

Tremellin, a Novel Symmetrical Compound, from the Basidiomycete *Tremella aurantialba*

by Ding Zhi-Hui, Li Jing-Ping, and Liu Ji-Kai*

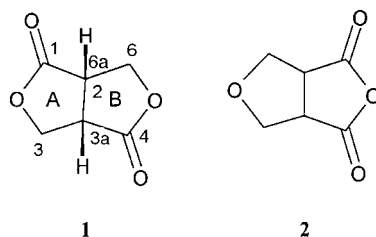
Kunming Institute of Botany, Chinese Academy of Sciences, Kunming 650204, P.R. China
(e-mail: jkliu@mail.kib.ac.cn)

and Lu Yang, Wang Cheng, and Zheng Qi-Tai

Institute of Materia Medica, Chinese Academy of Medical Sciences, Beijing 100050, P.R. China

A novel, highly symmetrical compound, named tremellin (**1**), was isolated from the fruiting bodies of the basidiomycete *Tremella aurantialba*. Its structure was established by spectroscopic means and X-ray analysis.

Introduction. – *Tremella aurantialba* is one of the edible mushrooms that has been used as antihepatitis agent and immunostimulant [1]. The fungus is distributed in Yunnan, Sichuan, Xizang, Gansu, and Jiangxi provinces of China [2]. As part of our studies on the active metabolites from higher fungi [3–9], the chemical constituents of *Tremella aurantialba* were investigated. The present report deals with the structure elucidation of a new compound **1**, named tremellin, which was isolated from the fruiting bodies of this fungus. Tremellin (**1**) has a simple, highly symmetric structure.



Results and Discussion. – *T. aurantialba* (dry weight 3.5 kg) was extracted with $\text{CHCl}_3/\text{MeOH}$ 1:1. Repeated chromatography afforded tremellin (**1**; 10 mg) as colorless needles with a molecular formula of $\text{C}_6\text{H}_6\text{O}_4$ (HR-MS: m/z 142.0264 (M^+ , $\text{C}_6\text{H}_6\text{O}_4^+$; calc. 142.0266). The ^1H - and ^{13}C -NMR and IR data established its structure as tetrahydro-1*H*,4*H*-furo[3,4-*c*]furan-1,4-dione (**1**).

Three signals in the ^{13}C -NMR (DEPT) spectrum of **1** were recognized (1 C, 1 CH_2 , 1 CH), which were assigned to a carbonyl (δ 175.5), an oxymethylene (δ 68.9), and a methine group (δ 40.7). The three signals in the ^1H -NMR spectrum at δ (H) 4.67 (*d*, $J=9.7$ Hz, $\text{H}_\alpha\text{-C}(3)$, $\text{H}_\alpha\text{-C}(6)$), 4.07 (*m*, $\text{H}_\beta\text{-C}(3)$, $\text{H}_\beta\text{-C}(6)$), and 3.51 (*m*, $\text{H-C}(3\text{a})$, $\text{H-C}(6\text{a})$) revealed a highly symmetrical structure and led to the deduction of two possible structures **1** and **2**. In the IR spectrum of tremellin, the carbonyl absorption at 1765 cm^{-1} indicated the presence of a γ -lactone moiety, and the fragment ion m/z 98 ($[M - \text{CO}_2]^+$) in the EI-MS confirmed the presence of this γ -lactone unit.

The relative configuration of tremellin (**1**) was established by a single-crystal X-ray analysis (Figs. 1 and 2).

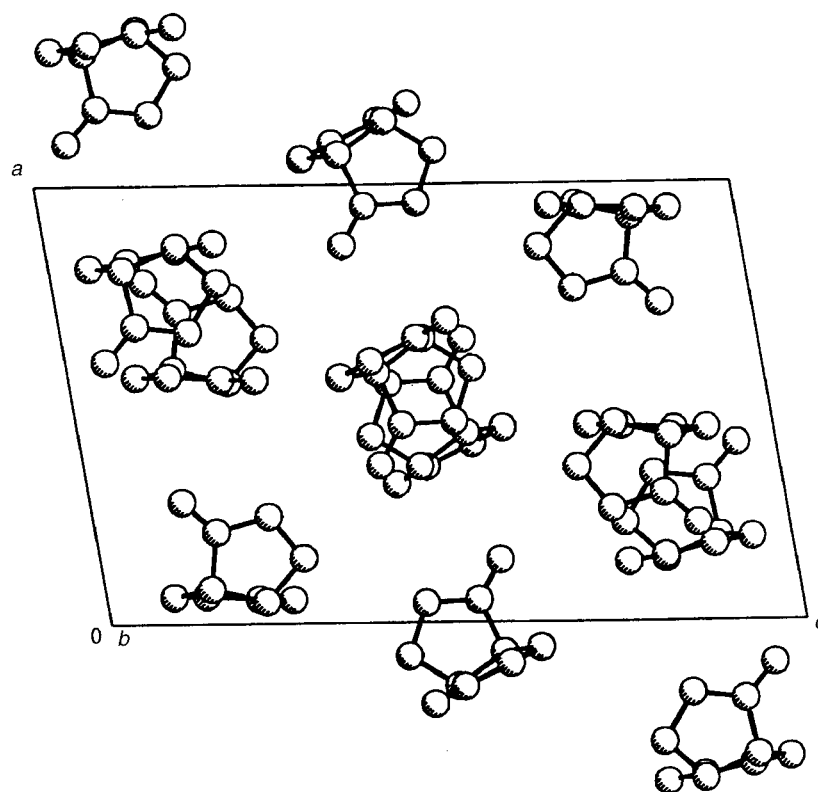


Fig. 1. Perspective views of the molecules in the unit cell along the *b* axis

Experimental Part

General. M.p.: uncorrected. IR: KBr pellets; in cm^{-1} . ^1H - and ^{13}C -NMR: Bruker DRX-500 spectrometers, δ in ppm, J in Hz. MS. VG Autospect-3000 spectrometer; m/z (rel. int).

Mushroom Material. The fruiting bodies of the basidiomycete *Tremella aurantialba* were provided by the Kunming Institute of Edible Mushroom.

Extraction and Isolation. The entire fruiting bodies of *Tremella aurantialba* (dry weight 3.5 kg) were extracted with $\text{CHCl}_3/\text{MeOH}$ 1:1 at r.t. (4 times). The residue was first extracted with petroleum ether and then with AcOEt. The AcOEt extract (36.5 g) was submitted to column chromatography (silica gel, gradient $\text{CHCl}_3/\text{AcOEt}$ 10:0, 9:1, 8:2). The combined fractions ($\text{CHCl}_3/\text{AcOEt}$ 9:1) were purified by recrystallization from petroleum ether/ Et_2O to give pure tremellin (**1**, 10 mg).

Tremellin (=rel-(3*a*R,6*a*R)-Tetrahydro-1*H*,4*H*-furo[3,4-*c*]furan-1,4-dione; **1**). Colorless crystals. M.p. 130–133° (petroleum ether/ Et_2O). IR (KBr): 3514, 2985, 1765, 1477, 1370, 1299, 1176. ^1H -NMR (CDCl_3): 3.51 (*m*, 2 H); 4.54 (*m*, 2 H); 4.67 (*d*, $J = 9.7, 2$ H). ^{13}C -NMR (CDCl_3): 40.7 (CH_2); 68.9 (CH); 175.5 (C=O). HR-EI-MS: 142.0264 ($\text{C}_6\text{H}_6\text{O}_4^+$, M^+ ; calc. 142.0266). EI-MS: 142 (48), 98 (23), 84 (34), 69 (100), 55 (72), 54 (76).

X-Ray Analysis. Crystal data: $\text{C}_6\text{H}_6\text{O}_4$, M 142, monoclinic, space group $P2_1/a$; $a = 11.3010(11)$, $b = 9.1700(10)$, $c = 17.6090(21)$ Å, $\beta = 99.785(7)^\circ$, $V = 1798.3(3)$ Å³, $Z = 12$. Final R_f and R_w values were 0.064 and 0.052, resp. A total of 2557 reflections were recorded in the ω scanning mode with a MAC-DIP-2030K

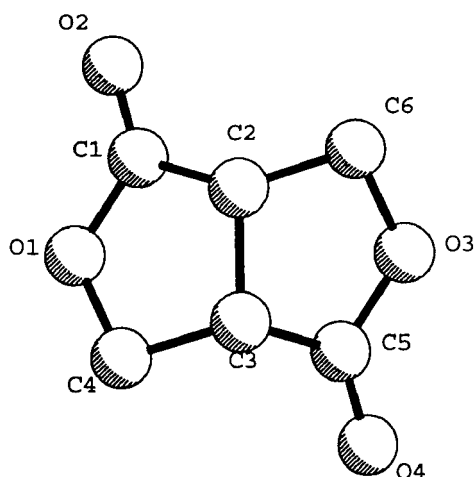


Fig. 2. Perspective view of tremellin (1) from the X-ray analysis. Arbitrary numbering.

diffractometer with graphite-monochromated Mo- $K\alpha$ scanning radiation. The structure was solved by the direct method (SHELXS-86).

Crystallographic data (excluding structure factors) have been deposited with the *Cambridge Crystallographic Data Centre*. Copies of the data can be obtained, free of charge, on application to the CCDC, 12 Union Road, Cambridge CB2 1EZ UK (fax: +44(1223) 336033; e-mail: deposit@cdc.ac.uk).

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