



# 28-Norcastasterone is biosynthesized from castasterone

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## Abstract

Metabolic experiments with deuterium-labeled castasterone in seedlings of *Arabidopsis thaliana*, *Oryza sativa* and *Lycopersicon esculentum*, and cultured cells of *Catharanthus roseus* were performed, and the metabolites were analyzed by GC–MS. In all the plant species examined, [<sup>2</sup>H<sub>3</sub>]28-norcastasterone was identified as a metabolite of [26,28-<sup>2</sup>H<sub>6</sub>]castasterone, indicating that castasterone is the biosynthetic origin of 28-norcastasterone. Moreover, the natural occurrence of 28-norcastasterone and 28-nortyphasterol in seedlings of *A. thaliana* has been demonstrated. This is the first report of the natural occurrence of 28-nortyphasterol in plants. © 2000 Elsevier Science Ltd. All rights reserved.

**Keywords:** *Arabidopsis thaliana*; *Catharanthus roseus*; Apocynaceae; Cruciferae; Biosynthesis; Cultured cells; Biosynthetic intermediates of brassinolide; Castasterone; 28-Norcastasterone; 28-Nortyphasterol

## 1. Introduction

Brassinosteroids (BRs) are now recognized as an important hormone controlling plant growth and development (Altmann, 1998; Clouse and Feldmann, 1999). Naturally occurring BRs so far identified are C<sub>27</sub>-, C<sub>28</sub>-, and C<sub>29</sub>-steroids (Fujioka, 1999). The carbon skeletons of the side chain are common in plant sterols, suggesting that BRs are biosynthesized from the corresponding plant sterols. Brassinolide belonging to C<sub>28</sub>-steroids is highly bioactive and is widely distributed in the plant kingdom. A biosynthetic pathway for brassinolide has been elucidated by feeding labeled compounds of possible brassinolide intermediates to cultured cells of *Catharanthus roseus*, followed by analyzing the metabolites produced by GC–MS. The present model includes parallel branched pathways, namely, early and late C-6 oxidation pathways (Fujioka and Sakurai, 1997a,b; Sakurai, 1999; Fujioka et al., 2000). By analogy with the biosynthesis of brassinolide, it is postulated that C<sub>27</sub>-BRs and C<sub>29</sub>-BRs could be biosynthesized from the corresponding C<sub>27</sub>-sterols and C<sub>29</sub>-sterols, respectively. Therefore, it is likely that 28-norcastasterone is synthe-

sized from cholesterol, because 28-norcastasterone and cholesterol have the same structural skeleton. However, no experimental evidence has been obtained relating to biosynthesis of 28-norcastasterone. To date, 28-norcastasterone has been found in as many as twelve plant species (Fujioka, 1999). Thus, it would be important to clarify the biosynthesis of 28-norcastasterone. Unexpectedly, during our study of BR biosynthesis, we found that castasterone was converted to 28-norcastasterone. Here, we report the evidence for castasterone being the biosynthetic precursor of 28-norcastasterone. In addition, we describe the identification of 28-norcastasterone and 28-nortyphasterol in seedlings of *Arabidopsis thaliana*, and their biological activity.

## 2. Results and discussion

### 2.1. Metabolism of [<sup>2</sup>H<sub>6</sub>]castasterone in seedlings of *Arabidopsis thaliana*

[26,28-<sup>2</sup>H<sub>6</sub>]Castasterone was applied to seedlings of *A. thaliana* (wild type: Ws-2). After incubation, the plant was extracted, and the metabolites of interest purified using both a silica gel column and an ODS cartridge column, repeatedly followed an ODS-HPLC. HPLC-

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purified fractions were analyzed by gas chromatography–mass spectrometry (GC–MS) after conversion to the methanoboronate derivative. A prominent peak of a metabolite [GC retention time (Rt): 11.58 min] was found in the HPLC fraction (Rt: 6–8 min). Its mass spectral data was as follows:  $m/z$  501 ( $M^+$ , 92), 486 (11), 459 (7), 399 (7), 358 (20), 287 (78), 144 (100). The mass spectral data was very similar to that of authentic 28-norcastasterone bismethanoboronate (28-norcastasterone bismethanoboronate; Rt: 11.60 min,  $m/z$  498 ( $M^+$ , 94), 483 (13), 456 (8), 399 (9), 358 (24), 287 (82), 141 (100)). A clear difference in the mass spectra of the metabolite and authentic 28-norcastasterone, however, was observed at  $m/z$  501 and 498, and 144 and 141. This difference must be derived from the side chain. As we used  $[26,28-^2H_6]$ castasterone as a substrate, we expected the corresponding  $[^2H_6]$ metabolite, if the C26 and C28 methyl groups were not involved in the metabolism. Unexpectedly, we observed a molecular ion of the metabolite at  $m/z$  501. This suggested the removal of either the 26-methyl or 28-methyl group. From the similarity of the mass spectra between the metabolite and 28-norcastasterone, it was concluded that the 28-methyl group was removed during its metabolism. Another possible candidate for the metabolite,  $[28-^2H_3]$ 28-norcastasterone was excluded, because the GC retention time (11.74 min) and the mass spectral data of 26-norcastasterone bismethanoboronate were different from those of the 28-norcastasterone bismethanoboronate. Thus, it was demonstrated that  $[26,28-^2H_6]$ castasterone was converted to  $[26-^2H_3]$ 28-norcastasterone in the seedlings of *A. thaliana* (Fig. 1). In this administration experiment, the conversion ratio (the percentage of the detected amount of the metabolite versus the amount of added substrate) was ca. 5%.

A similar experiment was performed using the seedlings of BR-insensitive mutants (*bri1-5*) of *A. thaliana*, where  $[^2H_3]$ 28-norcastasterone was identified as a metabolite of  $[26,28-^2H_6]$ castasterone by full-scan GC–MS analysis, confirming that castasterone is the biosynthetic origin of 28-norcastasterone in *A. thaliana*.

## 2.2. Metabolism of $[26,28-^2H_6]$ castasterone in cultured cells of *C. roseus*

To test whether or not our finding in *A. thaliana* occurs in other higher plants, we examined the metabolism of  $[26,28-^2H_6]$ castasterone in cultured cells of *Catharanthus roseus*.

$[26,28-^2H_6]$ castasterone was administered to cultured cells of *C. roseus*, and incubated for 2 days. The culture was extracted with methanol, and the extract was purified. The purified HPLC fractions were converted to methanoboronate derivatives and analyzed by GC–MS. A prominent peak, co-migrating with authentic 28-norcastasterone, was found in the HPLC fraction (Rt: 6–8 min).

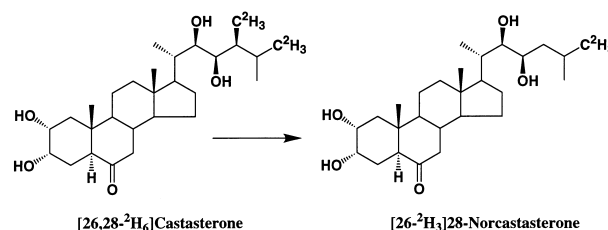


Fig. 1. Conversion of castasterone to 28-norcastasterone. In this study, this conversion was demonstrated in seedlings of *Arabidopsis thaliana*, rice and tomato, and cultured cells of *Catharanthus roseus*.

Mass spectral data of the metabolite was as follows:  $m/z$  501 ( $M^+$ , 90), 486 (12), 459 (6), 399 (7), 358 (22), 287 (76), 144 (100). Molecular and fragment ions such as  $m/z$  501 and 144 corresponding to  $m/z$  498 and 141 of non-labeled 28-norcastasterone-bismethanoboronate derivative was detected, along with characteristic ions of 28-norcastasterone bismethanoboronate (such as  $m/z$  399, 358 and 287). Thus,  $[^2H_3]$ 28-norcastasterone was identified as a metabolite of  $[26,28-^2H_6]$ castasterone, demonstrating that castasterone is converted to 28-norcastasterone in cultured cells of *C. roseus*. In the same HPLC fraction,  $[^2H_6]$ brassinolide was identified as a major metabolite of  $[^2H_6]$ castasterone by full-scan GC–MS analysis, confirming our previous work (Suzuki et al., 1993). Conversion ratios in this feeding were ca. 12% ( $[^2H_6]$ brassinolide) and ca. 4% ( $[^2H_3]$ 28-norcastasterone).

## 2.3. Metabolism of $[^2H_6]$ castasterone in seedlings of *Oryza sativa* and *Lycopersicon esculentum*

The metabolism of  $[^2H_6]$ castasterone in seedlings of rice (*Oryza sativa*) and tomato (*Lycopersicon esculentum*) was also investigated. In both plant species,  $[^2H_3]$ 28-norcastasterone was identified as a metabolite of  $[26,28-^2H_6]$ castasterone by full-scan GC–MS analysis. In tomato, ca. 4% of the substrate was converted to the metabolite following 2 day incubation, while lower conversion was found in rice (ca. 1%).

## 2.4. Identification of 28-norcastasterone and a novel brassinosteroid, 28-nortyphasterol in seedlings of *Arabidopsis thaliana*

In *A. thaliana*, the natural occurrence of 28-norcastasterone has not yet been demonstrated. Accordingly we have re-examined the natural occurrence of BRs in the seedlings of *A. thaliana*. Thus, *bri1-5* seedlings were grown and used for the identification of BRs, because *bri1-5* was shown to accumulate very high levels of endogenous BRs (Noguchi et al., 1999). The final HPLC-purified fractions were derivatized and analyzed by GC–MS. From the ODS-HPLC fraction (Rt: 8–10 min), 28-norcastasterone and brassinolide were identified by full-scan GC–MS analysis. Mass spectra and retention

Table 1  
GC–MS identification of Brassinosteroids in *Arabidopsis thaliana*

Brassinosteroid	Retention time (min)	Characteristic ions $m/z$ (relative intensity)
28-Norcastasterone		
standard	11.60	498 [ $M^+$ ] (80), 483 (6), 399 (4), 358 (12), 329 (11), 287 (51), 228 (19), 141 (100)
endogenous	11.60	498 [ $M^+$ ] (88), 483 (7), 399 (4), 358 (13), 329 (11), 287 (48), 228 (19), 141 (100)
Brassinolide		
standard	13.20	528 [ $M^+$ ] (4), 457 (2), 397 (7), 374 (21), 345 (11), 332 (22), 177 (82), 155 (100)
endogenous	13.22	582 [ $M^+$ ] (4), 457 (2), 397 (7), 374 (24), 345 (12), 332 (24), 177 (90), 155 (100)
Castasterone		
standard	12.23	512 [ $M^+$ ] (53), 497 (5), 441 (4), 399 (8), 358 (20), 329 (11), 287 (28), 155 (100)
endogenous	12.22	512 [ $M^+$ ] (56), 497 (6), 441 (4), 399 (10), 358 (22), 329 (13), 287 (31), 155 (100)
28-Nortyphasterol		
standard	11.05	530 [ $M^+$ ] (91), 515 (58), 501 (100), 440 (48), 425 (16), 229 (33), 141 (51)
endogenous	11.05	530 [ $M^+$ ] (88), 515 (56), 501 (100), 440 (42), 425 (15), 229 (30), 141 (50)

times on GC were identical to those of authentic specimens (Table 1). This is the first full-scan GC–MS identification of 28-norcastasterone and brassinolide in *A. thaliana*, although the presence of brassinolide has been already shown by GC–SIM analysis (Fujioka et al., 1998b; Noguchi et al., 1999). From the HPLC fractions (Rt: 10–12 min, 12–14 min), castasterone was identified by full-scan GC–MS analysis (Table 1). The natural co-occurrence of castasterone and 28-norcastasterone, and the conversion of castasterone to 28-norcastasterone support our assumption that castasterone is the biosynthetic origin of 28-norcastasterone.

In this series of experiments, a novel BR was identified. From the HPLC fraction (Rt: 20–23 min), the following mass spectral data was obtained as a methanoboronate-trimethylsilyl derivative; prominent ions at  $m/z$  530 [ $M^+$ ], 515, 501 (base peak) and 440. This mass spectral data is very similar to that of a typhasterol-methanoboronate-trimethylsilyl derivative (prominent ions at  $m/z$  544 [ $M^+$ ], 529, 515 (base peak) and 454), suggesting that this BR is a typhasterol analog lacking a methyl group. By direct comparison with an authentic specimen, this BR was shown to be 28-nortyphasterol (Table 1). This is the first report of the natural occurrence of 28-nortyphasterol in the plant kingdom.

## 2.5. Biological activity of 28-norcastasterone and castasterone, and 28-nortyphasterol and typhasterol

Biological activity was evaluated using the rice lamina inclination test (Fujioka et al., 1998a). Two assay methods were employed: single application of the sample (Fig. 2), and co-application with 5  $\mu$ g of indole-3-acetic acid (Fig. 3). In both assay methods, the activity of 28-norcastasterone was shown to be ca. 10% of that of castasterone (Figs. 2 and 3). Activity of 28-nortyphasterol was also shown to be ca. 10% of that of typhasterol (Figs. 2 and 3). Relative activities were as follows: castasterone (100), 28-norcastasterone (10), typhasterol (10), 28-nortyphasterol (1).

The present study revealed that castasterone is converted to 28-norcastasterone at least in four plant species, that is, *A. thaliana*, *C. roseus*, rice and tomato, indicating that castasterone is the biosynthetic origin of 28-norcastasterone. This was a totally unexpected result, because we originally thought that 28-norcastasterone would be biosynthesized from 28-nor-6-deoxocastasterone by analogy of the established step of 6-deoxocastasterone to castasterone. The present study clearly shows that 28-norcastasterone is biosynthesized from castasterone. This result explains the fact that when 28-norcastasterone was identified in a plant, castasterone was always found to co-occur in the same plant (Fujioka, 1999). However, castasterone may not be the only biosynthetic origin of 28-norcastasterone. As the natural occurrence of 28-nor-6-deoxocastasterone has been found in several plant species (Fujioka, 1999), this BR may be another biosynthetic precursor of 28-norcastasterone. In any case, the natural occurrence of 28-nortyphasterol is intriguing:

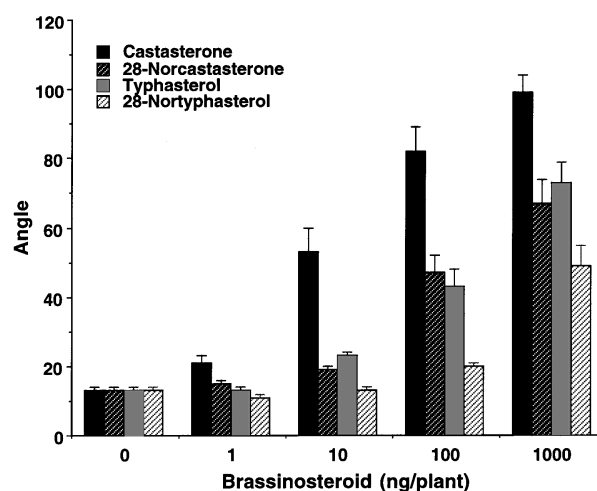


Fig. 2. Rice lamina inclination test of castasterone, 28-norcastasterone, typhasterol and 28-nortyphasterol by single application of the test sample. Each data point represents the mean of 30 replicates  $\pm$  SE.

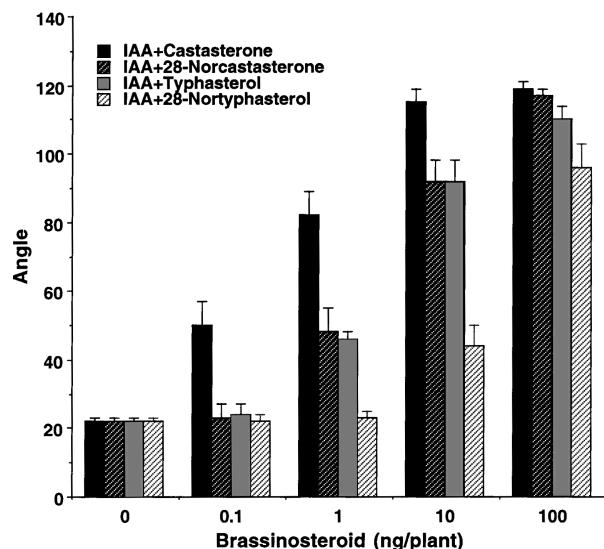


Fig. 3. Rice lamina inclination test of castasterone, 28-norcastasterone, typhasterol and 28-nortyphasterol by co-application of the test sample with IAA (5  $\mu$ g/plant). Each data point represents the mean of 30 replicates  $\pm$  SE.

the co-occurrence of 28-nortyphasterol and 28-norcastasterone suggests that 28-norcastasterone may be biosynthesized from 28-nortyphasterol. Furthermore, typhasterol may be predicted to be a precursor of 28-nortyphasterol by analogy of the findings in the present study. Furthermore, 28-nortyphasterol may be another precursor of 28-norcastasterone; detailed metabolic studies of these BRs will clarify these possibilities.

The biological activity of 28-norcastasterone was shown to be ca. 10% of that of castasterone. This means that the level of castasterone could be regulated by this metabolism. Thus, the conversion of castasterone to 28-norcastasterone may, in effect, have some regulatory role in maintaining the proper level of castasterone, an immediate precursor of brassinolide. This speculation must be also clarified by further studies.

### 3. Experimental

#### 3.1. General

GC–MS analysis was carried out on a JEOL Auto-mass JMS-AM 150 mass spectrometer connected to a Hewlett-Packard 5890A-II gas chromatograph with a capillary column DB-5 (0.25 mm  $\times$  15 m, 0.25  $\mu$ m film thickness). The analytical conditions were the same as previously described (Fujioka et al., 1997).

#### 3.2. Metabolism of [ $^2$ H $_6$ ]castasterone in seedlings of *A. thaliana*

[26,28- $^2$ H $_6$ ]Castasterone (Takatsuto and Ikekawa, 1986), 28-norcastasterone (Takatsuto et al., 1984), and

26-norcastasterone (Watanabe, Seto, Yokota and Takatsuto, unpublished work) were chemically synthesized.

Before precursor administration experiments, seven-day-old seedlings of *A. thaliana* (Ws-2: 20 seedlings, *bri1-5*: 50 seedlings) were transferred to a 200-ml flask containing 30 ml of half-strength MS medium supplemented with 1% sucrose. Seven days after transfer, a MeOH solution (10  $\mu$ l) of [26,28- $^2$ H $_6$ ]castasterone (10  $\mu$ g) was added. The seedlings were incubated for 2 days at 22°C in the light on a shaker (120 rpm), then extracted with MeOH. The MeOH extract was partitioned between CHCl $_3$  and H $_2$ O. The CHCl $_3$ -soluble fraction was purified with a cartridge of silica gel (Sep-Pak Vac 2 g; Waters, Milford, MA), which was eluted with 30 ml of chloroform, 2% MeOH in chloroform, and 7% MeOH in chloroform. The last fraction was purified with HPLC on a 150  $\times$  4.6-mm Senshu Pak ODS-1151-D column (Senshu Scientific Co., Ltd., Tokyo) by using 45% acetonitrile at a flow rate of 1.0 ml/min. The fractions were collected at two-minute intervals (Rt of 4 to 30 min). Each fraction was subjected to GC–MS analysis after derivatization.

#### 3.3. Metabolism of [ $^2$ H $_6$ ]castasterone in cultured cells of *C. roseus*

Cultured cells of *C. roseus* (V208) were grown in a MS medium supplemented with 3% sucrose at 27°C by shaking at 100 rpm in the dark. A MeOH solution (10  $\mu$ l) of [ $^2$ H $_6$ ]castasterone (10  $\mu$ g) was added to a 200-ml flask containing cultured cells, which were grown for 10 days (log phase) in 60 ml MS medium. The cells were allowed to grow for 2 days. After incubation, cultures were extracted with MeOH, and the extract was purified by the same method described above.

#### 3.4. Metabolism of [ $^2$ H $_6$ ]castasterone in seedlings of *O. sativa* and *L. esculentum*

Seedlings (*O. sativa* cv. Koshihikari and *L. esculentum* cv. Alisa Craig) were grown in pots containing vermiculite at 25°C under continuous light. Prior to precursor administration, they were transferred to a water culture in a 10-ml conical flask containing 10 ml of water. A MeOH solution (10  $\mu$ l) of [ $^2$ H $_6$ ]castasterone (10  $\mu$ g) was added into the water and allowed to grow for additional 2 days. Extraction and purification were performed by the same method described above.

#### 3.5. Identification of 28-norcastasterone and 28-nortyphasterol in *A. thaliana*

Thirty-two grams of five-week-old plants (*bri1-5*) were collected and extracted with MeOH. The MeOH extract was partitioned between CHCl $_3$  and H $_2$ O. The

CHCl<sub>3</sub>-soluble fraction was purified with a cartridge of silica gel (Sep-Pak Vac 10 g, Waters, Milford, MA), which was eluted with 100 ml of chloroform, 2% MeOH in chloroform, and 7% MeOH in chloroform. The last fraction was purified by ODS cartridge column (Sep-Pak Plus C18, Waters, Milford, MA), which was eluted with 20 ml of MeOH, and subjected to ODS-HPLC (Senshu Pak Pegasil ODS, 10×30 mm + Senshu Pak Pegasil ODS, 20×250 mm; Senshu Scientific Co., Ltd., Tokyo) by using 70% acetonitrile at a flow rate of 8 ml/min. The fractions were collected at 2-min intervals (Rt of 6–14 min) and 3-min intervals (Rt of 14 to 35 min). Each fraction was subjected to GC–MS analysis after derivatization. Authentic 28-nortyphasterol was chemically synthesized (Takatsuto and Ikekawa, 1987).

### 3.6. Rice lamina inclination test

Rice lamina inclination test was carried out according to the methods reported by Fujioka et al. (1998a).

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