



# Co-crystallization of *N'*-benzylidenepyridine-4-carbohydrazide and benzoic acid *via* autoxidation of benzaldehyde

Itumeleng B. Setshedi,<sup>a</sup> Andreas Lemmerer<sup>b</sup> and Mark G. Smith<sup>c\*</sup>

Received 8 May 2023

Accepted 28 June 2023

Edited by J. Ellena, Universidade de São Paulo, Brazil

This article is part of a collection of articles to commemorate the founding of the African Crystallographic Association and the 75th anniversary of the IUCr.

**Keywords:** crystal structure; autoxidation; benzoic acid; isoniazid.

**CCDC reference:** 2250754

**Supporting information:** this article has supporting information at journals.iucr.org/e

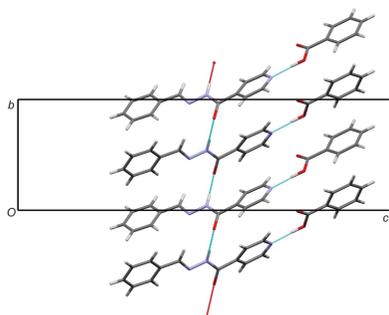
<sup>a</sup>University of South Africa, Department of Life Science, Unisa Science Campus, 28 Pioneer Avenue, Florida, Roodepoort, Gauteng, South Africa, <sup>b</sup>Molecular Sciences Institute, School of Chemistry, University of the Witwatersrand, Johannesburg, Gauteng, South Africa, and <sup>c</sup>University of South Africa, Chemistry Department, Unisa Science Campus, 28 Pioneer Avenue, Florida, Roodepoort, Gauteng, South Africa. \*Correspondence e-mail: smithm2@unisa.ac.za

The 1:1 co-crystal *N'*-[(2-methylphenyl)methylidene]pyridine-4-carbohydrazide–benzoic acid (1/1), C<sub>13</sub>H<sub>11</sub>N<sub>3</sub>O·C<sub>7</sub>H<sub>6</sub>O<sub>2</sub>, formed unexpectedly after autoxidation of benzaldehyde during the slow evaporation process of a solution of isoniazid in benzaldehyde. The original intent of the synthesis was to modify isoniazid with benzaldehyde and crystallize the product in order to improve efficacy against *Mycobacteria* species, but benzoic acid formed spontaneously and co-crystallized with the intended product, *N'*-benzylidenepyridine-4-carbohydrazide.

## 1. Chemical context

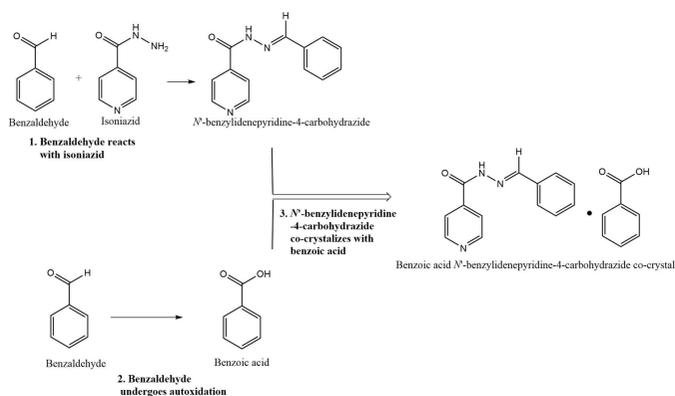
Mycobacterial infections are a historic tribulation to mankind, and are managed with an array of drugs ranging from natural to synthetic derivatives that possess antimicrobial properties. However, these strategies have failed over time due to the emergence of resistant *mycobacteria* (Cully, 2014). A number of constituents such as isoniazid (INH) have been modified to try and curb the scourge of tuberculosis (Cully, 2014). Some of the resulting modified INH derivatives have been shown to render the active pharmaceutical ingredients (API) a lot more active to the circulating resistant strains of TB (Hearn & Cynamon, 2004; Suarez *et al.*, 2009). It was for this reason that the covalent modification of API's was adopted to synthesize new analogues by modifying the NH<sub>2</sub> group of the hydrazide moiety of INH (Smith *et al.*, 2015), believed to assist in the evasion of the N-arylaminoacetyl transferases, an enzyme capable of reducing the efficacy of INH in particular, by acetylating the NH<sub>2</sub> position, thus ultimately preventing its reaction with nicotinamide adenine dinucleotide (NADH) (Vishweshwar *et al.*, 2006; Smith *et al.*, 2015).

Benzaldehyde is known to undergo autoxidation resulting in the formation of benzoic acid. The formation of benzoic acid occurs when benzaldehyde is exposed to air at room temperature (293 K) where the rate of the reaction is increased by the presence of a catalyst. However, this phenomenon can occur spontaneously without a catalyst over a prolonged period (Sankar *et al.*, 2014). The synthesis of this co-crystal was interesting as there were three separate processes that took place within the reaction mixture to create the final product. Firstly, benzaldehyde reacted with isoniazid to form *N'*-benzylidenepyridine-4-carbohydrazide. Secondly, excess benzaldehyde spontaneously autoxidized to form



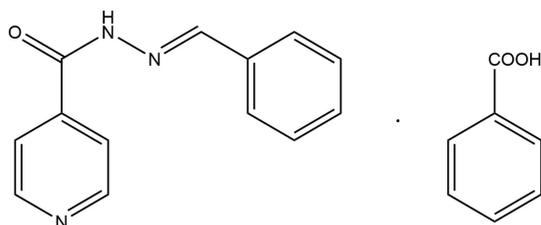
OPEN ACCESS

Published under a CC BY 4.0 licence



**Figure 1**  
Modification and autoxidative co-crystallization.

benzoic acid as described above (no benzoic acid was added to the reaction mixture). Lastly, the carbohydrazide moiety co-crystallized with the benzoic acid (as shown in Fig. 1) to form the product, *N'*-[(2-methylphenyl)methylidene]pyridine-4-carbohydrazide–benzoic acid (1/1).

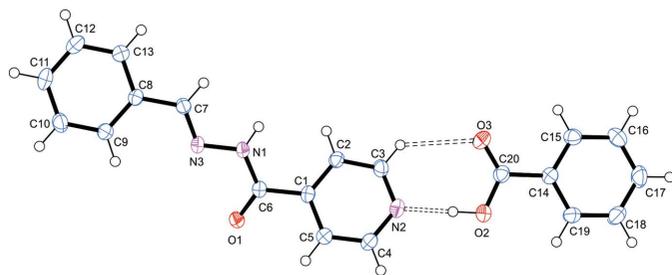


## 2. Structural commentary

The asymmetric unit contains one molecule of *N'*-benzylidenepyridine-4-carbohydrazide ( $C_{13}H_{11}N_3O_1 \cdot C_7H_6O_2$ ) and one molecule of benzoic acid (as shown in Fig. 2). This co-crystal crystallizes in the *Pbca* space group. The benzoic acid molecule lies in the plane of the pyridine ring of the benzylidene derivative. All bond lengths and angles are normal.

## 3. Supramolecular features

Each carbohydrazide moiety is hydrogen bonded by a strong  $O2-H2 \cdots N2$  hydrogen bond (Table 1) to a benzoic acid molecule to form a co-crystal. This interaction is supported by



**Figure 2**  
Asymmetric unit of *N'*-[(2-methylphenyl)methylidene]pyridine-4-carbohydrazide–benzoic acid (1/1).

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

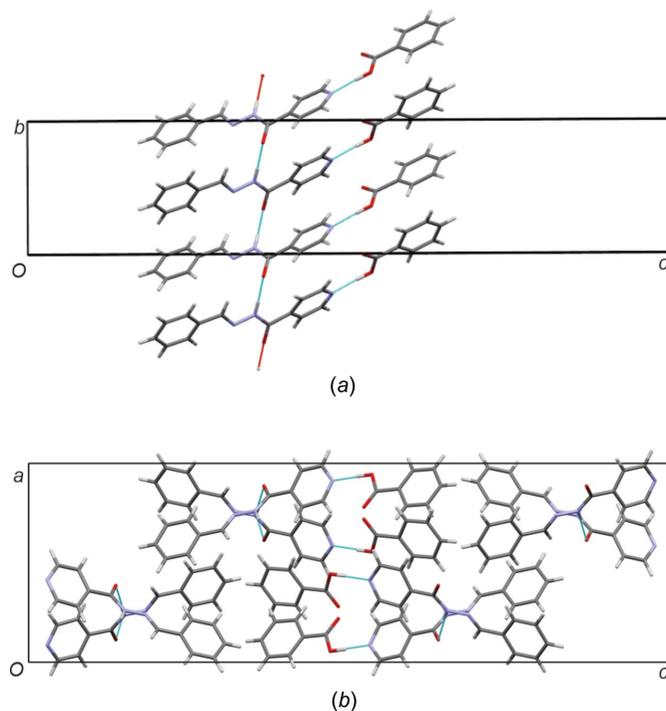
$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N1-H1 \cdots O1^i$	0.89 (2)	2.02 (2)	2.8981 (17)	171 (2)
$O2-H2 \cdots N2$	0.95 (2)	1.72 (3)	2.6693 (17)	176 (2)
$C3-H3 \cdots O3$	0.95	2.53	3.236 (2)	131

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

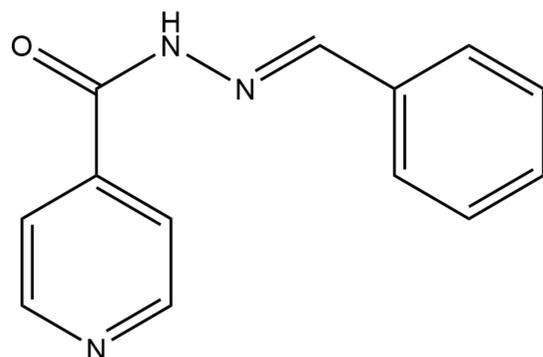
a weaker  $C-H \cdots O$  hydrogen bond that stabilizes the coplanar arrangement of the carboxylic acid moiety and the pyridine ring. The graph-set notation for this would be  $R_2^2(7)$  (Bernstein *et al.*, 1995), and is observed in other isoniazid co-crystals (Lemmerer *et al.*, 2010) (Fig. 2). This co-crystal is another example of the robust carboxylic acid  $\cdots$  pyridine heterosynthon (Shattock *et al.*, 2008; Aakerøy *et al.*, 2007). Each carbohydrazide moiety is also hydrogen bonded *via* its  $N1-H1$  donor to the carbonyl oxygen ( $O1$ ) acceptor of an adjacent carbohydrazide moiety. This results in a mono-periodic hydrogen-bonded chain along the *b*-axis direction, with graph-set notation  $C(4)$ . Overall, the combined carbohydrazide moiety with the benzoic acid forms a ribbon motif (as shown in Fig. 3*a*). Viewed along the *b*-axis, the ribbons forms a X-shaped motif seen in other carbohydrazide moieties (Hean *et al.* 2018) (Fig. 3*b*).

## 4. Database survey

ConQuest (Bruno *et al.*, 2002), Version 2022.1.0 of the CSD (Groom *et al.*, 2016) was used for the database survey, where



**Figure 3**  
(*a*) View of the strong hydrogen bonds that link the carbohydrazide molecules in a chain, and the benzoic acid molecules to the carbohydrazide forming a ribbon. (*b*) Packing diagram along the *b* axis.



**Figure 4**  
Benzylidene derivative of isoniazid.

only one similar structure was found. The survey consisted of structures consisting of isoniazid that had been modified with benzaldehyde and may have either a co-former or solvent molecule in the crystal structure. The structure of the anhydrous benzylidene derivative, *N'*-[(2-methylphenyl)methylidene]pyridine-4-carbohydrazide (as shown in Fig. 4), formed from the reaction of isoniazid and benzaldehyde, was reported by Wardell *et al.*, (2007) (YIQDEI). Several structures have been reported where substituted benzaldehyde reacted with isoniazid, for example, three polymorphs of the 4-methylbenzylidene derivative (WOGGOR, WOGGOR01 and WOGGUX) were reported by Purushothaman *et al.* (2019) and in 2016, Almeida and colleagues published the structure of a hydrate of the same 4-methylbenzylidene derivative (OLECOZ; Pereira Almeida *et al.*, 2016). However, there has not been any co-crystal of the benzylidene derivative (Fig. 4) reported in the literature to date.

## 5. Synthesis and crystallization

All reagents were commercially sourced and used without further purification. 1.00 g of isonicotinic acid hydrazide (isoniazid) (7.29 mmol) were dissolved in 15 ml of benzaldehyde in a 50 ml amber Schott bottle. The mixture was placed on a stirring heating block and heated to 333 K while stirring with a magnetic stirrer bar. Once the isoniazid had completely dissolved, the lid was tightly sealed. The solution was then allowed to react for 24 h. To maintain the temperature throughout the duration of the experiment, the amber Schott bottle was covered with an inverted round glass evaporation dish. After 24 h, the solution was allowed to cool to ambient temperature. The stirrer bar was retrieved and the sample was left to evaporate slowly for 6 weeks at ambient temperature without a lid. Over the 6 weeks, the temperature in the laboratory fluctuated between 298 and 300 K. Due to the fact that benzaldehyde evaporates extremely slowly, the Schott bottle was placed in the laminar flow biohazard safety level 2 cabinet to facilitate evaporation. Crystals (colourless blocks) started forming on the rim on the outside of the bottle as the benzaldehyde evaporated. One of these crystals was sampled for XRD analysis.

**Table 2**  
Experimental details.

Crystal data	
Chemical formula	$C_{13}H_{11}N_3O \cdot C_7H_6O_2$
$M_r$	347.36
Crystal system, space group	Orthorhombic, <i>Pbca</i>
Temperature (K)	173
$a, b, c$ (Å)	11.7044 (17), 7.8531 (10), 38.253 (5)
$V$ (Å <sup>3</sup> )	3516.1 (8)
$Z$	8
Radiation type	Mo $K\alpha$
$\mu$ (mm <sup>-1</sup> )	0.09
Crystal size (mm)	0.30 × 0.22 × 0.16
Data collection	
Diffractometer	Bruker D8 Venture Microfocus with Photon III CCD area- detector
Absorption correction	Multi-scan ( <i>SADABS</i> ; Krause <i>et al.</i> , 2015)
$T_{min}, T_{max}$	0.709, 0.746
No. of measured, independent and observed [ $I > 2\sigma(I)$ ] reflections	73794, 4239, 3902
$R_{int}$	0.036
$(\sin \theta/\lambda)_{max}$ (Å <sup>-1</sup> )	0.660
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.054, 0.112, 1.18
No. of reflections	4239
No. of parameters	243
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}, \Delta\rho_{min}$ (e Å <sup>-3</sup> )	0.33, -0.21

Computer programs: *APEX3*, *SAINT-Plus* and *XPREP* (Bruker 2016), *SHELXT2014* (Sheldrick, 2015a), *SHELXL2019/2* (Sheldrick, 2015b) and *ORTEP-3 for Windows* and *WinGX* (Farrugia, 2012).

## 6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. C-bound H atoms were first located in the difference map, then positioned geometrically and allowed to ride on their respective parent atoms, with thermal displacement parameters 1.2 times of the parent C atom. The coordinates and isotropic displacement parameters of the O and N-bound H atoms involved in hydrogen-bonding interactions (H1 and H2) were allowed to refine freely.

## Acknowledgements

The authors would like to acknowledge the University of South Africa for the Masters and Doctoral Grant award.

## Funding information

Funding for this research was provided by: National Research Foundation (grant No. 118127; grant No. 117850).

## References

- Aakeröy, C. B., Hussain, I., Forbes, S. & Desper, J. (2007). *CrystEngComm*, **9**, 46–54.  
Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.

- Bruker (2016). *APEX3, SAINT-Plus and XPREP*. Bruker AXS Inc., Madison, Wisconsin, USA.
- Bruno, I. J., Cole, J. C., Edgington, P. R., Kessler, M., Macrae, C. F., McCabe, P., Pearson, J. & Taylor, R. (2002). *Acta Cryst.* **B58**, 389–397.
- Cully, M. (2014). *Nat. Rev. Drug Discov.* **13**, 257. <https://doi.org/10.1038/nrd4287>
- Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849–854.
- Groom, C. R., Bruno, I. J., Lightfoot, M. P. & Ward, S. C. (2016). *Acta Cryst.* **B72**, 171–179.
- Hean, D., Michael, J. P. & Lemmerer, A. (2018). *J. Mol. Struct.* **1157**, 693–707.
- Hearn, M. J. & Cynamon, M. H. (2004). *J. Antimicrob. Chemother.* **53**, 185–191.
- Krause, L., Herbst-Irmer, R., Sheldrick, G. M. & Stalke, D. (2015). *J. Appl. Cryst.* **48**, 3–10.
- Lemmerer, A., Bernstein, J. & Kahlenberg, V. (2010). *CrystEngComm*, **12**, 2856–2864.
- Pereira Almeida, W., Paes Koury, I. & Simoni, D. de A. (2016). *IUCrData*, **1**, x160752.
- Purushothaman, G., Angira, D. & Thiruvankatam, V. (2019). *J. Mol. Struct.* **1197**, 34–44.
- Sankar, M., Nowicka, E., Carter, E., Murphy, D., Knight, D., Bethell, D. & Hutchings, G. (2014). *Nat. Commun.* **5**, 3332. <https://doi.org/10.1038/ncomms4332>
- Shattock, T. R., Arora, K. K., Vishweshwar, P. & Zaworotko, M. J. (2008). *Cryst. Growth Des.* **8**, 4533–4545.
- Sheldrick, G. M. (2015a). *Acta Cryst.* **A71**, 3–8.
- Sheldrick, G. M. (2015b). *Acta Cryst.* **C71**, 3–8.
- Smith, M. G., Forbes, R. P. & Lemmerer, A. (2015). *Cryst. Growth Des.* **15**, 3813–3821.
- Suarez, J., Ranguelova, K., Jarzecki, A. A., Manzerova, J., Krymov, V., Zhao, X., Yu, S., Metlitsky, L., Gerfen, G. J. & Magliozzo, R. S. (2009). *J. Biol. Chem.* **284**, 7017–7029.
- Vishweshwar, P., McMahon, J. A., Bis, J. A. & Zaworotko, M. J. (2006). *J. Pharm. Sci.* **95**, 499–516.
- Wardell, S. M. S. V., de Souza, M. V. N., Wardell, J. L., Low, J. N. & Glidewell, C. (2007). *Acta Cryst.* **B63**, 879–895.

## supporting information

*Acta Cryst.* (2023). E79, 682-685 [https://doi.org/10.1107/S2056989023005698]

## Co-crystallization of *N'*-benzylidenepyridine-4-carbohydrazide and benzoic acid *via* autoxidation of benzaldehyde

Itumeleng B. Setshedi, Andreas Lemmerer and Mark G. Smith

### Computing details

Data collection: *APEX3* (Bruker, 2016); cell refinement: *S SAINT-Plus* (Bruker, 2016); data reduction: *S SAINT-Plus* and *XPREP* (Bruker 2016); program(s) used to solve structure: *SHELXT2014* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2019/2* (Sheldrick, 2015b); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *WinGX* (Farrugia, 2012).

### *N'*-[(2-methylphenyl)methylidene]pyridine-4-carbohydrazide; benzoic acid

#### Crystal data

$C_{13}H_{11}N_3O \cdot C_7H_6O_2$

$M_r = 347.36$

Orthorhombic, *Pbca*

$a = 11.7044$  (17) Å

$b = 7.8531$  (10) Å

$c = 38.253$  (5) Å

$V = 3516.1$  (8) Å<sup>3</sup>

$Z = 8$

$F(000) = 1456$

$D_x = 1.312$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 9857 reflections

$\theta = 2.8$ – $28.1^\circ$

$\mu = 0.09$  mm<sup>-1</sup>

$T = 173$  K

Block, colourless

$0.30 \times 0.22 \times 0.16$  mm

#### Data collection

Bruker D8 Venture Microfocus with Photon III

CCD area-detector

diffractometer

Graphite monochromator

$\omega$  scans

Absorption correction: multi-scan

(SADABS; Krause *et al.*, 2015)

$T_{\min} = 0.709$ ,  $T_{\max} = 0.746$

73794 measured reflections

4239 independent reflections

3902 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.036$

$\theta_{\max} = 28.0^\circ$ ,  $\theta_{\min} = 3.2^\circ$

$h = -15 \rightarrow 15$

$k = -8 \rightarrow 10$

$l = -49 \rightarrow 50$

#### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.054$

$wR(F^2) = 0.112$

$S = 1.18$

4239 reflections

243 parameters

0 restraints

0 constraints

Primary atom site location: dual

Secondary atom site location: dual

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0274P)^2 + 2.3245P]$

where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.33$  e Å<sup>-3</sup>

$\Delta\rho_{\min} = -0.21$  e Å<sup>-3</sup>

*Special details*

**Experimental.** Absorption corrections were made using the program SADABS (Sheldrick, 1996)

**Geometry.** All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

**Refinement.** The crystal structure was solved through direct methods using *SHELXT*. Non-hydrogen atoms were first refined isotropically followed by anisotropic refinement by full matrix least-squares calculations based on  $F^2$  using *SHELXL*.

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.63979 (12)	0.05245 (18)	0.40684 (4)	0.0224 (3)
C2	0.71809 (13)	0.15589 (19)	0.42397 (4)	0.0252 (3)
H2A	0.79074	0.178612	0.413922	0.03*
C3	0.68809 (13)	0.2256 (2)	0.45612 (4)	0.0278 (3)
H3	0.742184	0.295037	0.467926	0.033*
C4	0.51195 (13)	0.0976 (2)	0.45471 (4)	0.0294 (3)
H4	0.439948	0.077268	0.465382	0.035*
C5	0.53502 (13)	0.0207 (2)	0.42278 (4)	0.0273 (3)
H5	0.48046	-0.052042	0.412045	0.033*
C6	0.66282 (12)	-0.03373 (18)	0.37245 (4)	0.0217 (3)
C7	0.83497 (12)	0.04298 (19)	0.29981 (4)	0.0241 (3)
H7	0.875011	0.138628	0.308817	0.029*
C8	0.85942 (12)	-0.01869 (18)	0.26424 (4)	0.0225 (3)
C9	0.78697 (13)	-0.13366 (19)	0.24744 (4)	0.0275 (3)
H9	0.72116	-0.175227	0.259209	0.033*
C10	0.81045 (14)	-0.1876 (2)	0.21367 (4)	0.0325 (3)
H10	0.761097	-0.266749	0.202491	0.039*
C11	0.90585 (15)	-0.1264 (2)	0.19618 (4)	0.0329 (4)
H11	0.921511	-0.16284	0.17299	0.039*
C12	0.97809 (14)	-0.0121 (2)	0.21258 (4)	0.0330 (4)
H12	1.043564	0.029452	0.200633	0.04*
C13	0.95532 (13)	0.0423 (2)	0.24649 (4)	0.0288 (3)
H13	1.005101	0.121063	0.257586	0.035*
N1	0.73940 (11)	0.04239 (16)	0.35104 (3)	0.0243 (3)
N2	0.58653 (11)	0.19911 (17)	0.47113 (3)	0.0284 (3)
N3	0.75993 (11)	-0.03198 (16)	0.31860 (3)	0.0235 (3)
O1	0.61404 (9)	-0.16790 (13)	0.36487 (3)	0.0264 (2)
H1	0.7778 (16)	0.136 (3)	0.3562 (5)	0.035 (5)*
C14	0.62773 (13)	0.54692 (19)	0.57434 (4)	0.0255 (3)
C15	0.71251 (13)	0.6581 (2)	0.58553 (4)	0.0296 (3)
H15	0.776244	0.680985	0.570885	0.036*
C16	0.70449 (15)	0.7358 (2)	0.61798 (4)	0.0359 (4)
H16	0.762819	0.81133	0.625584	0.043*
C17	0.61171 (16)	0.7034 (3)	0.63923 (4)	0.0415 (4)
H17	0.605726	0.757686	0.661344	0.05*

C18	0.52751 (16)	0.5920 (3)	0.62834 (5)	0.0463 (5)
H18	0.463906	0.569504	0.643068	0.056*
C19	0.53547 (14)	0.5127 (2)	0.59597 (4)	0.0365 (4)
H19	0.477851	0.435326	0.588664	0.044*
C20	0.63940 (13)	0.4638 (2)	0.53921 (4)	0.0281 (3)
O2	0.55589 (10)	0.35567 (16)	0.53235 (3)	0.0357 (3)
O3	0.71911 (11)	0.49208 (17)	0.51992 (3)	0.0428 (3)
H2	0.565 (2)	0.303 (3)	0.5101 (6)	0.070 (7)*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0269 (7)	0.0212 (6)	0.0192 (6)	0.0038 (6)	−0.0001 (5)	0.0006 (5)
C2	0.0264 (7)	0.0278 (7)	0.0212 (6)	−0.0004 (6)	0.0014 (5)	0.0009 (6)
C3	0.0313 (8)	0.0302 (8)	0.0220 (7)	−0.0009 (6)	−0.0012 (6)	−0.0021 (6)
C4	0.0280 (7)	0.0339 (8)	0.0261 (7)	0.0010 (6)	0.0051 (6)	0.0002 (6)
C5	0.0266 (7)	0.0280 (7)	0.0273 (7)	−0.0013 (6)	0.0017 (6)	−0.0017 (6)
C6	0.0229 (7)	0.0218 (6)	0.0205 (6)	0.0042 (5)	−0.0018 (5)	−0.0001 (5)
C7	0.0262 (7)	0.0235 (7)	0.0226 (7)	−0.0011 (6)	−0.0010 (5)	−0.0012 (5)
C8	0.0258 (7)	0.0214 (6)	0.0203 (6)	0.0027 (6)	0.0007 (5)	0.0017 (5)
C9	0.0296 (7)	0.0304 (7)	0.0226 (7)	−0.0043 (6)	0.0001 (6)	0.0015 (6)
C10	0.0383 (8)	0.0353 (8)	0.0240 (7)	−0.0029 (7)	−0.0047 (6)	−0.0032 (6)
C11	0.0403 (9)	0.0392 (9)	0.0192 (7)	0.0088 (7)	0.0023 (6)	−0.0003 (6)
C12	0.0298 (8)	0.0415 (9)	0.0276 (8)	0.0025 (7)	0.0075 (6)	0.0043 (7)
C13	0.0267 (7)	0.0317 (8)	0.0279 (7)	−0.0022 (6)	0.0018 (6)	−0.0009 (6)
N1	0.0289 (6)	0.0238 (6)	0.0201 (5)	−0.0027 (5)	0.0019 (5)	−0.0051 (5)
N2	0.0340 (7)	0.0300 (7)	0.0212 (6)	0.0026 (5)	0.0023 (5)	−0.0012 (5)
N3	0.0266 (6)	0.0254 (6)	0.0185 (5)	0.0017 (5)	0.0003 (5)	−0.0031 (5)
O1	0.0295 (5)	0.0244 (5)	0.0252 (5)	−0.0022 (4)	0.0007 (4)	−0.0028 (4)
C14	0.0261 (7)	0.0264 (7)	0.0240 (7)	0.0016 (6)	−0.0001 (6)	0.0018 (6)
C15	0.0294 (7)	0.0300 (8)	0.0294 (7)	−0.0038 (6)	0.0006 (6)	0.0032 (6)
C16	0.0382 (9)	0.0349 (8)	0.0348 (8)	−0.0053 (7)	−0.0086 (7)	−0.0029 (7)
C17	0.0439 (10)	0.0542 (11)	0.0263 (8)	0.0004 (9)	−0.0018 (7)	−0.0125 (8)
C18	0.0338 (9)	0.0726 (14)	0.0324 (9)	−0.0093 (9)	0.0096 (7)	−0.0118 (9)
C19	0.0277 (8)	0.0499 (10)	0.0318 (8)	−0.0110 (7)	0.0037 (6)	−0.0070 (7)
C20	0.0312 (7)	0.0282 (7)	0.0250 (7)	−0.0002 (6)	0.0023 (6)	0.0016 (6)
O2	0.0366 (6)	0.0436 (7)	0.0271 (6)	−0.0091 (5)	0.0055 (5)	−0.0098 (5)
O3	0.0474 (8)	0.0493 (7)	0.0316 (6)	−0.0129 (6)	0.0156 (5)	−0.0054 (6)

*Geometric parameters (Å, °)*

C1—C2	1.389 (2)	C11—C12	1.384 (2)
C1—C5	1.392 (2)	C11—H11	0.95
C1—C6	1.5039 (19)	C12—C13	1.391 (2)
C2—C3	1.391 (2)	C12—H12	0.95
C2—H2A	0.95	C13—H13	0.95
C3—N2	1.336 (2)	N1—N3	1.3922 (16)
C3—H3	0.95	N1—H1	0.89 (2)

C4—N2	1.338 (2)	C14—C19	1.387 (2)
C4—C5	1.389 (2)	C14—C15	1.389 (2)
C4—H4	0.95	C14—C20	1.500 (2)
C5—H5	0.95	C15—C16	1.386 (2)
C6—O1	1.2328 (17)	C15—H15	0.95
C6—N1	1.3533 (19)	C16—C17	1.380 (2)
C7—N3	1.2784 (19)	C16—H16	0.95
C7—C8	1.4725 (19)	C17—C18	1.382 (3)
C7—H7	0.95	C17—H17	0.95
C8—C9	1.395 (2)	C18—C19	1.389 (2)
C8—C13	1.397 (2)	C18—H18	0.95
C9—C10	1.387 (2)	C19—H19	0.95
C9—H9	0.95	C20—O3	1.2101 (19)
C10—C11	1.388 (2)	C20—O2	1.3210 (19)
C10—H10	0.95	O2—H2	0.95 (2)
C2—C1—C5	118.66 (13)	C11—C12—C13	120.31 (15)
C2—C1—C6	123.89 (13)	C11—C12—H12	119.8
C5—C1—C6	117.41 (13)	C13—C12—H12	119.8
C3—C2—C1	118.72 (14)	C12—C13—C8	120.14 (15)
C3—C2—H2A	120.6	C12—C13—H13	119.9
C1—C2—H2A	120.6	C8—C13—H13	119.9
N2—C3—C2	122.90 (14)	C6—N1—N3	117.93 (12)
N2—C3—H3	118.6	C6—N1—H1	124.6 (12)
C2—C3—H3	118.6	N3—N1—H1	117.5 (12)
N2—C4—C5	123.01 (14)	C3—N2—C4	118.11 (13)
N2—C4—H4	118.5	C7—N3—N1	115.24 (12)
C5—C4—H4	118.5	C19—C14—C15	119.61 (14)
C4—C5—C1	118.57 (14)	C19—C14—C20	121.40 (14)
C4—C5—H5	120.7	C15—C14—C20	118.98 (14)
C1—C5—H5	120.7	C16—C15—C14	120.27 (15)
O1—C6—N1	122.82 (13)	C16—C15—H15	119.9
O1—C6—C1	120.47 (13)	C14—C15—H15	119.9
N1—C6—C1	116.70 (12)	C17—C16—C15	119.97 (16)
N3—C7—C8	120.11 (13)	C17—C16—H16	120
N3—C7—H7	119.9	C15—C16—H16	120
C8—C7—H7	119.9	C16—C17—C18	120.03 (16)
C9—C8—C13	119.10 (13)	C16—C17—H17	120
C9—C8—C7	121.35 (13)	C18—C17—H17	120
C13—C8—C7	119.53 (13)	C17—C18—C19	120.28 (16)
C10—C9—C8	120.42 (14)	C17—C18—H18	119.9
C10—C9—H9	119.8	C19—C18—H18	119.9
C8—C9—H9	119.8	C14—C19—C18	119.84 (16)
C9—C10—C11	120.16 (15)	C14—C19—H19	120.1
C9—C10—H10	119.9	C18—C19—H19	120.1
C11—C10—H10	119.9	O3—C20—O2	124.57 (15)
C12—C11—C10	119.87 (14)	O3—C20—C14	122.44 (15)
C12—C11—H11	120.1	O2—C20—C14	112.98 (13)

C10—C11—H11	120.1	C20—O2—H2	112.0 (15)
C5—C1—C2—C3	-0.9 (2)	C7—C8—C13—C12	-178.61 (14)
C6—C1—C2—C3	-178.49 (13)	O1—C6—N1—N3	2.5 (2)
C1—C2—C3—N2	-0.8 (2)	C1—C6—N1—N3	-178.20 (11)
N2—C4—C5—C1	-1.0 (2)	C2—C3—N2—C4	1.6 (2)
C2—C1—C5—C4	1.7 (2)	C5—C4—N2—C3	-0.7 (2)
C6—C1—C5—C4	179.49 (13)	C8—C7—N3—N1	-176.72 (12)
C2—C1—C6—O1	152.30 (14)	C6—N1—N3—C7	-178.41 (13)
C5—C1—C6—O1	-25.3 (2)	C19—C14—C15—C16	-0.6 (2)
C2—C1—C6—N1	-27.0 (2)	C20—C14—C15—C16	-179.49 (15)
C5—C1—C6—N1	155.35 (13)	C14—C15—C16—C17	-0.3 (3)
N3—C7—C8—C9	13.8 (2)	C15—C16—C17—C18	0.8 (3)
N3—C7—C8—C13	-167.96 (14)	C16—C17—C18—C19	-0.3 (3)
C13—C8—C9—C10	0.5 (2)	C15—C14—C19—C18	1.1 (3)
C7—C8—C9—C10	178.77 (14)	C20—C14—C19—C18	179.94 (17)
C8—C9—C10—C11	-0.6 (2)	C17—C18—C19—C14	-0.7 (3)
C9—C10—C11—C12	0.5 (3)	C19—C14—C20—O3	179.85 (17)
C10—C11—C12—C13	-0.3 (3)	C15—C14—C20—O3	-1.3 (2)
C11—C12—C13—C8	0.2 (2)	C19—C14—C20—O2	-1.6 (2)
C9—C8—C13—C12	-0.3 (2)	C15—C14—C20—O2	177.25 (14)

*Hydrogen-bond geometry (Å, °)*

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...O1 <sup>i</sup>	0.89 (2)	2.02 (2)	2.8981 (17)	171 (2)
O2—H2...N2	0.95 (2)	1.72 (3)	2.6693 (17)	176 (2)
C3—H3...O3	0.95	2.53	3.236 (2)	131

Symmetry code: (i)  $-x+3/2, y+1/2, z$ .