

Mustafa Odabaşoğlu^a and Orhan
Büyükgüngör^{b*}^aDepartment of Chemistry, Faculty of Arts & Science, Ondokuz Mayıs University, TR-55139 Kurupelit Samsun, Turkey, and ^bDepartment of Physics, Faculty of Arts & Science, Ondokuz Mayıs University, TR-55139 Kurupelit Samsun, Turkey

Correspondence e-mail: muodabas@omu.edu.tr

Key indicators

Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(C-C) = 0.003$ Å
 R factor = 0.035
 wR factor = 0.084
Data-to-parameter ratio = 13.4For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.3-Anilinoisobenzofuran-1(3*H*)-oneCrystals of the title compound, $C_{14}H_{11}NO_2$, are stabilized by an intermolecular $N-H \cdots O$ hydrogen bond and a weak $\pi-\pi$ interaction. $N-H \cdots O$ hydrogen-bond interactions generate $C(6)$ chains. The phthalide section of the molecule is planar and the dihedral angle between the phthalide group and the benzene ring is $78.43(5)^\circ$.

Comment

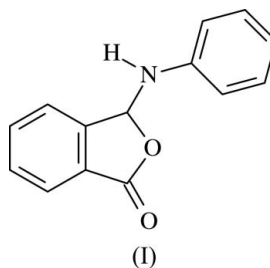
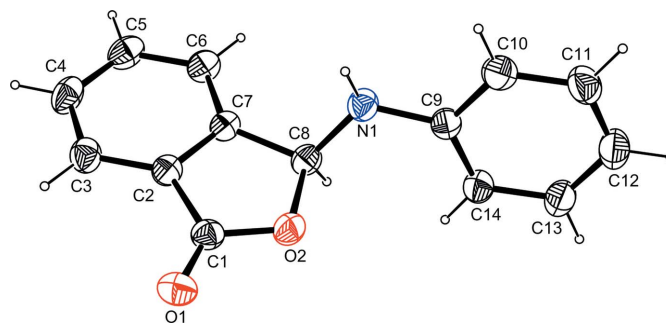
The present work is part of a structural study of compounds of 3-phenylsubstituted phthalides (Odabaşoğlu, & Büyükgüngör, 2006*a,b,c,d*; Büyükgüngör & Odabaşoğlu, 2006) and we report here the structure of 3-anilinoisobenzofuran-1(3*H*)-one, (I).The phthalide group (C1–C8/O2) is essentially planar, the largest deviation from the mean plane being $0.036(5)$ Å for atom C4 (Fig. 1). The dihedral angle between the mean planes of the phthalide group and the phenyl ring is $78.43(5)^\circ$.The crystal packing is stabilized by intermolecular $N-H \cdots O$ hydrogen bonds, which generate $C(6)$ chains (Etter, 1990) (Table 1). There are also weak $\pi-\pi$ interactions which occur between the C2–C7 six-membered ring and its symmetry-related counterpart at $(1-x, 1-y, 1-z)$, with a centroid-to-centroid distance of 3.646 Å and a plane-to-plane separation of 3.541 Å.

Figure 1

A view of (I), showing the atomic numbering scheme and displacement ellipsoids drawn at the 50% probability level. H atoms are drawn as spheres of arbitrary radii.

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Experimental

The title compound was prepared as described by Odabaşođlu & Büyükgüngör (2006a) using phthaldehydic acid and aniline as starting materials (yield 82%; m.p. 452–453 K). Crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of a dimethylformamide solution at room temperature.

Crystal data

$C_{14}H_{11}NO_2$	$Z = 4$
$M_r = 225.24$	$D_x = 1.374 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 12.5710 (14) \text{ \AA}$	$\mu = 0.09 \text{ mm}^{-1}$
$b = 7.0258 (6) \text{ \AA}$	$T = 296 \text{ K}$
$c = 15.7581 (15) \text{ \AA}$	Plate, light brown
$\beta = 128.525 (6)^\circ$	$0.38 \times 0.25 \times 0.07 \text{ mm}$
$V = 1088.8 (2) \text{ \AA}^3$	

Data collection

Stoe IPDS-2 diffractometer	7405 measured reflections
ω scans	2109 independent reflections
Absorption correction: integration (<i>X-RED32</i> ; Stoe & Cie, 2002)	1457 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.969$, $T_{\max} = 0.993$	$R_{\text{int}} = 0.038$
	$\theta_{\text{max}} = 26.0^\circ$

Refinement

Refinement on F^2	H atoms treated by a mixture of independent and constrained refinement
$R[F^2 > 2\sigma(F^2)] = 0.035$	$w = 1/[\sigma^2(F_o^2) + (0.043P)^2]$
$wR(F^2) = 0.084$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.01$	$(\Delta/\sigma)_{\text{max}} < 0.001$
2109 reflections	$\Delta\rho_{\text{max}} = 0.13 \text{ e \AA}^{-3}$
157 parameters	$\Delta\rho_{\text{min}} = -0.10 \text{ e \AA}^{-3}$

Table 1

 Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots O1^i$	0.880 (18)	2.169 (18)	3.0117 (17)	160.2 (14)

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

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

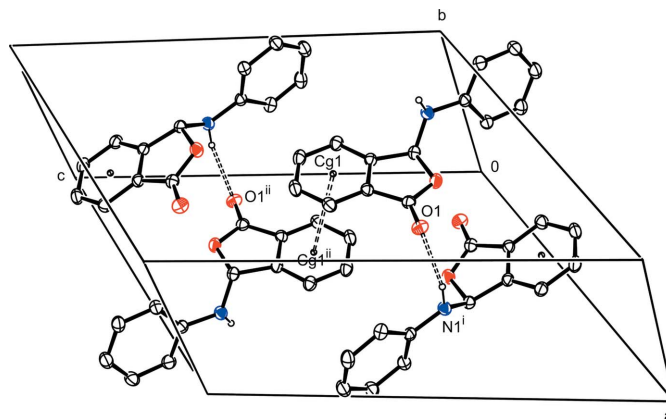


Figure 2

A packing diagram for (I), showing the $N-H\cdots O$ and $\pi-\pi$ interactions represented as dashed lines. H atoms not involved in hydrogen bonds have been omitted for clarity. Cg1 is the centroid of the phthalide benzene ring. [Symmetry codes: (i) $-x + 1, y - \frac{1}{2}, -z$; (ii) $1 - x, 1 - y, 1 - z$.]

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, 1997); 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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