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Is there creatine in plants? The true compound behind the ¹H NMR signal at 3.05 ppm in plant extracts

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Introduction: Some papers describe the presence of creatine in plants, based on a singlet signal at 3.02–3.05 ppm in the ¹H NMR spectra. Although is there creatine in plants? Therefore, to answer this question, a comprehensive NMR investigation has been performed aiming the unambiguous assignment of the compound responsible for that signal.

Objective: Determine whether the compound behind the signal at 3.05 ppm is truly creatine or if it was just a misassignment, instead.

Methods: Samples of leaves and cherries from *Eugenia uniflora* in their natural swollen state were submitted to HR-MAS NMR analysis.

Results: It was found that the signal at 3.05 ppm was misassigned to creatine. The exhaustive NMR investigation revealed that the signal is related to the amino acid 4-hydroxy-*N*-methyl proline, instead.

Conclusion: The comprehensive NMR investigation revealed that there is no creatine in plants, it was just a misassignment.

KEYWORDS

NMR, HR-MAS NMR, *Eugenia uniflora*, 4-hydroxy-*N*-methyl proline, creatine, Brazilian cherry

Introduction

High-Resolution Magic-Angle-Spinning (HR-MAS) is a versatile technique that allow the acquisition of Nuclear Magnetic Resonance (NMR) data directly from semisolid samples, such as vegetable (e.g., leaves, flowers, fruits, etc.) and animal tissues, in their natural swollen state, without the need of any sample pre-treatment, being a valuable tool for the rapid investigation of plant chemical composition (Broberg and Kenne, 2000; Kelleher et al., 2006; Bharti et al., 2011; Alam and Jenkins, 2012; Choze et al., 2013; Farooq et al., 2013; André et al., 2014; Santos et al., 2015; Flores et al., 2018; Santos et al., 2018; Ali et al., 2020).

In a previous work, while investigating the chemical composition of *Berberis laurina* (Berberidaceae) by means of HR-MAS NMR spectroscopy, it was faced up a straight singlet signal at 3.02 ppm (Ali et al., 2020). This signal has been described only in NMR studies of *Eugenia uniflora* (Myrtaceae) (Iqbal et al., 2013; Flores et al., 2018), which is a native plant in tropical South America's Atlantic coast countries, including Suriname, Guyana, French



Guyana, Brazil, Uruguay, Paraguay, and Northern Argentina, known as Surinam cherry or Brazilian cherry. While it is currently cultivated and naturalized worldwide, mainly in tropical and subtropical regions.

Its edible fruits are botanical cherries that look like small orangered pumpkins (Supplementary Figure S1). Known for their exotic flavor, which ranges from sweet to sour, they are primarily consumed fresh, as well as in juice, frozen pulp, jams, jellies, and other forms. Eugenia uniflora also has several significant pharmacological properties being widely used in folk medicine. Several diseases and disorders such as bronchitis, cold, cough, gout, sore throat, hypertension, headaches, influenza, hepatic diseases, painful urination, rheumatism, diarrhea, fever, stomach diseases, and other gastro-intestinal disorders are treated based on its extracts. Likewise, it is used in the treatment of obesity, and diabetes as well as to stimulate menstrual flow. Also, it was described to have diuretic, insect repelling, digestive, eupeptic, and carminative properties. Its tea is used to facilitate the process of childbirth, while its hot water extract of the fresh leaves and unripe fruits are used to treat malaria and fever (Schapoval et al., 1994; Consolini et al., 1999; Begossi et al., 2002; Bagetti et al., 2011; Celli et al., 2011; Moura et al., 2018). Besides its innumerable therapeutic properties, E. uniflora is used by the cosmetic industries (Melo et al., 2007) as well as an important colonizing species in restoration of disturbed forest areas, and to provide food for wildlife (Rodrigues and Nave, 2000; Botrel et al., 2002; Margis et al., 2002; Bianchini et al., 2003; Pinto et al., 2005).

To the best of our knowledge, in many papers regarding plant tissues investigation through HR-MAS NMR technique, any singlet signal has been described around 3.02-3.05 ppm, except for *E. uniflora* (Myrtaceae) (Iqbal et al., 2013; Flores et al., 2018) and *B. laurina* (Berberidaceae) (Ali et al., 2020). This signal has been assigned to the N-CH₃ group of the amino acid creatine (Iqbal et al., 2013; Ali et al., 2020). However, this compound is synthesized in the liver, pancreas, and kidney from the transamination of the amino acids arginine, glycine, and methionine. Therefore, it is not produced by the kingdom plantae. There are only a few reports in which creatine was described in kingdom plantae such as in *Phaseolus mungo* (Fabaceae), *Lens culinaris* (Fabaceae), *E. uniflora* (Myrtaceae) (Azmat et al., 2009; Iqbal et al., 2013), and *Tussilago farfara* (Asteraceae) (Zhi et al., 2012).

Considering the biosynthesis of creatine, this compound might not have been found in the kingdom plantae, thus there may have a misassignment for the signal around 3.02–3.05 ppm. Therefore, in this work, NMR spectroscopy was exhaustively used to identify, in an unequivocal way, the true compound which gives rise to the signal at 3.02–3.05 ppm found in *E. uniflora*.

Materials and methods

Plant material

Leaves and cherries from *E. uniflora* L. (Myrtaceae) were collected in November 2021 from a healthy tree specimen in the backyard of the Department of Chemistry at Polytechnic Centre campus of the Federal University of Paraná, Curitiba, PR, Brazil [coordinates: 25°27′01.1″ S, 49°14′05.5″ W].

Sample preparation for NMR analysis

Fresh leaves and pitless cherries of E. uniflora were frozen in liquid nitrogen in a mortar and powdered with aid of pestle. The resulting powder (~10.0 \pm 0.5 mg) of each sample was then inserted into a 50 µL volume 4-mm zirconium oxide HR-MAS rotor, followed by the addition of 40 μ L of deuterium oxide (D₂O, 99.9% D, containing 0.01% w/v TMSP-d4). The sample inside the HR-MAS rotor was then homogenized with aid of a syringe needle, which also made it possible to remove air bubbles. The rotors were sealed with a small upper Kel-F spacer/Kel-F sealing screw cap to prevent the leakage of fluids due to high-spinning speed during NMR experiments, followed by a Kel-F spinning cap (Alam and Jenkins, 2012; Flores et al., 2018), and then submitted to ¹H HR-MAS NMR analysis. In order to support the NMR chemical shift assignments in the HR-MAS NMR analysis, solution NMR experiments from deuterated solvent extracts of E. uniflora leaves were also performed and compared to the HR-MAS results. Therefore, each powdered plant material (i.e., 100.0 ± 1.0 mg) was submitted to extraction directly with deuterated solvent (i.e., 700 µL of D₂O) in a 1.5 mL microcentrifuge tube, assisted by sonication for 10 min at 298 K (De Aquino et al., 2019). After that, samples were centrifuged at 12,000 rpm for 10 min and 600 μ L of supernatant were directly transferred into a 5 mm NMR tube and submitted to solution NMR analysis.



NMR analysis

¹H HR-MAS NMR spectra were acquired at 298K on a Bruker AVANCE 400 NMR spectrometer operating at 9.4 T, observing ¹H and ¹³C nuclei at 400.13 and 100.62 MHz, respectively. The spectrometer was equipped with a 4-mm four-channel (¹H/²H/ ¹³C/¹⁵N) HR-MAS probe with an actively shielded gradient field along the magic angle direction. The acquisition was performed with 5 kHz spinning speed and the spectra were acquired with 64k timedomain data points distributed over a spectral width of 20 ppm, resulting in a digital resolution of 0.24 Hz by applying single 90° excitation pulses with aid of zgpr, a water suppression pulse sequence, 1.0 s recycle delay and averaging 256 scans. The spectra were processed by applying an exponential Lorentzian multiplication with a line broadening factor of 1.0 Hz to the FID, followed by Fourier transform using a zero-filling factor of 2. Phase and baseline corrections were manually performed, and the NMR chemical shifts were referenced to TMSP-d4 signal at 0.00 ppm. Prior to the NMR analysis, the magic angle was adjusted using the ⁷⁹Br signal from powdered KBr as reference. Samples were locked on the deuterium signal from D₂O, and the magnetic field homogeneity was optimized for each sample. The total experiment time was 35 min, including the time required for rotor packing. Direct and long-range 1H-13C as well as 1H-1H NMR correlations experiments were acquired directly from deuterated solvent extracts on the same NMR instrument as above, although it was equipped with a 5-mm broadband solution direct detection four-channel (¹H/²H/¹³C/³¹P) probe with actively shielded gradient field along z-direction, instead. The experiments were performed using standard pulse sequences from the Bruker library that include COSY, HSQC, and HMBC, and the correlation maps were processed as usual.

Results and discussion

The ¹H HR-MAS NMR spectra, acquired directly from both leaves and cherries of *E. uniflora* (Supplementary Figure S2) in their natural swollen state, showed a remarkable singlet signal at 3.05 ppm, mainly in its leaves (Figure 1). This signal has been previously assigned to creatine while investigating some plant species (Iqbal et al., 2013; Flores et al., 2018; Ali et al., 2020). However, creatine is a common amino acid found in the kingdom Animalia, but not in kingdom Plantae. NMR singlet signals around 3.00 ppm are typical of N-CH₃ groups, which might have been the reason it was assigned to creatine.

In order to confirm the identity of the compound in an unambiguous way, two-dimensional NMR experiments such as one-bond and long-range ${}^{1}H{-}{}^{13}C$ and ${}^{1}H{-}{}^{1}H$ correlation maps from HSQC, HMBC, and COSY NMR experiments were acquired from D₂O extracts from both leaves and cherries of *E. uniflora* (Supplementary Figures S3–S5). In turn, the HSQC correlation map revealed that the hydrogen nuclei at 3.05 ppm is directly linked to a carbon at 46.2 ppm, that is typical from carbons connected to nitrogen nuclei. Moreover, the long-range ${}^{1}H{-}{}^{13}C$ correlation map from HMBC NMR experiment showed, for the hydrogen nuclei at 3.05 ppm, only two hydrogen-carbon correlations at 65.8 and 73.2 ppm, instead of three as usual, supporting the presence of a N-CH₃ group rather than a C-CH₃ group (Table 1; Supplementary Figures S3, S4).

The presence of hydrogen nuclei at 3.05 ppm connected to a carbon at 46.2 ppm is in accordance with the structure of the amino acid creatine or creatinine (Supplementary Figure S6), although the two adjacent carbons at 65.8 and 73.2 ppm do not support this hypothesis. The one-bond ^{1}H - ^{13}C correlation map showed that there

4-hydroxy-N-methyl proline			
Position	δ _C	δ _H (J in Hz)	HMBC ^a
N-CH ₃	46.2	3.05 s	2 and 5
2	73.2	4.22 dd (11.1 and 7.5)	3, N-CH ₃ and COOH
3	41.1	2.27 ddd (14.1, 11.1 and 4.9) 2.51 dd (14.1 and 7.5)	4 and 5
4	72.4	4.64 m	
5	65.8	3.20 ddd (13.0, 2.0 and 2.0) 3.97 dd (13.0 and 4.9)	4 and N-CH ₃
2-COOH	176.0		

TABLE 1 NMR data (400 MHz, D₂O extract) for 4-hydroxy-N-methyl proline.

^aLong-range, ¹H-¹³C correlations in the HMBC NMR, experiments were optimized for a coupling constant of 8 Hz, connecting hydrogen signals to the corresponding indicated carbon positions.

are two hydrogen nuclei linked to the carbon at 65.8 ppm at 3.20 and 3.97 ppm (i.e., a CH_2 group), while for the carbon at 73.2 ppm there is only one hydrogen nucleus connect to it at 4.22 ppm (i.e., CH group). On the other hand, the hydrogen nuclei at 3.20 ppm showed long-range ¹H-¹³C correlation with the carbons at 46.2 and 72.4 ppm, and the hydrogen nuclei at 4.22 ppm presented correlation with the carbons at 41.1, 46.2 and 176.0 ppm, revealing to be a more complex structure, which includes a carbonyl group. In addition, the one-bond ¹H-¹³C correlation map showed that there are two hydrogen nuclei at 2.27 and 2.51 ppm connected to the carbon at 41.1 ppm, and just one hydrogen nucleus at 4.64 ppm connected to the carbon 72.4 ppm, revealing additional CH₂ and CH groups in the structure (Table 1).

The comprehensive exploration of the ${}^{1}H{-}{}^{1}H$ correlation map from COSY NMR experiment, also allowed to establish and support a spin system consisting of the hydrogen nuclei at 4.22 (2), 2.27 and 2.51 (3), 4.64 (4), 3.20 and 3.97 ppm (5) (Figure 2; Supplementary Figure S5).

Therefore, one can conclude in an unequivocally way that, instead of creatine, the amino acid 4-hydroxy-*N*-methyl proline is the true compound behind the singlet signal at 3.05 in the ¹H NMR spectra of *E. uniflora* (Figures 1, 2; Supplementary Figure S6). The misassignment may have occurred due to the hydrogen nuclei, except those for the N-CH₃ group, present NMR signals in crowed regions of the spectra. In addition, they are less intense, as well as are present as multiplets, being hard to be observed.

The 4-hydroxy-*N*-methyl proline, one of the hydroxylated proline-derivatives, has been reported in plant species of several families including Euphorbiaceae, Fabaceae, Mytaceae, Meliaceae, Leguminosae, Asteraceae, Rutaceae, Loranthaceae, Annonaceae, Berberidaceae, Lauraceae, Sapindaceae, Rhamnaceae, Tamaricaceae, Sapotaceae (Blunden et al., 2004; Blunden et al., 2006; Jones et al., 2006; Giacometti et al., 2018), as well as in marine red algae (De Aquino et al., 2017). However, to the best of our knowledge, it has not been described in *E. uniflora* (Myrtaceae). Indeed, it was misassigned as creatine (Supplementary Figure S7).

This amino acid is a very important compound from a biological point of view, presenting pharmacological activities, mainly antiinflammatory properties. In fact, the anti-inflammatory properties of several plant species are attributed to 4-hydroxy-*N*-methyl proline (Sciuto et al., 1983; Jones et al., 2006; Ribeiro et al.,



2018). Therefore, its presence in *E. uniflora* as well as in *B. laurina* (Ali et al., 2020) supports a large number of its use in folk medicine. Furthermore, HR-MAS NMR analysis unveiled substantial amounts of 4-hydroxy-*N*-methyl proline in the leaves of *E. uniflora*. In contrast, the fruits exhibited comparatively very lower amounts of the compound (Supplementary Figure S2). This information holds meaningful value for its application in traditional medicine and the cosmetic industries.

HR-MAS NMR was also used to explore some changes in the chemical composition of the cherries during ripening process. The analysis unveiled an increase in the levels of α - and β -glucose as the fruits underwent the ripening process. Conversely, stable amounts of 4-hydroxy-*N*-methyl proline and malic acid were consistently observed in the cherries throughout the ripening stages (Supplementary Figure S8).

Conclusion

In conclusion, the investigation of *E. uniflora* leaves and cherries by means of HR-MAS NMR has brought some significant insights. The initial misassignment of the singlet signal at 3.02–3.05 ppm, assigned to creatine, has been corrected through comprehensive NMR analysis. Conversely to the initial hypothesis of creatine, the compound was unequivocally identified as 4-hydroxy-*N*-methyl

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proline. This amino acid, previously unrecognized in *E. uniflora*, holds biological significance, known for its pharmacological activities, particularly its anti-inflammatory properties. Its presence in significant amounts in the leaves of *E. uniflora*, as revealed by HR-MAS NMR, underscores its potential applications in traditional medicine and cosmetic industries. Furthermore, the investigation into the fruit's chemical composition indicated a consistency in 4hydroxy-*N*-methyl proline content during ripening process. These findings not only enhance our understanding of the chemical composition of *E. uniflora* but also shed light on its potential applications in medicinal and cosmetic contexts. The use of HR-MAS NMR proves to be a valuable tool in unraveling such intricate chemical composition, offering important insights into the plant's pharmacological properties.

Data availability statement

All NMR raw data are available free of charge in the UFPR public repository under the DOI number 10.5380/bdc/93, available at 10.5380/bdc/93.

Author contributions

LR: Data curation, Formal Analysis, Investigation, Writing-original draft, Writing-review and editing. SA: Data curation, Formal Analysis, Investigation, Writing-original draft. CD'O: Data curation, Investigation, Writing-review and editing, Validation. KS: Conceptualization, Data curation, Validation, Writing-original draft, Writing-review and editing. AB: Conceptualization, Methodology, Project administration, Resources, Supervision, Writing-original draft, Writing-review and editing, Data curation.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fntpr.2024.1360175/ full#supplementary-material

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