

ACIDIC N^{α} -ACYLARGININE DERIVATIVES IN ARGININE-ACCUMULATING PLANT TISSUES

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(Received 10 May 1982)

Key Word Index—*Lilium maximowiczii*, *Smilax china*, *Vicia faba*, *Rumex obtusifolius*, Liliaceae, Leguminosae, Polygonaceae, acylarginine derivatives, N^{α} -(2-hydroxy-2-carboxymethylsuccinyl)-L-arginine, N^{α} -(2-hydroxy-succinyl)arginine

Abstract—Two new acidic N^{α} -acylarginine derivatives were isolated from arginine-accumulating plant tissues. The first, N^{α} -(2-hydroxy-2-carboxymethylsuccinyl)-L-arginine, was isolated from bulbs of *Lilium maximowiczii*, whilst the second, N^{α} -(2-hydroxysuccinyl)arginine, was obtained from tubers of *Smilax china* and seeds of *Vicia faba*. No acidic N^{α} -acylarginine derivative was detected, however, in roots of *Rumex obtusifolius* which contained fairly large amounts of arginine and malonic acid.

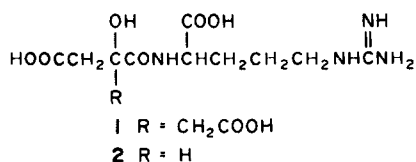
INTRODUCTION

Many plants utilize arginine as a main storage form of nitrogen [1]. It is considered that plant tissues accumulating a large amount of arginine overcome the high pI value of this amino acid and maintain the pH of the physiological fluid around neutral by forming an ionic and/or organic bond between the amino acid and acidic compounds. So far, however, only a few neutral or acidic derivatives of arginine such as γ -glutamylarginine [2] and arginosuccinic acid have been found in plant materials. This paper reports the isolation of two new acidic derivatives from arginine-accumulating plant tissues.

RESULTS AND DISCUSSION

Arginine was the most abundant amino acid in all four plant tissues studied (Table 1). The acidic amino acid fraction of three of them, bulbs of *L. maximowiczii*, tubers of *S. china* and seeds of *V. faba*, contained a compound which gave a positive reaction with Sakaguchi reagent, but did not react with ninhydrin. Each component was isolated using ion exchange chromatography and prep-PC.

The compound (1) from bulbs of *L. maximowiczii* was hydrolysed completely to L-arginine and citric acid (6 M



HCl, 120°, 3 hr). The L-arginine and citric acid were separated from each other by treatment with a Dowex 50(H^+) column, and the L-arginine identified by PC, HVE, FDMS, amino acid analysis and optical rotation. FDMS (m/z) 175 [$\text{M} + 1$]⁺ (100%), [α]_D²⁰ + 12.1° (H_2O , c 0.7). Lit. + 12.5° (H_2O , c 3.5) [3]. Citric acid was identified by PC, HVE and FDMS. FDMS (m/z) 193 [$\text{M} + 1$]⁺ (100%), 147 [$\text{M} - \text{COOH}$]⁺ (22). The ¹H NMR spectrum of 1 contained signals corresponding to arginyl and citryl residues. ¹H NMR (90 MHz in D_2O) δ 1.38–2.07 (4H, m, arginyl β - and γ -CH₂), 2.69 and 2.89 (1H each, both d, $J = 15$ Hz, citryl one CH₂), 2.75 and 2.93 (1H each, both d, $J = 15$ Hz, citryl another CH₂), 3.17 (2H, t, $J = 7$ Hz, arginyl δ -CH₂) and 4.34 (partially overlap with DOH, arginyl α -CH). FDMS of 1 gave a parent ion corresponding to arginine conjugated with citric acid, i.e. m/z 349 [$\text{M} + 1$]⁺ (100%). Since 1 gave a positive reaction with Sakaguchi reagent and did not react with

Table 1 The arginine content and the most abundant organic acids in some arginine-accumulating plant tissues

	Bulbs of <i>L. maximowiczii</i> (mg/100 g fr wt)	Tubers of <i>S. china</i> (mg/100 g fr wt)	Seeds of <i>V. faba</i> (mg/100 g dry wt)	Roots of <i>R. obtusifolius</i> (mg/100 g fr wt)
Arginine	333	63	192	46
Citric acid	645	nd*	19	+
Malic acid	nd*	258	41	+
Malonic acid	na†	na†	na†	100

*Not detected

†Not analysed

Table 2 Titration shift of the proton signals of the two methylene groups of the citrlyl residue in *N*^ε-(2-hydroxy-2-carboxymethylsuccinyl)arginine (1) and of the methylene and methyne groups of the malyl residue in *N*^ε-(2-hydroxysuccinyl)arginine (2)

	δ_{DSS}		Titration shift
	pD < 1*	pD > 13*	
Centre of one CH ₂ of the citrlyl residue in 1	2.95	2.63	0.32
Centre of the second CH ₂ of the citrlyl residue in 1	2.90	2.59	0.31
Centre of the CH ₂ of the malyl residue in 2	2.86	2.54	0.32
CH of the malyl residue in 2	4.56	4.48	0.08

*pD was adjusted by adding ca 37% DCl soln in D₂O or ca 40% NaOD soln in D₂O

ninhydrin, the α -amino group of arginine was acylated. Measurement of the ¹H NMR spectra of 1 in acidic and basic medium showed that the β -carboxyl group of citric acid formed an amide bond with the α -amino group of arginine. Both of the CH₂ groups of the citrlyl residue revealed a big titration shift when the acidic medium was rendered basic by addition of 40% NaOD in D₂O (Table 2). The structure of 1 was thus *N*^ε-(2-hydroxy-2-carboxymethylsuccinyl)-L-arginine.

The compound (2) from tubers of *S. china* was hydrolysed to arginine and malic acid (3 M HCl, 110°, 3 hr). The arginine and malic acid were separated from each other by treatment with a Dowex 50 (H⁺) column, and the arginine identified by PC, HVE and amino acid analysis. Malic acid was identified by HVE and GC/MS after esterification with diethyl ether-diazomethane. MS (*m/z*) 163 [*M* + 1]⁺ (1%), 103 [*M* - COOMe]⁺ (100%). The ¹H NMR spectrum of 2 contained signals corresponding to arginyl and malyl residues. ¹H NMR (100 MHz in D₂O) δ 1.40–2.00 (4H, *m*, arginyl β - and γ -CH₂), 2.63 and 2.77 (1H, *dd*, *J* = 7, 16 Hz and 1H, *dd*, *J* = 4, 16 Hz, respectively, malyl CH₂), 3.19 (2H, *t*, *J* = 6 Hz, arginyl δ -CH₂), 4.25 (1H, *dd*, *J* = 5, 7 Hz, arginyl α -CH) and 4.46 (1H, *dd*, *J* = 4, 7 Hz, malyl CH). FDMS of 2 gave a parent ion corresponding to arginine conjugated with malic acid, i.e. *m/z* 581 [2*M* + 1]⁺ (3%), 291 [*M* + 1]⁺ (100%), 273 [291 - H₂O]⁺ (72%). Since 2 showed a positive reaction with Sakaguchi reagent and did not react with ninhydrin, the α -amino group of arginine was acylated as in compound 1. It was shown by measuring the titration shift in the ¹H NMR spectrum that the α -carboxyl group of malic acid formed an amide bond with the α -amino group of arginine. Malyl CH₂ protons showed a big upfield shift when an acidic medium was turned basic by addition of 40% NaOD in D₂O, while the upfield shift of a malyl CH proton was slight (Table 2). The structure of 2 was established as *N*^ε-(2-hydroxysuccinyl)arginine. The configurations of the two chiral centres of 2 were not determined.

A compound isolated from the acidic amino acid fraction of mature seeds of *V. faba* which gave a positive reaction with Sakaguchi reagent and did not react with ninhydrin showed the same behaviour on PC and HVE as 2. ¹H NMR and FDMS spectra of the substance isolated from *V. faba* seeds were also consistent with those of 2. FDMS (*m/z*) 291 [*M* + 1]⁺ (100%). The compound gave arginine and malic acid by hydrolysis (3 M HCl, 130°, 3 hr) which were identified by amino acid analysis and GC after diethyl ether-diazomethane treatment, respectively.

The compound isolated from *V. faba* seeds was thus identified as 2.

In bulbs of *L. maximowiczii* and tubers of *S. china*, the content of organic acid is fairly high and the most abundant organic acids are citric acid and malic acid, respectively (Table 1). Therefore, 1 and 2 are the amides which are formed between the most abundant amino acid and organic acid in bulbs of *L. maximowiczii* and tubers of *S. china*, respectively. The content of organic acids is relatively low in seeds of *V. faba*. The difference in amount of malic acid and citric acid is not big (Table 2), but 1 could not be detected in seeds of *V. faba*. There was no spot which gave a positive reaction with Sakaguchi reagent and did not react with ninhydrin on PC and HVE of the acidic amino acid fraction from roots of *R. obtusifolius*, although the tissue contained plenty of arginine and organic acids, especially malonic acid (Table 1). Amino acids and peptides in seeds of *V. faba* [4, 5] and organic acids and peptides in *R. obtusifolius* have been reported [6, 7]. A survey of ninhydrin positive substances in the acidic amino acid fraction of *L. maximowiczii* and *S. china* is now in progress.

EXPERIMENTAL

General methods. PC: *n*-BuOH-HOAc-H₂O (4:1:2) (solvent 1), PhOH-H₂O-conc. NH₄OH (120:30:1, w/v/v) (solvent 2), HVE: pH 6.5 (pyridine-HOAc-H₂O, 25:1:500, 100 V/cm).

Plant materials. Bulbs of *L. maximowiczii* and mature seeds of *V. faba* were purchased at a local market. Tubers of *S. china* were collected in natural habitats on Shikoku Island, Japan, by Dr Tahara, Hokkaido University. Roots of *R. obtusifolius* were from the experimental farm of Hokkaido University.

Quantitative analysis of organic acids and amino acids. An aliquot of the effluent from a column of Amberlite IR-120 (H⁺) loaded with the plant extract was concd, dissolved in MeOH, esterified with excess Et₂O-CH₂N₂ and analysed by GC using tridecane as int. reference according to the method of ref. [8] with slight modification. GC conditions: FID, 5% Reoplex 400 (2 m × 3 mm, glass column), N₂ 60 ml/min, temp. programme 80–210° at 5°/min, injector temp. 240°, detector temp. 255°. Amino acids were analysed with an amino acid analyser.

Isolation of 1 from bulbs of *L. maximowiczii* and hydrolysis. Bulbs of *L. maximowiczii* (70 kg) were homogenized with 80% MeOH (20:1). The residue was placed in a column and extracted continuously with 70% MeOH (32:1). The extracts were combined, concd and dissolved in H₂O. After centrifugation, the supernatant was applied to a column of Amberlite IR-120.

(H⁺, 2 l), which was thoroughly washed with H₂O. The amino acid fraction was eluted with 2 M NH₄OH. Fractions of 1 l each were collected. Fractions 3–5 were concd and applied to a column of Dowex 1 × 4 (AcO⁻, 500 ml). After the neutral and basic fraction was washed out with H₂O, acidic amino acids were eluted with 2 M HOAc. Fractions of 500 ml each were collected. Fractions 4 and 5 were concd, applied to a column of Sephadex G-10 (450 ml) and developed with H₂O. Fractions of 15 ml each were collected. Compound 1 (85 mg) was isolated as a colourless evaporation residue from fractions 11–20 by prep. PC (solvent 1) followed by Dowex 50 (H⁺) treatment. Nearly the same amount of 1 was also contained in fraction 3 from Dowex 1 × 4 (AcO⁻) as in fractions 4 and 5 judging from the intensity of the spot of 1 on PC, but fraction 3 was lost by accident. The amount of 1, therefore, should be ca 150 mg/7 kg fr. wt. See Results and Discussion for ¹H NMR and FDMS spectra of 1. Hydrolysate of 1 (6 M HCl, 120°, 3 hr) was concd and treated with a column of Dowex 50 (H⁺). Citric acid was identified in the effluent and H₂O wash from the column by PC, HVE and FDMS. See Results and Discussion for FDMS spectrum. L-Arginine was identified in the 2 M NH₄OH eluate from the column by PC, HVE and amino acid analysis, FDMS and optical rotation. See Results and Discussion for FDMS spectrum and $[\alpha]_D$ value.

Isolation of 2 from tubers of *S. china* and hydrolysis Tubers of *S. china* (570 g) were sliced and extracted with 70% EtOH (9 l). The residue was extracted (twice) with 70% EtOH (5 l each). The extracts were combined, concd and dissolved in H₂O. After filtration, the filtrate was applied to a column of Amberlite IR-120 (H⁺, 250 ml), which was thoroughly washed with H₂O. The amino acid fraction was eluted with 2 M NH₄OH and fractions of 500 ml each were collected, except for fraction 1 which was 200 ml. Fractions 2–6 were concd and applied to a column of Dowex 1 × 4 (AcO⁻, 100 ml). After the neutral and basic fractions had been washed out with H₂O, acidic amino acids were eluted with 2 M HOAc. Fractions of 200 ml each were collected. Fraction 1 was concd and applied to a column of Dowex 1 × 8 (AcO⁻, 5 ml) and eluted with 0.2 M HOAc. Fractions of 5 ml each were collected. Compound 2 (49 mg) was obtained as a colourless evaporation residue from fractions 3–6 by prep. PC (solvent 2) followed by Dowex 50 (H⁺) treatment. See Results and Discussion for ¹H NMR and FDMS spectra of 2. The hydrolysate of 2 (3 M HCl, 110°, 3 hr) was concd and treated with a Dowex 50 (H⁺) column. Effluent and H₂O wash from the column was concd, esterified with excess Et₂O–CH₂N₂ and analysed for organic acid by GC/MS, 5% Peg-20 M (1 m × 2 mm) He 30 ml/min, temp. programme 110–150° at 8°/min, injector and jet separator at 230° and 260°, respectively, MS 24 eV. Dimethyl malate was eluted at 2.4 min. See Results and Discussion for MS. Arginine was identified in 2 M NH₄OH eluate from the column of Dowex 50 (H⁺) loaded with hydrolysate of 2 by PC, HVE and amino acid analysis.

Isolation of 2 from seeds of *V. faba* and hydrolysis Meal of *V. faba* seeds (1.3 kg) was extracted with 70% EtOH (10 l). The residue was extracted (twice) with 70% EtOH (9 and 5 l). The extracts were combined, concd and dissolved in H₂O. After centrifugation, the supernatant was applied to a column of Amberlite IR-120 (H⁺, 1 l), which was thoroughly washed with H₂O. The amino acid fraction was eluted with 2 M NH₄OH. Fractions of 1 l each were collected. Fractions 2–4 were concd and applied to a column of Dowex 1 × 4 (AcO⁻, 50 ml). After neutral and basic fractions were washed out with H₂O, acidic amino acids were eluted with 2 M HOAc. Fractions of 50 ml each were collected. Fractions 2–7 were combined, put on a column of

Sephadex G-10 (450 ml) and developed with H₂O. Fractions of 10 ml each were collected. Fractions 18–23 were concd, applied again to a column of Dowex 1 × 4 (AcO⁻, 10 ml) and eluted with 0.2 M HOAc. Fractions of 5 ml each were collected. Fractions 8–13 were concd, put on a column of cellulose powder (600 ml) and developed with solvent 1. Fractions of 15 ml each were collected. Compound 2 (19 mg) isolated as colourless evaporation residue from fractions 36–41 by prep. PC (solvent 1) followed by Dowex 50 (H⁺) treatment had the same behavior on PC and HVE as those of 2. Its ¹H NMR and FDMS spectra were also consistent with those of 2. See Results and Discussion for FDMS spectrum. The hydrolysate of the isolated material (3 M HCl, 130°, 3 hr) was concd and treated with a Dowex 50 (H⁺) column. Effluent from the column was concd, esterified with excess Et₂O–CH₂N₂ and analysed for organic acid by GC (by the method described above). A peak corresponding to dimethyl malate was detected (16 min). The presence of malic acid in the effluent from the Dowex 50 (H⁺) column was confirmed also by PC and HVE. Arginine was identified in the 2 M NH₄OH eluate from the Dowex 50 (H⁺) column by an amino acid analysis.

Survey of acidic arginine derivative in roots of *R. obtusifolius* Roots (1.9 kg) of *R. obtusifolius* were sliced and extracted with 70% EtOH (19 l). The residue was homogenized with 70% EtOH (19 l). The combined extracts were concd and centrifuged. The supernatant was applied to a column of Amberlite IR-120 (H⁺, 500 ml), which was thoroughly washed with H₂O. The amino acid fraction was eluted with 2 M NH₄OH and fractions of 1 l were collected. Fractions 1–7 were concd and applied to a column of Amberlite IR-120 (NH₄⁺, 500 ml), which was washed with H₂O until the wash gave a negative ninhydrin reaction. The effluent and H₂O wash was concd and applied to a column of Dowex 1 × 8 (AcO⁻, 50 ml). After neutral amino acids were washed out with H₂O, the acidic amino acid fraction was eluted with 2 M HOAc (1 l) and 8 M HOAc (200 ml). No spot giving a positive reaction to Sakaguchi reagent and a negative reaction to ninhydrin was detected on PC and HVE of the acidic amino acid fraction.

Acknowledgements—We express our gratitude to Dr S. Tahara and Mr K. Watanabe, Department of Agricultural Chemistry, Hokkaido University, for gifts of tubers of *Smilax china* and measurements of GC/MS and FDMS, respectively.

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