

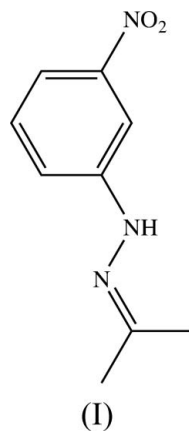
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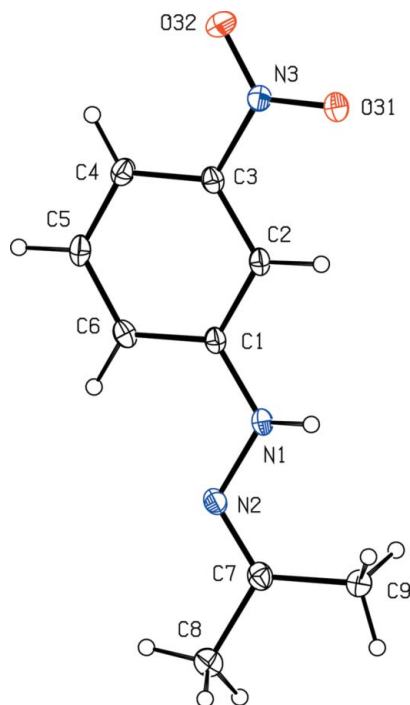
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## Key indicators

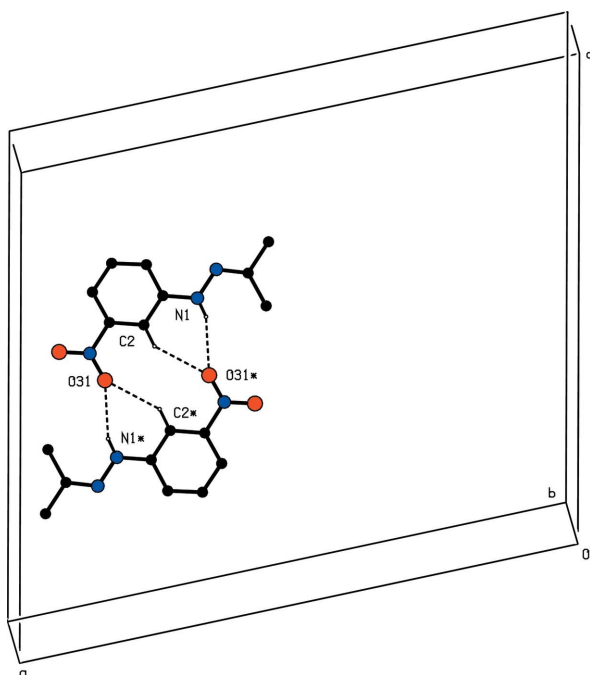
Single-crystal X-ray study  
 $T = 120\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$   
 $R$  factor = 0.058  
 $wR$  factor = 0.146  
Data-to-parameter ratio = 16.4For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.Acetone 3-nitrophenylhydrazone, redetermined at  
120 K: sheets built from N—H···O, C—H···O and  
C—H···N hydrogen bondsMolecules of the title compound,  $\text{C}_9\text{H}_{11}\text{N}_3\text{O}_2$ , are linked into  
sheets by a combination of N—H···O, C—H···O and C—  
H···N hydrogen bonds.Received 30 May 2006  
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## Comment

The structure of the title compound, (I) (Fig. 1), was deter-  
mined many years ago [Cambridge Structural Database,  
Version 7.27 (Allen 2002) refcode NPHYAC; Menczel, 1969],  
using diffraction data collected at ambient temperature; no H  
atom coordinates were reported, and the structure was refined  
only to  $R = 0.169$ . We have now redetermined this structure  
using diffraction data collected at 120 K, and we report here  
the details of the supramolecular aggregation.The molecules are linked into sheets by a combination of  
N—H···O, C—H···O and C—H···N hydrogen bonds  
(Table 1), and the sheet formation is readily analysed in terms  
of a hydrogen-bonded dimer as the basic building block.  
Atoms N1 and C2 in the molecule at  $(x, y, z)$  both act as  
hydrogen-bond donors to atom O31 in the molecule at  $(\frac{3}{2} - x,$   
 $\frac{1}{2} - y, 1 - z)$ , so forming a centrosymmetric dimer centred at  
 $(\frac{3}{4}, \frac{1}{4}, \frac{1}{2})$  containing three edge-fused rings, one of  $R_2^2(10)$  type  
(Bernstein *et al.*, 1995) flanked by two of  $R_2^2(6)$  type (Fig. 2).Atoms C5 in the molecules at  $(x, y, z)$  and  $(\frac{3}{2} - x, \frac{1}{2} - y,$   
 $1 - z)$ , which form the dimer centred at  $(\frac{3}{4}, \frac{1}{4}, \frac{1}{2})$ , act as  
hydrogen-bond donors to atoms N2 in the molecules at  $(\frac{3}{2} - x,$   
 $\frac{1}{2} + y, \frac{3}{2} - z)$  and  $(x, -y, -\frac{1}{2} + z)$ , which lie in the dimers centred  
at  $(\frac{3}{4}, \frac{3}{4}, 1)$  and  $(\frac{3}{4}, -\frac{1}{4}, 0)$ , respectively. Similarly, atoms N2 at  $(x,$   
 $y, z)$  and  $(\frac{3}{2} - x, \frac{1}{2} - y, 1 - z)$  accept hydrogen bonds from  
atoms C5 in the molecules at  $(\frac{3}{2} - x, -\frac{1}{2} + y, \frac{3}{2} - z)$  and  $(x, 1 - y,$   
 $-\frac{1}{2} + z)$ , which themselves form parts of the dimers centred at  
 $(\frac{3}{4}, -\frac{1}{4}, 1)$  and  $(\frac{3}{4}, \frac{3}{4}, 0)$ , respectively. Hence each dimer is linked  
by C—H···N hydrogen bonds to four other dimers and



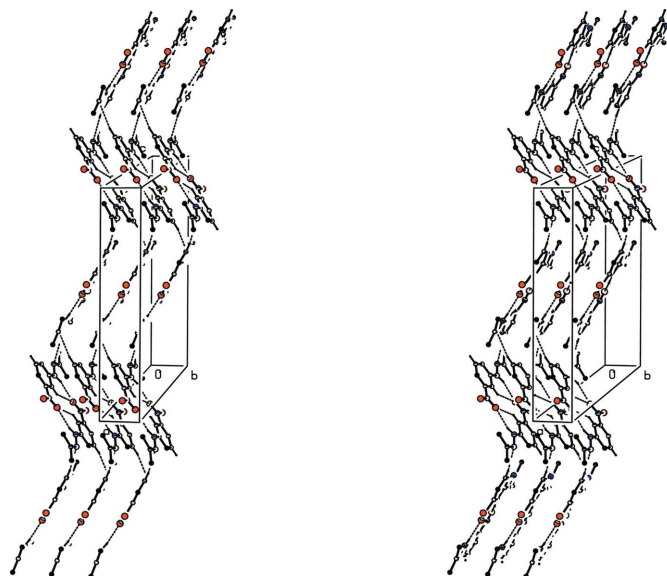
**Figure 1**  
A molecule of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.



**Figure 2**  
The molecular structure of (I), showing the formation of a centrosymmetric dimer containing three edge-fused hydrogen-bonded (dashed lines) rings. For the sake of clarity, H atoms not involved in the motifs shown have been omitted. The atoms marked with an asterisk (\*) are at the symmetry position ( $1-x, 1-y, 1-z$ ).

propagation of these interactions then generates a sheet parallel to (100) (Fig. 3).

This sheet is generated by centres of inversion with  $x = \frac{3}{4}$ , and it occupies the domain  $0.5 < x < 1.0$ ; a second sheet, related to the first by the *C*-centring operation, is generated by



**Figure 3**  
A stereoscopic view of part of the crystal structure of (I), showing the formation of a hydrogen-bonded (dashed lines) sheet parallel to (100). For the sake of clarity, H atoms not involved in the motifs shown have been omitted.

centres of inversion with  $x = \frac{1}{4}$ , and it occupies the domain  $0 < x < 0.5$ ; however, there are no direction-specific interactions between adjacent sheets.

## Experimental

3-Nitrohydrazine hydrochloride (3 mmol) was dissolved in acetone (30 ml) and the solution was then heated under reflux for 1 h. The solution was cooled and the excess solvent was removed under reduced pressure. The resulting solid product, (I), was crystallized from ethanol.

### Crystal data

$C_9H_{11}N_3O_2$   
 $M_r = 193.21$   
Monoclinic,  $C2/c$   
 $a = 22.7837$  (15) Å  
 $b = 3.8307$  (2) Å  
 $c = 21.7292$  (13) Å  
 $\beta = 100.129$  (3)°  
 $V = 1866.91$  (19) Å<sup>3</sup>

$Z = 8$   
 $D_x = 1.375$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
 $\mu = 0.10$  mm<sup>-1</sup>  
 $T = 120$  (2) K  
Lath, colourless  
 $0.20 \times 0.06 \times 0.02$  mm

### Data collection

Bruker-Nonius KappaCCD  
diffractometer  
 $\varphi$  and  $\omega$  scans  
Absorption correction: multi-scan  
(SADABS; Sheldrick, 2003)  
 $T_{\min} = 0.977$ ,  $T_{\max} = 0.998$

10187 measured reflections  
2117 independent reflections  
1392 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.079$   
 $\theta_{\max} = 27.5^\circ$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.058$   
 $wR(F^2) = 0.146$   
 $S = 1.03$   
2117 reflections  
129 parameters  
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0576P)^2 + 1.9052P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.28$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.30$  e Å<sup>-3</sup>

**Table 1**  
Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots O31^i$	0.84	2.35	3.144 (2)	158
$C2-H2\cdots O31^i$	0.95	2.49	3.305 (3)	143
$C5-H5\cdots N2^{ii}$	0.95	2.61	3.546 (3)	167

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

All H atoms were located in difference maps and then treated as riding atoms with distances  $C-H = 0.95 \text{ \AA}$  or  $0.98 \text{ \AA}$  and  $N-H = 0.84 \text{ \AA}$ , and with  $U_{iso}(H) = 1.2U_{eq}(C,N)$ .

Data collection: *COLLECT* (Hooft, 1999); cell refinement: *DENZO* (Otwinowski & Minor, 1997) and *COLLECT*; data reduction: *DENZO* and *COLLECT*; program(s) used to solve structure: *OSCAIL* (McArdle, 2003) and *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *OSCAIL* and *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PRPKAPPA* (Ferguson, 1999).

X-ray data were collected at the EPSRC X-ray Crystallographic Service, University of Southampton, England; the authors thank the staff of the Service for all their help and advice. JLW thanks CNPq and FAPERJ for financial support.

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