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Melatonin and its derivatives in red wine: contribution of fermenting microorganisms

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INTRODUCTION

✓ Melatonin (N-acetyl-5-methoxytryptamine; MEL) is an indoleamine produced mainly by the pineal gland in vertebrates [1].

✓ MEL carries out several functions: modulation of the circadian and circannual rhythms, reproductive function, bone metabolism and turnover via

- cell-receptor-mediated mechanisms. It shows an antioxidant activity directly scavenging the free radical species and stimulating the activity of antioxidant enzymes [2].
- ✓ MEL has been found in plant foods, such as seeds, fruits and fermented beverages, wine included [3-5].
- ✓ The circulating levels of MEL are very low (about 0.2 ng/mL at the maximum night peak and lower than 0.01 ng/mL during the day) [6] compared to MEL in grape products (about 1 ng/g in berry skin and 0.5 ng/mL in wine) and, therefore, the intake of grape products is of particular interest [4,7].
- ✓ The content of MEL increases during the fermentation step of the winemaking and other fermented beverages meaning the yeast plays a significant role in its biosynthesis [8-9].
- ✓ MEL isomers were also detected in wine and one of them was recently identified as tryptophan-ethyl ester (TEE) [10-11].

AIMS

Set up of a sample preparation procedure and the validation of an analytical method

for the simultaneous detection of MEL, TEE and tryptophan (TRP).

MATERIALS AND METHODS

Solid Phase Extraction (SPE) procedure

The final protocol included four steps:

- Sample loading (fraction A).
- Washing with formic acid 0.1% [v/v] (fraction B). Ο
- Washing with methanol 40% [v/v] (fraction C). Ο
- Elution with methanol 100% (fraction D) which was evaporated under vacuum.

Analytical method development and performances

The detection of the three compounds was carried out by HPLC coupled with both fluorescence (FL) and mass spectrometry (MS) detectors.

The validation was carried out in terms of selectivity, linearity, limit of detection (LOD), limit of quantification (LOQ), repeatability and recovery for both model wine solution (tartaric acid 5 g/L, ethanol 12% [v/v], pH 3.2) and red wine.

Re-suspension in methanol 10% acidified with formic acid 0.1% [v/v] Ο corresponding to a concentration folds of 10.

In order to prove the suitability of the method red wine samples (n=8) were analyzed.

RESULTS

| | Concentration range added (µg/L) | LOD (µg/L) (n=3) | LOQ (µg/L) (n=3) | Recovery (%) (n=6) | | Repeatability (%RSD) | |
|----------|--|------------------------|------------------------|--------------------|--------------------|-------------------------|--------------------|
| Compound | | | | SWS | Spiked red wine | SWS | Spiked red wine |
| TRP | 0.11-5500 | 0.75 | 1.25 | 89 | 84 | 9.1 | 10.5 |
| TEE | 5-250 | 0.038 | 0.12 | 88 | 76 | 6.5 | 7.9 |
| MEL | 5-250 | 0.0023 | 0.018 | 86 | 79 | 4.6 | 5.4 |

Table 1: Limits of detection (LOD) and quantification (LOQ), recovery (%) and repeatability (as Relative Standard Deviation, %RSD) for the analytical method developed in HPLC-MS.

- ✓ The SPE purification allowed both the removal of compounds potentially interfering the HPLC separation from the wine matrix and the concentration of the analytes of interest making possible their detection due to their presence in low amounts in wine.
- The analytes were revealed by both the detectors (FL and MS) that were investigated.
- ✓ Linear response was observed for both spiked model wine solution and red wine analyzed by MS detector.

- ✓ The repeatability of the SPE purification was evaluated its average values were: 5.4%, 7.4%, 4.0% and 4.2%, 5.2, 5.2% for TRP, TEE and MEL of spiked red wine and model wine solution, respectively.
- ✓ The intra-day precision (n = 12) corresponded to 3.8%, 2.6% and 3.1% for TRP, TEE and MEL, respectively.
- ✓ The developed analytical method resulted suitable for the determination of TRP, TEE and MEL in red wine (Table 2).
- ✓ TEE resulted over one thousand folds highly concentrated than MEL in the Nebbiolo wine samples analyzed.

| Cample code | TRP | TEE | MEL | |
|-------------|-----------|------------|-------------|--|
| Sample code | mg/L | μg/L | μg/L | |
| Red wine 1 | 3.85±0.40 | 172.2±13.6 | 0.057±0.003 | |
| Red wine 2 | 4.39±0.46 | 212.0±16.8 | 0.062±0.003 | |
| Red wine 3 | 1.56±0.16 | 256.2±20.2 | 0.063±0.004 | |
| Red wine 4 | 1.02±0.11 | 223.2±17.6 | 0.038±0.002 | |
| Red wine 5 | 0.98±0.10 | 113.0±8.9 | 0.046±0.003 | |
| Red wine 6 | 0.84±0.09 | 92.9±7.3 | 0.054±0.003 | |
| Red wine 7 | 0.44±0.05 | 71.7±5.7 | 0.063±0.004 | |
| Red wine 8 | 0.57±0.06 | 74.4±5.9 | 0.038±0.001 | |

✓ No significant difference was found between the two matrices.

Table 2: Levels of TRP (tryptophan), TEE (tryptophan ethyl ester) and MEL (melatonin) in Nebbiolo red wine. Data are expressed as mean \pm standard deviation.

CONCLUSIONS

- ✓ The proposed analytical method allowed the detection and the reliable quantification of TRP, TEE and MEL simultaneously.
- \checkmark Good linearity (r > 0.99), recovery (average 84%) and repeatability (average RSD 7.3%) were achieved.
- ✓ This method represents a useful tool for monitoring the release of MEL and TEE and their fate throughout the wine production and storage.
- ✓ The concentrations of TEE much higher than those of MEL in wine samples confirmed previous outcomes [10,12] even if the origin and nutritional role of this compound still need to be elucidated.

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ACKNOWLEDGMENT

The authors gratefully acknowledge the financial support provided by Piano di Sostegno alla Ricerca 2015/2017 – Linea 2 – Università degli Studi di Milano, Milan, Italy.



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