organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 100 KMean $\sigma(\text{C}-\text{C}) = 0.003 \text{ Å}$ R factor = 0.048 wR factor = 0.125 Data-to-parameter ratio = 39.5

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

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1-(4-Bromophenyl)-3-(4-methoxyphenyl)prop-2-en-1-one

Each of the two unique molecules of the title compound, $C_{16}H_{13}BrO_2$, is non-planar, the dihedral angle between the two benzene rings being 55.04 (7)° in one molecule and 51.45 (6)° in the other. In the crystal structure, the molecules are stabilized by short $Br \cdots O$ interactions which link the molecules into chains along the *a* axis. These chains are stacked down the *b* axis.

Comment

The title compound, (I), was prepared as part of our ongoing studies (Patil *et al.*, 2006*a*,*b*) of non-linear optical (NLO) properties of chalcone derivatives (Fichou *et al.*, 1988; Kitaoka *et al.*, 1990; Uchida *et al.*, 1998; Goto *et al.*, 1991; Zhang *et al.*, 1990; Zhao *et al.*, 2000). Crystals of (I) do not exhibit second-order non-linear optical properties as they crystallize in a centrosymmetric space group.



The asymmetric unit of (I) contains two molecules, A and B (Fig. 1). The bond lengths and angles in both molecules are similar, showing normal values (Allen *et al.*, 1987), and are comparable to those in related structures (Patil *et al.*, 2006*a*,*b*). The dihedral angle between the two benzene rings is 55.04 (7)° in molecule A and 51.45 (6)° in molecule B. The least-squares plane through the enone unit (O1/C7–C9) makes dihedral angles of 24.03 (13) and 31.22 (13)° with the planes of the C1–C6 and C10–C15 benzene rings, respectively, in molecule A and 25.56 (9) and 25.90 (10)°, respectively, in molecule B. The methoxy group attached at C13 is almost coplanar with the C10–C15 benzene ring, with C16–O2–C13–C12 torsion angles of –170.33 (19) and 171.05 (19)°, respectively, for A and B.

In each independent molecule, an intramolecular C– H···O interaction (Table 1) generates an S(5) ring motif (Bernstein *et al.*, 1995). The short Br1A···O2 A^{i} [3.056 (2) Å] and Br1B···O2 B^{i} [3.022 (2) Å] [symmetry code: (i) 1 + x, y, z] contacts form chains of molecules along the *a* axis. The chains are stacked down the *b* axis (Fig. 2). Received 18 August 2006 Accepted 22 August 2006

Experimental

Crystal data C₁₆H₁₃BrO₂

4-Bromoacetophenone (0.01 mol) in ethanol (25 ml) was mixed with 4-methoxybenzaldehyde (0.01 mol) in ethanol (25 ml) and the mixture was treated with an aqueous solution of sodium hydroxide (5 ml, 20%). This mixture was stirred well and left to stand for 12 h. The resulting crude solid mass was collected by filtration and recrystallized from acetone.

Z = 8

 $D_x = 1.612 \text{ Mg m}^{-3}$

 $0.42\,\times\,0.28\,\times\,0.24$ mm

92812 measured reflections

13630 independent reflections

10641 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 3.14 \text{ mm}^{-1}$

Block, vellow

 $R_{\rm int} = 0.056$

 $\theta_{\rm max} = 37.5^\circ$

T = 100.0 (1) K



Figure 1

The asymmetric unit of (I), showing 50% probability displacement ellipsoids and the atomic numbering.



Figure 2

Part of the crystal packing of (I), viewed down the *b* axis. Dashed lines represent $Br \cdots O$ short contacts.

(UGC), Bangalore and New Delhi, for the award of a teacher fellowship under the Faculty Improvement Programme (FIP) of X Plan period.

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 $M_r = 317.17$ Monoclinic, $P2_1/c$ a = 15.7797 (3) Å b = 5.8463 (1) Å c = 32.4291 (6) Å $\beta = 119.088$ (1)° V = 2614.35 (9) Å³ Data collection

Bruker SMART APEX2 CCD areadetector diffractometer ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2005) $T_{\min} = 0.241, T_{\max} = 0.522$ (expected range = 0.217–0.471)

Refinement

$w = 1/[\sigma^2(F_0^2) + (0.0399P)^2]$
+ 4.617 <i>P</i>]
where $P = (F_0^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} = 0.001$
$\Delta \rho_{\rm max} = 1.13 \text{ e } \text{\AA}^{-3}$
$\Delta \rho_{\rm min} = -2.11 \text{ e } \text{\AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$\overline{D-\mathrm{H}\cdots A}$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C9A - H9A \cdots O1A$ $C9B - H9B \cdots O1B$	0.95	2.45	2.800 (3)	102
	0.95	2.48	2.808 (3)	100

H atoms were placed in calculated positions, with C–H distances is in the range 0.95–0.98 Å, and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms and $1.2U_{eq}$ for the remaining H atoms. The highest peak is located 0.65 Å from atom C9*B* and the deepest hole is 1.77 Å from Br1*A*.

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

The authors thank the Malaysian Government and Universiti Sains Malaysia for the Scientific Advancement Grant Allocation (SAGA) grant No. 304/PFIZIK/653003/ A118. VS is grateful to the University Grants Commission