

Acetone 2-nitrophenylhydrazone

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Key indicators

Single-crystal X-ray study

$T = 120$ K

Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å

R factor = 0.054

wR factor = 0.153

Data-to-parameter ratio = 16.3

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

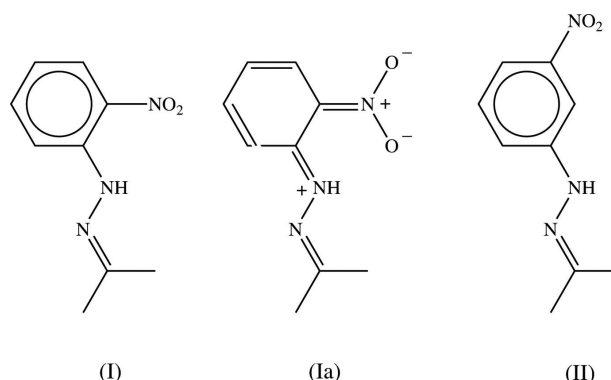
There are no direction-specific interactions between the almost-planar molecules of the title compound, $\text{C}_9\text{H}_{11}\text{N}_3\text{O}_2$.

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Comment

We report here the structure of acetone 2-nitrophenylhydrazone, (I) (Fig. 1), whose behaviour differs significantly from that of the isomeric compound acetone 3-nitrophenylhydrazone, (II) (Wardell *et al.*, 2006).



The non-H atoms in the molecule of (I) are virtually coplanar, as shown by the key torsion angles (Table 1). There is a short intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond (Table 2), which may assist in controlling the planar conformation. The bond distances (Table 1) show evidence for a significant contribution from the quinonoid form (Ia). In particular, the bonds $\text{C}3-\text{C}4$ and $\text{C}5-\text{C}6$ are shorter than the remaining bonds in the ring, while $\text{C}2-\text{N}21$ is very short for its type and the $\text{N}-\text{O}$ bonds are long (Allen *et al.*, 1987). In contrast, the bond distances in (II) show no unusual values (Wardell *et al.*, 2006).

Whereas the molecules of (II) are linked into complex sheets by a combination of $\text{N}-\text{H}\cdots\text{O}$, $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{N}$ hydrogen bonds, there are no direction-specific intermolecular interactions in the structure of (I). In particular, hydrogen bonds of all types and aromatic $\pi-\pi$ stacking interactions are absent.

Experimental

2-Nitrophenylhydrazine (3 mmol) was dissolved in acetone (30 ml) and the solution was heated under reflux for 1 h. The solution was then cooled and excess solvent was removed under reduced pressure. The resulting solid product, (I), was crystallized from ethanol (m.p. 339–341 K).

Crystal data

$C_9H_{11}N_3O_2$
 $M_r = 193.21$
 Monoclinic, $P2_1/c$
 $a = 3.8451$ (2) Å
 $b = 11.4926$ (8) Å
 $c = 21.3214$ (13) Å
 $\beta = 92.208$ (4)°
 $V = 941.50$ (10) Å³

$Z = 4$
 $D_x = 1.363$ Mg m⁻³
 Mo $K\alpha$ radiation
 $\mu = 0.10$ mm⁻¹
 $T = 120$ (2) K
 Needle, orange
 $0.40 \times 0.04 \times 0.04$ mm

Data collection

Bruker Nonius KappaCCD area-detector diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 2003)
 $T_{\min} = 0.971$, $T_{\max} = 0.996$

10829 measured reflections
 2104 independent reflections
 1738 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.046$
 $\theta_{\text{max}} = 27.8^\circ$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.054$
 $wR(F^2) = 0.153$
 $S = 1.06$
 2104 reflections
 129 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0854P)^2 + 0.2667P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.24$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.25$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

C1—C2	1.417 (2)	C1—N1	1.3637 (17)
C2—C3	1.4027 (19)	N1—N2	1.3799 (17)
C3—C4	1.370 (2)	N2—C7	1.2877 (18)
C4—C5	1.398 (2)	C2—N21	1.4387 (18)
C5—C6	1.3772 (19)	N21—O21	1.2513 (16)
C6—C1	1.4100 (19)	N21—O22	1.2267 (17)
C1—C2—N21—O21	−0.3 (2)	C1—N1—N2—C7	175.80 (12)
C1—C2—N21—O22	178.88 (13)	N1—N2—C7—C8	179.80 (12)
C2—C1—N1—N2	−175.52 (12)	N1—N2—C7—C9	0.2 (2)

Table 2

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1 \cdots O21	0.88	1.97	2.6010 (17)	128

All H atoms were located in a difference map and then treated as riding, with C—H distances of 0.95 or 0.98 Å and N—H distances of 0.88 Å, and with $U_{\text{iso}}(\text{H}) = kU_{\text{eq}}(\text{C,N})$ where $k = 1.5$ for methyl groups and 1.2 for all other H.

Data collection: COLLECT (Nonius, 1999); cell refinement: DENZO (Otwinowski & Minor, 1997) and COLLECT; data reduction: DENZO and COLLECT; program(s) used to solve structure: FLIPPER (Oszlányi & Sütő, 2004, 2005; Spek, 2003); program(s)

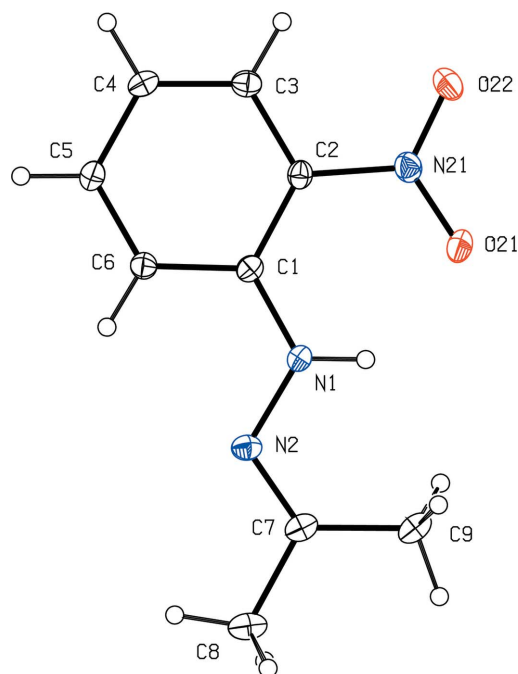


Figure 1

The molecular structure of compound (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

used to refine structure: OSCAIL (McArdle, 2003) and SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97 and PRPKAPPA (Ferguson, 1999).

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