Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Jia-Liang Zhong, Yang Lu* and Qi-Tai Zheng

Institute of Materia Medica, Chinese Academy of Medical Sciences, and Peking Union Medical College, 1 Xiannong Tan Street, Beijing 100050, People's Republic of China

Correspondence e-mail: luy@imm.ac.cn

Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(\text{C}-\text{C}) = 0.005 \text{ Å}$ R factor = 0.054 wR factor = 0.135 Data-to-parameter ratio = 8.8

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

5,7-Dihydroxy-8-methoxy-2-phenyl-4*H*-chromen-4-one monohydrate

In the crystal structure of the title compound, $C_{16}H_{12}O_5 \cdot H_2O_5$, the water O atom is involved in intermolecular hydrogen bonds which link the molecules into a three-dimensional network.

Received 3 January 2006 Accepted 17 January 2006

Comment

The title compound, (I), was extracted from *Scutellaria rehderiana Diels* with acetone (Su *et al.*, 2004; Li & Chen, 2005). Recently, the compound was successfully crystallized from ethyl acetate. The molecule is essentially planar, except for the methoxy methyl group. The dihedral angle between the two benzene rings is $1.5 (2)^{\circ}$ and between the C5–C10 and O1/C2/C3/C4/C9/C10 planes is only $0.2 (1)^{\circ}$. The torsion angle O1–C2–C1'–C6' is –179.8 (3)°.



Water atom OW acts as both a hydrogen-bond acceptor *via* HO4 from the hydroxy group O4 and as a donor *via* atom HWA to carbonyl atom O2 of the neighbouring molecule at $(-x - \frac{1}{2}, -y + 1, z - \frac{1}{2})$, as well as with the neighbouring OW atom at $(x + \frac{1}{2}, -y + \frac{1}{2}, -z)$.



Figure 1

The asymmetric unit of the title compound, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented by circles of arbitrary size.

© 2006 International Union of Crystallography All rights reserved

organic papers

The interaction of atoms O3 with O2 is intramolecular and completes a closed six-membered ring. Water molecules link to form chains which, in turn, link the chromenone molecules, forming a hydrogen-bonded three-dimensional network.

Experimental

The title compound was prepared according to the procedure for extracting *Scutellaria rehderiana Diels* (Su *et al.*, 2004; Li & Chen, 2005). At 283 K and under unventilated conditions, crystals appropriate for data collection were obtained by evaporation of an ethyl acetate solution over a period of one week.

Mo $K\alpha$ radiation Cell parameters from 1825 reflections $\theta = 3.5-27.3^{\circ}$ $\mu = 0.11 \text{ mm}^{-1}$ T = 298 (2) K Prism, yellow $0.40 \times 0.10 \times 0.10 \text{ mm}$

 $R_{\rm int} = 0.040$

 $\theta_{\rm max} = 27.3^{\circ}$

 $h = 0 \rightarrow 5$

 $k = -21 \rightarrow 21$

 $l = -23 \rightarrow 23$

Crystal data

$C_{16}H_{12}O_5 \cdot H_2O$
$M_r = 302.27$
Orthorhombic, $P2_12_12_1$
a = 4.7160 (9) Å
b = 16.551 (3) Å
c = 18.466 (4) Å
V = 1441.4 (5) Å ³
Z = 4
$D_x = 1.393 \text{ Mg m}^{-3}$

Data collection

MAC DIP 2030K diffractometer ω scans Absorption correction: none 4404 measured reflections 1825 independent reflections 1801 reflections with $I > 2\sigma(I)$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_0^2) + (0.0494P)^2$
$R[F^2 > 2\sigma(F^2)] = 0.054$	+ 0.6551P]
$wR(F^2) = 0.135$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.23	$(\Delta/\sigma)_{\rm max} = 0.001$
1825 reflections	$\Delta \rho_{\rm max} = 0.16 \ {\rm e} \ {\rm \AA}^{-3}$
208 parameters	$\Delta \rho_{\rm min} = -0.15 \text{ e} \text{ Å}^{-3}$
H-atom parameters constrained	Extinction correction: SHELXL97
-	Extinction coefficient: 0.059 (11)

Table 1

Selected geometric parameters (Å, °).

O2-C4	1.262 (4)	C9-C10	1.401 (4)
C2-C1′	1.466 (4)	C1′-C2′	1.394 (5)
C5-C6	1.378 (4)	C4′-C5′	1.383 (5)
C6-C7	1.398 (5)		
C2-O1-C9	119.6 (2)	C5-C6-C7	119.6 (3)
C8-O5-C11	113.4 (3)	C2' - C1' - C2	121.4 (3)
C3-C2-O1	121.7 (3)	C2'-C3'-C4'	120.1 (4)
C11-O5-C8-C9	83.0 (4)	O1-C2-C1'-C2'	-0.3(4)
C2-O1-C9-C8	-179.5 (3)	O1-C2-C1'-C6'	-179.8 (3)
C3 - C2 - C1' - C2'	178.5 (4)		

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
O3−H3A···O2	0.82	1.88	2.609 (3)	147
$O4-H4A\cdots OW$	0.82	2.02	2.766 (4)	151
$OW-HWA\cdots O2^{i}$	0.90(5)	1.93 (5)	2.828 (3)	177 (5)
$OW - HWB \cdots OW^{ii}$	0.90 (9)	2.03 (8)	2.903 (3)	161 (7)

Symmetry codes: (i) $-x - \frac{1}{2}, -y + 1, z - \frac{1}{2}$; (ii) $x + \frac{1}{2}, -y + \frac{1}{2}, -z$.



Figure 2

The molecular packing of the title compound. Dashed lines indicate the hydrogen-bonding interactions.

Water H atoms were initially located in a difference Fourier map and refined freely. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H distances in the range 0.92–0.98 Å and O-H distances of 0.82 Å and with $U_{iso}(H) = 1.2$ or 1.5 times (for methyl H) $U_{eq}(C)$ and $1.5U_{eq}(O)$. In the absence of significant anomalous dispersion effects, Friedel pairs were averaged.

Data collection: *DENZO* (Otwinowski & Minor, 1997); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

The authors acknowledge the financal support of the International Centre for Diffraction Data.

References

Johnson, C. K. (1976). ORTEPII. Report ORNL-5138. Oak Ridge National Laboratory, Tennessee, USA.

- Li, H.-B. & Chen, F. (2005). J. Chromatogr. A, 1074, 107-110.
- Otwinowski, Z. & Minor, W. (1997). Methods in Enzymology, Vol. 276, Macromolecular Crystallography, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.

Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.

Su, Y. L., Huang, Y. & Chen, Z. Y. (2004). China J. Chin. Mat. Med. 29, 863– 864.