Communications to the Editor

Chem. Pharm. Bull. 33(3)1323—1326(1985)

STEREOSTRUCTURE OF CURDIONE

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The stereostructure of curdione (1) isolated from <u>Curcuma wenyujin</u> together with curcumol (2a) was determined as <u>la</u> on the basis of its direct X-ray crystallography and transformation to <u>2a</u>. The preferred conformation of <u>la</u> in solution was found to be <u>CC/syn</u> (C(5)=O/C(1)-H) (3a) rather than <u>CC/anti</u> (3b) as in crystal based on ¹H-NMR variable-temperature and NOE measurements, especially at -70°C. A possible biogenetic pathway from <u>la</u> through conformational change (3a to <u>3b</u>) to <u>2a</u> and the absolute configuration were proposed.

KEYWORDS——curdione; germacranoid; Curcuma wenyujin; Zingiberaceae; stereostructure; $^{1}\text{H-NMR}$ (400 MHz); variable-temperature; NOE; X-ray diffraction

The crystal data are: $C_{15}H_{24}O_2$, MW 236, orthorhombic, space group $P2_1^2_1^2_1$, lattice constant a = 10.758(5), b = 13.164(7), c = 10.176(5), U = 1441.1 Å³, z = 4, D_{calc} = 1.0895 gm⁻³. The intensity data were collected on a Philips PW 1100

diffractometer using graphite monochromated CuK α radiation. 1229 Reflections up to 155° (20) were used. The structure was solved by the direct method using a MULTAN⁵) program and refined by the method of least-squares with block-diagonal matrix approximations. The final R value was 0.05 including 24 hydrogen atoms with isotropic temperature factors. The molecular structure in crystal and the relative stereochemistry of 1 thus depicted in Fig. 1 reveal that curdione (1) exists in a CC conformation⁶ possessing a trans relationship between C(4)-methyl and C(7)-isopropyl side chain. Moreover, this conformation is characteristic of the syn-arrangement⁷⁾ of C(5)=0 and C(1)-H <u>i.e.</u> oriented on the same side of the plane put through the ten-member ring.

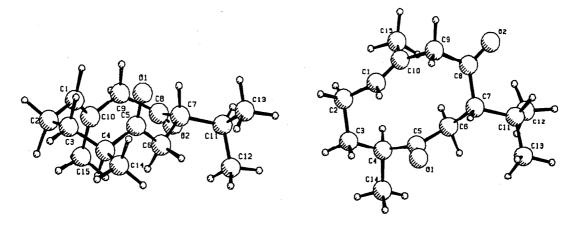


Fig. 1 Perspective Views of Curdione (la)

An examination of Dreiding models shows that two major conformations are possible for 1. In order to verify the co-existence of more than two diverse conformers, variable-temperature studies of $400~\text{MHz}^{-1}\text{H}-\text{NMR}$ spectroscopy of 1 were performed. The spectrum showed a set of well-separated peaks at room temperature in spite of broad signals at -30°C . At the temperatures less than -50°C , especially at -70°C , two sets of signals with an intensity ratio of approximately 5:1 (e.g. C(1)-H 5.06 and 5.63; $C(10)-\text{CH}_3$ 1.76 and 1.48 ppm) were observed. This indicates that each of the signals arose from the two CC conformers, i.e. conformation A (3a) and conformation B (3b), the latter of which takes the anti-arrangement of C(5)=0~and~C(1)-H. For the determination of the stereostructures of each conformer, intramolecular nuclear Overhauser effect (NOE) measurements were carried out at -70°C . When the frequency corresponding to C(1)-H of the major isomer was sat-

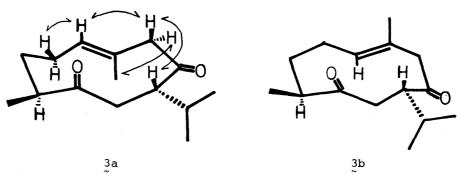


Fig. 2 Conformation A (3a) and B (3b) of Curdione (la)

urated, 6% of NOE was observed both in C(9)- βH and C(2)- βH . Irradiation of the frequency at C(10)- CH_3 was effected by 3% C(9)- αH , but not C(9)- βH . When C(9)- βH was irradiated, 6% of NOE appeared both in C(1)-H and C(7)- βH (Fig. 2). Due to the anisotropic effect of both carbonyl groups in 1, C(1)-H signal of the major conformer (δ 5.06) appears upward in comparison with the corresponding proton of the minor one (δ 5.63). On the contrary, C(10)- CH_3 of the former is observed at the lower field (δ 1.76) rather than that of the latter (δ 1.48). These results clearly suggest that the predominant conformer and the minority should be represented as conformation A (3a)(C(5)=O/C(10)- CH_3 : anti) and conformation B (3b)(C(5)=O/C(10)- CH_3 : syn), respectively, (Fig. 2). Furthermore, as the dominant spectrum of each set of signals measured at -70°C was closely similar to that observed at ambient temperature, 3a proves to be the preferred conformation in solution at room temperature as well as in the crystals shown in Fig. 1.

Since 2a was exclusively produced only by heating 1a at 200°C in ethanol, 9) it seems to be suggested that curdione (1a) mainly existing in conformation A (3a) could transform via conformation B (3b) to curcumol (2a) with retention of C(7)-configuration, not to the hypothetical C(7)- β -epimer (2b) 10 in a possible biogenetic pathway. As the absolute configuration of curcumol 3 has been established as 2a possessing C(7)- α -isopropyl side chain, which is more stable than 2b with C(7)- β -substituent, 10 the absolute stereostructure of curdione (1) should be presumably represented as 1a, rather than its mirror image (1b), providing that the transformation from 1a to 2a proceeded without epimerization.

Further studies obtaining the concrete proof about this point are now in progress.

ACKNOWLEDGEMENTS One of the authors (S.I.) acknowledges partial financial support from the Miyata academic prize for this work. The authors are grateful to Dr. H. Seto of the Applied Microbiological Institute of Tokyo University and Mr. Y. Watanabe of JEOL Co., Ltd. for measurements of 400 MHz NMR spectrum of curdione. Thanks are also due to Mr. H. Yamanaka and Miss S. Takei of the Joint Laboratory, School of Medicine, Keio University, for MS and IR spectral measurements.

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(Received February 9, 1985)