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Key indicators

Single-crystal X-ray study
 T = 120 K
 Mean $\sigma(\text{C}-\text{C}) = 0.003 \text{ \AA}$
 R factor = 0.040
 wR factor = 0.142
 Data-to-parameter ratio = 15.7

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

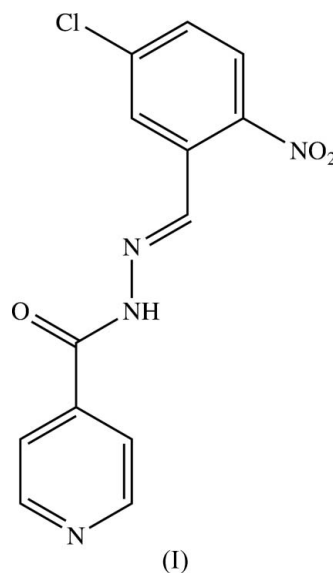
5-Chloro-2-nitrobenzaldehyde isonicotinoyl-hydrazone: a three-dimensional framework built from N—H···N and C—H···O hydrogen bonds

In the title compound, $\text{C}_{13}\text{H}_9\text{ClN}_4\text{O}_3$, the molecules are linked into a three-dimensional framework by one N—H···N hydrogen bond and three C—H···O hydrogen bonds.

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Comment

We report here the molecular and supramolecular structure of the title compound, (I) (Fig. 1), originally synthesized as a potential antimycobacterial agent (Junior *et al.*, 2005).



The coordination of the hydrazine atom N1 is planar and the central spacer unit between C1 and C21 is nearly planar, as shown by the leading torsion angles (Table 1); however, the two rings are significantly twisted out of this plane, although the two rings remain nearly parallel.

The molecules of (I) are linked by a combination of N—H···N and C—H···O hydrogen bonds (Table 2) into a three-dimensional framework structure, whose formation is readily analysed in terms of a number of very simple one-dimensional substructures, each formed by the action of a single hydrogen bond.

Amino atom N1 in the molecule at (x, y, z) acts as hydrogen-bond donor to pyridyl atom N4 in the molecule at $(-x, -\frac{1}{2} + y, \frac{3}{2} - z)$, so forming a $C(7)$ (Bernstein *et al.*, 1995) chain running parallel to the $[010]$ direction and generated by the 2_1 screw axis along $(0, y, \frac{3}{4})$ (Fig. 2). Atoms C3 and C24 in the molecule at (x, y, z) act as hydrogen-bond donors, respectively, to carbonyl atom O7 in the molecule at $(x, \frac{3}{2} - y, \frac{1}{2} + z)$, and nitro atom O221 in the molecule at $(x, \frac{1}{2} - y, -\frac{1}{2} + z)$, so forming two distinct $C(6)$ chains running parallel to the

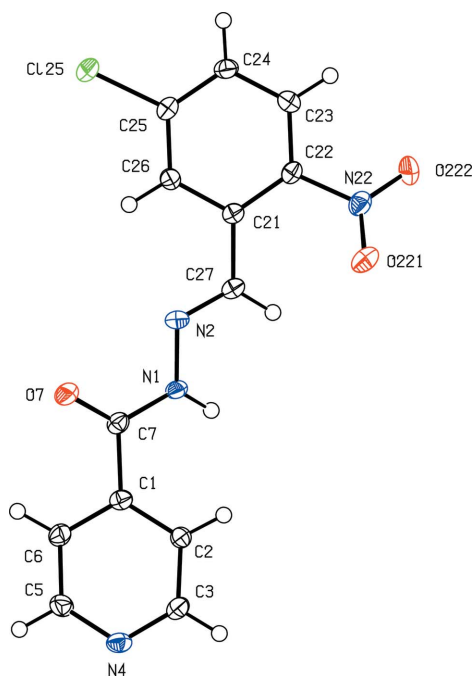


Figure 1
The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

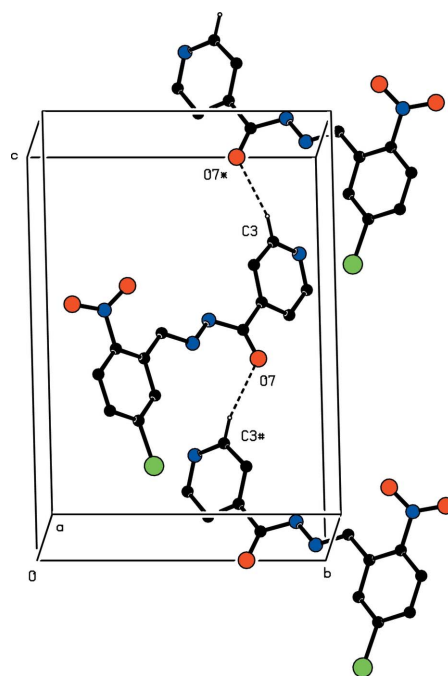


Figure 3
Part of the crystal structure of compound (I), showing the formation of a C(6) chain along [001] built from C—H...O(carbonyl) hydrogen bonds. For the sake of clarity, H atoms not involved in the motif shown have been omitted. Atoms marked with an asterisk (*) or a hash (#) are at the symmetry positions $(x, \frac{3}{2} - y, \frac{1}{2} + z)$ and $(x, \frac{3}{2} - y, -\frac{1}{2} + z)$, respectively.

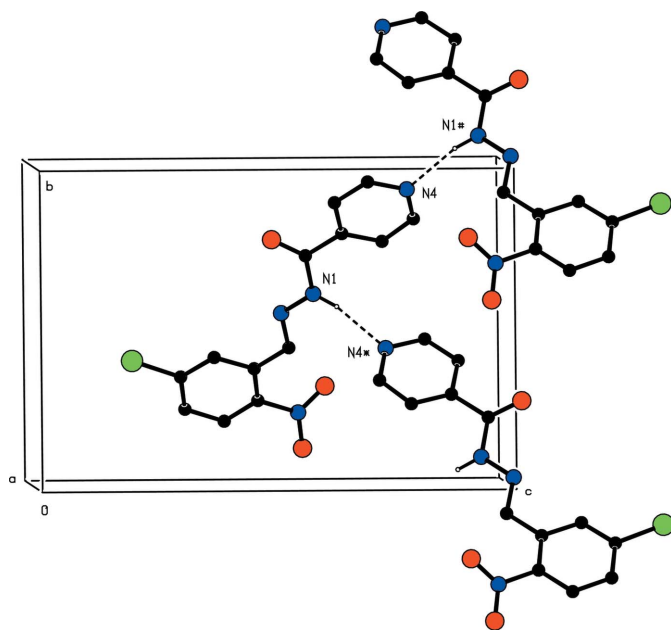


Figure 2
Part of the crystal structure of compound (I), showing the formation of a C(7) chain along [010] built from N—H...N hydrogen bonds. For the sake of clarity, H atoms bonded to C atoms have been omitted. Atoms marked with an asterisk (*) or a hash (#) are at the symmetry positions $(-x, -\frac{1}{2} + y, \frac{3}{2} - z)$ and $(-x, \frac{1}{2} + y, \frac{3}{2} - z)$, respectively.

[001] direction and generated respectively by the *c*-glide planes at $y = 0.75$ (Fig. 3) and $y = 0.25$ (Fig. 4). Finally, atom C5 in the molecule at (x, y, z) acts as hydrogen-bond donor to nitro atom O222 in the molecule at $(-1 + x, 1 + y, z)$, so generating by translation a C(12) chain running parallel to the $[1\bar{1}0]$ direction (Fig. 5).

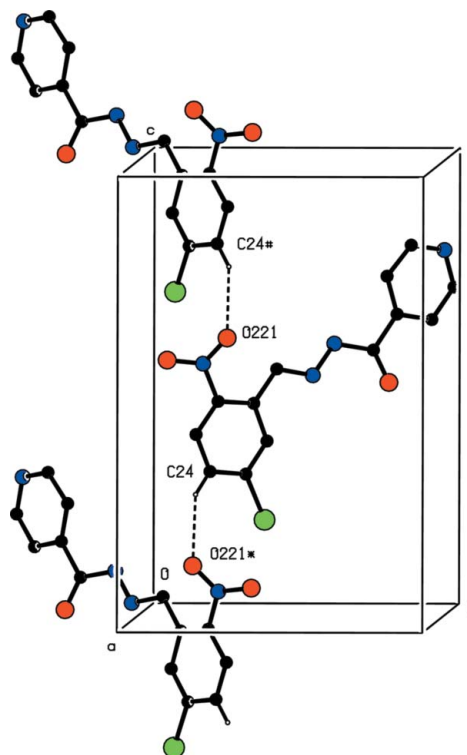


Figure 4
Part of the crystal structure of compound (I), showing the formation of a C(6) chain along [001] built from C—H...O(nitro) hydrogen bonds. For the sake of clarity, H atoms not involved in the motif shown have been omitted. Atoms marked with an asterisk (*) or a hash (#) are at the symmetry positions $(x, \frac{1}{2} - y, -\frac{1}{2} + z)$ and $(x, \frac{1}{2} - y, \frac{1}{2} + z)$, respectively.

The combination of the [010], [001] and $[1\bar{1}0]$ chains generates a single three-dimensional framework structure: it is notable that all three O atoms act as hydrogen-bond acceptors.

Experimental

Crystals of the title compound were prepared according to a published procedure (Junior *et al.*, 2005).

Crystal data

$C_{13}H_9ClN_4O_3$	$Z = 4$
$M_r = 304.69$	$D_x = 1.561 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 8.0597 (2) \text{ \AA}$	$\mu = 0.31 \text{ mm}^{-1}$
$b = 10.4797 (4) \text{ \AA}$	$T = 120 (2) \text{ K}$
$c = 15.4798 (6) \text{ \AA}$	Lath, colourless
$\beta = 97.383 (2)^\circ$	$0.34 \times 0.16 \times 0.08 \text{ mm}$
$V = 1296.63 (8) \text{ \AA}^3$	

Data collection

Bruker–Nonius KappaCCD diffractometer	17385 measured reflections
φ and ω scans	2980 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	2132 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.916$, $T_{\max} = 0.976$	$R_{\text{int}} = 0.053$
	$\theta_{\max} = 27.5^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0817P)^2 + 0.135P]$
$R[F^2 > 2\sigma(F^2)] = 0.040$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.143$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.09$	$\Delta\rho_{\max} = 0.36 \text{ e \AA}^{-3}$
2980 reflections	$\Delta\rho_{\min} = -0.36 \text{ e \AA}^{-3}$
190 parameters	
H-atom parameters constrained	

Table 1

Selected torsion angles ($^\circ$).

C2–C1–C7–N1	24.3 (3)	N1–N2–C27–C21	179.18 (16)
C1–C7–N1–N2	–174.60 (17)	N2–C27–C21–C22	163.1 (2)
C7–N1–N2–C27	175.55 (18)	C21–C22–N22–O221	–29.5 (3)

Table 2

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1–H1 \cdots N4 ⁱ	0.88	2.14	3.002 (3)	168
C3–H3 \cdots O7 ⁱⁱ	0.95	2.46	3.371 (3)	161
C5–H5 \cdots O222 ⁱⁱⁱ	0.95	2.35	3.175 (3)	145
C24–H24 \cdots O221 ^{iv}	0.95	2.47	3.327 (3)	151

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

All H atoms were located in difference maps, and then treated as riding atoms, with $C-H = 0.95 \text{ \AA}$, $N-H = 0.88 \text{ \AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$.

Data collection: COLLECT (Hooft, 1999); cell refinement: DENZO (Otwinowski & Minor, 1997) and COLLECT; data reduction: DENZO and COLLECT; program(s) used to solve structure:

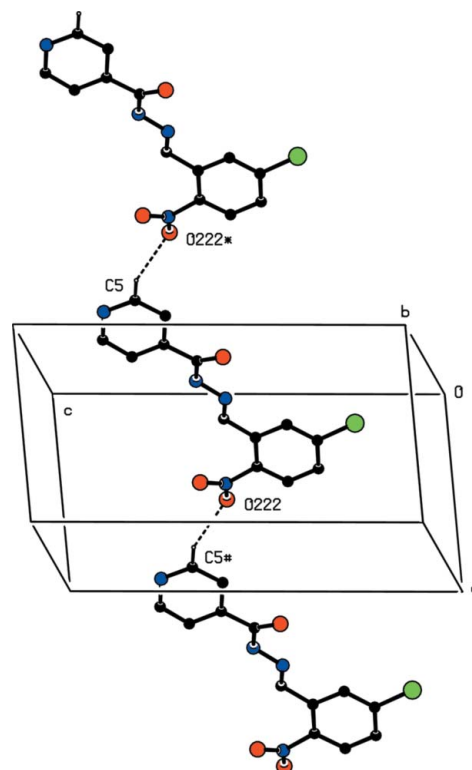


Figure 5

Part of the crystal structure of compound (I), showing the formation of a $C(12)$ chain along $[1\bar{1}0]$ built from $C-H\cdots O(\text{nitro})$ hydrogen bonds. For the sake of clarity, H atoms not involved in the motif shown have been omitted. Atoms marked with an asterisk (*) or a hash (#) are at the symmetry positions $(-1+x, 1+y, z)$ and $(1+x, -1+y, z)$, respectively.

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 National Crystallography Service, University of Southampton, England. The authors thank the staff of the Service for all their help and advice. JLW thanks CNPq for financial support.

References

- Bernstein, J., Davis, R. E., Shimoni, L. & Chang, N.-L. (1995). *Angew. Chem. Int. Ed. Engl.* **34**, 1555–1573.
- Ferguson, G. (1999). *PRPKAPPA*. University of Guelph, Canada.
- Hooft, R. W. W. (1999). *COLLECT*. Nonius BV, Delft, The Netherlands.
- Junior, I. N., Lourenco, M. C. S., das Gracas, M., Henriques, M. O., Ferreira, B., Vasconcelos, T. R. A., Peralta, M. A., de Oliveira, P. S. M., Wardell, S. M. S. V. & de Souza, M. V. N. (2005). *Lett. Drug. Des. Discov.* **2**, 563–566.
- McArdle, P. (2003). *OSCAIL for Windows*. Version 10. Crystallography Centre, Chemistry Department, NUI Galway, Ireland.
- Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
- Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.
- Sheldrick, G. M. (2003). *SADABS*. Version 2.10. University of Göttingen, Germany.
- Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.

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5-Chloro-2-nitrobenzaldehyde isonicotinoylhydrazone: a three-dimensional framework built from N—H⋯N and C—H⋯O hydrogen bonds

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5-Chloro-2-nitrobenzaldehyde isonicotinoylhydrazone

Crystal data

C₁₃H₉ClN₄O₃

M_r = 304.69

Monoclinic, *P*2₁/*c*

Hall symbol: -*P* 2ybc

a = 8.0597 (2) Å

b = 10.4797 (4) Å

c = 15.4798 (6) Å

β = 97.383 (2)°

V = 1296.63 (8) Å³

Z = 4

F(000) = 624

D_x = 1.561 Mg m⁻³

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 2980 reflections

θ = 2.4–27.5°

μ = 0.31 mm⁻¹

T = 120 K

Lath, colourless

0.34 × 0.16 × 0.08 mm

Data collection

Bruker–Nonius KappaCCD
diffractometer

Radiation source: Bruker–Nonius FR591
rotating anode

Graphite monochromator

Detector resolution: 9.091 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan
(SADABS; Sheldrick, 2003)

T_{min} = 0.916, *T_{max}* = 0.976

17385 measured reflections

2980 independent reflections

2132 reflections with *I* > 2σ(*I*)

R_{int} = 0.053

θ_{max} = 27.5°, θ_{min} = 2.4°

h = -10→10

k = -12→13

l = -20→20

Refinement

Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.040

wR(*F*²) = 0.143

S = 1.09

2980 reflections

190 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.0817*P*)² + 0.135*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} < 0.001

Δρ_{max} = 0.36 e Å⁻³

Δρ_{min} = -0.36 e Å⁻³

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.0301 (3)	0.7986 (2)	0.63712 (13)	0.0170 (5)
C2	0.0839 (3)	0.7718 (2)	0.72369 (13)	0.0182 (5)
C3	0.0234 (3)	0.8436 (2)	0.78804 (14)	0.0201 (5)
N4	-0.0852 (2)	0.93930 (19)	0.77059 (12)	0.0212 (4)
C5	-0.1344 (3)	0.9650 (2)	0.68648 (15)	0.0237 (5)
C6	-0.0799 (3)	0.8991 (2)	0.61819 (14)	0.0206 (5)
C7	0.0889 (3)	0.7284 (2)	0.56203 (13)	0.0188 (5)
O7	0.0865 (2)	0.77907 (16)	0.49088 (10)	0.0298 (4)
N1	0.1468 (2)	0.60745 (16)	0.57975 (11)	0.0179 (4)
N2	0.2145 (2)	0.54538 (17)	0.51433 (11)	0.0186 (4)
C27	0.2790 (2)	0.4353 (2)	0.53226 (13)	0.0184 (5)
C21	0.3495 (2)	0.3683 (2)	0.46136 (13)	0.0162 (4)
C22	0.4603 (3)	0.2650 (2)	0.47262 (13)	0.0174 (5)
N22	0.5337 (2)	0.22436 (19)	0.56006 (12)	0.0222 (4)
O221	0.55542 (19)	0.30530 (16)	0.61789 (10)	0.0272 (4)
O222	0.5726 (2)	0.11140 (16)	0.57015 (11)	0.0324 (4)
C23	0.5128 (3)	0.1994 (2)	0.40315 (14)	0.0200 (5)
C24	0.4612 (3)	0.2393 (2)	0.31871 (14)	0.0208 (5)
C25	0.3551 (2)	0.3436 (2)	0.30634 (13)	0.0192 (5)
Cl25	0.28646 (7)	0.39720 (6)	0.20175 (3)	0.0280 (2)
C26	0.2974 (3)	0.4059 (2)	0.37533 (13)	0.0178 (5)
H2	0.1616	0.7048	0.7389	0.022*
H3	0.0609	0.8239	0.8472	0.024*
H5	-0.2116	1.0328	0.6729	0.028*
H6	-0.1171	0.9223	0.5596	0.025*
H1	0.1352	0.5678	0.6287	0.022*
H27	0.2814	0.3985	0.5885	0.022*
H23	0.5839	0.1272	0.4135	0.024*
H24	0.4976	0.1964	0.2705	0.025*
H26	0.2214	0.4750	0.3643	0.021*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0200 (10)	0.0147 (11)	0.0167 (11)	-0.0042 (9)	0.0039 (8)	-0.0017 (8)
C2	0.0241 (11)	0.0135 (11)	0.0172 (11)	-0.0003 (9)	0.0025 (9)	0.0003 (9)
C3	0.0263 (11)	0.0186 (12)	0.0152 (10)	-0.0041 (10)	0.0017 (9)	0.0000 (9)
N4	0.0221 (9)	0.0224 (10)	0.0190 (10)	-0.0006 (8)	0.0022 (7)	-0.0066 (8)
C5	0.0239 (11)	0.0231 (13)	0.0227 (12)	0.0051 (10)	-0.0028 (9)	-0.0062 (10)
C6	0.0226 (11)	0.0205 (12)	0.0173 (11)	-0.0010 (9)	-0.0025 (9)	-0.0027 (9)
C7	0.0217 (11)	0.0186 (12)	0.0157 (11)	-0.0025 (9)	0.0011 (8)	-0.0007 (9)
O7	0.0481 (11)	0.0256 (9)	0.0169 (8)	0.0077 (8)	0.0085 (7)	0.0031 (7)
N1	0.0230 (9)	0.0185 (10)	0.0133 (9)	0.0005 (8)	0.0058 (7)	-0.0006 (7)
N2	0.0208 (9)	0.0190 (10)	0.0165 (9)	-0.0006 (8)	0.0049 (7)	-0.0043 (8)
C27	0.0205 (11)	0.0206 (12)	0.0146 (10)	-0.0028 (9)	0.0035 (8)	-0.0010 (9)

C21	0.0167 (10)	0.0162 (11)	0.0158 (10)	-0.0027 (9)	0.0023 (8)	-0.0009 (8)
C22	0.0185 (10)	0.0184 (11)	0.0150 (10)	-0.0031 (9)	0.0009 (8)	0.0015 (9)
N22	0.0203 (9)	0.0286 (12)	0.0180 (9)	0.0000 (8)	0.0034 (8)	0.0035 (9)
O221	0.0276 (9)	0.0361 (10)	0.0174 (8)	-0.0046 (7)	0.0013 (7)	-0.0014 (7)
O222	0.0414 (10)	0.0277 (10)	0.0281 (10)	0.0133 (8)	0.0039 (8)	0.0102 (7)
C23	0.0193 (11)	0.0182 (12)	0.0227 (12)	0.0018 (9)	0.0034 (9)	-0.0017 (9)
C24	0.0226 (11)	0.0234 (12)	0.0170 (11)	0.0015 (9)	0.0052 (9)	-0.0031 (9)
C25	0.0198 (11)	0.0225 (12)	0.0150 (10)	-0.0019 (9)	0.0012 (8)	0.0015 (9)
Cl25	0.0349 (3)	0.0344 (4)	0.0150 (3)	0.0088 (3)	0.0045 (2)	0.0037 (2)
C26	0.0180 (10)	0.0170 (12)	0.0183 (11)	0.0012 (9)	0.0023 (8)	0.0006 (9)

Geometric parameters (Å, °)

C1—C2	1.383 (3)	C27—C21	1.477 (3)
C1—C6	1.384 (3)	C27—H27	0.95
C1—C7	1.503 (3)	C21—C22	1.400 (3)
C2—C3	1.386 (3)	C21—C26	1.400 (3)
C2—H2	0.95	C22—C23	1.387 (3)
C3—N4	1.336 (3)	C22—N22	1.469 (3)
C3—H3	0.95	N22—O221	1.229 (2)
N4—C5	1.339 (3)	N22—O222	1.229 (2)
C5—C6	1.380 (3)	C23—C24	1.385 (3)
C5—H5	0.95	C23—H23	0.95
C6—H6	0.95	C24—C25	1.386 (3)
C7—O7	1.221 (3)	C24—H24	0.95
C7—N1	1.366 (3)	C25—C26	1.382 (3)
N1—N2	1.374 (2)	C25—Cl25	1.736 (2)
N1—H1	0.88	C26—H26	0.95
N2—C27	1.281 (3)		
C2—C1—C6	118.19 (19)	N2—C27—H27	121.6
C2—C1—C7	123.99 (19)	C21—C27—H27	121.6
C6—C1—C7	117.77 (19)	C22—C21—C26	116.40 (19)
C1—C2—C3	119.4 (2)	C22—C21—C27	125.13 (19)
C1—C2—H2	120.3	C26—C21—C27	118.42 (18)
C3—C2—H2	120.3	C23—C22—C21	122.67 (19)
N4—C3—C2	123.0 (2)	C23—C22—N22	116.36 (18)
N4—C3—H3	118.5	C21—C22—N22	120.91 (18)
C2—C3—H3	118.5	O221—N22—O222	124.18 (19)
C3—N4—C5	116.87 (18)	O221—N22—C22	118.29 (18)
N4—C5—C6	124.1 (2)	O222—N22—C22	117.52 (19)
N4—C5—H5	117.9	C24—C23—C22	119.8 (2)
C6—C5—H5	117.9	C24—C23—H23	120.1
C5—C6—C1	118.4 (2)	C22—C23—H23	120.1
C5—C6—H6	120.8	C23—C24—C25	118.30 (19)
C1—C6—H6	120.8	C23—C24—H24	120.9
O7—C7—N1	123.4 (2)	C25—C24—H24	120.9
O7—C7—C1	121.0 (2)	C26—C25—C24	121.9 (2)

N1—C7—C1	115.62 (18)	C26—C25—C125	117.99 (17)
C7—N1—N2	116.68 (17)	C24—C25—C125	120.07 (16)
C7—N1—H1	122.6	C25—C26—C21	120.81 (19)
N2—N1—H1	120.5	C25—C26—H26	119.6
C27—N2—N1	117.08 (18)	C21—C26—H26	119.6
N2—C27—C21	116.84 (19)		
C6—C1—C2—C3	1.6 (3)	C26—C21—C22—C23	-2.1 (3)
C7—C1—C2—C3	179.1 (2)	C27—C21—C22—C23	175.0 (2)
C1—C2—C3—N4	-0.3 (3)	C26—C21—C22—N22	174.93 (18)
C2—C3—N4—C5	-0.5 (3)	C27—C21—C22—N22	-8.0 (3)
C3—N4—C5—C6	0.0 (3)	C23—C22—N22—O221	147.72 (19)
N4—C5—C6—C1	1.2 (3)	C21—C22—N22—O221	-29.5 (3)
C2—C1—C6—C5	-2.0 (3)	C23—C22—N22—O222	-30.9 (3)
C7—C1—C6—C5	-179.66 (19)	C21—C22—N22—O222	151.9 (2)
C2—C1—C7—O7	-154.4 (2)	C21—C22—C23—C24	3.0 (3)
C6—C1—C7—O7	23.1 (3)	N22—C22—C23—C24	-174.15 (19)
C2—C1—C7—N1	24.3 (3)	C22—C23—C24—C25	-1.2 (3)
C1—C7—N1—N2	-174.60 (17)	C23—C24—C25—C26	-1.5 (3)
C7—N1—N2—C27	175.55 (18)	C23—C24—C25—C125	-179.84 (16)
N1—N2—C27—C21	179.18 (16)	C24—C25—C26—C21	2.4 (3)
N2—C27—C21—C22	163.1 (2)	C125—C25—C26—C21	-179.24 (16)
C6—C1—C7—N1	-158.12 (18)	C22—C21—C26—C25	-0.6 (3)
O7—C7—N1—N2	4.2 (3)	C27—C21—C26—C25	-177.83 (19)
N2—C27—C21—C26	-19.9 (3)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1 \cdots N4 ⁱ	0.88	2.14	3.002 (3)	168
C3—H3 \cdots O7 ⁱⁱ	0.95	2.46	3.371 (3)	161
C5—H5 \cdots O222 ⁱⁱⁱ	0.95	2.35	3.175 (3)	145
C24—H24 \cdots O221 ^{iv}	0.95	2.47	3.327 (3)	151

Symmetry codes: (i) $-x, y-1/2, -z+3/2$; (ii) $x, -y+3/2, z+1/2$; (iii) $x-1, y+1, z$; (iv) $x, -y+1/2, z-1/2$.