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2-Benzoyl-1,1-diethyl-3-phenylguanidine

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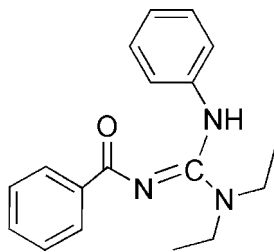
 Key indicators: single-crystal X-ray study; $T = 113$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å;

 R factor = 0.068; wR factor = 0.112; data-to-parameter ratio = 17.3.

In the title tetrasubstituted guanidine, $\text{C}_{18}\text{H}_{21}\text{N}_3\text{O}$, the guanidine and carbonyl groups are not coplanar, as reflected by the torsion angles involving the $\text{N}=\text{C}$ atoms [17.6 (3), -141.68 (17) and 42.2 (3) $^\circ$]. This is probably due to the absence of an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond, forming a six-membered ring, and is commonly observed in this class of compounds. In the crystal structure, centrosymmetric dimers are formed *via* pairs of intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds. The dihedral angles between the guanidine plane and the phenyl ring and benzoyl plane are 38.06 (9) and 41.54 (7) $^\circ$, respectively.

Related literature

For thiourea derivatives with biological activity, see: Berlinck (2002); Heys *et al.* (2000); Laeckmann *et al.* (2002); Kelley *et al.* (2001); Moroni *et al.* (2001); Ishikawa *et al.* (2002). For related structures, see: Murtaza *et al.* (2007, 2008); Cunha *et al.* (2005).



Experimental

Crystal data

 $\text{C}_{18}\text{H}_{21}\text{N}_3\text{O}$
 $M_r = 295.38$
 Monoclinic, $P2_1/c$
 $a = 10.472$ (6) Å
 $b = 15.010$ (8) Å

 $c = 10.154$ (6) Å
 $\beta = 102.992$ (6) $^\circ$
 $V = 1555.2$ (15) Å³
 $Z = 4$
 Mo $K\alpha$ radiation

 $\mu = 0.08$ mm⁻¹
 $T = 113$ (2) K

 $0.50 \times 0.40 \times 0.30$ mm

Data collection

 Rigaku/MSC Mercury CCD
 diffractometer
 Absorption correction: none
 12318 measured reflections

 3556 independent reflections
 3239 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.042$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.068$
 $wR(F^2) = 0.112$
 $S = 1.27$
 3556 reflections
 205 parameters

 H atoms treated by a mixture of
 independent and constrained
 refinement
 $\Delta\rho_{\text{max}} = 0.26$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.19$ e Å⁻³

Table 1

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N3}-\text{H3}\cdots\text{O1}^i$	0.90 (2)	1.97 (2)	2.852 (2)	168 (2)

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

Data collection: *CrystalClear* (Molecular Structure Corporation & Rigaku, 2001); cell refinement: *CrystalClear*; data reduction: *TEXSAN* (Molecular Structure Corporation & Rigaku, 2004); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97* and *TEXSAN*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2090).

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Acta Cryst. (2009). E65, o343 [doi:10.1107/S1600536809001469]

2-Benzoyl-1,1-diethyl-3-phenylguanidine

Ghulam Murtaza, Hanif-Ur-Rehman, M. Khawar Rauf, Masahiro Ebihara and Amin Badshah

S1. Comment

Guanidines are important compounds that have many biological, chemical and medicinal applications (Berlinck *et al.*, 2002; Heys *et al.*, 2000). They have received increasing interest as medicinal agents with antitumour, anti-hypertensive, anti-glaucoma and cardiotoxic activities (Laeckmann *et al.*, 2002; Kelley *et al.*, 2001; Moroni *et al.*, 2001). Due to their strongly basic character, they can be considered as super-bases that readily undergo protonation to generate resonance-stabilized guanidinium cations (Ishikawa *et al.*, 2002).

The molecular structure of the title compound, (I), is illustrated in Fig. 1. It is a typical *N',N,N,N'*-Tetrasubstituted guanidine with normal geometrical parameters (Murtaza *et al.*, 2007, 2008; Cunha *et al.*, 2005). The carbonyl bond (C2=O1) shows the expected full double bond character, while the shorter values for bonds C2—N1, N1—C1, C1—N2, and C1—N3 indicate partial double bond character. The dihedral angles between the guanidine mean plane (C1/N1/N2/N3), and the phenyl ring (C13—C18), the benzoyl ring (C3—C8,C2,O1) and the N2/C9/C11 plane, are 38.06 (9)°, 41.54 (7)°, and 11.97 (13)°, respectively. The guanidine moiety and the carbonyl group are not co-planar, as reflected by the torsion angles C1—N1—C2—O1 = 17.6 (3)°, N2—C1—N1—C2 = -141.68 (17)°, and N3—C1—N1—C2 = 42.2 (3)°. This is probably due to the absence of an intramolecular N—H···O hydrogen bond, forming a six-membered ring, and commonly observed in this class of compounds (Cunha *et al.*, 2005).

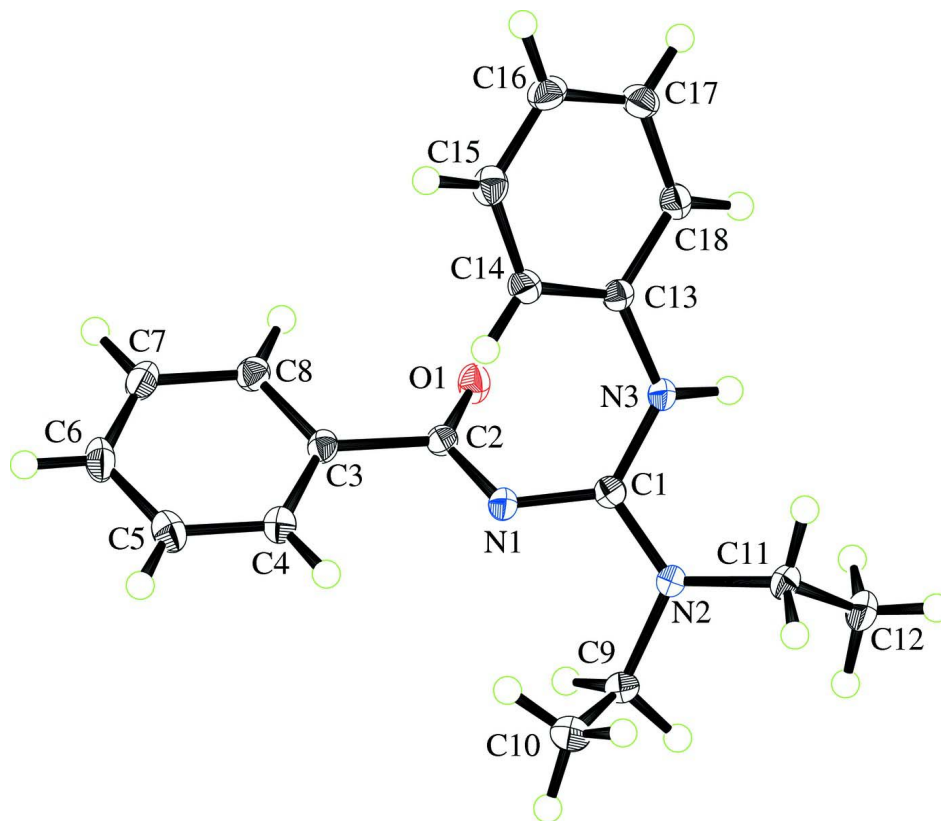
The crystal packing shows intermolecular N—H···O hydrogen bonds which result in the formation of centrosymmetric dimers (Fig. 2).

S2. Experimental

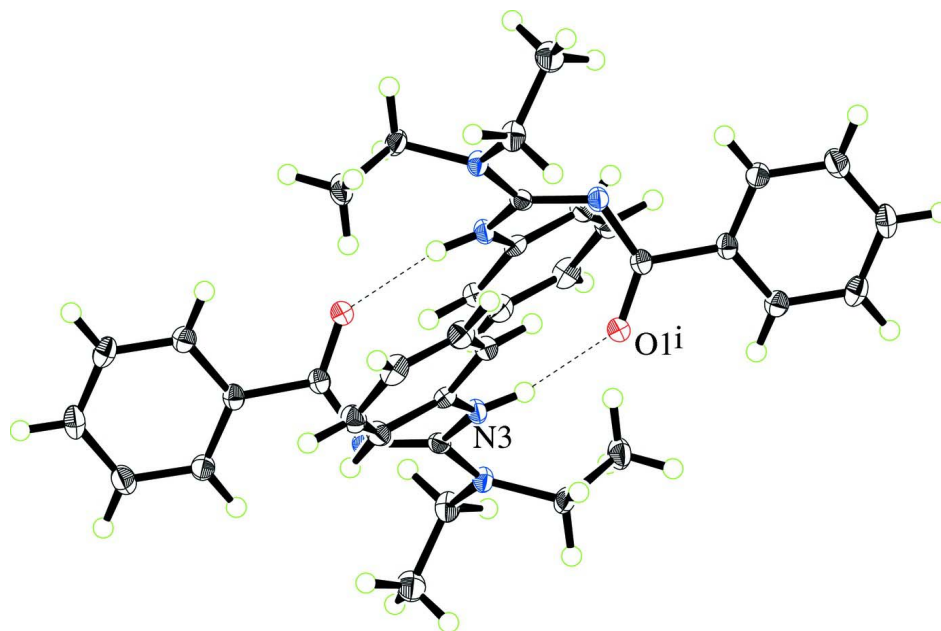
N-Benzoyl-*N'*-phenylthiourea (0.512 g, 2 mmol), triethyl amine (0.56 ml, 4 mmol) and diethyl amine (0.11 mL, 2 mmol) dissolved in 20 ml dimethylformamide, were mixed with vigorous stirring at 5°C. Mercuric chloride (0.544 g, 2 mmol) was then added and the mixture vigorously stirred for 12 h. The progress of the reaction was monitored by TLC. When all the thiourea had been consumed, 20 mL of chloroform were added and the suspension was filtered through a sintered glass funnel to remove any residue (HgS) formed as a byproduct during the reaction. The solvent was evaporated under reduced pressure and the residue was dissolved in 20 mL of CH₂Cl₂. Other byproducts were extracted out with water (4 × 30 mL). The organic phase was dried over anhydrous MgSO₄ and then filtered. After filtration the solvent was evaporated and compound (I) was recrystallized in ethanol. Full spectroscopic and physical characterization will be reported elsewhere.

S3. Refinement

The N-H hydrogen atom was located in a difference Fourier map and freely refined: N-H = 0.90 (2) Å. The C-bound H-atoms were included in calculated positions and treated as riding atoms: C—H = 0.95 - 0.98 Å with $U_{\text{iso}}(\text{H}) = 1.2$ or $1.5U_{\text{eq}}(\text{C})$.

**Figure 1**

Molecular structure of compound (I), showing the atom numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

**Figure 2**

A view of the centrosymmetric hydrogen-bonded dimer structure of compound (I) [Hydrogen bonds shown as dashed lines; symmetry code: (i) $-x, 1 - y, 1 - z$].

2-Benzoyl-1,1-diethyl-3-phenylguanidine

Crystal data

$C_{18}H_{21}N_3O$

$M_r = 295.38$

Monoclinic, $P2_1/c$

Hall symbol: $-P 2_1 ybc$

$a = 10.472$ (6) Å

$b = 15.010$ (8) Å

$c = 10.154$ (6) Å

$\beta = 102.992$ (6)°

$V = 1555.2$ (15) Å³

$Z = 4$

$F(000) = 632$

$D_x = 1.262$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71070$ Å

Cell parameters from 4060 reflections

$\theta = 6.3$ – 55.0 °

$\mu = 0.08$ mm⁻¹

$T = 113$ K

Block, colourless

$0.50 \times 0.40 \times 0.30$ mm

Data collection

Rigaku/MSC Mercury CCD
diffractometer

Graphite monochromator

Detector resolution: 14.62 pixels mm⁻¹

ω scans

12318 measured reflections

3556 independent reflections

3239 reflections with $I > 2\sigma(I)$

$R_{int} = 0.042$

$\theta_{max} = 27.5$ °, $\theta_{min} = 3.4$ °

$h = -13 \rightarrow 8$

$k = -19 \rightarrow 19$

$l = -13 \rightarrow 13$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.068$

$wR(F^2) = 0.112$

$S = 1.27$

3556 reflections

205 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

$$w = 1/[\sigma^2(F_o^2) + (0.0155P)^2 + 1.0509P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.26 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e } \text{Å}^{-3}$

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.

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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.21836 (16)	0.49684 (12)	0.59825 (17)	0.0158 (3)
N1	0.28409 (14)	0.54983 (10)	0.53093 (14)	0.0174 (3)
N2	0.24622 (14)	0.50194 (10)	0.73344 (14)	0.0172 (3)
N3	0.12913 (15)	0.43370 (10)	0.53944 (14)	0.0171 (3)
H3	0.063 (2)	0.4210 (15)	0.580 (2)	0.035 (6)*
C2	0.22449 (17)	0.58833 (11)	0.41264 (16)	0.0162 (3)
O1	0.10437 (12)	0.59798 (9)	0.36752 (12)	0.0208 (3)
C3	0.31721 (17)	0.62748 (11)	0.33424 (17)	0.0168 (4)
C4	0.45097 (18)	0.63592 (12)	0.39163 (18)	0.0201 (4)
H4	0.4854	0.6146	0.4806	0.024*
C5	0.53391 (19)	0.67529 (13)	0.31946 (19)	0.0248 (4)
H5	0.6247	0.6810	0.3593	0.030*
C6	0.4846 (2)	0.70640 (13)	0.18924 (19)	0.0249 (4)
H6	0.5414	0.7337	0.1402	0.030*
C7	0.35200 (19)	0.69752 (12)	0.13084 (18)	0.0228 (4)
H7	0.3182	0.7184	0.0414	0.027*
C8	0.26838 (18)	0.65820 (12)	0.20278 (17)	0.0195 (4)
H8	0.1777	0.6522	0.1623	0.023*
C9	0.32896 (18)	0.57493 (13)	0.80129 (18)	0.0222 (4)
H9A	0.3065	0.5871	0.8892	0.027*
H9B	0.3097	0.6294	0.7455	0.027*
C10	0.47450 (19)	0.55474 (14)	0.82538 (19)	0.0286 (4)
H10A	0.4928	0.4974	0.8718	0.043*
H10B	0.5243	0.6018	0.8814	0.043*
H10C	0.5003	0.5521	0.7385	0.043*
C11	0.20739 (18)	0.43504 (12)	0.82325 (17)	0.0202 (4)
H11A	0.2858	0.4169	0.8922	0.024*
H11B	0.1734	0.3816	0.7694	0.024*
C12	0.10328 (19)	0.46898 (14)	0.89432 (19)	0.0262 (4)

H12A	0.1380	0.5199	0.9518	0.039*
H12B	0.0793	0.4213	0.9502	0.039*
H12C	0.0256	0.4876	0.8267	0.039*
C13	0.12031 (17)	0.39230 (11)	0.41262 (16)	0.0156 (3)
C14	0.22270 (18)	0.39048 (12)	0.34509 (17)	0.0195 (4)
H14	0.3026	0.4205	0.3821	0.023*
C15	0.20751 (19)	0.34464 (12)	0.22372 (18)	0.0225 (4)
H15	0.2771	0.3444	0.1777	0.027*
C16	0.09264 (19)	0.29929 (12)	0.16861 (18)	0.0240 (4)
H16	0.0834	0.2680	0.0857	0.029*
C17	-0.00860 (19)	0.30017 (12)	0.23605 (18)	0.0227 (4)
H17	-0.0876	0.2691	0.1993	0.027*
C18	0.00458 (17)	0.34619 (12)	0.35696 (17)	0.0190 (4)
H18	-0.0655	0.3464	0.4023	0.023*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0136 (8)	0.0170 (8)	0.0167 (8)	0.0015 (7)	0.0033 (6)	0.0001 (6)
N1	0.0168 (7)	0.0203 (8)	0.0156 (7)	-0.0023 (6)	0.0049 (5)	0.0008 (6)
N2	0.0190 (7)	0.0188 (7)	0.0145 (7)	-0.0030 (6)	0.0049 (6)	0.0001 (6)
N3	0.0158 (7)	0.0209 (8)	0.0153 (7)	-0.0029 (6)	0.0053 (6)	0.0002 (6)
C2	0.0181 (9)	0.0161 (8)	0.0156 (8)	-0.0008 (7)	0.0062 (7)	-0.0028 (6)
O1	0.0167 (6)	0.0247 (7)	0.0217 (6)	0.0005 (5)	0.0058 (5)	0.0048 (5)
C3	0.0200 (9)	0.0151 (8)	0.0175 (8)	0.0000 (7)	0.0086 (7)	-0.0019 (6)
C4	0.0210 (9)	0.0198 (9)	0.0201 (9)	-0.0001 (7)	0.0059 (7)	0.0017 (7)
C5	0.0213 (9)	0.0243 (10)	0.0308 (10)	-0.0031 (8)	0.0098 (8)	0.0006 (8)
C6	0.0304 (10)	0.0207 (9)	0.0291 (10)	-0.0032 (8)	0.0180 (8)	0.0001 (8)
C7	0.0332 (11)	0.0197 (9)	0.0177 (8)	0.0006 (8)	0.0101 (8)	0.0015 (7)
C8	0.0221 (9)	0.0189 (9)	0.0181 (8)	0.0001 (7)	0.0060 (7)	-0.0009 (7)
C9	0.0272 (10)	0.0227 (9)	0.0168 (8)	-0.0064 (8)	0.0053 (7)	-0.0036 (7)
C10	0.0258 (10)	0.0346 (11)	0.0234 (9)	-0.0083 (9)	0.0011 (8)	-0.0010 (8)
C11	0.0228 (9)	0.0232 (9)	0.0146 (8)	-0.0021 (7)	0.0041 (7)	0.0041 (7)
C12	0.0269 (10)	0.0337 (11)	0.0205 (9)	-0.0046 (8)	0.0105 (8)	0.0014 (8)
C13	0.0176 (8)	0.0139 (8)	0.0145 (8)	0.0016 (7)	0.0022 (6)	0.0020 (6)
C14	0.0167 (9)	0.0217 (9)	0.0202 (9)	-0.0014 (7)	0.0044 (7)	-0.0008 (7)
C15	0.0231 (9)	0.0224 (9)	0.0242 (9)	0.0018 (7)	0.0102 (7)	-0.0009 (7)
C16	0.0339 (11)	0.0197 (9)	0.0183 (9)	-0.0010 (8)	0.0056 (8)	-0.0035 (7)
C17	0.0248 (10)	0.0193 (9)	0.0225 (9)	-0.0042 (7)	0.0023 (7)	-0.0011 (7)
C18	0.0172 (9)	0.0200 (9)	0.0201 (8)	-0.0007 (7)	0.0049 (7)	0.0022 (7)

Geometric parameters (Å, °)

C1—N1	1.336 (2)	C9—H9A	0.9900
C1—N2	1.340 (2)	C9—H9B	0.9900
C1—N3	1.370 (2)	C10—H10A	0.9800
N1—C2	1.352 (2)	C10—H10B	0.9800
N2—C9	1.469 (2)	C10—H10C	0.9800

N2—C11	1.474 (2)	C11—C12	1.524 (3)
N3—C13	1.414 (2)	C11—H11A	0.9900
N3—H3	0.90 (2)	C11—H11B	0.9900
C2—O1	1.247 (2)	C12—H12A	0.9800
C2—C3	1.506 (2)	C12—H12B	0.9800
C3—C8	1.396 (2)	C12—H12C	0.9800
C3—C4	1.397 (3)	C13—C14	1.397 (3)
C4—C5	1.388 (3)	C13—C18	1.400 (2)
C4—H4	0.9500	C14—C15	1.389 (3)
C5—C6	1.388 (3)	C14—H14	0.9500
C5—H5	0.9500	C15—C16	1.386 (3)
C6—C7	1.388 (3)	C15—H15	0.9500
C6—H6	0.9500	C16—C17	1.385 (3)
C7—C8	1.392 (3)	C16—H16	0.9500
C7—H7	0.9500	C17—C18	1.388 (3)
C8—H8	0.9500	C17—H17	0.9500
C9—C10	1.519 (3)	C18—H18	0.9500
N1—C1—N2	118.14 (15)	C9—C10—H10A	109.5
N1—C1—N3	124.62 (15)	C9—C10—H10B	109.5
N2—C1—N3	117.14 (15)	H10A—C10—H10B	109.5
C1—N1—C2	121.38 (15)	C9—C10—H10C	109.5
C1—N2—C9	119.52 (14)	H10A—C10—H10C	109.5
C1—N2—C11	124.62 (15)	H10B—C10—H10C	109.5
C9—N2—C11	115.71 (14)	N2—C11—C12	113.04 (15)
C1—N3—C13	126.83 (15)	N2—C11—H11A	109.0
C1—N3—H3	117.8 (15)	C12—C11—H11A	109.0
C13—N3—H3	114.9 (14)	N2—C11—H11B	109.0
O1—C2—N1	127.02 (16)	C12—C11—H11B	109.0
O1—C2—C3	118.52 (16)	H11A—C11—H11B	107.8
N1—C2—C3	114.36 (15)	C11—C12—H12A	109.5
C8—C3—C4	119.10 (16)	C11—C12—H12B	109.5
C8—C3—C2	119.54 (16)	H12A—C12—H12B	109.5
C4—C3—C2	121.33 (16)	C11—C12—H12C	109.5
C5—C4—C3	120.41 (17)	H12A—C12—H12C	109.5
C5—C4—H4	119.8	H12B—C12—H12C	109.5
C3—C4—H4	119.8	C14—C13—C18	118.86 (16)
C6—C5—C4	120.22 (18)	C14—C13—N3	123.74 (16)
C6—C5—H5	119.9	C18—C13—N3	117.28 (16)
C4—C5—H5	119.9	C15—C14—C13	119.85 (17)
C5—C6—C7	119.79 (17)	C15—C14—H14	120.1
C5—C6—H6	120.1	C13—C14—H14	120.1
C7—C6—H6	120.1	C16—C15—C14	121.16 (18)
C6—C7—C8	120.23 (17)	C16—C15—H15	119.4
C6—C7—H7	119.9	C14—C15—H15	119.4
C8—C7—H7	119.9	C17—C16—C15	119.13 (17)
C7—C8—C3	120.24 (17)	C17—C16—H16	120.4
C7—C8—H8	119.9	C15—C16—H16	120.4

C3—C8—H8	119.9	C16—C17—C18	120.46 (17)
N2—C9—C10	113.05 (16)	C16—C17—H17	119.8
N2—C9—H9A	109.0	C18—C17—H17	119.8
C10—C9—H9A	109.0	C17—C18—C13	120.53 (17)
N2—C9—H9B	109.0	C17—C18—H18	119.7
C10—C9—H9B	109.0	C13—C18—H18	119.7
H9A—C9—H9B	107.8		
N2—C1—N1—C2	-141.68 (17)	C5—C6—C7—C8	0.5 (3)
N3—C1—N1—C2	42.2 (3)	C6—C7—C8—C3	0.1 (3)
N1—C1—N2—C9	10.7 (2)	C4—C3—C8—C7	-0.7 (3)
N3—C1—N2—C9	-172.82 (15)	C2—C3—C8—C7	177.63 (16)
N1—C1—N2—C11	-164.64 (16)	C1—N2—C9—C10	-85.7 (2)
N3—C1—N2—C11	11.8 (2)	C11—N2—C9—C10	90.06 (19)
N1—C1—N3—C13	22.4 (3)	C1—N2—C11—C12	-110.70 (19)
N2—C1—N3—C13	-153.84 (16)	C9—N2—C11—C12	73.8 (2)
C1—N1—C2—O1	17.6 (3)	C1—N3—C13—C14	19.1 (3)
C1—N1—C2—C3	-166.17 (15)	C1—N3—C13—C18	-164.97 (16)
O1—C2—C3—C8	-12.0 (2)	C18—C13—C14—C15	1.1 (3)
N1—C2—C3—C8	171.40 (16)	N3—C13—C14—C15	176.97 (16)
O1—C2—C3—C4	166.25 (16)	C13—C14—C15—C16	-0.9 (3)
N1—C2—C3—C4	-10.3 (2)	C14—C15—C16—C17	0.2 (3)
C8—C3—C4—C5	0.8 (3)	C15—C16—C17—C18	0.3 (3)
C2—C3—C4—C5	-177.53 (16)	C16—C17—C18—C13	-0.1 (3)
C3—C4—C5—C6	-0.2 (3)	C14—C13—C18—C17	-0.6 (3)
C4—C5—C6—C7	-0.4 (3)	N3—C13—C18—C17	-176.78 (16)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N3—H3 \cdots O1 ⁱ	0.90 (2)	1.97 (2)	2.852 (2)	168 (2)

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