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Yu-Cheng Wang,* Xiang-Shan Wang, Zhao-Sen Zeng and Da-Qing Shi

Department of Chemistry, Xuzhou Normal University, Xuzhou 221116, People's Republic of China

Correspondence e-mail: xswang1974@yahoo.com

Key indicators

Single-crystal X-ray study T = 289 K Mean σ (C–C) = 0.004 Å R factor = 0.071 wR factor = 0.242 Data-to-parameter ratio = 12.2

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

organic papers

Diethyl 2,8-diamino-4,10-bis(3-nitrophenyl)naphtho[1,2-*b*;6,5-*b*']dipyran-3,9dicarboxylate dimethylformamide disolvate

The title compound, $C_{34}H_{28}N_4O_{10}\cdot 2C_3H_7NO$, was synthesized by the reaction of ethyl 2-cyano-3-(3-nitrophenyl)-1-acrylate and 1,5-naphthalenediol in the presence of KF–Al₂O₃ in refluxing ethanol. The structure is centrosymmetric and the fused pyran ring adopts a boat conformation. Received 26 May 2005 Accepted 31 May 2005 Online 10 June 2005

Comment

2-Aminochromenes are an important class of compounds found as the main components of many naturally occurring products employed as cosmetics and pigments and utilized as potential biodegradable agrochemicals (Morinaka & Takahashi, 1977; Witte *et al.*, 1986; Hafez *et al.*, 1987). The utility of fluoride salts as potential bases in a variety of synthetic reactions has been recognized in recent years. In particular, potassium fluoride coated with alumina (KF–alumina) has been a versatile base developed by Ando *et al.* (1982) for alkylation. Over the years, this reagent has found application in a large number of organic reactions (Clark, 1980). This background prompted us to synthesize compounds catalyzed by this solid-supported reagent. Here, we report the X-ray crystal structure analysis of the title compound, (I).



The structure of (I) is centrosymmetric. Each pyran ring is slightly distorted and adopts a boat conformation (Fig. 1). Atoms C3 and O1 deviate from the basal plane defined by atoms C1/C2/C4/C5 by 0.191 (2) Å and 0.084 (2) Å, respectively. A similar distortion was observed in methyl 2-amino-4- (4-methylphenyl)-5-oxo-5,6-dihydro-4H-pyrano[3,2-c]quino-line-3-carboxylate dimethylformamide solvate (Wang *et al.*, 2004) and 9-(2-hydroxy-4,4-dimethyl-6-oxocyclohex-1-enyl)-3,3,7-trimethyl-1,2,3,4-hexahydro-7H-xanthene (Li *et al.*, 2004). The basal plane of the pyran ring is nearly perpendicular to the C12–C17 benzene ring, forming a dihedral angle of 88.3 (2)°, and nearly parallel to the naphthalene ring system, forming a dihedral angle of 4.1 (2)°.

An intramolecular N1-H1A \cdots O3 hydrogen bond (Table 2) is formed between the amino and carbonyl groups (Fig. 2).

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Figure 1

The molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. Unlabelled atoms are related to labelled atoms by the symmetry operator (-x, 1 - y, 1 - z).

The amino group also forms an N1-H1B···O6 hydrogen bond with the solvent dimethylformamide solvent molecule.

Experimental

Compound (I) was prepared by the reaction of ethyl 2-cyano-3-(3nitrophenyl)-1-acrylate (0.98 g, 4 mmol) and 1,5-naphthalenediol (0.32 g, 2 mmol) in the presence of KF-Al₂O₃ (0.25 g) in EtOH (15 ml) at 363 K for 8 h (yield 84%, m.p. 559-561 K). Crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of a dimethylformamide solution. Elemental analysis, calculated: C 60.14, H 5.30, N 10.52%; found: C 60.25, H 5.45, N 10.43%. ¹H NMR (DMSO- d_6 , δ , p.p.m.): 1.09 (t, J = 7.2 Hz, 6H, 2CH₃), 2.74 (s, 6H, 2CH₃), 2.90 (*s*, 6H, 2CH₃), 4.01 (*q*, *J* = 7.2 Hz, 4H, 2CH₂), 5.29 (*s*, 2H, 2CH), 7.46-7.58 (m, 4H, ArH), 7.64-7.74 (m, 2H, ArH), 7.85 (s, 4H, 2NH2), 7.95 (s, 2H, 2CHO), 7.97-8.04 (m, 4H, ArH), 8.08-8.14 (m, 2H, ArH); IR (v, cm⁻¹): 3435, 3302 (NH₂), 3032 (Ar-H), 2953 (C-H), 1687 (C=O), 1609, 1525, 1495 (phenyl ring).

Crystal data

C ₃₄ H ₂₈ N ₄ O ₁₀ ·2C ₃ H ₇ NO	Z = 1
$M_r = 798.80$	$D_x = 1.377 \text{ Mg m}^{-3}$
Triclinic, $P\overline{1}$	Mo $K\alpha$ radiation
a = 8.355 (1) Å	Cell parameters from 26
b = 9.750 (1) Å	reflections
c = 12.202 (2) Å	$\theta = 3.3-29.6^{\circ}$
$\alpha = 99.292 \ (3)^{\circ}$	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 96.912 \ (3)^{\circ}$	T = 289 (2) K
$\gamma = 97.103 \ (3)^{\circ}$	Block, yellow
V = 963.4 (2) Å ³	$0.34 \times 0.24 \times 0.18 \text{ mm}$

Data collection

Siemens P4 diffractometer ω scans Absorption correction: none 5054 measured reflections 3340 independent reflections 2241 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.022$

 $\theta_{\rm max} = 25.0^\circ$ $h = -9 \rightarrow 9$ $k = -11 \rightarrow 11$ $l = -14 \rightarrow 8$ 3 standard reflections every 97 reflections intensity decay: 2.3%



Figure 2

A molecular packing diagram for (I). Dashed lines indicate hydrogen bonds. H atoms not involved in hydrogen bonding have been omitted.

Refinement

Refinement on F^2	H atoms treated by a mixture of
$R[F^2 > 2\sigma(F^2)] = 0.071$	independent and constrained
$vR(F^2) = 0.242$	refinement
S = 1.02	$w = 1/[\sigma^2(F_o^2) + (0.16P)^2]$
3340 reflections	where $P = (F_0^2 + 2F_c^2)/3$
273 parameters	$(\Delta/\sigma)_{\rm max} < 0.001$
	$\Delta \rho_{\rm max} = 0.42 \ {\rm e} \ {\rm \AA}^{-3}$
	$\Delta \rho_{\rm min} = -0.32 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

O1-C1	1.360 (3)	C2-C3	1.518 (4)
O1-C5	1.379 (3)	C3-C4	1.518 (3)
N1-C1	1.332 (4)	C4-C5	1.357 (4)
C1-C2	1.359 (4)	$C6-C6^{i}$	1.417 (5)
C1 - O1 - C5	119.3 (2)	$C_2 - C_3 - C_4$	110.4 (2)
C2-C1-O1	122.5 (2)	C5-C4-C3	120.7(2)
C1-C2-C3	121.7 (2)	C4-C5-O1	123.1 (2)
C5-O1-C1-C2	9.4 (4)	C2-C3-C4-C5	15.8 (3)
O1-C1-C2-C3	2.1 (4)	C3-C4-C5-O1	-6.3(4)
C1-C2-C3-C4	-13.9 (4)	C1-O1-C5-C4	-7.2 (4)

Symmetry code: (i) -x, -y + 1, -z + 1.

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

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1 - H1A \cdots O3$	0.86(1)	2.11 (3)	2.713 (3)	127 (3)
N1 - H1B \cdots O6	0.86(1)	2.02 (1)	2.866 (5)	170 (4)

Atoms H1A and H1B were refined isotropically, with the N-H bond lengths restrained to 0.86 Å. All other H atoms were positioned geometrically and refined as riding, with C-H distances in the range 0.93–0.98 Å and with $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: XSCANS (Siemens, 1994); cell refinement: XSCANS; data reduction: SHELXTL (Sheldrick, 1997); program(s) used to solve structure: SHELXTL; program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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