## organic papers

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## Jeannie Bee-Jan Teh,<sup>a</sup> P. S. Patil,<sup>b</sup> Hoong-Kun Fun,<sup>a</sup>\* Ibrahim Abdul Razak<sup>a</sup> and S. M. Dharmaprakash<sup>b</sup>

<sup>a</sup>X-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and <sup>b</sup>Department of Studies in Physics, Mangalore University, Mangalagangotri, Mangalore 574 199, India

Correspondence e-mail: hkfun@usm.my

#### **Key indicators**

Single-crystal X-ray study T = 100 K Mean  $\sigma$ (C–C) = 0.002 Å R factor = 0.030 wR factor = 0.097 Data-to-parameter ratio = 44.2

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

# 3-(4-Bromophenyl)-1-(2,4-dichlorophenyl)prop-2-en-1-one

In the title compound,  $C_{15}H_9BrCl_2$ , the dihedral angle between the two benzene rings is  $42.39 (4)^{\circ}$ . The crystal structure is stabilized by short intermolecular Br...Br contacts.

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### Comment

Our continuing research on NLO materials (Patil et al., 2006a,b,c; Chantrapromma et al., 2006) has led us to synthesize the title compound, (I). We report here the crystal structure of (I) (Fig. 1). Crystals of (I) do not exhibit second-order nonlinear optical properties as this compound crystallizes in a centrosymmetric space group.



Bond lengths and angles in (I) show normal values (Allen et al., 1987) and are comparable to those in related structures (Teh et al., 2006a,b; Patil et al., 2006a,b,c). The least-squares plane through the enone unit (C7-C9/O1) makes dihedral angles of 45.42 (6) and 5.98  $(11)^{\circ}$  with the C1–C6 and C10– C15 benzene rings, respectively. The dihedral angle between the two benzene rings is  $42.39 (4)^{\circ}$ . The relatively short distance [3.5923 (2) Å] between the Br1 and Br1<sup>i</sup> [symmetry code: (i) 1 - x, -y, 1 - z atoms indicates the presence of intermolecular Br...Br interactions, which contribute to the stabilization of the molecular packing (Fig. 2).

## **Experimental**

2,4-Dichloroacetophenone (0.01 mol) in ethanol (30 ml) was mixed with 4-bromobenzaldehyde (0.01 mol) in ethanol (30 ml) and the mixture was treated with an aqueous solution of sodium hydroxide (5 ml, 20%). This mixture was stirred well and left for 24 h. The resulting crude solid mass was collected by filtration and recrystallized from acetone.

Crystal data	
C <sub>15</sub> H <sub>9</sub> BrCl <sub>2</sub> O	Z = 4
$M_r = 356.03$	$D_x = 1.781 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
a = 24.9039 (3) Å	$\mu = 3.49 \text{ mm}^{-1}$
b = 3.8474 (1)  Å	T = 100.0 (1) K
c = 13.9069 (2) Å	Plate, yellow
$\beta = 94.923 \ (1)^{\circ}$	$0.21 \times 0.21 \times 0.09 \text{ mm}$
V = 1327.58 (4) Å <sup>3</sup>	

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Data collection

Brucker SMART APEX2 CCD area-detector diffractometer  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Bruker, 2005)  $T_{min} = 0.552, T_{max} = 0.738$ 

## Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.030$   $wR(F^2) = 0.098$  S = 1.057604 reflections 172 parameters 42784 measured reflections 7604 independent reflections 5633 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.039$  $\theta_{\text{max}} = 38.8^{\circ}$ 

H-atom parameters constrained  $w = 1/[\sigma^2(F_o^2) + (0.0553P)^2]$ where  $P = (F_o^2 + 2F_c^2)/3$   $(\Delta/\sigma)_{max} = 0.001$   $\Delta\rho_{max} = 0.68 \text{ e } \text{\AA}^{-3}$  $\Delta\rho_{min} = -0.57 \text{ e } \text{\AA}^{-3}$ 

H atoms were placed in calculated positions and refined as riding, with C-H = 0.93 Å and  $U_{iso}(H) = 1.2U_{eq}(C)$ .

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).

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#### Figure 1

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



#### Figure 2

The crystal packing of (I), viewed down the b axis. Dashed lines indicate Br $\cdots$ Br interactions.

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