

Non-targeted metabolomic approach as a tool to evaluate the chemical and volatile profile of sparkling wines fermented with autochthonous yeast strains

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Aim

Champenoise method The is based ``in bottle" on refermentation usually driven by few commercial strains belonging to the *S. cerevisiae* species. In this study, the impact of selected autochthonous yeast strains on the chemical profile of sparkling wines (SW) has been evaluated through nontargeted metabolomic approach based on HPLC-HRMS and GC-**MS** techniques. The HPLC-HRMS/GC-MS correlation analysis was permitted to draw a map that constitutes a useful tool to monitor the different patterns of aroma release operated by the indigenous strains.

Methods



The autochthonous cerevisiae strains employed were previously selected [1] and deposited in the ITEM Culture Collection of the CNR-ISPA (www.ispacnr.it/ collezioni-microbiche). The commercial *S. cerevisiae* DV10 (Lallemand, USA) was used as control.



Production of sparkling wine made using the was traditional method [2].

Results

Volatolomic Profile

- A total of **26 volatiles** belonging to higher alcohols, esters, terpenes, acids, were identified;
- Quantitative differences of the above compounds were detected in the produced wines;
- The selected strains produced higher concentrations of esters (ethyl and acetate), alcohols such as 2-phenylethanol and lower values (<400 ppm) of higher alcohols compared to the wine produced with the control veast;
- Moreover, the SW produced with selected yeast showed a **low volatile acidity** (<0.3 g/L), high glycerol content, a **good phenolic and acid profile** that influences the foam stability and the sensorial quality (body, bitterness and astringency);
- The Figure 1 shows the score plot reporting, for each sample, the Mean Standardized Concentration (MSC) as a function of the number of volatiles (NV) (reported in percentage) with a concentration higher or equal to the mean. This scatterplot allowed bettering outlining the differences among the samples that on the plane of the three PCs are clustered. **ITEM 9518**, 17294, 14077, 9520 samples differ from both the control and other samples by a higher number of volatiles in concentration above the mean.

HPLC-HRMS results

- 61 compounds belonging to several metabolite classes were identified;
- HPLC-HRMS analysis showed that **DV10** sample was characterized by high values of **gluconic** acid that adversely affects wine foamability, while lower concentrations of this molecule were detected in sparkling wines produced by the other strains selected in Apulia region. In particular, samples **9518**, **14077** and **17294** showed a high content of polysaccharide that promotes a better foam stability and improves the sensory quality.
- A HPLC-HRMS/GC-MS correlation analysis (Fig.2) was performed and permitted to obtain a map that constitutes a useful tool to monitor the different patterns of aroma release.
- Finally, this contribution provide information to modulate the quality of regional sparkling wines from minor autochthonous grape varieties using selected microbial resources.



References

- [1] Grieco et al. (2011). https://doi.org/10.1007/s13213-010-0091-7
- 2] Garofalo et al. (2018). https://doi.org/10.1016/j.ijfoodmicro.2018.07.004
- [3] Tufariello et al. (2019). https://doi.org/10.1016/j.lwt.2019.03.063
- [4] Rizzuti et al. (2015). https://doi.org/10.1016/j.foodcont.2021.108099





WineScan Flex was used to determine the principal chemical parameters of SW

Chemical analyses



The volatile profile was analyzed using SPME-GC/MS [3].



HPLC/High Resolution Mass Spectrometry analysis [4],

