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#### **Key indicators**

Single-crystal X-ray study T = 120 K Mean  $\sigma$ (C–C) = 0.006 Å R factor = 0.050 wR factor = 0.088 Data-to-parameter ratio = 8.5

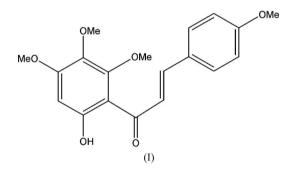
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the crystal structure of the title compound,  $C_{19}H_{20}O_6$ , the dihedral angle between the two aromatic planes is 31.8 (1)°. Intermolecular C-H···O hydrogen bonds serve to stabilize the structure.

2'-Hydroxy-4,4',5',6'-tetramethoxychalcone

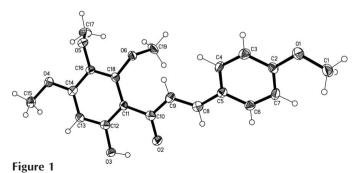
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### Comment

2'-Hydroxy-4,4',5',6'-tetramethoxychalcone (or odoratin) is a chalcone derivative that was isolated from the stem bark of *Eupatroium odoratum*, a plant used as a remedy for a variety of ailments, including malaria, fever and infections, and as a haemostatic agent (Shuib *et al.*, 1999). Chalcones represent an important group of natural compounds, showing a variety of biological activities including antifungal (Boeck *et al.*, 2005) and anticoagulant properties (Shuib *et al.*, 1999). The compounds are also used as depigmenting agents (Khatib *et al.*, 2005). Although known for long time (Bose *et al.*, 1973), the title compound, (I), has only now been characterized crystallographically.

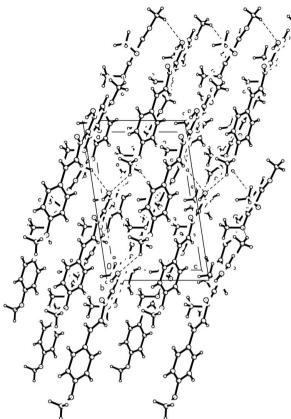


The molecular structure of (I) is closely related to that of the dimethoxychalcone [Cambridge Structural Database (*MOGUL*Version 1.7; Allen, 2002) refcode JEPJAP (Schmalle *et al.*, 1990)] with two methoxy groups at the 5'- and 6'-positions instead of H and OH. Most geometric parameters of the non-H atoms are not significantly different from those



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The molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level.





The crystal packing, viewed along [010], with the hydrogen-bonding pattern indicated by dashed lines.

of JEPJAP and need no further discussion. However, the dihedral angle of 31.8  $(1)^{\circ}$  between the two aromatic planes is considerably increased compared with  $13.1 (4)^{\circ}$  from the nearly planar JEPJAP molecule. With respect to the C8-C9-C10-O2 enone moiety, both aromatic rings are twisted out of this common plane by 14.7 (2) and 17.2 (3)°, respectively. The methoxy groups O1-C1 and O4-C15 lie in the planes of the attached aromatic rings, whereas both groups O5-C17 and O6-C19 are oriented almost perpendicular, with torsion angles C14-C16-O5-C17 of  $-78.3(5)^{\circ}$  and C11-C18-O6-C19 of 82.3 (4)°. As in JEPJAP, a strong intramolecular O3-H3···O2 hydrogen bond is observed, with  $H \cdots O = 1.64 \text{ Å}$  and  $O - H \cdots O = 144^{\circ}$ . The intramolecular C9-H9A...O6 bridge is less dominant, with H...O = 2.25 Å and C-H···O =  $111^{\circ}$ . Intermolecular C-H··O interactions are rather weak, with H...O distances greater than 2.45 Å. All these values are normalized for O-H =0.938 Å and C-H = 1.080 Å.

## **Experimental**

The leaves of Eupatorium odoratum Linn. were collected in Yaounde, Cameroon. Air-dried and powdered leaves (3 kg) were macerated in a mixture of CH<sub>2</sub>Cl<sub>2</sub>–MeOH (1:1) for 48 h. The extract was evaporated to yield the residue (200 g) which was then extracted with n-hexane. The extract (20 g) was subjected to column chromatography (CC) over a silica gel column using *n*-hexane with a gradient of acetone. The column eluates were monitored by thin-layer chro-

Crystal data

C19H20O6  $D_x = 1.423 \text{ Mg m}^{-3}$  $M_r = 344.35$ Mo  $K\alpha$  radiation Monoclinic, Cc Cell parameters from 1038 a = 13.092 (4) Å reflections b = 15.041 (5) Å  $\theta = 2.7 - 22.4^{\circ}$  $\mu = 0.11~\mathrm{mm}^{-1}$ c = 8.256 (3) Å  $\beta = 98.649 (7)^{\circ}$ T = 120 (2) K V = 1607.3 (9) Å<sup>3</sup> Prism, yellow  $0.48 \times 0.20 \times 0.18 \text{ mm}$ Z = 4

> 1954 independent reflections 1346 reflections with  $I > 2\sigma(I)$

 $R_{\text{int}} = 0.068$   $\theta_{\text{max}} = 28.1^{\circ}$   $h = -17 \rightarrow 17$   $k = -19 \rightarrow 19$  $l = -10 \rightarrow 10$ 

#### Data collection

Bruker SMART CCD area-detector
diffractometer
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Bruker, 2002)
$T_{\min} = 0.951, \ T_{\max} = 0.978$
7488 measured reflections

Refinement

Refinement on  $F^2$ H-atom parameters constrained $R[F^2 > 2\sigma(F^2)] = 0.050$  $w = 1/[\sigma^2(F_o^2) + (0.0277P)^2]$  $wR(F^2) = 0.088$ where  $P = (F_o^2 + 2F_c^2)/3$ S = 0.91 $(\Delta/\sigma)_{max} < 0.001$ 1954 reflections $\Delta\rho_{max} = 0.23$  e Å<sup>-3</sup>231 parameters $\Delta\rho_{min} = -0.29$  e Å<sup>-3</sup>

# Table 1

Selected geometric parameters (Å, °).

O1-C2	1.357 (4)	O5-C17	1.422 (4)
O1-C1	1.414 (5)	O6-C18	1.379 (4)
O2-C10	1.234 (4)	O6-C19	1.415 (4)
O3-C12	1.349 (4)	C5-C8	1.449 (5)
O4-C14	1.344 (5)	C8-C9	1.336 (5)
O4-C15	1.425 (5)	C9-C10	1.453 (5)
O5-C16	1.364 (4)	C10-C11	1.471 (5)
C2-O1-C1	116.9 (3)	C16-O5-C17	113.3 (3)
C14-O4-C15	116.8 (3)	C18-O6-C19	113.7 (3)

Table 2			
Hydrogen-bond	geometry	(Å,	°).

$D-\mathrm{H}\cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots \mathbf{A}$
O3−H3···O2	0.84	1.72	2.457 (4)	146
C9−H9A···O6	0.95	2.31	2.829 (4)	114
$C7-H7A\cdots O2^{i}$	0.95	2.57	3.329 (5)	137
$C1 - H1B \cdots O3^{ii}$	0.98	2.54	3.289 (5)	133
$C15-H15B\cdots O6^{iii}$	0.98	2.55	3.410 (5)	147
$C15-H15C\cdots O3^{iv}$	0.98	2.51	3.201 (5)	127

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

H atoms were refined at idealized positions (C–H = 0.95–0.98, O– H = 0.84 Å), riding on the parent C and O atoms with isotropic displacement parameters  $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$  or  $1.5 U_{\rm eq}({\rm methyl} {\rm C})$  and OH). All CH<sub>3</sub> and OH H atoms were allowed to rotate but not to tip. The title compound crystallizes in the non-centrosymmetric space group Cc; however, in the absence of significant anomalous scattering effects, the Flack (1983) parameter is essentially meaningless. Accordingly, Friedel pairs were merged.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2002); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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