## organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 296 KMean  $\sigma$ (C–C) = 0.007 Å R factor = 0.035 wR factor = 0.079 Data-to-parameter ratio = 14.6

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

## 3-(4-Bromoanilino)isobenzofuran-1(3H)-one

Crystals of the title compound,  $C_{14}H_{10}BrNO_2$ , contain N— H···O hydrogen bond interactions that generate C(6) chains; these chains are linked by C—H···O hydrogen bonds, generating an  $R_4^3(21)$  ring motif. Received 18 July 2006 Accepted 4 September 20063-Substituted phthalides, Part XIV

#### Comment

We report here the structure of 3-(4-bromoanilino)isobenzofuran-1(3*H*)-one, (I) (Fig. 1 and Table 1), and we briefly compare this with the structure of the chloro and fluoro analogues, which have been previously reported by us (Büyükgüngör & Odabaşoğlu, 2006; Odabaşoğlu & Büyükgüngör, 2006*a*). The dihedral angle between the *p*-bromophenyl ring and the mean plane of the phthalide group is  $62.2 (2)^{\circ}$ ; for comparison, this angles is  $75.58 (15)^{\circ}$  in 3-(4chloroanilino)phthalide and  $74.10 (9)^{\circ}$  in 3-(4-fluoroanilino)phthalide.



The phthalide group (C1–C8/O2) is essentially planar, the largest deviation from the mean plane being 0.030 (4) Å for atom C7. In (I), as in the chloro analogue, the crystal packing is stabilized by N–H···O intermolecular hydrogen bonds (Fig. 2 and Table 2), which generate a C(6) chain. The C(6) chains are linked by C–H···O interactions, generating an  $R_4^3(21)$  ring motif (Etter, 1990). The chloro and fluoro analogues have  $\pi$ - $\pi$  stacking interactions but (I) does not.



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# The molecular structure of (I), showing the atomic numbering scheme and displacement ellipsoids drawn at the 30% probability level.

#### **Experimental**

The title compound was prepared as described by Odabaşoğlu & Büyükgüngör (2006*b*), using phthalaldehydic acid and 4-bromoaniline as starting materials (yield 90%; m.p. 452–453 K). Crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of a DMF solution at room temperature.

Z = 4

T = 296 K

 $R_{\rm int} = 0.081$ 

 $\theta_{\rm max} = 26.0^{\circ}$ 

Plate, colourless

 $0.77 \times 0.31 \times 0.07 \text{ mm}$ 

11941 measured reflections

2452 independent reflections

1446 reflections with  $I > 2\sigma(I)$ 

 $D_x = 1.602 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation  $\mu = 3.25 \text{ mm}^{-1}$ 

#### Crystal data

$C_{14}H_{10}BrNO_2$
$M_r = 304.14$
Orthorhombic, Pna21
a = 8.1158 (8) Å
b = 27.3737 (18) Å
c = 5.6775 (4)  Å
$V = 1261.31 (17) \text{ Å}^3$

#### Data collection

Stoe IPDS-2 diffractometer  $\omega$  scans Absorption correction: integration (X-RED; Stoe & Cie, 2002)  $T_{\min} = 0.298, T_{\max} = 0.810$ 

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0347P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.035$	where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.079$	$(\Delta/\sigma)_{\rm max} = 0.002$
S = 0.92	$\Delta \rho_{\rm max} = 0.21 \text{ e } \text{\AA}^{-3}$
2452 reflections	$\Delta \rho_{\rm min} = -0.29 \text{ e } \text{\AA}^{-3}$
168 parameters	Absolute structure: Flack (1983),
H atoms treated by a mixture of	1089 Friedel pairs
independent and constrained	Flack parameter: 0.004 (15)
refinement	

#### Table 1

Selected geometric parameters (Å, °).

C1-O1	1.193 (6)	C2-C7	1.360 (6)
C1-O2	1.354 (4)	C9-N1	1.397 (5)
O1-C1-O2	121.5 (4)	N1-C8-O2	112.2 (3)
O1-C1-C2	131.1 (4)		

#### Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$N1 - H1 \cdots O1^{i}$	0.78 (4)	2.35 (4)	3.083 (5)	156 (4)
C5−H5···O1 <sup>ii</sup>	0.93	2.42	3.254 (7)	149

Symmetry codes: (i) x, y, z + 1; (ii)  $x - \frac{1}{2}, -y + \frac{1}{2}, z + 1$ .



Figure 2

A partial packing diagram for (I), showing the N-H·O and C-H···O interactions (dashed lines). H atoms not involved in hydrogen bonds have been omitted for clarity. [Symmetry codes: (i) x, y, 1 + z; (ii) x, y, z - 1; (iii)  $\frac{1}{2} + x, \frac{1}{2} - y, z - 1$ ].

All H atoms attached to C atoms were treated as riding on their parent atoms, with C-H = 0.93 Å for aromatic and 0.98 Å for methine H atoms, and with  $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C})$ . The H atom of the amino group was located in a Fourier difference map and freely refined.

Data collection: X-AREA (Stoe & Cie, 2002); cell refinement: X-AREA; data reduction: X-RED32 (Stoe & Cie, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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