organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Ming Yu, Zhi Fan, Zuo-Liang Jing,* Xin Chen, Chun-Hua Diao and Qi-Liang Deng

College of Sciences, Tianjin University of Science and Technology, Tianjin 300222, People's Republic of China

Correspondence e-mail: jzl74@tust.edu.cn

Key indicators

Single-crystal X-ray study T = 294 K Mean σ (C–C) = 0.003 Å R factor = 0.040 wR factor = 0.118 Data-to-parameter ratio = 13.2

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

N-(2,4-Dinitrophenyl)-*N*'-(2-pyridylmethylene)hydrazine

The title compound [alternative name: pyridine-2-carbaldehyde 2,4-dinitrophenylhydrazone], $C_{12}H_9N_5O_4$, was prepared using pyridine-2-carbaldehyde and *N*-(2,4-dinitrophenyl)hydrazine. In the crystal structure, weak intermolecular $C-H\cdots O$ hydrogen-bonding interactions link adjacent molecules to form a three-dimensional supramolecular network.

Comment

In order to establish control over the preparation of crystalline solid materials so that their architecture and properties are predictable (Belloni *et al.*, 2005; Tynan *et al.*, 2005; Parashar *et al.*, 1988),the synthesis of new and designed crystal structures has become a major strand of modern chemistry. In the present study, we report the synthesis and molecular structure of the title compound, (I).



A view of the molecule of (I) is shown in Fig. 1. In the molecular structure, the C6–N2 and N2–N3 bond lengths of 1.274 (2) and 1.369 (2) Å, respectively, are consistent with those found in a related structure (Jing *et al.*, 2005). The dinitrophenylhydrazone moiety and the pyridine ring are both planar, with r.m.s. deviations of the fitted atoms of 0.039 (2) and 0.005 (2) Å, respectively. The dihedral angle between the dinitrophenylhydrazone and pyridine planes is 3.88 (8)°. A strong intramolecular N–H···O hydrogen bond (Table 1) stabilizes the molecular conformation. In the crystal packing, two weak C–H···O hydrogen-bond interactions are observed (Table 1), resulting in the formation of a three-dimensional supramolecular network.

Experimental

An anhydrous ethanol solution of pyridine-2-carbaldehyde (1.22 g, 10 mmol) was added to an anhydrous ethanol solution of (2,4-dinitrophenyl)hydrazine (2.03 g, 10 mmol), and the mixture was stirred at 350 K for 8 h under nitrogen. The red-brown precipitate was isolated, recrystallized from ethanol, and dried *in vacuo* to give the pure compound (I) in 81% yield. Bright-red single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution.

© 2005 International Union of Crystallography All rights reserved Received 23 August 2005 Accepted 15 September 2005 Online 21 September 2005

Crystal data

 $C_{12}H_9N_5O_4$ $M_r = 287.24$ Monoclinic, $P2_1/c$ a = 8.5690 (18) Å b = 6.3333 (13) Å c = 23.340 (5) Å $\beta = 91.532$ (4)° V = 1266.2 (5) Å³ Z = 4

Data collection

Bruker SMART CCD area-detector
diffractometer2564 independent reflections
1567 reflections with $I > 2\sigma(I)$ φ and ω scans $R_{int} = 0.037$ Absorption correction: multi-scan
(SADABS; Bruker, 1999) $\theta_{max} = 26.4^{\circ}$ $T_{min} = 0.929, T_{max} = 0.977$ $k = -7 \rightarrow 7$ 6807 measured reflections $l = -21 \rightarrow 29$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.040$ $wR(F^2) = 0.118$ S = 0.992564 reflections 194 parameters H atoms treated by a mixture of independent and constrained refinement

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - H \cdots A$
N3-H3A···O1	0.89(1)	1.98 (2)	2.624 (2)	128 (2)
$C4-H4\cdots O4^{i}$	0.93	2.55	3.368 (2)	147
$C1-H1\cdots O2^{ii}$	0.93	2.54	3.402 (3)	154
a			. 2 . 1	

 $D_x = 1.507 \text{ Mg m}^{-3}$

Cell parameters from 1875

 $0.34 \times 0.26 \times 0.20 \text{ mm}$

 $w = 1/[\sigma^2(F_0^2) + (0.0543P)^2$

where $P = (F_0^2 + 2F_c^2)/3$

+ 0.1982P]

 $\begin{array}{l} (\Delta/\sigma)_{\rm max} < 0.001 \\ \Delta\rho_{\rm max} = 0.14 \ {\rm e} \ {\rm \AA}^{-3} \\ \Delta\rho_{\rm min} = -0.19 \ {\rm e} \ {\rm \AA}^{-3} \end{array}$

Mo $K\alpha$ radiation

reflections

 $\theta = 2.9-25.9^{\circ}$ $\mu = 0.12 \text{ mm}^{-1}$

T = 294 (2) K

Block, red

Symmetry codes: (i) -x + 1, -y + 1, -z + 1; (ii) -x, $y + \frac{3}{2}$, $-z + \frac{1}{2}$.

H atoms bonded to C atoms were included in calculated positions (C-H = 0.93-0.96 Å) and refined using a riding-model approximation, with $U_{\rm iso}(H) = 1.2U_{\rm eq}(C)$ or $1.5U_{\rm eq}(\text{methyl C})$. The H atom attached to N3 was located in a difference Fourier map and refined freely.



A view of the title compound, shown with 30% probability displacement ellipsoids.

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

This work was supported by Science Fund of Tianjin University of Science and Technology (Grant No. 118181), which is gratefully acknowledged.

References

Belloni, M., Kariuki, B. M., Manickam, M., Wilkie, J. & Preece, J. A. (2005). Cryst. Growth, 5, 1443–1449.

- Bruker (1997). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruker (1999). SADABS, SMART and SAINT, Bruker AXS Inc., Madison, Wisconsin, USA.
- Jing, Z.-L., Yu, M., Chen, X., Diao, C.-H., Deng, Q.-L. & Fan, Z. (2005). Acta Cryst. E61, 0534–0535.
- Parashar, R. K., Sharma, R. C., Kumar, A. & Mohan, G. (1988). Inorg. Chim. Acta, 151, 201–208.
- Sheldrick, G. M. (1997). SHELXS97 and SHELXL97. University of Göttingen, Germany.
- Tynan, E., Jensen, P., Lees, A. C., Moubaraki, B., Murray, K. S. & Kruger, P. E. (2005). CrystEngComm, 7, 90–95.