# CODISTEROL AND OTHER $\Delta^5$ -STEROLS IN THE SEEDS OF CUCURBITA MAXIMA

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**Key Word Index**—*Cucurbita maxima*; Cucurbitaceae;  $\Delta^5$ -sterols; 25(27)-dehydroporiferasterol; clerosterol; isofucosterol; stigmasterol; sitosterol; campesterol; codisterol; 24 $\beta$ -ethylcholesta-5,22,25(27)-trien-3 $\beta$ -ol; 24 $\beta$ -ethylcholesta-5,25(27)-dien-3 $\beta$ -ol; 24 $\alpha$ -ethylcholesta-5,22-dien-3 $\beta$ -ol; 24 $\alpha$ -methylcholest-5-en-3 $\beta$ -ol; 24 $\alpha$ -methylcholesta-5,25(27)-dien-3 $\beta$ -ol.

**Abstract**—A substantial amount (ca 18%) of the sterol found in the seeds of Cucurbita maxima had a  $\Delta^5$ -bond and consisted of seven components. They were identified as 25(27)-dehydroporiferasterol, clerosterol, isofucosterol, stigmasterol, sitosterol, campesterol and codisterol. The C-24 configuration of each of the sterols was unequivocally established by a <sup>1</sup>H NMR spectral comparison with authentic standards. This is the first time codisterol has been found in a higher plant and also the first time the structures and configurations of the  $\Delta^5$ -sterols from a Cucurbitaceae species have been clearly characterized.

### INTRODUCTION

Sterols bearing a  $\Delta^5$ -bond, e.g. cholesterol and sitosterol, are believed to be the most common biological types and seem to be either required or preferred by the great majority of eukaryotes. This may be associated with a superior biological role which the  $\Delta^5$ -structure allows [1, 2]. However, the existence of some higher plant species which contain predominately  $\Delta^7$ -sterols is well documented [1-7]. Among these are members of the family Cucurbitaceae in which the occurrence of  $\Delta^5$ - along with the  $\Delta^7$ -sterols is assumed to be rare [4, 8–11]. Only a few investigators have reported the presence of  $\Delta^5$ -sterols in Cucurbitaceae plants [12-14] (in seeds and seedlings), and in most cases, a sufficient amount of sterol was not isolated to allow proper identification. However, in one publication, a significant level (ca 25 %) of  $\Delta^5$ -sterols was reported to occur in the seed oil of the cucurbit Trichosanthes kirilowii [14]. The structures of the various  $\Delta^5$ -components were determined by UV. IR and mass spectrometry as well as by GLC on a variety of liquid phases [14]. Although the configuration at C-24 was not determined, based on a pattern in which a homologous series at C-24 (H, CH<sub>3</sub>, C<sub>2</sub>H<sub>5</sub>) occurred together with the  $\Delta^{22}$ -derivative of the 24-ethyl-sterol, the components probably were cholesterol, campesterol, 22-dihydrobrassicasterol, sitosterol and stigmasterol, reflecting a 'mainline' distribution in which the 24-ethyl components are  $24\alpha$  and the 24-methyl analogues  $24\alpha$  and  $24\beta$  [9, 15]. In addition, a  $\Delta^{5, 25(27)}$ -sterol\* was present. Since no 24 $\alpha$ ethyl- $\Delta^{25(27)}$ -sterol has yet been isolated from a plant, the  $\Delta^{25(27)}$ -compound from T. kirilowii was probably  $24\beta$ , i.e.

clerosterol previously isolated from Clerodendrum infortunatum [16] and Kalanchoe diagremontiana [9].

In view of the existence of  $\Delta^5$ -sterols in the cucurbits as a family, we made an exceptionally careful study of the sterols of *Cucurbita maxima* seeds to ascertain whether  $\Delta^5$ -sterols could be found in this species. A substantial amount of  $\Delta^5$ -sterols was actually found distributed among seven components. In this paper we report the identification of each of these by the use of GLC, HPLC, mass spectrometry and <sup>1</sup>H NMR techniques.

# RESULTS AND DISCUSSION

A significant level of  $\Delta^5$ -sterols (accounting for ca 18%) of the total 4-desmethyl fraction) was isolated from the seeds of C. maxima by TLC separation. These  $\Delta^5$ -sterols were further separated by preparative HPLC into four fractions, referred to as fractions 1 to 4 in order of polarity beginning with the least polar. Each one of these fractions was separately analysed by GLC, analytical HPLC, mass spectrometry, and high resolution <sup>1</sup>H NMR. GLC and analytical HPLC of fraction 1 indicated that it was a mixture of two sterols: a major sterol, 1a  $(RR_t 1.28, \alpha_c)$ 0.69), accounting for 90% of the mixture, and a minor sterol, 1b  $(RR_t 1.40, \alpha_c 0.72)$ , accounting for only 10% of the mixture. Components 1a and 1b were identified on the basis of their mobilities [17, 18] as 24ξ-methylcholesta-5,25(27)-dien-3 $\beta$ -ol (codisterol or 24-epicodisterol) and 24 $\xi$ -ethylcholesta-5,22,25(27)-trien-3 $\beta$ -ol, respectively. The mass spectrum of the major component (1a) gave peaks at m/z (fragment, relative intensity in %): 398 [M]<sup>+</sup> (69), 383 [M - Me]<sup>+</sup> (31), 380 [M - H<sub>2</sub>O]<sup>+</sup> (21), 365 [M - Me - H<sub>2</sub>O]<sup>+</sup> (13), 314 [M - C<sub>6</sub>H<sub>12</sub>]<sup>+</sup> (46), 300 [M - C<sub>6</sub>H<sub>10</sub>O]<sup>+</sup> (31), 299 [M - C<sub>6</sub>H<sub>12</sub> - Me]<sup>+</sup> (38), 273 [M - SC]<sup>+</sup> (12), 273 [M - SC]<sup>+</sup> (13), 27  $[M - SC]^+$  (13), 272  $[M - SC - H]^+$  (33), 271  $[M - SC]^+$  $-2H_1^+$  (92), 255 [M - SC -  $H_2O_1^+$  (28), 253 [M - SC -  $H_2O_1^+$  (31), 231 [M - SC -  $C_3H_6_1^+$  (33), 229  $[M-SC-C_3H_8]^+$  (41),  $2\bar{1}3[M-SC-C_3\bar{H}_6-H_2O]^+$ 

<sup>\*</sup>Throughout this work, all  $\Delta^{25}$ -sterols are considered to have the 25(27)-designation rather than 25(26), because C-26 is taken to be derived from C-2 of MVA [1, 2, 9] and [2-14C]MVA does not label the methylene carbon atom in the case examined [23].

(100) and 211  $[M - SC - C_3H_8 - H_2O]^+$  (33) (SC = side chain). The molecular ion peak at m/z 398 and other peaks at m/z 314, 273, 271 and 255 indicate a di-unsaturated  $C_{28}$ -sterol, which is consistent with the above structural deduction of **1a** [19-22].

The final confirmation of the structure of sterol 1a was provided by <sup>1</sup>H NMR analysis. The spectrum showed signals at  $\delta 0.672$  (3H, s, H-18), 0.912 (3H, d, J = 6.4 Hz, H-21), 0.992 (3H, d, J = 6.5 Hz, H-28), 1.006 (3H, s, H-19), 1.637 (3H, s, H-26), 3.500 (1H, m, H-3), 4.662 (2H, br s, H-27), 5.359 (1H, br s, H-6). These signals are consistent with a 24-methyl,  $\Delta^{5,25(27)}$ -sterol structure, and were observed with this sterol when isolated from marine invertebrates [20, 21] or the siphonous marine algae of the genus Codium [19, 22]. The configuration at C-24 was established by comparing the spectrum of 1a with those of the two possible epimers, codisterol (24 $\beta$ ) and 24-epicodisterol (24 $\alpha$ ). Both these epimers are known to occur in the sponge Verongia cauliformis, and their <sup>1</sup>H NMR spectra have been documented [20]. The <sup>1</sup>H NMR spectrum in the present study did not show the olefinic methyl signal (H-26) at  $\delta$ 1.651 which corresponds to 24-epicodisterol [20]. Instead, the signal appeared at 1.637, which is characteristic of codisterol (24β-epimer) [19-22]. Thus, sterol 1a was pure  $24\beta$ -methylcholesta-5,25(27)-dien-3 $\beta$ ol (codisterol) without any of its 24α-epimer (24-epicodisterol). Codisterol is a rare sterol in nature, and to our knowledge this work represents the first instance of the detection of this sterol in a higher plant.

The mass spectrum of 1b, the minor component, showed a molecular ion peak at m/z 410 and other characteristic peaks at m/z 381  $[M-C_2H_5]^+$  and 300  $[M - C_7 H_{10} O]^+$ . All the other fragments were the same as those of the major component. This fragmentation pattern was consistent with that reported for this sterol in the literature [9, 23]. The <sup>1</sup>H NMR analysis showed signals distinctly different from those of the major component at  $\delta 0.693$  (3H, s, H-18), 0.833 (3H, t, J = 7.3 Hz, H-29) and 5.214 (2H, m, H-22, 23). The H-18, H-22 and H-23 signals agree with the presence of a  $\Delta^{22}$ -bond, and the H-29 triplet with the presence of a 24-ethyl structure [15, 24]. In addition to the H-18 and H-29 signals, the signals for H-21 (3H, d), H-26 (3H, s) and H-27 (2H, s) at  $\delta$ 1.02, 1.64 and 4.70, respectively, were also observed as shoulders on the major component signals. All the other signals were completely masked by those of the major component and thus were apparently at the same positions for both sterols. These observations are consistent with a 24-ethyl,  $\Delta^{5,22,25(27)}$ -structural deduction [9, 23, 25, 26]. The configuration at C-24 was deduced on the basis of the H-29 (t) signals, which showed a chemical shift identical to that of an authentic  $24\beta$ -epimer isolated earlier in this laboratory from Clerodendrum splendens [26]. Thus, sterol 1b was 24β-ethylcholesta-5,22,25(27)-trien-3 $\beta$ -ol, i.e. 25(27)-dehydroporiferasterol. As with codisterol (1a), 25(27)-dehydroporiferasterol (1b) is also a rare sterol in higher plants, and in addition to the present report in C. maxima 1b has so far been reported only in some Clerodendrum species [23, 25, 26], Kalanchoe diagremontiana [9] and Enhydra fluctuans [27].

GLC and analytical HPLC of fraction **2** as with fraction **1** also indicated the presence of two sterols, which were identified by their mobilities as  $24\xi$ -ethylcholesta-5,25(27)-dien-3 $\beta$ -ol (**2a**, 72.8%;  $RR_t$  1.57,  $\alpha_c$  0.88) and 24-ethylidenecholest-5-en-3 $\beta$ -ol (**2b**, 27.2%;  $RR_t$  1.67,  $\alpha_c$  0.90). The mass spectral analysis of the mixture gave major

peaks at m/z (fragment, relative intensity in %): 412 [M]<sup>+</sup> (49), 397  $[M - Me]^+$  (11), 394  $[M - H_2O]^+$  (9), 379  $[M - Me - H_2O]^+$  (18), 314  $[M - C_7H_{14}]^+$  (62), 300  $[M - C_7H_{12}O]^+$  (20), 299  $[M - C_7H_{14} - Me]^+$  (38), 273  $[M - SC]^+$  (14), 271  $[M - SC - 2H]^+$  (100), 255  $[M - SC]^+$  $-SC - \vec{H}_2O$ ]<sup>+</sup> (17), 253 [M - SC -  $\vec{H}_2O$  - 2H]<sup>+</sup> (25), 231  $[M-SC-C_3H_6]^+$  (29), 229  $[M-SC-C_3H_8]^+$  (54), 213  $[M-SC-C_3H_6-H_2O]^+$  (66) and 211  $[M-SC-C_3H_8-H_2O]^+$  (42). As expected, both **2a** and **2b** produced a very similar fragmentation pattern, which is in agreement with the literature [9, 13, 28, 29]. On the other hand, the 1H NMR signals for each of the two sterols were quite distinct and could be assigned to the respective sterols on the basis of their relative amounts. The signals for sterol **2a** appeared at  $\delta$ 0.671 (3H, s, H-18), 0.800 (3H, t, J = 7.3 Hz, H-29), 0.905 (3H, d, J = 6.6 Hz, H-21), 1.007 (3H, s, H-19), 1.566 (3H, s, H-26), 3.495 (1H, m, H-3), 4.643 and 4.724 (2H, s, H-27, terminal methylene protons resonating at two distinct  $\delta$  values) and 5.356 (1H, br s, H-6). These signals are consistent with the structural deduction of 2a and were also obtained with this sterol previously isolated from Kalanchoe daigremontiana in our laboratory [9]. The C-24 configuration of the K. daigremontiana sterol was demonstrated to be  $24\beta$ -ethyl by comparing the <sup>1</sup>H NMR spectrum of its 25(27)-dihydro derivative with those of the corresponding 24x- (sitosterol) and  $24\beta$ - (clionasterol) epimers [9]. Since the <sup>1</sup>H NMR spectrum of 2a from the present work was identical to that of the above-mentioned K. daigremontiana sterol, the C-24 configuration of 2a was also deduced to be  $24\beta$ -ethyl. Thus, sterol **2a** was identified as  $24\beta$ -ethylcholesta-5,25(27)-dien-3 $\beta$ -ol (clerosterol). This is a rare sterol in higher plants, and to our knowledge, in addition to K. daigremontiana [9], C. maxima is only the second higher plant species in which the occurrence of 2a has been supported by NMR (for configuration). However, a 24-ethyl- $\Delta^{5.25(27)}$ -sterol, the C-24 $\beta$  configuration of which is yet to be established, has been found also in Clerodendrum infortunatum [16], Calendula officinalis [30], Enhydra fluctuans [31] and Brassica napus [29].

The <sup>1</sup>H NMR signals for sterol **2b** appeared at  $\delta$ 0.682 (3H, s, H-18), 0.947 (3H, d, J = 6.3 Hz, H-21), 0.976 (6H, 2d, J = 6.9 Hz, H-26 and H-27), 1.007 (3H, s, H-19), 2.819 (1H, m, H-25), 3.495 (1H, m, H-3), 5.115 (1H, m, H-28) and 5.356 (1H, br s, H-6). The H-29 signal (d at ca 1.58) was not clearly detected as it was masked by the H-26 signal for the major component (**2a**). These signals were consistent with the literature [28, 32], and matched those of an authentic sample of isofucosterol. The configuration of the  $\Delta$ 2.4(28)-bond was assigned on the basis of the signal at  $\delta$ 2.819 for the H-25 multiplet. This is the position for the Z-isomer [28, 32], whereas in the case of E-isomer this signal appears distinctly upfield at  $\delta$ 2.22 [32]. Thus, sterol **2b** was 24Z-ethylidenecholest-7-en-3 $\beta$ -ol (isofucosterol).

As with the first two fractions, GLC and analytical HPLC of fraction 3 indicated that two sterols were present. These were identified as  $24\xi$ -ethylcholesta-5,22-dien-3 $\beta$ -ol (3a, stigmasterol or poriferasterol, 65.65°,  $RR_t$ , 1.42,  $\alpha_c$ , 1.09) and  $24\xi$ -methylcholest-5-en-3 $\beta$ -ol (3b, campesterol or 22-dihydrobrassicasterol, 34.35°,  $RR_t$ , 1.30,  $RR_t$ , 1.31). The major component (3a) upon mass spectrometry showed peaks at m/z (fragment, relative intensity in °0): 412 [M]+ (42), 394 [M-H<sub>2</sub>O]+ (8), 379 [M-Me-H<sub>2</sub>O]+ (25), 369 [M-C<sub>3</sub>H<sub>7</sub>]+ (8), 327 [M-85]+ (8), 314 [M-98]+ (31), 300 [M-C<sub>7</sub>H<sub>12</sub>O]+ (39), 299 [M-113]+ (22), 273 [M-SC]+ (39), 272

 $\begin{bmatrix} M-SC-H \end{bmatrix}^+ & (32), \ 271 \ \begin{bmatrix} M-SC-2H \end{bmatrix}^+ & (56), \ 255 \\ \begin{bmatrix} M-SC-H_2O \end{bmatrix}^+ & (95), \ 253 \ \begin{bmatrix} M-SC-H_2O-2H \end{bmatrix}^+ \\ (24), \ 231 \ \begin{bmatrix} M-SC-C_3H_6 \end{bmatrix}^+ & (51), \ 229 \ \begin{bmatrix} M-SC-C_3H_8 \end{bmatrix}^+ & (51), \ 211 \ \begin{bmatrix} M-SC-C_3H_8-H_2O \end{bmatrix}^+ & (22). \ 229 \ \begin{bmatrix} M-SC-C_3H_8 \end{bmatrix}^+ & (22). \ 229 \ \begin{bmatrix} M-SC-C_3H_8 \end{bmatrix}^+ & (23) \ 230 \ \begin{bmatrix} M-M_2 \end{bmatrix}^+ & (23) \ 230 \ \begin{bmatrix} M-M_2 \end{bmatrix}^+ & (23) \ 230 \ \begin{bmatrix} M-M_2O \end{bmatrix}^+ & (24), \ 315 \ \begin{bmatrix} M-H_2O \end{bmatrix}^+ & (25), \ 367 \ \begin{bmatrix} M-M_2O \end{bmatrix}^+ & (26), \ 367 \ \begin{bmatrix} M-M_2O \end{bmatrix}^+ & (26), \ 315 \ \begin{bmatrix} M-85 \end{bmatrix}^+ & (26), \ 310 \ \end{bmatrix}$  were the same as those for 3a. These fragmentation patterns are in agreement with those described for 3a and 3b in the literature [9].

The final confirmation of the structures of sterols 3a and 3b was provided by  $^1H$  NMR analysis. In each case, the spectrum of the natural sterol was compared with those of the authentic samples of the two possible C-24 epimers  $(24\alpha$  and  $24\beta$ ). The data (Table 1) clearly show that the spectrum of 3a is practically identical to that of authentic stigmasterol  $(24\alpha$ -epimer) and is markedly different from that of authentic poriferasterol  $(24\beta$ -epimer). The chemical shifts of the epimeric standards were distinctly different in the H-29 methyl proton region and agree with those reported in the literature [15, 24].

Component 3a was therefore identified as  $24\alpha$ -ethylcholesta-5,22-dien-3 $\beta$ -ol (stigmasterol). The signals for component 3b (Table 2) are also identical to those of its corresponding  $24\alpha$ -epimer (campesterol) and are significantly different from those of the corresponding  $24\beta$ -epimer (22-dihydrobrassicasterol) [9, 15, 24]. However, a slight shoulder distinctly downfield (ca 0.01 ppm) from that of the  $24\alpha$ -epimer in the H-21 methyl proton region, indicated that the  $24\beta$ -epimer may be present as a very minor component. Thus, sterol 3b was mostly  $24\alpha$ -methylcholest-5-en-3 $\beta$ -ol (campesterol), with the copresence of a very small amount of  $24\beta$ -epimer (22-dihydrobrassicasterol).

In contrast to the first three fractions from the preparative HPLC, fraction 4 was shown to contain only one sterol by GLC and analytical HPLC, with an  $RR_t$  of 1.62 and  $\alpha_c$  of 1.26. This component was identified as  $24\xi$ -ethylcholest-5-en-3 $\beta$ -ol (sitosterol or clionasterol). The mass spectral analysis gave major peaks at m/z (fragment, relative intensity in %): 414 [M]<sup>+</sup> (78), 399 [M - Me]<sup>+</sup> (30), 396 [M - H<sub>2</sub>O]<sup>+</sup> (36), 381 [M - Me - H<sub>2</sub>O]<sup>+</sup> (26), 329 [M - 85]<sup>+</sup> (45), 303 [M - C<sub>7</sub>H<sub>11</sub>O]<sup>+</sup> (55), 273 [M - SC]<sup>+</sup> (36), 272 [M - SC - H]<sup>+</sup> (5), 271

Table 1. <sup>1</sup>H NMR chemical shifts ( $\delta$ ) of C<sub>29</sub>- $\Delta$ <sup>5,22</sup>-sterols\*

Proton	Sterol 3a (from C. maxima seeds)	Stigmasterol†	Poriferasterol‡	
H-18 (3H, s)	0.698	0.698	0.697	
H-19 (3H, s)	1.011	1.012	1.011	
H-21 $(3H, d)$	1.022 (6.5)	1.022 (6.5)	1.024 (6.5)	
H-26 or 27 (3H, d)	0.795 (6.6)	0.795 (6.6)	0.791 (6.7)	
H-26 or 27 (3H, d)	0.846 (6.5)	0.846 (6.5)	0.843 (6.5)	
H-29 (3H, t)	0.805 (7.1)	0.804 (7.1)	0.810 (7.2)	
H-3 (1H, m)	3.504	3.505	3.505	
H-6 (1H, br s)	5.359	5.357	5.356	
H-22 or 23 (1H, dd)	5.012 (ca 7.5)	5.015 (ca 7.5)	5.016 (ca 7.5	
H-22 or 23 (1H, dd)	5.156 (ca 7.5)	5.159 (ca 7.5)	5.162 (ca 7.5	

<sup>\*</sup>Values in parentheses are the coupling constants (J, in Hz).

Table 2. <sup>1</sup>H NMR chemical shifts (δ) of C<sub>28</sub>-Δ<sup>5</sup>-sterols\*

Proton	Sterol 3b (from C. maxima seeds)	Campesterol†	22-Dihydro- brassicasterol‡	
H-18 (3H, s)	0.680	0.680	0.680	
H-19 (3H, s)	1.011	1.011	1.010	
H-21 $(3H, d)$	0.911 (6.1)	0.910 (6.1)	0.921 (6.2)	
H-26 or 27 (3H, d)	0.800 (6.0)	0.800 (6.0)	0.784 (6.1)	
H-26 or 27 (3H, d)	0.851 (6.0)	0.850 (6.1)	0.862 (6.1)	
H-28(3H, d)	0.771 (6.0)	0.771 (6.0)	0.774 (6.0)	
H-3 (1H, m)	3.504	n.m.	n.m.	
H-6 (1H, $br$ s)	5.359	n.m.	n.m.	

<sup>\*</sup>Values in parentheses are the coupling constants (*J*, in Hz); the spectra of campesterol and 22-dihydrobrassicasterol were recorded at 220 MHz, and that of sterol 3b at 360 MHz; n.m. = not measured.

<sup>†</sup>Obtained from a commercial source.

<sup>‡</sup>Isolated from Chlorella ellipsoidea.

<sup>†</sup>Obtained from a commercial source.

<sup>‡</sup>Prepared from ergosterol.

[M-SC-2H]<sup>+</sup> (4), 255 [M-SC-H<sub>2</sub>O]<sup>+</sup> (46), 253 [M-SC-H<sub>2</sub>O-2H]<sup>+</sup> (3), 231 [M-SC-C<sub>3</sub>H<sub>6</sub>]<sup>+</sup> (50), 229 [M-SC-C<sub>3</sub>H<sub>8</sub>]<sup>+</sup> (20), 213 [M-SC-C<sub>3</sub>H<sub>6</sub>-H<sub>2</sub>O]<sup>+</sup> (100) and 211 [M-SC-C<sub>3</sub>H<sub>8</sub>-H<sub>2</sub>O]<sup>+</sup> (6). This fragmentation was similar to those of authentic sitosterol and clionasterol, and agreed with the literature [9]. The final proof of the structure of 4 was again provided by <sup>1</sup>H NMR analysis. When compared with the spectra of the authentic epimers (Table 3), the spectrum of 4 was found to be identical to that of the corresponding  $24\alpha$ -epimer (sitosterol), but was markedly different from that of the  $24\beta$ -epimer (clionasterol). The signals for protons at H-21 and H-29 were diagnostic for the two epimers [9, 15, 24]. Fraction 4 was therefore identified as  $24\alpha$ -ethylcholest-5-en-3 $\beta$ -ol (sitosterol).

In all, seven  $\Delta^5$ -4-desmethylsterols were identified in the seeds of *C. maxima*. The absolute amounts and percentages of each of these sterols are given in Table 4. This work represents the first time the structures and configurations of the  $\Delta^5$ -sterols from a Cucurbitaceae species have been clearly characterized. It is also the first report of the presence of  $\Delta^5$ -sterols in *C. maxima*.

report of the presence of  $\Delta^5$ -sterols in *C. maxima*. When compared with the  $\Delta^7$ -fraction [33], it is interesting to note that the structures of all seven  $\Delta^5$ -sterols were analogous to those of the  $\Delta^7$ -sterols. This would indicate that structurally similar sterols are being synthesized in

both the  $\Delta^5$ - and  $\Delta^7$ -series, and thus very similar biosynthetic pathways seem to exist. Although the biosynthetic route to  $\Delta^5$ -sterols is usually believed to operate from  $\Delta^7$ - to  $\Delta^{5,7}$ - to  $\Delta^5$ -sterols [1, 2], in the present study HPLC of the 4-desmethyl mixture using a 282 nm monitor failed to reveal the presence of any  $\Delta^{5,7}$ intermediates. In contrast to the structural similarity, the relative amounts of the various  $\Delta^5$ -sterols (Table 4) were markedly different from those of their corresponding  $\Delta^7$ analogues [33]. The unusual  $24\beta$ -methyl- $\Delta^{5.25(27)}$ -sterol, codisterol, was the most abundant (35.3 % of the total)  $\Delta^{5}$ sterol, whereas its  $\Delta^7$ -analogue, 25(27)-dehydrofungisterol, was present in the  $\Delta^7$ -fraction only in trace amounts [33]. Conversely, the  $\Delta^7$ -analogue of the minor  $\Delta^5$ -sterol, 25(27)-dehydroporiferasterol, was one of the major components in the  $\Delta^7$ -series [33]. Furthermore, it is interesting to note that in the  $\Delta^7$ -series [33], the 24methyl-sterols (24ξ-methyllathosterol and 25(27)-dehydrofungisterol) constituted a much smaller part of the mixture than did the corresponding analogues (campesterol and codisterol) in the  $\Delta^5$ -series. This difference has also been observed in other plants [1, 2] and thus seems to represent a general pattern. In addition, it has also been commonly noted in the literature that in the  $\Delta^7$ -series the percentage of the  $\Delta^{22}$ -24 $\alpha$ -ethyl-sterol (spinasterol) is higher than is usually found for its  $\Delta^5$ -analogue (stigmas-

Table 3. <sup>1</sup>H NMR chemical shifts ( $\delta$ ) of C<sub>29</sub>- $\Delta$ <sup>5</sup>-sterols\*

Proton	Sterol 4 (from C. maxima seeds)	Sitosterol†	Clionasterol‡
H-18 (3H, s)	0.680	0.680	0.680
H-19 (3H, s)	1.009	1.010	1.011
H-21 $(3H, d)$	0.921 (6.6)	0.921 (6.5)	0.924 (6.5)
H-26 or 27 (3H, d)	0.813 (6.5)	0.814 (6.5)	0.810 (6.5)
H-26 or 27 (3H, d)	0.834 (6.5)	0.833 (6.5)	0.827 (6.5)
H-29 $(3H, t)$	0.845 (7.3)	0.845 (7.3)	0.855 (7.3)
H-3 (1H, m)	3.504	3.502	n.m.
H-6 (1H, br s)	5.358	5.358	n.m.

<sup>\*</sup>Values in parentheses are the coupling constants (J, in Hz); the spectrum of clionasterol was recorded at 220 MHz, and the others at 360 MHz, n.m. = not measured

Table 4. The  $\Delta^5$ -4-desmethylsterol composition of C. maxima seeds

Sterol	Amount (mg/100 g of seeds)	% of total
$24\beta$ -Ethylcholesta-5,22,25(27)-trien-3β-ol (1b)		
(25(27)-dehydroporiferasterol)	0.31	4.3
$24\beta$ -Ethylcholesta-5,25(27)-dien-3 $\beta$ -ol (2a) (clerosterol)	1.43	19.8
24Z-Ethylidenecholest-5-en-3β-ol ( <b>2b</b> ) (isofucosterol)	0.54	7.4
24 $\alpha$ -Ethylcholesta-5,22-dien-3 $\beta$ -ol (3a) (stigmasterol)	0.79	10.9
24 $\alpha$ -Ethylcholest-5-en-3 $\beta$ -ol (4) (sitosterol)	1.20	16.6
24α-Methylcholest-5-en-3β-ol (3b) (campesterol)*	0.41	5.7
$24\beta$ -Methylcholesta-5,25(27)-dien-3 $\beta$ -ol (1a) (codisterol)	2.56	35.3

<sup>\*</sup>A minute amount of the  $24\beta$ -epimer, 22-dihydrobrassicasterol, was also present.

<sup>†</sup>Obtained from a commercial source.

<sup>‡</sup>Isolated from Chlorella ellipsoidea.

terol) [1, 2]. A similar pattern can be observed in C. maxima. In the  $\Delta^7$ -series, spinasterol was the most abundant sterol (32.0% of the total) [33], while its  $\Delta^5$ -analogue, stigmasterol, is considerably less in abundance (10.9% of the total; Table 4). Based on these observations, it seems likely that such distinct patterns in the relative ratios (of the various  $\Delta^5$ -sterols) may be a reflection of their functional significance.

Finally, it should be pointed out that as with the  $\Delta^7$ sterols [33], the  $\Delta^5$ -fraction of C. maxima also contains several unusual higher plant sterols which occur in addition to the usual ones of the 'main line' series, Of particular interest is the presence of a large amount of codisterol (1a); this has hitherto been found only in some marine invertebrates [20, 21] and the siphonous marine algae of the genus Codium [19, 22]. In fact this is the first time that such a high concentration of codisterol has been reported in any organism. While 1a probably plays a definite role in C. maxima, the precise function must remain an open question. The other two unusual higher plant sterols were 25(27)-dehydroporiferasterol (1b) and clerosterol (2a). They have previously been reported to occur only in a few plant species [9, 23, 25-27, 29-31]. Thus, together with the  $\Delta^7$ -sterols [33], the 4-desmethylsterols of C. maxima seeds represent an unusual array of sterols. This further strengthens the earlier suggestion [1, 2, 34] that the plants of the family Cucurbitaceae may show an evolutionary transition in terms of sterols.

## **EXPERIMENTAL**

Squash (Cucurbita maxima cv True Hubbard) seeds were obtained from W. Atlee Burpee, Co., U.S.A. Specialized standards were obtained and purified from the following sources: stigmasterol, campesterol and sitosterol from Applied Science Laboratories; poriferasterol and clionasterol from Chlorella ellipsoidea (courtesy of G. W. Patterson); and 22-dihydro-brassicasterol was prepared from ergosterol [35].

The 4-desmethylsterols were extracted and separated from the neutral lipids as previously described [33]. The  $\Delta^5$ - and  $\Delta^7$ mixtures were separated from each other by TLC. The plates (silica gel G coated; 0.25 and 1.00 mm thick) were developed 4 × in Et<sub>2</sub>O-C<sub>6</sub>H<sub>6</sub> (1:9) and the separated components were visualized under UV light after being sprayed with a 0.05 % soln of rhodamine 6G in Me<sub>2</sub>CO. Two bands, which were completely separated, were visualized, co-chromatographing with the standards containing authentic  $\Delta^7$ - (lathosterol-spinasterol mixture) and  $\Delta^5$ - (cholesterol-sitosterol mixture) sterols, respectively. The faster moving  $\Delta^5$ -region was separated into glass tubes and the sterols were extracted with several washes of Et2O. The sterol mixture thus obtained was analysed by GLC and HPLC, and finally separated into various components by prep. HPLC. The instrumentation and the techniques for the GLC and the HPLC have been described in ref. [33]. The term RR, is used for GLC and a for HPLC.

Mass spectra were obtained by direct probe (EIMS, ionizing energy 70 eV) on a Finnigan Model 4000 instrument with a series 6000 data system. <sup>1</sup>H NMR spectroscopy was performed at 360 MHz at ambient temp. on a Bruker instrument, Model WH-360, in CDCl<sub>3</sub> with TMS as internal standard. The <sup>1</sup>H NMR spectroscopy of some of the standards was performed at 220 MHz.

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