* · Red wine

Introduction

Wine production involves a succession of biological processes including alcoholic fermentation by yeast and malolactic fermentation (MLF) by lactic acid bacteria (LAB). MLF in wine is performed, preferably by inoculation with *Oenococcus oeni* in order to reduce wine acidity by biotransformation of the dicarboxylic L-malic acid to the monocarboxylic L-lactic acid by the malolactic enzyme [9, 15, 33, 36]. MLF improves biological stability and affects wine organoleptic properties such as aroma, flavour, and mouthfeel [4, 9, 29]. *O. oeni* is recognised as the species most tolerant to the harsh wine conditions of low pH, high sulfur dioxide (SO₂), and high alcohol content. For this reason, in addition to its favourable flavour profile, *O. oeni* is mostly selected as starter culture [33].

The complexity and diversity of the metabolic activity associated with the growth of LAB suggest that MLF may affect wine quality both positively and negatively [15, 35]. Wine aroma and flavour could be affected by LAB via several mechanisms including:

- 1 reduction of flavour compounds by metabolism and adsorption on the bacterial cell wall;

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- 2 production of new volatiles by metabolism of grape sugars, amino acids, and other nutrient compounds; and
- 3 metabolism or extracellular modification of grape and yeast secondary metabolites to either more or less flavour and aroma-active metabolites [2].

Wine-associated LAB have been shown to induce a variety of enzymatic activity which has the potential to affect or produce a range of volatile compounds [35, 42–44]. Different enzymatic activity has been observed among the LAB genera and strains evaluated [43]. The use of different bacterial strains in starter cultures during winemaking could, therefore, substantially affect the volatile composition and possibly the resulting sensory properties in a strain-dependent manner.

One of the most important aroma compounds synthesized during MLF, and the most frequently reported cause of aroma modification associated with LAB, is diacetyl (2,3-butanedione), which, when present at concentrations above its sensory threshold, contributes a buttery, nutty, and/or toasty aroma to wine [2, 9, 18, 32, 41]. It is well-known that diacetyl, acetic acid, acetoin, and 2,3-butanediol are formed by citric acid catabolism by LAB; these have been discussed in several comprehensive reviews [3–5, 37, 58]. According to previous reports, modification of wine aroma induced by MLF is far more complex and often involves changes in fruity, flowery, and nutty attributes, and reduction of vegetative, green, grassy, and herbaceous aromas [2, 28]. A significant increase in the concentration of several esters produced by bacterial metabolism has been reported [11, 39], whereas other studies reported reduced ester concentrations [24]. The catabolism of acetaldehyde by wine LAB was reported by Osborne et al. [48]. This illustrates the potential of LAB to metabolise aldehydes [35] and, consequently, reduce the associated herbaceous aroma; reports on these changes during MLF are few, however. Additional compounds such as higher alcohols, fatty acids, lactones, and sulfur and nitrogen-containing compounds may also be produced and could potentially contribute to or alter the wine aroma profile [56].

Different analytical procedures have been described for quantification of volatile compounds in wine, and comprehensive reviews are available [16, 47]. Gas chromatography (GC) in combination with a variety of extraction and detection techniques have been most extensively used for quantification of wine volatile compounds. Headspace SPME (HS-SPME) [1, 61] is an effective and solventless sampling technique especially suitable for quantification of volatile analytes, because it reduces interferences from other, non-volatile, wine constituents. Hayasaka et al. [26] described a simple and effective method for quantification of diacetyl by use of HS-SPME coupled to GC–MS. Other

workers have described methods for the quantification of diacetyl and other dicarbonyl compounds [14], and several aldehydes [20, 21, 59], by use of derivatisation procedures before chromatographic analysis. Because the effects of MLF on aldehydes have not yet been characterised, mainly because of a lack of analytical data, a robust method for simultaneous determination of diacetyl and aldehydes in wine would provide insight into changes associated with LAB during MLF.

The purpose of this study was to investigate the effect of MLF on the chemical composition, in terms of volatile aroma compounds and organic acids, of Pinotage and Shiraz wine from South Africa by using four commercial *O. oeni* starter cultures. The results would provide a better understanding of the contribution of MLF to the composition of wine and its potential contribution to wine aroma.

Materials and methods

Chemical standards and reagents

All standards (Table 1) were of analytical grade (purity 95–99.9%) and purchased from Fluka (Buchs, Switzerland), Sigma-Aldrich (Steinheim, Germany), Riedel-de-Häen (Seelze, Germany), and Merck (Darmstadt, Germany). Sodium chloride (HPLC quality) and diethyl ether (99.5%) were purchased from Merck (Darmstadt, Germany) and pure water was obtained from a Milli-Q purification system (Millipore, Bedford, MA, USA). The internal standards (2-pentanone and 4-methyl-2-pentanol, Fluka) and volatile standards were dissolved in a wine simulant (12% v/v ethanol) prepared according to Louw et al. [38] and used for the respective calibration curves and subsequent validation procedures as reported in the supplementary material.

Bacterial strains

The four commercial starter cultures used for this study are listed in Table 2. These cultures were selected because they are frequently used in the South African wine industry.

Experimental design of winemaking

The experimental design for the winemaking experiments was the same for both years (2008, 2009) and cultivars (Pinotage, Shiraz). Alcoholic fermentation (1,000 l) was followed by MLF performed in triplicate (4.5 l × 3) for each of the different treatments, namely control (no MLF), spontaneous (spon), Enoferm alpha[®] (A), Lalvin VP41[®] (V), Viniflora oenos[®] (O), and Viniflora CH16[®] (C). The control wine treatment (no MLF; lysozyme added to inhibit

Table 1 Odour threshold (OTH) values (mg/l) and descriptions as reported in the literature (the Ref. no. is given in parentheses)

Analyte	OTH (mg/L)	Odour description	Source
Esters			
Ethyl decanoate	0.2 [22]	Grape, floral, soap [23]	Aldrich, >99%
Ethyl hexanoate	0.014 [22]	Fruity, anise [18]	Fluka, 99%
Ethyl butyrate	0.02 [25]	Fruity [17], apple [23]	Fluka, >98%
Ethyl octanoate	0.005 [22]	Fruit, fat [23]	Fluka, >98%
Ethyl lactate	154.6 [19]	Butter, cream, fruit [23]	Fluka, 99%
Ethyl propionate	1.8 [19]	Fruity [17]	Fluka, >99.7%
Ethyl 2-methylpropanoate	0.015 [22]	Fruity [17]	Fluka, >98%
Ethyl 2-methylbutyrate	0.018 [22]	Fruity [17], apple [23]	Aldrich, >98%
Ethyl isovalerate	0.003 [22]	Fruity, anise [17]	Fluka, >99.7%
Ethyl 3-hydroxybutanoate	20	Strawberry, burnt marshmallow [56]	Fluka, >97%
Ethyl phenylacetate	0.65	Rose, floral [23]	Fluka, >99%
Ethyl acetate	12.26 [19]	Fruit, nail polish [23]	Sigma-Aldrich, >99.7%
Isoamyl acetate	0.03 [25]	Banana, pear [17]	Riedel de Haën, >98%
Hexyl acetate	1.5 [19]	Sweet, perfume [55]	Fluka, 99%
2-Phenylethyl acetate	0.25 [25]	Roses [17]	Fluka, >99%
Diethyl succinate	200 [19]	Berry [23]	Fluka, >98%
2-Methylpropyl acetate	1.6	Solvent [17]	Fluka, >99.8%
Alcohols			
Hexanol	8 [25]	Green, grass, resin [17]	Merck, >98%
Butanol	150 [19]	Fusel, spirituous [23]	Fluka, >99.5%
Methanol	500 [50]	Alcohol [50]	Sigma-Aldrich, >99.9%
2-Phenylethanol	14 [22]	Honey, spice, rose, lilac [23]	Merck, >99%
Propanol	306	Pungent, harsh [23]	Fluka, >99%
Isobutanol	40 [25]	Wine, solvent, bitter [23]	Fluka, >99.5%
Isoamyl alcohol	30 [25]	Fusel [17], whiskey, malt, burnt [23]	Aldrich, >99%
Pentanol	64 [19]		Fluka, >99.8%
4-Methyl-1-pentanol	1 [60]		Sigma-Aldrich, >95%
3-Methyl-1-pentanol	1 [60]		Sigma-Aldrich, >97%
3-Ethoxy-1-propanol	0.1 [50]	Fruity [50]	Sigma-Aldrich, >97%
Acids and fatty acids			
Acetic acid	200 [25]	Vinegar [17]	Saarchem, >98%
Propionic acid	20	Pungent, rancid, sweat [23]	Fluka, >99.5%
Isobutyric acid	2.3 [22]	Rancid, butter, cheese [23]	Fluka, >99.5%
Butyric acid	0.173 [22]	Cheese [17]	Fluka, >99.5%
Isovaleric acid	0.033 [22]	Cheese [17]	Fluka, >99%
Valeric acid			Fluka, >99%
Hexanoic acid	0.42 [22]	Sweat [23]	Aldrich, >99.5%
Octanoic acid	0.50 [22]	Sweat, cheese [23]	Aldrich, >99.5%
Decanoic acid	1 [22]	Rancid, fat [23]	Sigma, >98%
Carbonyl compounds			
Diacetyl (2,3-butanedione)	0.1 ^a [25]	Butter, cream [17]	Fluka, >99.5%
Acetoin (3-hydroxy-2-butanone)	150 [19]	Butter, cream [22]	Fluka, >97%
2,3-Pentanedione	0.9 [50]	Butter, cream [17]	Fluka, >95%
<i>E</i> -2-Hexenal	0.01 [8]	Herbaceous, green [12]	Fluka, >97%
<i>E</i> -2-Heptenal	0.013 [8]	Herbaceous [12]	Fluka, >96%
Octanal	0.05 [60]	Herbaceous [12], fatty, citrus [60]	Fluka, >98%
<i>E</i> -2-Octenal	0.0001 [8]	Lemon [17]; herbaceous [12]	Aldrich, >94%

Table 1 continued

Analyte	OTH (mg/L)	Odour description	Source
Nonanal	0.001 [20]	Herbal, floral	Fluka, >95%
<i>E</i> -2-Nonenal	0.000068 [7]	Sawdust, plank [7]	Aldrich, 97%
Decanal	0.0001 [20]	Citrus, fruity	Sigma, >98%
<i>trans</i> -2, <i>cis</i> -6-Nonadienal	0.00001	Cucumber, green	Aldrich, 95%

^a Shown to range from 0.2 mg/L to 2.8 mg/L depending on wine style [40]

Table 2 Sensory attributes of the commercial starter cultures, according to the respective manufacturers

Starter culture	Abbreviation	Company	Sensory contribution description in brief ^a
Enoferm alpha [®]	A	Lallemand	Mouthfeel, lower perception of green and vegetative flavours, positive effect on wine complexity
Lalvin VP41 [®]	V	Lallemand	Enhances complexity and mouthfeel, contributes to aroma and wine structure
Viniflora [®] oenos	O	Chr Hansen	Clean and classic flavour profile, low production of volatile acidity
Viniflora [®] CH16	C	Chr Hansen	Low production of volatile acidity

Abbreviations for the starter cultures used during this study are listed

^a Information obtained from the respective technical data sheets: www.chr-hansen.com; www.lallemandwine.com

LAB growth) and spontaneous treatment (no MLF inoculation) were subjected to the same experimental design and included for comparative purposes. Subsequent chemical analysis of each of the biological triplicates was performed in duplicate or triplicate, depending on the method of analysis.

One-hundred and seventy kilograms of Shiraz grapes were harvested at 25.0°B during the 2008 season from the Wellington region, Western Cape, South Africa. Pinotage grapes were harvested from the same region in South Africa at 28.4°B (170 kg) and 22.0°B (175 kg) during the 2008 and 2009 seasons, respectively. After crushing and destemming, 30 mg/l sulfur dioxide (SO₂) was added to the must to reduce possible growth of natural flora present on the grapes and to prevent oxidation. Alcoholic fermentation was performed in a 1,000 l stainless-steel tank at 25°C. A commercial strain of actively dried *Saccharomyces cerevisiae*, WE372 (Anchor Yeast, South Africa) was inoculated for alcoholic fermentation at 0.3 g/l after rehydration, in accordance with the manufacturer's specifications. Lysozyme (DSM Food Specialties, Oenology, France) was added at 0.25 g/l to inhibit indigenous LAB microflora. During alcoholic fermentation the skins were punched down manually twice a day. At 3°B the wine was removed from the skins by light pressing, using only the free-flow wine, to reduce hard tannins present in the wine. After completion of alcoholic fermentation (residual sugar less than 5 g/l) the wine was divided into 4.5-l glass bottles for MLF. MLF was performed at 20°C in triplicate for each of the respective treatments, namely spontaneous MLF, four commercial starter cultures, and a control treatment. The

spontaneous MLF treatment was not inoculated with a starter culture and no SO₂ was added. This treatment was included to evaluate whether any natural flora could have potentially contributed to the MLF process and whether the lysozyme treatment was effective. Commercial starter cultures were rehydrated and inoculated, in accordance with the manufacturers' specifications, at 0.01 g/l. For the control treatment, three of the 4.5-l glass bottles were racked and 50 mg/l SO₂ was added directly after alcoholic fermentation to inhibit microbial growth and enable capture of the chemical composition of the wines before MLF. Bacterial complex nutrients were added in accordance with each manufacturer's instructions: 0.2 g/l Optimalo (Lallemand, Stellenbosch, South Africa) for the Lalvin VP41 and Enoferm alpha cultures and 0.1 g/l Bactiv-aid (Chr Hansen, Hørsholm, Denmark) for the Viniflora oenos and Viniflora CH16 cultures. MLF was regarded as complete at malic acid concentrations less than 0.3 g/l. After MLF, all wines were racked, SO₂ levels adjusted to 50 mg/l, and the wines were bottled. Wines were stored at 15°C before all chemical analysis.

Microbial enumeration

Microbial populations for LAB were monitored to evaluate the effectiveness of the inoculated commercial cultures and to establish if other LAB species survived and could potentially contribute to MLF. For this purpose, LAB were determined by plating 100 µl of a dilution series of the wines, prepared in sterile distilled water, on selective media. MRST plates contained 50 g/l De Man, Rogosa, and Sharpe

(MRS; Biolab, Merck, Wadeville, South Africa), 20 g/l bacteriological agar (Biolab, Merck) supplemented with 10% preservative-free tomato juice (All Gold, South Africa) and pH adjusted to 5.0 with hydrochloric acid (HCl). MRS plates contained 50 g/l MRS broth (Biolab, Merck) and 15 g/l bacteriological agar (Biolab, Merck). All plates contained 50 mg/l Delvocid Instant (DSM Food Specialties, The Netherlands) to prevent yeast growth and 25 mg/l kanamycin sulfate (Roche Diagnostics, Mannheim, Germany) to suppress the growth of acetic acid bacteria. MRST, which favours the growth of *O. oeni*, was used for enumeration of *O. oeni* whereas MRS agar was used for enumeration of other wine LAB. Agar plates were incubated at 30°C for 5–7 days after which colony-forming units per ml (cfu/ml) were determined. All LAB were anaerobically cultivated by use of Microbiology Anaerocult pads in anaerobic jars (Merck, Darmstadt, Germany).

Fourier transform mid-infrared spectroscopy (FT-MIR)

FT-MIR spectra acquired by use of a Winescan FT120 instrument (FOSS Analytical software version 2.2.1) equipped with Winescan FT120 2001 version 2.2.1 software, in accordance with the method described elsewhere [38].

Quantified chemical data, including pH, ethanol, titratable acidity (TA), volatile acidity (VA), glycerol, and residual sugar (RS), were predicted from infrared spectra by use of commercial or in-house developed calibrations as described elsewhere [38].

Organic acid analysis

Malic acid, lactic acid, pyruvic acid, gluconic acid, acetic acid, succinic acid, and citric acid were quantified before and after MLF by use of a modified version of the certified OIV reference method [46]. The original OIV method was modified (running buffer containing 5% acetonitrile compared with 10% in the original) to include more analytes for quantification. Samples were diluted 1/25 in the running buffer before injection. A 3D CE instrument (Agilent Technologies, Waldbronn, Germany) equipped with Agilent Chemstation software version B.01.03 (204) was used for the analysis and data processing in accordance with the certified OIV method [46]. Calibration ranges were between 0.04 and 2 g/l for all compounds except pyruvic acid, for which the upper limit was 1 g/l.

Volatile aroma compound analysis

Major volatile aroma compounds

Volatile higher alcohols, esters, fatty acids, and carbonyl compounds were analysed in triplicate with a Hewlett–

Packard (Little Falls, USA) 6890 Plus gas chromatograph equipped with a split/splitless injector and flame-ionisation detection (FID), following a newly developed fast GC procedure. In brief, volatile compounds were isolated from 5 ml wine, after addition of 10 mg/l 4-methyl-2-pentanol ($\geq 97\%$) as internal standard, by liquid–liquid extraction with diethyl ether [38]. Analysis of the different compounds was achieved by separation using a J&W DB-FFAP capillary GC column (Agilent, Little Falls, Wilmington, USA) with dimensions 20 m length \times 0.1 mm inside diameter \times 0.2 μ m film thickness followed by FID. Analyte concentrations were calculated by comparing their respective peak areas with those from calibration standard curves, by use of a data-handling system (HP GC Chemstation, revision A.07.01 (682)).

Carbonyl compounds

A headspace solid-phase microextraction gas chromatography–mass spectrometry (HS-SPME GC–MS) method was developed to quantify the carbonyl compounds diacetyl, acetoin, and 2,3-pentanedione, and a selection of aldehydes including hexanal, *E*-2-hexenal, decanal, octanal, *E*-2-octenal, *E*-2-nonenal, and *cis*-2, *trans*-6-nonadienal. A selection of method-validation data is provided in the Supplementary Material available for this paper.

Headspace solid-phase microextraction was performed with a 60- μ m poly(ethylene glycol) (PEG) SPME fibre (Supelco, Bellefonte, PA, USA), specific for extraction of polar compounds from the headspace. Glass screw-cap vials with polytetrafluoroethylene (PTFE)/silicone septa (20 mm) (Agilent Technologies, Little Falls, Wilmington, USA) were used. After optimisation, SPME analysis was performed on a mixture of 1 ml wine, 9 ml distilled Milli-Q water (Millipore), and 2 g sodium chloride (NaCl; Sigma) in a 20-ml vial. The internal standard, 2-pentanone, was added at 10 mg/l to each vial. The wine was agitated to ensure that NaCl dissolved completely. Extraction of volatiles from the headspace was performed at 50°C for 10 min. Subsequently, the fibre was desorbed in the hot injection port of the GC–MS at 220°C for 2 min. The injector was operated in pulsed split mode (300 kPa, split ratio 10:1) at 220°C for 2 min and 171 kPa afterwards. Each wine was analysed in duplicate.

Separation was performed on a 60 m length \times 0.25 mm i.d. \times 0.25 μ m f.t. FFAP column (Agilent Technologies, Little Falls, Wilmington, USA) using a 6890 gas chromatograph coupled to a 5975C mass spectrometer (Agilent Technologies) and equipped with Enhanced Chemstation version D.01.02.16 software (Agilent Technologies). For SPME sample preparation and injection, the instrument was equipped with a CTC CombiPal autosampler (CTC Analytics, Switzerland) and used with the SPME option. The carrier

gas (helium) flow through the GC column was 1.7 ml/min and the oven was programmed from 35°C (2 min), ramped at 5°/min to 150°C (held for 2 min) and ramped at 15°/min to 240°C (held for 1 min). The mass spectrometer (MS) was operated in electron-impact (EI) mode (70 eV). Data acquisition was performed in SIM mode by monitoring the mass-to-charge (m/z) ratios representing unique ion fragments for the respective compounds: 2-pentanone (IS; 43, 86); diacetyl (43, 86); 2,3-pentanedione (57, 100); *E*-2-hexenal (69, 83, 98); octanal (69, 84, 110); acetoin (45, 88); nonanal (82, 98, 114); *E*-2-octenal (70, 83, 97); nonanal (82, 95, 112); decanal (82, 95, 112); *E*-2-nonenal (83, 70, 96); *cis*-2,*trans*-6-nonadienal (69, 70, 81). Peak identification of the volatile components was achieved by comparison of retention times after injection of pure, authentic standards.

Data analysis

Data were subjected to one-way analysis of variance (ANOVA) followed by Fisher's least significant difference (LSD) test to determine whether differences between samples were significant, using XLStat software version 2009.1.02 (Addinsoft, www.xlstat.com). Differences between samples with a significance level of 5% ($P \leq 0.05$) were considered significant [49; SAS, 2002]. To obtain a more comprehensible overview of the volatile aroma compounds and to investigate possible correlations amongst the analytes, multivariate data analysis techniques [45], including principal component analysis (PCA) and partial least-squares discriminant analysis (PLS-DA), were performed with the *Unscrambler* software (version 9.2.1, Camo ASA, Norway). Data were pretreated by autoscaling to eliminate differences in measurement units.

Results and discussion

Monitoring MLF

Malolactic fermentation in the wines was induced with four different LAB strains, after completion of alcoholic fermentation with *S. cerevisiae* WE372. The alcohol content, pH, and malic acid concentration after alcoholic fermentation were 13.20% (v/v), 3.95, and 2.50 g/l, respectively. MLF resulted in a decrease in titratable acidity (TA), and increases in pH and volatile acidity (VA), all of which are known to be typically associated with MLF, as reported widely [9, 27].

Organic acid profiles

After completion of MLF in the Pinotage 2008 wine, no statistically significant (95% confidence level) differences

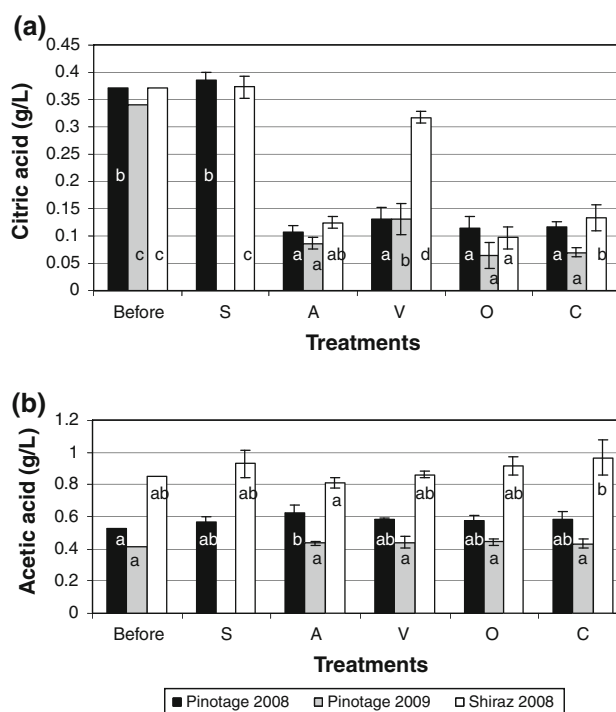


Fig. 1 Changes observed in the **a** citric acid and **b** acetic acid concentrations as a result of MLF with different starter cultures (A, V, O, C) compared to a spontaneous (S) and control wine (no MLF: indicated by the word “Before”). Enoferm alpha: A; Lalvin VP41: V; Viniflora oenos: O; Viniflora CH16: C. Different alphabetical letters indicate significant differences

were observed among the bacterial starter cultures for the malic, lactic, and citric acid (Fig. 1a) profiles. Citric acid consumption is directly involved in the production of diacetyl via the citric acid pathway [3]. No significant difference was observed among the four commercial starter cultures in terms of the citric acid concentrations left in the wine after MLF. Therefore it is not surprising that the diacetyl concentrations were similar (Table 3). No changes in the tartaric acid concentrations were observed during MLF (data not shown). In terms of acetic acid formation (Fig. 1b), the Enoferm alpha strain produced significantly more acetic acid than the control (in Pinotage 2008), although compared with the other bacterial strains there was no significant difference.

In the Pinotage 2009 and Shiraz 2008 wines, the Lalvin VP41 strain consumed significantly less citric acid than the other three strains during MLF (Fig. 1a). This could possibly suggest less metabolic activity towards citric acid and, consequently, lower diacetyl concentrations could be expected for this specific strain [3]. No significant changes in the acetic acid (Fig. 1b) and tartaric acid (data not shown) concentrations were observed in the Pinotage 2009 wine. After MLF in the Shiraz 2008 wine, significant differences were observed among the different bacterial

starter cultures in terms of citric acid (Fig. 1a) and acetic acid (Fig. 1b). Viniflora CH16 produced the highest acetic acid concentration and the Enoferm alpha strain produced the lowest acetic acid concentration. The other bacteria, control wine, and spontaneous fermentation did not differ significantly from each other in terms of acetic acid concentration.

Effect of MLF on volatile composition

The concentrations of the 48 volatile compounds determined in the control wine (before MLF) and after MLF are listed in Table 3 for the Pinotage 2008 and 2009 wines and in Table 4 for the Shiraz wine. Analysis of variance (ANOVA) showed concentration differences were significant ($P < 0.05$) for 30 compounds in the Pinotage 2008 wine (Table 3), 28 compounds in the Pinotage 2009 wine (Table 3), and 34 compounds in the Shiraz wine (Table 4) as a consequence of MLF. Different alphabetical letters indicate significant differences ($P < 0.05$) among the average values obtained for each of the LAB strains used to perform MLF. This outcome is in agreement with previous studies on other red grape varieties [39, 51].

The odour thresholds (OTH) reported in the literature, aroma descriptor, and supplier information for each compound are listed in Table 1. Odour activity values (OAVs), listed in Tables 3 and 4, were calculated by dividing the mean concentration of a compound by its odour threshold value as reported in the literature [25]. This indicates that the volatile compounds with $OAV > 1$ could potentially make an contribution to the odour of the wine [25]. Of the 48 volatile compounds quantified, 18 analytes had $OAVs > 1$ in the Pinotage 2008 wine (Table 3), 20 analytes had $OAVs > 1$ in the Pinotage 2009 wine (Table 3), and 23 analytes had $OAVs > 1$ in the Shiraz 2008 wine (Table 4). These compounds include ethyl hexanoate, ethyl butyrate, ethyl octanoate, ethyl 2-methylpropanoate, ethyl 2-methylbutyrate, ethyl isovalerate, ethyl acetate, isoamyl acetate, 2-phenylethanol, isoamyl alcohol, 3-ethoxy-1-propanol, acetic acid, butyric acid, isovaleric acid, hexanoic acid, octanoic acid, diacetyl (2,3-butanedione), 2,3-pentanedione, *E*-2-hexenal, *E*-2-octenal, *E*-2-nonenal, and *trans*-2,*cis*-6-nonadienal. However, volatile compounds with high OAVs do not always have an effect on the aroma of wine and this information only shows potential contribution to the aroma [17] by individual analytes.

The PCA scores plot and corresponding loadings plot in Fig. 2a, b provide an overview of the volatile profiles associated with the metabolic activity of the four starter cultures during MLF in the Pinotage 2008 wine in terms of esters, alcohols, and acids. Separation along the first principal component (PC1) seems to be driven by the association of the Viniflora CH16 strain (positioned to the left of the

scores plot in Fig. 2a) with a selection of ethyl esters, namely ethyl 2-methylpropanoate (fruity), ethyl propionate (fruity), ethyl isovalerate (fruity, anise), and ethyl butyrate (fruity, apple) (Fig. 2b). The Enoferm alpha and Lalvin VP41 strains, and the Viniflora oenos strain to some extent, are positioned toward the right of the scores plot (Fig. 2a) along PC3 and are associated with ethyl hexanoate (fruity, anise), ethyl lactate (butter, cream, fruit), 2-phenylethanol (honey, rose), 3-ethoxy-1-propanol (fruity), and diethyl succinate (berry). Along the second PC, Viniflora oenos, positioned toward the bottom of the scores plot (Fig. 2a), is separated from the other bacteria in terms of its association with acetic acid (vinegar), propionic acid (pungent, rancid, sweat), octanoic acid (sweat, cheese), isovaleric acid (cheese), hexanoic acid (sweat), decanoic acid (rancid, fat), butyric acid (cheese), isobutyric acid (rancid, butter, cheese), isobutanol (wine, solvent), propanol (pungent, harsh), butanol (fusel, spirituous), hexanol (green, grass, resin), 3-methyl-1-pentanol, isoamyl alcohol (fusel, whisky, malt), ethyl acetate (fruit, nail polish), ethyl decanoate (grape, floral, soap), isoamyl acetate (banana, pear), ethylphenyl acetate (rose, floral), and 2-methyl-propyl acetate (solvent). The VP41, CH16, and Enoferm alpha strains are positioned toward the top of the scores plot (Fig. 2a) along the second PC and associated with ethyl octanoate (fruit, fat), 2-phenylethyl acetate (roses), ethyl 2-methylbutyrate (fruity, apple), diethyl succinate (berry), and ethyl 3-hydroxybutanoate (strawberry, burnt marshmallow).

PCA results (Fig. 2c, d) for the volatile profiles obtained during MLF of the Pinotage 2009 wine illustrate less prominent strain discrimination in terms of the esters, higher alcohols, and acid profiles associated with the four different bacteria. The Enoferm alpha strain is positioned more to the right of the scores plot (Fig. 2c) along PC3 and toward the bottom of the plot along PC2. This position seems to be driven by association with acetoin (butter, cream), diethyl succinate (berry), ethyl propionate (fruity), ethyl decanoate (grape, floral, soap), valeric acid, and ethyl 2-methylbutyrate (fruity, apple) (Fig. 2d). The remaining three strains are positioned more to the left of the scores plot (Fig. 2c) along PC3 and slightly nearer the top of the plot along the second PC. These strains are associated with ethyl butyrate (fruity, apple), 4-methyl-1-pentanol, ethyl hexanoate (fruity, anise), ethyl octanoate (fruit, fat), ethyl 3-hydroxybutyrate (strawberry, burnt marshmallow), hexanol (green, grass), acetic acid (vinegar), isoamyl alcohol (fusel, whisky), isoamyl acetate (banana, pear), octanoic acid (sweat, cheese), hexanoic acid (sweat), decanoic acid (rancid, fat), hexyl acetate (sweet, perfume), and 2-methylpropyl acetate (solvent), with concentrations dependent on the strain used.

PCA results for the Shiraz 2008 wine show clear differentiation amongst the four different bacterial starter cultures (Fig. 2e). Separation along the second PC seems to

Table 3 Concentrations of different volatiles after use of different bacterial starter cultures, with odour activity values (OAV) calculated for the Pinotage 2008 and 2009 wines

Analyte	Pinotage 2008									
	Control		Bacteria A		Bacteria V		Bacteria O		Bacteria C	
	Average	OAV	Average	OAV	Average	OAV	Average	OAV	Average	OAV
Esters										
Ethyl decanoate	0.00c	0.0	0.08a	0.42	0.08a	0.4	0.08b	0.4	0.08b	0.4
Ethyl hexanoate	0.48b	33.9	0.56a	40.12	0.57a	40.6	0.51b	36.4	0.51b	36.2
Ethyl butyrate	0.48a	24.0	0.48a	24.18	0.49a	24.4	0.49a	24.5	0.49a	24.3
Ethyl octanoate	0.18d	36.7	0.29a	58.69	0.30a	59.5	0.22c	43.2	0.21c	41.3
Ethyl lactate	0.00e	0.0	33.05a	0.21	29.55b	0.2	22.35c	0.1	19.93d	0.1
Ethyl propionate	0.000b	0.0	0.355ab	0.20	0.706a	0.4	0.178ab	0.1	0.819a	0.5
Ethyl 2-methylpropanoate	0.000b	0.0	0.191b	12.71	0.099b	6.6	0.121b	8.0	0.697a	46.4
Ethyl 2-methylbutyrate	0.000b	0.0	0.267a	14.83	0.081ab	4.5	0.041ab	2.3	0.084ab	4.7
Ethyl isovalerate	0.407a	135.6	0.392a	130.55	0.406a	135.5	0.420a	139.9	0.437a	145.7
Ethyl 3-hydroxybutanoate	1.074b	0.1	2.871a	0.14	1.577ab	0.1	1.758ab	0.1	1.270ab	0.1
Ethyl phenylacetate	1.249b	1.9	1.226	1.89	1.308a	2.0	1.346a	2.1	1.234b	1.9
Ethyl acetate	59.56a	4.9	53.29a	4.35	52.73a	4.3	47.74a	3.9	47.37a	3.9
Isoamyl acetate	2.01a	66.9	1.93a	64.42	2.03a	67.6	1.99a	66.5	1.97a	65.6
Hexyl acetate	0.166a	0.1	0.120b	0.08	0.126ab	0.1	0.122b	0.1	0.110b	0.1
2-Phenylethyl acetate	0.07a	0.3	0.07a	0.28	0.07a	0.3	0.07a	0.3	0.06a	0.2
Diethyl succinate	0.54c	0.0	0.85a	0.00	0.70b	0.0	0.60c	0.0	0.54c	0.0
2-Methylpropyl acetate	0.622a	0.4	0.614a	0.38	0.661a	0.4	0.677a	0.4	0.659a	0.4
Alcohols										
Hexanol	0.59c	0.1	0.66ab	0.08	0.67a	0.1	0.59c	0.1	0.56c	0.1
Butanol	1.81ab	0.0	1.90a	0.01	1.87a	0.0	1.83ab	0.0	1.68b	0.0
Methanol	29.02ab	0.1	30.08a	0.06	25.65abc	0.1	22.36bc	0.0	21.28c	0.0
2-Phenylethanol	24.06ab	1.7	25.14a	1.80	24.93a	1.8	23.07b	1.6	21.38c	1.5
Propanol	79.80a	0.3	80.12a	0.26	76.32ab	0.2	76.35ab	0.2	69.37b	0.2
Isobutanol	24.34ab	0.6	24.42ab	0.61	23.72ab	0.6	22.43bc	0.6	20.90c	0.5
Isoamyl alcohol	171.35a	5.7	171.91a	5.73	171.28a	5.7	155.85b	5.2	151.61b	5.1
Pentanol	0.21a	0.0	0.17ab	0.00	0.19a	0.0	0.18ab	0.0	0.13ab	0.0
4-Methyl-1-pentanol	0.02a	0.0	0.02a	0.02	0.01a	0.0	0.01a	0.0	0.02a	0.0
3-Methyl-1-pentanol	0.04b	0.0	0.03b	0.03	0.05ab	0.0	0.08a	0.1	0.06 ab	0.1
3-Ethoxy-1-propanol	3.45b	34.5	4.34a	43.44	3.69ab	36.9	3.84ab	38.4	3.33b	33.3
Acids and fatty acids										
Acetic acid	214.96f	1.1	421.52a	2.11	395.97b	2.0	369.46c	1.8	344.92d	1.7
Propionic acid	11.46b	0.6	10.83bc	0.54	10.18bcd	0.5	9.83 cd	0.5	9.19d	0.5
Isobutyric acid	0.82bc	0.4	0.87ab	0.38	0.89a	0.4	0.79c	0.3	0.76c	0.3
Butyric acid	1.09ab	6.3	1.16a	6.71	1.15ab	6.6	1.08bc	6.2	1.00c	5.8
Isovaleric acid	0.71bc	21.5	0.77ab	23.35	0.78ab	23.7	0.71bc	21.6	0.65c	19.6
Valeric acid	0.42ab		0.43ab		0.45a		0.43ab		0.43ab	
Hexanoic acid	1.52bc	3.6	1.68a	3.99	1.68a	4.0	1.60ab	3.8	1.47c	3.5
Octanoic acid	0.95d	1.9	1.57 a	3.14	1.54a	3.1	1.44b	2.9	1.37b	2.7
Decanoic acid	0.23d	0.2	0.41a	0.41	0.40ab	0.4	0.40a	0.4	0.37b	0.4
Carbonyl compounds										
Diacetyl (2,3-butanedione)	7.45a	74.5	7.08a	70.83	6.55a	65.5	7.97a	79.7	6.71a	67.1
Acetoin (3-hydroxy-2-butanone)	4.51ab	0.0	4.35ab	0.03	3.69a	0.0	4.88b	0.0	4.20ab	0.0
2,3-Pentanedione	2.51e	2.8	1.29c	1.43	1.23bc	1.4	1.14a	1.3	1.15ab	1.3

Table 3 continued

Analyte	Pinotage 2008									
	Control		Bacteria A		Bacteria V		Bacteria O		Bacteria C	
	Average	OAV	Average	OAV	Average	OAV	Average	OAV	Average	OAV
<i>E</i> -2-Hexenal	0.002a	0.2	0.0081a	0.81	0.0a	0.0	0.002a	0.2	0.001a	0.1
<i>E</i> -2-Heptenal	nd		nd		nd		nd	nd		nd
Octanal	nd		nd		nd		nd	nd		nd
<i>E</i> -2-Octenal	nd		0.0003a	3.33	nd	nd	nd	nd	nd	nd
Nonanal	nd		0.0009b	0.88	nd	nd	nd	nd	nd	nd
<i>E</i> -2-Nonenal	nd		nd		nd	nd	nd	nd	nd	nd
Decanal	nd		0.0002a	2.27	nd	nd	nd	nd	nd	nd
<i>trans</i> -2, <i>cis</i> -6-Nonadienal	0.002a	237.8	0.0018a	181.98	0.0013a	132.9	0.002a	243.2	0.002a	199.8
Analyte	Pinotage 2009									
	Control		Bacteria A		Bacteria V		Bacteria O		Bacteria C	
	Average	OAV	Average	OAV	Average	OAV	Average	OAV	Average	OAV
Esters										
Ethyl decanoate	0.18a	0.9	0.18a	0.9	0.18a	0.9	0.18a	0.9	0.18a	0.9
Ethyl hexanoate	0.47b	33.5	0.44a	31.1	0.44a	31.6	0.44a	31.3	0.47b	33.5
Ethyl butyrate	0.11a	5.6	0.11a	5.6	0.11a	5.5	0.12ab	5.8	0.13b	6.3
Ethyl octanoate	0.25a	49.7	0.25a	49.7	0.25a	49.7	0.25a	49.2	0.28a	55.2
Ethyl lactate	13.73b	0.1	36.70a	0.2	40.27c	0.3	37.37a	0.2	36.54a	0.2
Ethyl propionate	0.00b	0.0	0.28ab	0.2	0.61a	0.3	0.03b	0.0	0.13b	0.1
Ethyl 2-methylpropanoate	0.19a	12.8	0.36a	23.7	0.47a	31.4	0.30a	19.8	0.30a	19.8
Ethyl 2-methylbutyrate	1.09a	60.5	0.13b	7.2	0.05b	2.6	0.04b	2.5	0.06b	3.3
Ethyl isovalerate	0.38a	125.2	0.29a	97.4	0.37a	124.4	0.40a	131.9	0.43a	141.7
Ethyl 3-hydroxybutanoate	3.72a	0.2	1.19b	0.1	1.01b	0.1	1.22b	0.1	1.94b	0.1
Ethyl phenylacetate	1.07a	1.7	1.06a	1.6	1.23a	1.9	1.00a	1.5	0.99a	1.5
Ethyl acetate	24.36a	2.0	25.01a	2.0	25.84ab	2.1	27.72b	2.3	27.93b	2.3
Isoamyl acetate	1.29b	43.1	1.22a	40.7	1.23a	40.9	1.25ab	41.6	1.29b	43.0
Hexyl acetate	0.45a	0.3	0.32a	0.2	0.20a	0.1	0.24a	0.2	0.20a	0.1
2-Phenylethyl acetate	0.59b	2.4	0.58a	2.3	0.58a	2.3	0.58a	2.3	0.58a	2.3
Diethyl succinate	0.35c	0.0	0.38ab	0.0	0.38ab	0.0	0.38a	0.0	0.41b	0.0
2-Methylpropyl acetate	0.59a	0.4	0.52a	0.3	0.53a	0.3	0.60a	0.4	0.64a	0.4
Alcohols										
Hexanol	1.50a	0.2	1.52ab	0.2	1.50a	0.2	1.51ab	0.2	1.54b	0.2
Butanol	0.96a	0.0	0.97a	0.0	0.97a	0.0	1.00a	0.0	0.99a	0.0
Methanol	43.59c	0.1	40.36bc	0.1	36.89abc	0.1	33.60ab	0.1	31.35a	0.1
2-Phenylethanol	32.20a	2.3	32.29a	2.3	31.72a	2.3	31.57a	2.3	32.02a	2.3
Propanol	43.93a	0.1	42.86a	0.1	44.27ab	0.1	46.29b	0.2	44.75ab	0.1
Isobutanol	24.92ab	0.6	24.46a	0.6	24.77ab	0.6	25.69b	0.6	25.02ab	0.6
Isoamyl alcohol	175.29a	5.8	174.34a	5.8	173.77a	5.8	176.81a	5.9	176.11a	5.9
Pentanol	0.11a	0.0	0.04a	0.0	0.12a	0.0	0.13a	0.0	0.18a	0.0
4-Methyl-1-pentanol	0.15a	0.1	0.01b	0.0	0.05b	0.0	0.04b	0.0	0.05b	0.0
3-Methyl-1-pentanol	0.11ab	0.1	0.06b	0.1	0.09ab	0.1	0.07 b	0.1	0.16a	0.2
3-Ethoxy-1-propanol	3.91a	39.1	3.48a	34.8	3.88a	38.8	3.59a	35.9	5.01a	50.1

Table 3 continued

Analyte	Pinotage 2009									
	Control		Bacteria A		Bacteria V		Bacteria O		Bacteria C	
	Average	OAV	Average	OAV	Average	OAV	Average	OAV	Average	OAV
Acids and fatty acids										
Acetic acid	160.15b	0.8	256.72a	1.3	257.21a	1.3	269.29a	1.3	267.93a	1.3
Propionic acid	3.55a	0.2	3.65ab	0.2	3.89ab	0.2	3.98b	0.2	3.85ab	0.2
Isobutyric acid	1.59a	0.7	1.77a	0.8	1.77ab	0.8	2.17c	0.9	1.95b	0.8
Butyric acid	0.68a	3.9	0.71b	4.1	0.70ab	4.0	0.68a	3.9	0.70ab	4.0
Isovaleric acid	1.83a	55.6	2.00b	60.7	1.77a	53.7	1.79a	54.2	1.79a	54.1
Valeric acid	0.43ab		0.72a		0.72a		0.72a		0.72a	
Hexanoic acid	2.44a	5.8	2.51ab	6.0	2.45ab	5.8	2.47ab	5.9	2.54b	6.0
Octanoic acid	3.81c	7.6	4.05ab	8.1	3.95 a	7.9	3.99ab	8.0	4.11b	8.2
Decanoic acid	0.97a	1.0	0.95a	1.0	0.94a	0.9	0.94a	0.9	0.97a	1.0
Carbonyl compounds										
Diacetyl (2,3-butanedione)	5.19b	51.9	14.45a	144.5	7.19b	71.9	14.99a	149.9	13.35a	133.5
Acetoin (3-hydroxy-2-butanone)	3.08a	0.0	12.01b	0.1	2.94a	0.0	5.81a	0.0	4.82a	0.0
2,3-Pentanedione	1.40b	1.6	1.29a	1.4	1.33ab	1.5	1.28a	1.4	1.31a	1.5
<i>E</i> -2-Hexenal	0.0015a	0.1	0.0005a	0.1	0.0009a	0.1	0.0024a	0.2	nd	
<i>E</i> -2-Heptenal	nd	nd	nd		nd		nd		nd	
Octanal	nd	nd	nd		nd		nd		nd	
<i>E</i> -2-Octenal	0.0009ab	8.6	nd		0.0014b	14.2	nd		nd	
Nonanal	nd		nd		nd		nd		nd	
<i>E</i> -2-Nonenal	nd		nd		nd		nd		nd	
Decanal	nd		0.0003a	3.5	0.0001a	1.4	nd		nd	
<i>trans</i> -2, <i>cis</i> -6-Nonadienal	0.0017a	172.8	0.0006a	57.6	0.0012a	121.9	0.0007a	73.5	0.0011a	111.8

Averages are expressed as milligrams per litre (mg/l). Different alphabetic letters in the same row indicate significant differences ($P < 0.05$)

discriminate VP41 and Viniflora CH16 from Viniflora oenos and Enoferm alpha. Along PC3, the Viniflora CH16 and oenos strains are positioned toward the left of the scores plot whereas the VP41 and Enoferm alpha strains are positioned more to the right (Fig. 2e). The corresponding loadings plot (Fig. 2f) represents the volatile profiles associated with the respective bacteria. The Viniflora oenos strain is positioned toward the bottom of the scores plot (Fig. 2e) and is associated with higher concentrations of acetoin, acetic acid, ethyl lactate, butanol, isobutanol, propanol, ethyl 3-hydroxybutanoate, ethyl 2-methylbutyrate, ethyl 2-methylpropanoate, hexanoic acid, and isoamyl alcohol. The VP41 strain is separated from Viniflora oenos along PC2 as a result of its association with a selection of esters, higher alcohols, and fatty acids including ethyl octanoate, ethyl hexanoate, ethyl isovalerate, 2-methylpropyl acetate, diethyl succinate hexyl acetate isoamyl acetate, 2-phenylethyl acetate, 2-phenylethanol, 4-methyl-1-pentanol, hexanol, 3-ethoxy-1-propanol, isobutyric acid, decanoic acid, and octanoic acid. Enoferm alpha is positioned toward the right of the scores plot along PC3 (Fig. 2e) and is associated with decanoic acid, 2-phenylethyl acetate, hexyl acetate, isoamyl acetate, and ethyl

acetate. This strain is positioned in the bottom half of the score plot as a result of higher concentrations of ethyl 2-methylbutyrate, ethyl 2-methylpropanoate, ethyl 3-hydroxybutanoate, and ethyl lactate produced during MLF. The CH16 strain was positioned to the left of the scores plot (Fig. 2e) as a result of increased amounts of isovaleric acid, propionic acid, butyric acid, hexanoic acid, octanoic acid, acetic acid, 3-methyl-1-pentanol, hexanol, acetoin, ethyl lactate, ethyl hexanoate, ethyl decanoate, diethyl succinate, and 2-methylpropyl acetate.

General observations in terms of the changes within the different chemical groups including esters, higher alcohols, volatile fatty acids, and carbonyl compounds are discussed in the following sections.

Esters

Changes observed in ester concentrations after the completion of MLF are illustrated in Fig. 3. Synthesis and hydrolysis of esters during MLF were evident, as the results indicate. Ethyl lactate, diethyl succinate, ethyl octanoate, ethyl 2-methylpropanoate, and ethyl propionate concentra-

Table 4 Concentrations of different volatiles after use of different bacterial starter cultures, with odour activity values (OAV) calculated for the Shiraz wines

	Control		Bacteria A		Bacteria V		Bacteria O		Bacteria C	
	Average	OAV	Average	OAV	Average	OAV	Average	OAV	Average	OAV
Esters										
Ethyl decanoate	0.038a	0.2	0.028a	0.1	0.032a	0.2	0.025a	0.1	0.040a	0.2
Ethyl hexanoate	0.402c	28.7	0.454ab	32.4	0.444b	31.7	0.461a	32.9	0.447b	32.0
Ethyl butyrate	0.443b	22.1	0.469a	23.4	0.464ab	23.2	0.478a	23.9	0.464ab	23.2
Ethyl octanoate	0.127d	25.3	0.171ab	34.1	0.163bc	32.5	0.177a	35.3	0.166b	33.1
Ethyl lactate	0.000e	0.0	25.893b	0.2	16.372d	0.1	31.529a	0.2	24.490c	0.2
Ethyl propionate	0.202a	0.1	1.184a	0.7	1.155a	0.6	0.384a	0.2	1.098a	0.6
Ethyl 2-methylpropanoate	0.305b	20.3	0.811a	54.1	0.413ab	27.5	0.430ab	28.7	0.261b	17.4
Ethyl 2-methylbutyrate	0.013a	0.7	0.158a	8.8	0.040a	2.2	0.095a	5.3	0.036a	2.0
Ethyl isovalerate	0.375b	125.2	0.328b	109.4	0.384b	127.9	0.345b	115.1	0.337b	112.4
Ethyl 3-hydroxybutanoate	1.971ab	0.1	1.738b	0.1	1.478b	0.1	1.582b	0.1	1.447b	0.1
Ethyl phenylacetate	2.271bc	3.5	2.235c	3.4	2.444a	3.8	2.410ab	3.7	2.420ab	3.7
Ethyl acetate	43.957a	3.6	46.475a	3.8	43.417a	3.5	46.859a	3.8	46.667a	3.8
Isoamyl acetate	1.486a	49.5	1.476ab	49.2	1.436b	47.9	1.507a	50.2	1.506a	50.2
Hexyl acetate	0.108ab	0.1	0.109ab	0.1	0.124a	0.1	0.094b	0.1	0.099ab	0.1
2-Phenylethyl acetate	0.028a	0.1	0.030a	0.1	0.033a	0.1	0.030a	0.1	0.027a	0.1
Diethyl succinate	0.334b	0.0	0.519a	0.0	0.490a	0.0	0.493a	0.0	0.463a	0.0
2-Methylpropyl acetate	0.654b	0.4	0.515d	0.3	0.584c	0.4	0.548 cd	0.3	0.561 cd	0.4
Alcohols										
Hexanol	1.215c	0.2	1.406a	0.2	1.365b	0.2	1.441a	0.2	1.412a	0.2
Butanol	3.532c	0.0	3.868ab	0.0	3.824ab	0.0	3.926a	0.0	3.873ab	0.0
Methanol	114.136b	0.2	128.928a	0.3	123.490ab	0.2	128.292a	0.3	130.038a	0.3
2-Phenylethanol	36.787b	2.6	40.939a	2.9	39.661ab	2.8	40.906a	2.9	38.761ab	2.8
Propanol	68.792b	0.2	76.253a	0.2	74.049a	0.2	74.242a	0.2	75.362a	0.2
Isobutanol	25.897b	0.6	28.209a	0.7	27.890a	0.7	28.104a	0.7	28.291a	0.7
Isoamyl alcohol	225.917b	7.5	243.408a	8.1	243.053a	8.1	246.364a	8.2	245.353a	8.2
Pentanol	0.166a	0.0	0.139a	0.0	0.172a	0.0	0.173a	0.0	0.167a	0.0
4-Methyl-1-pentanol	0.024ab	0.0	0.000c	0.0	0.034a	0.0	0.006c	0.0	0.000c	0.0
3-Methyl-1-pentanol	0.113b	0.1	0.112b	0.1	0.134a	0.1	0.129ab	0.1	0.137a	0.1
3-Ethoxy-1-propanol	3.004b	30.0	3.296ab	33.0	3.936a	39.4	3.808ab	38.1	3.208ab	32.1
Acids and fatty acids										
Acetic acid	186.742e	0.9	309.976c	1.5	238.173d	1.2	389.174a	1.9	348.797b	1.7
Propionic acid	6.102c	0.3	11.983a	0.6	11.244a	0.6	11.982a	0.6	11.292a	0.6
Isobutyric acid	0.772c	0.3	0.879ab	0.4	0.866ab	0.4	0.878ab	0.4	0.851b	0.4
Butyric acid	0.683b	3.9	0.741a	4.3	0.760a	4.4	0.757a	4.4	0.735a	4.2
Isovaleric acid	1.022c	31.0	1.220ab	37.0	1.195b	36.2	1.291a	39.1	1.212ab	36.7
Valeric acid	0.607a	0.436 cd	0.480bc	0.475bc	0.398d					
Hexanoic acid	0.987a	2.3	0.997a	2.4	0.974a	2.3	1.015a	2.4	0.962a	2.3
Octanoic acid	1.721a	3.4	1.043a	2.1	1.034a	2.1	1.032a	2.1	1.013a	2.0
Decanoic acid	8.189a	8.2	0.492b	0.5	0.491b	0.5	0.423b	0.4	0.422b	0.4
Carbonyl compounds										
Diacetyl (2,3-butanedione)	7.62a	76.2	21.34d	213.4	10.72b	107.2	8.10a	81.0	12.42c	124.2
Acetoin (3-hydroxy-2-butanone)	3.32a	0.0	10.80 cd	0.1	4.53b	0.0	10.08c	0.1	11.21d	0.1
2,3-Pentanedione	2.32b	2.6	1.62c	1.8	1.88d	2.1	1.28a	1.4	1.34a	1.5

Table 4 continued

	Control		Bacteria A		Bacteria V		Bacteria O		Bacteria C	
	Average	OAV	Average	OAV	Average	OAV	Average	OAV	Average	OAV
<i>E</i> -2-Hexenal	0.343b	34.3	0.0005a	0.0	0.0000a	0.0	0.083a	8.3	0.107a	10.7
<i>E</i> -2-Heptenal	nd	nd	nd		nd	nd	nd		nd	
Octanal	nd	nd	nd		nd	nd	nd		nd	
<i>E</i> -2-Octenal	nd a		nd	0.0007a	6.7	0.0003a	2.5	0.0007a	7.0	0.00002a
Nonanal	nd	nd	nd		nd		nd		nd	
<i>E</i> -2-Nonenal	nd a	nd	nd a		0.0004a	5.9	nd		nd a	
Decanal	nd a	nd	nd a		nd a		0.0001a	0.6	0.00008a	
<i>trans</i> -2, <i>cis</i> -6-Nonadienal	0.003a	327.2	0.0041a	410.0	0.0029a	293.1	0.003a	343.4	0.003a	288.9

Averages are expressed as milligrams per litre (mg/l). Different alphabetic letters in the same row indicate significant differences ($P < 0.05$)

tions were increased during MLF in comparison with the control wine, irrespective of the cultivar or bacterial strain evaluated (Fig. 3a, b). For interpretation of the graphs, ethyl lactate is excluded because of its much higher concentration in comparison with the other esters.

In the Pinotage wines, the *Viniflora* CH16 strain produced consistently lower concentrations of ethyl lactate (Table 3) whereas the *Enoferm* alpha strain seems to produce consistently higher concentrations of diethyl succinate, irrespective of the cultivar tested (Tables 3, 4). Ethyl lactate and diethyl succinate are the most important esters typically associated with MLF [30, 39, 56]. The increased concentrations are the result of succinic acid and lactic acid produced during *O. oeni* metabolism followed by subsequent esterification of succinic acid and lactic acid, respectively, with ethanol present in the wine [39]. Although the increase in ethyl lactate concentration was quantitatively the largest, this compound was far below its aroma threshold and is, therefore, not necessarily contributing to wine aroma. Ethyl propionate was consistently produced at higher concentrations by the *Lalvin* VP41 strain and at lower concentrations by the *Viniflora* oenos in the Pinotage and Shiraz wines.

Changes of ethyl hexanoate, ethyl decanoate, and ethyl butyrate concentrations depended on both cultivar and bacterial strain used during MLF, although the *Viniflora* oenos strain tended to produce higher concentrations ethyl butyrate in general. Similarly, ethyl 2-methylbutyrate and ethyl 3-hydroxybutyrate were either increased or reduced with *Enoferm* alpha producing higher levels of ethyl 2-methylbutyrate (fruity, apple) across the wines tested and increased levels of ethyl 3-hydroxybutyrate (strawberry or burnt marshmallow) in the Pinotage 2008 wine. After MLF, ethyl 2-methylbutyrate had an OAV > 1 in all wines, indicative of potential contribution to the resulting fruity wine aroma. Ethyl isovalerate concentrations were reduced in the Shiraz wine whereas a slight increase was observed

for VP41. However, in the Pinotage wines characteristic behaviour was observed, with the *Viniflora* oenos and *Viniflora* CH16 strains increasing the ethyl isovalerate concentrations, *Enoferm* alpha resulting in a decrease, and VP41 not affecting the concentration of this compound during both vintages.

The concentrations of hexyl acetate, isoamyl acetate, 2-phenylethyl acetate, and ethyl acetate were generally decreased or remained unchanged, depending on the strain used or the cultivar tested. Ethyl 2-phenylacetate and 2-methylphenyl acetate concentrations were increased or reduced depending on the cultivar tested, although a general trend for the *Enoferm* alpha strain to produce lower concentrations was observed. Contradictory to previous results [32, 39, 56], the concentration of the powerful odourant isoamyl acetate, characterised by banana attributes, was generally decreased after MLF in all three data sets, with the exception of a strain-specific increase observed in the Pinotage 2009 wine. Isoamyl acetate concentrations found in this study were far above its aroma threshold (0.03 mg/l; [25]) and could, therefore, potentially contribute to the aroma of the wines. 2-Phenylethyl acetate was also decreased or not affected throughout, although the final concentration of this compound was well below its odour threshold (0.25 mg/l; [25]). Pozo-Bayón et al. [51] also reported no differences in hexyl acetate and 2-phenylethyl acetate concentrations as a result of MLF.

Ethyl hexanoate, ethyl butyrate, ethyl octanoate, ethyl 2-methylpropanoate, ethyl 2-methylbutyrate, ethyl isovalerate, ethyl acetate, and isoamyl acetate were all present at OAVs > 1 and could therefore contribute to the fruity aroma of the wines. Total ester production was increased by the bacterial starter cultures and it seems that two of the cultures (bacteria A and V) produced higher ester concentrations than the other two bacteria (bacteria O and C). The total amounts of esters found in these wines after MLF suggest their beneficial contribution to the wines' final aroma.

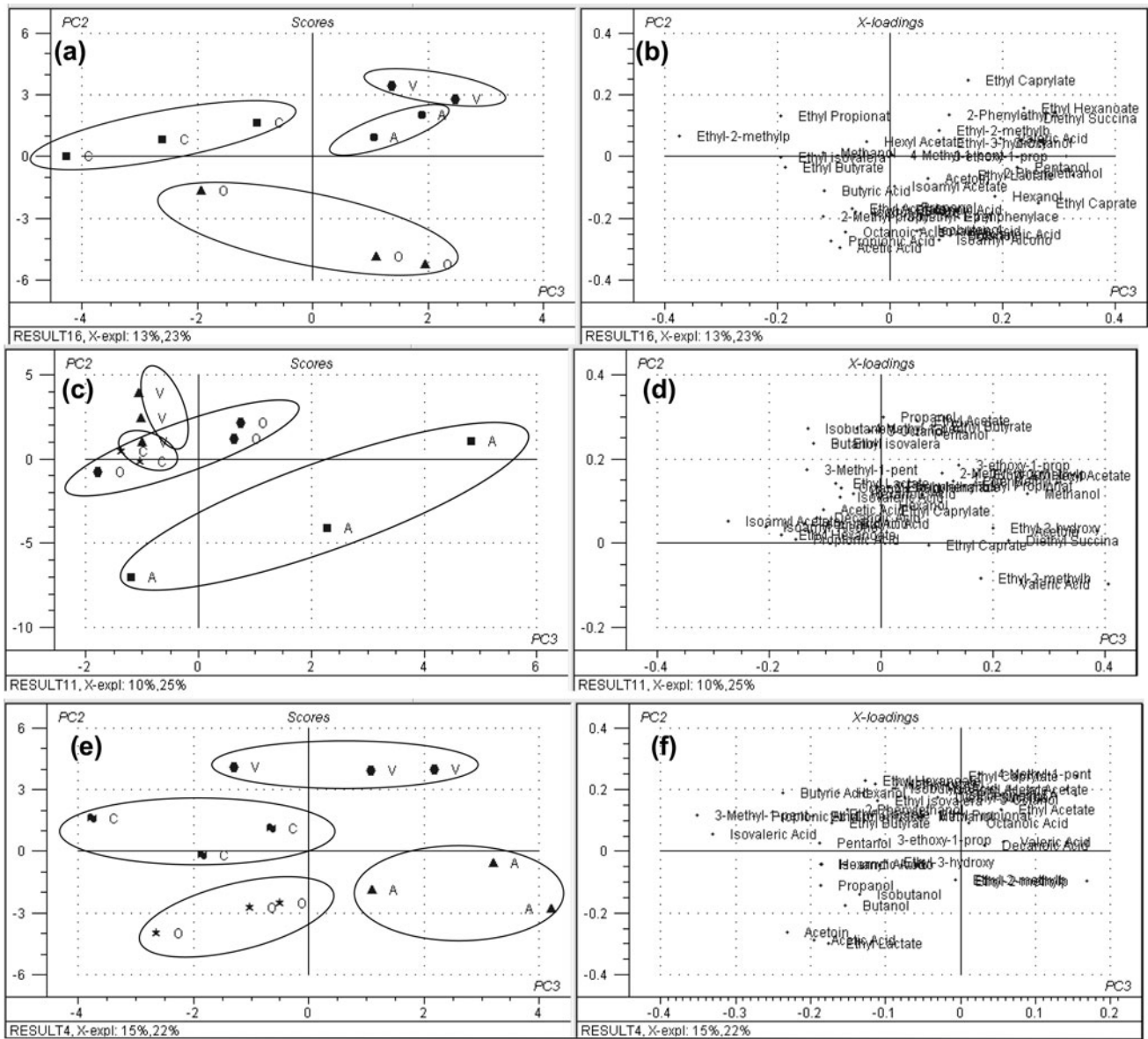


Fig. 2 PCA providing a visual overview of changes in the ester, higher alcohol, and acid composition imparted by bacterial metabolism during MLF. **a** Pinotage 2008 wine scores plot and **b** the corresponding loadings plot; **c** Pinotage 2009 wine scores plot and **d** the corresponding

loadings plot; **e** Shiraz 2008 scores plot and **f** the corresponding loadings plot. The different starter cultures were: Enoferm alpha (indicated by A); Lalvin VP41 (indicated by V); Viniflora oenos (indicated by O), and Viniflora CH16 (indicated by C) are compared

Higher alcohols

Increments in the concentrations of most of the higher alcohols were observed in comparison with control wines in which MLF was suppressed (Tables 3, 4). Higher alcohols are synthesized by yeasts by degradation of amino acids and are considered to affect the aroma and flavour of wine [55]. Isoamyl alcohol, isobutanol, 2-phenylethanol, propanol, butanol, hexanol, 3-methyl-1-pentanol, and 3-ethoxy-1-propanol concentrations were significantly increased by MLF, with characteristic results depending on the strain used to perform MLF. For isoamyl alcohol and isobutanol,

the effect of the bacterial strain selected seems to be more profound in the Pinotage wines than in the Shiraz wine. Maicas et al. [39] have also found production of isobutanol, propanol, butanol, and isoamyl alcohol to be strain dependant. In contrast, other studies found no change in the isoamyl alcohol, 2-phenylethanol, isobutanol, and propanol concentration after MLF [30, 32]. Other authors [13, 31] found that MLF had no significant effect on higher alcohol concentrations in wine, except for significant increases in isoamyl alcohol [13] and isobutanol and 2-phenylethanol, respectively [31]. The observed increase in hexanol and 3-methyl-1-pentanol concentrations as a result of MLF is in

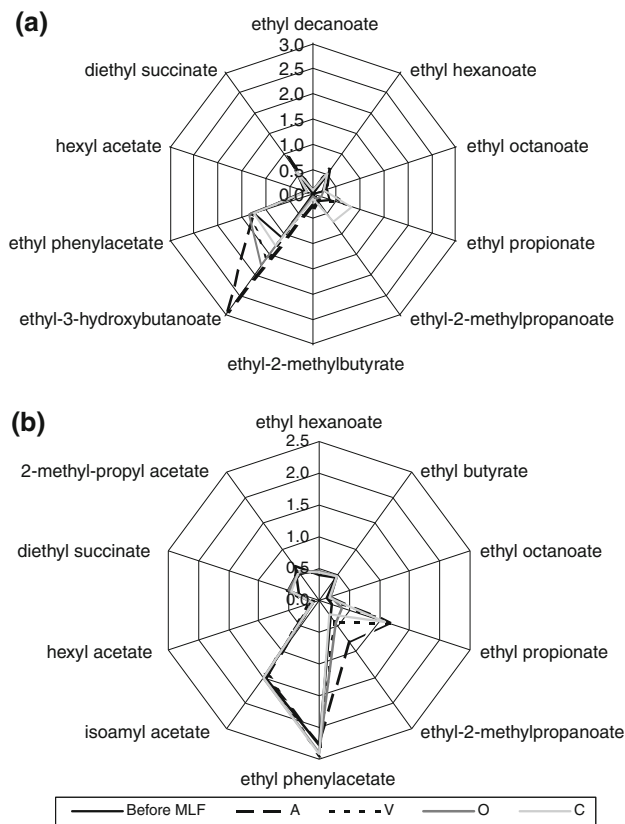
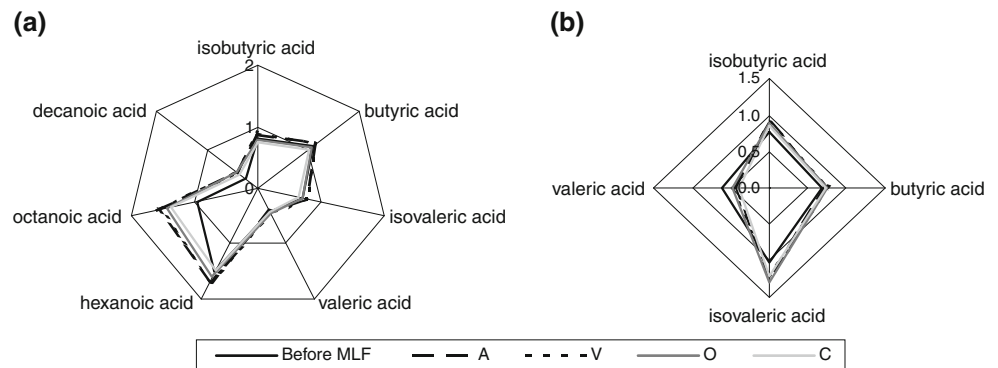


Fig. 3 Changes in the esters profiles associated with MLF by different bacterial starter cultures in **a** Pinotage 2008 and **b** Shiraz 2008 wines

agreement with a previous report by Ugliano and Moio [56]. Pozo-Bayón et al. [51] also found MLF to increase the levels of higher alcohols, however, none of these increases was significant.

In general, the concentrations of these alcohols, depending on the specific alcohol, were in agreement with levels found in young red wines [23]. It is interesting to note that only 2-phenylethanol, isoamyl alcohol, and 3-ethoxy-1-propanol, characterised by honey, spice, rose, lilac, fusel, whisky, malt, and fruity aroma notes, had OAVs > 1 after the completion of MLF. This indicates the potential of higher alcohols to contribute to the complexity and fruity

Fig. 4 Changes in the volatile fatty acid profiles associated with MLF by different bacterial starter cultures in **a** Pinotage 2008 and **b** Shiraz 2008 wines



aromas of wine; at higher concentrations (above 400 mg/l), however, these compounds are detrimental to wine aroma because of their harsh chemical-like aromas [55].

Volatile fatty acids

All four commercial strains of *O. oeni* tested in this study caused significant increases in the concentrations of short-chain fatty acids (Tables 3, 4). Volatile short-chain fatty acids are produced by yeast and bacteria as a result of fatty acid metabolism and, despite their low concentrations in wine, these compounds can negatively affect the aroma quality of wine because of their low perception threshold values and odours reminiscent of cheese and rancid cheese [52]. However, in this study, the extent to which these compounds were affected differed significantly from each other and was strain-dependant for some compounds. Hexanoic, decanoic, and octanoic acids were increased by MLF (Fig. 4), although the magnitudes of the concentration changes were more strain-dependant in the Pinotage wine (Fig. 4a) than in the Shiraz wine (Fig. 4b). In agreement with these results, Maicas et al. [39] found increased levels of decanoic and octanoic acid after MLF. Herjavec et al. [30] found significant increases in octanoic, hexanoic, and decanoic acid concentrations, and Pozo-Bayón et al. [51] found significant differences for octanoic and decanoic acids depending on the MLF culture used.

The other measured fatty acids were either increased or unchanged by MLF (Fig. 4). In a recent metabolic profiling study, Lee et al. [34] reported differentiation between wines according to LAB strain used with regard to, among other compounds, differences observed in isobutyric and octanoic acids. In our study, butyric acid, isovaleric acid, hexanoic acid, and octanoic acid were the only fatty acids present at concentrations above their reported threshold values. This observation is in accordance with previous reports regarding the unlikely contribution of volatile fatty acids to flavour changes or cheesy off-flavours during MLF with *O. oeni* [52, 56]. It has been proposed that wine LAB have the metabolic capacity to produce volatile fatty acids

through lipase activity [10] but lipolytic systems in wine LAB are not well studied and further work on this topic is needed [35, 42].

Carbonyl compounds

Changes related to aldehyde metabolism and citric acid metabolism in terms of the formation of carbonyl compounds in the Pinotage 2008 wine are listed in Table 3 and displayed graphically in Fig. 5a. No significant difference was found between the control and the wines fermented with commercial starter cultures in terms of diacetyl concentration, although the Viniflora CH16 strain produced increased concentrations compared with the other bacteria (Table 3). Similarly, no significant difference was observed in acetoin concentrations, although the Viniflora CH16 strain produced the highest concentration. PCA provides a summary of the changes observed (Fig. 5a). Diacetyl and acetoin were correlated with each other and strongly associated with the Viniflora oenos bacteria and, to a lesser extent, with the Viniflora CH16 and VP41 strains. Enoferm alpha was positioned toward the bottom of the plot and was associated with increased concentrations of *E*-2-octenal (herbaceous, lemon), decanal (citrus, fruity), nonanal (herbal, floral), and *E*-2-hexenal (herbaceous, green), all of which are associated with green or herbaceous aromas in wine [12]. However, none of these compounds was present at concentrations above their individual aroma thresholds, but possible contribution to wine aroma should not be excluded, because their cumulative effect might contribute to perceived wine aroma. The presence of *trans*-2,*cis*-6-nonadienal and 2,3-pentanedione was associated more with the control and spontaneous treatments during this experiment. It is interesting to note the negative correlation between diacetyl and 2,3-pentanedione. The chemical oxidation of diacetyl results in the formation of 2,3-pentanedione whereas reduction of diacetyl by yeasts and LAB results in the formation of acetoin. Diacetyl was present at concentrations above its range of reported aroma threshold values (0.2–2.3 mg/l; [41]), as was 2,3-pentanedione, whereas acetoin was present at levels below its reported threshold and would be less likely to contribute to wine aroma. All three compounds contribute very similar aroma attributes to wine, therefore the cumulative effect on wine aroma should not be excluded.

During MLF of the Pinotage 2009 vintage, separation along the first PC was driven by the strong association of the Enoferm alpha (indicated by A in Fig. 5c and Table 3) strain with diacetyl and acetoin positioned toward the far left of the plot. Viniflora oenos and Viniflora CH16 also produced significant levels of diacetyl and acetoin whereas the VP41 strain (indicated by V in the graph) produced slightly lower levels of these compounds. The control treat-

ment was strongly associated with higher concentrations of the aldehydes and was positioned toward the far right of the plot, and the VP41 strain was also associated with these compounds.

Prominent discrimination, by use of PCA, among the different MLF treatments in terms of carbonyl compounds for the Shiraz 2008 experiment is shown in Fig. 5e. The control, spontaneous, and VP41 treatments were positioned to the left of the PCA scores plot and correlated with each other (Fig. 5e) as a result of their association with higher concentrations of 2,3-pentanedione, *E*-2-hexenal, and *E*-2-nonanal (Fig. 5f), and with lower concentrations of diacetyl and acetoin. Conversely, the Enoferm alpha, Viniflora oenos, and Viniflora CH16 strains were positioned to the right of the scores plot and showed strong association with diacetyl and acetoin (Fig. 5f). The Enoferm alpha strain is slightly separated toward the top right of the scores plot (Fig. 5e) as a result of its association with *trans*-2,*cis*-6-nonadienal and *E*-2-octenal. The Viniflora oenos and CH16 strains are more correlated and positioned slightly toward the bottom of the scores plot, because of the presence of decanal. The Enoferm alpha strain produced significantly more diacetyl than the other three bacteria (Table 3) whereas VP41 imparted the smallest increase in diacetyl concentrations. All four strains produced diacetyl concentrations with OAVs > 1, potentially contributing to the buttery aroma of the wine [3]. Acetoin was produced at higher concentrations by Enoferm alpha and Viniflora CH16 strains than by VP41; these concentrations did not exceed the reported threshold level of 150 mg/l, however [18].

In summary, MLF increased diacetyl concentrations significantly with the exception of the Pinotage 2008 wine. Strain-dependant differences were observed with the VP41 strain generally producing lower concentrations of diacetyl, irrespective of the cultivar tested. Acetoin concentrations were always increased.

Significant differences in organic acid profiles corresponding to different bacterial starter cultures were shown to be a useful means of depicting possible differences in terms of specific metabolites such as ethyl lactate and diacetyl. As previously reported [35], differences in lactic acid production and citric acid metabolism could indicate different metabolic requirements and resulting volatile metabolites.

In terms of volatile composition, general increases in the ester, higher alcohol, and volatile fatty acid concentrations of all the wines were observed after the completion of MLF, irrespective of the bacterial strain and grape cultivar used. However, specific strain-dependent differences were observed for some compounds. A large portion of the esters, which are important for the fruity aroma notes of wine, were found to have OAVs > 1, indicating their

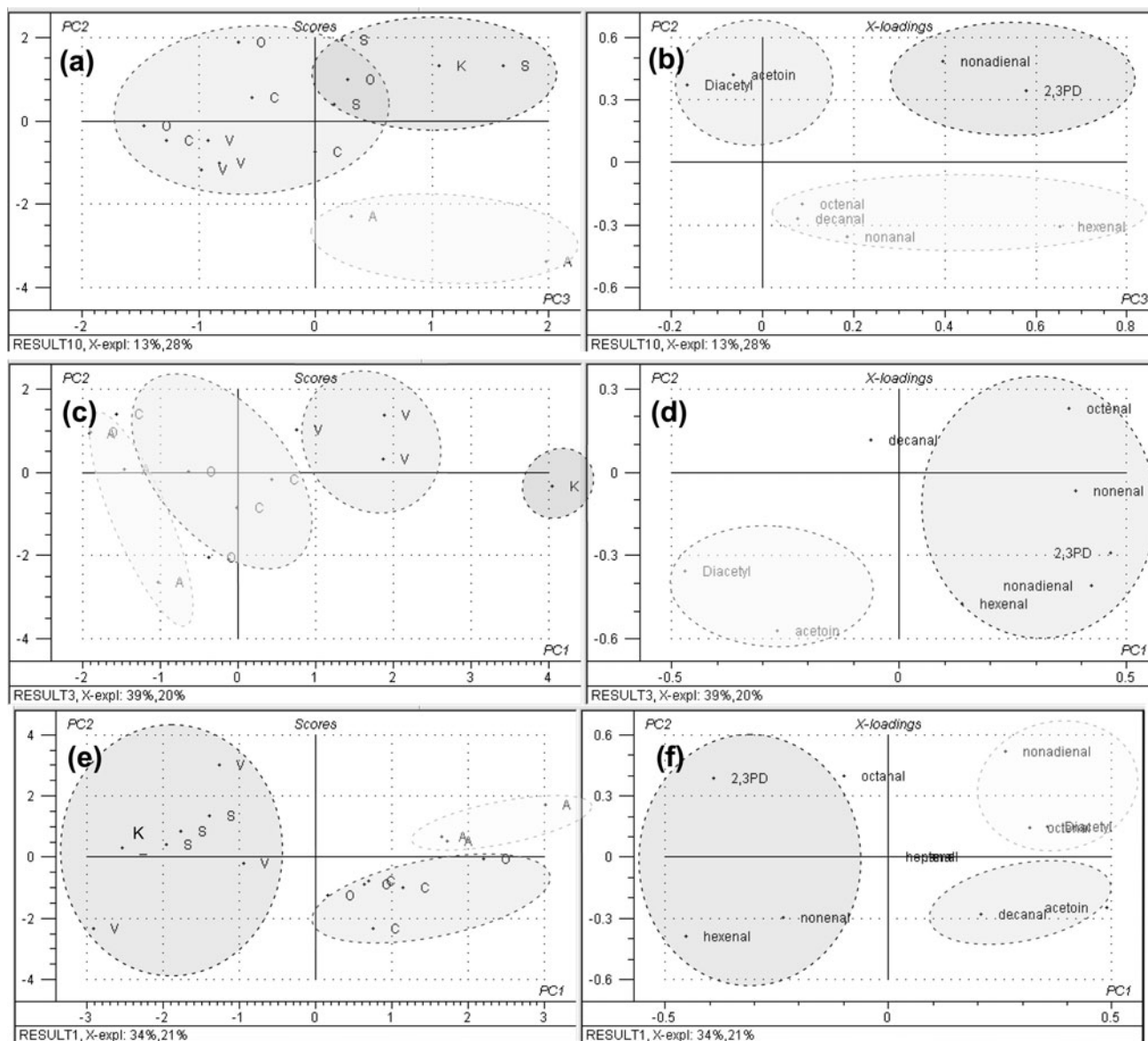


Fig. 5 PCA illustrating differences among the four strains (indicated by A, V, O, C) in terms of their effect on a selection of carbonyl compounds in the Pinotage 2008 wine (a, b), Pinotage 2009 wine (c, d), and Shiraz 2008 wine (e, f). Bacteria are indicated by A (Enoferm al-

pha), V (Lalvin VP41), O (Viniflora oenos), and C (Viniflora CH16), and the control wine is indicated by the letter K. The letter S represents the spontaneous MLF treatment included in the study

potential aroma contribution. It could therefore be concluded that MLF using any of these four strains may contribute to wine quality by modifying the concentrations of some of the aroma compounds. However, the effect of chemical changes on perceived aroma should further be investigated by use of sensory evaluation techniques.

Comparative studies of the effect of different commercial MLF bacteria on the concentration of wine volatiles often focus on selected groups of compounds whereas the cultivars and strains tested are often very specific to countries and regions. The effect of MLF activity in Tannat, the most important red wine in Uruguayan viticulture, has been investigated [6], and focussed on comparison of different

major volatile compounds. In other studies, the potential of four commercial MLF starter cultures to hydrolyse glycosides and release volatile compounds, and the effect on yeast-derived volatile compounds during MLF in Aglianico grapes from Southern Italy, were evaluated [56, 57]. Pozo-Bayón et al. [51] found significant differences in the wine volatile and amino acid composition of Tempranillo wine, one of the most important Spanish red grape cultivars, after MLF with *O. oeni* and *Lactobacillus plantarum* starter cultures. Recently, metabolic profiling studies revealed significant differences among major volatile compounds after MLF by use of different starter cultures in Meoru wine, made from a wild Korean grape [34, 54]. Some studies

have reported the sensory effects of MLF for Chardonnay [52] and Pinot noir [53] by comparing the effect of different bacterial inoculations. However, these studies lack supporting volatile composition data. Volatile aroma constituents including esters, aldehydes, alcohols, ketones, acids, and sulfur-containing compounds in Chancellor wine after MLF have been investigated by use of two commercial starter cultures [11]. De Revel et al. [13] found increased concentrations of wood-derived volatile compounds after MLF performed in barrels with Sauvignon blanc must. Studies specifically focussing on the effect of different MLF conditions on diacetyl concentrations in Chardonnay have also been reported [32].

In conclusion, this research contributes to our current knowledge of malolactic fermentation and, more specifically, the potential contribution to wine composition and aroma. The results therefore illustrate and reiterate that MLF and, specifically, the use of different starter cultures, affects wine aroma and flavour. However, the contribution of these starter cultures was different depending on seasonal variation and precursors present in the wine as a result of the cultivar used. Future studies should include recently developed starter cultures, combination starter cultures, and additional inoculation strategies in order to further expand our current knowledge on MLF and wine aroma under different conditions.

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