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Key indicators

Single-crystal X-ray study T = 296 KMean $\sigma(C-C) = 0.002 \text{ Å}$ R factor = 0.042 wR factor = 0.115 Data-to-parameter ratio = 12.7

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

3-[(2-Hydroxy-5-nitrophenyl)amino]-2-benzofuran-1(3*H*)-one monohydrate

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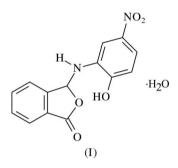
3-Substituted phthalides.

Part XI.

The crystal structure of the title compound, $C_{14}H_{10}N_2O_5 \cdot H_2O_5$, is stabilized by inversion-related $O-H \cdots O$ and $C-H \cdots O$ intermolecular hydrogen bonds and also by $\pi-\pi$ interactions. The dihedral angle between the phthalide group and the benzene ring is 51.45 (8)°.

Comment

The present work is part of a structural study of compounds of 3-substituted phthalides and we report here the structure of 3-(2-hydroxy-5-nitrophenyl)aminoisobenzofuran-1(3*H*)-one, (I) (Fig. 1 and Table 1).



The phthalide group (C1–C8/O2) is planar, the largest deviation from the mean plane being 0.037 (1) Å for atom O2. The dihedral angle between the mean planes of the phthalide group and the benzene ring is 51.45 (8)°; that between the nitro group and the benzene ring is 2.52 (16)°.

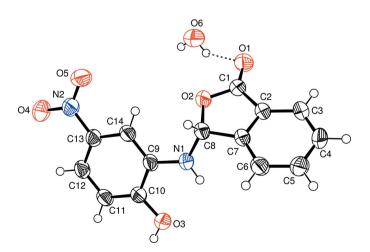


Figure 1

© 2006 International Union of Crystallography All rights reserved The asymmetric unit of (I), showing the atomic numbering scheme, with displacement ellipsoids drawn at the 50% probability level and a hydrogen bond shown as a dashed line.

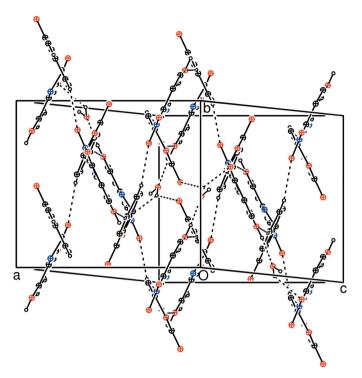


Figure 2

A packing diagram for (I), with hydrogen-bond interactions drawn as dashed lines. H atoms not involved in hydrogen bonding have been omitted.

The crystal packing is stabilized by inversion-related O- $H \cdots O$ and $C - H \cdots O$ intermolecular hydrogen bonds (Fig. 2 and Table 2) and $\pi - \pi$ interactions $[Cg1 \cdots Cg1^{vi} = 3.563 (1) \text{ Å};$ symmetry code: (vi) 2 - x, 1 - y, 1 - z; perpendicular distance = 3.400(12) Å; Cg1 is the centroid of the C9–C14 ring].

Experimental

The title compound was prepared as described by Odabaşoğlu & Büyükgüngör (2006), using phthalaldehydic acid and 2-hydroxy-4nitroaniline as starting materials (yield 83%, m.p. 526-527 K). Crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of an ethanol (95%) solution at room temperature.

Crystal data

$C_{14}H_{10}N_2O_5 \cdot H_2O$	Z = 4
$M_r = 304.26$	$D_x = 1.508 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
a = 10.8202 (7) Å	$\mu = 0.12 \text{ mm}^{-1}$
b = 9.8139 (4) Å	T = 296 K
c = 15.1475 (10) Å	Prism, light brown
$\beta = 123.588 \ (4)^{\circ}$	$0.45 \times 0.34 \times 0.14 \text{ mm}$
V = 1339.93 (14) Å ³	

18837 measured reflections

 $R_{\rm int} = 0.090$

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

2628 independent reflections

2077 reflections with $I > 2\sigma(I)$

Data collection

Stoe IPDS-II diffractometer ω scans Absorption correction: integration (X-RED32; Stoe & Cie, 2002) $T_{\rm min}=0.952,\ T_{\rm max}=0.984$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0646P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.042$	+ 0.1854P]
$wR(F^2) = 0.115$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} < 0.001$
2628 reflections	$\Delta \rho_{\rm max} = 0.24 \text{ e } \text{\AA}^{-3}$
207 parameters	$\Delta \rho_{\rm min} = -0.22 \text{ e } \text{\AA}^{-3}$
H atoms treated by a mixture of	
independent and constrained	
refinement	
Tabla 1	

Table 1

Selected geometric parameters (Å, °).

C1-O1 C1-O2 C2-C7	1.207 (2) 1.3502 (19) 1.377 (2)	C7-C8 C9-N1 C13-N2	1.501 (2) 1.3949 (19) 1.450 (2)
O1-C1-O2 O1-C1-C2	121.19 (14) 130.00 (15)	N1-C8-O2	112.19 (13)
C14-C13-N2-O4	179.99 (15)		

Table 2			
Hydrogen-bond	geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
O3−H3A…O6 ⁱ	0.82	1.84	2.643 (2)	166
$O6-H6A\cdots O1$	0.790 (19)	2.34 (4)	2.936 (3)	133 (4)
$O6-H6A\cdots O4^{ii}$	0.790 (19)	2.36 (4)	3.011 (2)	141 (5)
$O6-H6B\cdots O6^{iii}$	0.794 (19)	2.34 (4)	2.937 (4)	133 (5)
$C4-H4\cdots O3^{iv}$	0.93	2.60	3.500 (2)	164
$C5\!-\!H5\!\cdots\!O5^v$	0.93	2.54	3.367 (2)	148

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

The water H atoms were refined with distance restraints O-H =0.83 (2) Å and H6A···H6B = 1.20 (2) Å. All other H atoms were refined using the riding model approximation, with C-H = 0.93 Å for aromatic and 0.98 Å for methine, N-H = 0.86 Å, and O-H = 0.82 Å, with $U_{iso}(H) = 1.2U_{eq}(C,N)$ and $1.5U_{eq}(O)$.

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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