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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.002 Å R factor = 0.043 wR factor = 0.118 Data-to-parameter ratio = 10.6

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. (*E*)-1-(2-Hydroxy-3,4-dimethoxyphenyl)-3-(4-hydroxyphenyl)prop-2-en-1-one

The title compound, $C_{17}H_{16}O_5$, crystallizes with two independent molecules in the asymmetric unit. These are oriented approximately orthogonal to one another. The keto group adopts an *s*-*cis* conformation; the six-membered ring formed by the O-H···O hydrogen bond and the adjacent benzene ring are approximately coplanar. In addition to an intramolecular O-H···O hydrogen bond, intermolecular O-H···O interactions link adjacent molecules into chains.

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Comment

The title compound (I), also referred to as heliannone A (Macias *et al.*, 1997), is one of the bioactive flavonoids from *Helianthus annuus* cultivars which possibly play a role in the allelopathic activity of sunflowers. It was synthesized by Koteswara Rao *et al.* (2001) and we report the crystal structure here.



The two independent molecules, A and B, in the asymmetric unit of (I) are oriented approximately orthogonal to one another at an angle of 85.44 (5)°. Bond lengths and angles in the central enone unit agree with those found in chalcone derivatives with different para substituents (Jevabharathi et al., 2002; Rabinovich & Shakked, 1974). The keto group adopts an s-cis conformation, with C7-C8-C9-O5 torsion angles of 5.8 (2)° for molecule A and -0.4 (2)° for molecule B. The keto system is not strictly planar, as can be seen from the torsion angles in Table 1. The least-squares mean planes of the hydroxyphenyl and dimethoxyphenyl rings make dihedral angles with the keto group of 13.93 (5) and 7.32 (5) $^{\circ}$, respectively, for molecule A, and 8.34 (5) and 8.42 (5)°, respectively, for molecule B. The dihedral angles between the rings themselves are 20.61 (5) for molecule A and 9.35 (5)° for molecule B. The enol rings formed by hydrogen bonding, comprising atoms O4A, O5A, C9A, C1', C2' and H30 in molecule A, and atoms O4B, O5B, C9B, C1", C2" and H30' in molecule B, are

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Figure 1

The structure of the asymmetric unit of (I), with 30% probability displacement ellipsoids.



Figure 2

The molecular packing of (I), viewed down the b axis. H atoms bonded to C atoms have been omitted for clarity. Dashed lines indicate hydrogen bonds.

slightly distorted. The fused benzene and enol rings are approximately coplanar, with dihedral angles between the least-squares mean planes of the benzene and enol rings of 4.44 (5)° for molecule A and 2.36 (5)° for molecule B.

The structure of (I) is stabilized by intra- and intermolecular hydrogen bonds. The intramolecular interaction involves the hydroxyl group and the adjacent ketone O atom, with O4A -H30···O5A 2.5534 (17) Å for molecule A and O4B- $0H30' \cdots O5B$ 2.5784 (16) Å for molecule B. This classic interaction is the likely cause of the lengthening of the C2'-O4A [1.340 (16) Å] and C2"-O4B bonds [1.3391 (16) Å] and shortening of the corresponding C9A = O5Athe [1.2538(17) Å] and C9B=O5B bonds [1.2406(17) Å] in comparison with normal C-O and C=O distances (Allen et al., 1987). In addition, atom O5A forms an intermolecular hydrogen bond with a hydroxyl group from a neighbouring molecule B $[O1B-H10' \cdots O5A^{ii} 2.8093 (16);$ symmetry code (ii): x - 1, y - 1, z - 1, to generate an infinite chain parallel to the b axis (Fig. 2). Further stabilization derives from hydrogen bonds between the H atom of the hydroxyl group O1A and the methoxy atoms O2B and O3B. These form additional links between the primary chains [O1A- $H10 \cdots O3B^{i} 3.0616 (17) \text{ and } O1A - H10 \cdots O2B 2.8855 (19) \text{ Å};$ symmetry code (i): -x + 2, -y + 2, -z + 2].

Experimental

The title compound was obtained from a solution of 2-hydroxy-3,4dimethoxyacetophenone (98 mg, 0.5 mmol) and p-hydroxybenzaldehyde (61 mg, 0.5 mmol) in EtOH (10 ml). This was added dropwise to 50% aqueous KOH (10 ml). The resulting mixture was stirred at room temperature for 72 h. The reaction mixture was poured into ice-cold water, its pH was adjusted to 3-4 with 10% aqueous HCl and it was extracted with Et_2O (4 \times 20 ml). The organic extract was washed with water and brine, dried over anhydrous MgSO₄ and evaporated under reduced pressure. The resulting residue was purified on a silica-gel column eluted with hexane-EtOAc (7:3) to give yellow crystals of the title chalcone, (I) (72 mg, 48%).

Crystal data

C ₁₇ H ₁₆ O ₅	Z = 4
$M_r = 300.30$	$D_x = 1.318 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 11.6831 (13) Å	Cell parameters from 8782
b = 11.8477 (13) Å	reflections
c = 12.9212 (14) Å	$\theta = 1.9-25.0^{\circ}$
$\alpha = 111.911 \ (2)^{\circ}$	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 111.296 \ (2)^{\circ}$	T = 293 (2) K
$\gamma = 92.437 \ (2)^{\circ}$	Prism, yellow
V = 1512.9 (3) Å ³	$0.25 \times 0.25 \times 0.13 \text{ mm}$

Data collection

Siemens SMART CCD area	4667
detector diffractometer	$R_{\rm int}$
ω scans	$\theta_{\rm max}$
Absorption correction: none	h =
14 357 measured reflections	<i>k</i> =
5298 independent reflections	l = -
Absorption correction: none 14 357 measured reflections 5298 independent reflections	h = k = l = l

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.043$ $wR(F^2) = 0.118$ S = 1.065298 reflections 502 parameters H atoms treated by a mixture of independent and constrained refinement

7 reflections with $I > 2\sigma(I)$ = 0.019 $= 25.0^{\circ}$ $-13 \rightarrow 13$

 $-14 \rightarrow 14$ $-15 \rightarrow 15$

 $w = 1/[\sigma^2(F_0^2) + (0.0699P)^2]$ + 0.2158P] where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} = 0.001$ $\Delta \rho_{\rm max} = 0.24 \text{ e} \text{ Å}^{-3}$ $\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$ Extinction correction: SHELXL97 (Sheldrick, 1997) Extinction coefficient: 0.025 (3)

Table 1Selected geometric parameters (Å, °).

O4B-C2"	1.3391 (16)	C8A-C9A	1.4650 (19)
O4A - C2'	1.3401 (16)	C1' - C9A	1.4644 (19)
O5B-C9B	1.2406 (17)	C8B-C7B	1.323 (2)
O5A-C9A	1.2538 (17)	C8B-C9B	1.4693 (18)
C1''-C9B	1.4711 (18)	C1B-C7B	1.4575 (19)
C8A-C7A	1.325 (2)	C1A-C7A	1.453 (2)
C6'' - C1'' - C9B	121.83 (12)	O5B-C9B-C8B	120.69 (12)
C2'' - C1'' - C9B	120.01 (11)	O5B-C9B-C1"	120.46 (11)
C7A-C8A-C9A	121.14 (14)	C8B-C9B-C1"	118.79 (12)
C6'-C1'-C9A	122.74 (12)	C2B-C1B-C7B	122.77 (12)
C2' - C1' - C9A	120.15 (12)	C6B - C1B - C7B	120.06 (13)
O5A-C9A-C1′	119.11 (12)	C8B-C7B-C1B	127.06 (14)
O5A-C9A-C8A	119.97 (12)	C2A - C1A - C7A	124.26 (13)
C1′-C9A-C8A	120.92 (12)	C6A - C1A - C7A	118.17 (14)
C7B-C8B-C9B	122.84 (14)	C8A-C7A-C1A	129.86 (15)
C2' $C1'$ $C0A$ $C8A$	174 55 (12)	$C^{\prime\prime\prime}$ $C^{\prime\prime\prime}$ $C^{\prime\prime}$ C^{0} R C^{0}	D 171 00 (12)
$C_2 = C_1 = C_9 A = C_8 A = -1/4.55$ (12)		$C_2 = C_1 = C_3 B = C_3 B$	B = -1/1.80(12)
C/A - C8A - C9A - O5A = 5.8 (2)		$C_{9B} = C_{8B} = C_{7B} = C_{1}$	D = 179.01(13)
C/A - C8A - C9A - C1' - 173.69 (15)		COB - CIB - C/B - CB	5B = 1/3.54(1/)
C/B = C8B = C9B = O3	BB = -0.4 (2)	C9A - C8A - C/A - C	1A = 1/8.6/(16)
C/B = C8B = C9B = C1	1/6.69 (13)	COA - CIA - C/A - C	8A - 1/0.5(2)

Table 2Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O4A−H30···O5A	0.92 (2)	1.70 (3)	2.5534 (17)	153 (2)
$O4B - H30' \cdots O5B$	0.94 (2)	1.74 (2)	2.5784 (16)	147 (2)
$O1A - H10 \cdots O3B^{i}$	0.83 (2)	2.27 (2)	3.0616 (17)	160 (2)
$O1A - H10 \cdots O2B^{i}$	0.83 (2)	2.30 (3)	2.8855 (19)	128 (2)
$O1B-H10' \cdots O5A^{ii}$	0.90 (2)	1.92 (2)	2.8093 (16)	169 (2)

Symmetry codes: (i) -x + 2, -y + 2, -z + 2; (ii) x - 1, y - 1, z - 1.

All H atoms, except those on the methyl groups of molecule *B*, were located in a difference Fourier map and only their positional parameters were refined. The remainder were included in calculated positions, with C-H distances constrained to 0.96 Å, and allowed to ride on their attached C atoms, with $U_{iso}(H) = 1.5U_{eq}(C)$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ZORTEP* (Zsolnai, 1997); software used to prepare material for publication: *enCIFer* (CCDC, 2003) and *PARST* (Nardelli, 1995).

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