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## Structure Reports

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# 3-Phenyl-2-(prop-2-ynoxy)-1-benzofuro[3,2-*d*]pyrimidin-4(3*H*)-one

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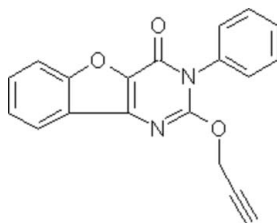
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 Key indicators: single-crystal X-ray study;  $T = 292$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.046;  $wR$  factor = 0.119; data-to-parameter ratio = 16.1.

In the title compound,  $\text{C}_{19}\text{H}_{12}\text{N}_2\text{O}_3$ , the 1-benzofuro[3,2-*d*]pyrimidinone unit is approximately planar, the maximum deviation from the mean plane being 0.045 (1) Å. The attached phenyl ring makes a dihedral angle of 86.73 (6)° with the fused ring system. The packing of the molecules in the crystal structure is mainly governed by  $\text{C}-\text{H}\cdots\pi$  hydrogen-bonding interactions.

## Related literature

For related preparation and biological activity, see: Bodke & Sangapure (2003). For related literature, see: Ding *et al.*, 2004. For the crystal structures of other fused pyrimidinone derivatives, see: Hu *et al.* (2005, 2006, 2007).



## Experimental

### Crystal data

$\text{C}_{19}\text{H}_{12}\text{N}_2\text{O}_3$   
 $M_r = 316.31$   
 Monoclinic,  $P2_1/n$   
 $a = 12.6748$  (13) Å  
 $b = 7.1531$  (7) Å  
 $c = 17.5793$  (17) Å  
 $\beta = 104.645$  (2)°

$V = 1542.0$  (3) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 292$  (2) K  
 $0.30 \times 0.20 \times 0.06$  mm

### Data collection

Bruker SMART 4K CCD area-detector diffractometer  
 Absorption correction: multi-scan (*SADABS*; Sheldrick, 2003)  
 $T_{\min} = 0.972$ ,  $T_{\max} = 0.994$   
 9340 measured reflections  
 3496 independent reflections  
 2681 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.041$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$   
 $wR(F^2) = 0.119$   
 $S = 1.02$   
 3496 reflections  
 217 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\text{max}} = 0.16$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.22$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> — <i>H</i> ··· <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> — <i>H</i> ··· <i>A</i>
C3—H3···Cg1 <sup>i</sup>	0.93	2.78	3.529 (1)	138
C4—H4···Cg2 <sup>ii</sup>	0.93	2.81	3.641 (1)	150
C11—H11A···Cg2 <sup>iii</sup>	0.97	2.76	3.402 (1)	124
C31—H13···Cg1 <sup>iv</sup>	0.93	2.77	3.521 (1)	139

Symmetry codes: (i)  $x + \frac{1}{2}, -y + \frac{3}{2}, z + \frac{1}{2}$ ; (ii)  $-x + \frac{5}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$ ; (iii)  $x, y - 1, z$ ; (iv)  $-x + \frac{3}{2}, y + \frac{1}{2}, -z + \frac{1}{2}$ . Cg1 is the centroid of the C14–C19 phenyl ring and Cg2 is the centroid of the C1–C6 ring.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2001).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT2655).

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## supporting information

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**3-Phenyl-2-(prop-2-ynyloxy)-1-benzofuro[3,2-*d*]pyrimidin-4(3*H*)-one****Yong-Nian Qu, Long-Rui Pan and Yang-Gen Hu****S1. Comment**

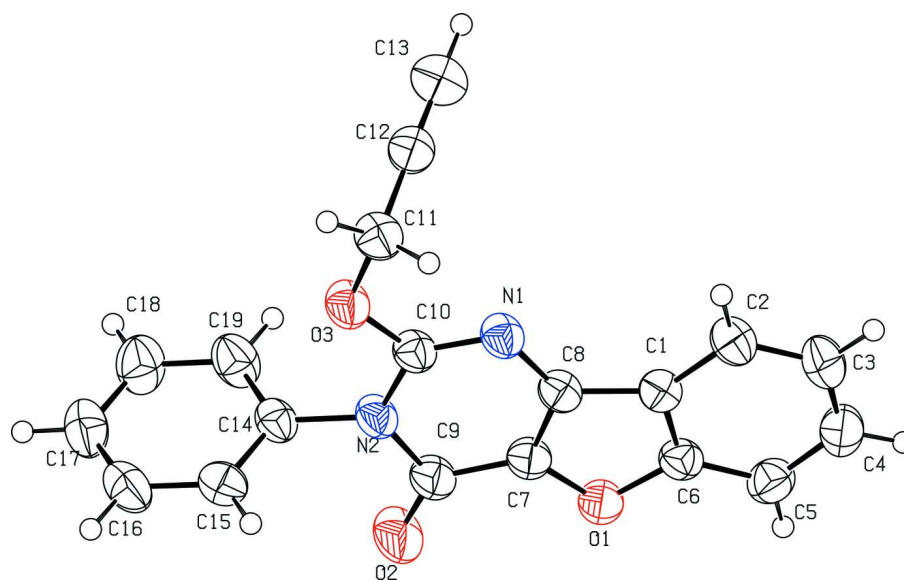
Benzofuopyrimidine derivatives are of interest as possible antiviral agents, and because of their other biological properties, including antibacterial, antifungal, antiallergic and antiinflammatory activities (Bodke & Sangapure, 2003). We have recently focused on the synthesis of the fused heterocyclic systems containing pyrimidinone *via* aza-Wittig reactions at room temperature (Ding *et al.*, 2004). We present here the structure of one such benzofuopyrimidinone derivative. Fig. 1 shows the molecular structure of (I) with the atomic numbering scheme. Bond lengths and angles are unexceptional (Hu *et al.*, 2005, 2006, 2007). The benzofuopyrimidine ring system is almost planar, with a maximum deviation of 0.045 (6) Å for atom C7; the C14—C19 phenyl ring is twisted with respect to it, with a dihedral angle of 86.73 (6)°. In the crystal, intermolecular C—H $\cdots$  $\pi$  hydrogen bonds (Table 1) stabilize the crystal structure (Fig. 2).

**S2. Experimental**

To a solution of ethyl 3-((phenylimino)methyleneamino)benzofuran-2-carboxylate (3 mmol) in dichloromethane (5 ml) was added sodium prop-2-yn-1-oxide (3 mmol) in prop-2-yn-1-ol (5 ml). After stirring the reaction mixture for 2 h, the solvent was removed under reduced pressure and the residue was recrystallized from ethanol to give the title compound, in a yield of 89%. Crystals suitable for single-crystal X-ray diffraction were obtained by recrystallization from a mixed solvent of ethanol and dichloromethane (1:1 *v/v*) at room temperature.

**S3. Refinement**

All C-bound H atoms were positioned geometrically, with C—H = 0.93 Å,  $U_{\text{iso}} = 1.2U_{\text{eq}}$  (C) for  $C_{\text{sp}}$  and  $C_{\text{sp}^2}$ , C—H = 0.97 Å,  $U_{\text{iso}} = 1.2U_{\text{eq}}$  (C) for  $\text{CH}_2$ .



**Figure 1**

The molecular structure of the title compound, showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 50% probability level.

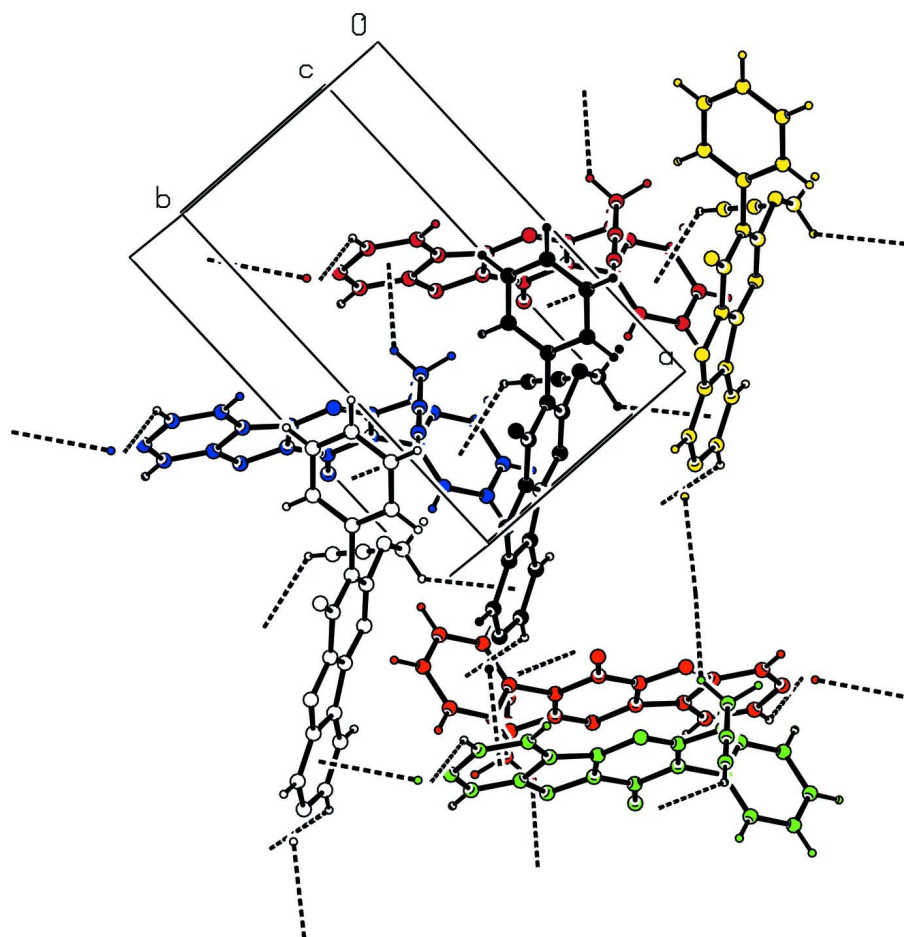


Figure 2

The packing in the crystal structure, showing C—H... $\pi$  hydrogen bonding interactions as dashed lines.

### 3-Phenyl-2-(prop-2-ynyloxy)-1-benzofuro[3,2-*d*]pyrimidin-4(3*H*)-one

#### Crystal data

$C_{19}H_{12}N_2O_3$

$M_r = 316.31$

Monoclinic,  $P2_1/n$

Hall symbol:  $-P\ 2_1n$

$a = 12.6748$  (13) Å

$b = 7.1531$  (7) Å

$c = 17.5793$  (17) Å

$\beta = 104.645$  (2)°

$V = 1542.0$  (3) Å<sup>3</sup>

$Z = 4$

$F(000) = 656$

$D_x = 1.362$  Mg m<sup>-3</sup>

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

Cell parameters from 3101 reflections

$\theta = 2.4$ – $27.4$ °

$\mu = 0.09$  mm<sup>-1</sup>

$T = 292$  K

Block, colorless

$0.30 \times 0.20 \times 0.06$  mm

#### Data collection

Bruker SMART 4K CCD area-detector  
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\varphi$  and  $\omega$  scans

Absorption correction: multi-scan  
(*SADABS*; Sheldrick, 2003)

$T_{\min} = 0.972$ ,  $T_{\max} = 0.994$

9340 measured reflections

3496 independent reflections

2681 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.041$   
 $\theta_{\text{max}} = 27.5^\circ$ ,  $\theta_{\text{min}} = 1.8^\circ$   
 $h = -8 \rightarrow 16$

$k = -9 \rightarrow 8$   
 $l = -22 \rightarrow 22$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.046$   
 $wR(F^2) = 0.119$   
 $S = 1.02$   
 3496 reflections  
 217 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0641P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.16 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.22 \text{ e } \text{\AA}^{-3}$

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	1.05495 (11)	0.81800 (18)	0.22716 (7)	0.0377 (3)
C2	1.11240 (12)	0.85724 (19)	0.30421 (8)	0.0455 (4)
H2	1.1116	0.7745	0.3449	0.055*
C3	1.17039 (12)	1.0220 (2)	0.31838 (9)	0.0510 (4)
H3	1.2099	1.0502	0.3693	0.061*
C4	1.17078 (13)	1.1476 (2)	0.25748 (10)	0.0522 (4)
H4	1.2099	1.2584	0.2691	0.063*
C5	1.11500 (12)	1.1122 (2)	0.18077 (9)	0.0482 (4)
H5	1.1156	1.1953	0.1402	0.058*
C6	1.05806 (11)	0.9455 (2)	0.16777 (7)	0.0403 (3)
C7	0.95602 (11)	0.7144 (2)	0.11079 (7)	0.0409 (3)
C8	0.98690 (11)	0.66798 (18)	0.18824 (7)	0.0380 (3)
C9	0.88390 (12)	0.6046 (2)	0.05294 (8)	0.0454 (4)
C10	0.88737 (11)	0.40745 (19)	0.16784 (7)	0.0397 (3)
C11	0.88258 (13)	0.1884 (2)	0.26800 (8)	0.0467 (4)
H11A	0.9608	0.2064	0.2860	0.056*
H11B	0.8678	0.0561	0.2716	0.056*
C12	0.82912 (12)	0.2919 (2)	0.31909 (8)	0.0456 (4)
C13	0.78913 (15)	0.3726 (2)	0.36209 (10)	0.0625 (5)
H13	0.7572	0.4371	0.3964	0.075*
C14	0.76773 (12)	0.32631 (18)	0.03895 (7)	0.0400 (3)
C15	0.79825 (13)	0.1850 (2)	-0.00388 (8)	0.0468 (4)

H15	0.8715	0.1633	-0.0009	0.056*
C16	0.71813 (15)	0.0755 (2)	-0.05158 (9)	0.0555 (4)
H16	0.7378	-0.0211	-0.0806	0.067*
C17	0.60970 (15)	0.1081 (2)	-0.05642 (9)	0.0583 (4)
H17	0.5563	0.0355	-0.0893	0.070*
C18	0.58085 (13)	0.2486 (3)	-0.01233 (10)	0.0612 (5)
H18	0.5077	0.2693	-0.0146	0.073*
C19	0.65962 (13)	0.3592 (2)	0.03527 (9)	0.0555 (4)
H19	0.6399	0.4551	0.0646	0.067*
N1	0.95387 (9)	0.50832 (16)	0.21918 (6)	0.0416 (3)
N2	0.85008 (9)	0.44489 (16)	0.08854 (6)	0.0413 (3)
O1	0.99762 (8)	0.88383 (13)	0.09524 (5)	0.0461 (3)
O2	0.85057 (11)	0.63467 (16)	-0.01731 (6)	0.0670 (4)
O3	0.84625 (9)	0.24665 (14)	0.18664 (5)	0.0495 (3)

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0361 (7)	0.0393 (7)	0.0367 (7)	0.0037 (6)	0.0071 (6)	0.0010 (5)
C2	0.0483 (8)	0.0460 (9)	0.0378 (7)	0.0050 (6)	0.0027 (6)	0.0011 (6)
C3	0.0485 (9)	0.0527 (9)	0.0455 (8)	0.0006 (7)	0.0003 (7)	-0.0085 (7)
C4	0.0470 (9)	0.0486 (9)	0.0597 (9)	-0.0061 (7)	0.0112 (7)	-0.0075 (7)
C5	0.0487 (9)	0.0489 (9)	0.0490 (8)	-0.0044 (7)	0.0161 (7)	0.0043 (7)
C6	0.0378 (7)	0.0472 (8)	0.0356 (7)	0.0017 (6)	0.0087 (6)	-0.0002 (6)
C7	0.0459 (8)	0.0434 (8)	0.0327 (7)	-0.0017 (6)	0.0086 (6)	0.0042 (6)
C8	0.0382 (7)	0.0410 (8)	0.0331 (7)	0.0049 (6)	0.0055 (6)	0.0014 (5)
C9	0.0517 (9)	0.0499 (9)	0.0326 (7)	-0.0026 (7)	0.0070 (6)	0.0031 (6)
C10	0.0450 (8)	0.0407 (7)	0.0319 (7)	0.0003 (6)	0.0070 (6)	0.0028 (6)
C11	0.0588 (9)	0.0415 (8)	0.0377 (7)	0.0021 (7)	0.0082 (7)	0.0091 (6)
C12	0.0475 (9)	0.0474 (9)	0.0390 (8)	-0.0010 (7)	0.0055 (6)	0.0067 (6)
C13	0.0673 (11)	0.0661 (11)	0.0560 (10)	0.0056 (9)	0.0194 (9)	-0.0024 (8)
C14	0.0468 (8)	0.0405 (8)	0.0295 (6)	0.0001 (6)	0.0040 (6)	0.0004 (5)
C15	0.0527 (9)	0.0495 (9)	0.0385 (7)	0.0036 (7)	0.0124 (7)	0.0015 (6)
C16	0.0757 (12)	0.0456 (9)	0.0444 (8)	0.0005 (8)	0.0135 (8)	-0.0092 (7)
C17	0.0628 (11)	0.0528 (10)	0.0509 (9)	-0.0088 (8)	-0.0011 (8)	-0.0049 (7)
C18	0.0433 (9)	0.0667 (11)	0.0666 (11)	0.0016 (8)	0.0009 (8)	-0.0073 (9)
C19	0.0515 (10)	0.0567 (10)	0.0551 (9)	0.0076 (7)	0.0075 (7)	-0.0141 (7)
N1	0.0489 (7)	0.0410 (7)	0.0319 (6)	-0.0009 (5)	0.0048 (5)	0.0028 (5)
N2	0.0475 (7)	0.0443 (7)	0.0290 (5)	-0.0030 (5)	0.0041 (5)	-0.0005 (5)
O1	0.0533 (6)	0.0497 (6)	0.0342 (5)	-0.0078 (5)	0.0087 (4)	0.0047 (4)
O2	0.0928 (9)	0.0695 (8)	0.0301 (5)	-0.0207 (7)	-0.0005 (5)	0.0078 (5)
O3	0.0649 (7)	0.0460 (6)	0.0339 (5)	-0.0101 (5)	0.0058 (5)	0.0040 (4)

*Geometric parameters (Å, °)*

C1—C6	1.3941 (19)	C10—N2	1.3804 (16)
C1—C2	1.3946 (18)	C11—O3	1.4486 (15)
C1—C8	1.4371 (18)	C11—C12	1.457 (2)

C2—C3	1.378 (2)	C11—H11A	0.9700
C2—H2	0.9300	C11—H11B	0.9700
C3—C4	1.398 (2)	C12—C13	1.164 (2)
C3—H3	0.9300	C13—H13	0.9300
C4—C5	1.379 (2)	C14—C15	1.3731 (19)
C4—H4	0.9300	C14—C19	1.376 (2)
C5—C6	1.383 (2)	C14—N2	1.4522 (17)
C5—H5	0.9300	C15—C16	1.385 (2)
C6—O1	1.3832 (15)	C15—H15	0.9300
C7—C8	1.3593 (18)	C16—C17	1.375 (2)
C7—O1	1.3766 (16)	C16—H16	0.9300
C7—C9	1.4200 (19)	C17—C18	1.374 (2)
C8—N1	1.3749 (16)	C17—H17	0.9300
C9—O2	1.2190 (16)	C18—C19	1.378 (2)
C9—N2	1.4199 (18)	C18—H18	0.9300
C10—N1	1.2888 (17)	C19—H19	0.9300
C10—O3	1.3382 (16)		
C6—C1—C2	119.28 (13)	O3—C11—H11A	109.1
C6—C1—C8	105.01 (11)	C12—C11—H11A	109.1
C2—C1—C8	135.71 (12)	O3—C11—H11B	109.1
C3—C2—C1	118.12 (13)	C12—C11—H11B	109.1
C3—C2—H2	120.9	H11A—C11—H11B	107.9
C1—C2—H2	120.9	C13—C12—C11	177.69 (16)
C2—C3—C4	121.05 (14)	C12—C13—H13	180.0
C2—C3—H3	119.5	C15—C14—C19	121.04 (13)
C4—C3—H3	119.5	C15—C14—N2	119.98 (13)
C5—C4—C3	122.07 (14)	C19—C14—N2	118.98 (12)
C5—C4—H4	119.0	C14—C15—C16	118.90 (15)
C3—C4—H4	119.0	C14—C15—H15	120.6
C4—C5—C6	115.92 (13)	C16—C15—H15	120.6
C4—C5—H5	122.0	C17—C16—C15	120.66 (14)
C6—C5—H5	122.0	C17—C16—H16	119.7
C5—C6—O1	124.85 (12)	C15—C16—H16	119.7
C5—C6—C1	123.55 (12)	C18—C17—C16	119.57 (15)
O1—C6—C1	111.59 (12)	C18—C17—H17	120.2
C8—C7—O1	112.72 (12)	C16—C17—H17	120.2
C8—C7—C9	123.55 (13)	C17—C18—C19	120.47 (16)
O1—C7—C9	123.65 (12)	C17—C18—H18	119.8
C7—C8—N1	124.03 (12)	C19—C18—H18	119.8
C7—C8—C1	106.30 (12)	C14—C19—C18	119.36 (14)
N1—C8—C1	129.65 (12)	C14—C19—H19	120.3
O2—C9—N2	121.50 (13)	C18—C19—H19	120.3
O2—C9—C7	128.57 (14)	C10—N1—C8	113.50 (11)
N2—C9—C7	109.92 (11)	C10—N2—C9	122.46 (11)
N1—C10—O3	122.34 (12)	C10—N2—C14	120.23 (11)
N1—C10—N2	126.51 (13)	C9—N2—C14	117.18 (10)
O3—C10—N2	111.15 (11)	C7—O1—C6	104.37 (10)

O3—C11—C12	112.40 (12)	C10—O3—C11	116.41 (11)
C6—C1—C2—C3	0.0 (2)	C16—C17—C18—C19	1.4 (3)
C8—C1—C2—C3	179.90 (15)	C15—C14—C19—C18	-0.1 (2)
C1—C2—C3—C4	0.7 (2)	N2—C14—C19—C18	179.27 (14)
C2—C3—C4—C5	-0.9 (2)	C17—C18—C19—C14	-0.7 (3)
C3—C4—C5—C6	0.4 (2)	O3—C10—N1—C8	179.94 (12)
C4—C5—C6—O1	-179.66 (13)	N2—C10—N1—C8	0.8 (2)
C4—C5—C6—C1	0.3 (2)	C7—C8—N1—C10	-1.97 (19)
C2—C1—C6—C5	-0.5 (2)	C1—C8—N1—C10	176.24 (13)
C8—C1—C6—C5	179.57 (13)	N1—C10—N2—C9	1.1 (2)
C2—C1—C6—O1	179.48 (12)	O3—C10—N2—C9	-178.16 (12)
C8—C1—C6—O1	-0.48 (14)	N1—C10—N2—C14	-174.64 (13)
O1—C7—C8—N1	178.20 (12)	O3—C10—N2—C14	6.10 (17)
C9—C7—C8—N1	1.4 (2)	O2—C9—N2—C10	179.54 (13)
O1—C7—C8—C1	-0.37 (16)	C7—C9—N2—C10	-1.62 (19)
C9—C7—C8—C1	-177.17 (13)	O2—C9—N2—C14	-4.6 (2)
C6—C1—C8—C7	0.50 (14)	C7—C9—N2—C14	174.25 (12)
C2—C1—C8—C7	-179.45 (15)	C15—C14—N2—C10	-96.71 (16)
C6—C1—C8—N1	-177.95 (13)	C19—C14—N2—C10	83.87 (17)
C2—C1—C8—N1	2.1 (3)	C15—C14—N2—C9	87.32 (15)
C8—C7—C9—O2	179.20 (15)	C19—C14—N2—C9	-92.09 (16)
O1—C7—C9—O2	2.7 (3)	C8—C7—O1—C6	0.08 (15)
C8—C7—C9—N2	0.5 (2)	C9—C7—O1—C6	176.88 (13)
O1—C7—C9—N2	-176.00 (12)	C5—C6—O1—C7	-179.79 (13)
O3—C11—C12—C13	-164 (4)	C1—C6—O1—C7	0.26 (14)
C19—C14—C15—C16	0.3 (2)	N1—C10—O3—C11	-1.5 (2)
N2—C14—C15—C16	-179.10 (12)	N2—C10—O3—C11	177.81 (11)
C14—C15—C16—C17	0.4 (2)	C12—C11—O3—C10	77.20 (16)
C15—C16—C17—C18	-1.2 (2)		

*Hydrogen-bond geometry (Å, °)*

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C3—H3...Cg1 <sup>i</sup>	0.93	2.78	3.529 (1)	138
C4—H4...Cg2 <sup>ii</sup>	0.93	2.81	3.641 (1)	150
C11—H11A...Cg2 <sup>iii</sup>	0.97	2.76	3.402 (1)	124
C31—H13...Cg1 <sup>iv</sup>	0.93	2.77	3.521 (1)	139

Symmetry codes: (i)  $x+1/2, -y+3/2, z+1/2$ ; (ii)  $-x+5/2, y+1/2, -z+1/2$ ; (iii)  $x, y-1, z$ ; (iv)  $-x+3/2, y+1/2, -z+1/2$ .