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Sterol Composition of the Corn Root Lesion Nematode, *pratylenchus agilis*, and Corn Root Cultures

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ABSTRACT: Sterols from mixed stages of the corn root lesion nematode, *Pratylenchus agilis*, and uninfected corn root cultures were analyzed by gas chromatography-mass spectrometry. Twenty-eight sterols were identified in *P. agilis*, including 9 not previously detected in nematodes. The major sterols were 24-ethylcholest-22-enol, 24-ethylcholesta-5,22-dienol, 24-methylcholestanol, 24-ethylcholestanol, isofucostanol, and 24-ethylcholesterol. The principal corn root sterols were 24-ethylcholesta-5,22-dienol, 24-methylcholesterol, 24-ethylcholesterol, isofucosterol, and cycloartenol. Therefore, the major metabolic transformation of sterols by *P. agilis* was saturation of the sterol nucleus. In addition, very small amounts of 4α -methylsterols were biosynthesized by *P. agilis*. The 4-methylation pathway is unique to nematodes and was previously demonstrated only in free-living Rhabditida. The sterol composition of *P. agilis* is remarkably different from that of other nematodes in its very small relative percentage of cholesterol (0.4% of the total nematode sterol).

KEY WORDS: corn, lesion nematode, lipid, maize, nematode, Pratylenchus agilis, sterol, Zea mays.

The development of new control strategies for parasitic nematodes could exploit biochemical differences between them and their vertebrate or plant hosts. One of the few such areas is steroid metabolism; unlike higher animals and plants, nematodes possess a nutritional requirement for sterol as a result of their lack of de novo sterol biosynthesis (for review: Chitwood et al., 1986). Sterols have 2 major functions; they are integral components of cell membranes as well as metabolic precursors of steroid hormones (Nes and McKean, 1977). Interest in nematode sterol biochemistry has intensified in recent years because of the discovery that several azasteroids and aliphatic amines strongly disrupt growth and development and also inhibit sterol metabolism in nematodes (Chitwood, 1987).

Because free-living nematodes are easily propagated in artificial media, they have been utilized in most investigations involving metabolism of radiolabeled dietary sterols by nematodes (Chitwood et al., 1986). The difficulty in in vitro culture of parasitic nematodes has restricted investigation of sterol metabolism in these organisms to comparison of sterol compositions of host and parasite. As expected, cholesterol is the major sterol of mammalian parasites (Cole and Krusberg, 1967b; Castro and Fairbairn, 1969; Barrett et al., 1970; Fulk and Shorb, 1971; Chung et al., 1986).

A significant biochemical difference between higher animals and plants is that plant sterols

usually contain alkyl (i.e., methyl or ethyl) substituents at C-24 of the sterol side chain (Nes and McKean, 1977). Comparison of the sterols of plant-parasitic nematodes and their hosts has indicated that plant parasites are similar to freeliving nematodes in that both groups can remove the C-24 alkyl groups of plant sterols (Chitwood, 1987). Unequivocal proof in the case of plant parasites, however, would involve experiments with radiolabeled plant sterols in an in vitro system. Because large quantities of the corn root lesion nematode, Pratylenchus agilis Thorne and Malek, 1968, can be readily obtained from corn root explant cultures, we examined the sterols of this plant parasite and its host in order to determine if it was a suitable candidate for radiotracer studies of dealkylation of plant sterols by phytoparasitic nematodes.

Materials and Methods

Seeds of Zea mays cv. Iochief were surface sterilized by immersion in distilled water for 2 hr and then 1.3% sodium hypochlorite containing 0.1% (v:v) Tween 80 for 25 min. The sterilized seeds were germinated for 4 days on 1.5% water agar in culture dishes, and then 2.5-cm root pieces were transferred to culture dishes containing Gamborg's B5 medium formulated with 1.5% agar and 2.0% sucrose, and lacking cytokinis or auxins (Huettel, 1990). The agar had been extracted previously with methanol and chloroform : methanol 2:1 (v:v, twice each) to reduce the high amounts of endogenous cholesterol in commercial agar (Nes, 1987) to an acceptable level (20.6 ng of cholesterol per gram dry weight as determined by subsequently described Table 1. Sterol content of *Pratylenchus agilis* and root cultures of *Z. mays*.

	% dry weight		- % sterol	
	Lipid	Sterol	esterified	
P. agilis	14.6	0.03	26.8	
Z. mays	3.9	0.83	_*	

* Not determined.

methods). One-week-old root cultures were inoculated with agar plugs from similar cultures containing sterile *P. agilis*. Nematodes from 8-week-old cultures maintained at 25°C were harvested with modified Baermann funnels, floated on 30% sucrose, rinsed 5 times with distilled water, frozen immediately, and lyophilized.

Lipids were extracted from lyophilized nematodes (2.0-2.2 g dry wt) by homogenization in a Ten-Broeck tissue grinder 3 times with chloroform : methanol 2:1 and were purified by partition against 0.85% NaCl (Folch et al., 1957). Neutral lipids were separated from polar lipids and were fractionated into various lipid classes by column chromatography on silica (Chitwood et al., 1985). The lipid fractions containing steryl esters and free sterols were saponified separately in methanolic KOH (Chitwood et al., 1985); the liberated 4-desmethylsterols and 4-methylsterols were purified by Florisil column chromatography (Chitwood et al., 1987).

Roots from uninfected cultures were gently removed with forceps, frozen, and lyophilized. Extraction of roots (0.3-0.5 g dry wt) was identical to that of nematodes, except a Virtis homogenizer was used. The root lipids were saponified directly without prior fractionation on a silica column. The 4-methylsterols and 4-desmethylsterols were isolated as described above. As a precautionary measure, any potentially tightly bound sterols were removed from the extraction residue by saponification of the residue and analyzed identically to the root extracts.

Sterols were analyzed by gas-liquid chromatography (GLC) on a DB-1 fused silica capillary column and an OV-17 packed glass column as described previously (Chitwood et al., 1987c). Following preliminary analysis, the sterols were acetylated, the steryl acetates were purified on Florisil columns and analyzed by GLC, and then tentative identifications were confirmed by gas chromatography-mass spectrometry (GC-MS, Chitwood et al., 1987c). The steryl acetates were also fractionated according to degree and position of unsaturation by chromatography on columns of silica impregnated with AgNO₃ and analyzed by GLC (Chitwood et al., 1987c).

Results are the means of two replicates.

Results

Gravimetric analysis indicated that lipid constituted approximately 15% of the dry weight of *P. agilis*; subsequent quantification of sterols by GLC revealed that sterol comprised only a very small proportion of the lipid extract (Table 1). Table 2. Gas-liquid chromatographic relative retention times (RRT's) of sterols from *Pratylenchus agilis* and *Zea mays*, expressed relative to cholesterol. GLC was performed on a temperature-programmed DB-1 fused silica capillary column (14 m \times 0.25-mm i.d., 0.25-µm film) and on an isothermally operated packed glass column containing 2.0% OV-17 stationary phase,

Sterol	DB-1	OV-17
Cholesterol	1.00	1.00
Cholestanol	1.02	1.02
24-Methylcholesta-5,22-dienol	1.11	1.14
Lathosterol	1.12	1.19
24-Methylcholest-22-enol	1.14	1.17
24-Methylcholesta-5,23-dienol	1.25	1.38
24-Methylenecholesterol	1.26	1.37
24-Methylenecholestanol	1.30	1.39
24-Methylcholesterol	1.30	1.33
24-Methylcholestanol	1.33	1.35
24-Ethylcholesta-5,22E-dienol	1.40	1.45
24-Ethylcholest-22E-enol	1.43	1.47
24-Methylcholesta-5,24(25)-dienol	1.47	1.67
24-Ethylcholesta-5,23-dienol	1.53	1.67
24-Ethylcholesta-7,22-dienol	1.57	1.70
Fucosterol	1.59	1.76
24-Ethylcholesterol	1.60	1.67
24-Ethylcholestanol	1.63	1.69
Isofucosterol	1.66	1.85
Isofucostanol	1.69	1.88
24-Ethylcholesta-5,24(25)-dienol	1.77	1.98
24-Ethyllathosterol	1.81	1.96
24-Ethylcholest-24(25)-enol	1.81	2.04
24Z-Ethylidenelathosterol	1.89	2.20
Obtusifoliol	1.52	1.56
	(1.49)*	(1.50)
4α,24-Dimethylcholestanol	1.58	1.52
	(1.53)	(1.48)
4α-Methyl-24-ethylcholest-22-enol	1.71	1.66
	(1.67)	(1.61)
4α-Methyl-24-ethylcholestanol	2.10	1.87
	(1.91)	(1.87)
4α-Methylisofucostanol	2.07	2.11
	(1.95)	(2.06)
4α-Methyl-24Z-ethylidenelathosterol	2.26	2.52
	(2.12)	(2.44)
Cycloartanol	1.71	1.67
	(1.60)	(1.53)
Cycloeucalenol	1.74	1.88
	(1.69)	(1.78)
Cycloartenol	1.86	2.05
	(1.73)	(1.85)
24-Methylenecycloartanol	2.14	2.28
	(1.99)	(2.06)

• Values in parentheses are RRT's of steryl acetates relative to cholesteryl acetate. Trivial and systematic names: cholesterol, cholest-5-en-3 β -ol; cholestanol, 5α -cholestan-3 β -ol; lathosterol, 5α -cholest-7-en-3 β -ol; fucosterol, 24E-ethylidenecholest-5-en-3 β -ol; isofucosterol, 24Z-ethylidenecholest-5-en- 3β -ol; isofucostanol, 24Z-ethylidene-5 α -cholestan-3 β -ol; obtusifoliol, 4α , 14-dimethyl-24-methylene- 5α -cholesta-8-en- 3β -ol; cycloartanol, 4,4,14-trimethyl-9 β , 19-cyclo- 5α -cholestan- 3β -ol; cycloeucalenol, 4α , 14-dimethyl-24-methylene- 9β , 19-cyclo- 5α cholestan- 3β -ol; cycloartenol, 4,4,14-trimethyl- 9β , 19-cyclo- 5α cholest-24-en- 3β -ol.

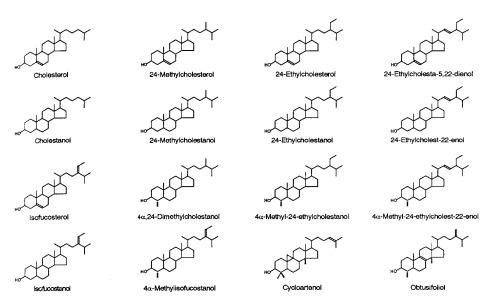


Figure 1. Some representative sterols from Pratylenchus agilis or Zea mays.

Most of the nematode sterol was unesterified. Total sterols were much more abundant in Z. mays. Saponification of the root extraction residue yielded only an additional 0.1% sterol.

All sterols from P. agilis and Z. mays (Fig. 1) were identical to authentic standards (when available) by GLC relative retention times (RRT's, Table 2) of both the free sterols and steryl acetate derivatives, argentation column chromatographic behavior, and GC-MS. Authentic 24-methylcholesta-5,24(25)-dienol was prepared by isomerization of 24-methylenecholesterol by iodine (Salt et al., 1986). Its GLC RRT's were similar to literature values (Itoh et al., 1982) and the mass spectrum of the steryl acetate (mass/charge m/z [structure and relative intensity]: 380 [molecular ion M+-CH₃COOH, 27], 365 [M-CH₃COOH-CH₃, 5], 296 [M-CH₃COOH-C₆H₁₁-H, 53], 281 [296-CH₃, 17], 253 [M-CH₃COOH-C₉H₁₇ side chain-2H, 19], 213 [M-CH₃COOH-side chain-C₃H₆, 17], and 83 [C₆H₁₁, 100]) was nearly identical to that of 24methylenecholesterol but contained an abundant side chain fragment at m/z 83 produced by allylic cleavage at C-22 due to the $\Delta^{24(25)}$ -bond. We lacked authentic standards for several compounds, including 24-ethylcholest-22E-enol, 24Z-ethylidenelathosterol, 4α , 24-dimethylcholestanol, 4α -methyl-24-ethylcholestanol, 4α - methyl-24Z-ethylidenelathosterol, cycloartanol, and 24-methylenecycloartanol. These were indistinguishable by GLC and GC-MS from the same compounds previously identified by us in *Heterodera zeae* and *Meloidogyne incognita* (Chitwood et al., 1985, 1987c).

Other sterols for which we lacked authentic standards were 24-methylcholest-22-enol, 24-ethylcholesta-5,23-dienol, 24-ethylcholesta-5,24(25)-dienol, 24-ethylcholest-24(25)-dienol, obtusifoliol, 4α -methyl-24-ethylcholest-22-enol, 4α -methylisofucostanol, and cycloeucalenol. These were identified as follows.

As is typical for other sterols with saturated ring systems (Patterson, 1971), the GLC RRT's for 24-methylcholest-22-enol were characteristically slightly greater than those of the corresponding sterol containing a Δ^5 -bond, i.e., 24methylcholesta-5,22-dienol. The mass spectrum of the steryl acetate (442 [M+, 18], 344 [M-C7H13-H, 29], 329 [344-CH₃, 3], 315 [M-C₉H₁₇ side chain-2H, 36], 257 [M-CH₃COOH-side chain, 57], 215 [M-CH₃COOH-side chain-C₃H₆, 14], and 55 [C4H7, 100]) was very similar to that of 24-ethylcholest-22-enyl acetate (Chitwood et al., 1985), but all ions containing an intact side chain were 14 mass units smaller. The GLC RRT's of 24-ethylcholesta-5,23-dienol were similar to literature values (Itoh et al., 1982); the mass spectrum of the acetate (394 [M-CH₃COOH, 100], 379 [M-CH₃COOH-CH₃, 7], 283 [M-CH₃COOH-C₈H₁₅, 16], 253 [M-CH₃COOH-C10H19 side chain-2H, 22], and 213 [M-CH₃COOH-side chain-C₃H₆, 11]) included a diagnostic loss of C8H15 resulting from allylic cleavage of the C-20(22) bond. The GLC RRT's for 24-ethylcholesta-5,24(25)-dienol were similar to literature values (Itoh et al., 1982). The mass spectrum of the steryl acetate (394 [M-CH₃COOH, 32], 379 [M-CH₃COOH-CH₃, 6], 296 [M-CH₃COOH-C₇H₁₃-H, 100], 281 [296-CH₃, 23], 253 [M-CH₃COOH-C₁₀H₁₉ side chain-2H, 18], and 213 [M-CH₃COOH-side chain- C_3H_6 , 9]) was similar to a literature spectrum (Kim et al., 1988), contained a more abundant side chain fragment at m/z 97 (analogous to the m/z 83 fragment in the spectrum of 24-methylcholesta-5,24(25)-dienyl acetate), but was otherwise indistinguishable from the spectra of fucosteryl and isofucosteryl acetates. The RRT's of 24-ethylcholest-24(25)-enol were slightly greater than RRT's of 24-ethylcholesta-5,24(25)dienol. The mass spectrum of the $\Delta^{24(25)}$ -acetate was indistinguishable from that of its $\Delta^{24(28)}$ -isomer, i.e., authentic isofucostanyl acetate (Chitwood et al., 1987c).

The fact that the RRT's of the newly identified 4α -methylsterols were greater as free sterols than as the acetate derivatives is indicative of the presence of a 4α -methyl substituent (Patterson, 1971). The RRT's of obtusifoliyl acetate (Table 2) and its mass spectrum (468 [M⁺, 28], 453 [M-CH₃, 64], 393 [M-CH₃COOH-CH₃, 22], 369 [M-CH₃-C₆H₁₁-H, 9], 343 [M-C₉H₁₇ side chain, 4], 309 [M-CH₃COOH-CH₃-C₆H₁₁-H, 11], 287 [M-side chain-CH₃-D ring, 19], 227 [287-CH₃COOH, 14], and 55 $[C_4H_7, 100]$) were similar to literature values (Itoh et al., 1978, 1982; Staphylakis and Gegiou, 1985; Rahier et al., 1989). The mass spectrum included ions produced by loss of a C_6H_{11} fragment resulting from allylic cleavage of the C-22 bond. The mass spectrum of 4α -methyl-24-ethylcholest-22-enyl acetate (470 [M+, 28], 455 [M-CH₃, 2], 427 [M-C₃H₇, 2], 367 [M-CH₃COOH-C₃H₇, 17], 358 [M-C₈H₁₅-H, 24], 343 [M-C₈H₁₅-H-CH₃, 4], 329 [M-C₁₀H₁₉ side chain-2H, 23], 271 [M-CH₃COOH-side chain, 33], 229 [M-CH₃COOH-side chain-C₃H₆, 8], and 55 $[C_4H_7, 100]$) included ions produced by the side chain fragmentations characteristic of Δ^{22} -sterols (Lenfant et al., 1967; Hutchins et al., 1970) and nuclear ions (229, 271) characteristic of 4α - methylsterols. The mass spectrum of 4α -methvlisofucostanyl acetate (470 [M+, 4], 455 [M-CH, 1], 372 [M-C₇H₁₃-H, 100], 357 [M-C₇H₁₃-H-CH 12], 329 [M-C₁₀H₁₉ side chain-2H, 6], 312 [M-CH3COOH-C7H13-H, 4], 269 [M-CH3COOH-side chain-2H,3],230[M-CH3COOH-sidechain-C3H, 27], and 229 [M-CH₃COOH-side chain-C₃H₆, 18]) was similar to a literature spectrum (Piretti and Viviani, 1989) and virtually identical to that of authentic isofucostanyl acetate (Chitwood et al., 1987b), but all ions containing an intact steroid nucleus were 14 mass units higher. The GLC RRT's of cycloeucalenyl acetate (Table 2) and its mass spectrum (468 [M+, 4], 453 [M-CH₃, 7], 408 [M-CH₃COOH, 62], 393 [M-CH₃COOH-CH₃, 37], 300 [M-CH₃COOH-A ring, 6], 283 [M-CH₃COOH-C₉H₁₇ side chain, 8], 281 [M-CH₃COOH-side chain-2H, 7], and 55 [C₄H₂, 100]) agreed with literature values (Itoh et al., 1978, 1982; Staphylakis and Gegiou, 1985; Rahier et al., 1989). The mass spectrum included the loss of the A ring characteristic of 9β , 19cyclopropyl sterols (Rahier and Benveniste, 1989).

The major sterols of corn root cultures were 24-ethylcholesta-5,22E-dienol, 24-methylcholesterol, and 24-ethylcholesterol (Table 3). In addition to these 3 sterols, *P. agilis* contained 24-ethylcholest-22E-enol as its major sterol and substantial quantities of 24-methylcholestanol, 24-ethylcholestanol, and isofucostanol. The free and esterified sterol fractions of *P. agilis* were qualitatively similar but quantitatively different; the steryl esters contained a much greater proportion of cycloartenol and 24-methylenecy-cloartenol.

Discussion

The lipid content of *P. agilis* (15%) was less than that found by Krusberg (1967) in *Pratylenchus penetrans* (25%) propagated on alfalfa callus. The small percentage of dry weight of *P. agilis* as sterol (0.03%) is within the range reported in other phytoparasitic nematodes: Globodera tabacum solanacearum (0.01%), Meloidogyne incognita (0.01%), M. arenaria (0.01%), Rotylenchulus reniformis, (0.02%), Heterodera zeae (0.05%), and Ditylenchus dipsaci (0.06%) (Cole and Krusberg, 1967a; Svoboda and Rebois, 1977; Orcutt et al., 1978; Chitwood et al., 1985, 1987c). These values are lower than the range of 0.2%-0.6% reported in the animal-par-

Table 3.	Relative percentages of	f sterols in roo	t cultures of 2	Zea mays and	in free sterol a	nd steryl ester fractions	
from Pra	tylenchus agilis.						

	Р. а	P. agilis	
	Free sterols	Steryl esters	Z. mays
Cholesterol	0.3	0.7	0.2
Cholestanol	0.4	0.4	Trace*
24-Methylcholesta-5,22-dienol	0.1	Trace	0.3
Lathosterol	0.1	Trace	Trace
24-Methylcholest-22E-enol	0.1	Trace	-+
24-Methylcholesta-5,23-dienol		-	0.1
24-Methylenecholesterol	0.2	0.1	1.2
24-Methylenecholestanol	0.4	0.2	0.1
24-Methylcholesterol	2.3	4.7	14.2
24-Methylcholestanol	11.0	4.8	0.5
24-Ethylcholesta-5,22E-dienol	24.6	21.9	60.3
24-Ethylcholest-22E-enol	34.2	18.4	1.4
24-Methylcholesta-5,24(25)-dienol		_	0.1
24-Ethylcholesta-5,23-dienol	_	_	0.2
24-Ethylcholesta-7,22E-dienol	0.4	0.1	0.3
Fucosterol	_	-	0.1
24-Ethylcholesterol	4.5	4.8	9.0
24-Ethylcholestanol	10.2	11.9	0.6
Isofucosterol	1.8	2.1	4.3
Isofucostanol	5.2	5.1	0.1
24-Ethylcholesta-5,24(25)-dienol	Trace		0.2
24-Ethyllathosterol	0.3	0.9	0.1
24-Ethylcholest-24(25)-enol	0.1	_	_
24Z-Ethylidenelathosterol	0.8	3.0	0.2
Obtusifoliol	0.1	0.2	0.1
4α ,24-Dimethylcholestanol	0.6	0.3	_
4α-Methyl-24-ethylcholest-22E-enol	0.4	0.4	
4α-Methyl-24-ethylcholestanol	0.6	0.2	_
4α-Methylisofucostanol	0.2	0.4	_
4α-Methyl-24Z-ethylidenelathosterol	0.1	Trace	0.1
Cycloartanol	-	-	0.3
Cycloeucalenol	0.1	_	0.4
Cycloartenol	0.7	16.1	3.9
24-Methylenecycloartanol	0.2	3.3	1.7

* Less than 0.05%.

† None detected.

asitic nematodes Ascaris suum, Nippostrongylus brasiliensis, Strongyloides ratti, and Trichostrongylus colubriformis (Cole and Krusberg, 1967b; Barrett, 1968, 1969; Tarr and Fairbairn, 1973; Chung et al., 1986). A small fraction of the sterols in P. agilis are esterified, as in H. zeae, Meloidogyne spp., and the free-living nematodes Caenorhabditis elegans and Panagrellus redivivus (Chitwood et al., 1984, 1985, 1987c; Salt et al., 1989). The composition of the free and esterified sterol fractions from P. agilis were qualitatively identical but quantitatively different. Most notably, the steryl esters were rich in cycloartenol and 24-methylenecycloartanol, 2 important intermediates in de novo sterol biosynthesis in higher plants (Nes and McKean, 1977). These

two 4,4,14-trimethylsterols are probably not utilized by nematodes, as a related 4,4,14-trimethylsterol, lanosterol, does not support growth and reproduction in *C. elegans* or the insect-parasitic nematode, *Steinernema feltiae* (Chitwood et al., 1987a; Ritter, 1988). Perhaps *P. agilis* and other nematodes isolate nonutilizable sterols by sequestering them as steryl esters.

The major sterols of *P. agilis* are 24-ethylcholest-22-enol, 24-ethylcholesta-5,22-dienol, 24methylcholestanol, 24-ethylcholestanol, isofucostanol, and 24-ethylcholesterol. The principal sterols of corn root cultures are 24-ethylcholesta-5,22-dienol, 24-methylcholesterol, 24-ethylcholesterol, and isofucosterol. Our results support other reports of 24-ethylcholesta-5,22-dienol (stigmasterol) as the predominant sterol of Z. mays roots and 24-ethylcholesterol (sitosterol) and 24-methylcholesterol as other abundant sterols (Rohmer et al., 1972; Bladocha and Benveniste, 1983). The most striking difference between P. agilis and its host is in the percentage of stanols (i.e., sterols without double bonds in the ring system). Less than 3% of the corn root sterols were stanols, whereas approximately twothirds of the sterols from P. agilis were stanols. Although selective uptake of the small quantities of plant stanols cannot be excluded, it seems likely that *P. agilis* can hydrogenate the Δ^5 -bonds of corn sterols to produce a variety of stanols. Two free-living nematodes, Turbatrix aceti and P. redivivus, reduce radiolabeled dietary cholesterol to form cholestanol (Chitwood et al., 1987b), and comparison of sterols of hosts and parasite indicate that S. feltiae, G. tabacum solanacearum, H. zeae, and Meloidogyne spp. also saturate Δ^{5} bonds (Chitwood et al., 1985, 1987b; Morrison and Ritter, 1986; Orcutt et al., 1978).

One interesting aspect of sterol metabolism in free-living Rhabditida is their production of significant quantities of 4α -methylsterols (Chitwood et al., 1984, 1987b). This 4-methylation pathway does not occur in any other class of organisms. Interestingly, we identified very small amounts of several 4α -methylstanols in *P. agilis*: 4α -methyl-24-ethylcholest-22-enol, 4α , 24-dimethylcholestanol, 4α -methyl-24-ethylcholestanol, and 4α -methylisofucostanol. Their small relative percentage (less than 2% of the total P. agilis sterol) makes any conclusion about their biosynthetic origin speculative. However, their endogenous biosynthesis by P. agilis is suggested by the facts that we were unable to isolate them from corn roots and that they are the 4-methyl derivatives of the 4 major products of sterol biosynthesis in P. agilis, i.e., the 4 abundant stanols.

Many compounds that inhibit sterol metabolism and growth and development in free-living nematodes inhibit the C-24 dealkylation pathway (Chitwood, 1987). Because free-living nematodes remove the C-24 methyl and ethyl groups of plant sterols and because the sterols of all other plant-parasitic nematodes examined consist of 5%-50% cholesterol, plant-parasitic nematodes are thought similarly to dealkylate phytosterols (Chitwood, 1987). *P. agilis*, however, is distinguished from other nematodes by the small (0.4%) relative percentage of cholesterol and may lack the ability to dealkylate phytosterols. The following sterols previously undetected in nematodes were identified in P. agilis in this investigation: 24-methylcholest-22-enol, 24methylcholesta-5,24(25)-dienol, 24-ethylcholesta-5,23-dienol, 24-ethylcholesta-5,24(25)-dienol, 24-ethylcholest-24(25)-enol, 4α -methyl-24ethylcholest-22-enol, 4α -methylisofucostanol. obtusifoliol, and cycloeucalenol. We are unaware of identification in Z. mays of the following: 24-methylenecholestanol, 24-methylcholesta-5,24(25)-dienol, 24-ethylcholesta-5,23-dienol. and 24-ethylcholesta-5,24(25)-dienol. Sterols with $\Delta^{24(25)}$ - or Δ^{23} -bonds have been implicated in the biosynthesis of the major sterols of corn (Zakelj and Goad, 1983), although few of these have been identified previously in Z. mays.

The sterol composition of *P. agilis* reflects the diversity of sterol metabolism in nematodes. There is no evidence for phytosterol dealkylation in *P. agilis*, in contrast to other plant-parasitic and free-living nematodes. Instead, *P. agilis* saturates the nucleus of its dietary sterols; the specific value to nematodes of stanol production is presently unknown. Further examination of sterol metabolism in taxonomically distant orders could reveal even greater diversity in sterol biochemistry between nematodes and their hosts.

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