organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 150 KMean σ (C–C) = 0.004 Å R factor = 0.048 wR factor = 0.093 Data-to-parameter ratio = 8.9

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

2-Phenylpyrrolo[2,3-h]quinoline dihydrate

The crystal structure of the title compound, $C_{17}H_{12}N_2 \cdot 2H_2O$, comprises an essentially flat molecule, with the phenyl ring being inclined at 18.7 (1)° to the plane of the pyrrolo[2,3-h]quinoline moiety, packed with two water molecules per asymmetric unit, all in a tetragonal lattice. One water molecule is involved in hydrogen-bonding associations with both pyrrolo-NH and quinoline-N sites while, as part of the overall hydrogen-bonding network, both water molecules and their symmetry equivalents construct a 'Chinese lantern' arrangement.

Comment

The title compound, (I), was prepared from a two-stage procedure starting with the reaction of 8-hydrazinoquinoline dihydrochloride hydrate and acetophenone, followed by the cyclization of the intermediate product (E)-2-acetylbenzene-8-quinonylhydrozone. A structural example of this intermediate, as the thiophene derivative, has been previously published (Lynch & McClenaghan, 2001*a*). Furthermore, the structure of 2-(4-pyridyl)pyrrolo[3,2-*h*]quinoline, a product analogous to the title compound, has also been reported (Lynch & McClenaghan, 2001*b*). We are currently studying the structural aspects of derivatives of both 8-quinonyl-hydrozone and pyrrolo[3,2-*h*]quinoline before studying their potential as metal-chelating agents.



The molecule of (I) (Fig. 1) is essentially flat, with the phenyl ring being inclined at $18.7 (1)^{\circ}$ to the plane of the pyrrolo[2,3-*h*]quinoline moiety. It crystallizes with two water molecules per asymmetric unit. One O—H site from one water molecule resides in a hydrogen-bonded triangular arrangement with both pyrrolo-NH and quinoline-N sites. The other O—H site, in conjunction with the second water molecule and their symmetry-related analogues, then forms a hydrogen-bonded 'Chinese lantern' arrangement (Fig. 2).

Experimental

© 2002 International Union of Crystallography Printed in Great Britain – all rights reserved The title compound was obtained from Key Organics Ltd and crystals were grown from an ethanol solution.

Received 28 August 2002 Accepted 23 September 2002 Online 27 September 2002

Crystal data

 $C_{17}H_{12}N_2 \cdot 2H_2O$ $M_r = 280.32$ Tetragonal, $I\overline{4}$ a = 24.758 (4) Å c = 4.6952 (9) Å V = 2878.1 (8) Å³ Z = 8 $D_x = 1.294$ Mg m⁻³

Data collection

Bruker-Nonius KappaCCD area-
detector diffractometer
φ and ω scans
Absorption correction: multi-scan
(SORTAV; Blessing, 1995)
$T_{\min} = 0.983, T_{\max} = 0.991$
8208 measured reflections

Refinement

Refinement on F^2	H atoms treated by a mixture of
$R[F^2 > 2\sigma(F^2)] = 0.048$	independent and constrained
$wR(F^2) = 0.093$	refinement
S = 1.05	$w = 1/[\sigma^2(F_o^2) + (0.0339P)^2]$
1838 reflections	where $P = (F_o^2 + 2F_c^2)/3$
206 parameters	$(\Delta/\sigma)_{\rm max} < 0.001$
	$\Delta \rho_{-} = 0.16 \text{ e} \text{ Å}^{-3}$

Mo $K\alpha$ radiation

reflections

 $\theta = 2.9-27.5^{\circ}$ $\mu = 0.09 \text{ mm}^{-1}$

T = 150 (2) K

 $R_{\rm int} = 0.079$ $\theta_{\rm max} = 27.5^{\circ}$

 $h = -32 \rightarrow 32$ $k = -32 \rightarrow 32$ $l = -6 \rightarrow 5$

 $\Delta \rho_{\rm min} = -0.20 \text{ e } \text{\AA}^{-3}$

Block, colourless

 $0.20 \times 0.10 \times 0.10 \ \mathrm{mm}$

1838 independent reflections

1205 reflections with $I > 2\sigma(I)$

Cell parameters from 5959

Table 1

Hydrogen-bonding geometry (Å, °).

<i>D</i> -H··· <i>A</i>	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
N9 $-H1 \cdots O1W$	0.88 (3)	2.36 (3)	3.145 (3)	148 (3)
$O2W - H3W \cdot \cdot \cdot O2W^{i}$	0.83(2)	1.99 (2)	2.807 (3)	168 (3)
$O2W - H4W \cdot \cdot \cdot O1W^{ii}$	0.85(2)	1.90 (2)	2.744 (3)	175 (3)
$O1W - H1W \cdot \cdot \cdot N1$	0.87(2)	1.94 (2)	2.801 (3)	172 (3)
$O1W - H2W \cdots O2W$	0.83 (2)	2.02 (2)	2.840 (3)	170 (3)

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

All aromatic H atoms were included in the refinement, at calculated positions, as riding models, with C–H set to 0.95 Å. All water H atoms were initially located in a difference synthesis, but were then restrained to a distance of 0.83 Å, with riding displacement parameters. The remaining N-attached H atom was located in a difference synthesis; its positional and displacement parameters were refined. In the absence of significant anomalous scattering effects, the absolute structure can not be determined; Friedel pairs were merged.

Data collection: *DENZO* (Otwinowski & Minor, 1997) and *COLLECT* (Hooft, 1998); cell refinement: *DENZO* and *COLLECT*; data reduction: *DENZO* and *COLLECT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLUTON94* (Spek, 1994) and *PLATON97* (Spek, 1997); software used to prepare material for publication: *SHELXL97*.

The authors thank the EPSRC National Crystallography Service (Southampton).



Figure 1

The molecular configuration and atom-numbering scheme for the title compound, showing 50% probability ellipsoids.



Figure 2

Partial packing diagram, showing the hydrogen-bonded 'Chinese lantern' arrangement of the water molecules. Hydrogen-bonding associations are shown as dotted lines.

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