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Key indicators

Single-crystal X-ray study T = 183 K Mean σ (C–C) = 0.002 Å Disorder in main residue R factor = 0.049 wR factor = 0.144 Data-to-parameter ratio = 16.2

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

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1-(2-Hydroxy-4,6-dimethoxy-3-methylphenyl)-3-(4-methoxyphenyl)-2-methyl-1,3-propanedione

The crystal structure of the title compound, $C_{20}H_{22}O_6$, an intermediate in the synthesis of the flavone 2-(4-methoxy-phenyl)-5,7-dimethoxy-3,8-dimethyl-4*H*-1-benzopyran-4-one, has been determined. The molecules are held together by van der Waals forces. There is an intramolecular hydrogen bond between the hydroxy group and the neighbouring keto O atom. The molecules in the crystal structure are disordered, with significantly different orientations of the methyl group in the disordered 4-methoxyphenyl substituent.

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Comment

In an investigation of the selective inhibition of certain rat heart enzymes caused by a series of lipophilic flavones, the inhibition effect of eight flavones, which are derivatives of either *O*-methylluteolin or *O*-methylapigenin, was studied (Gaillard *et al.*, 1996). The crystal structure of one of the *O*methylluteolin derivatives, 2-(3,4-dimethoxyphenyl)-5,7-dimethoxy-3,8-dimethyl-4*H*-1-benzopyran-4-one, has recently been determined by Stomberg *et al.* (2002). A second one of these flavones, the *O*-methylapigenin derivative 2-(4methoxyphenyl)-5,7-dimethoxy-3,8-dimethyl-4*H*-1-benzopyran-4-one, was prepared by acid-catalysed cyclization and dehydration of 1-(2-hydroxy-4,6-dimethoxy-3-methylphenyl)-3-(4-methoxyphenyl)-2-methyl-1,3-propanedione, (I). This paper describes the crystal structure of this synthetic intermediate, (I).



A perspective view of (I), together with the atomnumbering scheme, is shown in Fig. 1. The disorder in the methoxyphenyl group is shown in Fig. 2, and the packing of the molecules in Fig. 3. The molecules are held together by van der Waals forces. There is an intramolecular hydrogen bond between the hydroxy H atom H(O1) and the O atom O4 in the neighbouring carbonyl group. The benzylic C atom C7, the hydroxy O atom O1, and the methyl carbon atom C10 are twisted out of the C1/C2/C3/C4/C5/C6 ring plane by -0.094 (2), -0.014 (2) and 0.018 (3) Å, respectively. The methyl C atoms in the methoxy groups are situated close to the respective benzene ring planes [torsion angles: C5–C4–



Figure 1

The atom-numbering scheme for (I). Displacement ellipsoids are shown at the 50% probability level. The disorder of the methyl group C10 is shown. A dashed line indicates the intramolecular hydrogen bond.



Figure 2

The disorder of the 4-methoxyphenyl group. The minor part, with occupancy 0.194 (3), is shown with smaller radii of atoms and dashed bonds.



Figure 3 A packing diagram, in projection along the *c* axis.

O2-C11 -4.8 (2)°, C5-C6-O3-C12 1.6 (2)°, C16-C17-O6-C20 11.4 (4)° and C16A-C17A-O6A-C20A 172.6 (10)°].

Experimental

The synthesis of 1-(2-hydroxy-4,6-dimethoxy-3-methylphenyl)-3-(4-methoxyphenyl)-2-methyl-1,3-propanedione, (I), has been described by Gaillard *et al.* (1996), who also published ¹H-NMR data for the substance. Crystals suitable for X-ray analysis were obtained from benzene (m.p. 445–446 K).

Crystal data

$C_{20}H_{22}O_6$	$D_x = 1.332 \text{ Mg m}^{-3}$		
$M_r = 358.38$	Mo Ka radiation		
Monoclinic, I2/c	Cell parameters from 8192		
a = 12.2304 (2) Å	reflections		
b = 16.4299 (2) Å	$\theta = 1.7 - 30.6^{\circ}$		
c = 17.8325 (2) Å	$\mu = 0.10 \text{ mm}^{-1}$		
$\beta = 93.999 (1)^{\circ}$	T = 183 (2) K		
V = 3574.61 (8) Å ³	Needle, colourless		
<i>Z</i> = 8	$0.60 \times 0.12 \times 0.10 \text{ mm}$		
Data collection			
Siemens SMART CCD	5495 independent reflections		
diffractometer	3717 reflections with $I > 2\sigma(I)$		
ω scans	$R_{\rm int} = 0.041$		
Absorption correction: multi-scan	$\theta_{\rm max} = 30.6^{\circ}$		
(SADABS; Sheldrick, 2001)	$h = -17 \rightarrow 17$		
$T_{\min} = 0.943, T_{\max} = 0.990$	$k = -23 \rightarrow 23$		
24968 measured reflections	$l = -25 \rightarrow 25$		

Refinement	
Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0721P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.049$	+ 1.3054P]
$wR(F^2) = 0.144$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.99	$(\Delta/\sigma)_{\rm max} = 0.001$
5495 reflections	$\Delta \rho_{\rm max} = 0.24 \text{ e} \text{ Å}^{-3}$
340 parameters	$\Delta \rho_{\rm min} = -0.28 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	
-	

Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
O1−H1···O4	0.84	1.74	2.4899 (14)	148

Initial refinement of the structure converged with unacceptably high *R*-values (R = 0.095 for observed reflections and $wR^2 = 0.306$ for all reflections), and there were seven residual peaks in the electron density difference map larger than 0.42 e Å⁻³ (maximum 1.64 e Å⁻³). These peaks appeared close to the 4-methoxyphenyl group and indicated disorder. Six of the seven largest peaks formed a planar ring and they were taken to be the phenyl C atoms, while the largest residual peak could be attributed to the methoxy O atom. The methoxy C atom was also found (peak height 0.26 e Å⁻³). No other atom, except the benzylic C atom attached to the 4-methoxyphenyl group, could be resolved into two parts. Another kind of disorder was found in the methyl group C10. Refinement with geometrical similarity restraints on disordered atoms led to occupancies of 0.806:0.194(3).

H atoms were refined isotropically and were constrained to the ideal geometry, using an appropriate riding model. For the hydroxyl

group, the O–H distance (0.84 Å) and C–O–H angle (109.5°) were kept fixed, and the torsion angle was chosen to match the observed electron density. For methyl groups, the C–H distances (0.96 Å) and C–C–H angles (109.5°) were kept fixed, while the torsion angles were allowed to refine, with the starting position based on the threefold averaged circular Fourier synthesis. For methyl group C10, an idealized disordered methyl group was assumed, with two positions rotated from each other by 60 ° and with a half-occupancy for H atoms.

Data collection: *SMART* (Siemens, 1995); cell refinement: *SAINT* (Siemens, 1995); data reduction: *SAINT* and *SADABS* (Sheldrick, 2001); program(s) used to solve structure: *SHELXTL* (Bruker, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *DIAMOND* (Brandenburg, 2000); software used to prepare material for publication: *SHELXTL*.

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