

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

5-Bromo-2-[5-(4-nitrophenyl)-3-phenyl-4,5-dihydro-1H-pyrazol-1-yl]pyrimidine

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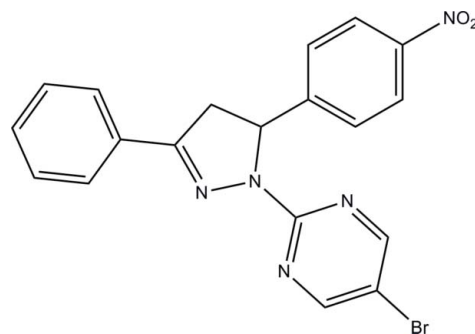
Received 13 November 2009; accepted 16 November 2009

Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.024; wR factor = 0.060; data-to-parameter ratio = 12.7.

In the title pyrazoline compound, $\text{C}_{19}\text{H}_{14}\text{BrN}_5\text{O}_2$, the essentially planar pyrazoline and pyrimidine rings [maximum deviations = 0.013 (1) and 0.009 (1) Å, respectively] are inclined slightly to one another, making a dihedral angle of 10.81 (10)°. The nitrobenzene unit is almost perpendicular to the attached pyrazoline ring, as indicated by the dihedral angle of 84.61 (8)°. In the crystal structure, intermolecular C—H...N contacts link the molecules into dimers in an antiparallel manner. These dimers are further linked into one-dimensional chains along the b axis via C—H...O contacts. The crystal structure is consolidated by three different intermolecular π — π interactions [range of centroid-centroid distances = 3.5160 (11)—3.6912 (11) Å].

Related literature

For general background to and applications of the title compound, see: Hegde *et al.* (2006); Kalluraya & Chimbalkar (2001); Kalluraya *et al.* (2001); Rai *et al.* (2008); Rathish *et al.* (2009); Tawab *et al.* (1960). For closely related structures, see: Goh *et al.* (2009*a,b*). For the stability of the temperature controller used for the data collection, see: Cosier & Glazer (1986).



Experimental

Crystal data

$\text{C}_{19}\text{H}_{14}\text{BrN}_5\text{O}_2$	$\gamma = 91.560$ (1)°
$M_r = 424.26$	$V = 882.12$ (2) Å ³
Triclinic, $P\bar{1}$	$Z = 2$
$a = 6.9709$ (1) Å	Mo $K\alpha$ radiation
$b = 11.6500$ (2) Å	$\mu = 2.36$ mm ⁻¹
$c = 12.4365$ (2) Å	$T = 100$ K
$\alpha = 114.969$ (1)°	$0.33 \times 0.22 \times 0.12$ mm
$\beta = 103.303$ (1)°	

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	28197 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2005)	3804 independent reflections
$T_{\min} = 0.510$, $T_{\max} = 0.760$	3431 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.026$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.024$	300 parameters
$wR(F^2) = 0.060$	All H-atom parameters refined
$S = 1.05$	$\Delta\rho_{\text{max}} = 0.49$ e Å ⁻³
3804 reflections	$\Delta\rho_{\text{min}} = -0.30$ e Å ⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C8—H8B...O2 ⁱ	0.95 (2)	2.41 (2)	3.352 (2)	176.0 (17)
C11—H11A...N4 ⁱⁱ	0.92 (2)	2.56 (2)	3.431 (2)	160.5 (18)
C19—H19A...O2 ⁱⁱⁱ	0.98 (2)	2.58 (2)	3.412 (3)	143.3 (17)

Symmetry codes: (i) $-x, -y, -z + 1$; (ii) $-x, -y + 1, -z + 1$; (iii) $x, y + 1, z$.

Data collection: APEX2 (Bruker, 2005); cell refinement: SAINT (Bruker, 2005); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2009).

HKF and JHG thank Universiti Sains Malaysia (USM) for the Research University Golden Goose grant (No. 1001/PFIZIK/811012). JHG also thanks USM for the award of a USM fellowship.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2577).

‡ Thomson Reuters ResearcherID: C-7576-2009.

§ Thomson Reuters ResearcherID: A-3561-2009.

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supporting information

Acta Cryst. (2009). E65, o3134–o3135 [doi:10.1107/S1600536809048600]

5-Bromo-2-[5-(4-nitrophenyl)-3-phenyl-4,5-dihydro-1H-pyrazol-1-yl]pyrimidine

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S1. Comment

Pyrazoles and pyrazoline derivatives are an important class of heterocyclic compounds (Rai *et al.*, 2008; Hegde *et al.*, 2006). The addition of aliphatic diazo compounds to olefins leads to pyrazolines. Also, the addition of hydrazine or its derivatives to α , β -unsaturated aldehydes or ketones yields pyrazoline (Kalluraya & Chimbalkar, 2001). Pyrazoline derivatives have been found to possess potential anti-pyretic, analgesic (Tawab *et al.*, 1960), anti-inflammatory (Rathish *et al.*, 2009), and anti-microbial (Kalluraya *et al.*, 2001) properties. In the present work, an X-ray crystal structure analysis has been undertaken in order to determine the 3D chemical structure and also the crystal packing of the molecules. We herein report the synthesis and crystal structure of the title compound, (I).

In (I), Fig. 1, the pyrazoline (C7-C9/N1/N2) and pyrimidine (C16-C19/N3/N4) rings are essentially planar, with maximum deviations of 0.013 (1) and 0.009 (1) Å, respectively, for atoms N1 and C16. These two rings are slightly inclined to one another, making a dihedral angle of 10.81 (10)°. The nitrobenzene moiety is almost perpendicular to the attached pyrazoline ring, as indicated by the dihedral angle formed between the mean plane through C10-C15/N5/O1/O2 and the pyrazoline ring of 84.61 (8)°. The bond lengths and angles are consistent with those closely related structures (Goh *et al.*, 2009*a,b*).

In the crystal structure (Fig. 2), intermolecular C11—H11A \cdots N4 contacts (Table 1) link the molecules into dimers in an anti-parallel manner. These dimers are further linked into a 1-D chain along the *b* axis by intermolecular C8—H8B \cdots O2 and C19—H19A \cdots O2 contacts (Table 1). The crystal structure is consolidated by three different weak π - π interactions involving the pyrazoline (*Cg*1), pyrimidine (*Cg*2) and C1-C6 benzene (*Cg*3) rings [*Cg*1 \cdots *Cg*1^{iv} = 3.5160 (11) Å, *Cg*2 \cdots *Cg*3ⁱⁱ = 3.6912 (11) Å and *Cg*2 \cdots *Cg*3^{iv} = 3.5779 (11) Å, respectively; (ii) -x, 1-y, 1-z and (iv) 1-x, 1-y, 1-z].

S2. Experimental

A mixture of 5-bromo-2-hydrazinopyrimidine (0.01 mol) and 3-(4-nitrophenyl)-1-phenyl-prop-2-en-1-one (0.01 mol) was taken in acetic acid (20 ml), and two drops of concentrated H₂SO₄ added. The mixture was refluxed for 4 h. The precipitated solids were filtered, dried and recrystallized from ethanol. The single crystals were obtained from a mixture of ethanol and DMF by slow evaporation.

S3. Refinement

All the H atoms were located from difference Fourier map [range of C—H = 0.91 (2) – 0.995 (19) Å] and allowed to refine freely. The reflections (001) and (0 $\bar{1}$ 1) were omitted as the intensities were affected by the beam-stop.

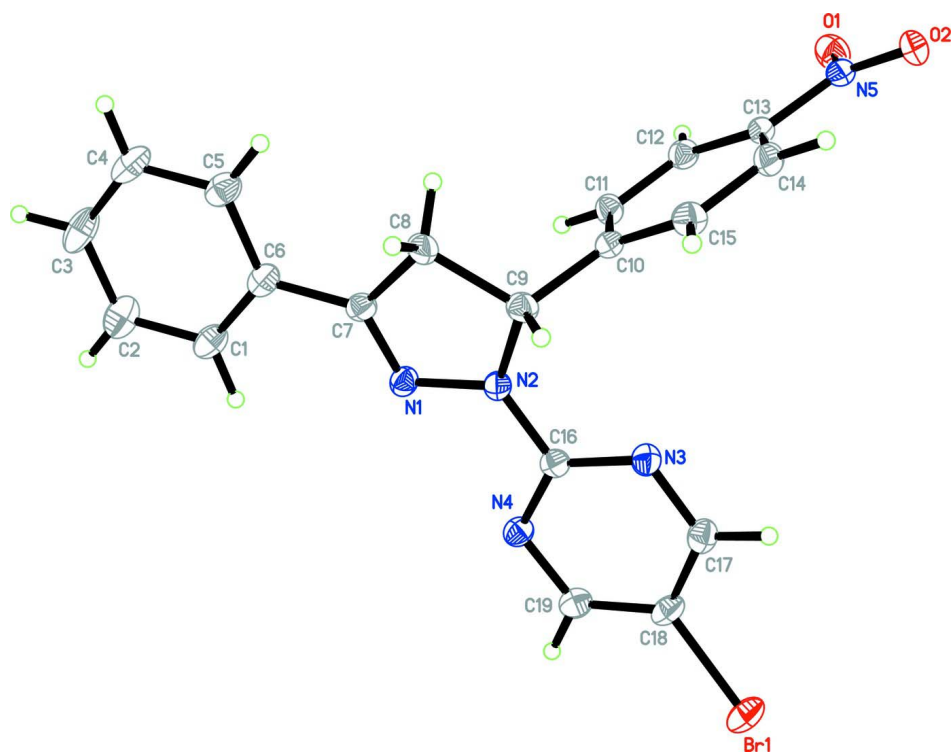


Figure 1

The molecular structure of (I), showing 50% probability displacement ellipsoids for non-H atoms and the atom-numbering scheme.

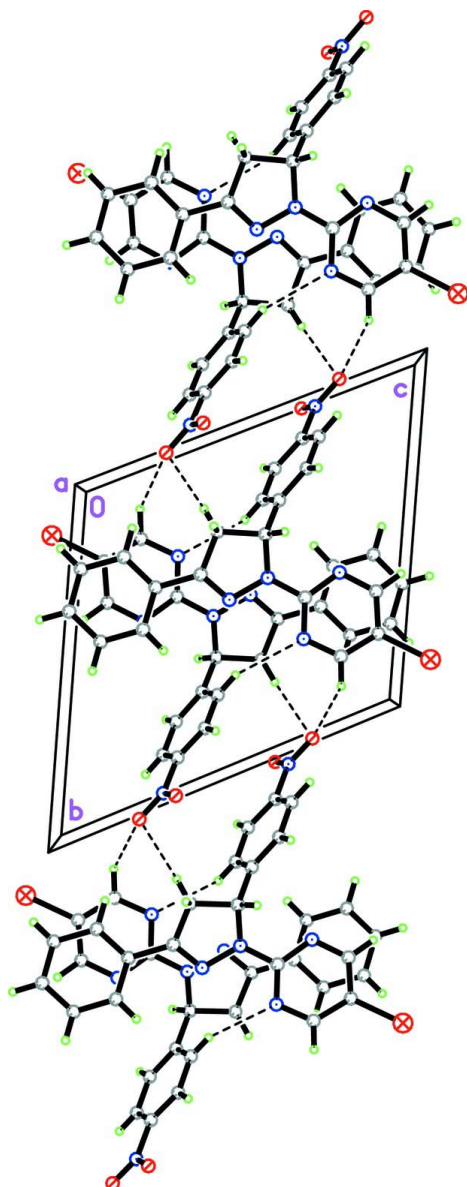


Figure 2

Part of the crystal structure of (I), viewed along the a axis, showing the dimers being linked into a 1-D chain along the b axis. Intermolecular contacts are shown as dashed lines.

5-Bromo-2-[5-(4-nitrophenyl)-3-phenyl-4,5-dihydro-1H-pyrazol-1-yl]pyrimidine

Crystal data

$C_{19}H_{14}BrN_5O_2$

$M_r = 424.26$

Triclinic, $P\bar{1}$

Hall symbol: $-P\ 1$

$a = 6.9709$ (1) Å

$b = 11.6500$ (2) Å

$c = 12.4365$ (2) Å

$\alpha = 114.969$ (1)°

$\beta = 103.303$ (1)°

$\gamma = 91.560$ (1)°

$V = 882.12$ (2) Å³

$Z = 2$

$F(000) = 428$

$D_x = 1.597$ Mg m⁻³

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

Cell parameters from 9873 reflections

$\theta = 3.0$ – 33.9 °

$\mu = 2.36$ mm⁻¹

$T = 100$ K $0.33 \times 0.22 \times 0.12$ mm
 Block, green

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	28197 measured reflections 3804 independent reflections
Radiation source: fine-focus sealed tube	3431 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\text{int}} = 0.026$
φ and ω scans	$\theta_{\text{max}} = 27.0^\circ$, $\theta_{\text{min}} = 2.0^\circ$
Absorption correction: multi-scan (SADABS; Bruker, 2005)	$h = -8 \rightarrow 8$ $k = -14 \rightarrow 14$ $l = -15 \rightarrow 15$
$T_{\text{min}} = 0.510$, $T_{\text{max}} = 0.760$	

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.024$	All H-atom parameters refined
$wR(F^2) = 0.060$	$w = 1/[\sigma^2(F_o^2) + (0.0288P)^2 + 0.5594P]$
$S = 1.05$	where $P = (F_o^2 + 2F_c^2)/3$
3804 reflections	$(\Delta/\sigma)_{\text{max}} = 0.001$
300 parameters	$\Delta\rho_{\text{max}} = 0.49 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.30 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

Special details

Experimental. The crystal was placed in the cold stream of an Oxford Cyrosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1)K.

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\sigma(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.32778 (3)	0.891599 (17)	1.068122 (15)	0.02770 (7)
O1	-0.3115 (2)	0.02211 (12)	0.64236 (12)	0.0274 (3)
O2	-0.0404 (2)	-0.01117 (12)	0.74511 (11)	0.0258 (3)
N1	0.2287 (2)	0.49488 (13)	0.46957 (12)	0.0175 (3)
N2	0.2681 (2)	0.47661 (13)	0.57546 (12)	0.0177 (3)
N3	0.3203 (2)	0.54232 (13)	0.78176 (13)	0.0191 (3)
N4	0.2364 (2)	0.68890 (13)	0.69321 (13)	0.0182 (3)
N5	-0.1294 (2)	0.03714 (13)	0.68034 (13)	0.0205 (3)
C1	0.1859 (3)	0.48220 (17)	0.22862 (16)	0.0224 (4)
C2	0.1604 (3)	0.4683 (2)	0.10980 (17)	0.0282 (4)
C3	0.1682 (3)	0.3501 (2)	0.01564 (17)	0.0300 (4)
C4	0.2012 (3)	0.24641 (19)	0.04084 (17)	0.0275 (4)

C5	0.2323 (3)	0.26060 (18)	0.16071 (16)	0.0236 (4)
C6	0.2252 (2)	0.37903 (16)	0.25600 (15)	0.0201 (3)
C7	0.2607 (2)	0.39338 (15)	0.38231 (15)	0.0176 (3)
C8	0.3349 (3)	0.29328 (16)	0.42148 (17)	0.0228 (4)
C9	0.3406 (3)	0.35347 (16)	0.55889 (15)	0.0191 (3)
C10	0.2129 (3)	0.27153 (15)	0.59130 (14)	0.0175 (3)
C11	0.0070 (3)	0.24832 (16)	0.54497 (15)	0.0188 (3)
C12	-0.1072 (3)	0.17094 (16)	0.57339 (15)	0.0191 (3)
C13	-0.0101 (3)	0.11782 (15)	0.64817 (15)	0.0178 (3)
C14	0.1942 (3)	0.13677 (16)	0.69325 (16)	0.0211 (3)
C15	0.3060 (3)	0.21486 (16)	0.66433 (16)	0.0213 (3)
C16	0.2747 (2)	0.57397 (15)	0.68747 (15)	0.0165 (3)
C17	0.3326 (3)	0.63670 (17)	0.89244 (16)	0.0207 (3)
C18	0.2999 (3)	0.75881 (16)	0.90934 (15)	0.0204 (3)
C19	0.2496 (3)	0.78065 (16)	0.80562 (16)	0.0201 (3)
H1A	0.177 (3)	0.563 (2)	0.2909 (19)	0.025 (5)*
H2A	0.138 (4)	0.540 (2)	0.092 (2)	0.038 (6)*
H3A	0.154 (3)	0.340 (2)	-0.066 (2)	0.037 (6)*
H4A	0.206 (3)	0.169 (2)	-0.020 (2)	0.024 (5)*
H5A	0.257 (3)	0.190 (2)	0.178 (2)	0.028 (5)*
H8A	0.467 (3)	0.2771 (19)	0.4110 (19)	0.024 (5)*
H8B	0.250 (3)	0.214 (2)	0.378 (2)	0.031 (6)*
H9A	0.478 (3)	0.3691 (17)	0.6123 (17)	0.013 (4)*
H11A	-0.055 (3)	0.2853 (18)	0.4970 (18)	0.019 (5)*
H12A	-0.245 (3)	0.1549 (19)	0.5429 (19)	0.024 (5)*
H14A	0.258 (3)	0.0991 (19)	0.7416 (19)	0.025 (5)*
H15A	0.446 (3)	0.2291 (19)	0.6952 (19)	0.025 (5)*
H17A	0.372 (3)	0.6149 (19)	0.9598 (19)	0.022 (5)*
H19A	0.224 (3)	0.8658 (19)	0.8141 (18)	0.019 (5)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.02884 (11)	0.02882 (11)	0.01659 (9)	0.00509 (7)	0.00679 (7)	0.00134 (7)
O1	0.0263 (7)	0.0275 (7)	0.0323 (7)	0.0020 (5)	0.0112 (6)	0.0154 (6)
O2	0.0386 (8)	0.0213 (6)	0.0223 (6)	0.0044 (5)	0.0086 (6)	0.0139 (5)
N1	0.0171 (7)	0.0197 (7)	0.0145 (6)	0.0012 (5)	0.0029 (5)	0.0070 (5)
N2	0.0231 (8)	0.0155 (6)	0.0160 (7)	0.0037 (5)	0.0070 (6)	0.0073 (5)
N3	0.0199 (7)	0.0202 (7)	0.0180 (7)	0.0023 (6)	0.0060 (6)	0.0087 (6)
N4	0.0185 (7)	0.0179 (7)	0.0174 (7)	0.0043 (5)	0.0047 (6)	0.0068 (6)
N5	0.0302 (9)	0.0147 (6)	0.0173 (7)	0.0036 (6)	0.0107 (6)	0.0055 (6)
C1	0.0190 (9)	0.0260 (9)	0.0173 (8)	0.0058 (7)	0.0023 (7)	0.0061 (7)
C2	0.0244 (10)	0.0390 (11)	0.0209 (9)	0.0122 (8)	0.0031 (7)	0.0140 (8)
C3	0.0235 (10)	0.0459 (12)	0.0152 (8)	0.0112 (8)	0.0028 (7)	0.0091 (8)
C4	0.0202 (9)	0.0321 (10)	0.0173 (8)	0.0036 (8)	0.0049 (7)	-0.0012 (8)
C5	0.0185 (9)	0.0247 (9)	0.0219 (9)	0.0023 (7)	0.0063 (7)	0.0045 (7)
C6	0.0137 (8)	0.0250 (9)	0.0169 (8)	0.0010 (6)	0.0040 (6)	0.0049 (7)
C7	0.0149 (8)	0.0172 (7)	0.0179 (8)	-0.0004 (6)	0.0050 (6)	0.0051 (6)

C8	0.0315 (10)	0.0164 (8)	0.0246 (9)	0.0043 (7)	0.0156 (8)	0.0086 (7)
C9	0.0212 (9)	0.0161 (7)	0.0209 (8)	0.0036 (6)	0.0081 (7)	0.0078 (6)
C10	0.0236 (9)	0.0140 (7)	0.0149 (7)	0.0033 (6)	0.0077 (6)	0.0050 (6)
C11	0.0225 (9)	0.0184 (8)	0.0179 (8)	0.0055 (7)	0.0056 (7)	0.0099 (7)
C12	0.0204 (9)	0.0187 (8)	0.0183 (8)	0.0046 (7)	0.0060 (7)	0.0075 (7)
C13	0.0252 (9)	0.0137 (7)	0.0153 (7)	0.0028 (6)	0.0085 (7)	0.0057 (6)
C14	0.0278 (10)	0.0197 (8)	0.0179 (8)	0.0066 (7)	0.0054 (7)	0.0103 (7)
C15	0.0194 (9)	0.0216 (8)	0.0212 (8)	0.0032 (7)	0.0030 (7)	0.0092 (7)
C16	0.0136 (8)	0.0174 (8)	0.0173 (8)	0.0013 (6)	0.0048 (6)	0.0064 (6)
C17	0.0191 (9)	0.0255 (9)	0.0180 (8)	0.0028 (7)	0.0059 (7)	0.0095 (7)
C18	0.0176 (8)	0.0222 (8)	0.0152 (8)	0.0020 (7)	0.0049 (6)	0.0023 (7)
C19	0.0187 (9)	0.0192 (8)	0.0204 (8)	0.0046 (7)	0.0051 (7)	0.0067 (7)

Geometric parameters (Å, °)

Br1—C18	1.8863 (16)	C5—H5A	0.94 (2)
O1—N5	1.228 (2)	C6—C7	1.468 (2)
O2—N5	1.2337 (19)	C7—C8	1.504 (2)
N1—C7	1.292 (2)	C8—C9	1.539 (2)
N1—N2	1.3886 (19)	C8—H8A	0.97 (2)
N2—C16	1.365 (2)	C8—H8B	0.95 (2)
N2—C9	1.481 (2)	C9—C10	1.519 (2)
N3—C17	1.332 (2)	C9—H9A	0.995 (19)
N3—C16	1.346 (2)	C10—C11	1.391 (2)
N4—C19	1.334 (2)	C10—C15	1.393 (2)
N4—C16	1.348 (2)	C11—C12	1.389 (2)
N5—C13	1.471 (2)	C11—H11A	0.91 (2)
C1—C2	1.384 (3)	C12—C13	1.387 (2)
C1—C6	1.399 (3)	C12—H12A	0.93 (2)
C1—H1A	0.95 (2)	C13—C14	1.379 (3)
C2—C3	1.394 (3)	C14—C15	1.390 (2)
C2—H2A	0.96 (2)	C14—H14A	0.93 (2)
C3—C4	1.383 (3)	C15—H15A	0.94 (2)
C3—H3A	0.95 (2)	C17—C18	1.381 (3)
C4—C5	1.392 (3)	C17—H17A	0.96 (2)
C4—H4A	0.91 (2)	C18—C19	1.388 (2)
C5—C6	1.400 (2)	C19—H19A	0.98 (2)
C7—N1—N2	107.71 (14)	N2—C9—C10	113.23 (14)
C16—N2—N1	121.50 (13)	N2—C9—C8	101.32 (13)
C16—N2—C9	123.73 (14)	C10—C9—C8	113.11 (14)
N1—N2—C9	113.65 (13)	N2—C9—H9A	110.2 (10)
C17—N3—C16	115.56 (15)	C10—C9—H9A	107.2 (10)
C19—N4—C16	115.45 (14)	C8—C9—H9A	111.9 (10)
O1—N5—O2	123.49 (14)	C11—C10—C15	119.99 (15)
O1—N5—C13	118.57 (14)	C11—C10—C9	121.01 (15)
O2—N5—C13	117.94 (15)	C15—C10—C9	118.95 (15)
C2—C1—C6	120.44 (17)	C12—C11—C10	120.27 (16)

C2—C1—H1A	118.6 (12)	C12—C11—H11A	119.2 (12)
C6—C1—H1A	121.0 (12)	C10—C11—H11A	120.5 (12)
C1—C2—C3	120.15 (19)	C13—C12—C11	118.28 (16)
C1—C2—H2A	120.0 (14)	C13—C12—H12A	120.8 (12)
C3—C2—H2A	119.8 (14)	C11—C12—H12A	121.0 (12)
C4—C3—C2	119.95 (17)	C14—C13—C12	122.77 (15)
C4—C3—H3A	119.4 (14)	C14—C13—N5	118.41 (15)
C2—C3—H3A	120.6 (14)	C12—C13—N5	118.82 (15)
C3—C4—C5	120.16 (17)	C13—C14—C15	118.17 (16)
C3—C4—H4A	120.9 (13)	C13—C14—H14A	122.0 (13)
C5—C4—H4A	118.9 (13)	C15—C14—H14A	119.8 (13)
C4—C5—C6	120.28 (18)	C14—C15—C10	120.49 (17)
C4—C5—H5A	120.2 (13)	C14—C15—H15A	119.0 (12)
C6—C5—H5A	119.5 (13)	C10—C15—H15A	120.5 (12)
C1—C6—C5	118.96 (16)	N3—C16—N4	127.09 (15)
C1—C6—C7	121.19 (15)	N3—C16—N2	114.49 (14)
C5—C6—C7	119.85 (16)	N4—C16—N2	118.43 (14)
N1—C7—C6	121.79 (16)	N3—C17—C18	122.24 (16)
N1—C7—C8	114.40 (15)	N3—C17—H17A	115.3 (12)
C6—C7—C8	123.81 (15)	C18—C17—H17A	122.4 (12)
C7—C8—C9	102.87 (13)	C17—C18—C19	117.58 (15)
C7—C8—H8A	112.2 (12)	C17—C18—Br1	121.08 (13)
C9—C8—H8A	110.4 (12)	C19—C18—Br1	121.33 (13)
C7—C8—H8B	112.8 (13)	N4—C19—C18	122.05 (16)
C9—C8—H8B	111.1 (13)	N4—C19—H19A	118.1 (11)
H8A—C8—H8B	107.5 (18)	C18—C19—H19A	119.8 (11)
C7—N1—N2—C16	170.72 (14)	C15—C10—C11—C12	1.4 (2)
C7—N1—N2—C9	2.42 (18)	C9—C10—C11—C12	178.62 (15)
C6—C1—C2—C3	1.8 (3)	C10—C11—C12—C13	-0.2 (2)
C1—C2—C3—C4	0.1 (3)	C11—C12—C13—C14	-1.2 (2)
C2—C3—C4—C5	-1.8 (3)	C11—C12—C13—N5	179.18 (14)
C3—C4—C5—C6	1.6 (3)	O1—N5—C13—C14	178.59 (15)
C2—C1—C6—C5	-1.9 (3)	O2—N5—C13—C14	-0.9 (2)
C2—C1—C6—C7	177.49 (16)	O1—N5—C13—C12	-1.8 (2)
C4—C5—C6—C1	0.2 (3)	O2—N5—C13—C12	178.73 (14)
C4—C5—C6—C7	-179.20 (16)	C12—C13—C14—C15	1.5 (3)
N2—N1—C7—C6	177.49 (14)	N5—C13—C14—C15	-178.92 (15)
N2—N1—C7—C8	-2.15 (19)	C13—C14—C15—C10	-0.3 (3)
C1—C6—C7—N1	10.4 (2)	C11—C10—C15—C14	-1.1 (2)
C5—C6—C7—N1	-170.23 (16)	C9—C10—C15—C14	-178.41 (15)
C1—C6—C7—C8	-170.01 (16)	C17—N3—C16—N4	-1.6 (2)
C5—C6—C7—C8	9.4 (2)	C17—N3—C16—N2	178.34 (14)
N1—C7—C8—C9	1.11 (19)	C19—N4—C16—N3	1.4 (2)
C6—C7—C8—C9	-178.52 (15)	C19—N4—C16—N2	-178.54 (14)
C16—N2—C9—C10	68.9 (2)	N1—N2—C16—N3	-178.10 (14)
N1—N2—C9—C10	-123.07 (15)	C9—N2—C16—N3	-11.0 (2)
C16—N2—C9—C8	-169.64 (15)	N1—N2—C16—N4	1.8 (2)

N1—N2—C9—C8	-1.64 (17)	C9—N2—C16—N4	168.92 (14)
C7—C8—C9—N2	0.35 (16)	C16—N3—C17—C18	0.3 (2)
C7—C8—C9—C10	121.86 (15)	N3—C17—C18—C19	1.0 (3)
N2—C9—C10—C11	49.0 (2)	N3—C17—C18—Br1	-177.75 (12)
C8—C9—C10—C11	-65.5 (2)	C16—N4—C19—C18	0.1 (2)
N2—C9—C10—C15	-133.70 (16)	C17—C18—C19—N4	-1.2 (3)
C8—C9—C10—C15	111.76 (18)	Br1—C18—C19—N4	177.53 (13)

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>
C8—H8B...O2 ⁱ	0.95 (2)	2.41 (2)	3.352 (2)	176.0 (17)
C11—H11A...N4 ⁱⁱ	0.92 (2)	2.56 (2)	3.431 (2)	160.5 (18)
C19—H19A...O2 ⁱⁱⁱ	0.98 (2)	2.58 (2)	3.412 (3)	143.3 (17)

Symmetry codes: (i) $-x, -y, -z+1$; (ii) $-x, -y+1, -z+1$; (iii) $x, y+1, z$.