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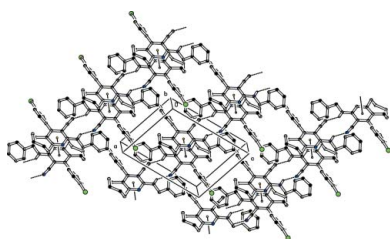
Crystal structures of 2-benzylamino-4-(4-bromo-phenyl)-6,7,8,9-tetrahydro-5H-cyclohepta[b]-pyridine-3-carbonitrile and 2-benzylamino-4-(4-chlorophenyl)-6,7,8,9-tetrahydro-5H-cyclohepta[b]pyridine-3-carbonitrile

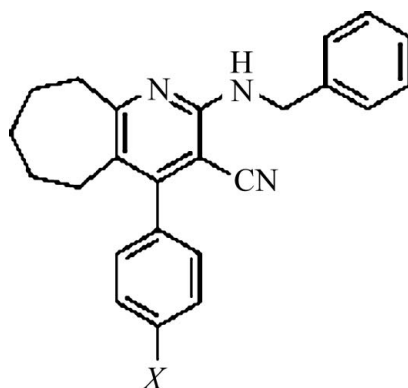
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In the title compounds, $C_{24}H_{22}BrN_3$, (I), and $C_{24}H_{22}ClN_3$, (II), the 2-aminopyridine ring is fused with a cycloheptane ring, which adopts a half-chair conformation. The planes of the phenyl and benzene rings are inclined to that of the central pyridine ring [r.m.s. deviations = 0.0083 (1) and 0.0093 (1) Å for (I) and (II), respectively] by 62.47 (17) and 72.51 (14)°, respectively, in (I), and by 71.44 (9) and 54.90 (8)°, respectively, in (II). The planes of the aromatic rings are inclined to one another by 53.82 (17)° in (I) and by 58.04 (9)° in (II). In the crystals of both (I) and (II), pairs of $N-H \cdots N_{\text{nitrile}}$ hydrogen bonds link the molecules, forming inversion dimers with $R_2^2(12)$ ring motifs. In (I), the resulting dimers are connected through $C-H \cdots Br$ hydrogen bonds, forming sheets parallel to (10 $\bar{1}$), and π - π interactions [inter-centroid distance = 3.7821 (16) Å] involving inversion-related pyridine rings, forming a three-dimensional network. In (II), the resulting dimers are connected through π - π interactions [inter-centroid distance = 3.771 (2) Å] involving inversion-related pyridine rings, forming a two-dimensional network lying parallel to (001).

1. Chemical context

The heterocyclic skeleton containing a nitrogen atom is the basis of many essential pharmaceuticals and of many physiologically active natural products. Molecules containing heterocyclic substructures continue to be attractive targets for synthesis since they often exhibit diverse and important biological properties. Pyridine is used in the pharmaceutical industry as a raw material for various drugs, vitamins and fungicides, and as a solvent (Shinkai *et al.*, 2000; Jansen *et al.*, 2001; Amr *et al.*, 2006). Pyridines are also omnipresent in medicaments and in agrochemicals (Tomlin, 1994). Pyridine derivatives have occupied a unique position in medicinal chemistry. Among them, 2-amino-3-cyanopyridines have been identified as IKK- β inhibitors (Murata *et al.*, 2003). Many fused cyanopyridines have also been shown to have a wide spectrum of biological activity (Boschelli *et al.*, 2004). Our interest in the preparation of pharmacologically active 3-cyanopyridine compounds led us to synthesize the title compounds and we report herein on their crystal structures.



(I) $X = \text{Br}$ (II) $X = \text{Cl}$

2. Structural commentary

The molecular structures of the title compounds, (I) and (II), are shown in Figs. 1 and 2, respectively. The bromo derivative (I), crystallizes in the monoclinic space group $P2_1/n$ while the chloro derivative (II), crystallizes in the triclinic space group $P\bar{1}$.

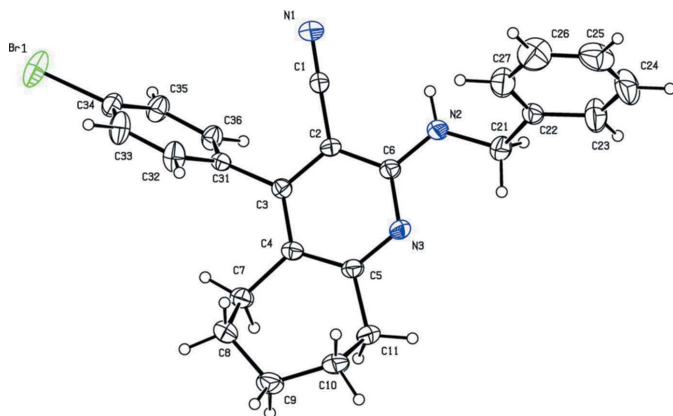


Figure 1

The molecular structure of compound (I), showing 50% probability displacement ellipsoids and the atom labelling.

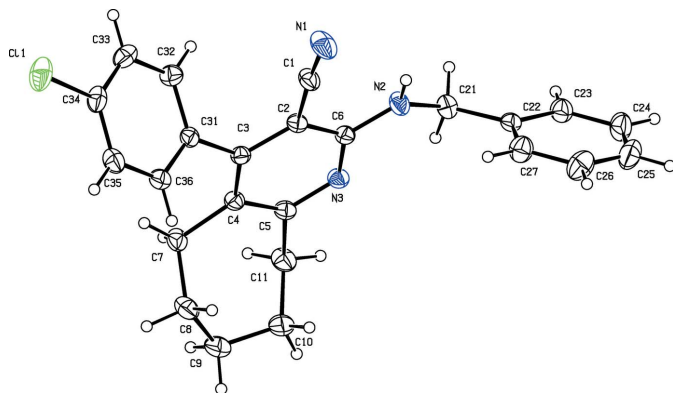


Figure 2

The molecular structure of (II), showing 50% probability displacement ellipsoids and the atom labelling.

Table 1

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N2-H2\cdots N1^i$	0.86	2.28	3.010 (3)	143
$C21-H21B\cdots Br1^{ii}$	0.97	2.90	3.703 (3)	141

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

In both compounds, the pyridine ring is connected to a benzene ring by a $-\text{CH}_2-\text{NH}_2-$ chain, as found in a similar structure N^6 -(4-fluorobenzyl)-3-nitropyridine-2,6-diamine (Ge & Qian, 2011). As expected, the pyridine ring ($\text{C}2-\text{C}6/\text{N}3$) is planar with r.m.s. deviations of 0.0083 and 0.0093 \AA in compounds (I) and (II), respectively. In both compounds, the cycloheptane ring adopts a half-chair conformation, with puckering parameters $Q2 = 0.415$ (3) \AA , $\varphi2 = 310.1$ (4) $^\circ$ and $Q3 = 0.637$ (3) \AA and $\varphi3 = 283.4$ (3) $^\circ$ for compound (I) and $Q2 = 0.475$ (2) \AA , $\varphi2 = 310.3$ (2) $^\circ$ and $Q3 = 0.635$ (2) \AA and $\varphi3 = 283.58$ (17) $^\circ$ for compound (II). The amine N atom, N2, attached to the pyridine ring ($\text{N}3/\text{C}2-\text{C}6$) deviates by only 0.0107 (1) and 0.0073 (1) \AA from the ring plane in (I) and (II), respectively. Steric hindrance rotates the benzene ring ($\text{C}31-\text{C}36$) out of the plane of the central pyridine ring by 72.51 (14) $^\circ$ in compound (I) and by only 54.90 (8) $^\circ$ in compound (II). The benzene ring is inclined to the phenyl ring ($\text{C}22-\text{C}27$) by 53.82 (17) in (I) and by 58.04 (9) $^\circ$ in (II).

3. Supramolecular features

In the crystal of (I), molecules are linked by pairs of $\text{N}-\text{H}\cdots\text{N}_{\text{nitrile}}$ hydrogen bonds, forming inversion dimers with $R_2^2(12)$ ring motifs (Table 1 and Fig. 3). The resulting dimers are connected through $\text{C}-\text{H}\cdots\text{Br}$ hydrogen bonds, forming sheets lying parallel to $(10\bar{1})$. The sheets are connected by weak $\pi-\pi$ stacking interactions involving adjacent inversion-related pyridine rings with a centroid-to-centroid distance of 3.7710 (7) \AA , as shown in Fig. 3. These interactions lead to the formation of a three-dimensional network.

In the crystal of (II), molecules are also linked by pairs of $\text{N}-\text{H}\cdots\text{N}_{\text{nitrile}}$ hydrogen bonds, forming inversion dimers with $R_2^2(12)$ ring motifs (Table 2 and Fig. 4). The dimers are connected through weak $\pi-\pi$ interactions involving inversion-related pyridine rings with a centroid-to-centroid distance of

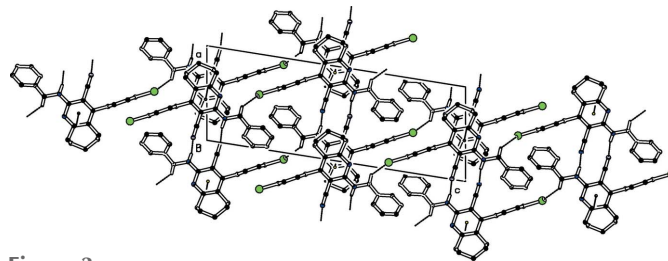


Figure 3

Crystal packing diagram of compound (I), viewed along the b axis. Hydrogen bonds (see Table 1 for details) and $\pi-\pi$ interactions are shown as dashed lines (centroids are shown as small circles). H atoms not involved in hydrogen bonding have been omitted for clarity.

Table 2

Hydrogen-bond geometry (Å, °) for (II).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N2-H2\cdots N1^i$	0.86	2.26	3.007 (2)	145

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

3.7818 (2) Å (Fig. 4). The resulting structure is a two-dimensional network lying parallel to (001).

4. Synthesis and crystallization

Compounds (I) and (II) were prepared in a similar manner using 4-bromo aldehyde (1 mmol) for compound (I) and 4-chloro aldehyde (1 mmol) for compound (II). A mixture of cycloheptanone (1 mmol), aromatic aldehyde (1 mmol), malononitrile (1 mmol) and benzylamine (1 mmol) were taken in ethanol (10 ml) to which *p*-TSA (*p*-toluenesulfonic acid) (1.0 mmol) was added. The reaction mixture was heated under reflux for 2–3 h. On completion of the reaction, verified by thin-layer chromatography (TLC), the mixture was poured into crushed ice and extracted with ethyl acetate. The excess solvent was removed under vacuum and the residue was subjected to column chromatography using a petroleum ether/ethyl acetate mixture (97:3 v/v) as eluent to afford the pure products. They were recrystallized from ethyl acetate, giving colourless crystals of compounds (I) [m.p. 417 K; yield 74%] and (II) [m.p. 397 K; yield 75%].

Table 3

Experimental details.

	(I)	(II)
Crystal data		
Chemical formula	$C_{24}H_{22}BrN_3$	$C_{24}H_{22}ClN_3$
M_r	432.35	387.89
Crystal system, space group	Monoclinic, $P2_1/n$	Triclinic, $P\bar{1}$
Temperature (K)	293	293
a, b, c (Å)	8.9710 (3), 9.3794 (4), 24.9788 (9)	9.002 (5), 10.097 (5), 11.856 (5)
α, β, γ (°)	90, 99.002 (2), 90	94.939 (5), 108.204 (5), 101.272 (5)
V (Å ³)	2075.89 (14)	991.3 (8)
Z	4	2
Radiation type	Mo $K\alpha$	Mo $K\alpha$
μ (mm ⁻¹)	1.99	0.21
Crystal size (mm)	0.21 × 0.19 × 0.18	0.21 × 0.19 × 0.18
Data collection		
Diffractionmeter	Bruker Kappa APEXII	Bruker Kappa APEXII
Absorption correction	Multi-scan (SADABS; Bruker, 2004)	Multi-scan (SADABS; Bruker, 2004)
T_{min} – T_{max}	0.967, 0.974	0.967, 0.974
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	51599, 3863, 2927	24808, 3685, 2918
R_{int}	0.040	0.026
$(\sin \theta/\lambda)_{max}$ (Å ⁻¹)	0.606	0.606
Refinement		
$R[F^2 > 2\sigma(F^2)]$, $wR(F^2)$, S	0.041, 0.099, 1.10	0.037, 0.105, 1.05
No. of reflections	3863	3685
No. of parameters	253	253
No. of restraints	0	1
H-atom treatment	H-atom parameters constrained	H-atom parameters constrained
$\Delta\rho_{max}$, $\Delta\rho_{min}$ (e Å ⁻³)	0.42, -0.58	0.19, -0.33

Computer programs: APEX2 and SAINT (Bruker, 2004), SHELXS97, SHELXL97 and SHELXL2014 (Sheldrick, 2008) and PLATON (Spek, 2009).

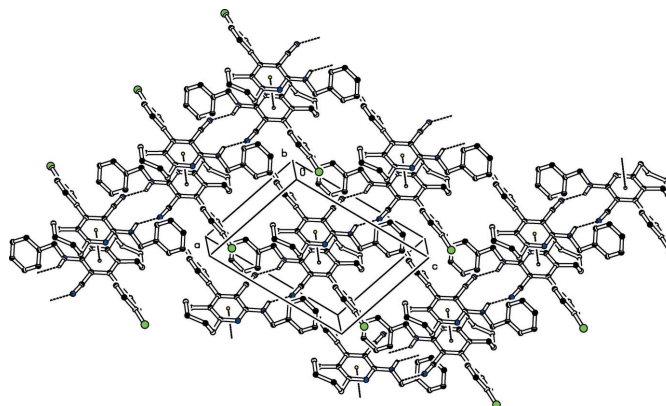


Figure 4

Crystal packing diagram of compound (II), viewed along the *b* axis. Hydrogen bonds (see Table 2 for details) and π – π interactions are shown as dashed lines (centroids are shown as small circles). H atoms not involved in hydrogen bonding have been omitted for clarity.

5. Database survey

A similar structure reported in the literature, 2-(4-bromophenyl)-4-(4-methoxyphenyl)-6,7,8,9-tetrahydro-5*H*-cyclohepta[*b*]pyridine (Çelik *et al.*, 2013) also has a chair conformation of the cycloheptane ring and a planar conformation of the pyridine ring, as found for (I) and (II). In compounds (I) and (II) the C–N bond lengths in the –CH₂–NH₂– chain, *viz.* C6–N2 and C21–N2, are 1.350 (3) and 1.441 (3) Å, respectively, in (I) and 1.354 (2) and 1.442 (2) Å, respectively, in (II). These distances are similar to those reported for

N^6 -(4-fluorobenzyl)-3-nitropyridine-2,6-diamine (Ge & Qian, 2011), viz. 1.341 (3) and 1.454 (3) Å, respectively.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. The NH and C-bound H atoms were placed in calculated positions and allowed to ride on their carrier atoms: N–H = 0.86 Å and C–H = 0.93–0.97 Å with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and $= 1.2U_{\text{eq}}(\text{N,C})$ for other H atoms.

Acknowledgements

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supporting information

Acta Cryst. (2015). E71, 12-15 [https://doi.org/10.1107/S2056989014025936]

Crystal structures of 2-benzylamino-4-(4-bromophenyl)-6,7,8,9-tetrahydro-5H-cyclohepta[b]pyridine-3-carbonitrile and 2-benzylamino-4-(4-chlorophenyl)-6,7,8,9-tetrahydro-5H-cyclohepta[b]pyridine-3-carbonitrile

R. A. Nagalakshmi, J. Suresh, S. Maharani, R. Ranjith Kumar and P. L. Nilantha Lakshman

Computing details

For both compounds, data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT* (Bruker, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2014* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009). Software used to prepare material for publication: *SHELXL2014* (Sheldrick, 2008) for (I); *SHELXL97* (Sheldrick, 2008) for (II).

(I) 2-Benzylamino-4-(4-bromophenyl)-6,7,8,9-tetrahydro-5H-cyclohepta[b]pyridine-3-carbonitrile

Crystal data

$C_{24}H_{22}BrN_3$	$F(000) = 888$
$M_r = 432.35$	$D_x = 1.383 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
$a = 8.9710 (3) \text{ \AA}$	Cell parameters from 2000 reflections
$b = 9.3794 (4) \text{ \AA}$	$\theta = 2-31^\circ$
$c = 24.9788 (9) \text{ \AA}$	$\mu = 1.99 \text{ mm}^{-1}$
$\beta = 99.002 (2)^\circ$	$T = 293 \text{ K}$
$V = 2075.89 (14) \text{ \AA}^3$	Block, colourless
$Z = 4$	$0.21 \times 0.19 \times 0.18 \text{ mm}$

Data collection

Bruker Kappa APEXII diffractometer	3863 independent reflections
Radiation source: fine-focus sealed tube	2927 reflections with $I > 2\sigma(I)$
ω and ϕ scans	$R_{\text{int}} = 0.040$
Absorption correction: multi-scan (SADABS; Bruker, 2004)	$\theta_{\text{max}} = 25.5^\circ$, $\theta_{\text{min}} = 2.3^\circ$
$T_{\text{min}} = 0.967$, $T_{\text{max}} = 0.974$	$h = -10 \rightarrow 10$
51599 measured reflections	$k = -11 \rightarrow 11$
	$l = -30 \rightarrow 30$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.041$	$w = 1/[\sigma^2(F_o^2) + (0.0313P)^2 + 2.0872P]$
$wR(F^2) = 0.099$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.10$	$(\Delta/\sigma)_{\text{max}} < 0.001$
3863 reflections	$\Delta\rho_{\text{max}} = 0.42 \text{ e \AA}^{-3}$
253 parameters	$\Delta\rho_{\text{min}} = -0.58 \text{ e \AA}^{-3}$
0 restraints	

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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.1307 (3)	0.3845 (3)	−0.03963 (10)	0.0396 (6)
C2	0.2856 (3)	0.3506 (3)	−0.02221 (10)	0.0328 (6)
C3	0.3766 (3)	0.3123 (3)	−0.06035 (10)	0.0315 (5)
C4	0.5237 (3)	0.2684 (3)	−0.04248 (10)	0.0340 (6)
C5	0.5720 (3)	0.2670 (3)	0.01347 (11)	0.0357 (6)
C6	0.3445 (3)	0.3496 (3)	0.03339 (10)	0.0344 (6)
C7	0.6278 (3)	0.2236 (3)	−0.08138 (11)	0.0440 (7)
H7A	0.7181	0.2819	−0.0748	0.053*
H7B	0.5783	0.2430	−0.1180	0.053*
C8	0.6743 (3)	0.0676 (3)	−0.07771 (12)	0.0515 (8)
H8A	0.5864	0.0101	−0.0742	0.062*
H8B	0.7083	0.0404	−0.1113	0.062*
C9	0.7981 (3)	0.0338 (3)	−0.03080 (12)	0.0519 (8)
H9A	0.8247	−0.0660	−0.0330	0.062*
H9B	0.8866	0.0894	−0.0351	0.062*
C10	0.7601 (3)	0.0622 (3)	0.02499 (12)	0.0508 (8)
H10A	0.8436	0.0298	0.0517	0.061*
H10B	0.6724	0.0058	0.0296	0.061*
C11	0.7278 (3)	0.2176 (3)	0.03676 (12)	0.0470 (7)
H11A	0.7424	0.2313	0.0757	0.056*
H11B	0.8007	0.2772	0.0224	0.056*
C21	0.3110 (3)	0.3992 (3)	0.12794 (11)	0.0481 (7)
H21A	0.3218	0.4992	0.1377	0.058*
H21B	0.4101	0.3557	0.1358	0.058*
C22	0.2093 (3)	0.3296 (3)	0.16277 (11)	0.0399 (6)
C23	0.2260 (4)	0.3649 (4)	0.21649 (13)	0.0708 (10)
H23	0.2957	0.4342	0.2302	0.085*
C24	0.1401 (5)	0.2983 (5)	0.25061 (15)	0.0878 (13)
H24	0.1530	0.3231	0.2871	0.105*
C25	0.0380 (5)	0.1977 (5)	0.23151 (18)	0.0812 (12)
H25	−0.0183	0.1521	0.2547	0.097*
C26	0.0187 (5)	0.1640 (5)	0.17855 (19)	0.0843 (12)
H26	−0.0527	0.0959	0.1650	0.101*
C27	0.1039 (4)	0.2295 (4)	0.14421 (14)	0.0629 (9)
H27	0.0890	0.2049	0.1077	0.075*
C31	0.3078 (3)	0.3224 (3)	−0.11850 (10)	0.0320 (6)
C32	0.2628 (4)	0.2034 (3)	−0.14877 (11)	0.0512 (8)
H32	0.2815	0.1134	−0.1336	0.061*
C33	0.1904 (4)	0.2159 (3)	−0.20130 (12)	0.0564 (8)

H33	0.1598	0.1348	−0.2215	0.068*
C34	0.1638 (3)	0.3475 (3)	−0.22362 (10)	0.0427 (7)
C35	0.2063 (3)	0.4684 (3)	−0.19455 (11)	0.0473 (7)
H35	0.1864	0.5580	−0.2099	0.057*
C36	0.2793 (3)	0.4546 (3)	−0.14193 (11)	0.0420 (6)
H36	0.3099	0.5360	−0.1219	0.050*
N1	0.0060 (3)	0.4098 (3)	−0.05266 (10)	0.0611 (8)
N2	0.2581 (3)	0.3886 (3)	0.07055 (9)	0.0475 (6)
H2	0.1649	0.4085	0.0592	0.057*
N3	0.4865 (2)	0.3065 (2)	0.05039 (8)	0.0364 (5)
Br1	0.07272 (5)	0.36307 (5)	−0.29690 (2)	0.07749 (17)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0387 (16)	0.0493 (17)	0.0303 (13)	0.0095 (13)	0.0039 (11)	−0.0035 (12)
C2	0.0278 (12)	0.0349 (14)	0.0347 (13)	0.0055 (11)	0.0013 (10)	0.0019 (11)
C3	0.0329 (13)	0.0272 (13)	0.0341 (13)	0.0021 (10)	0.0047 (10)	−0.0001 (10)
C4	0.0289 (13)	0.0318 (13)	0.0414 (14)	0.0030 (11)	0.0058 (11)	−0.0005 (11)
C5	0.0287 (13)	0.0345 (14)	0.0424 (14)	0.0036 (11)	0.0010 (11)	−0.0025 (11)
C6	0.0329 (13)	0.0355 (14)	0.0345 (13)	0.0051 (11)	0.0041 (10)	0.0025 (11)
C7	0.0355 (14)	0.0557 (18)	0.0424 (15)	0.0074 (13)	0.0105 (12)	0.0055 (13)
C8	0.0460 (17)	0.0580 (19)	0.0516 (17)	0.0139 (15)	0.0106 (14)	−0.0072 (15)
C9	0.0439 (17)	0.0483 (18)	0.0636 (19)	0.0153 (14)	0.0085 (14)	−0.0014 (15)
C10	0.0396 (16)	0.0541 (19)	0.0566 (18)	0.0157 (14)	0.0010 (13)	0.0057 (15)
C11	0.0321 (14)	0.0582 (19)	0.0474 (16)	0.0094 (13)	−0.0036 (12)	−0.0064 (14)
C21	0.0487 (17)	0.0607 (19)	0.0340 (14)	0.0015 (14)	0.0035 (12)	0.0033 (13)
C22	0.0400 (15)	0.0438 (16)	0.0360 (14)	0.0105 (12)	0.0066 (11)	0.0013 (12)
C23	0.086 (3)	0.087 (3)	0.0424 (18)	−0.019 (2)	0.0176 (17)	−0.0114 (18)
C24	0.106 (3)	0.118 (4)	0.047 (2)	0.006 (3)	0.037 (2)	0.002 (2)
C25	0.068 (3)	0.095 (3)	0.087 (3)	0.008 (2)	0.035 (2)	0.033 (3)
C26	0.077 (3)	0.087 (3)	0.090 (3)	−0.023 (2)	0.019 (2)	0.012 (2)
C27	0.068 (2)	0.071 (2)	0.0505 (19)	−0.0095 (19)	0.0094 (16)	−0.0065 (17)
C31	0.0276 (13)	0.0357 (14)	0.0334 (13)	0.0045 (11)	0.0071 (10)	−0.0002 (11)
C32	0.078 (2)	0.0347 (15)	0.0397 (15)	−0.0015 (15)	0.0057 (15)	0.0002 (13)
C33	0.084 (2)	0.0457 (19)	0.0381 (16)	−0.0168 (17)	0.0037 (15)	−0.0081 (14)
C34	0.0402 (15)	0.0544 (18)	0.0320 (13)	−0.0076 (13)	0.0006 (11)	0.0001 (13)
C35	0.0580 (18)	0.0404 (16)	0.0399 (15)	0.0015 (14)	−0.0037 (13)	0.0031 (13)
C36	0.0502 (16)	0.0357 (15)	0.0378 (14)	0.0007 (13)	−0.0002 (12)	−0.0044 (12)
N1	0.0367 (14)	0.094 (2)	0.0506 (15)	0.0198 (14)	−0.0004 (11)	−0.0073 (14)
N2	0.0377 (12)	0.0729 (18)	0.0316 (11)	0.0193 (12)	0.0044 (10)	0.0047 (11)
N3	0.0324 (11)	0.0394 (12)	0.0352 (11)	0.0067 (10)	−0.0012 (9)	0.0002 (10)
Br1	0.0937 (3)	0.0883 (3)	0.04077 (19)	−0.0216 (2)	−0.01969 (17)	0.00315 (18)

Geometric parameters (Å, °)

C1—N1	1.140 (3)	C21—C22	1.505 (4)
C1—C2	1.426 (4)	C21—H21A	0.9700

C2—C3	1.395 (3)	C21—H21B	0.9700
C2—C6	1.406 (3)	C22—C27	1.361 (4)
C3—C4	1.388 (3)	C22—C23	1.367 (4)
C3—C31	1.490 (3)	C23—C24	1.384 (5)
C4—C5	1.397 (4)	C23—H23	0.9300
C4—C7	1.508 (4)	C24—C25	1.349 (6)
C5—N3	1.341 (3)	C24—H24	0.9300
C5—C11	1.501 (3)	C25—C26	1.345 (6)
C6—N3	1.340 (3)	C25—H25	0.9300
C6—N2	1.350 (3)	C26—C27	1.379 (5)
C7—C8	1.521 (4)	C26—H26	0.9300
C7—H7A	0.9700	C27—H27	0.9300
C7—H7B	0.9700	C31—C32	1.372 (4)
C8—C9	1.517 (4)	C31—C36	1.378 (4)
C8—H8A	0.9700	C32—C33	1.375 (4)
C8—H8B	0.9700	C32—H32	0.9300
C9—C10	1.509 (4)	C33—C34	1.359 (4)
C9—H9A	0.9700	C33—H33	0.9300
C9—H9B	0.9700	C34—C35	1.368 (4)
C10—C11	1.524 (4)	C34—Br1	1.890 (3)
C10—H10A	0.9700	C35—C36	1.380 (4)
C10—H10B	0.9700	C35—H35	0.9300
C11—H11A	0.9700	C36—H36	0.9300
C11—H11B	0.9700	N2—H2	0.8600
C21—N2	1.441 (3)		
N1—C1—C2	178.5 (3)	N2—C21—C22	114.2 (2)
C3—C2—C6	120.2 (2)	N2—C21—H21A	108.7
C3—C2—C1	119.7 (2)	C22—C21—H21A	108.7
C6—C2—C1	120.0 (2)	N2—C21—H21B	108.7
C4—C3—C2	119.0 (2)	C22—C21—H21B	108.7
C4—C3—C31	124.1 (2)	H21A—C21—H21B	107.6
C2—C3—C31	116.8 (2)	C27—C22—C23	117.7 (3)
C3—C4—C5	117.0 (2)	C27—C22—C21	123.6 (3)
C3—C4—C7	121.9 (2)	C23—C22—C21	118.6 (3)
C5—C4—C7	121.1 (2)	C22—C23—C24	120.6 (4)
N3—C5—C4	124.4 (2)	C22—C23—H23	119.7
N3—C5—C11	114.6 (2)	C24—C23—H23	119.7
C4—C5—C11	121.0 (2)	C25—C24—C23	120.7 (4)
N3—C6—N2	118.9 (2)	C25—C24—H24	119.7
N3—C6—C2	120.4 (2)	C23—C24—H24	119.7
N2—C6—C2	120.6 (2)	C26—C25—C24	119.2 (4)
C4—C7—C8	114.9 (2)	C26—C25—H25	120.4
C4—C7—H7A	108.6	C24—C25—H25	120.4
C8—C7—H7A	108.6	C25—C26—C27	120.6 (4)
C4—C7—H7B	108.6	C25—C26—H26	119.7
C8—C7—H7B	108.6	C27—C26—H26	119.7
H7A—C7—H7B	107.5	C22—C27—C26	121.2 (3)

C9—C8—C7	114.1 (3)	C22—C27—H27	119.4
C9—C8—H8A	108.7	C26—C27—H27	119.4
C7—C8—H8A	108.7	C32—C31—C36	118.6 (2)
C9—C8—H8B	108.7	C32—C31—C3	121.8 (2)
C7—C8—H8B	108.7	C36—C31—C3	119.5 (2)
H8A—C8—H8B	107.6	C31—C32—C33	120.7 (3)
C10—C9—C8	115.6 (2)	C31—C32—H32	119.7
C10—C9—H9A	108.4	C33—C32—H32	119.7
C8—C9—H9A	108.4	C34—C33—C32	119.7 (3)
C10—C9—H9B	108.4	C34—C33—H33	120.2
C8—C9—H9B	108.4	C32—C33—H33	120.2
H9A—C9—H9B	107.4	C33—C34—C35	121.2 (3)
C9—C10—C11	115.1 (3)	C33—C34—Br1	119.3 (2)
C9—C10—H10A	108.5	C35—C34—Br1	119.5 (2)
C11—C10—H10A	108.5	C34—C35—C36	118.6 (3)
C9—C10—H10B	108.5	C34—C35—H35	120.7
C11—C10—H10B	108.5	C36—C35—H35	120.7
H10A—C10—H10B	107.5	C31—C36—C35	121.2 (3)
C5—C11—C10	114.5 (2)	C31—C36—H36	119.4
C5—C11—H11A	108.6	C35—C36—H36	119.4
C10—C11—H11A	108.6	C6—N2—C21	124.6 (2)
C5—C11—H11B	108.6	C6—N2—H2	117.7
C10—C11—H11B	108.6	C21—N2—H2	117.7
H11A—C11—H11B	107.6	C6—N3—C5	118.9 (2)
C6—C2—C3—C4	2.3 (4)	C22—C23—C24—C25	−0.2 (7)
C1—C2—C3—C4	−174.8 (2)	C23—C24—C25—C26	−1.0 (7)
C6—C2—C3—C31	−177.4 (2)	C24—C25—C26—C27	1.1 (7)
C1—C2—C3—C31	5.5 (4)	C23—C22—C27—C26	−1.1 (5)
C2—C3—C4—C5	−0.7 (4)	C21—C22—C27—C26	176.7 (3)
C31—C3—C4—C5	179.0 (2)	C25—C26—C27—C22	0.0 (6)
C2—C3—C4—C7	179.2 (2)	C4—C3—C31—C32	75.4 (4)
C31—C3—C4—C7	−1.1 (4)	C2—C3—C31—C32	−104.9 (3)
C3—C4—C5—N3	−0.7 (4)	C4—C3—C31—C36	−108.7 (3)
C7—C4—C5—N3	179.4 (3)	C2—C3—C31—C36	71.0 (3)
C3—C4—C5—C11	178.4 (2)	C36—C31—C32—C33	−0.2 (4)
C7—C4—C5—C11	−1.5 (4)	C3—C31—C32—C33	175.6 (3)
C3—C2—C6—N3	−2.6 (4)	C31—C32—C33—C34	0.4 (5)
C1—C2—C6—N3	174.5 (2)	C32—C33—C34—C35	−0.7 (5)
C3—C2—C6—N2	179.0 (2)	C32—C33—C34—Br1	177.1 (2)
C1—C2—C6—N2	−3.9 (4)	C33—C34—C35—C36	0.9 (5)
C3—C4—C7—C8	−114.6 (3)	Br1—C34—C35—C36	−176.9 (2)
C5—C4—C7—C8	65.3 (3)	C32—C31—C36—C35	0.5 (4)
C4—C7—C8—C9	−78.9 (3)	C3—C31—C36—C35	−175.5 (2)
C7—C8—C9—C10	61.7 (4)	C34—C35—C36—C31	−0.8 (4)
C8—C9—C10—C11	−62.9 (4)	N3—C6—N2—C21	5.5 (4)
N3—C5—C11—C10	116.6 (3)	C2—C6—N2—C21	−176.0 (3)
C4—C5—C11—C10	−62.6 (4)	C22—C21—N2—C6	−133.1 (3)

C9—C10—C11—C5	79.7 (3)	N2—C6—N3—C5	179.6 (2)
N2—C21—C22—C27	19.3 (4)	C2—C6—N3—C5	1.2 (4)
N2—C21—C22—C23	−162.8 (3)	C4—C5—N3—C6	0.5 (4)
C27—C22—C23—C24	1.2 (5)	C11—C5—N3—C6	−178.7 (2)
C21—C22—C23—C24	−176.7 (4)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N2—H2 \cdots N1 ⁱ	0.86	2.28	3.010 (3)	143
C21—H21B \cdots Br1 ⁱⁱ	0.97	2.90	3.703 (3)	141

Symmetry codes: (i) $-x, -y+1, -z$; (ii) $x+1/2, -y+1/2, z+1/2$.**(II) 2-Benzylamino-4-(4-chlorophenyl)-6,7,8,9-tetrahydro-5H-cyclohepta[b]pyridine-3-carbonitrile***Crystal data*C₂₄H₂₂ClN₃ $M_r = 387.89$ Triclinic, $P\bar{1}$ $a = 9.002$ (5) Å $b = 10.097$ (5) Å $c = 11.856$ (5) Å $\alpha = 94.939$ (5)° $\beta = 108.204$ (5)° $\gamma = 101.272$ (5)° $V = 991.3$ (8) Å³ $Z = 2$ $F(000) = 408$ $D_x = 1.299$ Mg m^{−3}Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 2000 reflections

 $\theta = 2\text{--}31^\circ$ $\mu = 0.21$ mm^{−1} $T = 293$ K

Block, colourless

 $0.21 \times 0.19 \times 0.18$ mm*Data collection*

Bruker Kappa APEXII

diffractometer

Radiation source: fine-focus sealed tube

 ω and ϕ scans

Absorption correction: multi-scan

(SADABS; Bruker, 2004)

 $T_{\min} = 0.967$, $T_{\max} = 0.974$

24808 measured reflections

3685 independent reflections

2918 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.026$ $\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 2.1^\circ$ $h = -10 \rightarrow 10$ $k = -12 \rightarrow 12$ $l = -14 \rightarrow 14$ *Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.105$ $S = 1.05$

3685 reflections

253 parameters

1 restraint

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0486P)^2 + 0.3383P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.19$ e Å^{−3} $\Delta\rho_{\min} = -0.33$ e Å^{−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.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.1170 (2)	0.47342 (18)	0.36822 (15)	0.0401 (4)
C2	0.24864 (18)	0.41014 (15)	0.38041 (14)	0.0327 (3)
C3	0.32569 (17)	0.40885 (15)	0.29430 (13)	0.0304 (3)
C4	0.44508 (17)	0.33539 (15)	0.30821 (14)	0.0321 (3)
C5	0.48147 (17)	0.26963 (15)	0.40915 (14)	0.0331 (3)
C6	0.29611 (17)	0.34315 (15)	0.48044 (14)	0.0320 (3)
C7	0.53288 (19)	0.32240 (17)	0.21988 (15)	0.0395 (4)
H7A	0.6478	0.3489	0.2631	0.047*
H7B	0.5074	0.3854	0.1633	0.047*
C8	0.4910 (2)	0.17808 (19)	0.14983 (16)	0.0477 (4)
H8A	0.3760	0.1411	0.1274	0.057*
H8B	0.5158	0.1839	0.0762	0.057*
C9	0.5782 (3)	0.0790 (2)	0.21741 (18)	0.0540 (5)
H9A	0.5446	−0.0086	0.1652	0.065*
H9B	0.6926	0.1124	0.2336	0.065*
C10	0.5522 (2)	0.05584 (19)	0.33498 (18)	0.0516 (5)
H10A	0.6113	−0.0104	0.3685	0.062*
H10B	0.4389	0.0163	0.3184	0.062*
C11	0.6037 (2)	0.18367 (19)	0.42923 (16)	0.0451 (4)
H11A	0.6223	0.1565	0.5081	0.054*
H11B	0.7046	0.2386	0.4285	0.054*
C21	0.2814 (2)	0.29636 (18)	0.67846 (15)	0.0416 (4)
H21A	0.3240	0.3740	0.7428	0.050*
H21B	0.3692	0.2540	0.6781	0.050*
C22	0.15556 (19)	0.19495 (16)	0.70552 (14)	0.0347 (3)
C23	0.1853 (2)	0.16979 (18)	0.82224 (16)	0.0449 (4)
H23	0.2789	0.2189	0.8823	0.054*
C24	0.0778 (3)	0.0727 (2)	0.85090 (19)	0.0589 (5)
H24	0.0996	0.0564	0.9298	0.071*
C25	−0.0605 (3)	0.0004 (2)	0.7635 (2)	0.0629 (6)
H25	−0.1323	−0.0658	0.7826	0.075*
C26	−0.0929 (2)	0.0256 (2)	0.6480 (2)	0.0617 (5)
H26	−0.1877	−0.0228	0.5886	0.074*
C27	0.0142 (2)	0.12280 (19)	0.61870 (17)	0.0503 (4)
H27	−0.0092	0.1396	0.5399	0.060*
C31	0.28089 (17)	0.49110 (15)	0.19592 (14)	0.0321 (3)
C32	0.2876 (2)	0.62890 (17)	0.22518 (15)	0.0396 (4)
H32	0.3223	0.6683	0.3056	0.047*
C33	0.2437 (2)	0.70820 (18)	0.13681 (17)	0.0459 (4)
H33	0.2491	0.8004	0.1574	0.055*
C34	0.19207 (19)	0.64947 (18)	0.01821 (16)	0.0428 (4)
C35	0.1840 (2)	0.51340 (19)	−0.01437 (16)	0.0454 (4)
H35	0.1493	0.4748	−0.0950	0.055*
C36	0.2284 (2)	0.43515 (17)	0.07516 (15)	0.0395 (4)
H36	0.2231	0.3430	0.0540	0.047*

N1	0.0094 (2)	0.51850 (19)	0.36511 (16)	0.0616 (5)
N2	0.22527 (17)	0.34558 (15)	0.56572 (13)	0.0425 (3)
H2	0.1413	0.3786	0.5517	0.051*
N3	0.41191 (15)	0.27423 (13)	0.49346 (11)	0.0349 (3)
Cl1	0.13087 (7)	0.74901 (6)	−0.09170 (5)	0.06667 (19)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0457 (9)	0.0481 (10)	0.0398 (9)	0.0242 (8)	0.0226 (7)	0.0144 (7)
C2	0.0336 (7)	0.0323 (8)	0.0369 (8)	0.0132 (6)	0.0151 (6)	0.0062 (6)
C3	0.0315 (7)	0.0271 (8)	0.0337 (8)	0.0084 (6)	0.0118 (6)	0.0038 (6)
C4	0.0294 (7)	0.0317 (8)	0.0382 (8)	0.0096 (6)	0.0141 (6)	0.0050 (6)
C5	0.0307 (7)	0.0328 (8)	0.0362 (8)	0.0110 (6)	0.0104 (6)	0.0028 (6)
C6	0.0328 (8)	0.0297 (8)	0.0369 (8)	0.0093 (6)	0.0153 (6)	0.0047 (6)
C7	0.0368 (8)	0.0452 (9)	0.0472 (9)	0.0170 (7)	0.0224 (7)	0.0144 (8)
C8	0.0523 (10)	0.0567 (11)	0.0441 (10)	0.0229 (9)	0.0246 (8)	0.0062 (8)
C9	0.0662 (12)	0.0486 (11)	0.0619 (12)	0.0279 (9)	0.0335 (10)	0.0071 (9)
C10	0.0646 (12)	0.0436 (10)	0.0629 (12)	0.0302 (9)	0.0318 (10)	0.0148 (9)
C11	0.0454 (9)	0.0549 (11)	0.0453 (10)	0.0294 (8)	0.0172 (8)	0.0146 (8)
C21	0.0435 (9)	0.0471 (10)	0.0368 (9)	0.0101 (8)	0.0168 (7)	0.0088 (7)
C22	0.0398 (8)	0.0331 (8)	0.0381 (8)	0.0154 (7)	0.0184 (7)	0.0066 (6)
C23	0.0513 (10)	0.0473 (10)	0.0398 (9)	0.0150 (8)	0.0171 (8)	0.0111 (8)
C24	0.0745 (14)	0.0608 (13)	0.0570 (12)	0.0228 (11)	0.0345 (11)	0.0288 (10)
C25	0.0589 (12)	0.0536 (12)	0.0923 (16)	0.0156 (10)	0.0404 (12)	0.0339 (12)
C26	0.0479 (11)	0.0535 (12)	0.0764 (14)	0.0035 (9)	0.0144 (10)	0.0156 (11)
C27	0.0498 (10)	0.0527 (11)	0.0455 (10)	0.0092 (9)	0.0127 (8)	0.0119 (8)
C31	0.0298 (7)	0.0341 (8)	0.0386 (8)	0.0116 (6)	0.0167 (6)	0.0094 (6)
C32	0.0413 (9)	0.0368 (9)	0.0408 (9)	0.0121 (7)	0.0123 (7)	0.0060 (7)
C33	0.0460 (9)	0.0332 (9)	0.0594 (11)	0.0111 (7)	0.0164 (8)	0.0136 (8)
C34	0.0384 (9)	0.0483 (10)	0.0500 (10)	0.0146 (8)	0.0195 (8)	0.0240 (8)
C35	0.0499 (10)	0.0558 (11)	0.0370 (9)	0.0171 (8)	0.0196 (8)	0.0115 (8)
C36	0.0457 (9)	0.0380 (9)	0.0416 (9)	0.0158 (7)	0.0206 (7)	0.0072 (7)
N1	0.0655 (10)	0.0839 (13)	0.0638 (11)	0.0493 (10)	0.0377 (9)	0.0285 (9)
N2	0.0472 (8)	0.0511 (9)	0.0466 (8)	0.0259 (7)	0.0281 (7)	0.0202 (7)
N3	0.0366 (7)	0.0351 (7)	0.0366 (7)	0.0145 (6)	0.0135 (6)	0.0072 (6)
Cl1	0.0687 (3)	0.0744 (4)	0.0699 (3)	0.0251 (3)	0.0271 (3)	0.0466 (3)

Geometric parameters (\AA , $^\circ$)

C1—N1	1.140 (2)	C21—C22	1.505 (2)
C1—C2	1.428 (2)	C21—H21A	0.9700
C2—C3	1.403 (2)	C21—H21B	0.9700
C2—C6	1.408 (2)	C22—C27	1.378 (3)
C3—C4	1.398 (2)	C22—C23	1.381 (2)
C3—C31	1.489 (2)	C23—C24	1.380 (3)
C4—C5	1.399 (2)	C23—H23	0.9300
C4—C7	1.508 (2)	C24—C25	1.366 (3)

C5—N3	1.337 (2)	C24—H24	0.9300
C5—C11	1.505 (2)	C25—C26	1.366 (3)
C6—N3	1.340 (2)	C25—H25	0.9300
C6—N2	1.354 (2)	C26—C27	1.382 (3)
C7—C8	1.529 (3)	C26—H26	0.9300
C7—H7A	0.9700	C27—H27	0.9300
C7—H7B	0.9700	C31—C36	1.388 (2)
C8—C9	1.520 (3)	C31—C32	1.389 (2)
C8—H8A	0.9700	C32—C33	1.380 (2)
C8—H8B	0.9700	C32—H32	0.9300
C9—C10	1.513 (3)	C33—C34	1.373 (3)
C9—H9A	0.9700	C33—H33	0.9300
C9—H9B	0.9700	C34—C35	1.376 (3)
C10—C11	1.524 (3)	C34—C11	1.7334 (17)
C10—H10A	0.9700	C35—C36	1.383 (2)
C10—H10B	0.9700	C35—H35	0.9300
C11—H11A	0.9700	C36—H36	0.9300
C11—H11B	0.9700	N2—H2	0.8600
C21—N2	1.442 (2)		
N1—C1—C2	174.73 (18)	N2—C21—C22	114.79 (14)
C3—C2—C6	120.15 (13)	N2—C21—H21A	108.6
C3—C2—C1	122.07 (14)	C22—C21—H21A	108.6
C6—C2—C1	117.73 (14)	N2—C21—H21B	108.6
C4—C3—C2	118.39 (14)	C22—C21—H21B	108.6
C4—C3—C31	123.49 (13)	H21A—C21—H21B	107.5
C2—C3—C31	118.06 (13)	C27—C22—C23	118.37 (16)
C3—C4—C5	117.26 (13)	C27—C22—C21	123.14 (15)
C3—C4—C7	123.47 (14)	C23—C22—C21	118.46 (15)
C5—C4—C7	119.26 (13)	C24—C23—C22	120.88 (18)
N3—C5—C4	124.52 (14)	C24—C23—H23	119.6
N3—C5—C11	114.38 (14)	C22—C23—H23	119.6
C4—C5—C11	121.08 (14)	C25—C24—C23	120.05 (19)
N3—C6—N2	118.13 (14)	C25—C24—H24	120.0
N3—C6—C2	120.89 (13)	C23—C24—H24	120.0
N2—C6—C2	120.98 (13)	C24—C25—C26	119.80 (19)
C4—C7—C8	113.51 (14)	C24—C25—H25	120.1
C4—C7—H7A	108.9	C26—C25—H25	120.1
C8—C7—H7A	108.9	C25—C26—C27	120.40 (19)
C4—C7—H7B	108.9	C25—C26—H26	119.8
C8—C7—H7B	108.9	C27—C26—H26	119.8
H7A—C7—H7B	107.7	C22—C27—C26	120.48 (18)
C9—C8—C7	114.77 (15)	C22—C27—H27	119.8
C9—C8—H8A	108.6	C26—C27—H27	119.8
C7—C8—H8A	108.6	C36—C31—C32	118.14 (14)
C9—C8—H8B	108.6	C36—C31—C3	122.71 (14)
C7—C8—H8B	108.6	C32—C31—C3	119.13 (14)
H8A—C8—H8B	107.6	C33—C32—C31	121.05 (16)

C10—C9—C8	116.06 (15)	C33—C32—H32	119.5
C10—C9—H9A	108.3	C31—C32—H32	119.5
C8—C9—H9A	108.3	C34—C33—C32	119.28 (16)
C10—C9—H9B	108.3	C34—C33—H33	120.4
C8—C9—H9B	108.3	C32—C33—H33	120.4
H9A—C9—H9B	107.4	C33—C34—C35	121.39 (16)
C9—C10—C11	115.01 (16)	C33—C34—C11	118.72 (14)
C9—C10—H10A	108.5	C35—C34—C11	119.86 (14)
C11—C10—H10A	108.5	C34—C35—C36	118.70 (16)
C9—C10—H10B	108.5	C34—C35—H35	120.6
C11—C10—H10B	108.5	C36—C35—H35	120.6
H10A—C10—H10B	107.5	C35—C36—C31	121.43 (16)
C5—C11—C10	113.24 (15)	C35—C36—H36	119.3
C5—C11—H11A	108.9	C31—C36—H36	119.3
C10—C11—H11A	108.9	C6—N2—C21	124.26 (14)
C5—C11—H11B	108.9	C6—N2—H2	117.9
C10—C11—H11B	108.9	C21—N2—H2	117.9
H11A—C11—H11B	107.7	C5—N3—C6	118.73 (13)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N2—H2 \cdots N1 ⁱ	0.86	2.26	3.007 (2)	145

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