SHORT REPORTS

IDENTIFICATION OF GA₃₆ IN PSILOTUM NUDUM

MASAHIRO TAKAHASHI, HISAKAZU YAMANE, YOSHIO SATOH, NOBUTAKA TAKAHASHI and KUNIO IWATSUKI*

Department of Agricultural Chemistry, Faculty of Agriculture, The University of Tokyo, Bunkyo-ku, Tokyo 113, Japan; *Botanical Gardens, University of Tokyo, Hakusan Bunkyo-ku, Tokyo 112, Japan

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Key Word Index-Psilotum nudum; Psilotaceae; pteridophyte; GA₃₆.

Abstract—GA₃₆ was identified by GC/MS analysis from the sporangia-bearing fronds of *Psilotum nudum*. This is the first chemical identification of a free gibberellin in a pteridophyte.

INTRODUCTION

We have been investigating phytohormones in pteridophytes and the occurrence of cytokinins and ABA was reported in the previous papers [1, 2]. We wish to report here the identification by GC/MS analysis of GA₃₆ from the pteridophyte *Psilotum nudum*.

RESULTS AND DISCUSSION

An acidic ethyl acetate fraction from sporangia-bearing fronds (250 g fr.wt) of P. nudum was subjected to silica gel partition column chromatography (stationary phase; 0.5 M aqueous formic acid) eluted with ethyl acetate—nhexane. The GA active eluate with 40% ethyl acetate in nhexane was further purified by HPLC using a Nucleosil N(Me)₂ column eluted with 0.05% acetic acid in methanol [3] to afford the active eluate at R_t 10–14 min. The finding that the active fraction showed GA activity on cv. Waito-C as well as cv. Tan-ginbozu in the rice seedling test, suggested that it contained a C-3 hydroxy GA as an active component [4]. Finally, we succeeded in identifying GA₃₆ by GC/MS analysis of its trimethylsilyl ester trimethylsilyl ether derivative. The content of GA₃₆ in P. nudum was estimated to be 2.34 μ g/kg fr. wt.

This is not only the first chemical identification of a free gibberellin from a pteridophyte, but also the first example for the occurrence of GA_{36} as an endogenous GA in vascular plants. GA_{36} is considered to be a direct precursor of GA_4 which is one of the most common GA_5 . The accumulation of GA_{36} in a pteridophyte may suggest that biological conversion of GA_{36} to GA_4 is either very slow as compared with that of higher plants or blocked.

EXPERIMENTAL

Plant material. Sporangia-bearing fronds of Psilotum nudum were harvested at the greenhouse of Kyoto University, Kyoto city, Japan, in November 1982.

Extraction and fractionation. Fresh plant material (250 g fr. wt) was homogenized and extracted with 500 ml of Me₂CO (×2). The mixture was filtered and the Me₂CO removed in vacuo. The resultant aq. residue was fractionated by the same method as that of Yamane et al. [5] to give an acidic EtOAc fraction (317.30 mg).

Partition column chromatography. The acidic EtOAc fraction adsorbed on celite (2 g) was placed on a column packed with silica gel (16.0 g) impregnated with 0.5 M aq. HCOOH (10 ml) and eluted with EtOAc-n-hexane (0-100% EtOAc in 10% steps/50 ml).

HPLC. Nucleosil 5N(Me)₂ column (6 mm i.d. × 100 mm) eluted with MeOH containing 0.05% AcOH was used. The flow rate was 1.5 ml/min, and the column temp. was 50°.

Bioassay. The dwarf rice test was carried out at a concn of 10 g fr. wt eq./ml according to the water culture method used by Yokota et al. [6] at 30° under continuous illumination (fluorescent light). Seeds of Oryza sativa L., dwarf cv. Tan-ginbozu and Waito-C were used for the assay.

GC/MS analysis. A Hitachi M80-A equipped with a glass column (3 mm i.d. \times 1 m) packed with 2 % OV-1 on Chromosorb W was used. Ionizing voltage, 20 eV; oven temp, 210°; He, 50 ml/min. A sample for GC/MS was silylated with pyridine-BSA-TMCS (1:2:1) at room temp. R_t of GA₃₆-TMS-TMS in GC: 4.3 min. GA₃₆-TMS-TMS from P. nudum showed the following mass spectrum in GC/MS: 578 [M] + (12%), 563 (51), 550 (19), 488 (79), 460 (58), 432 (45), 398 (55), 370 (84), 342 (100), 314 (38), 280 (46), 224 (89). The mass spectrum of authentic GA₃₆-TMS-TMS in GC/MS: 578 [M] + (20), 563 (51), 550 (18), 488 (75), 460 (64), 432 (54), 398 (66), 370 (82), 342 (100), 314 (51), 280 (58), 224 (99).

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