

## Application Potential of Baijiu Non-Saccharomyces Yeast in Winemaking Through Sequential Fermentation With Saccharomyces cerevisiae

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Li R-R, Xu M, Zheng J, Liu Y-J, Sun C-H, Wang H, Guo X-W, Xiao D-G, Wu X-L and Chen Y-F (2022) Application Potential of Baijiu Non-Saccharomyces Yeast in Winemaking Through Sequential Fermentation With Saccharomyces cerevisiae. Front. Microbiol. 13:902597. doi: 10.3389/fmicb.2022.902597 To explore the potential application of non-Saccharomyces yeasts screened from Baijiu fermentation environment in winemaking, the effect of four Baijiu non-Saccharomyces veasts (two Zvgosaccharomyces bailii and two Pichia kudriavzevii) sequentially fermented with Saccharomyces cerevisiae on the physicochemical parameters and volatile compounds of wine was analyzed. The results indicated that there was no obvious antagonism between S. cerevisiae and Z. bailli or P. kudriavzevii in sequential fermentations, and all strains could be detected at the end of alcoholic fermentation. Compare with S. cerevisiae pure fermentation, Z. bailii/S. cerevisiae sequential fermentations significantly reduced higher alcohols, fatty acids, and ethyl esters and increased acetate esters; P. kudriavzevii/S. cerevisiae sequential fermentations reduced the contents of C6 alcohols, total higher alcohols, fatty acids, and ethyl esters and significantly increased the contents of acetate esters (especially ethyl acetate and 3-methylbutyl acetate). Sequential fermentation of Baijiu non-Saccharomyces yeast and S. cerevisiae improved the flavor and quality of wine due to the higher ester content and lower concentration of higher alcohols and fatty acids, non-Saccharomyces yeasts selected from Baijiu fermentation environment have potential applications in winemaking, which could provide a new strategy to improve wine flavor and quality.

Keywords: S. cerevisiae, Zygosaccharomyces bailii, Pichia kudriavzevii, sequential fermentation, volatile compounds

## INTRODUCTION

Studies have reported that some non-*Saccharomyces* yeasts can improve the organoleptic quality and sensory notes of wine, depending on the specific yeast species and strains used (Padilla et al., 2016; Benito et al., 2019). For example, most species from the *Hanseniospora* genus can improve the color and polyphenolic composition in red wines (Leixà et al., 2016), *Torulaspora delbrueckii* and *Saccharomyces cerevisiae* sequential inoculation can increase total esters concentration such as 3-methylbutyl acetate, ethyl propanoate, and ethyl 2-methylpropanoate (Renault et al., 2015), *Pichia anomala* and *S. cerevisiae* mixed fermentation can increase the content of 3-methylbutyl acetate and

ethyl esters (Kurita, 2008). However non-*Saccharomyces* yeasts selected from vineyards (including the grapes) and wineries (including the winery equipment) have a low capacity to metabolize sugar to ethanol, and low resistance to sulfur dioxide and ethanol in most cases, they have often been inhibited by *S. cerevisiae* inoculations at the industrial level (Benito, 2018). Therefore, it is necessary to look for excellent non-*Saccharomyces* yeasts with high ability to ferment sugar, high tolerance to various stresses, and can produce high yield aroma compounds to be used in winemaking.

Baijiu, a traditional fermented alcoholic beverage in China, is rich in many flavor components, including esters, terpenes, organic acids, lactones, phenols, heterocycles, and aromatic compounds (Fan and Qian, 2006). Baijiu fermentation is generally under an open or semi-open fermentation environment (Xu et al., 2022), a variety of microorganisms from the Daqu (the fermentation starter), water, air, tools, and operators participate in the fermentation process (Xu et al., 2017a). Among these fermentation microbial communities, non-Saccharomyces yeasts can produce aldehydes, esters, higher alcohols, and other flavor substances during Baijiu fermentation, giving Baijiu its typical aroma characteristics. For example, Zygosaccharomyces bailii is a type of yeast with high tolerance to various stresses (Stratford et al., 2013; Palma et al., 2015). It was found to be a dominant species in Maotai-flavor liquor fermentation and the contributors to ethanol and various flavor compounds in Baijiu making (Xu et al., 2017b). In addition, it was found that the Zygosaccharomyces strains isolated from grape musts were described as strains with low higher alcohol production (Romano and Suzzi, 1993). In wine fermentation, the high ester-producing ability of Z. bailii was used to increase the content of ethyl ester in wine by mixed fermentation with S. cerevisiae (Ciani et al., 2010; Garavaglia et al., 2015). Pichia kudriavzevii is the dominant species of the genus Pichia in Baijiu fermentation (Jiang et al., 2019), which can generate esters, higher alcohols, and volatile acids (Liu et al., 2017). In addition, P. kudriavzevii can produce higher glycerol, ethyl acetate, and 3-methylbutyl acetate in mixed fermentation of wine but lower contents of fatty acids, higher alcohols, and phenylethyl alcohol (Luan et al., 2018). Shi et al. (2019) found that the wine fermented by P. kudriavzevii and S. cerevisiae had lower concentrations of volatile acids, higher alcohols, fatty acids, benzene derivatives, and C6 compounds, and higher concentrations of esters, which improved the aroma and overall flavor characteristics of the wine.

The production process of Baijiu is accompanied by special extreme environments, such as high ethanol, high temperature, and high acidity (Xu et al., 2017a; Wang et al., 2019). After long-term domestication in an extreme environment, Baijiu non-*Saccharomyces* yeast strains may have stronger ability to adapt to the winemaking environment. The non-*Saccharomyces* yeast strains selected from Baijiu may not be easily inhibited by *S. cerevisiae* in wine fermentation, which is more beneficial to improving the quality of wine. Therefore, the co-fermentation of non-*Saccharomyces* yeast strains selected from Baijiu fermentation environments and *S. cerevisiae* may be a new strategy to improve wine flavor and quality. To the best of our knowledge, one non-*Saccharomyces* yeast strain (*Pichia*)

*fermentans*) selected from Baijiu fermentation environments has been applied to wine fermentation, and it showed a positive contribution to wine aroma (Ma et al., 2017; Li et al., 2020).

In the previous work at our laboratory, several non-Saccharomyces yeasts were isolated from the fermented grains of a sauce-flavor Baijiu in China. Previous studies demonstrated that they have good fermentation characteristics and show high ester-producing ability (unpublished results). To explore the application potential of Baijiu non-Saccharomyces yeast strains in wine fermentation, in this study, four non-Saccharomyces yeast strains (two Z. bailii and two P. kudriavzevii) selected from the fermented grains of a sauce-flavor Baijiu in China were inoculated sequentially with S. cerevisiae EC1118, and their effects on the physicochemical parameters and volatile compositions of wine were evaluated.

#### MATERIALS AND METHODS

#### **Yeast Strains and Preculture Conditions**

Four indigenous non-*Saccharomyces* yeast strains were used in this study, including *Z. bailii* (BJII45005 and BJVI11007) and *P. kudriavzevii* (BJIV53006 and BJII44006). The four strains were isolated from the fermented grains of a sauceflavor Baijiu in China. They were identified by 26S rDNA analysis (**Supplementary Table 1**) and kept in our lab. The commercial wine strain *S. cerevisiae* Lalvin EC1118 (Lallemand Inc., Montreal, QC, Canada) was used as the control. These strains were stored at  $-80^{\circ}$ C in YPD medium with glycerol (20% v/v final concentration).

Starter cultures of all yeast strains were prepared by inoculating a single colony in 5 mL of YPD medium broth for each strain. The cultures were incubated in a test tube rotating overnight (30°C and 180 rpm). These starter cultures were transferred to 500-mL shake flasks containing 300 mL of YPD medium for 18 h (30°C and 180 rpm). The commercial strain EC1118 was prepared under the same conditions as non-*Saccharomyces* yeast. The cultured yeast strains were counted by the blood cell counting method, and each sample was counted in triplicate. The yeast cells were centrifuged and washed twice with sterile water. The inoculum ratio of non-*Saccharomyces* yeast and *S. cerevisiae* was 10:1, and the initial active population of non-*Saccharomyces* yeast and *S. cerevisiae* was  $1.0 \times 10^7$  cells/mL and  $1.0 \times 10^6$  cells/mL (Zhang et al., 2018), respectively.

#### **Fermentation Conditions and Sampling**

*Cabernet Sauvignon* grapes (227.54 g/L of sugar, pH 3.38, 259.81 mg/L yeast assimilable nitrogen) were harvested from the Qinhuangdao region vineyard (Hebei, China) in the 2020 vintage (October 3rd). The grapes were in good sanitary conditions and came from the same vineyard. The grapes were harvested by hand and immediately transported to the laboratory in the same box. After stemming and crushing, 400 g of grape musts were added into 500-mL Erlenmeyer flasks and pasteurized at 68°C for 30 min. The grape musts were macerated at 4°C for 12 h after 60 mg/L total SO<sub>2</sub> was added. When the temperature returned to 25°C, yeast was added. Flasks were sealed with hydrophobic





membranes to create anaerobic conditions (carbon dioxide was released through an air outlet membrane).

Mixed fermentation trials were performed with four indigenous non-*Saccharomyces* yeasts and EC1118 sequential inoculation. Nine trials were therefore set: (1) single inoculation with EC1118 (SC); (2) sequential inoculation with BJVI11007, followed by inoculation with EC1118 after 2 days (ZBI-2); (3) sequential inoculation with BJVI11007, followed by inoculation with EC1118 after 3 days (ZBI-3); (4) sequential inoculation with BJII45005, followed by inoculation with EC1118 after 2 days

(ZBII-2); (5) sequential inoculation with BJII45005, followed by inoculation with EC1118 after 3 days (ZBII-3); (6) sequential inoculation with BJIV53006, followed by inoculation with EC1118 after 2 days (PKI-2); (7) sequential inoculation with BJIV53006, followed by inoculation with EC1118 after 3 days (PKI-3); (8) sequential inoculation with BJII44006, followed by inoculation with EC1118 after 2 days (PKII-2); and (9) sequential inoculation with BJII44006, followed by inoculation with EC1118 after 3 days (PKII-3). Fermentations were carried out in triplicate for each treatment at a controlled temperature of 25°C and included punching skins down twice a day to improve extraction. A total of 2 mL of fermenting musts were sampled daily for counting the yeast population and for HPLC analysis.

### The Fermentation Process and Yeast Enumeration

The fermentation process was monitored twice per day by measuring the weight loss of the bottles due to the  $CO_2$  release. The fermentations were stopped when the weight loss was less than 0.1 g in 12 h. After alcoholic fermentation, grape pomace was separated from wine carefully, centrifuged, and stored at  $-20^{\circ}C$  for further analysis.

The viable cell count was performed by identifying colony colors using 100 mg/L of chloramphenicol (Solarbio, Beijing, China) that was added to Wallerstein Laboratory (WLN) nutrient agar (Qingdao Hope Bio-Technology Co., Ltd., China). One hundred microliter aliquots were plated onto WLN plates. After 48 h of incubation at 30°C, the cells could be differentially counted based on the morphological particularities presented by the non-*Saccharomyces* yeasts that distinguished them from *S. cerevisiae*.

#### **Analysis of Physicochemical Parameters**

Ethanol, glucose, fructose, glycerol, citric acid, tartaric acid, malic acid, succinic acid, lactic acid, and acetic acid were determined by HPLC (Agilent 1260 Infinity) system consisting of a quaternary gradient pump (1260 Iso Pump-G1310B), an autosampler (1260 ALS-C1328B) and a refractive index detector (1260 TCC-G1316A). Separations were performed on a Silgreen Ca/H column operating at 65°C, the mobile phase was 5 mM sulfuric acid at a flow rate of 0.6 mL min<sup>-1</sup>, and the running time of the program was 23 min. The samples were diluted and filtered (0.22 µm RC syringe filters, Tianjin Jinteng Experimental Equipment Equipment Co., Ltd., Tianjin, China) and 20 µL were injected. The concentration of each metabolite was calculated with an external standard method using peak areas. The detailed quantitation information about quantitative standards, calibration curves, and  $R^2$  for the quantification compounds used in this study was provided in Supplementary Table 2. The pH of the wine samples was measured using a pHSJ-4A model pH meter (Shanghai Scientific Instruments and Materials Co., Ltd.).

## **Analysis of Volatile Compounds**

The volatile compounds of wines after alcohol fermentation were quantified using headspace solid-phase microextraction coupled with gas chromatography-mass spectrometry (HS-SPME-GC-MS) according to our previous study (Li and Sun, 2019). Identification was based on ion fragment and mass spectra matching against the standard NIST 14 library. 4-Methyl-2-pentanol was used as the internal standard (20  $\mu$ L of 2.00 g/L sample solution). Subsequently, all standard stock solutions were combined, and this mixed standard solution was diluted into several levels in succession with a 12% (v/v) alcohol solution. Under the same conditions, the standards at all levels were extracted and analyzed. For quantification, calibration curves were obtained with regression coefficients above 98%

(**Supplementary Table 3**). In addition, the concentration of volatile compounds without pure standard was estimated based on the calibration curves of standard compounds with the most similar chemical structures and/or numbers of carbon atoms (Cai et al., 2014).

## **Sensory Evaluation**

The sensory evaluation was performed as described by Zhang et al. (2022) with modification. Twenty milliliters of wine samples were poured into wine glasses and presented in random order. Wine sensory evaluation was classified into nine attributes, including aromatic intensity, floral, fruity, sweet, herbaceous, fatty, solvent, acidity, and wine body. The descriptive sensory analysis was carried out by a well-trained sensory panel comprised of 12 assessors from the College of Biotechnology, Tianjin University of Science and Technology (six females and six males, ranging in ages from 22 to 36, an average of 25). Panelists were required to rate the intensity of the wine parameters using a 9-point scale (1 = extremely low, 5 = moderate intensity, 9 = extremely high). The final score of each sensory characteristic was the mean value of scores given by 12 assessors.

## **Statistical Analysis**

One way analysis of variance (ANOVA) using the Duncan test at the significance level of P < 0.05 was performed using SPSS 17.0 (Chicago, IL, United States), Principal component analysis (PCA) was conducted using the SIMCA-13.0 software (Umetrics, Sweden); Heat map was conducted using the TBtools software (Chen et al., 2020); Upset plot was conducted using the R software (version 3.3.2) in UpSetR package; others were conducted using the OriginPro81 (OriginLab, United States).

## RESULTS

# The Fermentation Process and Yeast Growth During Fermentation

The fermentation process (represented by  $CO_2$  release) and yeast growth kinetics in pure fermentation and sequential fermentations are shown in **Figure 1**. Alcohol fermentation lasted for 8 days, and no stuck or sluggish fermentations were found in any of the inoculation strategies. *S. cerevisiae* EC1118 single inoculation finished alcohol fermentation faster, showing that sequential inoculations slowed down the end of the process. Compared with *S. cerevisiae* pure fermentation, sequential fermentation delayed the time of alcohol fermentation (**Figure 1A**). In sequential fermentations, the inoculation time of *S. cerevisiae* had a great effect on the fermentation alcohol process, and the delayed inoculation for 2 days was faster than that for 3 days. Similarly, SC consumed glucose and fructose faster than sequential fermentations (**Supplementary Figures 1A,B**).

The influence of yeast strains on the final flavor characteristics of the wine is largely determined by the yeast cell number during the fermentation process (Zhang et al., 2018). In *S. cerevisiae* pure fermentation (**Figure 1B**), EC1118 grew rapidly

Compounds	SC	ZBI-2	ZBI-3	ZBII-2	ZBII-3	PKI-2	PKI-3	PKII-2	PKII-3
Ethanol	$92.14 \pm 5.52a$	$90.41 \pm 2.00a$	88.82 ± 2.33a	$89.83\pm2.03a$	90.13 ± 1.51a	$92.95 \pm 2.07a$	$91.25 \pm 2.72a$	91.86 ± 1.77a	$91.77 \pm 3.01a$
Glycerol	$11.14\pm0.09cd$	$11.58\pm0.04\text{ab}$	$11.74 \pm 0.03a$	$11.66\pm0.59\text{ab}$	$10.89\pm0.16d$	$11.22\pm0.23bcd$	$11.14\pm0.04cd$	$11.05\pm0.19\text{cd}$	$11.40 \pm 0.19$ abc
Glucose	$0.21\pm0.04 \text{f}$	$0.87\pm0.04b$	$0.99\pm0.07a$	$0.51\pm0.04$ d	$0.22\pm0.07\text{f}$	$0.36\pm0.08e$	$0.80\pm0.02b$	$0.68\pm0.17c$	$0.33\pm0.15\mathrm{e}$
Fructose	$0.27\pm0.06\mathrm{c}$	$0.51 \pm 0.10 b$	$0.33\pm0.17\text{bc}$	$1.05 \pm 0.13a$	$1.09 \pm 0.17a$	$1.17 \pm 0.32a$	$1.05 \pm 0.18a$	$0.30\pm0.07c$	$1.16 \pm 0.13a$
рН	$3.63\pm0.05\text{bc}$	$3.72\pm0.08ab$	$3.56\pm0.02c$	$3.68\pm0.06b$	$3.65\pm0.01\mathrm{b}$	$3.77 \pm 0.06a$	$3.71\pm0.03 \text{ab}$	$3.65\pm0.04b$	$3.65 \pm 0.02b$
Citric acid	$1.61\pm0.01c$	$1.10\pm0.01e$	$1.29\pm0.09d$	$1.19\pm0.06\text{de}$	$0.76\pm0.02 \text{f}$	$1.80\pm0.07b$	$1.95 \pm 0.12a$	$1.27\pm0.08d$	$1.08\pm0.04e$
Tartaric acid	$2.13\pm0.08cd$	$2.08\pm0.08d$	$2.14\pm0.05cd$	$2.21\pm0.09 \text{bc}$	$2.02\pm0.01d$	$2.35 \pm 0.09a$	$2.39\pm0.08a$	$2.27\pm0.03 \text{ab}$	$2.35 \pm 0.06a$
Malic acid	$1.88\pm0.08\text{f}$	$1.93\pm0.04\text{ef}$	$3.16\pm0.10a$	$2.12\pm0.08d$	$2.77 \pm 0.12b$	$1.98\pm0.05\text{ef}$	$2.32\pm0.10c$	$2.04\pm0.03\text{de}$	$2.14 \pm 0.01 d$
Succinic acid	$2.02\pm0.03d$	$3.62\pm0.08b$	$3.88\pm0.10a$	$3.51 \pm 0.14b$	$3.36\pm0.04c$	$1.30\pm0.01e$	$1.35\pm0.03e$	$1.38\pm0.03e$	$1.37\pm0.04e$
Lactic acid	$0.42 \pm 0.01 a$	$0.18\pm0.03e$	$0.14\pm0.01 f$	$0.25\pm0.02b$	$0.17\pm0.01e$	$0.21\pm0.01$ cd	$0.23\pm0.01\text{bc}$	$0.16\pm0.01\text{ef}$	$0.19\pm0.01$ de
Acetic acid	$0.40\pm0.03c$	$0.61\pm0.11b$	$0.73\pm0.07a$	$0.30\pm0.05d$	$0.26\pm0.02d$	$0.26\pm0.02d$	$0.26\pm0.01d$	$0.24\pm0.02d$	$0.25\pm0.01d$

SC, single inoculation with S. cerevisiae EC1118; ZBI-2, sequential inoculation with Z. bailii BJVI11007, followed by inoculation with EC1118 after 2 days; ZBI-3, sequential inoculation with Z. bailii BJVI11007, followed by inoculation with EC1118 after 3 days; ZBI-2, sequential inoculation with Z. bailii BJII45005, followed by inoculation with EC1118 after 2 days; ZBI-3, sequential inoculation with Z. bailii BJII45005, followed by inoculation with EC1118 after 2 days; ZBI-3, sequential inoculation with Z. bailii BJII45005, followed by inoculation with P. kudriavzevii BJIV53006, followed by inoculation with P. kudriavzevii BJIV53006, followed by inoculation with P. kudriavzevii BJIV53006, followed by inoculation with P. kudriavzevii BJIV4006, followed by inoculation with P. kudriavzevii BJIV43006, followed by inoculation with P. kudriavzevii BJIV44066, followed by inoculation with CI118 after 3 days. Values are given as mean  $\pm$  standard deviation of three replicates. Data with different letters (a, b, c, d, e, f, g, h) within each row are different according to Duncan tests ( $p \le 0.05$ ).

on the first day of inoculation before reaching a maximum  $(4.07 \times 10^8 \text{ CFU/mL})$  and then gradually decreased until the end of alcohol fermentation. In sequential fermentations (Figures 1C-F), two non-Saccharomyces yeasts of the same species showed a similar growth curve. In the sequential fermentation of Z. bailli and S. cerevisiae (ZB) (Figures 1C,D), the population of Z. bailli increased slightly on the first day of inoculation and then decreased gradually in the subsequent fermentation process. The number of EC1118 increased rapidly after inoculation, and the population of EC1118 was higher than that of Z. bailli in the subsequent fermentation process. In the sequential fermentation of P. kudriavzevii and S. cerevisiae (PK) (Figures 1E,F), P. kudriavzevii grew rapidly on the first day and then slightly decreased, and the quantity was maintained in the first 6 days. Compared with the inoculation of S. cerevisiae on the second day, the number of S. cerevisiae inoculated on the third day was lower than that of P. kudriavzevii. In summary, sequential fermentation delayed the time of alcohol fermentation, and no obvious antagonism between S. cerevisiae and Z. bailli or P. kudriavzevii, and all strains could be detected at the end of alcohol fermentation.

#### **Physicochemical Parameters**

The physicochemical parameters of wines produced by pure and sequential fermentations are shown in **Table 1**. At the end of alcohol fermentation, all treatments were completely fermented (residual sugars were less than 2 g/L). There was no significant difference in the content of ethanol, and no regular difference in the pH value among the samples. In this study, the acetic acid content in all wine samples was below 0.80 g/L, and PK could significantly decrease the acetic acid content. In terms of non-volatile acid, the citric acid content in ZB was significantly lower than that in SC, and the effects of *P. kudriavzevii* were slightly different between the two strains. PK could significantly increase the content of tartaric acid. Compared with SC, sequential fermentations could significantly increase the content of malic acid and reduce the content of lactic acid. It was worth noting that ZB could significantly increase the content of succinic acid.

## **Volatile Compositions**

The volatile compositions of the wines produced by different inoculation strategies were determined. Forty-six volatile compositions were identified in all samples, including 17 alcohols (three C6 alcohols and 14 higher alcohols), three fatty acids, 22 esters (four acetate esters, 13 ethyl esters, and five other esters), and four other compounds (Table 2). Compared with SC, PK increased the total content of volatile compositions, especially BJII44006. ZB could significantly reduce the total content of volatile compositions. The kinds of volatile compounds affect the complexity of the wine aroma, the Upset plot (Figure 2) was used to visualize the difference in the kinds of volatile compounds among different inoculation strategies. Among them, 28 kinds of volatile compositions were detected in all samples (Figure 2); SC had the most kinds (44 kinds) of volatile compositions, followed by ZBI-2 (43 kinds), and PKI-3 (31 kinds) had the least kinds of volatile compositions. In general, compared with SC, sequential fermentations could reduce the kinds of volatile compositions; the inoculation of S. cerevisiae, which was delayed by 3 days, reconfirmed this result. The odor activity value (OAV) was calculated as the ratio between the concentration of each volatile compound and its perception threshold. The compound that OAV over one has a high contribution to wine aroma (Guth, 1997). In recent years, some researchers (Escudero et al., 2007; Ryan et al., 2008) found that compounds with relatively low OAVs can have an unexpectedly high effect on the aroma. So, the ratios of volatile compound contents (OAV > 0.1)after alcoholic fermentation were calculated and are shown in Supplementary Table 4.

#### C6 Alcohols and Higher Alcohols

Three C6 alcohols (1-hexanol, E-3-hexen-1-ol, Z-3-hexen-1-ol) were identified (**Table 2**), and their OAVs > 0.1 (**Supplementary Table 4**). In general, ZB could increase the total content of C6

Aroma compounds	SC	ZBI-2	ZBI-3	ZBII-2	ZBII-3	PKI-2	PKI-3	PKII-2	PKII-3
1-Hexanol	$629.71 \pm 19.85 ab$	$611.50 \pm 46.49$ ab	738.30 ± 34.61a	735.14 ± 129.86a	738.55 ± 95.45a	$407.66 \pm 23.63 d$	$574.76 \pm 56.58 \text{bc}$	$604.32 \pm 75.30b$	$470.00 \pm 55.43$ cd
(E)-3-Hexen-1-ol	$182.08 \pm 25.13b$	$193.76 \pm 2.65 ab$	$195.88 \pm 2.07 ab$	$205.66 \pm 2.82a$	$202.13 \pm 5.77a$	$200.28 \pm 3.09a$	$194.64 \pm 3.61 ab$	210.47 ± 5.77a	$196.98 \pm 1.39 \mathrm{ab}$
(Z)-3-Hexen-1-ol	$196.51 \pm 7.41 {\rm abc}$	$187.28\pm5.51cd$	$200.03\pm3.87\text{ab}$	$201.34 \pm 10.94a$	$193.23\pm3.20\text{abcd}$	$184.54 \pm 3.95 d$	$189.89\pm2.71\text{abc}$	$192.04\pm1.60\text{abcd}$	$187.41\pm2.04cd$
Total C6 alcohols	$1008.30 \pm 50.41$ ab	$992.54\pm53.05b$	$1134.22 \pm 32.38a$	1142.14 ± 136.84a	1133.91 ± 103.31a	$792.48 \pm 27.96d$	$959.30\pm60.92\text{bc}$	$1006.82\pm76.54ab$	$854.38\pm58.75cd$
1-Butanol	$4040.30 \pm 345.55a$	$772.79 \pm 37.30 \text{c}$	$124.96 \pm 8.20e$	$1329.87 \pm 241.77b$	$409.16 \pm 43.42 d$	$441.79 \pm 36.29 d$	$362.64\pm38.73\text{de}$	$581.47\pm57.34cd$	$564.28\pm46.96\text{cd}$
2-Methyl-1-propanol (mg/L)	$47.59\pm2.20 f$	$54.95\pm3.62\text{ef}$	$80.88\pm8.33d$	$64.47\pm5.81\mathrm{e}$	$66.86\pm1.49e$	$145.93 \pm 11.17 ab$	$152.29 \pm 13.66a$	$123.62 \pm 10.36 \mathrm{c}$	$133.97\pm6.64\text{bc}$
3-Methyl-1-butanol (mg/L)	$270.47 \pm 9.58a$	$161.82 \pm 1.70e$	$170.89\pm3.68\text{de}$	$230.98\pm26.34b$	$154.01 \pm 5.77 e$	$195.54 \pm 11.70 \mathrm{c}$	$185.78\pm6.02cd$	$220.04 \pm 10.95b$	$215.61 \pm 5.02b$
3-Methyl-1-pentanol	$506.55 \pm 24.01a$	$240.49 \pm 18.72 \text{c}$	$209.25\pm4.01d$	$312.51 \pm 29.96b$	$223.79\pm6.12cd$	$207.11 \pm 1.35 d$	$204.71 \pm 1.73d$	$216.63\pm2.10\text{cd}$	$214.85\pm0.93\text{cd}$
4-Methyl-1-pentanol	248.78 ± 1.97a	$202.89\pm3.45c$	Nd	$217.59 \pm 5.42b$	$199.11 \pm 0.89c$	Nd	Nd	$202.63\pm0.21\mathrm{c}$	$201.92 \pm 1.09c$
1-Heptanol	$115.14 \pm 20.22a$	$16.20\pm0.58c$	Nd	$30.05\pm8.22b$	Nd	Nd	Nd	Nd	Nd
1-Octanol	$4.90 \pm 0.24a$	$0.80\pm0.19c$	Nd	$2.09\pm0.09b$	Nd	Nd	Nd	Nd	Nd
1-Non-anol	Nd	Nd	$18.61 \pm 0.01a$	Nd	Nd	Nd	$18.00\pm0.12b$	Nd	$16.51\pm0.03c$
2-Non-anol	$2.90\pm0.01d$	$3.77\pm0.47\mathrm{c}$	$4.16\pm0.37b$	$4.40\pm0.14b$	$5.56 \pm 0.16a$	$3.14\pm0.02d$	Nd	$3.11\pm0.01d$	Nd
1-Decanol	$18.15 \pm 0.23a$	$16.88\pm0.11c$	$16.33\pm0.30\text{de}$	$17.54\pm0.65b$	$16.51\pm0.07 \text{cd}$	$15.88\pm0.19\mathrm{e}$	Nd	$16.10\pm0.09\text{de}$	Nd
Benzyl alcohol	$171.68 \pm 19.13a$	$88.70\pm7.45d$	$90.69\pm5.37 \text{cd}$	$117.96 \pm 15.45b$	$108.60\pm0.63\text{bc}$	$98.38 \pm 16.30 \text{cd}$	$27.59\pm2.67e$	$83.01\pm6.64\mathrm{d}$	$12.32\pm2.68e$
Phenylethyl alcohol (mg/L)	$248.34 \pm 7.74a$	$57.14\pm6.52cd$	$40.10\pm4.42d$	$157.63 \pm 28.75b$	$51.84\pm7.65cd$	$53.12\pm4.73cd$	$50.63\pm2.62cd$	$66.87\pm3.52\mathrm{c}$	$59.91\pm4.55cd$
Leavo-2,3-Butanediol (mg/L)	$44.51 \pm 1.23a$	$25.48\pm2.91\text{cd}$	$33.84\pm6.05b$	$35.91\pm6.26\mathrm{b}$	$31.02\pm3.02 \text{bc}$	$44.36\pm0.93a$	$23.54\pm1.13d$	$26.53\pm1.39\text{cd}$	$25.13\pm2.07 \text{cd}$
Meso-2,3-Butanediol (mg/L)	$15.91 \pm 0.94 d$	$7.80\pm0.92e$	$13.54\pm1.02d$	$8.64\pm0.71\mathrm{e}$	$9.88\pm0.76\mathrm{e}$	$26.02\pm2.77c$	$27.52\pm1.50\text{bc}$	$30.22\pm2.13ab$	$31.36 \pm 3.13a$
Total higher alcohols (mg/L)	631.93 ± 4.59a	$\textbf{308.54} \pm \textbf{5.83d}$	$339.71 \pm 2.67d$	$499.66 \pm 67.47b$	$314.57 \pm 16.83d$	$465.72\pm23.78bc$	$440.37 \pm 20.59c$	$468.38\pm24.88bc$	467.00 ± 8.76bc
Hexanoic acid	$739.90 \pm 72.57a$	$184.39 \pm 15.85 \text{bc}$	$117.29 \pm 1.53 d$	$233.68 \pm 60.55 \text{b}$	$138.13 \pm 1.00$ cd	Nd	Nd	Nd	Nd
Octanoic acid	$366.86 \pm 48.47a$	$49.60\pm5.31\mathrm{c}$	$16.13\pm2.00d$	$98.97\pm9.07\mathrm{b}$	$47.21 \pm 7.77c$	Nd	Nd	$25.60\pm6.17 \text{cd}$	$12.58\pm1.00d$
Decanoic acid	$8.65\pm0.50\text{b}$	$13.03\pm0.02a$	$8.24\pm0.15\mathrm{c}$	$8.24\pm0.12c$	$8.30\pm0.04c$	Nd	Nd	Nd	Nd
Total fatty acids	1115.41 ± 85.16a	$\textbf{247.03} \pm \textbf{20.90c}$	$141.66 \pm 1.20d$	$340.88 \pm \mathbf{68.32b}$	$193.64\pm8.39cd$	Nd	Nd	$25.60\pm6.17e$	$12.58 \pm 1.00 e$
Ethyl acetate (mg/L)	$46.96 \pm 2.26h$	$67.27 \pm 6.11$ fg	$155.85 \pm 17.39e$	$48.87 \pm 8.18$ gh	$72.29\pm2.11\mathrm{f}$	$248.01 \pm 10.57 d$	$270.84 \pm 16.48 c$	$318.55 \pm 15.36b$	$369.25 \pm 8.98a$

TABLE 2 | Volatile composition ( $\mu$ g/L) of the final wines after alcoholic fermentation.

(Continued)

#### TABLE 2 | (Continued)

Aroma compounds	SC	ZBI-2	ZBI-3	ZBII-2	ZBII-3	PKI-2	PKI-3	PKII-2	PKII-3
3-Methylbutyl acetate	$294.91 \pm 30.54$ d	$324.97 \pm 57.40d$	$364.40 \pm 81.11d$	$274.68 \pm 9.30d$	$304.95 \pm 19.12d$	$1181.00 \pm 76.21c$	1473.47 ± 295.29b	1540.21 ± 289.05b	2160.99 ± 265.52a
Phenethyl acetate	154.45 ± 16.87ab	c $110.91 \pm 7.54 f$	$119.07 \pm 7.66 f$	145.19 ± 11.82bcd	123.82 ± 6.14 ef	$128.46\pm6.57 def$	$138.14\pm9.24cde$	162.23 ± 10.04ab	169.34 ± 11.37a
2-Methylpropyl acetate	$60.01 \pm 12.29 d$	$78.67\pm6.67 \text{cd}$	$90.02\pm6.45\mathrm{c}$	$58.89 \pm 11.64 d$	$69.73\pm7.78cd$	$248.05 \pm 17.27 b$	$267.54\pm6.62b$	$246.62\pm30.37b$	$311.34 \pm 16.62a$
Total acetate esters (mg/L)	47.47 ± 2.24h	$\textbf{67.79} \pm \textbf{6.17fg}$	$156.42 \pm 17.36e$	$49.35\pm8.17\text{gh}$	$72.78 \pm 2.14 f$	$249.57 \pm 10.50 d$	$272.72 \pm 16.78c$	$320.50\pm15.07b$	371.89 ± 8.98a
Ethyl propanoate	$224.67 \pm 27.15 e$	$274.57 \pm 24.17 de$	$557.89 \pm 32.82 \mathrm{c}$	$328.29 \pm 19.76 d$	$312.35 \pm 15.21 d$	$684.09 \pm 32.09a$	710.11 ± 51.24a	$619.23\pm24.19b$	$686.04 \pm 36.38a$
Ethyl butanoate	173.29 ± 11.29ab	$197.58 \pm 9.60a$	184.81 ± 16.06ab	$167.82 \pm 28.91 \mathrm{b}$	$192.84 \pm 17.40 ab$	$139.24 \pm 8.88c$	$113.80 \pm 0.44 d$	$106.07 \pm 6.61 d$	$103.37 \pm 1.42d$
Ethyl 2-methylpropanoate	$39.10\pm1.20\mathrm{f}$	$42.64 \pm 1.12 f$	$58.02 \pm 5.23 e$	$41.40\pm2.15 \mathrm{f}$	$43.42\pm1.21\mathrm{f}$	$133.53 \pm 12.51 d$	$149.09\pm2.87c$	$161.57 \pm 14.62b$	$200.12 \pm 5.64a$
Ethyl hexanoate	$369.72 \pm 26.62a$	$145.28 \pm 17.59 b$	$41.68\pm8.30d$	$160.70 \pm 22.63b$	$77.96 \pm 12.59c$	$73.45\pm9.03\mathrm{c}$	$61.74\pm9.29 \text{cd}$	$84.15 \pm 10.06 \mathrm{c}$	$39.80\pm8.03d$
Ethyl 2-hexenoate	Nd	Nd	$6.31 \pm 0.06b$	Nd	Nd	$7.97 \pm 0.73 a$	$8.41 \pm 1.21a$	$8.24 \pm 0.84a$	$7.42\pm0.50a$
Ethyl heptanoate	$6.87 \pm 0.75 a$	$3.69 \pm 0.15c$	$4.09\pm0.96 \rm bc$	$4.04\pm0.36\text{bc}$	$3.57\pm0.35\mathrm{c}$	$4.97\pm0.71b$	$4.19\pm0.28 \text{bc}$	$4.54\pm0.20 \mathrm{bc}$	$3.84\pm0.17\mathrm{c}$
Ethyl octanoate	$748.83 \pm 70.80a$	$204.24 \pm 25.37b$	$63.74 \pm 7.33e$	$219.55 \pm 75.42b$	$122.01 \pm 9.26$ cde	$162.39 \pm 58.66 { m bc}$	$74.11 \pm 9.14$ de	$146.79\pm9.75bcd$	$73.63\pm8.30\text{de}$
Ethyl decanoate	$567.55 \pm 66.16a$	$107.02 \pm 12.05c$	$33.20\pm3.18d$	$173.12 \pm 22.76b$	$74.29 \pm 11.92 \text{cd}$	$56.32\pm3.63d$	$33.00\pm0.71d$	$62.37\pm7.98\mathrm{d}$	$34.96\pm2.33d$
Ethyl phenylacetate	$50.17 \pm 1.13a$	$46.65 \pm 1.00$ de	$46.20 \pm 0.27e$	$49.28 \pm 1.26 ab$	Nd	$48.06\pm0.23 \text{bc}$	$47.65\pm0.13cd$	$48.80\pm0.25\text{bc}$	$49.17\pm0.30\text{ab}$
Ethyl undecanoate	$34.54 \pm 0.02a$	$33.95 \pm 0.58b$	Nd	$34.40 \pm 0.12a$	Nd	Nd	Nd	$34.28\pm0.06\text{ab}$	Nd
Ethyl dodecanoate	$138.47 \pm 16.41a$	$43.62\pm1.54\text{bc}$	$38.63 \pm 4.15c$	$52.26\pm8.44b$	$40.39\pm1.82c$	$37.77 \pm 0.77c$	$35.97 \pm 0.08 \mathrm{c}$	$37.90\pm0.73\mathrm{c}$	$36.11\pm0.31\mathrm{c}$
Ethyl tetradecanoate	$36.96 \pm 0.28a$	$34.24 \pm 1.00b$	$34.27\pm0.01b$	$34.08\pm0.91\mathrm{b}$	Nd	Nd	Nd	Nd	Nd
Diethyl succinate	$22.32 \pm 2.40a$	$15.73\pm0.68 \text{bc}$	$10.54\pm2.39d$	$16.63 \pm 1.87 b$	$13.64 \pm 1.57c$	Nd	Nd	Nd	Nd
Total ethyl esters	2412.48 ± 212.00a	1149.19 ± 27.95cd	1079.40 ± 63.74d	1270.48 ± 165.29bc	880.47 ± 37.57e	1347.80 ± 32.03b	1238.07 ± 64.83bcd	1313.95 ± 34.76bc	1234.47 ± 56.66bcd
Methyl decanoate	$54.25 \pm 0.33a$	Nd	$53.51 \pm 0.01 \mathrm{b}$	$53.54\pm0.07b$	$53.50\pm0.06\mathrm{b}$	Nd	Nd	Nd	Nd
2-Methylpropyl decanoate	$34.95 \pm 0.09a$	$34.18\pm0.02b$	Nd	$34.37\pm0.02ab$	Nd	$33.86\pm0.58b$	Nd	$34.26\pm0.75b$	Nd
Octanoic acid, 3-methylbutyl ester	$54.83 \pm 1.46a$	$36.47\pm0.44c$	$36.05\pm0.01\text{cd}$	$38.52\pm1.69b$	$35.68\pm0.32cd$	$35.24\pm0.29\text{cd}$	Nd	$35.62\pm0.28cd$	$34.69\pm0.22d$
Ethyl 9-decenoate	$68.72 \pm 7.09a$	$24.64\pm0.66 \text{bc}$	$21.33 \pm 1.10c$	$26.44\pm2.70b$	$21.78\pm0.83 \text{bc}$	$22.10\pm1.02\text{bc}$	$20.99\pm0.13c$	$21.55\pm0.16\text{bc}$	$20.99\pm1.00c$
3-Methylbutyl hexanoate	$15.96 \pm 0.71a$	$9.26\pm0.27\mathrm{c}$	$9.27\pm0.10c$	$10.59\pm1.03b$	$8.91\pm0.21\mathrm{c}$	$9.19\pm0.51\mathrm{c}$	$8.87\pm0.08\mathrm{c}$	$9.58\pm0.03$ bc	$8.36 \pm 1.35 \mathrm{c}$
Total other esters	$228.71 \pm 9.16a$	$104.55\pm1.27d$	$120.16\pm1.11\mathrm{c}$	163.46 ± 5.44b	$119.87 \pm 1.24c$	$100.40\pm2.21d$	$29.86 \pm 0.21$ g	$\textbf{78.16} \pm \textbf{20.20e}$	$64.05\pm2.41\mathrm{f}$
β-Damascenone	$15.56 \pm 0.95a$	$10.05\pm0.66bcd$	$9.99 \pm 1.13$ bcd	$11.35 \pm 1.05b$	$11.11 \pm 0.37 bc$	$9.28\pm0.97\text{d}$	$9.02\pm0.73d$	$9.76\pm0.23$ cd	$9.99\pm0.46\text{bcd}$
Benzaldehyde	$4.72 \pm 0.92a$	$1.10\pm0.25b$	$0.85\pm0.24$ bc	$1.21\pm0.11b$	$1.19 \pm 0.11b$	$0.66\pm0.02 \mathrm{bc}$	$0.58\pm0.02 \text{bc}$	$0.71\pm0.10$ bc	$0.41\pm0.02c$
2-Non-anone	$1.32\pm0.08a$	$0.51\pm0.06\text{bc}$	$0.41\pm0.01\mathrm{c}$	$0.52\pm0.03\text{bc}$	$0.50\pm0.09\text{bc}$	$0.55\pm0.01\text{b}$	Nd	Nd	$0.12\pm0.01d$
4-Heptanone, 2,6-dimethyl-	$7.82\pm0.99\mathrm{bc}$	$10.06 \pm 0.35 \mathrm{abc}$	$7.65\pm0.70\mathrm{c}$	$11.64 \pm 1.38a$	$11.06 \pm 2.26a$	10.25 ± 1.20ab	$11.29 \pm 1.84a$	$9.24 \pm 1.34 \mathrm{abc}$	$10.05 \pm 0.94$ abc
Others	$43.28 \pm 3.21a$	$\textbf{33.40} \pm \textbf{0.86bcd}$	$\textbf{27.82} \pm \textbf{2.65d}$	$\textbf{38.39} \pm \textbf{2.61ab}$	$\textbf{36.61} \pm \textbf{4.61bc}$	$\textbf{32.22} \pm \textbf{3.22cd}$	$\textbf{37.75} \pm \textbf{3.18cd}$	$\textbf{29.66} \pm \textbf{2.62d}$	$\textbf{31.17} \pm \textbf{2.21cd}$
Total Volatile composition (mg/L)	$684.20\pm2.61c$	$378.85\pm8.24 \mathrm{f}$	$498.62 \pm 17.66e$	$551.95 \pm 59.62d$	$389.71 \pm 15.43f$	$717.55 \pm 33.60c$	$715.34\pm6.78c$	$\textbf{791.42} \pm \textbf{39.92b}$	841.07 ± 17.80a

SC, single inoculation with S. cerevisiae EC1118; ZBI-2, sequential inoculation with Z. bailii BJV11007, followed by inoculation with EC1118 after 2 days; ZBI-3, sequential inoculation with Z. bailii BJV11007, followed by inoculation with EC1118 after 3 days; ZBI-3, sequential inoculation with Z. bailii BJV11007, followed by inoculation with EC1118 after 2 days; ZBI-3, sequential inoculation with Z. bailii BJV11007, followed by inoculation with EC1118 after 2 days; ZBI-3, sequential inoculation with Z. bailii BJV11007, followed by inoculation with EC1118 after 2 days; ZBI-3, sequential inoculation with Z. bailii BJV15005, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJIV53006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJIV53006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJIV45006, followed by inoculation with EC1118 after 2 days; PKI-2, sequential inoculation with P. kudriavzevii BJII44006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJII44006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJII44006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJII44006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJII44006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJII44006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJII44006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJIV53006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential inoculation with P. kudriavzevii BJIV53006, followed by inoculation with EC1118 after 2 days; PKI-3, sequential ino

alcohols. In C6 alcohol, the content of 1-hexanol (contributing to "herbaceous," "grass," and "woody" notes for wine) was the highest. PK could significantly reduce the content of 1-hexanol.

In this work, the total higher alcohols ranged from 308.54 mg/L (ZBI-2) to 631.93 mg/L (SC) (Table 2). Sequential fermentations could reduce the total higher alcohol content. 2-Methyl-1-propanol, 3-methyl-1-butanol, 3-methyl-1-pentanol, and phenylethyl alcohol were detected above their thresholds, and 1-heptanol, leavo-2,3-butanediol, and meso-2,3-butanediol were detected above their sub-thresholds (Supplementary Table 4). Sequentially fermented wines were characterized by significantly higher concentrations of 2-methyl-1-propanol (contributing to "alcohol," "solvent," "green," and "bitter" notes for wine) and significantly lower amounts of 3-methyl-1butanol, 3-methyl-1-pentanol, 4-methyl-1-pentanol, 1-heptanol, and phenylethyl alcohol. ZB could reduce the content of leavo-2,3-butanediol and meso-2,3-butanediol, while PK could significantly increase meso-2,3-butanediol content and decrease leavo-2,3-butanediol content, except PKI-2.

#### **Fatty Acids**

Three fatty acids (hexanoic acid, octanoic acid, and decanoic acid) were detected in this study (**Table 2**); among them, hexanoic acid and octanoic acid were the only two with OAVs above 0.1, and the former was up to 1 (**Supplementary Table 4**). The total fatty acid content was significantly lower in sequential fermentations. In PK, only a small amount of octanoic acid was detected in PKII-2 and PKII-3.

#### Esters

Esters can generally be categorized into acetate esters, fatty acid ethyl esters, and other esters. In this study, 22 esters were identified, and eight esters exceeded their thresholds (**Table 2** and **Supplementary Table 4**), including ethyl acetate, 3-methylbutyl acetate, ethyl 2-methylpropanoate, ethyl hexanoate, ethyl heptanoate, ethyl octanoate, ethyl decanoate, and methyl decanoate.

Four acetate esters were detected in this study, and compared with SC, PK, and ZB significantly increased the content of acetate esters (Table 2). This result was reconfirmed on the third day of inoculation of S. cerevisiae. In esters, the content of ethyl acetate was the highest. P. kudriavzevii and Z. bailli could significantly increase the content of ethyl acetate in sequential fermentation, especially for P. kudriavzevii where the highest content of ethyl acetate could be increased to 369.25 mg/L. In addition, PK significantly increased the content of 3-methylbutyl acetate and 2-methylpropyl acetate. Compared with SC, sequential fermentation significantly reduced the total ethyl ester content. In all the sequentially fermented wines, the content of mediumchain fatty acid ethyl esters such as ethyl hexanoate, ethyl heptanoate, ethyl octanoate, and ethyl decanoate significantly decreased, while the content of ethyl propanoate significantly increased. Z. Bailli could increase the content of ethyl butanoate and ethyl 2-methylpropanoate. P. kudriavzevii could significantly increase the content of ethyl 2-methylpropanoate and decrease the content of ethyl butanoate. In addition to acetate and ethyl esters, five other esters were detected in this study (Table 2). The OAV of methyl decanoate was above 1 in some samples, and ethyl 9-decenoate content was higher than its subthreshold (**Supplementary Table 4**).

#### Others

A total of four other volatile compounds were detected in this study, including  $\beta$ -damascenone, benzaldehyde, 2-nonanone, and 4-heptanone, 2,6-dimethyl- (**Table 2**).  $\beta$ -Damascenone contributed the "sweet," "exotic flower," and "stewed apple" aromatic notes to the wine. Compared with SC, sequential fermentation reduced the content of  $\beta$ -damascenone, but the OAV value in pure fermentation was still much higher than 1 (**Supplementary Table 4**). In addition, 2-nonanone content was higher than its subthreshold, which contributed to the fruity aroma of the wine.

#### Principal Component Analysis and Hierarchical Cluster Analysis of Wine Aroma Compounds in Different Fermentations

To highlight the differences in the fermentation of different inoculation strategies and to identify the effects of these treatments on volatile compounds, PCA was performed on 27 volatile compounds with OAV above 0.1. As shown in Figures 3A,B, the first two components accounted for 81% (58% for PC1 and 23% for PC2) of the total variance. The PCs were roughly distinguished by wine samples fermented by different inoculation strategies. Wines produced by SC were separated from the other wines by PC1. The main components responsible for this separation were 3-methyl-1butanol, leavo-2,3-butanediol, 1-heptanol, phenylethyl alcohol, 3-methyl-1-pentanol, octanoic acid, hexanoic acid, ethyl heptanoate, ethyl octanoate, ethyl hexanoate, ethyl decanoate, and β-damascenone. PC2 separated PK from ZB, mainly by 2-methyl-1-propanol, meso-2,3-butanediol, ethyl acetate, 2methylpropyl acetate, 3-methylbutyl acetate, ethyl propanoate, and ethyl 2-methylpropanoate. These data indicate that the volatile compounds of S. cerevisiae can be further affected by sequential fermentation with non-Saccharomyces yeast.

The dendrogram of Hierarchical Cluster Analysis (HCA) was used to visualize the differences in volatile compounds among different inoculation strategies (**Figure 3C**). The results showed that the samples can be divided into two categories: pure fermentation and sequential fermentations. Notably, the sequential fermentation of two non-*Saccharomyces* yeast strains of the same species was divided into two groups according to the inoculation time of *S. cerevisiae*, which indicates the inoculation time of *S. cerevisiae* had a significant effect on the volatile compounds of sequentially fermented wine.

#### **Sensory Evaluation**

The sensory evaluation results of the wine samples were derived from 12 assessors, and the aroma radar map was drawn as shown in **Figure 4**. It can be noticed that yeast strains and inoculation strategies have a great influence on the aroma characteristics of wine. The score of "aromatic intensity" and "wine body" in



sequential fermented wines was higher than those of SC (except ZBI-2 and ZBII-2). Compared with SC, PKI-3 and PKII-3 had a high aroma note for floral, fruity and sweet, and a low aroma note for herbaceous and fatty. In addition, PK had a high aroma note for solvent, in agreement with their high levels of the ethyl acetate; ZB had a low aroma note for fruity, in agreement with their low levels of higher alcohols and esters. It was noteworthy that, PKII-3 had a high aroma note for floral, fruity, sweet, and aromatic intensity, and it also had a high score of "wine body." These results showed that non-*Saccharomyces* yeasts selected from the Baijiu fermentation environment have the potential to improve the flavor and quality of wine.

## DISCUSSION

To explore the application potential for Baijiu non-Saccharomyces yeast strains in wine fermentation, the effect of four Baijiu non-Saccharomyces yeast strains (two Z. bailii and two P. kudriavzevii) inoculated with one S. cerevisiae (commercial strain EC1118) on the physicochemical parameters and volatile compositions of wine was investigated in this study.

The contribution of yeast to wine volatile compounds is largely dependent on the persistence of strains and the number

of yeast cells during alcoholic fermentation. In this study, it was found that there was no obvious antagonism between S. cerevisiae and Z. bailli or P. kudriavzevii, and all strains could be detected at the end of alcohol fermentation. The fermentation process of sequential fermentations was significantly longer than that of S. cerevisiae pure fermentation, because Z. bailli and P. kudriavzevii consumed glucose and fructose significantly slower than S. cerevisiae (Supplementary Figure 1). The acidity of grape juice and wine directly affects their sensory quality and physical, biochemical, and microbial stability (Swiegers and Pretorius, 2005). Acetic acid usually accounts for about 90% of the volatile acids in wine (Swiegers and Pretorius, 2005). At the concentration of 0.7-1.1 g/L, acetic acid imparts an unpleasant smell to wine, and the best concentration is 0.2-0.7 g/L (Swiegers and Pretorius, 2005). In this study, sequential fermentation (except for ZBI-2 and ZBI-3) could significantly reduce the acetic acid content. Previous studies have found that Z. bailii can produce acetic acid (Xu et al., 2017b). In this study, the two Z. bailii strains showed different characteristics of acetic acid production, which may be related to the differences between strains. Non-volatile organic acids have a direct impact on the quality of wine; the imbalance of this component will affect its physicochemical and sensory properties (Gawel et al., 2007) and change its microbial properties (Delcourt



et al., 1995; Pretorius, 2000). In this study, the malic acid content of sequential fermentation was higher and the lactic acid content was lower, which may lower the quality of wine but can be improved by malolactic fermentation. Our results clearly demonstrate that sequential fermentation of *Z. bailli* and *S. cerevisiae* could significantly increase the content of succinic acid in agreement with the research results of Zhu et al. (2020). It may be due to the non-*Saccharomyces* yeast can exhibit low activity through acetaldehyde pathway which trigger an important redistribution of fluxes through the central metabolic network (Englezos et al., 2018). The high level of acetyl-CoA in *Z. bailli*, leads to an increase in the level of  $\alpha$ -ketoglutarate and a significant increase in succinic acid content.

Yeast is one of the important factors affecting wine fermentation aroma. Our results demonstrate that two non-*Saccharomyces* yeast strains of the same species had common effects on wine volatile compounds. Consistent with previous literature (Shi et al., 2019), we found that the inoculation time of *S. cerevisiae* affected the volatile compositions of the wines. Notably, through HCA (**Figure 3C**), we found in sequential fermentations, compared with the difference between the two non-*Saccharomyces* yeast strains of the same species, the inoculation time of *S. cerevisiae* had a greater effect on

wine volatile compounds. Higher alcohols are produced by the decarboxylation and dehydrogenation of  $\alpha$ -ketoacids through the Ehrlich pathway and Harris pathway (Wang et al., 2020). When the concentration is low (<300 mg/L), they can increase the complexity of wine aroma (Swiegers and Pretorius, 2005), while at higher concentrations (> 400 mg/L), higher alcohols weaken the fresh fruit aroma, enhance the pepper characteristics of young red wine, and harm the overall flavor of wine (Aznar et al., 2003; San-Juan et al., 2011). In this study, the content of total higher alcohols in SC was as high as 631.93 mg/L (Table 2), which mainly consisted of 2-methyl-1-propanol (47.59 mg/L), isoamyl alcohol (270.47 mg/L), and phenylethyl alcohol (248.34 mg/L). Sequential fermentation of non-Saccharomyces yeasts and S. cerevisiae significantly reduced the content of higher alcohols (308.54-499.66 mg/L). The total amounts of higher alcohols in all treatments were above 300 mg/L (Table 2), which may be related to the grapes used in this experiment. 2-Methyl-1-propanol and 3-methyl-1-butanol are produced by yeasts during alcoholic fermentation through the conversion of valine and isoleucine, respectively, via the Ehrlich pathway (Hazelwood et al., 2008). It was worth noting that compared with SC, the content of 2-methyl-1-propanol in all sequentially fermented wines significantly increased, while the



content of isoamyl alcohol significantly decreased. It may be due to changes in acetyl-coenzyme (acetyl-CoA) availability, which is required for the conversion of  $\alpha$ -ketoisovalerate, the precursor of 2-methyl-1-propanol, into  $\alpha$ -ketoisocaproic, the precursor of isoamyl alcohol (Hazelwood et al., 2008). The concentration of phenylethyl alcohol is higher than its threshold, which contributes to the rose aroma of wine (Tristezza et al., 2016). In this study, the OAV of phenylethyl alcohol in all samples was above 1. Sequential fermentation significantly reduced the content of phenylethyl alcohol and may reduce the negative effect of high higher alcohol content on wine.

Volatile fatty acids are essential to the aroma of the wine. When the content of volatile fatty acids is at the subthreshold, it will have a positive effect on the aroma of wine, and when it exceeds the threshold, it will spoil the wine aroma (Swiegers and Pretorius, 2005). In this study, the OAV of hexanoic acid > 1 in SC may bring a fatty flavor to the wine and adversely affect the aroma of the wine. Sequential fermentation reduced the content of hexanoic acid, and the OAV of hexanoic acid in ZB was more than 0.1, which could render a "cheese" note to the wine. In addition, the volatile fatty acid is the precursor of fatty acid ethyl ester synthesis (Hernández-Orte et al., 2006). Sequential fermentation reduces the content of volatile fatty acids, leading to the decrease of fatty acid ethyl ester content.

Yeast strains showed high specificity in total ester yield and acetate and ethyl ester patterns, which led to sensory differences in wine (Soles et al., 1982). In this study, *Z. bailli* and *P. kudriavzevii* were associated with a higher production of acetate esters and lower production of ethyl esters. Ethyl acetate is the main ester in wine (Moreira et al., 2011), and it can be biosynthesized by acetyl-CoA and ethanol through the reaction catalyzed by alcohol acyltransferase (Shi et al., 2021), which adds a pleasant fruit aroma to wine at low concentrations. However, when the concentration of ethyl acetate is higher than 150 mg/L, the chemical odor of varnish may damage the aroma of wine (Peinado et al., 2004). In this study, the ethyl acetate concentration of ZB was between 48.87 mg/L and 155.85 mg/L, and the ethyl acetate concentration of PK was between 248.01 mg/L and 369.25 mg/L, which was much higher than 150 mg/L and would thus have a negative effect on wine aroma (Comitini et al., 2011; Mateo and Maicas, 2016). In the future, when making wine with P. kudriavzevii, we can try to reduce the negative effects of very high concentrations of ethyl acetate by reducing the inoculation amount of *P. kudriavzevii* or adopting the strategy of simultaneous inoculation. In addition, compared with SC, PK could also significantly increase the content of 3-methylbutyl acetate and 2-methylpropyl acetate in agreement with the observations of Padilla et al. (2016) and Luan et al. (2018). Ethyl ester is another important group of esters in wine, and they are produced during yeast fermentation through ethanolysis of acyl-CoA that is formed during fatty acid synthesis or degradation (Swiegers and Pretorius, 2005). In this study, compared with SC, the ethyl ester content of wine fermented in the sequence was lower, mainly because the content of mediumchain fatty acid esters (ethyl esters of fatty acids with 6-12 carbon atoms) was reduced by sequential fermentation. It may be caused by the decrease of fatty acid content as its precursor (Saerens et al., 2010). In addition, compared with SC, sequential fermentation could significantly increase the content of ethyl propionate and ethyl 2-methylpropanoate, and ZB could also significantly increase the content of ethyl butanoate, rendering "banana" and "pear" notes to the wine.

#### CONCLUSION

In summary, these experiments indicate that there was no obvious antagonism between S. cerevisiae and Z. bailli or P. kudriavzevii in sequential fermentations, and all strains could be detected at the end of alcoholic fermentation. Compared with the pure fermentation of S. cerevisiae, the sequential fermentation of Z. bailii and S. cerevisiae could significantly reduce the content of higher alcohols and ethyl esters and increase the content of acetate esters; the sequential fermentation of P. kudriavzevii and S. cerevisiae could significantly reduce the content of C6 alcohols, total higher alcohols, and ethyl esters and significantly increased the contents of acetate esters (especially ethyl acetate and 3-methylbutyl acetate). Sequential fermentation of Baijiu non-Saccharomyces yeast and S. cerevisiae improved the flavor and quality of wine due to the higher ester content and lower concentration of higher alcohols and fatty acids, non-Saccharomyces yeasts selected from the Baijiu fermentation environment have potential applications in winemaking, which could provide a new strategy to improve wine flavor and quality.

#### DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author/s.

#### **AUTHOR CONTRIBUTIONS**

X-LW and Y-FC designed the experiments. R-RL, MX, JZ, Y-JL, and C-HS conducted the experiments. R-RL, MX, JZ, Y-JL, C-HS, and HW analyzed the experimental data. R-RL wrote the manuscript. X-WG and D-GX contributed to data curation. All authors contributed to the article and approved the submitted version.

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#### SUPPLEMENTARY MATERIAL

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