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5,7,2',4'-TETRAHYDROXY-8-(3"-HYDROXY-3"-METHYL-BUTYL) ISOFLAVANONE, A METABOLITE OF KIEVITONE PRODUCED BY FUSARIUM SOLANI f.sp. PHASEOLI

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Key Word Index—Phaseolus vulgaris; Fusarium solani f.sp. phaseoli; Leguminosae; fungal metabolism; isoflavonoids; phytoalexins; kievitone; 5,7,2',4'-tetrahydroxy-8-(3"-hydroxy-3"-methyl-butyl) isoflavanone; kievitone hydrate; detoxification.

INTRODUCTION

Fusarium solani (Mart.) Sacc. f.sp. phaseoli (Burk.) Snyd. and Hans., a pathogen of French bean (Phaseolus vulgaris) is relatively insensitive to the phytoalexins produced by P. vulgaris [1, 2]. The tolerance of F. solani f.sp. phaseoli to the bean phytoalexin phaseollin, a pterocarpan [3, 4], has been attributed to its ability to metabolize phaseollin to a less fungitoxic product, 1ahydroxyphaseollone [5, 6]. The metabolism of phytoalexins by plant pathogenic fungi is now well established but, to date, pterocarpans are the only isoflavonoid phytoalexins for which structures of fungal metabolites are known [7]. Kievitone (1), an isoflavanone [8, 9]

kievitone (21.6 mg of kievitone yielded 14.3 mg of kievitone hydrate). The purified material had lower R_f values than kievitone on Si gel chromatograms [10], but yielded a similar orange-brown reaction product with diazotized p-nitroaniline. Its $UV[\lambda_{max}^{ENOH}$ (log ε) 214 (4.39) and 293 (4.22) nm; shoulder 340 (3.5) nm] and IR [KCl disc: OH band, 3350 cm⁻¹; carbonyl band, 1642 cm⁻¹] spectra were virtually identical to those of kievitone [9]. Low and high resolution MS (obtained using a heated direct insertion probe) revealed a M^+ at m/e 374.1373 (15%, $C_{20}H_{22}O_7$), 18 amu higher than kievitone [8, 9], indicating the addition of water to kievitone. Other major peaks in the MS occurred at m/e 356 (6%), 300 (64%), 221 (65%), 205 (15%), 192

and another of the five phytoalexins produced by P. vulgaris [7], is also metabolized by F. solani f.sp. phaseoli to a less fungitoxic compound [10]. The metabolite is a hydrated product of kievitone and this paper presents evidence that its structure is (2).

RESULTS AND DISCUSSION

The metabolite, 'kievitone hydrate', was isolated in mg amounts from liquid cultures of F. solani containing

(14%), 177 (26%) and 165 (100%), all consistent with the fragmentation observed for kievitone [9].

The structure of (2) was unequivocally established as 5,7,2',4'-tetrahydroxy-8-(3"-hydroxy-3"-methyl-butyl) isoflavanone from a comparison of the 100 MHz PMR spectra in DMSO-d₆ and MeOH-d₄ with the reported data for kievitone [9] (Table 1). The olefinic proton H-2" and the methylene proton H-1" of kievitone were totally absent in the spectrum of kievitone hydrate and were replaced by two multiplets which showed the typical AA'XX' structure expected for a X-CH₂-CH₂-Y

fragment. The chemical shifts were typical for methylene in an aliphatic chain (1.5 ppm) and methylene substituted by an aromatic ring (2.5 ppm). In MeOH-d₄ this ethane fragment probably has an exclusively trans arrangement of substituents since solvation of the C-7 and C-3"

Table 1. Chemical shifts (ppm) of kievitone and kievitone hydrate

Proton	Kievitone hydrate		Kievitone [9]
	MeOH-d₄	DMSO-d ₆	DMSO-d ₆ /CDCl ₃
H-6	5.95 (s)	5.98 (s)	6.04 (s)
H-1"	2.8 (m)	2.5(m)	3.21 (d)
H-2"	1.6 (m)	1.5(m)	5.18(t)
H-4"	1.25 (s)	1.15 (s)	1.64(s)
Me-3"			1.74 (s)
H-2a	4.5 (m)	4.45 (m)	4.49(m)
Н-2Ь			4.64 (m)
H-3	4.15 (dd)	4.15 (dd)	4.16 (dd)
H-3'	6.31 (d)	$6.3(\hat{d})$	6.45 (d)
H-5'	6.25 (dd)	6.15 (dd)	6.32 (dd)
H-6'	6.97 (d)	6.78(d)	6.91 (d)

hydroxyl groups will destroy any hydrogen bonding between these OH groups; such hydrogen bonding is the only factor which could offset the normal steric repulsions which reduce the population of the gauche conformer. A trans conformation about the bond between C-1" and C-2" implies a molecular shape for kievitone hydrate closely similar to that of its precursor kievitone. The nonequivalent Me groups of kievitone (1) were replaced by a 6-proton singlet in the spectrum of (2), indicating that hydration of the C=C group of kievitone had occurred in a Markovnikov fashion to yield a 3"-OH rather than a 2"-OH side chain. The presumably enzymatically-catalysed hydration was highly specific since no evidence of the system with a 2"-OH side chain was found in the spectrum.

In MeOH-d₄ all OH protons appeared as a single peak at 4.7 ppm due to rapid exchange. Some exchange was also evident in DMSO-d₆ since the phenolic protons for the 7-OH, 2'-OH and 4'-OH groups appeared as a very broad peak at ca 9.5 ppm. The aliphatic tertiary OH group was present as a very broad absorption around 3.8 ppm. A sharp singlet at 12.3 ppm (compared with 12.2 ppm in kievitone) confirmed the presence of the strongly intramolecular-hydrogen-bonded OH at C-5.

Preliminary bioassays of kievitone hydrate against Cladosporium herbarum (Pers.) Link. [10], Aphanomyces euteiches Drechs. and Rhizoctonia solani Kühn indicated that this metabolite was considerably less fungitoxic than kievitone. The metabolic conversion of kievitone to kievitone hydrate by F. solani f.sp. phaseoli, therefore, may well explain the comparative insensitivity of this fungus to the phytoalexin.

EXPERIMENTAL

Kievitone was obtained following procedures outlined previously [2]. Actively growing mycelium of F. solani f.sp. phaseoli was obtained as described elsewhere [5]. Kievitone dissolved in EtOH was added to fungal cultures at concus of 25–50 µg/ml; the final EtOH cone in the medium was 0.5%. The cultures were incubated at 25° on a reciprocal shaker for 6 hr, after which kievitone hydrate was isolated by slight modification of methods reported elsewhere [5]. The metabolite was purified by sequential TLC on Si gel G developed with C_6H_6 -MeOH (9:1) (R_f 0.07). Further purification was obtained by gel filtration through Sephadex LH-20 employing EtOH as the eluant. Bioassays were carried out as described in refs. [2, 10].

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ANTHOCYANIN COMPOSITION OF BRASSICA OLERACEA CV. RED DANISH

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Key Word Index—Brassica oleracea cv. Red Danish; Cruciferae; anthocyanins; malonyl-, p-coumaryl-, ferulyl-, sinapyl esters of cyanidin-3-sophoroside-5-glucoside.

Abstract—Eight anthocyanins were isolated from illuminated red cabbage seedlings. They were identified as: cyanidin-3-sophoroside-5-glucoside, cyanidin-3-p-coumaryl-sophoroside-5-glucoside, cyanidin-3-p-coumaryl-sophoroside-5-glucoside, cyanidin-3-ferulyl-sophoroside-5-glucoside, cyanidin-3-(diferulyl)sophoroside-5-glucoside, cyanidin-3-ferulyl-sophoroside-5-glucoside, cy