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## X-Ray Crystal and Molecular Structure of Kodo-cytochalasin-1

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**Summary** Kodo-cytochalasin-1 and cytochalasin H have been shown to be identical in structure and stereochemistry by single-crystal X-ray diffraction analyses; an inadvertent error in the stereochemistry at C(18) reported previously for cytochalasin H is corrected.

*Phomopsis paspalli*, Pendse and Kanitkar, a fungal pathogen of kodo millet (*Paspalum scrobiculatum* Linn. Syn. *P. commerionii* Lam), a minor food grain crop of India, elaborates two major toxins. These compounds, essentially on the basis of spectral data, have been assigned gross structures (1) and (2), and have been designated kodo-cytochalasin-1 and kodo-cytochalasin-2, respectively.<sup>1</sup> Kodo-cytochalasin-1 is obtained as white needles, m.p. 258—263 °C, from a CHCl<sub>3</sub>-Et<sub>2</sub>O solution. The present work confirms these gross structures and establishes the stereochemistry as depicted in (3) for kodo-cytochalasin-1. Further, we have

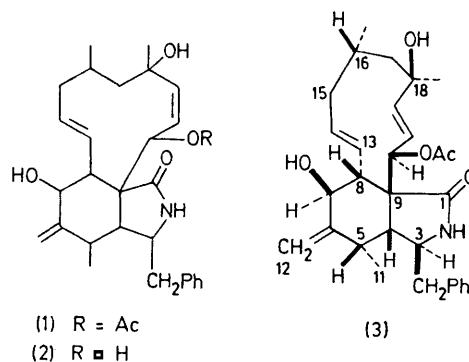
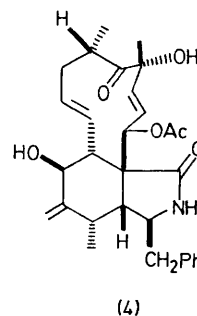


TABLE	
Unit cell constants	
Kodo-cytochalasin-1	Cytochalasin H (ref. 3)
a 15.348(4)	c 15.330(4)
b 13.089(3)	b 13.053(6)
c 7.347(3)	a 7.338(2)
$\beta$ 97.23(3)°	$\beta$ 97.02(2)°
$\lambda(\text{Cu-K}\alpha)$ 1.54178 Å	$\lambda(\text{Mo-K}\alpha)$ 0.71069 Å

demonstrated by independent crystal structure investigations that kodo-cytochalasin-1 and a fungal toxin recently



isolated from *Phomopsis* sp. infected pecans,<sup>2</sup> described as cytochalasin H,<sup>3</sup> are in fact identical. Some uncertainty on this point was caused by a clerical error in the formulation of the stereochemistry of cytochalasin H at C(18) in the structural formula (I) in ref. 3.

The full name of this toxin using the nomenclature proposed by Tamm *et al.*,<sup>4a</sup> is (7*S*,16*S*,18*S*,21*R*)-21-acetoxy-7,18-dihydroxy-16,18-dimethyl-10-phenyl[11]cytochalasa-

gations was found to be  $P2_1$  with two molecules of  $C_{30}H_{39}NO_5$  per unit cell. The structure of kodo-cytochalasin-1 was solved using X-ray crystallographic direct methods.<sup>6</sup> The co-ordinates of all atoms were refined by full-matrix least-squares methods, with all non-hydrogen atoms having anisotropic thermal parameters, to an *R*-factor of 0.043 for 1945 non-zero reflections collected on a Syntex  $P2_1$  diffractometer. A drawing of a molecule of kodo-cytochalasin-

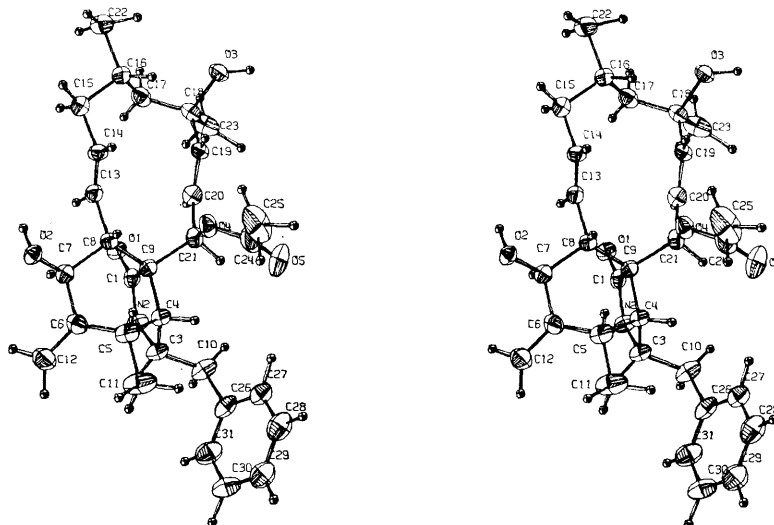


FIGURE. Stereoscopic view of the structure of a molecule of kodo-cytochalasin-1 (3).

6(12),13<sup>t</sup>,19<sup>t</sup>-triene-1-one.<sup>4b</sup> It is of interest that the relative stereochemistry at C(18) in this compound is different from that established for the same functionality in other closely related cytochalasins, *e.g.* zygospurin-A (4).<sup>5</sup>

The cell constants for kodo-cytochalasin-1, determined using a small, colourless, irregularly shaped crystal 0.3 mm in largest dimension, are compared with those for cytochalasin H in the Table. The space group in both investi-

1 is shown in the Figure. The structure determined from this X-ray work and of that reported in ref. 3 for cytochalasin H conform to that depicted in (3).

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<sup>4</sup> (a) M. Binder, C. Tamm, W. B. Turner, and H. Minato, *J.C.S. Perkin I*, 1973, 1146; (b) note should be taken that the name in ref. 3 was incorrect, reflecting the improper labelling of the ORTEP figure in that report. Full details of that work will appear elsewhere: M. A. Beno, R. H. Cox, J. M. Wells, R. J. Cole, J. W. Kirksey and G. G. Christoph, *J. Amer. Chem. Soc.*, submitted for publication.

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