

2-(5-Methoxy-2-methyl-1*H*-indol-3-yl)-*N'*-[(1*E*,2*E*)-3-phenylprop-2-en-1-yl-*idene*]acetohydrazide

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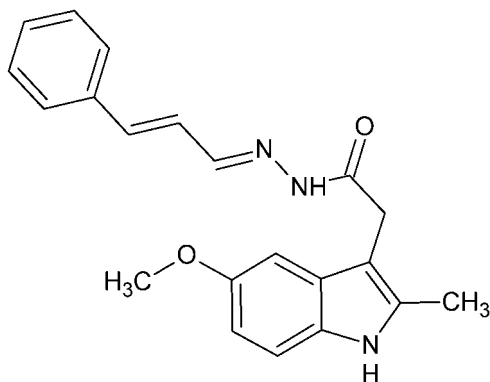
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Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.045; wR factor = 0.126; data-to-parameter ratio = 18.7.

The title compound, $\text{C}_{21}\text{H}_{21}\text{N}_3\text{O}_2$, adopts a J-shaped conformation which appears to be at least partially directed by a weak intramolecular $\text{C}-\text{H}\cdots\text{N}$ hydrogen bond. In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds into $R_2^2(8)$ and $R_2^2(14)$ cyclic dimers, which form a chain running parallel to the b axis.

Related literature

For general background to side-effect toxicity of non-steroidal anti-inflammatory drugs (NSAIDs), see: Agrawal *et al.* (2010); Champion *et al.* (1997); Allan & Fletcher (1990). For reduction of GI toxicity attributed to NSAIDs, see: Halen *et al.* (2009); Schoen & Vender (1989); Mitchell & Warner (1999). For hydrogen-bond motifs, see: Etter *et al.* (1990).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{21}\text{N}_3\text{O}_2$
 $M_r = 347.41$
 Triclinic, $P\bar{1}$
 $a = 8.2786$ (9) Å
 $b = 10.1194$ (11) Å
 $c = 11.7739$ (13) Å
 $\alpha = 93.001$ (2)°
 $\beta = 108.993$ (2)°
 $\gamma = 105.578$ (2)°
 $V = 887.76$ (17) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 150$ K
 $0.26 \times 0.13 \times 0.08$ mm

Data collection

Bruker SMART APEX CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2013)
 $T_{\min} = 0.78$, $T_{\max} = 0.99$
 16303 measured reflections
 4574 independent reflections
 3675 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$
 $wR(F^2) = 0.126$
 $S = 1.08$
 4574 reflections
 245 parameters
 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.36$ e Å⁻³
 $\Delta\rho_{\min} = -0.25$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{O2}^{\text{i}}$	0.891 (18)	2.093 (18)	2.9841 (14)	178.1 (9)
$\text{N2}-\text{H2}\cdots\text{O2}^{\text{ii}}$	0.903 (16)	2.012 (16)	2.9055 (13)	170.0 (14)
$\text{C7}-\text{H7}\cdots\text{N3}$	0.95	2.54	3.3609 (15)	145

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

Data collection: APEX2 (Bruker, 2013); cell refinement: SAINT (Bruker, 2013); 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, 2012) and PLATON (Spek, 2009); software used to prepare material for publication: WinGX (Farrugia, 2012) and PLATON.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: QM2100).

References

- Agrawal, N., Chandrasekar, M. J. N., Sara, U. V. S. & Rohini, A. (2010). *Int. J. Drug Deliv. Tech.* **2**, 12–17.
 Allan, H. P. & Fletcher, M. (1990). *Drugs*, **40**, 1–11.
 Bruker (2013). APEX2, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.
 Champion, G. D., Feng, P. H., Azuma, T., Caughey, D. E., Chan, K. H., Kashiwazaki, S., Liu, H.-C., Nasution, A. R., Hobunaga, M., Prichanon, S., Torralba, T. P., Udom, V. & Yoo, M. C. (1997). *Drugs*, **53**, 61–69.
 Etter, M. C., MacDonald, J. C. & Bernstein, J. (1990). *Acta Cryst.* **B46**, 256–262.
 Farrugia, L. J. (2012). *J. Appl. Cryst.* **45**, 849–854.
 Halen, P. K., Murumkar, P. R., Giridhar, R. & Yadav, M. R. (2009). *Mini Rev. Med. Chem.* **9**, 124–139.
 Mitchell, J. A. & Warner, T. D. (1999). *Br. J. Pharmacol.* **128**, 1121–1132.
 Schoen, R. T. & Vender, R. J. (1989). *Am. J. Med.* **86**, 449–458.
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
 Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.

supporting information

Acta Cryst. (2013). E69, o1493 [doi:10.1107/S1600536813023805]

2-(5-Methoxy-2-methyl-1*H*-indol-3-yl)-*N'*-[(1*E*,2*E*)-3-phenylprop-2-en-1-yl-*idene*]acetohydrazide

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S1. Comment

Indomethacin as other common anti-inflammatory drugs (NSAIDs) which are widely employed in the treatment of pain and inflammation has been reported to be associated with a number of undesirable effects, in particular gastrointestinal (GI) toxicity and ulceration (Agrawal *et al.*, 2010; Champion *et al.*, 1997; Allan & Fletcher, 1990) which represent a still unsolved therapeutic problem. Topical irritation by the free carboxylic group of Indomethacin is considered an important factor in establishing superficial stomach erosion (Schoen & Vender, 1989; Mitchell & Warner, 1999). Considerable attention has been focused on the development of bio-reversible derivatives of such pro-drugs to temporarily mask the acidic group as a promising means of reducing or abolishing the GI toxicity due to the local action mechanism (Halen *et al.*, 2009). Based on such facts and continue to our on-going study in functionalization of NSAIDs we herein report the synthesis and crystal structure of the title compound.

The molecular conformation adopted by I in the crystal is "J" shaped (Fig. 1) and appears to be at least partially directed by a weak, intramolecular C7—H7···N3 hydrogen bond. The indole ring system is almost planar [maximum deviations = -0.046 (1) Å for N1, -0.036 (1) Å for C2 and 0.035 (1) Å for C4] and the dihedral angle between it and the terminal phenyl ring is 79.10 (5)°.

In the crystal structure, the N—H···O hydrogen bonding consists of $R^2_2(8)$ rings (Etter *et al.*, 1990) with 2 N2—H2···O2 contacts and $R^2_2(14)$ rings with 2 N1—H1···O2 contacts which form a chain running parallel to the *b* axis (Table 1, Figs. 2 & 3).

S2. Experimental

A mixture of 233 mg (1 mmol) 2-(5-methoxy-2-methyl-1*H*-indole-3-yl)acetohydrazide and 132 mg (1 mmol) of (2*E*)-3-phenylprop-2-enal in 50 ml of ethanol containing a few drops of glacial acetic acid was refluxed for 6 hrs. The mixture was cooled to room temperature and the excess solvent was evaporated under *vacuum*. The resulting solid was collected, washed with ethanol and recrystallized from dioxan to give colourless tablets (*M.p.* 410–413 K) suitable for X-ray analysis.

IR (KBr cm^{-1}): (C=O amide 1661), (NH 3301), (C=N 1606) (C—H, Ar 3021–3072), (C—H aliphatic 2836–2957). $^1\text{H-NMR}$: (DMSO- d_6) δ at 3.6(s, 3H, -OCH₃), 2.3(s, 3H, CH₃), 3.4(s, 2H, -CH₂), 6.5(d, 1*H*, -CH= alkene), 6.6(d, 1*H*, -CH= alkene), 8.2(d, 1*H*, -CH=N), 11.8(s, 1*H*, -NH amide), the aromatic protons of indole nuclei and benzene ring were appeared in the range of 7.0–7.9. $^{13}\text{C-NMR}$: 161(C=O amide), 149(-CH=N), 55(-OCH₃), 10(1 C, CH₃), 136(-C=C alkene), 125(-C=C alkene).

S3. Refinement

C-bound H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with $C-H = 0.95-0.99 \text{ \AA}$, with $U_{iso}(H) = 1.5 U_{iso}(C)$ for methyl H atoms and $U_{iso}(H) = 1.2 U_{iso}(C)$ for other H atoms. H atoms bonded to N atoms were located in difference Fourier maps [$N1-H1 = 0.891 (18) \text{ \AA}$ and $N2-H2 = 0.903 (16) \text{ \AA}$] and refined isotropically.

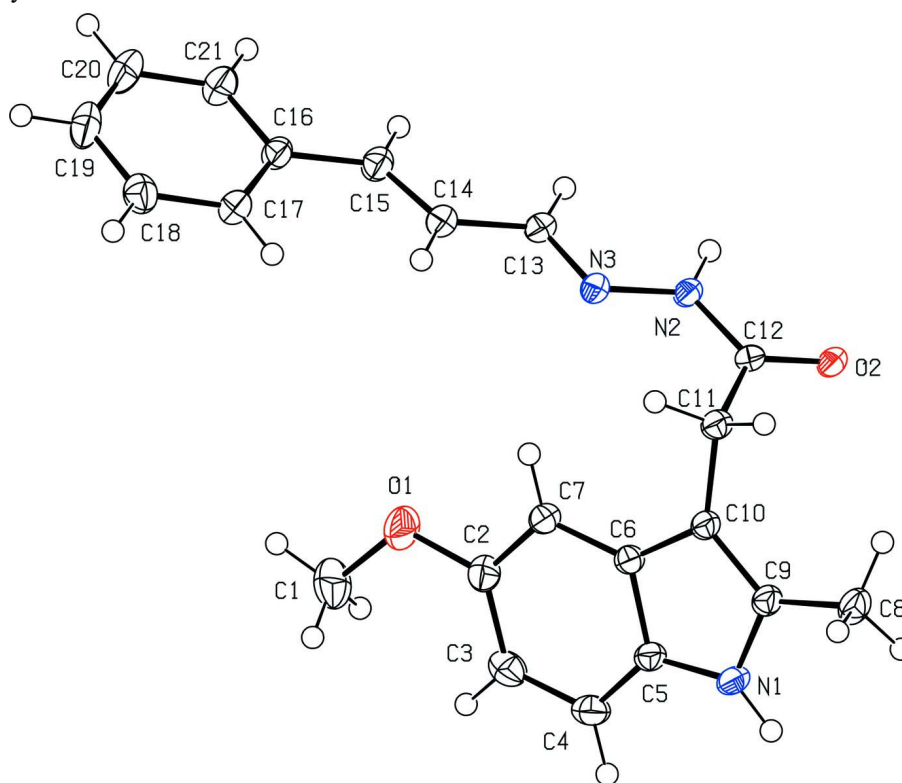


Figure 1

Perspective view of the title molecule with 50% probability displacement ellipsoids.

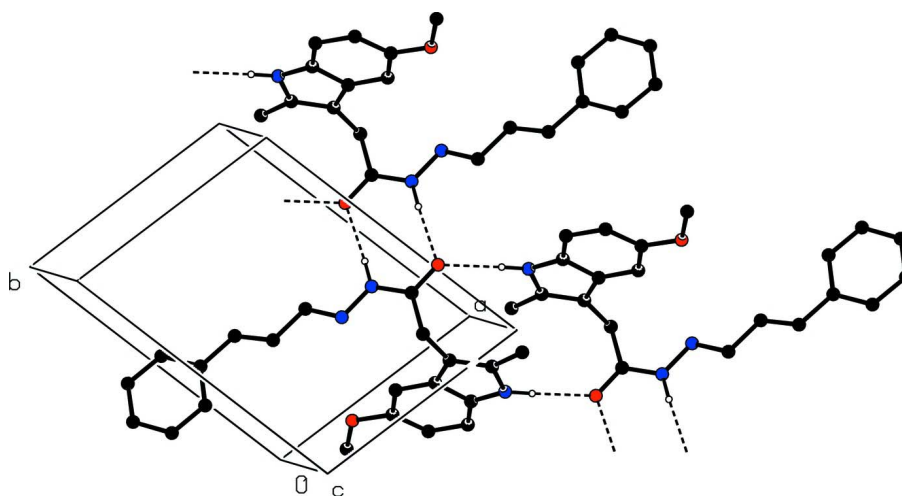
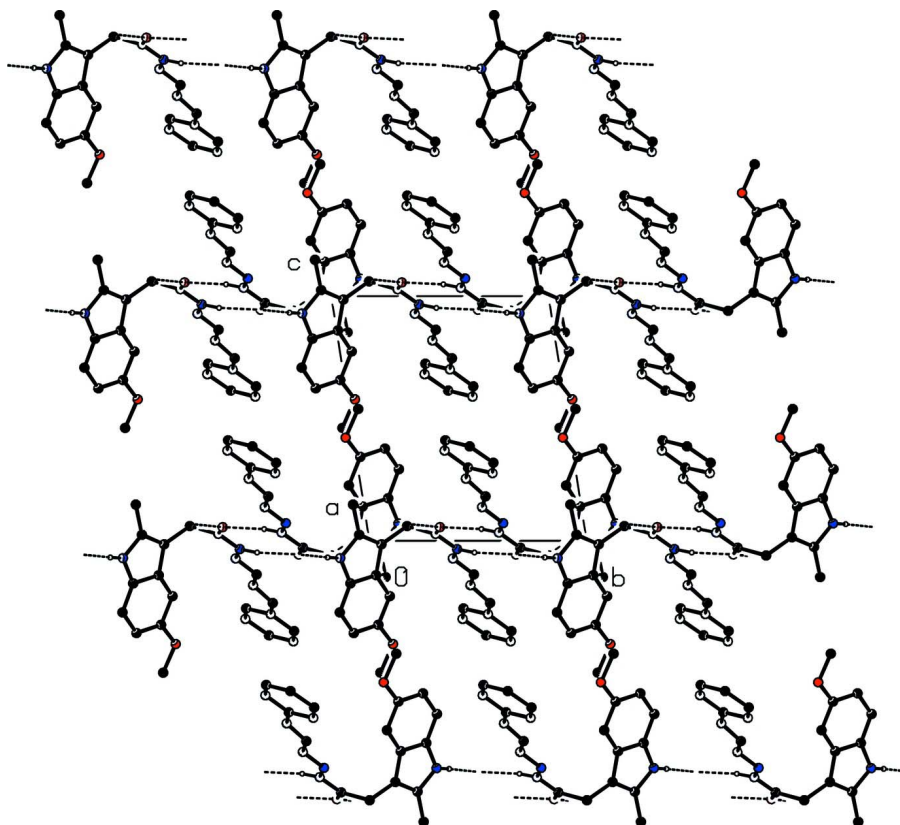


Figure 2

Partial view of the $R_2^2(8)$ and $R_2^2(14)$ cyclic dimers, down the a axis.

**Figure 3**

Packing of the title molecule viewed down a with the hydrogen bonds shown by dotted lines.

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Crystal data

$C_{21}H_{21}N_3O_2$

$M_r = 347.41$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 8.2786$ (9) Å

$b = 10.1194$ (11) Å

$c = 11.7739$ (13) Å

$\alpha = 93.001$ (2)°

$\beta = 108.993$ (2)°

$\gamma = 105.578$ (2)°

$V = 887.76$ (17) Å³

$Z = 2$

$F(000) = 368$

$D_x = 1.300$ Mg m⁻³

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

Cell parameters from 8428 reflections

$\theta = 2.6$ – 29.1 °

$\mu = 0.09$ mm⁻¹

$T = 150$ K

Tablet, clear colourless

$0.26 \times 0.13 \times 0.08$ mm

Data collection

Bruker SMART APEX CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 8.3660 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2013)

$T_{\min} = 0.78$, $T_{\max} = 0.99$

16303 measured reflections

4574 independent reflections
 3675 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$
 $\theta_{\text{max}} = 29.1^\circ$, $\theta_{\text{min}} = 1.9^\circ$

$h = -11 \rightarrow 10$
 $k = -13 \rightarrow 13$
 $l = -16 \rightarrow 15$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.045$
 $wR(F^2) = 0.126$
 $S = 1.08$
 4574 reflections
 245 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 H atoms treated by a mixture of independent
 and constrained refinement
 $W = 1/[\Sigma^2(FO^2) + (0.0677P)^2 + 0.0983P]$
 WHERE $P = (FO^2 + 2FC^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.36 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.25 \text{ e } \text{\AA}^{-3}$

Special details

Experimental. The diffraction data were obtained from 3 sets of 400 frames, each of width 0.5° in ω , collected at $\varphi = 0.00, 90.00$ and 180.00° and 2 sets of 800 frames, each of width 0.45° in φ , collected at $\omega = -30.00$ and 210.00° . The scan time was 15 sec/frame.

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 on F^2 for ALL reflections except those flagged by the user for potential systematic errors. Weighted R -factors wR and all goodnesses 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 observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating $-R$ -factor-obs *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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.28193 (13)	0.03509 (10)	0.57833 (8)	0.0383 (3)
O2	1.01220 (10)	0.33474 (8)	1.05582 (8)	0.0235 (3)
N1	0.76159 (13)	-0.14643 (9)	0.92901 (9)	0.0233 (3)
N2	0.76767 (12)	0.39782 (9)	0.96609 (9)	0.0198 (3)
N3	0.58483 (12)	0.37175 (9)	0.92778 (9)	0.0213 (3)
C1	0.1780 (2)	-0.04316 (16)	0.46126 (12)	0.0406 (4)
C2	0.40353 (16)	-0.01850 (12)	0.65687 (11)	0.0261 (3)
C3	0.44402 (17)	-0.13776 (12)	0.62313 (11)	0.0289 (3)
C4	0.56565 (16)	-0.18727 (11)	0.70752 (11)	0.0264 (3)
C5	0.64546 (14)	-0.11740 (10)	0.82602 (11)	0.0210 (3)
C6	0.61147 (14)	0.00623 (10)	0.85962 (10)	0.0190 (3)
C7	0.48756 (14)	0.05444 (11)	0.77385 (10)	0.0216 (3)
C8	0.92662 (16)	-0.04883 (12)	1.14821 (11)	0.0266 (3)
C9	0.80675 (14)	-0.04222 (11)	1.02468 (11)	0.0212 (3)
C10	0.71796 (14)	0.05319 (10)	0.98586 (10)	0.0190 (3)
C11	0.73269 (15)	0.18427 (10)	1.06001 (10)	0.0203 (3)
C12	0.84755 (14)	0.31009 (10)	1.02798 (10)	0.0184 (3)
C13	0.52187 (15)	0.45707 (11)	0.86269 (10)	0.0217 (3)
C14	0.33089 (15)	0.43345 (11)	0.81838 (11)	0.0235 (3)

C15	0.24928 (15)	0.51098 (12)	0.74611 (11)	0.0245 (3)
C16	0.05663 (15)	0.49356 (11)	0.70035 (10)	0.0223 (3)
C17	-0.06817 (16)	0.38488 (12)	0.72362 (11)	0.0257 (3)
C18	-0.24894 (16)	0.37336 (13)	0.68079 (11)	0.0289 (3)
C19	-0.30851 (17)	0.47052 (15)	0.61408 (12)	0.0342 (4)
C20	-0.18776 (18)	0.57806 (15)	0.58911 (12)	0.0350 (4)
C21	-0.00671 (17)	0.58889 (13)	0.63135 (11)	0.0291 (3)
H1	0.827 (2)	-0.2041 (17)	0.9322 (15)	0.044 (4)*
H1A	0.25730	-0.05190	0.41690	0.0610*
H1B	0.09370	0.00410	0.41640	0.0610*
H1C	0.11120	-0.13570	0.46960	0.0610*
H2	0.838 (2)	0.4754 (16)	0.9519 (13)	0.032 (4)*
H3	0.38760	-0.18480	0.54180	0.0350*
H4	0.59380	-0.26740	0.68470	0.0320*
H7	0.46140	0.13600	0.79550	0.0260*
H8A	0.98240	0.04490	1.19500	0.0400*
H8B	1.02000	-0.08780	1.14140	0.0400*
H8C	0.85640	-0.10790	1.18960	0.0400*
H11A	0.78640	0.17950	1.14750	0.0240*
H11B	0.61160	0.19340	1.04460	0.0240*
H13	0.59940	0.53380	0.84430	0.0260*
H14	0.25880	0.35810	0.84210	0.0280*
H15	0.32320	0.58470	0.72200	0.0290*
H17	-0.02820	0.31800	0.76950	0.0310*
H18	-0.33210	0.29880	0.69720	0.0350*
H19	-0.43240	0.46330	0.58550	0.0410*
H20	-0.22870	0.64450	0.54300	0.0420*
H21	0.07530	0.66240	0.61290	0.0350*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0382 (5)	0.0443 (5)	0.0258 (5)	0.0207 (4)	-0.0034 (4)	0.0015 (4)
O2	0.0174 (4)	0.0182 (4)	0.0346 (5)	0.0081 (3)	0.0066 (3)	0.0048 (3)
N1	0.0215 (5)	0.0171 (4)	0.0333 (5)	0.0093 (4)	0.0093 (4)	0.0057 (4)
N2	0.0159 (4)	0.0161 (4)	0.0280 (5)	0.0064 (3)	0.0072 (4)	0.0047 (4)
N3	0.0161 (4)	0.0197 (4)	0.0277 (5)	0.0069 (3)	0.0064 (4)	0.0016 (4)
C1	0.0359 (8)	0.0526 (8)	0.0244 (7)	0.0121 (6)	0.0006 (6)	0.0028 (6)
C2	0.0228 (6)	0.0271 (5)	0.0261 (6)	0.0082 (4)	0.0052 (5)	0.0048 (5)
C3	0.0302 (6)	0.0265 (6)	0.0257 (6)	0.0048 (5)	0.0083 (5)	-0.0029 (5)
C4	0.0277 (6)	0.0190 (5)	0.0338 (7)	0.0063 (4)	0.0138 (5)	0.0002 (4)
C5	0.0181 (5)	0.0161 (5)	0.0297 (6)	0.0054 (4)	0.0095 (4)	0.0040 (4)
C6	0.0162 (5)	0.0162 (4)	0.0252 (6)	0.0045 (4)	0.0084 (4)	0.0034 (4)
C7	0.0201 (5)	0.0191 (5)	0.0264 (6)	0.0074 (4)	0.0078 (5)	0.0041 (4)
C8	0.0227 (6)	0.0250 (5)	0.0329 (6)	0.0098 (4)	0.0076 (5)	0.0126 (5)
C9	0.0174 (5)	0.0180 (5)	0.0285 (6)	0.0057 (4)	0.0075 (4)	0.0071 (4)
C10	0.0165 (5)	0.0163 (4)	0.0244 (5)	0.0054 (4)	0.0068 (4)	0.0048 (4)
C11	0.0210 (5)	0.0183 (5)	0.0227 (5)	0.0075 (4)	0.0076 (4)	0.0044 (4)

C12	0.0190 (5)	0.0159 (4)	0.0206 (5)	0.0075 (4)	0.0059 (4)	0.0004 (4)
C13	0.0205 (5)	0.0191 (5)	0.0265 (6)	0.0085 (4)	0.0076 (5)	0.0024 (4)
C14	0.0199 (5)	0.0220 (5)	0.0278 (6)	0.0080 (4)	0.0065 (5)	0.0021 (4)
C15	0.0204 (5)	0.0242 (5)	0.0301 (6)	0.0088 (4)	0.0084 (5)	0.0059 (4)
C16	0.0212 (5)	0.0248 (5)	0.0223 (5)	0.0115 (4)	0.0059 (4)	0.0019 (4)
C17	0.0239 (6)	0.0260 (5)	0.0281 (6)	0.0116 (4)	0.0071 (5)	0.0043 (5)
C18	0.0230 (6)	0.0335 (6)	0.0282 (6)	0.0074 (5)	0.0082 (5)	0.0003 (5)
C19	0.0224 (6)	0.0487 (8)	0.0317 (7)	0.0172 (5)	0.0049 (5)	0.0051 (6)
C20	0.0320 (7)	0.0429 (7)	0.0341 (7)	0.0226 (6)	0.0066 (6)	0.0131 (6)
C21	0.0272 (6)	0.0319 (6)	0.0307 (6)	0.0136 (5)	0.0089 (5)	0.0095 (5)

Geometric parameters (Å, °)

O1—C1	1.4195 (17)	C16—C17	1.3991 (17)
O1—C2	1.3793 (16)	C17—C18	1.384 (2)
O2—C12	1.2432 (15)	C18—C19	1.3842 (19)
N1—C5	1.3824 (16)	C19—C20	1.382 (2)
N1—C9	1.3822 (15)	C20—C21	1.388 (2)
N2—N3	1.3758 (15)	C1—H1A	0.9800
N2—C12	1.3512 (15)	C1—H1B	0.9800
N3—C13	1.2866 (15)	C1—H1C	0.9800
N1—H1	0.891 (18)	C3—H3	0.9500
N2—H2	0.903 (16)	C4—H4	0.9500
C2—C7	1.3856 (16)	C7—H7	0.9500
C2—C3	1.4073 (18)	C8—H8A	0.9800
C3—C4	1.3871 (18)	C8—H8B	0.9800
C4—C5	1.3888 (17)	C8—H8C	0.9800
C5—C6	1.4179 (15)	C11—H11A	0.9900
C6—C10	1.4354 (16)	C11—H11B	0.9900
C6—C7	1.3986 (16)	C13—H13	0.9500
C8—C9	1.4869 (17)	C14—H14	0.9500
C9—C10	1.3748 (16)	C15—H15	0.9500
C10—C11	1.5030 (15)	C17—H17	0.9500
C11—C12	1.5169 (15)	C18—H18	0.9500
C13—C14	1.4409 (19)	C19—H19	0.9500
C14—C15	1.3328 (17)	C20—H20	0.9500
C15—C16	1.4645 (19)	C21—H21	0.9500
C16—C21	1.3936 (17)		
C1—O1—C2	117.79 (11)	C16—C21—C20	121.01 (13)
C5—N1—C9	109.03 (9)	O1—C1—H1A	109.00
N3—N2—C12	121.25 (9)	O1—C1—H1B	109.00
N2—N3—C13	116.02 (10)	O1—C1—H1C	109.00
C9—N1—H1	121.1 (11)	H1A—C1—H1B	109.00
C5—N1—H1	127.1 (11)	H1A—C1—H1C	109.00
N3—N2—H2	120.7 (11)	H1B—C1—H1C	109.00
C12—N2—H2	118.1 (11)	C2—C3—H3	120.00
O1—C2—C3	123.67 (11)	C4—C3—H3	120.00

O1—C2—C7	115.35 (11)	C3—C4—H4	121.00
C3—C2—C7	120.98 (12)	C5—C4—H4	121.00
C2—C3—C4	120.58 (11)	C2—C7—H7	121.00
C3—C4—C5	118.66 (11)	C6—C7—H7	120.00
C4—C5—C6	121.15 (11)	C9—C8—H8A	109.00
N1—C5—C6	107.51 (10)	C9—C8—H8B	109.00
N1—C5—C4	131.34 (10)	C9—C8—H8C	109.00
C5—C6—C10	106.84 (10)	H8A—C8—H8B	109.00
C7—C6—C10	133.62 (10)	H8A—C8—H8C	109.00
C5—C6—C7	119.50 (10)	H8B—C8—H8C	109.00
C2—C7—C6	119.01 (11)	C10—C11—H11A	109.00
N1—C9—C8	120.64 (10)	C10—C11—H11B	109.00
N1—C9—C10	109.46 (10)	C12—C11—H11A	109.00
C8—C9—C10	129.81 (11)	C12—C11—H11B	109.00
C6—C10—C11	126.11 (10)	H11A—C11—H11B	108.00
C9—C10—C11	126.77 (10)	N3—C13—H13	121.00
C6—C10—C9	107.10 (10)	C14—C13—H13	121.00
C10—C11—C12	110.76 (10)	C13—C14—H14	118.00
N2—C12—C11	118.76 (11)	C15—C14—H14	118.00
O2—C12—N2	118.88 (10)	C14—C15—H15	117.00
O2—C12—C11	122.36 (10)	C16—C15—H15	117.00
N3—C13—C14	118.40 (11)	C16—C17—H17	119.00
C13—C14—C15	123.95 (11)	C18—C17—H17	120.00
C14—C15—C16	126.31 (11)	C17—C18—H18	120.00
C15—C16—C17	122.69 (11)	C19—C18—H18	120.00
C15—C16—C21	119.29 (11)	C18—C19—H19	120.00
C17—C16—C21	118.02 (12)	C20—C19—H19	120.00
C16—C17—C18	121.03 (11)	C19—C20—H20	120.00
C17—C18—C19	119.96 (12)	C21—C20—H20	120.00
C18—C19—C20	119.99 (14)	C16—C21—H21	119.00
C19—C20—C21	119.99 (13)	C20—C21—H21	120.00
C1—O1—C2—C3	7.23 (19)	C5—C6—C10—C11	176.91 (11)
C1—O1—C2—C7	-173.47 (12)	C7—C6—C10—C9	176.05 (13)
C9—N1—C5—C4	178.38 (13)	C7—C6—C10—C11	-5.5 (2)
C9—N1—C5—C6	-2.32 (13)	N1—C9—C10—C6	0.16 (14)
C5—N1—C9—C8	178.24 (11)	N1—C9—C10—C11	-178.30 (11)
C5—N1—C9—C10	1.36 (14)	C8—C9—C10—C6	-176.35 (12)
C12—N2—N3—C13	-175.51 (10)	C8—C9—C10—C11	5.2 (2)
N3—N2—C12—O2	176.74 (10)	C6—C10—C11—C12	-74.62 (15)
N3—N2—C12—C11	-2.72 (16)	C9—C10—C11—C12	103.55 (13)
N2—N3—C13—C14	179.00 (10)	C10—C11—C12—O2	-70.23 (14)
O1—C2—C3—C4	-178.89 (12)	C10—C11—C12—N2	109.21 (12)
C7—C2—C3—C4	1.8 (2)	N3—C13—C14—C15	-177.08 (12)
O1—C2—C7—C6	179.33 (11)	C13—C14—C15—C16	-178.72 (11)
C3—C2—C7—C6	-1.35 (19)	C14—C15—C16—C17	-3.5 (2)
C2—C3—C4—C5	0.5 (2)	C14—C15—C16—C21	176.09 (12)
C3—C4—C5—N1	175.97 (13)	C15—C16—C17—C18	178.73 (11)

C3—C4—C5—C6	-3.25 (19)	C21—C16—C17—C18	-0.85 (18)
N1—C5—C6—C7	-175.64 (11)	C15—C16—C21—C20	-178.26 (12)
N1—C5—C6—C10	2.37 (13)	C17—C16—C21—C20	1.33 (18)
C4—C5—C6—C7	3.75 (18)	C16—C17—C18—C19	-0.17 (19)
C4—C5—C6—C10	-178.25 (11)	C17—C18—C19—C20	0.7 (2)
C5—C6—C7—C2	-1.38 (17)	C18—C19—C20—C21	-0.3 (2)
C10—C6—C7—C2	-178.75 (13)	C19—C20—C21—C16	-0.8 (2)
C5—C6—C10—C9	-1.55 (13)		

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—H1...O2 ⁱ	0.891 (18)	2.093 (18)	2.9841 (14)	178.1 (9)
N2—H2...O2 ⁱⁱ	0.903 (16)	2.012 (16)	2.9055 (13)	170.0 (14)
C7—H7...N3	0.95	2.54	3.3609 (15)	145
C11—H11B...N3	0.99	2.34	2.7996 (15)	107

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