

Identification of Endogenous Gibberellins by Feeding of [¹⁴C] GA₁₂ in Chinese Yam, *Dioscorea opposita*

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ABSTRACT: The metabolism of [¹⁴C] GA₁₂ in the Chinese yam (*Dioscorea opposita* Thunb. var. Tsukune) was examined to determine the identification of endogenous gibberellins. [¹⁴C] GA₁₂ was metabolized to GA₅₃, GA₄₄, GA₁₉, GA₂₀, GA₁, GA₈, GA₁₅, GA₂₄, GA₉, GA₃₆ and GA₄. Radioactivity of GAs in non C-13 hydroxylation route was five-fold higher than that of early C-13 hydroxylation in analyzed GA-metabolites. Radioactivity of GA₄ was always four times higher than that of GA₁ at every feeding time. GA₁ radioactivity has always a lower level to below 200 DPM. The major pathway of endogenous GA metabolism in seedlings of the Chinese yam might be the non C-13 hydroxylation pathway.

Keywords: Chinese yam, *Dioscorea opposita*, gibberellins, identification.

Gibberellins (GAs) play an important physiological role in the Chinese yam (*Dioscorea opposita* Thunb.) plants. They affect seed germination, tuber dormancy (Okagami and Tanno, 1993), and tuber enlargement (Kim et al., 2003). In particular, the application of gibberellins strengthens or deepens the dormancy of the tubers and bulbils in Chinese yams (Okagami and Nagao, 1971). Furthermore, bulbil dormancy in *Begonia evansiana* (Okagami 1972) and the genus *Dioscorea* (Okagami and Tanno, 1977) is also induced by endogenous and exogenous gibberellins. GA₁₂ is an early precursor in the GA biosynthesis and its biosynthesis pathway is important to elucidate the physiological role named to gibberellin induced dormancy in Chinese yam'. If the bioactive GAs, GA₁ or GA₄, may be involved in the controlling the dormancy and sprouting of tubers and bulbils in the Chinese yam, several GA biosynthesis inhibitors such as ancymidol, mepiquat chloride and trinexapac-ethyl will be applied to the Chinese yams for modulating the sprouting and dormancy periods.

In general, the predominant GA biosynthesis pathway in

most vegetative plant tissues seems to be the early C-13 hydroxylation route (Sponsel, 1995), meanwhile intermediates of other pathways, especially the non C-13 hydroxylation pathway, are often also present (Zhu and Davies, 1991). Till now, several endogenous gibberellins have been identified in dormant bulbils of Chinese yam (*Dioscorea opposita* Thunb.) (Tanno *et al.*, 1992). However, it was not found that which GA biosynthetic pathway is operated and also dominant, GA biosynthesis pathway has not been understood in the Chinese yam plants.

Thus, this study focuses principally on the identification of endogenous gibberellins and biosynthesis pathways active in vegetative Chinese yam, as determined by the identification of endogenous GA-metabolites following the feeding of [¹⁴C]GA₁₂ as a precursor of gibberellin in the Chinese yam seedlings.

MATERIALS AND METHODS

Plant material and feeding of [¹⁴C]gibberellin A₁₂

Sprouted tubers of the Chinese yam (*Dioscorea opposita* Thunb. var. Tsukune) were grown in a sand-vermiculite mixture. After the seedlings had fully expanded leaves and grown for 4 weeks, and then they were fed to determine the GA biosynthesis pathway. The substrate of [¹⁴C]GA₁₂ was purchased from L. N. Mander (Australian National University, Canberra, Australia). The compound was about 96% radioactive. Specific activity was 10 μCi mmol⁻¹. The substrate was dissolved in EtOH:water (1:1, v/v). 10 μl (370 Bq) for each twenty plant was fed to shoot tips and surface of whole leaves below apex of the plants. The application of [¹⁴C]GA₁₂ was performed by injection using a microsyringe. After fed, seedlings were grown at 25°C in a constant temperature room with continuous light at 255 μmolm⁻²s⁻¹ provided by fluorescent lamps. Seedlings were harvested at 6, 12, 18 and 24 h after feeding (HAF). Seedlings were frozen in liquid nitrogen immediately after harvesting and stored at -70°C for analysis of GAs.

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<Received February 7, 2003>

Extraction of endogenous gibberellins

Extraction and HPLC of GA-metabolites followed the reference (Lee *et al.*, 1998). Lyophilized tissue samples were ground to a fine powder in a mortar and pestle with the aid of acid-washed sea sand. The powdered tissue was extracted with 80% (v/v) methanol (MeOH). The 80% MeOH was removed by filtration, and the tissue was then extracted with 100% MeOH until the extract was clear. The volume of the 80% and 100% extracts were recorded, the two extracts were combined, and water was added to bring the combined MeOH extract concentration to 60%. This solution was chilled for 1 h at -70°C, and precipitated chlorophyll was removed by filtration through a GF/A filter. The extract was adjusted to pH 8.0-8.3 using 2N NH₄OH and passed through a 3 g column of Davisil C₁₈ (90-130 µm, 60Å pore size, Alltech). The eluant was reduced to near dryness at 40°C *in vacuo*. The sample was dried onto 1 g celite and then loaded onto a 4 g SiO₂ (ICN Silica 32-100, active 60Å) partitioning column (deactivated with 20% water) to separate the GAs as a group from more polar impurities. GAs were eluted with 80 ml of 95:5 ethyl acetate (EtOAc): hexane saturated with formic acid. This solution was dried at 40°C *in vacuo*, redissolved in 4 ml EtOAc and partitioned 3 times against 4 ml of 0.1 M phosphate buffer (pH 8.0). Dropwise addition of 2N NaOH was required during the first partitioning to neutralize residual formic acid. Polyvinylpyrrolidone (PVPP) 1 g was added to the combined aqueous phases, and this mixture was slurried for 1 h. Following the removal of the PVPP by filtration, 6N HCl was added to reduce the pH 2.5. The extract was partitioned 3 times against equal volumes of EtOAc. The combined EtOAc fraction was dried *in vacuo*, and the residue was dissolved in 3 ml of 100% MeOH. This solution was dried on a Savant Automatic Environmental Speedvac (Model AES 2000). The dried sample was subjected to reverse-phase C₁₈-HPLC.

HPLC condition of endogenous gibberellins

The GAs were chromatographed on a 3.9×300 mm µ-Bonda Pak C₁₈ column (Waters) and eluted at 1.5 ml min⁻¹ with the following gradient: 0 to 5 min, isocratic 28% MeOH in 1% aqueous acetic acid; 5 to 35 min, linear gradient from 28 to 86% MeOH; 35 to 36 min, 86 to 100% MeOH; 36 to 40 min, isocratic 100% MeOH. Up to fifty fractions of 1.5 ml each were collected. Small aliquots (15 µl) from each fraction were taken, and radioactivity was measured with liquid scintillation spectrometry (Beckman, LS 1801) to determine radioactivity and accurate retention times of each GA based upon the elution of ³H-GA standards.

Identification of endogenous gibberellins

Radioactivity for each concentrated GA fraction was counted for 15 min in liquid scintillation counter. Each GA fraction was redissolved in 100% methanol, transferred to a 1 ml vial and dried under N₂ at 40°C. The sample was dissolved in 35 µl of methanol, and the GA methyl ester was prepared with ethereal diazomethane. The sample was dried under N₂, redissolved in methanol and methylated one more time. The sample was dissolved in 35 µl pyridine, and silylated for 30 min at 65°C with the same amount of N, O-Bis (trimethylsilyl)-trifluoroacetamide (BSTFA) with 1% TMCS (Pierce Chemical Co.). The sample was then reduced to dryness with N₂ and solubilized in anhydrous dichloromethane. 1 µl of each sample was injected on-column on a 30 m×0.25 mm (i.d.), 0.25 µm film thickness DB-1 capillary column (J & W Co.). The GC (Finnigan Mat GCQ) oven temperature was programmed for a 1 min hold at 60°C, then to rise at 15°C min⁻¹ to 200°C followed by 5°C min⁻¹ to 285°C. Helium carrier gas was maintained at a head pressure of 30 kPa. The GC was directly interfaced to a Mass Selective Detector with an interface and source temperature of 280°C, an ionising voltage of 70 eV and a dwell time of 100 ms. Full scan mode (the first trial) and three major ions of the supplemented [²H₂]GA internal standards (the second trial) and the endogenous gibberellins were monitored simultaneously. Retention time was determined by using the hydrocarbon standards (C₂₃, C₂₄, C₂₅, C₂₆, C₂₇ and C₂₈) to calculate the KRI (Kovats retention indices) value.

RESULTS AND DISCUSSION

To date, several endogenous GAs have been identified in the bulbils of the Chinese yam (Tanno *et al.*, 1992). Many metabolic products were formed following the feeding of [¹⁴C]gibberellin A₁₂ as the primary gibberellin precursor. To check authentic GA-metabolites in seedlings of Chinese yam, the deuterated GAs, KRI values, retention times and parent ion from the each HPLC fraction of seedlings were evaluated by GC-MS (Table 1).

Radioactivity of GA-metabolites, GA₅₃, GA₄₄, GA₁₉, GA₂₀, GA₁, GA₁₅, GA₂₄, GA₉, GA₃₆ and GA₄ was also showed in Table 3 and 4. [¹⁴C] gibberellin A₁₂ as substrate was rapidly metabolized for 18h after feeding to seedlings of the Chinese yam, otherwise, [¹⁴C] gibberellin A₁₂ was slowly metabolized at 18 h to 24 h in time-course feeding (Table 2).

Table 3 shows the radioactivity of endogenous gibberellins for an early C-13 hydroxylation (ECH) pathway. In the radioactivity of an early C-13 hydroxylation route, GA₅₃ was one of the first compound committed to the GA pathway. After 6 HAF, GA₁₉ was metabolized sharply, and then,

Table 1. HPLC fractions, KRI, relative ion intensity from acidic ethyl acetate fractions of gibberellins in seedlings of the Chinese yam.

Fraction no.	GAs	KRI ^a	Source	<i>m/z</i> (% relative intensity of base peak) ^b				
6-8	GA ₈	2818	¹² C	594(100)	448(25)	379(20)	375(15)	238(28)
		2818	¹⁴ C	596(100)	450(24)	381(21)	375(11)	240(26)
12-14	GA ₁	2674	¹² C	506(100)	448(20)	313(17)	491(13)	377(12)
		2674	¹⁴ C	508(100)	450(19)	315(14)	493(11)	379(13)
24,25	GA ₂₀	2485	¹² C	418(100)	375(45)	403(14)	359(12)	301(13)
		2485	¹⁴ C	420(100)	377(45)	405(13)	361(10)	303(11)
26-28	GA ₄₄	2789	¹² C	432 (63)	238(41)	417(12)	373(17)	207(100)
		2789	¹⁴ C	434 (62)	240(39)	419(10)	375(16)	209(100)
26-28	GA ₃₆	2600	¹² C	284(100)	430(58)	312(47)	462(11)	402(38)
		2600	¹⁴ C	286(100)	432(56)	314(45)	464(10)	404(40)
29-31	GA ₁₉	2600	¹² C	434(100)	374(59)	402(41)	462(10)	375(57)
		2600	¹⁴ C	436(100)	376(57)	404(40)	464(9)	377(55)
32,33	GA ₄	2506	¹² C	284(100)	225(80)	289(70)	224(76)	418(26)
		2506	¹⁴ C	286(100)	227(76)	291(71)	226(75)	420(23)
34-36	GA ₂₄	2444	¹² C	314(100)	226(89)	286(77)	342(42)	374(4)
		2444	¹⁴ C	316(100)	228(87)	288(75)	344(40)	376(3)
37,38	GA ₅₃	2450	¹² C	448 (47)	251(30)	235(30)	389(25)	241(18)
		2450	¹⁴ C	450 (47)	253(28)	237(28)	391(25)	243(19)
37,38	GA ₉	2305	¹² C	298(100)	270(78)	227(48)	243(43)	330(6)
		2305	¹⁴ C	300(100)	272(77)	229(48)	245(42)	332(6)
39,40	GA ₁₅	2608	¹² C	239(100)	284(50)	344(23)	312(22)	298(13)
		2608	¹⁴ C	241(100)	286(48)	346(19)	314(17)	300(11)
42-44	GA ₁₂	2335	¹² C	300(100)	240(31)	328(31)	360(2)	285(19)
		2335	¹⁴ C	302(100)	242(32)	330(29)	362(2)	287(20)

^aKRI, Kovats retention indices. ^bIdentified as methyl ester trimethylsilyl ether derivatives by comparison with reference spectra and KRI data (Gaskin and MacMillan, 1991). Gibberellin is quantified with comparisons of peak area ratio of prominent ions.

Table 2. Reductive radioactivity of [¹⁴C] gibberellin A₁₂ in time-course feeding in seedlings of the Chinese yam.

Time after feeding (h)	Radioactivity (DPM)
6	11,395 800
12	8,945 500
18	6,248 450
24	4,955 490

DPM: disintegration per 10 min. Data are means values of two replicates SE.

GA₈, GA₄₄, GA₅₃ and GA₂₀ were metabolized respectively. Radioactivity of GA₄₄ including GA₅₃, GA₁₉ and GA₂₀ was decreased greatly at 12 HAF after feeding except for GA₈. At 18 HAF, radioactivity of GA₁₉, GA₄₄ and GA₅₃ was increased dramatically, however, radioactivity of GA₂₀ was smaller than that of 12 HAF. At 24h as end feeding, GA₁₉ was only increased dominantly and radioactivity of GA₅₃, GA₄₄ and GA₈ was decreased compared to 18 HAF, GA₁₉ was the highest GA metabolite at 6 HAF, meanwhile, GA₂₀ as precursor of bioactive GA₁ was the lowest among GA

Table 3. Radioactivity of endogenous gibberellins for an early C-13 hydroxylation in time-course feeding of [¹⁴C] gibberellin A₁₂ in the seedlings of Chinese yam.

GAs	Time after feeding (h)			
	6	12	18	24
	-----DPM-----			
GA ₅₃	184±21	146±7	205±14	184±9
GA ₄₄	233±9	130±11	270±23	197±14
GA ₁₉	361±11	187±15	379±20	439±11
GA ₂₀	147±15	138±18	127±16	159±26
GA ₁	169±8	182±22	208±11	188±18
GA ₈	255±24	266±10	301±25	257±23

DPM: disintegration per 10 min. Data are means values of two replicates±SE.

members of the ECH. Radioactivity of GA₂₀ was always lowest in time-course feeding except for 12 HAF. Radioactivity of GA₁₉ was linearly increased after 12 HAF suggesting one of the most abundant compounds in ECH route. The most abundant GA-metabolite in ECH route might be GA₁₉-

Table 4. Radioactivity of endogenous gibberellins for non C-13 hydroxylation in time-course feeding of [¹⁴C] gibberellin A₁₂ in the seedlings of Chinese yam.

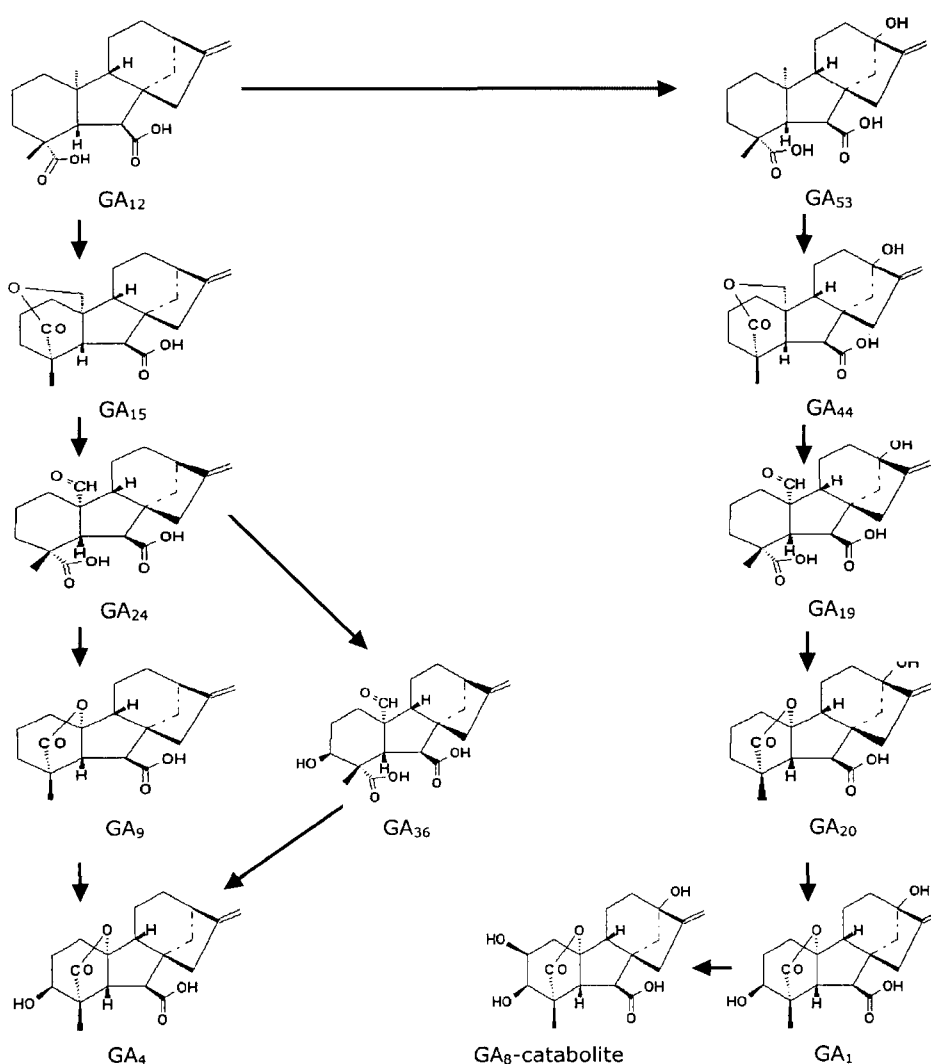
GAs	Time after feeding (h)			
	6	12	18	24
	-----DPM-----			
GA ₁₅	292±33	227±13	286±20	377±11
GA ₂₄	727±27	769±21	1,614±54	2,118±46
GA ₉	606±19	221±11	812±28	378±13
GA ₃₆	271±4	229±17	357±18	511±25
GA ₄	463±21	353±5	535±26	612±10

DPM: disintegration per 10 min. Data are means values of two replicates±SE.

Table 5. Radioactivity of endogenous gibberellins for an early C-13 hydroxylation (ECH), non C-13 hydroxylation (NCH) and total endogenous gibberellins (TGA) in time-course feeding of [¹⁴C] gibberellin A₁₂ in the seedlings of Chinese yam.

GA groups	Time after feeding (h)			
	6	12	18	24
	-----DPM-----			
ECH	1,349	1,049	1,490	1,424
NCH	2,359	1,799	3,604	3,996
TGA	3,708	2,848	5,094	5,420

DPM: disintegration per 10 min. Data for ECH, NCH and TGA are aggregated from Table 3 and 4.

**Fig. 1.** Non C-13 (left) hydroxylation and early C-13 hydroxylation (right) routes as the two GA biosynthesis pathways operated in the Chinese yam plants.

metabolite. Radioactivity of GA₈ as catabolite of GA₁ was always higher than that of GA₁. Radioactivity in GA mem-

bers of ECH route was below 450 DPM. This was consistent with GA metabolism in vegetative tissues of most other

plant species (Sponsel, 1995), including another member of the Solanaceae, *Lycopersicon esculentum* (Bohner *et al.*, 1988).

Table 4 shows the radioactivity of endogenous gibberellins for non C-13 hydroxylation (NCH) pathway. In radioactivity of NCH, radioactivity of GA members except for GA₂₄ was lowest at 12 HAF, in addition, radioactivity of GA members, GA₁₅, GA₉, GA₃₆ and GA₄ was always lower than that of GA₂₄ in all time-course feeding. Radioactivity of GA₂₄ was not changed at 6 and 12 HAF, however, GA₂₄ was highly increased at 18 and 24 HAF. GA₁₅ and GA₃₆ were smaller radioactivity than GA₂₄ and GA₉ except for 24 HAF. In addition, radioactivity of GA₉ was always higher than that of GA₃₆.

We also compared the change of bioactive GAs, GA₁ and GA₄ during time-course feeding and also compared GA members of each ECH and NCH and total GAs in time-course feeding (Table 3, 4 and 5). Radioactivity of GAs in NCH group was five-fold higher than that of ECH in analyzed GA-metabolites. It was a major difference between the metabolites of two GA metabolic pathways obtained by feeding of [¹⁴C] gibberellin A₁₂. In comparing the bioactive GAs, GA₄ and GA₁, radioactivity of GA₄ was always four times higher than that of GA₁ at every feeding time. GA₁ radioactivity has always a lower level to below 200 DPM.

Thus, we concluded that otherwise most other species which have been studied, the major pathway of endogenous GA metabolism in seedlings of the Chinese yam is the non C-13 hydroxylation pathway, and the early C-13 hydroxylation pathway is also presented (Fig. 1). Furthermore, detection of most of the members of the early C-13 hydroxylation (ECH) route, GA₅₃, GA₄₄, GA₁₉, GA₂₀, GA₁ and GA₈, and the member of the non C-13 hydroxylation (NCH) route, GA₁₅, GA₂₄, GA₉, GA₃₆ and GA₄, strongly suggested that two GA metabolic pathways active in the Chinese yam.

ACKNOWLEDGEMENTS

This work supported by a grant from the Agricultural R&D Promotion Center (ARPC 2001). We are also grateful

to Prof. Lewis N. Mander (Australian National University, Research School of Chemistry, Canberra, Australia) for providing labeled gibberellins.

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