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***N'*-[*E*-(1-Methyl-1*H*-pyrrol-2-yl)methylidene]pyridine-4-carbohydrazide. Corrigendum**

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The name of one of the authors in the paper by Hussain *et al.* [*Acta Cryst.* (2010), E66, o1881] is corrected.

In the paper by Hussain *et al.* (2010), the last author is incorrectly given as 'Muhammad Mazhar'. The correct name of the last author should be 'Mazhar Hussain' as above.

References

Hussain, A., Tahir, M. N., Shafiq, Z., Yaqub, M. & Mazhar, M. (2010). *Acta Cryst.* E66, o1881.

N'-[*E*-(1-Methyl-1*H*-pyrrol-2-yl)methylidene]pyridine-4-carbohydrazide

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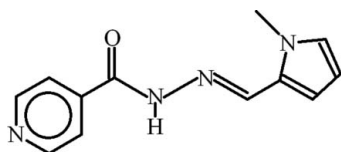
Received 27 June 2010; accepted 28 June 2010

Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.045; wR factor = 0.127; data-to-parameter ratio = 18.1.

In the title compound, $\text{C}_{12}\text{H}_{12}\text{N}_4\text{O}$, the pyridine and pyrrole rings are inclined at an angle of 29.22 (8) $^\circ$ and an intramolecular $\text{C}-\text{H}\cdots\text{N}$ interaction generates a $S(6)$ ring. In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds, forming (010) $C(7)$ chains. The chains are cross-linked by weak $\text{C}-\text{H}\cdots\text{O}$ interactions, which generate $R_2^2(18)$ ring motifs within an infinite sheet. Finally, two $\text{C}-\text{H}\cdots\pi$ interactions are present, where the $\text{C}-\text{H}$ groups are from the pyridine ring and π is the pyrrole ring.

Related literature

For background information on Schiff bases containing heterocyclic rings and for related structures, see: Shafiq *et al.*, (2009*a,b*); Hussain *et al.* (2010) For graph-set notation, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{12}\text{N}_4\text{O}$

$M_r = 228.26$

Monoclinic, $P2_1/n$

$a = 8.2134$ (3) Å

$b = 10.6740$ (4) Å

$c = 13.1332$ (4) Å

$\beta = 96.938$ (2) $^\circ$

$V = 1142.95$ (7) Å³

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 0.09$ mm⁻¹

$T = 296$ K

$0.24 \times 0.18 \times 0.15$ mm

Data collection

Bruker Kappa APEXII CCD

diffractometer

Absorption correction: multi-scan

(*SADABS*; Bruker, 2005)

$T_{\min} = 0.980$, $T_{\max} = 0.985$

12030 measured reflections

2803 independent reflections

2023 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.029$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.127$

$S = 1.04$

2803 reflections

155 parameters

H-atom parameters constrained

$\Delta\rho_{\text{max}} = 0.28$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, $^\circ$).

Cg1 is the centroid of the C8–C11/N4 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N2}-\text{H2}\cdots\text{N1}^{\text{i}}$	0.86	2.19	3.0205 (18)	163
$\text{C4}-\text{H4}\cdots\text{O1}^{\text{ii}}$	0.93	2.54	3.3821 (19)	150
$\text{C12}-\text{H12B}\cdots\text{O1}^{\text{iii}}$	0.96	2.55	3.450 (2)	156
$\text{C12}-\text{H12C}\cdots\text{N3}$	0.96	2.36	3.025 (2)	126
$\text{C2}-\text{H2A}\cdots\text{Cg1}^{\text{iv}}$	0.93	2.83	3.3258 (16)	114
$\text{C5}-\text{H5}\cdots\text{Cg1}^{\text{v}}$	0.93	2.71	3.4669 (17)	139

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

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

The authors acknowledge the provision of funds for the purchase of diffractometer and encouragement by Dr Muhammad Akram Chaudhary, Vice Chancellor, University of Sargodha, Pakistan.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5530).

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supporting information

Acta Cryst. (2010). E66, o1881 [doi:10.1107/S1600536810025341]

***N'*-[*E*]-[1-Methyl-1*H*-pyrrol-2-yl)methylidene]pyridine-4-carbohydrazide**

Abid Hussain, M. Nawaz Tahir, Zahid Shafiq, Muhammad Yaqub and Muhammad Mazhar

S1. Comment

We have reported crystal structures of Schiff bases with N-containing aromatic ring (Shafiq *et al.*, 2009*a*, 2009*b*), (Hussain *et al.*, 2010) and as a part of this project, we report herein the structure and synthesis of the title compound (I, Fig. 1).

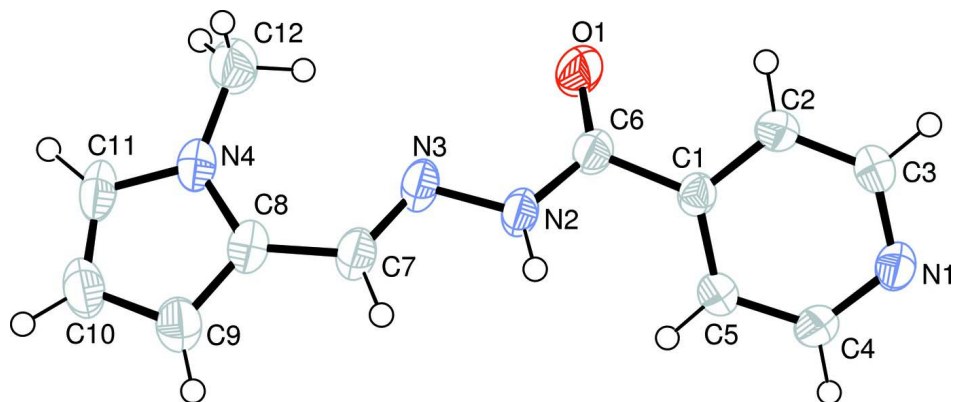
In (I) the pyridine ring A (C1–C5/N1), the central moiety B (O1/C6/N2/N3/C7) and the pyrrol moiety C (C8—C11/N4/C12) are planar with r. m. s. deviations of 0.0345, 0.0285 and 0.0276 Å, respectively. The dihedral angle between A/B, A/C and B/C is 38.32 (8)°, 29.22 (8)° and 9.44 (13)°, respectively. In title molecule, there exist intra as well inter-molecular H-bondings (Table 1). The molecules form infinite one dimensional polymeric chains extending along the *b* axis (Fig. 2), if only strong H-bondings are considered. If the strong as well as weak H-bondings are considered then the molecules form two-dimensional polymeric chains with $R_2^2(18)$ (Bernstein *et al.*, 1995) ring motifs (Fig. 3). The C—H $\cdots\pi$ interactions (Table 1) also play important role in stabilizing the molecules.

S2. Experimental

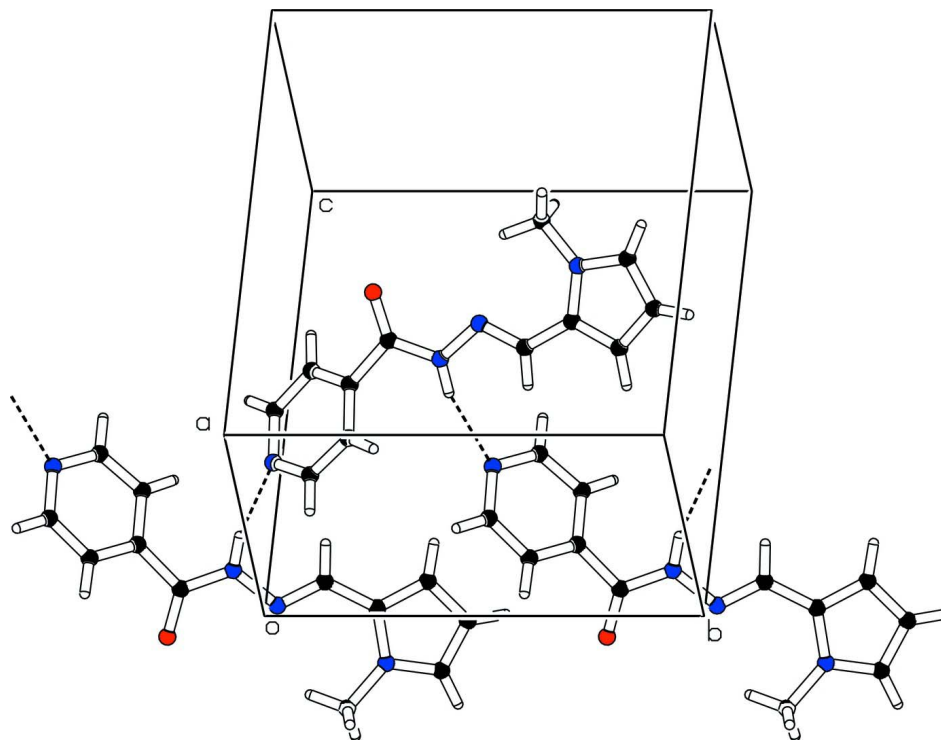
To a hot stirred solution of isoniazid (1.37 g, 0.01 mole) in ethanol 15 ml was added *N*-methylpyrrol-2-carboxaldehyde (1.1 ml, 0.01 mol). The resultant mixture was then heated under reflux. The reaction was monitored through TLC. After an hour, the precipitate were formed. The reaction mixture was further heated for 30 min. The resultant crude material was recrystallized in 1,4-dioxane:ethanol (1:4) to afford red prisms of (I).

S3. Refinement

The H-atoms were positioned geometrically (N–H = 0.86 Å, C–H = 0.93–0.96 Å) and refined as riding with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C}, \text{N})$, where $x = 1.5$ for methyl and $x = 1.2$ for all other H-atoms.

**Figure 1**

View of (I) with displacement ellipsoids drawn at the 50% probability level. H-atoms are shown by small circles of arbitrary radius.

**Figure 2**

The partial packing of (I), which shows that molecules form infinite one dimensional polymeric chains extending along *b* axis.

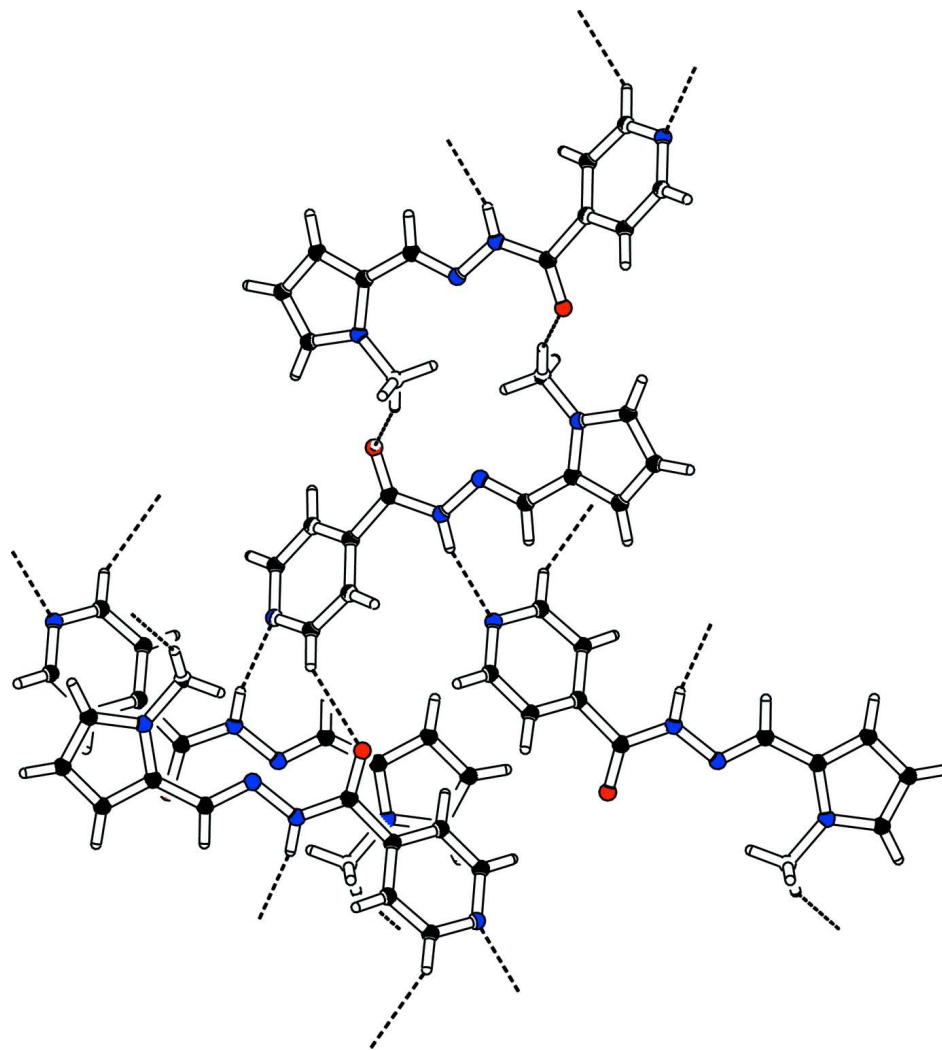


Figure 3

The partial packing (*PLATON*; Spek, 2009) which shows that molecules form $R_2^2(18)$ ring motifs in the infinite polymeric chains.

***N'*-[*(E)*-(1-Methyl-1*H*-pyrrol-2-yl)methylidene]pyridine-4-carbohydrazide**

Crystal data

$C_{12}H_{12}N_4O$

$M_r = 228.26$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 8.2134$ (3) Å

$b = 10.6740$ (4) Å

$c = 13.1332$ (4) Å

$\beta = 96.938$ (2)°

$V = 1142.95$ (7) Å³

$Z = 4$

$F(000) = 480$

$D_x = 1.326$ Mg m⁻³

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

Cell parameters from 1770 reflections

$\theta = 2.6$ – 28.4 °

$\mu = 0.09$ mm⁻¹

$T = 296$ K

Prism, red

$0.24 \times 0.18 \times 0.15$ mm

Data collection

Bruker Kappa APEXII CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 Detector resolution: 7.50 pixels mm⁻¹
 ω scans
 Absorption correction: multi-scan
 (SADABS; Bruker, 2005)
 $T_{\min} = 0.980$, $T_{\max} = 0.985$

12030 measured reflections
 2803 independent reflections
 2023 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$
 $\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 2.5^\circ$
 $h = -10 \rightarrow 10$
 $k = -14 \rightarrow 14$
 $l = -17 \rightarrow 17$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.045$
 $wR(F^2) = 0.127$
 $S = 1.04$
 2803 reflections
 155 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
 $w = 1/[\sigma^2(F_o^2) + (0.0618P)^2 + 0.235P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.28 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.68517 (16)	0.25900 (11)	0.46989 (10)	0.0596 (4)
N1	0.25031 (16)	0.01556 (12)	0.25463 (10)	0.0424 (4)
N2	0.55170 (15)	0.41010 (11)	0.37015 (10)	0.0388 (4)
N3	0.65784 (16)	0.50489 (12)	0.40795 (10)	0.0417 (4)
N4	0.85194 (16)	0.74061 (12)	0.46127 (10)	0.0411 (4)
C1	0.46019 (16)	0.19827 (12)	0.34909 (10)	0.0315 (4)
C2	0.40449 (19)	0.10059 (13)	0.40461 (11)	0.0374 (4)
C3	0.30152 (19)	0.01259 (14)	0.35500 (12)	0.0414 (5)
C4	0.3056 (2)	0.11012 (14)	0.20128 (11)	0.0421 (5)
C5	0.40918 (19)	0.20229 (14)	0.24441 (11)	0.0378 (4)
C6	0.57736 (18)	0.29160 (14)	0.40312 (11)	0.0371 (4)
C7	0.62947 (19)	0.61063 (14)	0.36395 (12)	0.0416 (5)
C8	0.72195 (19)	0.72388 (14)	0.38651 (12)	0.0404 (5)
C9	0.6991 (2)	0.83500 (15)	0.33312 (13)	0.0497 (5)
C10	0.8163 (2)	0.92006 (16)	0.37628 (13)	0.0539 (6)
C11	0.9081 (2)	0.85957 (15)	0.45424 (13)	0.0496 (6)
C12	0.9127 (2)	0.65484 (17)	0.54194 (14)	0.0574 (6)

H2	0.46924	0.42717	0.32551	0.0466*
H2A	0.43635	0.09445	0.47482	0.0449*
H3	0.26551	-0.05268	0.39352	0.0497*
H4	0.27213	0.11375	0.13112	0.0505*
H5	0.44438	0.26614	0.20412	0.0454*
H7	0.54104	0.61488	0.31268	0.0500*
H9	0.61937	0.85044	0.27801	0.0597*
H10	0.82924	1.00246	0.35569	0.0646*
H11	0.99575	0.89429	0.49628	0.0595*
H12A	0.86108	0.67234	0.60217	0.0860*
H12B	1.02928	0.66469	0.55737	0.0860*
H12C	0.88828	0.57043	0.51995	0.0860*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0566 (8)	0.0471 (7)	0.0666 (8)	-0.0071 (6)	-0.0275 (6)	0.0054 (6)
N1	0.0438 (8)	0.0334 (7)	0.0475 (7)	-0.0070 (6)	-0.0048 (5)	-0.0019 (5)
N2	0.0359 (7)	0.0296 (6)	0.0480 (7)	-0.0057 (5)	-0.0070 (5)	-0.0040 (5)
N3	0.0389 (7)	0.0336 (7)	0.0511 (7)	-0.0081 (5)	-0.0010 (5)	-0.0087 (5)
N4	0.0427 (7)	0.0335 (7)	0.0474 (7)	-0.0083 (5)	0.0062 (6)	-0.0074 (5)
C1	0.0297 (7)	0.0266 (7)	0.0374 (7)	0.0006 (5)	0.0003 (5)	-0.0018 (5)
C2	0.0437 (8)	0.0331 (8)	0.0340 (7)	-0.0001 (6)	-0.0012 (6)	0.0025 (6)
C3	0.0458 (9)	0.0331 (8)	0.0449 (8)	-0.0071 (7)	0.0035 (6)	0.0052 (6)
C4	0.0509 (9)	0.0385 (8)	0.0341 (7)	-0.0045 (7)	-0.0061 (6)	-0.0011 (6)
C5	0.0453 (9)	0.0318 (7)	0.0359 (7)	-0.0047 (6)	0.0030 (6)	0.0032 (6)
C6	0.0348 (8)	0.0343 (8)	0.0406 (7)	-0.0037 (6)	-0.0022 (6)	-0.0022 (6)
C7	0.0380 (8)	0.0344 (8)	0.0509 (9)	-0.0047 (7)	-0.0012 (7)	-0.0074 (7)
C8	0.0396 (8)	0.0351 (8)	0.0470 (8)	-0.0051 (6)	0.0069 (7)	-0.0087 (6)
C9	0.0580 (10)	0.0381 (9)	0.0529 (9)	-0.0047 (8)	0.0064 (8)	-0.0029 (7)
C10	0.0729 (12)	0.0331 (8)	0.0580 (10)	-0.0118 (8)	0.0178 (9)	-0.0040 (7)
C11	0.0560 (10)	0.0380 (9)	0.0570 (10)	-0.0198 (8)	0.0162 (8)	-0.0155 (8)
C12	0.0538 (11)	0.0461 (10)	0.0680 (11)	-0.0058 (8)	-0.0098 (8)	0.0025 (9)

Geometric parameters (Å, °)

O1—C6	1.219 (2)	C7—C8	1.439 (2)
N1—C3	1.335 (2)	C8—C9	1.379 (2)
N1—C4	1.339 (2)	C9—C10	1.393 (2)
N2—N3	1.3877 (18)	C10—C11	1.359 (2)
N2—C6	1.3453 (19)	C2—H2A	0.9300
N3—C7	1.277 (2)	C3—H3	0.9300
N4—C8	1.372 (2)	C4—H4	0.9300
N4—C11	1.358 (2)	C5—H5	0.9300
N4—C12	1.443 (2)	C7—H7	0.9300
N2—H2	0.8600	C9—H9	0.9300
C1—C2	1.3815 (19)	C10—H10	0.9300
C1—C5	1.3886 (19)	C11—H11	0.9300

C1—C6	1.502 (2)	C12—H12A	0.9600
C2—C3	1.374 (2)	C12—H12B	0.9600
C4—C5	1.377 (2)	C12—H12C	0.9600
C3—N1—C4	116.64 (13)	N4—C11—C10	109.47 (15)
N3—N2—C6	120.21 (12)	C1—C2—H2A	120.00
N2—N3—C7	114.22 (13)	C3—C2—H2A	120.00
C8—N4—C11	108.33 (13)	N1—C3—H3	118.00
C8—N4—C12	127.81 (13)	C2—C3—H3	118.00
C11—N4—C12	123.58 (14)	N1—C4—H4	118.00
C6—N2—H2	120.00	C5—C4—H4	118.00
N3—N2—H2	120.00	C1—C5—H5	121.00
C2—C1—C6	119.03 (12)	C4—C5—H5	121.00
C2—C1—C5	117.80 (13)	N3—C7—H7	117.00
C5—C1—C6	123.13 (12)	C8—C7—H7	117.00
C1—C2—C3	119.32 (13)	C8—C9—H9	126.00
N1—C3—C2	123.67 (14)	C10—C9—H9	126.00
N1—C4—C5	123.72 (14)	C9—C10—H10	127.00
C1—C5—C4	118.86 (13)	C11—C10—H10	127.00
N2—C6—C1	113.89 (12)	N4—C11—H11	125.00
O1—C6—N2	124.87 (14)	C10—C11—H11	125.00
O1—C6—C1	121.24 (13)	N4—C12—H12A	109.00
N3—C7—C8	125.91 (15)	N4—C12—H12B	109.00
N4—C8—C9	107.35 (13)	N4—C12—H12C	109.00
C7—C8—C9	125.57 (15)	H12A—C12—H12B	109.00
N4—C8—C7	127.04 (14)	H12A—C12—H12C	109.00
C8—C9—C10	107.99 (15)	H12B—C12—H12C	109.00
C9—C10—C11	106.87 (15)		
C4—N1—C3—C2	0.6 (2)	C2—C1—C5—C4	0.4 (2)
C3—N1—C4—C5	-0.5 (2)	C6—C1—C5—C4	177.90 (14)
C6—N2—N3—C7	173.81 (14)	C2—C1—C6—O1	38.1 (2)
N3—N2—C6—O1	4.1 (2)	C2—C1—C6—N2	-142.47 (14)
N3—N2—C6—C1	-175.38 (12)	C5—C1—C6—O1	-139.38 (16)
N2—N3—C7—C8	-178.82 (14)	C5—C1—C6—N2	40.08 (19)
C11—N4—C8—C7	177.64 (15)	C1—C2—C3—N1	-0.2 (2)
C11—N4—C8—C9	0.04 (17)	N1—C4—C5—C1	0.0 (2)
C12—N4—C8—C7	-8.4 (3)	N3—C7—C8—N4	-2.9 (3)
C12—N4—C8—C9	174.00 (15)	N3—C7—C8—C9	174.33 (16)
C8—N4—C11—C10	0.12 (19)	N4—C8—C9—C10	-0.18 (18)
C12—N4—C11—C10	-174.15 (15)	C7—C8—C9—C10	-177.82 (15)
C5—C1—C2—C3	-0.3 (2)	C8—C9—C10—C11	0.25 (19)
C6—C1—C2—C3	-177.89 (14)	C9—C10—C11—N4	-0.23 (19)

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

Cg1 is the centroid of the C8—C11/N4 ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N2—H2 \cdots N1 ⁱ	0.86	2.19	3.0205 (18)	163
C4—H4 \cdots O1 ⁱⁱ	0.93	2.54	3.3821 (19)	150
C12—H12B \cdots O1 ⁱⁱⁱ	0.96	2.55	3.450 (2)	156
C12—H12C \cdots N3	0.96	2.36	3.025 (2)	126
C2—H2A \cdots Cg1 ^{iv}	0.93	2.83	3.3258 (16)	114
C5—H5 \cdots Cg1 ^v	0.93	2.71	3.4669 (17)	139

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