

Received 25 May 2018
Accepted 30 May 2018

Edited by J. Simpson, University of Otago, New Zealand

Keywords: crystal structure; 2-aminopyrimidine; hydrogen bonds; intermolecular dimers.

CCDC reference: 1846182

Structural data: full structural data are available from iucrdata.iucr.org

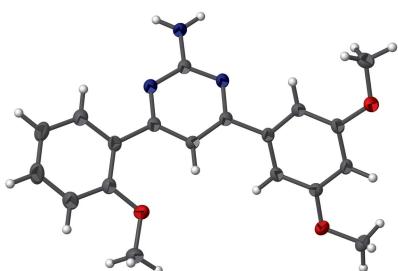
4-(3,5-Dimethoxyphenyl)-6-(2-methoxyphenyl)-pyrimidin-2-amine

Dongsoo Koh* and Ji Hye Lee

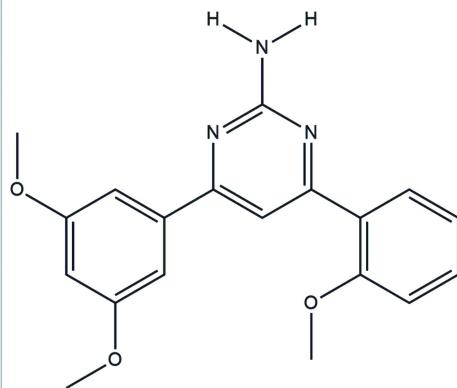
Department of Applied Chemistry, Dongduk Women's University, Seoul 136-714, Republic of Korea. *Correspondence e-mail: dskoh@dongduk.ac.kr

In the title molecule, $C_{19}H_{19}N_3O_3$, the 2-methoxyphenyl and 3,5-dimethoxyphenyl rings are attached at the 4- and 6-positions, respectively, of the central 2-aminopyrimidine ring. The dihedral angles between the planes of the benzene rings and that of the 2-aminopyrimidine ring are $17.31(9)$ and $44.39(6)^\circ$, respectively. In the crystal, pairs of $N-H \cdots N$ hydrogen bonds form inversion dimers enclosing $R_2^2(8)$ rings. Pairs of $N-H \cdots O$ hydrogen bonds link the dimers into chains along [010].

3D view



Chemical scheme

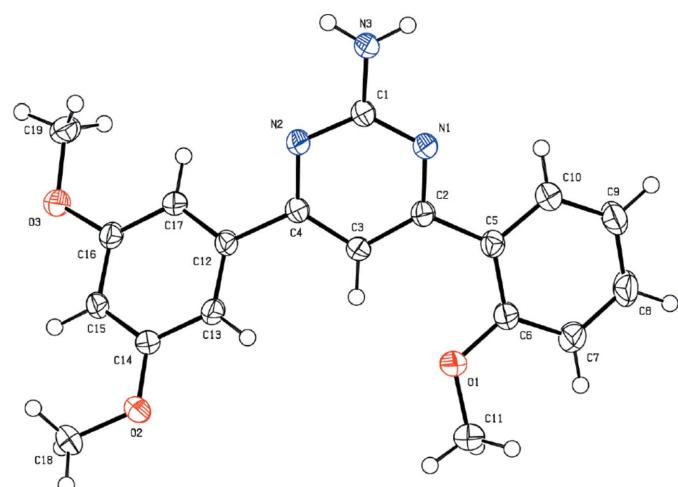


Structure description

2-Aminopyrimidine pharmacophores have shown a broad spectrum of biological activities including use against Parkinson's disease (Robinson *et al.*, 2016) and displaying anti-bacterial (Nagarajan *et al.*, 2014), anti-platelet (Giridhar *et al.*, 2012), antidiabetic (Singh *et al.*, 2011) and antitumor properties (Lee *et al.*, 2011). The title 2-aminopyrimidine compound was synthesized in a continuation of our research program to expand the use of novel synthetic chalcones (Lee *et al.* 2016), and its crystal structure was determined and is reported here.

The molecular structure of the title compound is shown in Fig. 1. The central 2-aminopyrimidine ring contains two benzene rings at the C4 and C6 positions respectively. The dihedral angles between central 2-aminopyrimidine ring and the C5–C10 and C12–C17 benzene rings are $17.31(9)$ and $44.39(6)^\circ$, respectively. All three methoxy groups on the benzene rings are slightly twisted from the ring plane [$C17-C16-O3-C19 = -4.9(2)$, $C15-C14-O2-C18 = -4.5(2)$ and $C7-C6-O1-C11 = 3.7(2)^\circ$].

In the crystal, pairs of $N3-H3B \cdots N2$ hydrogen bonds form inversion dimers that enclose $R_2^2(8)$ rings. These dimers are linked into chains along the *b*-axis direction by pairs of $N3-H3A \cdots O3$ hydrogen bonds (Table 1, Fig. 2).

**Figure 1**

The molecular structure of the title compound, showing the atom-labelling scheme with displacement ellipsoids drawn at the 30% probability level.

Some examples of other 2-aminopyrimidine structures have also been published recently (Sangeetha *et al.*, 2016; Thangaimani *et al.*, 2012).

Synthesis and crystallization

A synthetic scheme is shown in Fig. 3. The chalcone (*E*)-1-(3,5-dimethoxyphenyl)-3-(2-methoxyphenyl)prop-2-en-1-one was prepared as a starting material by a previously reported method (Koh *et al.* 2016). 2-Amino pyrimidine was obtained by a cyclization reaction of this chalcone with guanidine hydrochloride in basic solution. To an ethanol solution of 3,5-dimethoxyacetophenone (I) and 2-methoxybenzaldehyde (II) an excess amount of 50% aqueous KOH was added and the mixture was stirred at room temperature for 20 h. After completion of reaction, the reaction mixture was poured into 6M HCl in an ice-bath to give a precipitate of the chalcone (III). The solid was filtered and washed with ethanol and was used for the next reaction without further purification. The chalcone (III, 1 eq.) and the guanidine HCl salt (1.5 eq.) were

Table 1
Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N3—H3B···N2 ⁱ	0.87	2.28	3.1374 (19)	167
N3—H3A···O3 ⁱⁱ	0.87	2.39	3.2545 (17)	175

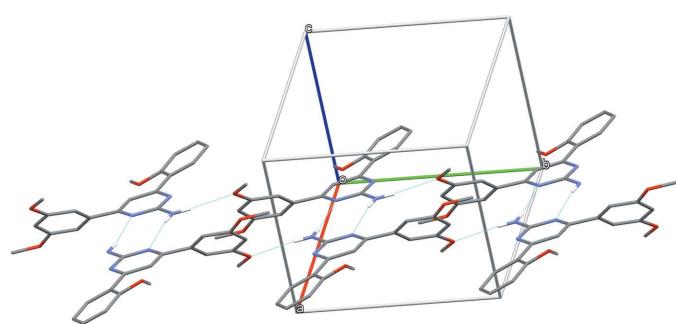
Symmetry codes: (i) $-x + 3, -y + 1, -z + 2$; (ii) $x, y + 1, z$.

Table 2
Experimental details.

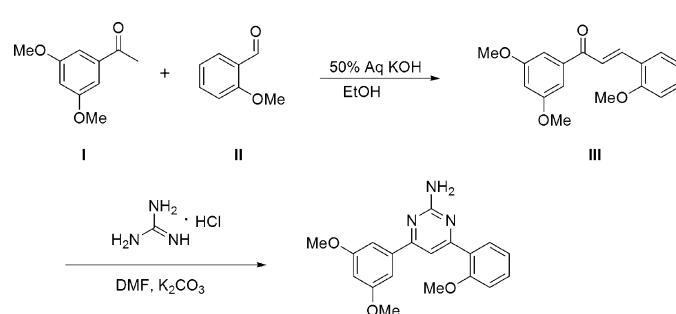
Crystal data	
Chemical formula	C ₁₉ H ₁₉ N ₃ O ₃
M_r	337.37
Crystal system, space group	Triclinic, $P\bar{1}$
Temperature (K)	223
a, b, c (Å)	7.5867 (7), 10.2680 (7), 11.4684 (8)
α, β, γ (°)	95.592 (4), 102.147 (4), 109.987 (3)
V (Å ³)	806.71 (11)
Z	2
Radiation type	Mo $K\alpha$
μ (mm ⁻¹)	0.10
Crystal size (mm)	0.18 × 0.10 × 0.06
Data collection	
Diffractometer	Bruker PHOTON 100 CMOS
Absorption correction	Multi-scan (<i>SADABS</i> ; Bruker, 2012)
T_{\min}, T_{\max}	0.690, 0.746
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	26671, 4065, 2736
R_{int}	0.063
(sin θ/λ) _{max} (Å ⁻¹)	0.671
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.046, 0.112, 1.02
No. of reflections	4065
No. of parameters	229
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{\text{max}}, \Delta\rho_{\text{min}}$ (e Å ⁻³)	0.24, -0.21

Computer programs: *APEX2* and *SAINT* (Bruker, 2012), *SHELXS* and *SHELXTL* (Sheldrick, 2008), *SHELXL2014* (Sheldrick, 2015) and *publCIF* (Westrip, 2010).

dissolved in a DMF solution to which was added solid K₂CO₃ (3 eq.). The reaction mixture was refluxed for 2 h and cooled to room temperature. The reaction mixture was then poured into 3M HCl in an ice bath to give a precipitate of the title 2-aminopyrimidine compound, which was purified by recrystallization from ethanol.

**Figure 2**

Part of the crystal structure with intermolecular hydrogen bonds shown as dashed lines. For clarity only those H atoms involved in hydrogen bonding are shown.

**Figure 3**

Synthetic scheme for the preparation of the title compound.

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2.

Funding information

The authors acknowledge financial support from the Basic Science Research Program (award No. NRF-2016R1D1A1B03931623).

References

- Bruker (2012). *APEX2, SAINT and SADABS*, Bruker AXS Inc. Madison, Wisconsin, USA.
- Giridhar, R., Tamboli, R. S., Ramajayam, R., Prajapati, D. G. & Yadav, M. R. (2012). *Eur. J. Med. Chem.* **50**, 428–432.
- Koh, D., Jung, Y., Kim, B. S., Ahn, S. & Lim, Y. (2016). *Magn. Reson. Chem.* **54**, 842–851.
- Lee, Y., Kim, B. S., Ahn, S., Koh, D., Lee, Y. H., Shin, S. Y. & Lim, Y. (2016). *Bioorg. Chem.* **68**, 166–176.
- Lee, J., Kim, K.-H. & Jeong, S. (2011). *Bioorg. Med. Chem. Lett.* **21**, 4203–4205.
- Nagarajan, S., Shanmugavelan, P., Sathishkumar, M., Selvi, R., PonnuSwamy, A., Harikrishnan, H., Shanmugaiah, V. & Murugavel, S. (2014). *Bioorg. Med. Chem. Lett.* **24**, 4999–5007.
- Robinson, S. J., Petzer, J. P., Rousseau, A. L., Terre'Blanche, G., Petzer, A. & Lourens, A. C. U. (2016). *Bioorg. Med. Chem. Lett.* **26**, 734–738.
- Sangeetha, R., Edison, B., Thanikasalam, K., Kavitha, S. J. & Balasubramani, K. (2016). *IUCrData*, **1**, x160794.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Sheldrick, G. M. (2015). *Acta Cryst. C* **71**, 3–8.
- Singh, N., Pandey, S. K., Anand, N., Dwivedi, R., Singh, S., Sinha, S. K., Chaturvedi, V., Jaiswal, N., Srivastava, A. K., Shah, P., Siddiqui, M. I. & Tripathi, R. P. (2011). *Bioorg. Med. Chem. Lett.* **21**, 4404–4408.
- Thanigaimani, K., Khalib, N. C., Arshad, S. & Razak, I. A. (2012). *Acta Cryst. E* **68**, o3318.
- Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.

full crystallographic data

IUCrData (2018). **3**, x180796 [https://doi.org/10.1107/S2414314618007964]

4-(3,5-Dimethoxyphenyl)-6-(2-methoxyphenyl)pyrimidin-2-amine

Dongsoo Koh and Ji Hye Lee

4-(3,5-Dimethoxyphenyl)-6-(2-methoxyphenyl)pyrimidin-2-amine

Crystal data

$C_{19}H_{19}N_3O_3$
 $M_r = 337.37$
Triclinic, $P\bar{1}$
 $a = 7.5867 (7)$ Å
 $b = 10.2680 (7)$ Å
 $c = 11.4684 (8)$ Å
 $\alpha = 95.592 (4)^\circ$
 $\beta = 102.147 (4)^\circ$
 $\gamma = 109.987 (3)^\circ$
 $V = 806.71 (11)$ Å³

$Z = 2$
 $F(000) = 356$
 $D_x = 1.389$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 5933 reflections
 $\theta = 2.6\text{--}27.9^\circ$
 $\mu = 0.10$ mm⁻¹
 $T = 223$ K
Block, yellow
 $0.18 \times 0.10 \times 0.06$ mm

Data collection

Bruker PHOTON 100 CMOS
diffractometer
 φ and ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2012)
 $T_{\min} = 0.690$, $T_{\max} = 0.746$
26671 measured reflections

4065 independent reflections
2736 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.063$
 $\theta_{\max} = 28.5^\circ$, $\theta_{\min} = 2.9^\circ$
 $h = -10 \rightarrow 10$
 $k = -13 \rightarrow 13$
 $l = -15 \rightarrow 15$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.112$
 $S = 1.02$
4065 reflections
229 parameters
0 restraints

Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0434P)^2 + 0.2728P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.24$ e Å⁻³
 $\Delta\rho_{\min} = -0.21$ e Å⁻³

Special details

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.

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	1.2100 (2)	0.51141 (15)	0.91277 (13)	0.0264 (3)
N1	1.04986 (18)	0.52945 (13)	0.85350 (11)	0.0277 (3)
C2	0.9093 (2)	0.41773 (15)	0.77721 (13)	0.0240 (3)
C3	0.9345 (2)	0.29009 (16)	0.75753 (14)	0.0266 (3)
H3	0.8409	0.2128	0.7001	0.032*
C4	1.1029 (2)	0.28096 (15)	0.82566 (13)	0.0251 (3)
N2	1.24219 (18)	0.38985 (13)	0.90484 (11)	0.0277 (3)
N3	1.3510 (2)	0.62546 (13)	0.98728 (12)	0.0346 (3)
H3A	1.3357	0.7058	0.9951	0.052*
H3B	1.4575	0.6189	1.0276	0.052*
C5	0.7267 (2)	0.44012 (15)	0.72488 (13)	0.0250 (3)
C6	0.5714 (2)	0.34812 (16)	0.62920 (14)	0.0282 (3)
C7	0.4022 (2)	0.37506 (19)	0.59430 (17)	0.0387 (4)
H7	0.2980	0.3120	0.5314	0.046*
C8	0.3857 (3)	0.49308 (19)	0.65090 (18)	0.0425 (4)
H8	0.2704	0.5100	0.6267	0.051*
C9	0.5371 (3)	0.58643 (18)	0.74266 (17)	0.0382 (4)
H9	0.5269	0.6681	0.7803	0.046*
C10	0.7041 (2)	0.55922 (17)	0.77906 (15)	0.0311 (4)
H10	0.8063	0.6230	0.8426	0.037*
O1	0.59265 (17)	0.23350 (12)	0.57163 (11)	0.0385 (3)
C11	0.4395 (3)	0.14655 (19)	0.46928 (17)	0.0456 (5)
H11A	0.3258	0.0969	0.4961	0.068*
H11B	0.4809	0.0788	0.4292	0.068*
H11C	0.4079	0.2049	0.4130	0.068*
C12	1.1292 (2)	0.14340 (15)	0.81773 (14)	0.0253 (3)
C13	1.0781 (2)	0.05546 (16)	0.70723 (14)	0.0269 (3)
H13	1.0340	0.0843	0.6351	0.032*
C14	1.0924 (2)	-0.07653 (16)	0.70354 (14)	0.0259 (3)
C15	1.1509 (2)	-0.12181 (16)	0.80938 (14)	0.0268 (3)
H15	1.1533	-0.2130	0.8067	0.032*
C16	1.2062 (2)	-0.03059 (15)	0.91990 (13)	0.0260 (3)
C17	1.1975 (2)	0.10205 (15)	0.92550 (14)	0.0259 (3)
H17	1.2368	0.1635	1.0004	0.031*
O2	1.04676 (17)	-0.15512 (12)	0.59013 (10)	0.0349 (3)
C18	1.0753 (3)	-0.28546 (17)	0.58342 (16)	0.0358 (4)
H18A	1.2073	-0.2692	0.6273	0.054*
H18B	1.0534	-0.3259	0.4991	0.054*
H18C	0.9848	-0.3500	0.6193	0.054*
O3	1.26833 (17)	-0.08393 (11)	1.01866 (10)	0.0348 (3)
C19	1.3421 (2)	0.00789 (18)	1.13319 (14)	0.0347 (4)
H19A	1.4454	0.0932	1.1287	0.052*
H19B	1.3926	-0.0386	1.1942	0.052*
H19C	1.2386	0.0318	1.1549	0.052*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0294 (8)	0.0204 (8)	0.0277 (8)	0.0096 (6)	0.0031 (6)	0.0041 (6)
N1	0.0291 (7)	0.0207 (7)	0.0298 (7)	0.0095 (5)	0.0008 (5)	0.0032 (5)
C2	0.0261 (8)	0.0215 (7)	0.0248 (7)	0.0094 (6)	0.0052 (6)	0.0062 (6)
C3	0.0278 (8)	0.0191 (7)	0.0279 (8)	0.0082 (6)	-0.0010 (6)	0.0012 (6)
C4	0.0281 (8)	0.0193 (7)	0.0263 (8)	0.0091 (6)	0.0031 (6)	0.0044 (6)
N2	0.0287 (7)	0.0199 (6)	0.0308 (7)	0.0098 (5)	-0.0007 (5)	0.0034 (5)
N3	0.0339 (8)	0.0200 (7)	0.0398 (8)	0.0102 (6)	-0.0078 (6)	-0.0023 (6)
C5	0.0266 (8)	0.0226 (8)	0.0273 (8)	0.0102 (6)	0.0069 (6)	0.0081 (6)
C6	0.0277 (8)	0.0245 (8)	0.0325 (8)	0.0116 (6)	0.0051 (7)	0.0044 (7)
C7	0.0282 (9)	0.0333 (9)	0.0487 (11)	0.0128 (7)	-0.0018 (8)	0.0019 (8)
C8	0.0303 (9)	0.0382 (10)	0.0622 (12)	0.0209 (8)	0.0055 (8)	0.0074 (9)
C9	0.0394 (10)	0.0306 (9)	0.0494 (11)	0.0201 (8)	0.0113 (8)	0.0036 (8)
C10	0.0333 (9)	0.0261 (8)	0.0336 (9)	0.0129 (7)	0.0058 (7)	0.0043 (7)
O1	0.0339 (7)	0.0347 (7)	0.0396 (7)	0.0182 (5)	-0.0071 (5)	-0.0095 (5)
C11	0.0466 (11)	0.0345 (10)	0.0433 (10)	0.0166 (9)	-0.0108 (9)	-0.0062 (8)
C12	0.0223 (7)	0.0183 (7)	0.0320 (8)	0.0073 (6)	0.0007 (6)	0.0039 (6)
C13	0.0273 (8)	0.0232 (8)	0.0279 (8)	0.0102 (6)	0.0001 (6)	0.0061 (6)
C14	0.0236 (8)	0.0228 (8)	0.0276 (8)	0.0089 (6)	0.0006 (6)	0.0003 (6)
C15	0.0264 (8)	0.0189 (7)	0.0334 (8)	0.0103 (6)	0.0022 (6)	0.0032 (6)
C16	0.0248 (8)	0.0226 (8)	0.0289 (8)	0.0092 (6)	0.0021 (6)	0.0068 (6)
C17	0.0242 (7)	0.0215 (8)	0.0274 (8)	0.0078 (6)	-0.0002 (6)	0.0010 (6)
O2	0.0477 (7)	0.0272 (6)	0.0286 (6)	0.0197 (5)	0.0010 (5)	-0.0009 (5)
C18	0.0431 (10)	0.0292 (9)	0.0380 (9)	0.0204 (8)	0.0075 (8)	-0.0001 (7)
O3	0.0471 (7)	0.0255 (6)	0.0279 (6)	0.0162 (5)	-0.0027 (5)	0.0050 (5)
C19	0.0373 (9)	0.0333 (9)	0.0275 (8)	0.0111 (7)	0.0001 (7)	0.0047 (7)

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

C1—N1	1.3428 (19)	O1—C11	1.427 (2)
C1—N3	1.3453 (19)	C11—H11A	0.9700
C1—N2	1.3497 (19)	C11—H11B	0.9700
N1—C2	1.3395 (19)	C11—H11C	0.9700
C2—C3	1.392 (2)	C12—C13	1.379 (2)
C2—C5	1.492 (2)	C12—C17	1.400 (2)
C3—C4	1.388 (2)	C13—C14	1.394 (2)
C3—H3	0.9400	C13—H13	0.9400
C4—N2	1.3349 (19)	C14—O2	1.3702 (18)
C4—C12	1.489 (2)	C14—C15	1.382 (2)
N3—H3A	0.8700	C15—C16	1.391 (2)
N3—H3B	0.8700	C15—H15	0.9400
C5—C10	1.397 (2)	C16—O3	1.3704 (17)
C5—C6	1.406 (2)	C16—C17	1.382 (2)
C6—O1	1.3675 (19)	C17—H17	0.9400
C6—C7	1.390 (2)	O2—C18	1.4246 (18)
C7—C8	1.375 (2)	C18—H18A	0.9700

C7—H7	0.9400	C18—H18B	0.9700
C8—C9	1.374 (3)	C18—H18C	0.9700
C8—H8	0.9400	O3—C19	1.4199 (19)
C9—C10	1.378 (2)	C19—H19A	0.9700
C9—H9	0.9400	C19—H19B	0.9700
C10—H10	0.9400	C19—H19C	0.9700
N1—C1—N3	116.69 (13)	O1—C11—H11B	109.5
N1—C1—N2	126.16 (14)	H11A—C11—H11B	109.5
N3—C1—N2	117.15 (13)	O1—C11—H11C	109.5
C2—N1—C1	117.58 (13)	H11A—C11—H11C	109.5
N1—C2—C3	120.33 (13)	H11B—C11—H11C	109.5
N1—C2—C5	114.92 (13)	C13—C12—C17	120.67 (14)
C3—C2—C5	124.63 (14)	C13—C12—C4	120.77 (13)
C4—C3—C2	117.65 (14)	C17—C12—C4	118.48 (14)
C4—C3—H3	121.2	C12—C13—C14	119.40 (14)
C2—C3—H3	121.2	C12—C13—H13	120.3
N2—C4—C3	122.85 (14)	C14—C13—H13	120.3
N2—C4—C12	116.92 (13)	O2—C14—C15	123.43 (13)
C3—C4—C12	120.12 (13)	O2—C14—C13	115.83 (13)
C4—N2—C1	115.26 (13)	C15—C14—C13	120.73 (14)
C1—N3—H3A	120.0	C14—C15—C16	119.08 (14)
C1—N3—H3B	120.0	C14—C15—H15	120.5
H3A—N3—H3B	120.0	C16—C15—H15	120.5
C10—C5—C6	117.11 (14)	O3—C16—C17	124.54 (14)
C10—C5—C2	117.29 (14)	O3—C16—C15	114.35 (13)
C6—C5—C2	125.53 (14)	C17—C16—C15	121.11 (13)
O1—C6—C7	121.48 (14)	C16—C17—C12	118.91 (14)
O1—C6—C5	118.28 (13)	C16—C17—H17	120.5
C7—C6—C5	120.23 (15)	C12—C17—H17	120.5
C8—C7—C6	120.64 (16)	C14—O2—C18	116.94 (12)
C8—C7—H7	119.7	O2—C18—H18A	109.5
C6—C7—H7	119.7	O2—C18—H18B	109.5
C9—C8—C7	120.26 (16)	H18A—C18—H18B	109.5
C9—C8—H8	119.9	O2—C18—H18C	109.5
C7—C8—H8	119.9	H18A—C18—H18C	109.5
C8—C9—C10	119.40 (16)	H18B—C18—H18C	109.5
C8—C9—H9	120.3	C16—O3—C19	117.23 (12)
C10—C9—H9	120.3	O3—C19—H19A	109.5
C9—C10—C5	122.32 (16)	O3—C19—H19B	109.5
C9—C10—H10	118.8	H19A—C19—H19B	109.5
C5—C10—H10	118.8	O3—C19—H19C	109.5
C6—O1—C11	117.68 (13)	H19A—C19—H19C	109.5
O1—C11—H11A	109.5	H19B—C19—H19C	109.5
N3—C1—N1—C2	178.82 (14)	C6—C5—C10—C9	0.8 (2)
N2—C1—N1—C2	-1.4 (2)	C2—C5—C10—C9	-176.38 (15)
C1—N1—C2—C3	-2.6 (2)	C7—C6—O1—C11	3.7 (2)

C1—N1—C2—C5	173.46 (13)	C5—C6—O1—C11	−175.69 (15)
N1—C2—C3—C4	4.4 (2)	N2—C4—C12—C13	−140.95 (15)
C5—C2—C3—C4	−171.27 (14)	C3—C4—C12—C13	42.5 (2)
C2—C3—C4—N2	−2.4 (2)	N2—C4—C12—C17	42.3 (2)
C2—C3—C4—C12	173.89 (14)	C3—C4—C12—C17	−134.23 (16)
C3—C4—N2—C1	−1.2 (2)	C17—C12—C13—C14	0.9 (2)
C12—C4—N2—C1	−177.64 (13)	C4—C12—C13—C14	−175.80 (14)
N1—C1—N2—C4	3.3 (2)	C12—C13—C14—O2	−177.38 (13)
N3—C1—N2—C4	−176.92 (14)	C12—C13—C14—C15	2.2 (2)
N1—C2—C5—C10	−13.77 (19)	O2—C14—C15—C16	175.90 (14)
C3—C2—C5—C10	162.12 (14)	C13—C14—C15—C16	−3.6 (2)
N1—C2—C5—C6	169.36 (14)	C14—C15—C16—O3	−177.70 (14)
C3—C2—C5—C6	−14.8 (2)	C14—C15—C16—C17	2.1 (2)
C10—C5—C6—O1	177.62 (14)	O3—C16—C17—C12	−179.36 (14)
C2—C5—C6—O1	−5.5 (2)	C15—C16—C17—C12	0.9 (2)
C10—C5—C6—C7	−1.8 (2)	C13—C12—C17—C16	−2.4 (2)
C2—C5—C6—C7	175.08 (15)	C4—C12—C17—C16	174.36 (14)
O1—C6—C7—C8	−178.07 (16)	C15—C14—O2—C18	−4.5 (2)
C5—C6—C7—C8	1.3 (3)	C13—C14—O2—C18	175.04 (14)
C6—C7—C8—C9	0.3 (3)	C17—C16—O3—C19	−4.9 (2)
C7—C8—C9—C10	−1.3 (3)	C15—C16—O3—C19	174.89 (13)
C8—C9—C10—C5	0.8 (3)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N3—H3B···N2 ⁱ	0.87	2.28	3.1374 (19)	167
N3—H3A···O3 ⁱⁱ	0.87	2.39	3.2545 (17)	175

Symmetry codes: (i) $-x+3, -y+1, -z+2$; (ii) $x, y+1, z$.