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**(Z)-N-tert-Butyl-2-(4-methoxyanilino)-
N'-(4-methoxyphenyl)-2-phenyl-
acetimidamide**

Sue A. Roberts,^{a*} Biswajit Saha,^{b,c} Brendan Frett^{b,c} and
Hong-Yu Li^{b,c}

^aDepartment of Chemistry and Biochemistry, 1306 E University Boulevard, The University of Arizona, Tucson, AZ 85721, USA, ^bCollege of Pharmacy, Department of Pharmacology and Toxicology, The University of Arizona, Tucson, AZ 85721, USA, and ^cBio5 Oro Valley, The University of Arizona, Oro Valley, AZ 85737, USA
Correspondence e-mail: suer@email.arizona.edu

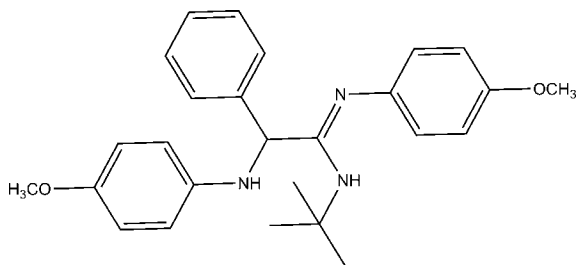
Received 10 April 2013; accepted 9 May 2013

Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.043; wR factor = 0.106; data-to-parameter ratio = 22.9.

In the crystal of the title compound, $\text{C}_{26}\text{H}_{31}\text{N}_3\text{O}_2$, pairs of $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds link molecules, forming inversion dimers, which enclose an $R_2^2(20)$ ring motif. One N atom does not form hydrogen bonds and lies in a hydrophobic pocket with closest intermolecular contacts of 4.196 (2) and 4.262 (2) Å.

Related literature

For the synthesis of the title compound, and a discussion of the use of the three-component Ugi reaction to synthesize amidines, see: Saha *et al.* (2013).



Experimental

Crystal data

$\text{C}_{26}\text{H}_{31}\text{N}_3\text{O}_2$
 $M_r = 417.54$

Triclinic, $P\bar{1}$
 $a = 10.0804$ (17) Å

$b = 10.5784$ (17) Å
 $c = 11.1573$ (18) Å
 $\alpha = 80.982$ (4)°
 $\beta = 85.152$ (4)°
 $\gamma = 80.270$ (4)°
 $V = 1156.1$ (3) Å³

$Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 100$ K
 $0.20 \times 0.20 \times 0.10$ mm

Data collection

Bruker Kappa APEXII DUO CCD diffractometer
Absorption correction: multi-scan (SADABS; Bruker, 2009)
 $T_{\min} = 0.89$, $T_{\max} = 0.99$

32612 measured reflections
6607 independent reflections
5476 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.019$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.106$
 $S = 0.94$
6607 reflections
289 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.36$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.38$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2}\cdots\text{O1}^i$	0.887 (15)	2.328 (15)	3.1698 (13)	158.5 (12)

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

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) and OLEX2 (Dolomanov *et al.*, 2009); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012) and Mercury (Macrae *et al.*, 2006); software used to prepare material for publication: pubCIF (Westrip, 2010).

The Bruker Kappa APEXII DUO was purchased with funding from NSF grant CHE-0741837. The work was supported by start-up funds from the University of Arizona to HYL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PK2477).

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supporting information

Acta Cryst. (2013). E69, o902 [doi:10.1107/S1600536813012877]

(Z)-N-tert-Butyl-2-(4-methoxyanilino)-N'-(4-methoxyphenyl)-2-phenyl-acetimidamide

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S1. Comment

The title compound was prepared as a secondary product of the three component Ugi reaction (Saha *et al.*, 2013). The central C1—C22 single bond has four bulky substituents arranged, roughly, as if pointing at the corners of a tetrahedron. The angle between the planes containing the substituents, defined by N1—C22—N3 and C2—C1—C20, is 72.41 (1)°. Pairs of hydrogen bonds between N2 and O1 [3.170 (1) Å] connect two molecules across an inversion center, creating a cavity around the inversion center with a diameter of about 5 Å. The closest contact across the inversion center involving atoms not involved in the hydrogen bonding is 3.151 (3) Å between the hydrogen bonded to C3 and its symmetry equivalent (symmetry transformation $-x + 1, -y + 2, -z + 2$). The hydrogen bonding graph set is $R^2_2(20)$. Nitrogen N3 is unusual in that, although protonated, it does not form hydrogen bonds with acceptor atoms. Instead, it lies in a hydrophobic cavity with closest intermolecular contacts of 4.196 (2) Å and 4.262 (2) Å to C15 and C16 respectively.

S2. Experimental

The compound was synthesized as previously reported [compound 5a in (Saha *et al.*, 2013)]. The crude residue was purified by silica gel column chromatography using (10–30%) ethylacetate-hexane to obtain the pure product. The pure compound was dissolved in 50% ethylacetate-hexane and kept at room temperature for 2 days during which crystals formed.

S3. Refinement

All hydrogen atoms were visible in a difference Fourier map and, except for H2 were added at calculated positions. Bond distances are set to 0.95 Å for aromatic carbon-hydrogen bonds, 0.98 Å for methyl group carbon-hydrogen bonds and 0.88 Å for nitrogen-hydrogen bonds. Thermal parameters for hydrogen atoms were set to 1.2 times the isotropic equivalent thermal parameter of the atom to which the hydrogen atom is bonded, except for methyl group hydrogen atoms where the thermal parameter was set to 1.5 times the isotropic equivalent thermal parameter of the carbon atom the hydrogen atom is bonded to. The positional and isotropic thermal parameters of H2 were refined.

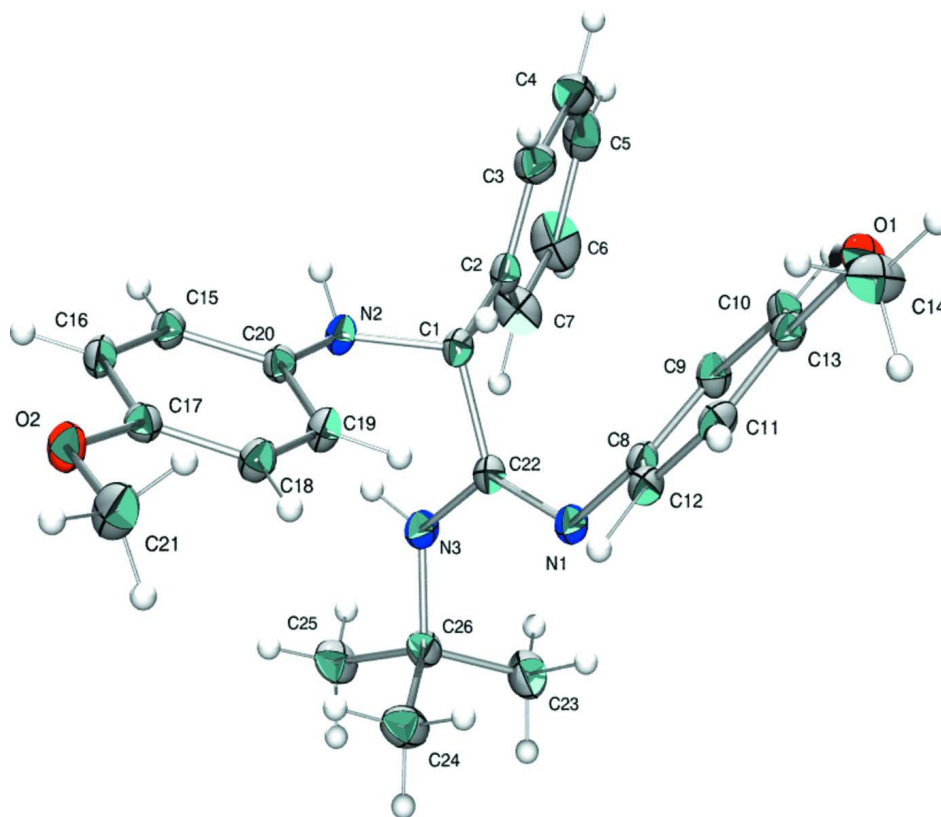
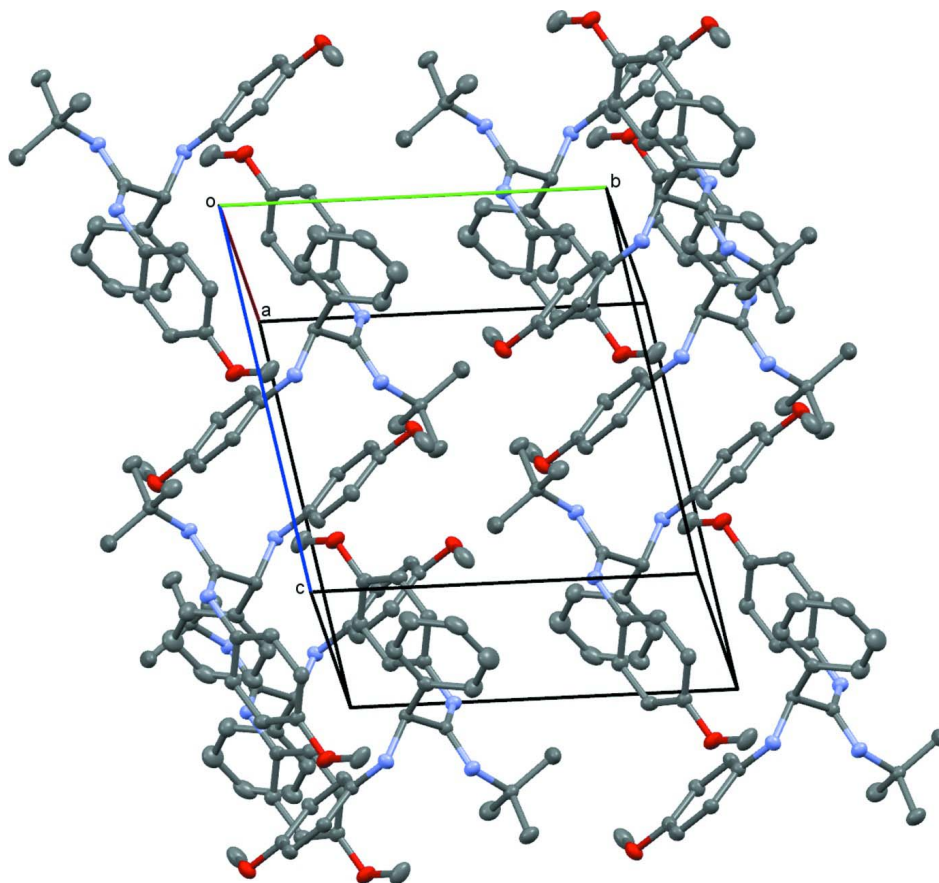


Figure 1

The molecular structure of the title compound. Non-hydrogen atoms are shown as 50% probability ellipsoids.

**Figure 2**

View showing unit cell packing. Unit cell axes are labeled in the figure.

(Z)-N-tert-Butyl-2-(4-methoxyanilino)-N'-(4-methoxyphenyl)-2-phenylacetimidamide

Crystal data

$C_{26}H_{31}N_3O_2$

$M_r = 417.54$

Triclinic, $P\bar{1}$

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$c = 11.1573$ (18) Å

$\alpha = 80.982$ (4)°

$\beta = 85.152$ (4)°

$\gamma = 80.270$ (4)°

$V = 1156.1$ (3) Å³

$Z = 2$

$F(000) = 448$

$D_x = 1.199$ Mg m⁻³

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

Cell parameters from 9952 reflections

$\theta = 2.9$ – 29.8 °

$\mu = 0.08$ mm⁻¹

$T = 100$ K

Block, colourless

$0.20 \times 0.20 \times 0.10$ mm

Data collection

Bruker Kappa APEXII DUO CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 8.3333 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2009)

$T_{\min} = 0.89$, $T_{\max} = 0.99$

32612 measured reflections

6607 independent reflections

5476 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.019$

$\theta_{\max} = 29.9^\circ$, $\theta_{\min} = 1.9^\circ$
 $h = -14 \rightarrow 11$

$k = -14 \rightarrow 12$
 $l = -15 \rightarrow 13$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.106$
 $S = 0.94$
 6607 reflections
 289 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 atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0397P)^2 + 0.6486P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.36 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.38 \text{ e } \text{\AA}^{-3}$

Special details

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

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
C20	0.37947 (10)	1.03405 (9)	0.68022 (9)	0.01884 (19)
C11	0.15660 (10)	0.92039 (10)	1.16518 (9)	0.0213 (2)
H11	0.0822	0.9836	1.1842	0.026*
C18	0.15655 (11)	1.15884 (10)	0.70473 (10)	0.0226 (2)
H18	0.0708	1.1695	0.7475	0.027*
C15	0.40664 (11)	1.12635 (10)	0.57996 (9)	0.02047 (19)
H15	0.4919	1.1155	0.5363	0.025*
C16	0.31125 (11)	1.23257 (10)	0.54406 (9)	0.0219 (2)
H16	0.3313	1.2940	0.4761	0.026*
C22	0.35777 (10)	0.75795 (9)	0.82814 (9)	0.01868 (19)
C10	0.36795 (11)	0.78673 (11)	1.21611 (9)	0.0228 (2)
H10	0.4377	0.7582	1.2709	0.027*
C12	0.16277 (10)	0.86653 (10)	1.05826 (9)	0.0212 (2)
H12	0.0906	0.8921	1.0057	0.025*
C19	0.25281 (11)	1.05135 (10)	0.74063 (10)	0.0224 (2)
H19	0.2315	0.9890	0.8074	0.027*
C8	0.27219 (10)	0.77613 (10)	1.02648 (9)	0.01957 (19)
C1	0.46163 (10)	0.84821 (9)	0.83240 (9)	0.01839 (18)
H1	0.4260	0.9082	0.8927	0.022*
C17	0.18548 (10)	1.25005 (10)	0.60709 (9)	0.0206 (2)
C9	0.37371 (11)	0.73460 (10)	1.10887 (9)	0.0223 (2)
H9	0.4473	0.6700	1.0910	0.027*

C2	0.59808 (10)	0.77504 (10)	0.87123 (9)	0.0209 (2)
C26	0.27437 (11)	0.63246 (10)	0.68176 (9)	0.0220 (2)
C3	0.67295 (12)	0.83256 (11)	0.93977 (11)	0.0280 (2)
H3	0.6367	0.9144	0.9636	0.034*
C24	0.12887 (12)	0.70234 (13)	0.67994 (12)	0.0318 (3)
H24A	0.0962	0.7183	0.7623	0.048*
H24B	0.0722	0.6484	0.6505	0.048*
H24C	0.1249	0.7852	0.6258	0.048*
C7	0.65156 (13)	0.65363 (12)	0.83952 (12)	0.0329 (3)
H7	0.6010	0.6121	0.7938	0.039*
C25	0.32629 (13)	0.61040 (12)	0.55297 (10)	0.0299 (2)
H25A	0.3167	0.6936	0.4991	0.045*
H25B	0.2739	0.5524	0.5239	0.045*
H25C	0.4215	0.5710	0.5532	0.045*
C4	0.80027 (13)	0.77209 (13)	0.97403 (12)	0.0336 (3)
H4	0.8505	0.8127	1.0209	0.040*
C21	-0.02644 (12)	1.38154 (12)	0.63778 (12)	0.0333 (3)
H21A	-0.0784	1.3119	0.6344	0.050*
H21B	-0.0785	1.4649	0.6052	0.050*
H21C	-0.0078	1.3829	0.7223	0.050*
C14	0.15403 (12)	1.02124 (13)	1.38420 (12)	0.0335 (3)
H14A	0.1460	1.0989	1.3229	0.050*
H14B	0.1674	1.0449	1.4633	0.050*
H14C	0.0715	0.9827	1.3897	0.050*
C5	0.85386 (12)	0.65322 (13)	0.94015 (12)	0.0350 (3)
H5	0.9419	0.6128	0.9616	0.042*
C23	0.28606 (16)	0.50314 (12)	0.76560 (12)	0.0366 (3)
H23A	0.3800	0.4597	0.7629	0.055*
H23B	0.2285	0.4483	0.7388	0.055*
H23C	0.2573	0.5183	0.8490	0.055*
C6	0.77862 (14)	0.59305 (14)	0.87460 (13)	0.0405 (3)
H6	0.8141	0.5098	0.8535	0.049*
O2	0.09729 (8)	1.35968 (7)	0.56724 (7)	0.02582 (17)
O1	0.26631 (8)	0.92977 (8)	1.35034 (7)	0.02582 (17)
N2	0.47953 (9)	0.92663 (8)	0.71356 (8)	0.01953 (17)
N3	0.36308 (9)	0.71424 (9)	0.71966 (8)	0.02124 (18)
H3A	0.4268	0.7374	0.6658	0.025*
N1	0.27336 (9)	0.72595 (9)	0.91628 (8)	0.02123 (18)
C13	0.26038 (10)	0.88065 (10)	1.24380 (9)	0.0204 (2)
H2	0.5625 (15)	0.9449 (14)	0.6977 (13)	0.027 (3)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C20	0.0210 (4)	0.0187 (4)	0.0171 (4)	-0.0043 (4)	-0.0002 (3)	-0.0025 (3)
C11	0.0189 (4)	0.0221 (5)	0.0226 (5)	-0.0038 (4)	0.0033 (4)	-0.0041 (4)
C18	0.0203 (5)	0.0226 (5)	0.0236 (5)	-0.0037 (4)	0.0019 (4)	-0.0008 (4)
C15	0.0230 (5)	0.0221 (5)	0.0164 (4)	-0.0057 (4)	0.0020 (4)	-0.0025 (3)

C16	0.0275 (5)	0.0212 (5)	0.0168 (4)	-0.0066 (4)	0.0000 (4)	0.0002 (3)
C22	0.0206 (4)	0.0177 (4)	0.0170 (4)	-0.0019 (3)	-0.0004 (3)	-0.0022 (3)
C10	0.0227 (5)	0.0274 (5)	0.0177 (4)	-0.0019 (4)	-0.0007 (4)	-0.0033 (4)
C12	0.0198 (4)	0.0246 (5)	0.0191 (4)	-0.0052 (4)	0.0003 (4)	-0.0017 (4)
C19	0.0229 (5)	0.0210 (5)	0.0212 (5)	-0.0034 (4)	0.0025 (4)	0.0016 (4)
C8	0.0219 (5)	0.0212 (5)	0.0160 (4)	-0.0072 (4)	0.0026 (4)	-0.0018 (3)
C1	0.0202 (4)	0.0185 (4)	0.0160 (4)	-0.0032 (3)	0.0014 (3)	-0.0021 (3)
C17	0.0228 (5)	0.0189 (5)	0.0205 (4)	-0.0036 (4)	-0.0036 (4)	-0.0022 (4)
C9	0.0231 (5)	0.0236 (5)	0.0186 (4)	-0.0005 (4)	0.0014 (4)	-0.0035 (4)
C2	0.0204 (4)	0.0227 (5)	0.0176 (4)	-0.0023 (4)	0.0017 (4)	0.0003 (4)
C26	0.0253 (5)	0.0227 (5)	0.0190 (4)	-0.0051 (4)	-0.0019 (4)	-0.0050 (4)
C3	0.0265 (5)	0.0246 (5)	0.0331 (6)	-0.0050 (4)	-0.0047 (4)	-0.0024 (4)
C24	0.0237 (5)	0.0407 (7)	0.0322 (6)	-0.0036 (5)	-0.0011 (4)	-0.0112 (5)
C7	0.0319 (6)	0.0317 (6)	0.0344 (6)	0.0050 (5)	-0.0052 (5)	-0.0120 (5)
C25	0.0323 (6)	0.0362 (6)	0.0235 (5)	-0.0045 (5)	-0.0005 (4)	-0.0132 (4)
C4	0.0283 (6)	0.0347 (6)	0.0371 (6)	-0.0080 (5)	-0.0088 (5)	0.0036 (5)
C21	0.0233 (5)	0.0299 (6)	0.0406 (7)	0.0017 (4)	-0.0003 (5)	0.0062 (5)
C14	0.0255 (5)	0.0433 (7)	0.0359 (6)	-0.0052 (5)	0.0066 (5)	-0.0234 (5)
C5	0.0249 (5)	0.0396 (7)	0.0336 (6)	0.0028 (5)	-0.0021 (5)	0.0080 (5)
C23	0.0594 (8)	0.0243 (6)	0.0291 (6)	-0.0136 (6)	-0.0126 (6)	-0.0005 (4)
C6	0.0363 (7)	0.0366 (7)	0.0433 (7)	0.0128 (5)	-0.0028 (6)	-0.0091 (6)
O2	0.0252 (4)	0.0223 (4)	0.0267 (4)	-0.0005 (3)	-0.0026 (3)	0.0034 (3)
O1	0.0240 (4)	0.0337 (4)	0.0215 (4)	-0.0035 (3)	0.0026 (3)	-0.0124 (3)
N2	0.0194 (4)	0.0206 (4)	0.0172 (4)	-0.0035 (3)	0.0028 (3)	-0.0002 (3)
N3	0.0237 (4)	0.0240 (4)	0.0171 (4)	-0.0065 (3)	0.0030 (3)	-0.0056 (3)
N1	0.0246 (4)	0.0233 (4)	0.0164 (4)	-0.0062 (3)	0.0010 (3)	-0.0033 (3)
C13	0.0213 (5)	0.0235 (5)	0.0173 (4)	-0.0069 (4)	0.0035 (4)	-0.0047 (4)

Geometric parameters (Å, °)

C20—C15	1.4057 (14)	C26—C25	1.5264 (15)
C20—C19	1.3901 (14)	C26—C23	1.5243 (16)
C20—N2	1.4108 (13)	C26—N3	1.4721 (13)
C11—H11	0.9500	C3—H3	0.9500
C11—C12	1.3930 (14)	C3—C4	1.3897 (17)
C11—C13	1.3897 (15)	C24—H24A	0.9800
C18—H18	0.9500	C24—H24B	0.9800
C18—C19	1.3946 (15)	C24—H24C	0.9800
C18—C17	1.3836 (14)	C7—H7	0.9500
C15—H15	0.9500	C7—C6	1.3903 (18)
C15—C16	1.3807 (15)	C25—H25A	0.9800
C16—H16	0.9500	C25—H25B	0.9800
C16—C17	1.3953 (15)	C25—H25C	0.9800
C22—C1	1.5397 (14)	C4—H4	0.9500
C22—N3	1.3556 (12)	C4—C5	1.3787 (19)
C22—N1	1.2873 (13)	C21—H21A	0.9800
C10—H10	0.9500	C21—H21B	0.9800
C10—C9	1.3869 (14)	C21—H21C	0.9800

C10—C13	1.3906 (15)	C21—O2	1.4220 (15)
C12—H12	0.9500	C14—H14A	0.9800
C12—C8	1.3930 (15)	C14—H14B	0.9800
C19—H19	0.9500	C14—H14C	0.9800
C8—C9	1.4021 (14)	C14—O1	1.4256 (14)
C8—N1	1.4120 (13)	C5—H5	0.9500
C1—H1	1.0000	C5—C6	1.385 (2)
C1—C2	1.5213 (14)	C23—H23A	0.9800
C1—N2	1.4643 (13)	C23—H23B	0.9800
C17—O2	1.3747 (12)	C23—H23C	0.9800
C9—H9	0.9500	C6—H6	0.9500
C2—C3	1.3868 (15)	O1—C13	1.3794 (12)
C2—C7	1.3934 (16)	N2—H2	0.887 (15)
C26—C24	1.5270 (16)	N3—H3A	0.8800
C15—C20—N2	119.16 (9)	C26—C24—H24A	109.5
C19—C20—C15	118.08 (9)	C26—C24—H24B	109.5
C19—C20—N2	122.74 (9)	C26—C24—H24C	109.5
C12—C11—H11	120.3	H24A—C24—H24B	109.5
C13—C11—H11	120.3	H24A—C24—H24C	109.5
C13—C11—C12	119.31 (9)	H24B—C24—H24C	109.5
C19—C18—H18	119.9	C2—C7—H7	119.9
C17—C18—H18	119.9	C6—C7—C2	120.22 (12)
C17—C18—C19	120.28 (10)	C6—C7—H7	119.9
C20—C15—H15	119.5	C26—C25—H25A	109.5
C16—C15—C20	120.96 (10)	C26—C25—H25B	109.5
C16—C15—H15	119.5	C26—C25—H25C	109.5
C15—C16—H16	119.8	H25A—C25—H25B	109.5
C15—C16—C17	120.31 (9)	H25A—C25—H25C	109.5
C17—C16—H16	119.8	H25B—C25—H25C	109.5
N3—C22—C1	112.76 (8)	C3—C4—H4	119.9
N1—C22—C1	124.98 (9)	C5—C4—C3	120.11 (12)
N1—C22—N3	122.26 (9)	C5—C4—H4	119.9
C9—C10—H10	119.9	H21A—C21—H21B	109.5
C9—C10—C13	120.25 (10)	H21A—C21—H21C	109.5
C13—C10—H10	119.9	H21B—C21—H21C	109.5
C11—C12—H12	119.2	O2—C21—H21A	109.5
C11—C12—C8	121.63 (10)	O2—C21—H21B	109.5
C8—C12—H12	119.2	O2—C21—H21C	109.5
C20—C19—C18	120.99 (9)	H14A—C14—H14B	109.5
C20—C19—H19	119.5	H14A—C14—H14C	109.5
C18—C19—H19	119.5	H14B—C14—H14C	109.5
C12—C8—C9	118.01 (9)	O1—C14—H14A	109.5
C12—C8—N1	119.03 (9)	O1—C14—H14B	109.5
C9—C8—N1	122.89 (9)	O1—C14—H14C	109.5
C22—C1—H1	108.1	C4—C5—H5	120.2
C2—C1—C22	112.85 (8)	C4—C5—C6	119.59 (11)
C2—C1—H1	108.1	C6—C5—H5	120.2

N2—C1—C22	110.37 (8)	C26—C23—H23A	109.5
N2—C1—H1	108.1	C26—C23—H23B	109.5
N2—C1—C2	109.12 (8)	C26—C23—H23C	109.5
C18—C17—C16	119.35 (9)	H23A—C23—H23B	109.5
O2—C17—C18	123.94 (10)	H23A—C23—H23C	109.5
O2—C17—C16	116.71 (9)	H23B—C23—H23C	109.5
C10—C9—C8	120.74 (10)	C7—C6—H6	119.8
C10—C9—H9	119.6	C5—C6—C7	120.43 (12)
C8—C9—H9	119.6	C5—C6—H6	119.8
C3—C2—C1	118.49 (9)	C17—O2—C21	116.18 (8)
C3—C2—C7	118.73 (10)	C13—O1—C14	117.04 (9)
C7—C2—C1	122.78 (10)	C20—N2—C1	118.01 (8)
C25—C26—C24	109.45 (9)	C20—N2—H2	112.8 (9)
C23—C26—C24	111.49 (10)	C1—N2—H2	113.0 (9)
C23—C26—C25	109.80 (10)	C22—N3—C26	127.34 (9)
N3—C26—C24	110.61 (9)	C22—N3—H3A	116.3
N3—C26—C25	105.50 (9)	C26—N3—H3A	116.3
N3—C26—C23	109.82 (9)	C22—N1—C8	119.35 (9)
C2—C3—H3	119.6	C11—C13—C10	119.96 (9)
C2—C3—C4	120.89 (11)	O1—C13—C11	124.18 (9)
C4—C3—H3	119.6	O1—C13—C10	115.85 (9)
C20—C15—C16—C17	-0.05 (16)	C9—C8—N1—C22	-72.76 (14)
C11—C12—C8—C9	3.56 (15)	C2—C1—N2—C20	-157.97 (9)
C11—C12—C8—N1	-179.35 (9)	C2—C3—C4—C5	-0.11 (19)
C18—C17—O2—C21	4.27 (15)	C2—C7—C6—C5	-0.7 (2)
C15—C20—C19—C18	-1.56 (16)	C3—C2—C7—C6	-1.07 (19)
C15—C20—N2—C1	166.59 (9)	C3—C4—C5—C6	-1.7 (2)
C15—C16—C17—C18	-1.01 (16)	C24—C26—N3—C22	62.76 (14)
C15—C16—C17—O2	178.62 (9)	C7—C2—C3—C4	1.49 (17)
C16—C17—O2—C21	-175.34 (10)	C25—C26—N3—C22	-178.98 (10)
C22—C1—C2—C3	-145.84 (10)	C4—C5—C6—C7	2.1 (2)
C22—C1—C2—C7	35.00 (14)	C14—O1—C13—C11	3.97 (15)
C22—C1—N2—C20	77.49 (11)	C14—O1—C13—C10	-176.82 (10)
C12—C11—C13—C10	-0.89 (15)	C23—C26—N3—C22	-60.72 (14)
C12—C11—C13—O1	178.29 (9)	N2—C20—C15—C16	179.73 (9)
C12—C8—C9—C10	-2.92 (15)	N2—C20—C19—C18	-179.91 (10)
C12—C8—N1—C22	110.29 (12)	N2—C1—C2—C3	91.07 (11)
C19—C20—C15—C16	1.32 (15)	N2—C1—C2—C7	-88.08 (12)
C19—C20—N2—C1	-15.07 (14)	N3—C22—C1—C2	-87.41 (10)
C19—C18—C17—C16	0.77 (16)	N3—C22—C1—N2	34.97 (11)
C19—C18—C17—O2	-178.83 (10)	N3—C22—N1—C8	-179.30 (9)
C1—C22—N3—C26	-176.49 (9)	N1—C22—C1—C2	92.83 (12)
C1—C22—N1—C8	0.44 (15)	N1—C22—C1—N2	-144.79 (10)
C1—C2—C3—C4	-177.70 (10)	N1—C22—N3—C26	3.28 (16)
C1—C2—C7—C6	178.08 (11)	N1—C8—C9—C10	-179.89 (10)
C17—C18—C19—C20	0.53 (16)	C13—C11—C12—C8	-1.69 (15)
C9—C10—C13—C11	1.50 (16)	C13—C10—C9—C8	0.45 (16)

C9—C10—C13—O1 -177.75 (9)

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>
N2—H2...O1 ⁱ	0.887 (15)	2.328 (15)	3.1698 (13)	158.5 (12)

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