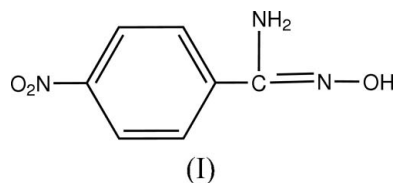
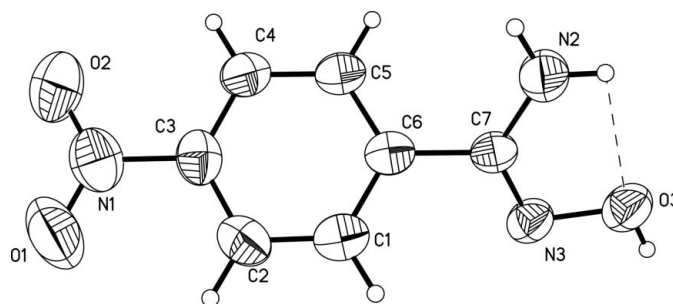


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wanghaibo@njut.edu.cn**Key indicators**Single-crystal X-ray study
 $T = 293$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.073
 wR factor = 0.200
Data-to-parameter ratio = 13.6For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.**4-Nitrobenzamidoxime**In the title compound, $\text{C}_7\text{H}_7\text{N}_3\text{O}_3$, which is a derivative of benzonitrile, the molecular conformation is stabilized by an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond.Received 22 December 2006
Accepted 22 December 2006**Comment**As part of our ongoing studies of benzonitrile derivatives (Wang *et al.*, 2007), we report here the crystal structure of the title compound, (I).The dihedral angle between the mean planes of the C1–C6 benzene ring and the C6/C7/N2/N3/O3 grouping is $41.0(2)^\circ$. In the related 2-methylbenzamidoxime the equivalent dihedral angle is $81.68(11)^\circ$ (Wang *et al.*, 2007).The molecular structure of (I) is shown in Fig. 1. Its conformation is stabilized by an intramolecular $\text{N2}-\text{H2A}\cdots\text{O3}$ hydrogen bond. The same H atom also participates in an intermolecular interaction (Table 1). Further intermolecular hydrogen bonds result in a three-dimensional network.**Experimental**

4-Nitrobenzonitrile (20 mmol) was dissolved in ethanol (8 ml), hydroxylamine hydrochloride (20 mmol) was dissolved in ethanol (6 ml) and potassium carbonate (10 mmol) was dissolved in water (10 ml); the three separate solutions were mixed. The resulting

**Figure 1**A view of the molecular structure of (I), showing displacement ellipsoids at the 50% probability level (arbitrary spheres for the H atoms). The dashed line indicates the intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond.

mixture was refluxed for 24 h. After cooling and filtering, crude compound (I) was obtained by crystallizing from a mixture of ethanol (6 ml) and water (2 ml). Crystals of (I) suitable for X-ray diffraction were obtained by slow evaporation of an ethanol solution.

Crystal data

$C_7H_7N_3O_3$ $Z = 8$
 $M_r = 181.16$ $D_x = 1.454 \text{ Mg m}^{-3}$
 Orthorhombic, *Pbca* Mo $K\alpha$ radiation
 $a = 11.320 (2) \text{ \AA}$ $\mu = 0.12 \text{ mm}^{-1}$
 $b = 7.7760 (16) \text{ \AA}$ $T = 293 (2) \text{ K}$
 $c = 18.802 (4) \text{ \AA}$ Needle, yellow
 $V = 1655.0 (6) \text{ \AA}^3$ $0.20 \times 0.10 \times 0.10 \text{ mm}$

Data collection

Enraf–Nonius CAD-4 1609 independent reflections
 diffractometer 671 reflections with $I > 2\sigma(I)$
 $\omega/2\theta$ scans $\theta_{\text{max}} = 26.0^\circ$
 Absorption correction: ψ scan 3 standard reflections
 (North *et al.*, 1968) every 200 reflections
 $T_{\text{min}} = 0.977$, $T_{\text{max}} = 0.988$ intensity decay: none
 1609 measured reflections

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0689P)^2]$
 $R[F^2 > 2\sigma(F^2)] = 0.073$ where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.200$ $(\Delta/\sigma)_{\text{max}} = 0.002$
 $S = 0.98$ $\Delta\rho_{\text{max}} = 0.21 \text{ e \AA}^{-3}$
 1609 reflections $\Delta\rho_{\text{min}} = -0.29 \text{ e \AA}^{-3}$
 118 parameters Extinction correction:
 H-atom parameters constrained *SHELXL97*
 Extinction coefficient: none

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N2-H2A\cdots O3$	0.86	2.23	2.544 (5)	102
$N2-H2A\cdots O2^i$	0.86	2.45	3.071 (5)	130
$N2-H2B\cdots O3^{ii}$	0.86	2.27	3.102 (6)	163
$O3-H3A\cdots N3^{iii}$	0.82	2.11	2.758 (5)	136

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

The H atoms were positioned geometrically ($C-H = 0.93 \text{ \AA}$, $O-H = 0.82 \text{ \AA}$, $N-H = 0.86 \text{ \AA}$) and refined as riding with $U_{\text{iso}}(H) = 1.2U_{\text{eq}}(C,N)$ or $1.5U_{\text{eq}}(O)$.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Siemens, 1996); software used to prepare material for publication: *SHELXL97*.

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