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Crystal structure of 3,5-bis(4-chlorophenyl)-1-propyl-1,3,5-triazacyclohexane

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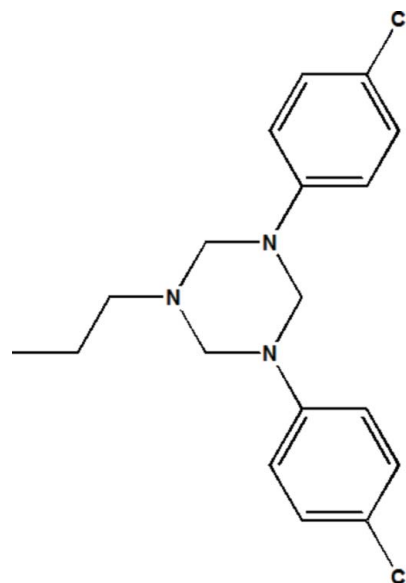
In the title molecule, C₁₈H₂₁Cl₂N₃, the triazacyclohexane ring adopts a chair conformation with both 4-chlorophenyl substituents in axial positions and the propyl group in an equatorial site. The dihedral angle between the planes of the benzene rings is 49.5 (1)°. In the crystal, molecules are arranged in a head-to-tail fashion, forming columns along [010], and pairs of weak C—H... π interactions form inversion dimers between columns.

Keywords: crystal structure; triazacyclohexane; C—H... π interactions.

CCDC reference: 1020727

1. Related literature

For conformations of 1,3,5-triaryl derivatives of 1,3,5-triazacyclohexane, see: Baker *et al.* (1978); Bouchemma *et al.* (1989, 1990); Bushweller (1995); Kleinpeter *et al.* (2005); Duke *et al.* (1973); Gilardi *et al.* (2003); Giumanini *et al.* (1985); Latreche *et al.* (2006); Mloston *et al.* (2006); Freeman *et al.* (2005); Wiberg *et al.* (1999).



2. Experimental

2.1. Crystal data

C ₁₈ H ₂₁ Cl ₂ N ₃	$\gamma = 99.055 (3)^\circ$
$M_r = 350.28$	$V = 893.19 (8) \text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 2$
$a = 6.0785 (3) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 10.3190 (6) \text{ \AA}$	$\mu = 0.37 \text{ mm}^{-1}$
$c = 14.4360 (8) \text{ \AA}$	$T = 295 \text{ K}$
$\alpha = 91.570 (3)^\circ$	$0.22 \times 0.13 \times 0.07 \text{ mm}$
$\beta = 91.946 (2)^\circ$	

2.2. Data collection

Nonius KappaCCD diffractometer	13643 measured reflections
Absorption correction: multi-scan (Blessing, 1995)	3448 independent reflections
$T_{\min} = 0.274$, $T_{\max} = 0.467$	2751 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.016$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.049$	209 parameters
$wR(F^2) = 0.142$	H-atom parameters constrained
$S = 1.03$	$\Delta\rho_{\text{max}} = 0.45 \text{ e \AA}^{-3}$
3448 reflections	$\Delta\rho_{\text{min}} = -0.25 \text{ e \AA}^{-3}$

Table 1

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

C_g is centroid of C21–C26 ring.

D—H...A	D—H	H...A	D...A	D—H...A
C2—H2A...C _g ⁱ	0.96	2.92	3.668 (3)	134

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

Data collection: COLLECT (Nonius, 1997); cell refinement: SCALEPACK (Otwinowski & Minor, 1997); data reduction: DENZO (Otwinowski & Minor, 1997) and SCALEPACK; program(s) used to solve structure: SIR2002 (Burla *et al.*, 2005); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012) and

DIAMOND (Brandenburg & Berndt, 2001); software used to prepare material for publication: *WinGX* (Farrugia, 2012).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: LH5723).

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supporting information

Acta Cryst. (2014). E70, o1061–o1062 [doi:10.1107/S1600536814019060]

Crystal structure of 3,5-bis(4-chlorophenyl)-1-propyl-1,3,5-triazacyclohexane

Leila Lefrada, Ahcene Bouchemma, Sofiane Bouacida, Nicolas Claiser and Mohamed Souhassou

S1. Comment

The conformational behaviour of substituted cyclohexanes as well as heterocyclohexanes has been the subject of numerous studies (Bushweller, 1995). The ring normally adopts the chair conformation unless specific intramolecular interactions stabilize the twist (Kleinpeter *et al.*, 2005) or boat conformers (Freeman *et al.*, 2005). Saturated six-membered rings are prevalent in organic chemistry. For cyclohexane, experimental and computational studies have established that the chair conformation is 5.5 kcal/mol more stable than the twist form (Wiberg *et al.*, 1999). *N,N,N'*-Trisubstituted 1,3,5-triazinanes are of interest as precursors for the preparation of different N-substituted imidazoles (Mloston *et al.*, 2006). The heterocyclic nucleus is expected to adopt a chair conformation and four distinct patterns of substituent orientation have to be considered, eee, eea, eaa and aaa, where e = equatorial and a = axial, with each of the conformers having axial repulsions involving the substituents or lone pairs of electrons on the N atoms. Several 1,3,5-tri-alkyl derivatives have been investigated in solution by dipole moment measurements and the results interpreted in terms of the eee conformer, the eea conformer (Baker *et al.*, 1978), and varying amounts of the eee, eea and eaa conformers (Duke *et al.*, 1973). Various 1,3,5-triaryl-1,3,5-triazacyclohexanes adopt the diaxial-equatorial orientation of substituents in the solid state thus avoiding 1,3-diaxial lone-pair repulsions (Giumanini *et al.* 1985; Gilardi *et al.* 2003 Bouchemma *et al.* 1989; 1990).

In the present work, a new derivate (I) of triazacyclohexane is reported and molecular structure is shown in Fig. 1. The 1,3,5-triazacyclohexane ring is in a chair conformation which is typical of this ring (Gilardi *et al.* 2003). The structure of a similar compound *viz* 1-propyl-3,5-bis-(4-fluorophenyl)-1,3,5-triazacyclohexane (II) has been reported (Latreche *et al.* 2006). In both (I) and (II) the heterocyclic rings adopt chair conformations with two fluorophenyl substituents situated in axial positions and a third group (propyl) equatorial. The dihedral angle between the benzene rings (C11-C16/C21-C26) is 49.5 (1)°. In the crystal, molecules are arranged in a 'head to tail' fashion forming columns along [010] (see Fig. 2) and pairs weak C—H... π interactions form inversion dimers between columns.

S2. Experimental

The title compound was obtained by mixing a 2:1:1 stoichiometric ratio of propylamine and 4-chloroaniline with formalin in ethanol (25 ml) at 293K. The resulting solution was evaporated on a rotary evaporator to dryness and the white residue was recrystallized from cyclohexane.

S3. Refinement

All non-H atoms were refined with anisotropic atomic displacement parameters. All H atoms were located in difference Fourier maps but introduced in calculated positions and treated as riding on their parent C atom, with C—H distances of 0.93 Å (C_{aromatic}), 0.97 Å ($C_{\text{methylene}}$) and 0.96 Å (C_{methyl}) with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(C_{\text{aromatic}} \text{ and } C_{\text{methylene}})$ or $1.5 U_{\text{eq}}(C_{\text{methyl}})$.

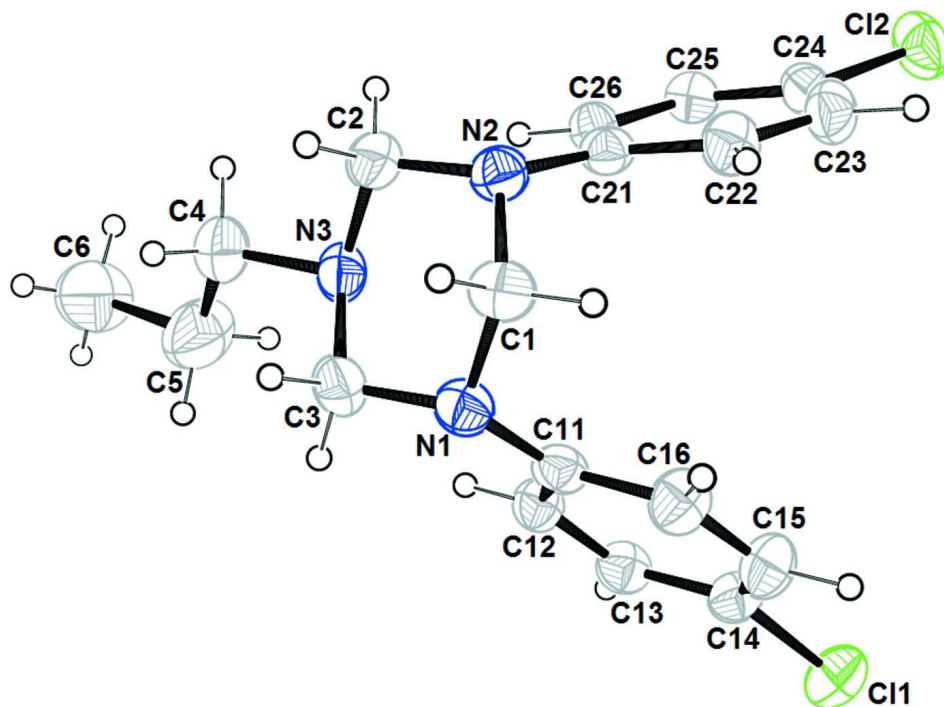
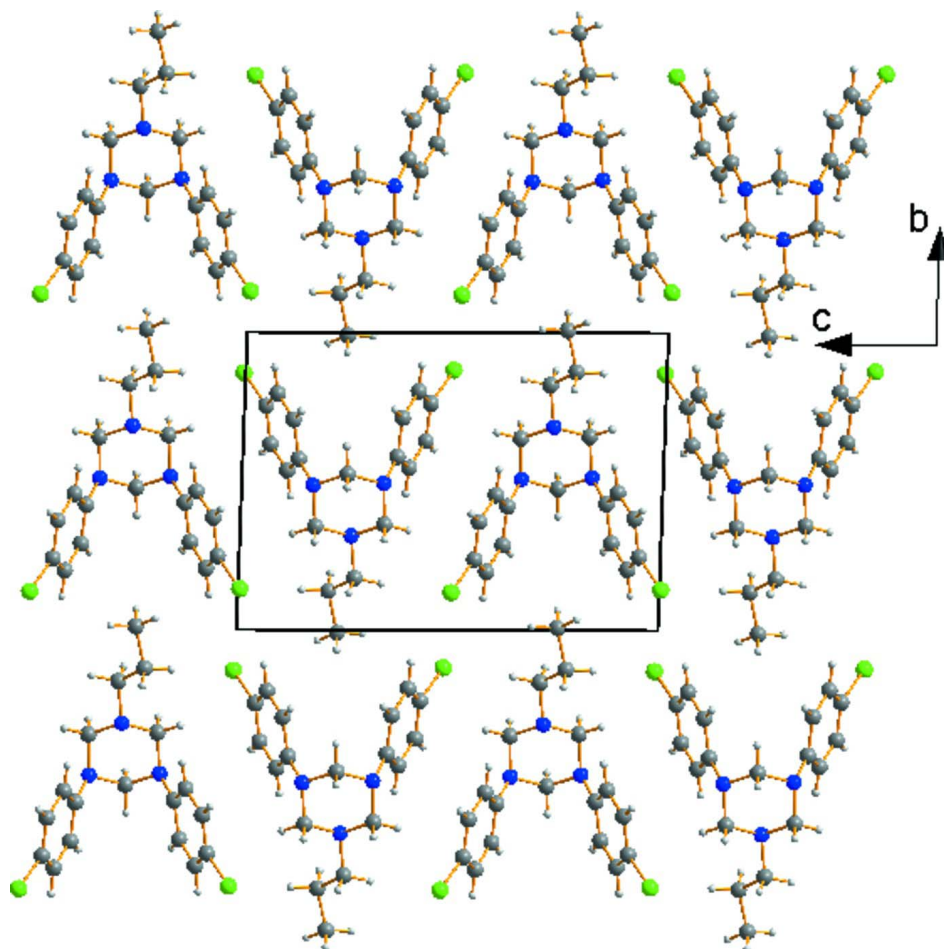


Figure 1

The molecular structure of the title compound with displacement ellipsoids drawn at the 50% probability level.

**Figure 2**

Part of the crystal structure of the title compound showing the 'head to tail' arrangement of molecules arranged in columns.

3,5-Bis(4-chlorophenyl)-1-propyl-1,3,5-triazacyclohexane

Crystal data

$C_{18}H_{21}Cl_2N_3$

$M_r = 350.28$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 6.0785$ (3) Å

$b = 10.3190$ (6) Å

$c = 14.4360$ (8) Å

$\alpha = 91.570$ (3)°

$\beta = 91.946$ (2)°

$\gamma = 99.055$ (3)°

$V = 893.19$ (8) Å³

$Z = 2$

$F(000) = 368$

$D_x = 1.302$ Mg m⁻³

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

Cell parameters from 13346 reflections

$\theta = 1.5$ – 27.3 °

$\mu = 0.37$ mm⁻¹

$T = 295$ K

Prism, colourless

$0.22 \times 0.13 \times 0.07$ mm

Data collection

Nonius KappaCCD
diffractometer

Graphite monochromator

ω and φ scans

Absorption correction: multi-scan
(Blessing, 1995)

$T_{\min} = 0.274$, $T_{\max} = 0.467$

13643 measured reflections

3448 independent reflections

2751 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.016$

$\theta_{\max} = 27.3^\circ$, $\theta_{\min} = 1.4^\circ$

$h = -7 \rightarrow 7$

$k = -12 \rightarrow 12$

$l = -17 \rightarrow 17$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.049$

$wR(F^2) = 0.142$

$S = 1.03$

3448 reflections

209 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

$w = 1/[\sigma^2(F_o^2) + (0.0652P)^2 + 0.413P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.45 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.25 \text{ e } \text{\AA}^{-3}$

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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	-0.3001 (4)	0.4766 (3)	0.25020 (16)	0.0613 (6)
H1A	-0.3547	0.383	0.2459	0.074*
H1B	-0.4281	0.5221	0.2505	0.074*
C2	-0.0783 (4)	0.6456 (2)	0.34207 (17)	0.0633 (6)
H2A	0.0103	0.6653	0.3993	0.076*
H2B	-0.1997	0.6964	0.3432	0.076*
C3	-0.0772 (5)	0.6525 (2)	0.17830 (18)	0.0652 (6)
H3A	-0.1979	0.7039	0.1778	0.078*
H3B	0.013	0.6767	0.1254	0.078*
C4	0.1485 (6)	0.8253 (3)	0.2756 (2)	0.0847 (8)
H4A	0.0283	0.8751	0.2638	0.102*
H4B	0.2019	0.8438	0.3394	0.102*
C5	0.3281 (8)	0.8689 (4)	0.2147 (3)	0.1185 (14)
H5A	0.2688	0.865	0.1513	0.142*
H5B	0.4368	0.8096	0.2185	0.142*
C6	0.4450 (8)	1.0093 (4)	0.2390 (3)	0.1321 (17)
H6A	0.3366	1.0677	0.2395	0.198*

H6B	0.5531	1.0364	0.1935	0.198*
H6C	0.5184	1.0117	0.2991	0.198*
C11	-0.0360 (4)	0.4255 (2)	0.13332 (14)	0.0505 (5)
C12	0.1801 (4)	0.4642 (2)	0.10651 (16)	0.0556 (5)
H12	0.2467	0.5511	0.1166	0.067*
C13	0.2992 (4)	0.3762 (2)	0.06492 (16)	0.0590 (6)
H13	0.4436	0.4041	0.0465	0.071*
C14	0.2027 (4)	0.2473 (2)	0.05101 (16)	0.0585 (6)
C15	-0.0100 (4)	0.2056 (2)	0.07759 (19)	0.0691 (7)
H15	-0.0742	0.1181	0.0685	0.083*
C16	-0.1281 (4)	0.2940 (2)	0.11776 (19)	0.0666 (6)
H16	-0.2732	0.2653	0.135	0.08*
C21	-0.0324 (4)	0.4175 (2)	0.36755 (14)	0.0491 (5)
C22	-0.1257 (4)	0.2865 (2)	0.37598 (18)	0.0622 (6)
H22	-0.2735	0.2589	0.3566	0.075*
C23	-0.0036 (4)	0.1967 (2)	0.41250 (19)	0.0660 (6)
H23	-0.0682	0.1094	0.4173	0.079*
C24	0.2138 (4)	0.2375 (2)	0.44168 (16)	0.0583 (6)
C25	0.3124 (4)	0.3650 (2)	0.43325 (16)	0.0565 (5)
H25	0.4605	0.3913	0.4526	0.068*
C26	0.1900 (4)	0.4544 (2)	0.39570 (15)	0.0536 (5)
H26	0.2579	0.5408	0.3892	0.064*
N1	-0.1697 (3)	0.51418 (19)	0.16948 (13)	0.0578 (5)
N2	-0.1687 (3)	0.50789 (19)	0.33683 (13)	0.0561 (5)
N3	0.0600 (4)	0.68403 (18)	0.26379 (14)	0.0615 (5)
Cl1	0.34990 (13)	0.13680 (7)	-0.00400 (5)	0.0841 (3)
Cl2	0.36408 (14)	0.12627 (7)	0.49405 (6)	0.0865 (3)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0497 (13)	0.0749 (16)	0.0606 (14)	0.0138 (11)	0.0016 (10)	0.0033 (11)
C2	0.0733 (16)	0.0592 (14)	0.0614 (14)	0.0235 (12)	0.0035 (12)	-0.0017 (11)
C3	0.0750 (16)	0.0586 (14)	0.0659 (15)	0.0211 (12)	0.0029 (12)	0.0117 (11)
C4	0.108 (2)	0.0557 (15)	0.090 (2)	0.0095 (15)	0.0082 (18)	0.0044 (14)
C5	0.146 (4)	0.088 (2)	0.115 (3)	-0.010 (2)	0.040 (3)	-0.009 (2)
C6	0.173 (4)	0.088 (3)	0.115 (3)	-0.044 (3)	0.017 (3)	0.006 (2)
C11	0.0516 (12)	0.0559 (12)	0.0421 (10)	0.0039 (10)	-0.0054 (9)	0.0039 (9)
C12	0.0574 (13)	0.0499 (12)	0.0562 (12)	-0.0013 (10)	-0.0019 (10)	0.0039 (10)
C13	0.0545 (13)	0.0635 (14)	0.0572 (13)	0.0030 (11)	0.0037 (10)	0.0052 (11)
C14	0.0639 (14)	0.0586 (13)	0.0519 (12)	0.0084 (11)	-0.0040 (10)	-0.0039 (10)
C15	0.0691 (16)	0.0530 (14)	0.0799 (17)	-0.0045 (12)	-0.0021 (13)	-0.0076 (12)
C16	0.0536 (13)	0.0641 (15)	0.0778 (16)	-0.0036 (11)	0.0046 (12)	-0.0022 (12)
C21	0.0510 (12)	0.0533 (12)	0.0431 (11)	0.0077 (9)	0.0077 (9)	-0.0002 (9)
C22	0.0524 (13)	0.0587 (14)	0.0725 (15)	-0.0012 (11)	0.0051 (11)	0.0020 (11)
C23	0.0690 (16)	0.0492 (13)	0.0782 (16)	0.0015 (11)	0.0125 (13)	0.0071 (11)
C24	0.0664 (15)	0.0554 (13)	0.0564 (13)	0.0164 (11)	0.0126 (11)	0.0087 (10)
C25	0.0528 (12)	0.0610 (13)	0.0561 (13)	0.0102 (10)	0.0020 (10)	0.0024 (10)

C26	0.0562 (13)	0.0486 (12)	0.0541 (12)	0.0018 (10)	0.0059 (10)	0.0023 (9)
N1	0.0583 (11)	0.0619 (11)	0.0538 (11)	0.0113 (9)	-0.0016 (8)	0.0046 (9)
N2	0.0567 (11)	0.0603 (11)	0.0525 (10)	0.0130 (9)	0.0024 (8)	0.0000 (8)
N3	0.0734 (13)	0.0446 (10)	0.0665 (12)	0.0104 (9)	-0.0022 (10)	0.0023 (9)
C11	0.0883 (5)	0.0780 (5)	0.0872 (5)	0.0207 (4)	0.0049 (4)	-0.0199 (4)
C12	0.0940 (5)	0.0765 (5)	0.0981 (6)	0.0357 (4)	0.0131 (4)	0.0278 (4)

Geometric parameters (Å, °)

C1—N1	1.458 (3)	C11—C16	1.393 (3)
C1—N2	1.461 (3)	C11—N1	1.417 (3)
C1—H1A	0.97	C12—C13	1.384 (3)
C1—H1B	0.97	C12—H12	0.93
C2—N2	1.439 (3)	C13—C14	1.372 (3)
C2—N3	1.458 (3)	C13—H13	0.93
C2—H2A	0.97	C14—C15	1.368 (4)
C2—H2B	0.97	C14—C11	1.747 (2)
C3—N1	1.450 (3)	C15—C16	1.374 (4)
C3—N3	1.465 (3)	C15—H15	0.93
C3—H3A	0.97	C16—H16	0.93
C3—H3B	0.97	C21—C26	1.389 (3)
C4—C5	1.448 (5)	C21—C22	1.392 (3)
C4—N3	1.475 (3)	C21—N2	1.412 (3)
C4—H4A	0.97	C22—C23	1.380 (4)
C4—H4B	0.97	C22—H22	0.93
C5—C6	1.536 (4)	C23—C24	1.371 (4)
C5—H5A	0.97	C23—H23	0.93
C5—H5B	0.97	C24—C25	1.367 (3)
C6—H6A	0.96	C24—C12	1.747 (2)
C6—H6B	0.96	C25—C26	1.384 (3)
C6—H6C	0.96	C25—H25	0.93
C11—C12	1.383 (3)	C26—H26	0.93
N1—C1—N2	111.86 (19)	C13—C12—H12	119.3
N1—C1—H1A	109.2	C14—C13—C12	119.6 (2)
N2—C1—H1A	109.2	C14—C13—H13	120.2
N1—C1—H1B	109.2	C12—C13—H13	120.2
N2—C1—H1B	109.2	C15—C14—C13	120.5 (2)
H1A—C1—H1B	107.9	C15—C14—C11	119.76 (19)
N2—C2—N3	111.81 (18)	C13—C14—C11	119.70 (19)
N2—C2—H2A	109.3	C14—C15—C16	119.6 (2)
N3—C2—H2A	109.3	C14—C15—H15	120.2
N2—C2—H2B	109.3	C16—C15—H15	120.2
N3—C2—H2B	109.3	C15—C16—C11	121.7 (2)
H2A—C2—H2B	107.9	C15—C16—H16	119.1
N1—C3—N3	112.04 (19)	C11—C16—H16	119.1
N1—C3—H3A	109.2	C26—C21—C22	117.6 (2)
N3—C3—H3A	109.2	C26—C21—N2	123.0 (2)

N1—C3—H3B	109.2	C22—C21—N2	119.3 (2)
N3—C3—H3B	109.2	C23—C22—C21	121.4 (2)
H3A—C3—H3B	107.9	C23—C22—H22	119.3
C5—C4—N3	113.4 (3)	C21—C22—H22	119.3
C5—C4—H4A	108.9	C24—C23—C22	119.4 (2)
N3—C4—H4A	108.9	C24—C23—H23	120.3
C5—C4—H4B	108.9	C22—C23—H23	120.3
N3—C4—H4B	108.9	C25—C24—C23	120.9 (2)
H4A—C4—H4B	107.7	C25—C24—C12	119.54 (19)
C4—C5—C6	112.8 (3)	C23—C24—C12	119.48 (19)
C4—C5—H5A	109	C24—C25—C26	119.5 (2)
C6—C5—H5A	109	C24—C25—H25	120.2
C4—C5—H5B	109	C26—C25—H25	120.2
C6—C5—H5B	109	C25—C26—C21	121.2 (2)
H5A—C5—H5B	107.8	C25—C26—H26	119.4
C5—C6—H6A	109.5	C21—C26—H26	119.4
C5—C6—H6B	109.5	C11—N1—C3	118.73 (19)
H6A—C6—H6B	109.5	C11—N1—C1	118.44 (19)
C5—C6—H6C	109.5	C3—N1—C1	109.38 (19)
H6A—C6—H6C	109.5	C21—N2—C2	118.41 (19)
H6B—C6—H6C	109.5	C21—N2—C1	118.07 (18)
C12—C11—C16	117.3 (2)	C2—N2—C1	109.76 (19)
C12—C11—N1	123.2 (2)	C2—N3—C3	108.2 (2)
C16—C11—N1	119.4 (2)	C2—N3—C4	108.2 (2)
C11—C12—C13	121.3 (2)	C3—N3—C4	112.7 (2)
C11—C12—H12	119.3		
N3—C4—C5—C6	-170.1 (3)	C16—C11—N1—C3	-175.1 (2)
C16—C11—C12—C13	0.7 (3)	C12—C11—N1—C1	-136.1 (2)
N1—C11—C12—C13	-175.0 (2)	C16—C11—N1—C1	48.3 (3)
C11—C12—C13—C14	-0.9 (3)	N3—C3—N1—C11	-83.0 (3)
C12—C13—C14—C15	0.2 (4)	N3—C3—N1—C1	57.2 (3)
C12—C13—C14—C11	178.45 (18)	N2—C1—N1—C11	84.7 (2)
C13—C14—C15—C16	0.6 (4)	N2—C1—N1—C3	-55.7 (3)
C11—C14—C15—C16	-177.6 (2)	C26—C21—N2—C2	-5.2 (3)
C14—C15—C16—C11	-0.8 (4)	C22—C21—N2—C2	170.6 (2)
C12—C11—C16—C15	0.1 (4)	C26—C21—N2—C1	131.2 (2)
N1—C11—C16—C15	176.0 (2)	C22—C21—N2—C1	-53.0 (3)
C26—C21—C22—C23	1.2 (3)	N3—C2—N2—C21	81.6 (2)
N2—C21—C22—C23	-174.9 (2)	N3—C2—N2—C1	-58.1 (3)
C21—C22—C23—C24	0.4 (4)	N1—C1—N2—C21	-83.5 (3)
C22—C23—C24—C25	-1.4 (4)	N1—C1—N2—C2	56.4 (3)
C22—C23—C24—C12	176.34 (19)	N2—C2—N3—C3	58.6 (2)
C23—C24—C25—C26	0.7 (4)	N2—C2—N3—C4	-179.0 (2)
C12—C24—C25—C26	-177.02 (17)	N1—C3—N3—C2	-58.3 (3)
C24—C25—C26—C21	0.9 (3)	N1—C3—N3—C4	-177.8 (2)
C22—C21—C26—C25	-1.8 (3)	C5—C4—N3—C2	166.4 (3)
N2—C21—C26—C25	174.05 (19)	C5—C4—N3—C3	-74.0 (4)

C12—C11—N1—C3 0.5 (3)

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>
C2—H2A···Cg ⁱ	0.96	2.92	3.668 (3)	134

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