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6-Chloro-3-nitro-*N*-(propan-2-yl)pyridin-2-amineXiao-Yu Qing,^a Yun-Chuang Huang,^a Ling-Ling Yang^b and Yong-Mei Xie^{a*}

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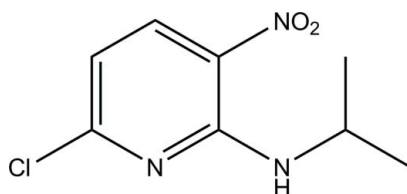
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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.061; wR factor = 0.149; data-to-parameter ratio = 15.8.

There are two molecules in the asymmetric unit molecule of the title compound, $\text{C}_8\text{H}_{10}\text{ClN}_3\text{O}_2$. Intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds stabilize the molecular structure. There are no classical intermolecular hydrogen bonds in the crystal structure.

Related literature

For the biological activity of 6-chloro-*N*-isopropyl-3-nitropyridin-2-amine derivatives, see: Lan *et al.* (2010); Bavetsias *et al.* (2010). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

 $\text{C}_8\text{H}_{10}\text{ClN}_3\text{O}_2$ $M_r = 215.64$ Triclinic, $P\bar{1}$ $a = 7.4283$ (8) Å $b = 8.9573$ (10) Å $c = 15.4301$ (17) Å $\alpha = 89.672$ (9)° $\beta = 86.252$ (9)° $\gamma = 78.860$ (9)° $V = 1005.16$ (19) Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.36$ mm⁻¹ $T = 295$ K $0.28 \times 0.23 \times 0.18$ mm

Data collection

Oxford Diffraction Xcalibur Eos diffractometer

Absorption correction: multi-scan

(CrysAlis PRO; Oxford

Diffraction, 2006)

 $T_{\min} = 0.971$, $T_{\max} = 1.0$

8239 measured reflections

4073 independent reflections

2773 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.022$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.061$ $wR(F^2) = 0.149$ $S = 1.07$

4073 reflections

257 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.24$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2}\cdots\text{O2}$	0.86	2.02	2.660 (3)	130
$\text{N5}-\text{H5}\cdots\text{O3}$	0.86	2.01	2.653 (3)	130

Data collection: CrysAlis PRO (Oxford Diffraction, 2006); cell refinement: CrysAlis PRO; data reduction: CrysAlis PRO; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: OLEX2 (Dolomanov *et al.*, 2009); software used to prepare material for publication: OLEX2.

We thank the Analytical and Testing Center of Sichuan University for the X-ray measurements.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RK2268).

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supporting information

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6-Chloro-3-nitro-*N*-(propan-2-yl)pyridin-2-amine

Xiao-Yu Qing, Yun-Chuang Huang, Ling-Ling Yang and Yong-Mei Xie

S1. Comment

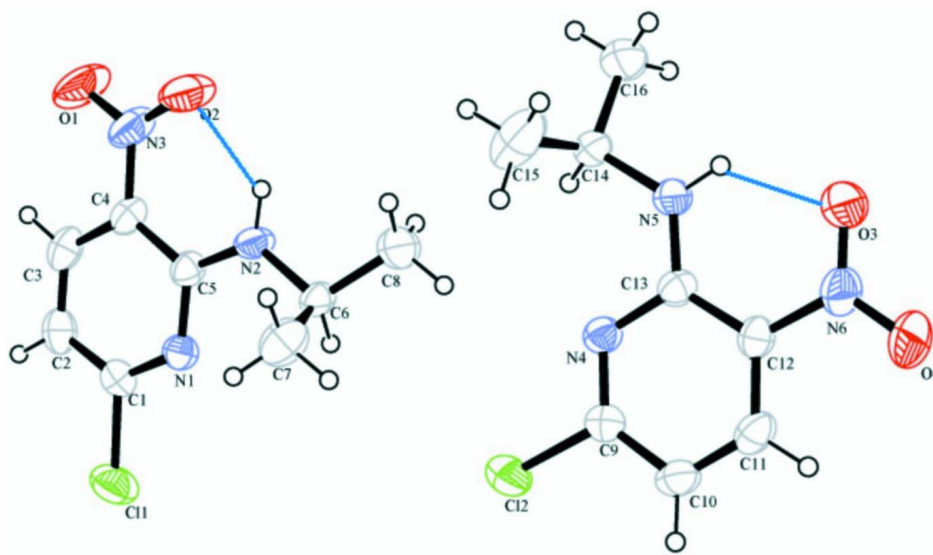
6-Chloro-*N*-isopropyl-3-nitropyridin-2-amine derivatives are of great importance owing to their anticancer activity (Lan *et al.*, 2010; Bavetsias *et al.*, 2010). The title compound is one of the key intermediates in our synthetic investigations of anticancer drugs. Now we synthesized the title compound and report here its molecular and crystal structures. In the title compound, C₈H₁₀ClN₃O₂, (Fig. 1), the bond lengths and angles are within normal ranges (Allen *et al.*, 1987). In crystal, there are two molecules of the compound observed in the asymmetric unit. The intramolecular hydrogen bond N2—H2···O2 (N2···O2 = 2.660 Å) stabilizes the almost coplanar arrangement of the N2—C5—C4—N3—O2 plane and pyridine ring. And there are no classical hydrogen bonds observed in the crystal packing (Fig. 2).

S2. Experimental

A solution of 0.58 g (3.0 mmol) of 2,6-dichloro-3-nitropyridine in 20 ml of dichloromethane was stirred in the ice-water bath for a few minutes, then 0.38 ml (4.5 mmol) isopropylamine was added dropwise. The reaction was stirred in the ice-water bath for 4 h, concentrated under reduced pressure and purified by silica gel column chromatography. Crystals suitable for X-ray analysis were obtained by slow evaporation from a solution of dichloromethane.

S3. Refinement

All H atoms were positioned geometrically and refined as riding (C—H = 0.93 Å–0.98 Å, N—H = 0.86 Å) and allowed to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5U_{\text{eq}}(\text{parent})$.

**Figure 1**

The molecular structure of the title compound with the atom numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. Intramolecular hydrogen bonds are presented by blue lines.

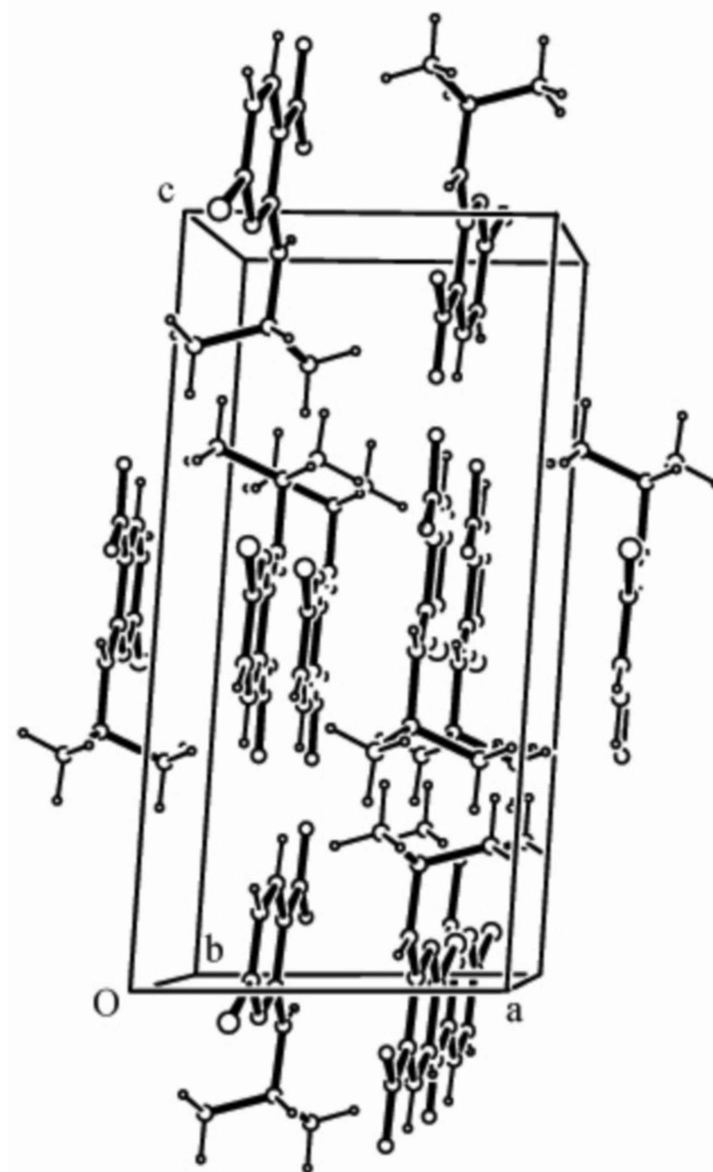


Figure 2

A packing diagram of the title compound.

6-Chloro-3-nitro-N-(propan-2-yl)pyridin-2-amine

Crystal data

$C_8H_{10}ClN_3O_2$

$M_r = 215.64$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 7.4283$ (8) Å

$b = 8.9573$ (10) Å

$c = 15.4301$ (17) Å

$\alpha = 89.672$ (9)°

$\beta = 86.252$ (9)°

$\gamma = 78.860$ (9)°

$V = 1005.16$ (19) Å³

$Z = 4$

$F(000) = 448$

$D_x = 1.425$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.7107$ Å

Cell parameters from 2782 reflections

$\theta = 3.0$ – 29.2 °

$\mu = 0.36$ mm⁻¹

$T = 295$ K $0.28 \times 0.23 \times 0.18$ mm
 Block, colourless

Data collection

Oxford Diffraction Xcalibur Eos diffractometer	8239 measured reflections
Radiation source: fine-focus sealed tube	4073 independent reflections
Graphite monochromator	2773 reflections with $I > 2\sigma(I)$
Detector resolution: 16.0874 pixels mm ⁻¹	$R_{\text{int}} = 0.022$
ω scans	$\theta_{\text{max}} = 26.4^\circ$, $\theta_{\text{min}} = 3.0^\circ$
Absorption correction: multi-scan (<i>CrysAlis PRO</i> ; Oxford Diffraction, 2006)	$h = -9 \rightarrow 8$
$T_{\text{min}} = 0.971$, $T_{\text{max}} = 1.0$	$k = -11 \rightarrow 11$
	$l = -19 \rightarrow 19$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.061$	H-atom parameters constrained
$wR(F^2) = 0.149$	$w = 1/[\sigma^2(F_o^2) + (0.0511P)^2 + 0.4285P]$
$S = 1.07$	where $P = (F_o^2 + 2F_c^2)/3$
4073 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
257 parameters	$\Delta\rho_{\text{max}} = 0.24 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.21 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C11	0.27337 (14)	-0.24866 (9)	0.55983 (6)	0.0755 (3)
C12	0.22248 (13)	0.27284 (9)	0.95299 (6)	0.0649 (3)
O1	0.2452 (5)	0.3317 (3)	0.30153 (16)	0.1036 (10)
O2	0.2403 (4)	0.4488 (3)	0.42175 (16)	0.0911 (9)
O3	0.3152 (4)	0.9485 (3)	1.08072 (16)	0.0773 (7)
O4	0.3156 (4)	0.8329 (3)	1.20304 (15)	0.0806 (8)
N1	0.2670 (3)	0.0404 (3)	0.55475 (14)	0.0438 (6)
N2	0.2640 (4)	0.2945 (3)	0.56967 (15)	0.0510 (6)
H2	0.2646	0.3818	0.5464	0.061*
N3	0.2445 (4)	0.3306 (4)	0.38070 (18)	0.0692 (8)
N4	0.2595 (3)	0.5533 (2)	0.95342 (14)	0.0420 (5)
N5	0.2928 (4)	0.8005 (3)	0.93459 (15)	0.0504 (6)
H5	0.3087	0.8849	0.9562	0.060*
N6	0.3082 (4)	0.8333 (3)	1.12380 (17)	0.0573 (7)

C1	0.2645 (4)	-0.0767 (3)	0.5049 (2)	0.0476 (7)
C2	0.2568 (4)	-0.0768 (4)	0.4162 (2)	0.0576 (8)
H2A	0.2560	-0.1652	0.3850	0.069*
C3	0.2504 (4)	0.0606 (4)	0.3764 (2)	0.0563 (8)
H3	0.2456	0.0676	0.3164	0.068*
C4	0.2512 (4)	0.1893 (3)	0.42550 (18)	0.0487 (7)
C5	0.2595 (4)	0.1783 (3)	0.51714 (17)	0.0421 (6)
C6	0.2678 (4)	0.2846 (3)	0.66435 (17)	0.0452 (7)
H6	0.3556	0.1926	0.6783	0.054*
C7	0.0831 (4)	0.2722 (4)	0.7059 (2)	0.0684 (9)
H7B	-0.0062	0.3597	0.6912	0.103*
H7C	0.0894	0.2673	0.7679	0.103*
H7A	0.0479	0.1818	0.6853	0.103*
C8	0.3374 (6)	0.4205 (4)	0.6968 (2)	0.0769 (11)
H8A	0.4551	0.4237	0.6680	0.115*
H8C	0.3493	0.4121	0.7583	0.115*
H8B	0.2520	0.5120	0.6847	0.115*
C9	0.2499 (4)	0.4388 (3)	1.00420 (19)	0.0439 (7)
C10	0.2613 (4)	0.4363 (3)	1.0939 (2)	0.0503 (7)
H10	0.2556	0.3494	1.1264	0.060*
C11	0.2814 (4)	0.5690 (4)	1.13095 (19)	0.0493 (7)
H11	0.2892	0.5746	1.1907	0.059*
C12	0.2901 (4)	0.6956 (3)	1.08053 (18)	0.0428 (6)
C13	0.2804 (4)	0.6865 (3)	0.98920 (18)	0.0405 (6)
C14	0.2811 (4)	0.7920 (3)	0.84014 (18)	0.0481 (7)
H14	0.3504	0.6925	0.8201	0.058*
C15	0.0870 (5)	0.8048 (6)	0.8159 (2)	0.0928 (14)
H15C	0.0166	0.9022	0.8341	0.139*
H15B	0.0344	0.7258	0.8439	0.139*
H15A	0.0854	0.7946	0.7540	0.139*
C16	0.3737 (5)	0.9121 (4)	0.7985 (2)	0.0737 (10)
H16A	0.3745	0.9037	0.7365	0.111*
H16B	0.4978	0.8982	0.8156	0.111*
H16C	0.3078	1.0110	0.8169	0.111*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.1077 (7)	0.0471 (5)	0.0766 (7)	-0.0291 (5)	0.0014 (5)	-0.0021 (4)
Cl2	0.0873 (6)	0.0453 (5)	0.0662 (6)	-0.0228 (4)	-0.0058 (4)	0.0016 (4)
O1	0.177 (3)	0.092 (2)	0.0360 (15)	-0.0075 (19)	-0.0197 (17)	0.0146 (14)
O2	0.164 (3)	0.0536 (15)	0.0507 (16)	-0.0076 (16)	-0.0119 (16)	0.0113 (13)
O3	0.128 (2)	0.0505 (14)	0.0553 (15)	-0.0203 (14)	-0.0114 (14)	-0.0059 (12)
O4	0.118 (2)	0.0906 (19)	0.0381 (14)	-0.0327 (15)	-0.0006 (13)	-0.0173 (13)
N1	0.0527 (14)	0.0420 (13)	0.0369 (13)	-0.0108 (10)	-0.0001 (11)	-0.0027 (11)
N2	0.0850 (18)	0.0355 (12)	0.0317 (13)	-0.0100 (12)	-0.0043 (12)	0.0031 (10)
N3	0.096 (2)	0.0656 (19)	0.0395 (17)	0.0025 (16)	-0.0126 (15)	0.0071 (15)
N4	0.0463 (13)	0.0424 (13)	0.0370 (13)	-0.0089 (10)	-0.0009 (10)	0.0026 (11)

N5	0.0793 (18)	0.0376 (13)	0.0359 (13)	-0.0146 (12)	-0.0054 (12)	-0.0011 (11)
N6	0.0691 (18)	0.0601 (17)	0.0418 (16)	-0.0106 (13)	-0.0013 (13)	-0.0112 (14)
C1	0.0488 (17)	0.0441 (16)	0.0514 (19)	-0.0130 (13)	-0.0023 (13)	-0.0034 (14)
C2	0.061 (2)	0.061 (2)	0.051 (2)	-0.0123 (15)	-0.0064 (15)	-0.0181 (17)
C3	0.059 (2)	0.071 (2)	0.0363 (17)	-0.0060 (16)	-0.0079 (14)	-0.0083 (16)
C4	0.0536 (18)	0.0546 (18)	0.0356 (16)	-0.0038 (14)	-0.0066 (13)	0.0015 (14)
C5	0.0476 (16)	0.0429 (16)	0.0344 (15)	-0.0051 (12)	-0.0022 (12)	-0.0028 (12)
C6	0.0665 (19)	0.0372 (15)	0.0311 (15)	-0.0067 (13)	-0.0063 (13)	-0.0017 (12)
C7	0.069 (2)	0.083 (2)	0.049 (2)	-0.0039 (18)	0.0027 (17)	0.0019 (18)
C8	0.129 (3)	0.054 (2)	0.053 (2)	-0.028 (2)	-0.016 (2)	-0.0034 (17)
C9	0.0434 (16)	0.0430 (16)	0.0451 (17)	-0.0083 (12)	-0.0017 (13)	0.0028 (13)
C10	0.0507 (18)	0.0546 (18)	0.0471 (18)	-0.0143 (14)	-0.0029 (14)	0.0154 (15)
C11	0.0485 (17)	0.067 (2)	0.0323 (16)	-0.0125 (14)	0.0005 (12)	0.0043 (14)
C12	0.0435 (16)	0.0474 (16)	0.0367 (16)	-0.0073 (12)	-0.0008 (12)	-0.0055 (13)
C13	0.0426 (15)	0.0404 (15)	0.0367 (15)	-0.0037 (11)	-0.0007 (12)	0.0014 (12)
C14	0.0665 (19)	0.0433 (16)	0.0363 (16)	-0.0139 (14)	-0.0075 (14)	0.0038 (13)
C15	0.072 (3)	0.149 (4)	0.059 (2)	-0.024 (3)	-0.011 (2)	0.024 (3)
C16	0.110 (3)	0.068 (2)	0.051 (2)	-0.036 (2)	-0.005 (2)	0.0099 (18)

Geometric parameters (Å, °)

C11—C1	1.746 (3)	C6—H6	0.9800
C12—C9	1.740 (3)	C6—C7	1.500 (4)
O1—N3	1.221 (3)	C6—C8	1.511 (4)
O2—N3	1.231 (3)	C7—H7B	0.9600
O3—N6	1.232 (3)	C7—H7C	0.9600
O4—N6	1.228 (3)	C7—H7A	0.9600
N1—C1	1.308 (3)	C8—H8A	0.9600
N1—C5	1.355 (3)	C8—H8C	0.9600
N2—H2	0.8600	C8—H8B	0.9600
N2—C5	1.329 (3)	C9—C10	1.392 (4)
N2—C6	1.465 (3)	C10—H10	0.9300
N3—C4	1.432 (4)	C10—C11	1.360 (4)
N4—C9	1.297 (3)	C11—H11	0.9300
N4—C13	1.357 (3)	C11—C12	1.381 (4)
N5—H5	0.8600	C12—C13	1.420 (4)
N5—C13	1.333 (3)	C14—H14	0.9800
N5—C14	1.469 (3)	C14—C15	1.496 (4)
N6—C12	1.438 (4)	C14—C16	1.504 (4)
C1—C2	1.374 (4)	C15—H15C	0.9600
C2—H2A	0.9300	C15—H15B	0.9600
C2—C3	1.365 (4)	C15—H15A	0.9600
C3—H3	0.9300	C16—H16A	0.9600
C3—C4	1.384 (4)	C16—H16B	0.9600
C4—C5	1.421 (4)	C16—H16C	0.9600
O1—N3—O2	120.7 (3)	C6—C8—H8C	109.5
O1—N3—C4	119.2 (3)	C6—C8—H8B	109.5

O2—N3—C4	120.1 (3)	C7—C6—H6	108.1
O3—N6—C12	119.4 (2)	C7—C6—C8	112.7 (3)
O4—N6—O3	121.6 (3)	H7B—C7—H7C	109.5
O4—N6—C12	118.9 (3)	H7B—C7—H7A	109.5
N1—C1—C11	114.6 (2)	H7C—C7—H7A	109.5
N1—C1—C2	126.9 (3)	C8—C6—H6	108.1
N1—C5—C4	118.8 (2)	H8A—C8—H8C	109.5
N2—C5—N1	116.6 (2)	H8A—C8—H8B	109.5
N2—C5—C4	124.6 (3)	H8C—C8—H8B	109.5
N2—C6—H6	108.1	C9—N4—C13	118.6 (2)
N2—C6—C7	111.4 (2)	C9—C10—H10	122.1
N2—C6—C8	108.3 (2)	C10—C9—C12	118.0 (2)
N4—C9—C12	115.5 (2)	C10—C11—H11	119.7
N4—C9—C10	126.5 (3)	C10—C11—C12	120.5 (3)
N4—C13—C12	118.9 (2)	C11—C10—C9	115.7 (3)
N5—C13—N4	116.6 (2)	C11—C10—H10	122.1
N5—C13—C12	124.4 (2)	C11—C12—N6	117.8 (3)
N5—C14—H14	107.9	C11—C12—C13	119.7 (3)
N5—C14—C15	111.9 (3)	C12—C11—H11	119.7
N5—C14—C16	108.4 (2)	C13—N5—H5	117.7
C1—N1—C5	118.3 (2)	C13—N5—C14	124.6 (2)
C1—C2—H2A	122.0	C13—C12—N6	122.5 (3)
C2—C1—C11	118.4 (2)	C14—N5—H5	117.7
C2—C3—H3	120.0	C14—C15—H15C	109.5
C2—C3—C4	120.0 (3)	C14—C15—H15B	109.5
C3—C2—C1	116.0 (3)	C14—C15—H15A	109.5
C3—C2—H2A	122.0	C14—C16—H16A	109.5
C3—C4—N3	117.8 (3)	C14—C16—H16B	109.5
C3—C4—C5	120.0 (3)	C14—C16—H16C	109.5
C4—C3—H3	120.0	C15—C14—H14	107.9
C5—N2—H2	117.6	C15—C14—C16	112.8 (3)
C5—N2—C6	124.8 (2)	H15C—C15—H15B	109.5
C5—C4—N3	122.2 (3)	H15C—C15—H15A	109.5
C6—N2—H2	117.6	H15B—C15—H15A	109.5
C6—C7—H7B	109.5	C16—C14—H14	107.9
C6—C7—H7C	109.5	H16A—C16—H16B	109.5
C6—C7—H7A	109.5	H16A—C16—H16C	109.5
C6—C8—H8A	109.5	H16B—C16—H16C	109.5

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>
N2—H2···O2	0.86	2.02	2.660 (3)	130
N5—H5···O3	0.86	2.01	2.653 (3)	130