

Hoong-Kun Fun,^{a*} Suchada
Chantrapromma,^{b,‡}
Shyamaprosad Goswami^c and
Subrata Jana^c^aX-ray Crystallography Unit, School of Physics,
Universiti Sains Malaysia, 11800 USM, Penang,
Malaysia, ^bDepartment of Chemistry, Faculty of
Science, Prince of Songkla University, Hat-Yai,
Songkhla 90112, Thailand, and ^cDepartment of
Chemistry, Bengal Engineering and Science
University, Shibpur, Howrah, India 711 103‡ Additional correspondence author, email:
suchada.c@psu.ac.th

Correspondence e-mail: hkfun@usm.my

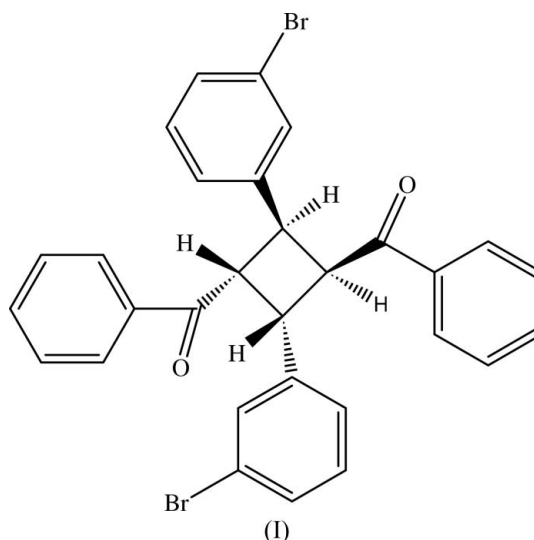
Key indicators

Single-crystal X-ray study
 $T = 100$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.026
 wR factor = 0.062
Data-to-parameter ratio = 20.8For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.[3-Benzoyl-2,4-bis(3-bromophenyl)cyclobutyl]-
(phenyl)methanoneMolecules of the title compound, $\text{C}_{30}\text{H}_{22}\text{Br}_2\text{O}_2$, lie across
crystallographic inversion centres. The dihedral angle between
the phenyl and bromobenzene rings is $23.88(8)^\circ$. Weak $\text{C}-\text{H}\cdots\text{O}$
interactions link the molecules into ribbons along the a
axis.

Received 10 January 2007

Accepted 25 January 2007

Comment

Cycloaddition reactions can be used to produce rings
containing different substituents as part of the total synthesis
of complex natural products (Nicolaou & Sorensen, 1996).
Cycloaddition reactions using microwave techniques are now
well established (Kappe, 2004).

The title compound, (I), formed by [2 + 2] cycloaddition using microwaves, lies across a crystallographic inversion centre; the asymmetric unit therefore contains one half-molecule (Fig. 1). The cyclobutane ring is planar. The dihedral angle between the phenyl and bromobenzene rings is $23.88(8)^\circ$, both rings being approximately perpendicular to the cyclobutane ring with dihedral angles of $89.18(11)^\circ$ and $78.26(10)^\circ$ for the phenyl and bromobenzene rings, respectively. The value of the dihedral angle [$30.44(10)^\circ$] between the $\text{C}7-\text{C}9/\text{O}1$ unit and phenyl ring shows that the carbonyl group is twisted out of the plane of the phenyl ring. The $\text{C}5-\text{C}6-\text{C}7-\text{O}1$ torsion angle is $-28.5(2)^\circ$. All bond lengths and angles in (I) show normal values (Allen *et al.*, 1987).

The crystal structure of (I) (Fig. 2) shows that the molecules are arranged in a zigzag manner along the b axis, with weak $\text{C}-\text{H}\cdots\text{O}$ interactions (Table 2) linking the molecules into ribbons along the a axis.

Experimental

Compound (I) was isolated from a reaction containing 3-(3-bromophenyl)-1-phenylpropenone (290 mg, 1 mmol) and 2-amino-6-methylpyridine (108 mg, 1 mmol) during optimization of the process to synthesize substituted naphthyridine-type heterocyclic compounds (Goswami, Jana, Dey & Adak, 2007; Goswami, Jana, Hazra & Adak, 2007) under microwave irradiation (450 W, 20 min) using a domestic microwave oven (BPL 800 G). One of the components, namely 3-(3-bromophenyl)-1-phenylpropenone, dimerized under these conditions to afford compound (I) as a minor product; (I) (yield 10 mg, 7%) was purified by preparative thin-layer chromatography using petroleum ether–ethyl acetate (9:1 v/v) as eluent. Single crystals were grown by slow evaporation of a CHCl₃/EtOH solution (1:1 v/v) (m.p. 337–340 K).

Crystal data

C₃₀H₂₂Br₂O₂ $Z = 2$
M_r = 574.28 $D_x = 1.615 \text{ Mg m}^{-3}$
 Monoclinic, *P*2₁/*c* Mo *K*α radiation
a = 5.7813 (1) Å $\mu = 3.46 \text{ mm}^{-1}$
b = 14.6794 (2) Å $T = 100.0 \text{ (1) K}$
c = 14.2828 (2) Å Needle, colourless
 $\beta = 102.988 \text{ (1) }^\circ$ $0.46 \times 0.14 \times 0.09 \text{ mm}$
V = 1181.11 (3) Å³

Data collection

Bruker SMART APEX2 CCD area-detector diffractometer 18746 measured reflections
 3429 independent reflections
 2869 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$
 $\theta_{\text{max}} = 30.0^\circ$
 Absorption correction: multi-scan (SADABS; Bruker, 2005)
 $T_{\text{min}} = 0.300, T_{\text{max}} = 0.737$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0261P)^2 + 0.7106P]$
 $R[F^2 > 2\sigma(F^2)] = 0.026$ where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.062$ $(\Delta/\sigma)_{\text{max}} = 0.003$
 $S = 1.04$ $\Delta\rho_{\text{max}} = 0.55 \text{ e } \text{Å}^{-3}$
 3429 reflections $\Delta\rho_{\text{min}} = -0.35 \text{ e } \text{Å}^{-3}$
 165 parameters
 H-atom parameters constrained

Table 1

Selected torsion angles (°).

C5–C6–C7–O1	–28.5 (2)	C8–C9–C10–C15	–30.8 (2)
C1–C6–C7–C8	–30.7 (2)		

Table 2

Hydrogen-bond geometry (Å, °).

D–H...A	D–H	H...A	D...A	D–H...A
C9–H9...O1 ⁱ	0.96	2.61	3.408 (2)	140

Symmetry code: (i) $-x + 3, -y + 1, -z + 1$.

All H atoms were positioned geometrically and refined using a riding model, with C–H = 0.93 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic, and C–H = 0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for CH H atoms.

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: SAINT (Bruker, 2005); program(s) used to solve

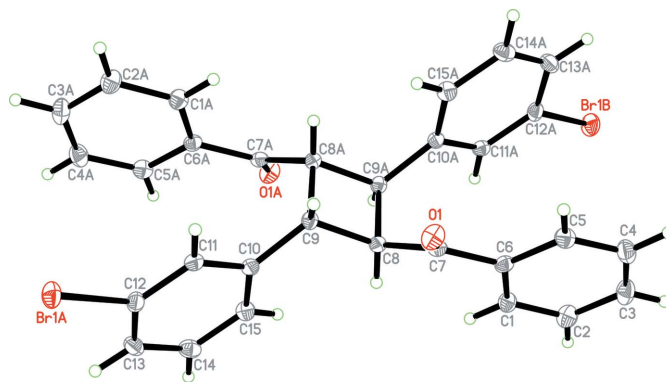


Figure 1

The molecular structure of (I), showing 50% probability displacement ellipsoids and the atomic numbering. [Symmetry code: (A) $2 - x, 1 - y, 1 - z$.]

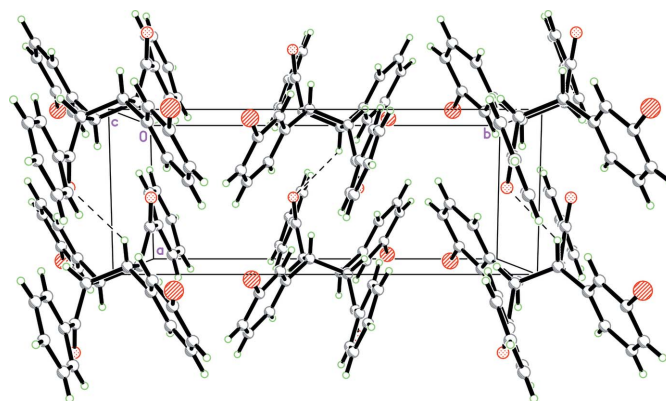


Figure 2

The crystal packing of (I), viewed down the *c* axis. C–H...O interactions are shown as dashed line.

structure: SHELXTL (Sheldrick, 1998); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2003).

SPG, SJ acknowledge the DST (SR/S1/OC-13/2005) and CSIR [01(1913)/04/EMR-II], Government of India, for financial support. SJ thanks the CSIR, Government of India, for a research fellowship. The authors also thank the Malaysian Government and Universiti Sains Malaysia for the Scientific Advancement Grant Allocation (SAGA) grant No. 304/PFIZIK/653003/A118.

References

Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
 Bruker (2005). APEX2 (Version 1.27), SAINT (Version V7.12a) and SADABS (Version 2004/1). Bruker AXS Inc., Madison, Wisconsin, USA.
 Goswami, S., Jana, S., Dey, A. & Adak, A. K. (2007). *Aust. J. Chem.* 60. In the press. [DOI: 10.1071/CH06388]
 Goswami, S., Jana, S., Hazra, A. & Adak, A. K. (2007). Unpublished results.
 Kappe, C. O. (2004). *Angew. Chem. Int. Ed.* 43, 6250–6284.
 Nicolaou, K. C. & Sorensen, E. J. (1996). *Classics in Total Synthesis*, pp. 265–283. Weinheim, New York, Cambridge, Tokyo: VCH.
 Sheldrick, G. M. (1998). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.
 Spek, A. L. (2003). *J. Appl. Cryst.* 36, 7–13.