## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=294 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.040$
$w R$ 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.

[^0]
## $N$-(2,4-Dinitrophenyl)- $N^{\prime}$-(2-pyridylmethylene)hydrazine

The title compound [alternative name: pyridine-2-carbaldehyde 2,4-dinitrophenylhydrazone], $\mathrm{C}_{12} \mathrm{H}_{9} \mathrm{~N}_{5} \mathrm{O}_{4}$, was prepared using pyridine-2-carbaldehyde and N -(2,4-dinitrophenyl)hydrazine. In the crystal structure, weak intermolecular $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ 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).

(I)

A view of the molecule of (I) is shown in Fig. 1. In the molecular structure, the $\mathrm{C} 6-\mathrm{N} 2$ and $\mathrm{N} 2-\mathrm{N} 3$ bond lengths of 1.274 (2) and 1.369 (2) $\AA$, 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) $\AA$, respectively. The dihedral angle between the dinitrophenylhydrazone and pyridine planes is $3.88(8)^{\circ}$. A strong intramolecular $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bond (Table 1) stabilizes the molecular conformation. In the crystal packing, two weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ 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 ( $2.03 \mathrm{~g}, 10 \mathrm{mmol}$ ), 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.

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## Crystal data

$$
\begin{aligned}
& \mathrm{C}_{12} \mathrm{H}_{9} \mathrm{~N}_{5} \mathrm{O}_{4} \\
& M_{r}=287.24 \\
& \text { Monoclinic, } P 2_{1} / c \\
& a=8.5690(18) \AA \\
& b=6.3333(13) \AA \\
& c=23.340(5) \AA \\
& \beta=91.532(4)^{\circ} \\
& V=1266.2(5) \AA^{3} \\
& Z=4
\end{aligned}
$$

## Data collection

Bruker SMART CCD area-detector diffractometer

## $\varphi$ and $\omega$ scans

Absorption correction: multi-scan (SADABS; Bruker, 1999) $T_{\text {min }}=0.929, T_{\text {max }}=0.977$
6807 measured reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.040$
$w R\left(F^{2}\right)=0.118$
$S=0.99$
2564 reflections
194 parameters
H atoms treated by a mixture of independent and constrained refinement

## $D_{x}=1.507 \mathrm{Mg} \mathrm{m}^{-3}$

Mo $K \alpha$ radiation
Cell parameters from 1875
reflections
$\theta=2.9-25.9^{\circ}$
$\mu=0.12 \mathrm{~mm}^{-1}$
$T=294$ (2) K
Block, red
$0.34 \times 0.26 \times 0.20 \mathrm{~mm}$

2564 independent reflections
1567 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.037$
$\theta_{\text {max }}=26.4^{\circ}$
$h=-8 \rightarrow 10$
$k=-7 \rightarrow 7$
$l=-21 \rightarrow 29$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0543 P)^{2}\right. \\
& \quad+0.1982 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.14 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }= \\
& \hline
\end{aligned}
$$

Table 1
Hydrogen-bond geometry ( $\AA{ }^{\circ}{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 3-\mathrm{H} 3 A \cdots \mathrm{O} 1$ | $0.89(1)$ | $1.98(2)$ | $2.624(2)$ | $128(2)$ |
| $\mathrm{C} 4-\mathrm{H} 4 \cdots \mathrm{O}^{\mathrm{i}}$ | 0.93 | 2.55 | $3.368(2)$ | 147 |
| $\mathrm{C} 1-\mathrm{H} 1 \cdots \mathrm{O}^{\text {ii }}$ | 0.93 | 2.54 | $3.402(3)$ | 154 |

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 ( $\mathrm{C}-\mathrm{H}=0.93-0.96 \AA$ ) and refined using a riding-model approximation, with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$ or $1.5 U_{\text {eq }}$ (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.

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