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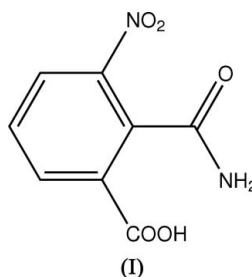
## Key indicators

Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
 $R$  factor = 0.057  
 $wR$  factor = 0.170  
Data-to-parameter ratio = 11.8For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

## 2-Carbamoyl-3-nitrobenzoic acid

In the title compound,  $\text{C}_8\text{H}_6\text{N}_2\text{O}_5$ , molecules are linked into  
ribbons along the  $b$  axis by  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds. The  
ribbons are interlinked into a three-dimensional framework.Received 15 May 2006  
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## Comment

We have previously reported the structure of 2-(2-nitrophenyl)-1*H*-benzimidazole (Li *et al.*, 2005). In our ongoing studies of benzimidazole derivatives, the title compound, (I), was obtained as an intermediate.

The bond lengths and angles in (I) show normal values (Allen *et al.*, 1987). The benzene ring, nitro and carboxyl groups are essentially coplanar, while atoms O3 and N2 lie above and below the molecular plane, with dihedral angles of 6.9 (2), 5.6 (1) and 84.5 (2)° between the C1–C6 ring plane and the mean planes through the C8/O4/O5, N1/O1/O2 and C7/N2/O3 groups, respectively. An intramolecular C4–H4A $\cdots$ O5 hydrogen bond forms a five-membered ring and contributes to the planarity of the molecule.

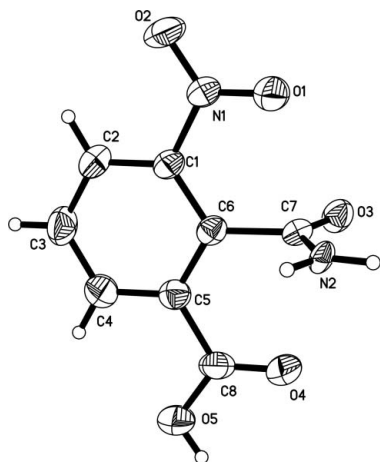
Molecules of (I) are linked into ribbons along the  $b$  axis (Fig. 2) via  $\text{N}-\text{H}\cdots\text{O}$  and  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds. These ribbons are further interlinked into a three-dimensional framework by  $\text{O5}-\text{H5A}\cdots\text{O4}$  and  $\text{C2}-\text{H2C}\cdots\text{O2}$  hydrogen bonds (Table 1).

## Experimental

Compound (I) was synthesized according to the method of Chapman & Stephen (1925). Single crystals were obtained by slow evaporation of an aqueous solution at room temperature over a period of 2 d.

## Crystal data

$\text{C}_8\text{H}_6\text{N}_2\text{O}_5$	$Z = 4$
$M_r = 210.15$	$D_x = 1.693$ Mg m $^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 9.6080$ (12) Å	$\mu = 0.14$ mm $^{-1}$
$b = 6.8998$ (9) Å	$T = 293$ (2) K
$c = 15.3291$ (14) Å	Plate, yellow
$\beta = 125.755$ (5)°	$0.25 \times 0.20 \times 0.06$ mm
$V = 824.68$ (17) Å $^3$	



**Figure 1**  
The structure of the compound (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.

*Data collection*

Siemens SMART 1000 CCD area-detector diffractometer	4196 measured reflections
$\omega$ scans	1606 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	1368 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.965, T_{\max} = 0.991$	$R_{\text{int}} = 0.019$
	$\theta_{\max} = 26.0^\circ$

*Refinement*

Refinement on $F^2$	$w = 1/[\sigma^2(F_o^2) + (0.0923P)^2 + 0.663P]$
$R[F^2 > 2\sigma(F^2)] = 0.057$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.170$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.07$	$\Delta\rho_{\max} = 0.51 \text{ e } \text{\AA}^{-3}$
1606 reflections	$\Delta\rho_{\min} = -0.62 \text{ e } \text{\AA}^{-3}$
136 parameters	
H-atom parameters constrained	

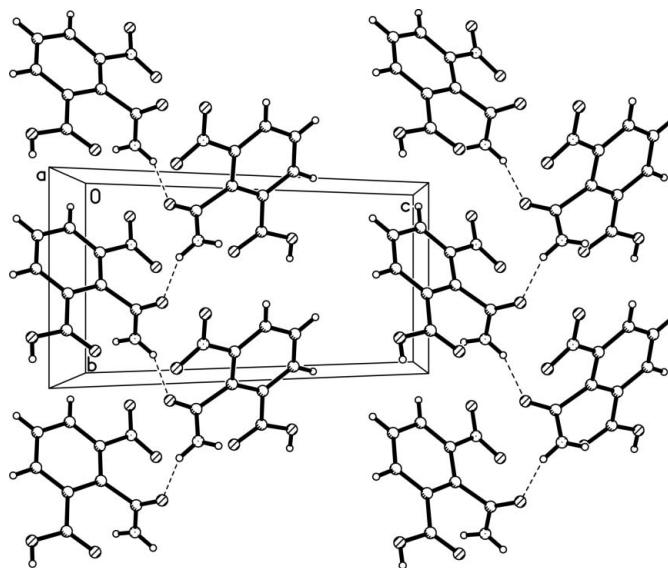
**Table 1**

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

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N2-H2A \cdots O3^i$	0.86	1.99	2.697 (3)	138
$O5-H5A \cdots O4^{ii}$	0.82	1.90	2.716 (3)	172
$C2-H2C \cdots O2^{iii}$	0.93	2.48	3.323 (4)	150
$C4-H4A \cdots O5$	0.93	2.41	2.733 (4)	100

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

All H atoms were located in difference Fourier maps and constrained to ride on their parent atoms with  $C-H = 0.93 \text{ \AA}$ ,  $O-H = 0.82 \text{ \AA}$  and  $N-H = 0.86 \text{ \AA}$ , and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$  and  $1.5U_{\text{eq}}(\text{O})$ .



**Figure 2**

The packing of (I), showing ribbons running along the  $b$  axis. Hydrogen bonds are indicated by dashed lines.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 1997); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2003).

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