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## Structure Reports

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# N-(1,4-Dioxo-1,4-dihydronaphthalen-2-yl)benzamide

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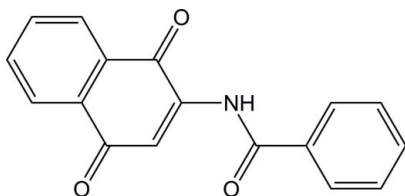
Received 2 July 2012; accepted 31 July 2012

 Key indicators: single-crystal X-ray study;  $T = 123$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.052;  $wR$  factor = 0.148; data-to-parameter ratio = 13.4.

The title compound,  $\text{C}_{17}\text{H}_{11}\text{NO}_3$ , was an intermediate synthesized during bisacylation of 2-amino-1,4-naphthoquinone with benzoyl chloride. A mixture of block- and needle-shaped crystals were obtained after column chromatography. The block-shaped crystals were identified as the imide and the needles were the title amide. The naphthoquinone scaffold is roughly planar (r.m.s. deviation = 0.047 Å for the C atoms). The N—H and C=O bonds of the amide group are *anti* to each other. A dihedral angle between the naphthoquinone ring system and the amide group of 3.56 (3)°, accompanied by a dihedral angle between the amide group and the phenyl group of 9.51 (3)°, makes the naphthoquinone ring essentially coplanar with the phenyl ring [dihedral angle = 7.12 (1)°]. In the crystal, molecules are linked by a weak N—H···O hydrogen bond and by two weak C—H···O interactions leading to the formation of zigzag chains along [010].

## Related literature

For similar crystal structures, see: Brandy *et al.* (2009, 2012); Akinboye *et al.* (2009a,b). For the pharmacological properties of related compounds, see: Bakare *et al.* (2003); Berhe *et al.* (2008); Lien *et al.* (1997); Huang, *et al.* (2005); Khraiweh *et al.* (2011).



## Experimental

### Crystal data

 $\text{C}_{17}\text{H}_{11}\text{NO}_3$ 
 $M_r = 277.27$ 

 Monoclinic,  $P2_1/n$ 
 $a = 6.9433$  (3) Å

 $b = 12.0112$  (4) Å

 $c = 15.2129$  (5) Å

 $\beta = 94.129$  (3)°

 $V = 1265.42$  (8) Å<sup>3</sup>
 $Z = 4$ 

 Cu  $K\alpha$  radiation

 $\mu = 0.83$  mm<sup>-1</sup>
 $T = 123$  K

 $0.67 \times 0.12 \times 0.08$  mm

### Data collection

Oxford Diffraction Xcalibur Ruby

Gemini diffractometer

Absorption correction: multi-scan

 (*CrysAlis PRO*; Oxford

Diffraction, 2007)

 $T_{\min} = 0.819$ ,  $T_{\max} = 1.000$ 

4550 measured reflections

2553 independent reflections

 2123 reflections with  $I > 2\sigma(I)$ 
 $R_{\text{int}} = 0.037$ 

### Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.052$ 
 $wR(F^2) = 0.148$ 
 $S = 1.05$ 

2553 reflections

190 parameters

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.25$  e Å<sup>-3</sup>
 $\Delta\rho_{\text{min}} = -0.25$  e Å<sup>-3</sup>
**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1A}\cdots\text{O2}^i$	0.88	2.65	3.3299 (19)	135
$\text{C4}-\text{H4A}\cdots\text{O3}^i$	0.95	2.49	3.149 (2)	127
$\text{C17}-\text{H17A}\cdots\text{O2}^i$	0.95	2.57	3.404 (2)	146

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2007); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

RJB wishes to acknowledge the NSF-MRI program (grant CHE-0619278) for funds to purchase the diffractometer. We also acknowledge MRI grant No. CHE-1126533 from the National Science Foundation for the purchase of a TOF LC/MS system used in this study and also funded in part by grant No. 5-U54-CA914-31 (Howard University/Johns Hopkins Cancer Center Partnership).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5965).

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## supplementary materials

*Acta Cryst.* (2012). E68, o2775–o2776 [doi:10.1107/S1600536812034150]

***N*-(1,4-Dioxo-1,4-dihydronaphthalen-2-yl)benzamide****Yakini Brandy, Ray J. Butcher and Oladapo Bakare****Comment**

Amido and imidonaphthoquinones are known for their anti-inflammatory, antiplatelet, antiallergic, antiparasitic and anticancer activities (Lien *et al.* (1997); Huang *et al.* (2005); Bakare *et al.* (2003), Khraiwesh *et al.* (2011)). During imide synthesis (2-*N*-bis(benzoyl)amino-1,4-naphthoquinone (Brandy *et al.* (2012)), we obtained the intermediate amido analog, 2-*N*-benzoylamino-1,4-naphthoquinone. In order to eventually perform an anticancer SAR (structure-activity relationship) study, these intermediate amido analogs were isolated, crystallized and subjected to an X-ray diffraction study.

This showed that the naphthoquinone scaffold was planar with N—H and C=O bonds anti to each other. A dihedral angle between the naphthoquinone ring and the amide group of  $-2.6(3)^\circ$ , accompanied with the dihedral angle between the amide group and the phenyl group of  $-8.8(2)^\circ$  makes the naphthoquinone ring coplanar to the phenyl group. The bond distances and angles are similar to those found in related structures (Brandy *et al.*, 2009, 2012; Akinboye *et al.*, 2009a, 2009b). The crystal packing pattern results from N—H $\cdots$ O hydrogen bonds along with two weak intermolecular C—H $\cdots$ O interactions.

**Experimental**

2-Amino-1,4-naphthoquinone (318 mg, 1.83 mmol) was dissolved in freshly distilled THF (15 ml). NaH (115 mg, 4.78 mmol) was added and the mixture was stirred at room temperature for 15 min. The appropriate benzoyl chloride (0.55 ml, 4.74 mmol) was added, drop wise, and the mixture was stirred for 24 h. THF was evaporated under vacuum and the mixture was washed with ice-water (10 g ice in 10 ml water). The ice-water mixture was extracted with CH<sub>2</sub>Cl<sub>2</sub> (30 ml, 20 ml consecutively) and the combined organic phase washed with water (3 x 20 ml), saturated NaCl solution (20 ml), then dried over anhydrous MgSO<sub>4</sub>. The crude was purified *via* triturating in ethanol (2 ml) and column chromatography with an eluent mixture of ethyl acetate and hexane to furnish the amide (39 mg, 7.7%). A mixture of block and needle crystals were obtained from column chromatography. The block crystals were identified as the imide (Brandy *et al.*, 2012) and the needles were identified as the amide. The needle crystals were hand-picked from the mixture and analyzed by X-ray diffraction.

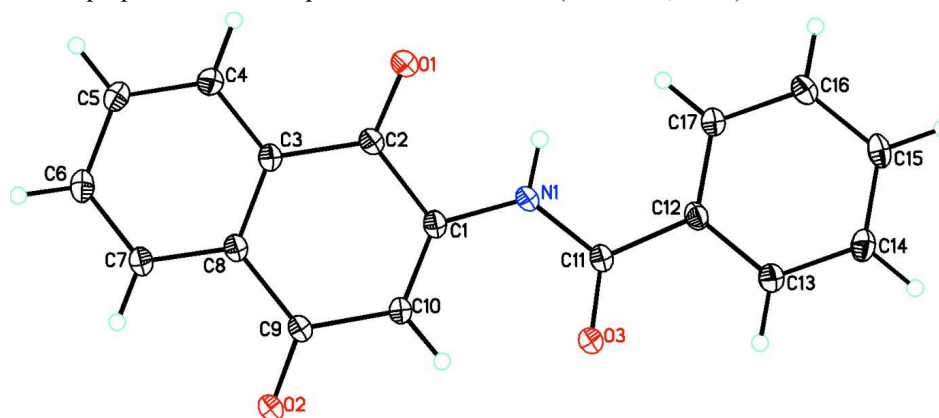
**Refinement**

H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with N—H = 0.88 Å and C—H = 0.95 Å and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ .

**Computing details**

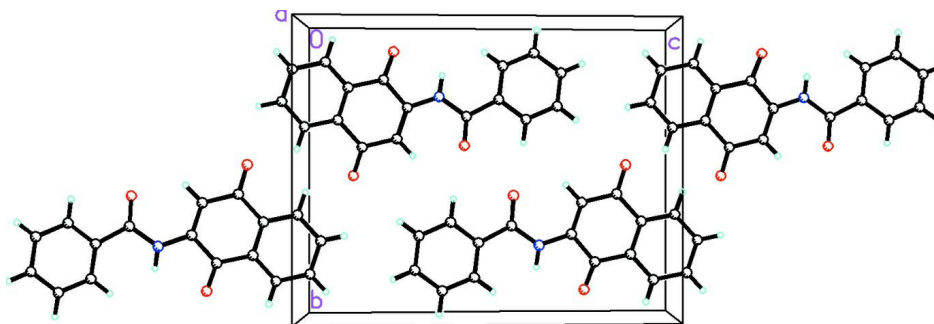
Data collection: *CrysAlis PRO* (Oxford Diffraction, 2007); cell refinement: *CrysAlis PRO* (Oxford Diffraction, 2007); data reduction: *CrysAlis PRO* (Oxford Diffraction, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick,

2008); software used to prepare material for publication: *SHELXTL* (Sheldrick, 2008).



**Figure 1**

Diagram of  $C_{17}H_{11}NO_3$  showing the atom labeling. Displacement ellipsoids are at the 30% probability level.



**Figure 2**

The molecular packing for  $C_{17}H_{11}NO_3$  viewed along the  $a$  axis.

### *N*-(1,4-Dioxo-1,4-dihydronaphthalen-2-yl)benzamide

#### Crystal data

$C_{17}H_{11}NO_3$

$M_r = 277.27$

Monoclinic,  $P2_1/n$

$a = 6.9433$  (3) Å

$b = 12.0112$  (4) Å

$c = 15.2129$  (5) Å

$\beta = 94.129$  (3)°

$V = 1265.42$  (8) Å<sup>3</sup>

$Z = 4$

$F(000) = 576$

$D_x = 1.455$  Mg m<sup>-3</sup>

Cu  $K\alpha$  radiation,  $\lambda = 1.54178$  Å

Cell parameters from 1585 reflections

$\theta = 2.9\text{--}75.6^\circ$

$\mu = 0.83$  mm<sup>-1</sup>

$T = 123$  K

Needle, pale yellow orange

$0.67 \times 0.12 \times 0.08$  mm

#### Data collection

Oxford Diffraction Xcalibur Ruby Gemini diffractometer

Radiation source: Enhance (Cu) X-ray Source

Graphite monochromator

Detector resolution: 10.5081 pixels mm<sup>-1</sup>

$\omega$  scans

Absorption correction: multi-scan

(*CrysAlis PRO*; Oxford Diffraction, 2007)

$T_{\min} = 0.819$ ,  $T_{\max} = 1.000$

4550 measured reflections

2553 independent reflections

2123 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.037$

$\theta_{\max} = 75.7^\circ$ ,  $\theta_{\min} = 4.7^\circ$   
 $h = -7 \rightarrow 8$

$k = -14 \rightarrow 14$   
 $l = -18 \rightarrow 18$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.052$   
 $wR(F^2) = 0.148$   
 $S = 1.05$   
 2553 reflections  
 190 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0937P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.25 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.25 \text{ e } \text{\AA}^{-3}$

*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>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.4411 (2)	0.09807 (11)	0.24463 (8)	0.0373 (3)
O2	0.3847 (2)	0.51857 (10)	0.13486 (8)	0.0348 (3)
O3	0.3674 (2)	0.41588 (11)	0.43416 (8)	0.0354 (3)
N1	0.3825 (2)	0.24883 (12)	0.36581 (9)	0.0273 (3)
H1A	0.3821	0.1763	0.3741	0.033*
C1	0.3905 (2)	0.28621 (14)	0.27989 (10)	0.0253 (4)
C2	0.4119 (2)	0.19196 (14)	0.21661 (10)	0.0269 (4)
C3	0.3957 (2)	0.21822 (14)	0.12102 (10)	0.0252 (4)
C4	0.3886 (2)	0.13177 (15)	0.05992 (11)	0.0299 (4)
H4A	0.3960	0.0566	0.0793	0.036*
C5	0.3707 (2)	0.15588 (15)	-0.02938 (11)	0.0314 (4)
H5A	0.3629	0.0971	-0.0713	0.038*
C6	0.3641 (2)	0.26586 (16)	-0.05777 (11)	0.0320 (4)
H6A	0.3547	0.2820	-0.1191	0.038*
C7	0.3713 (2)	0.35219 (15)	0.00297 (11)	0.0290 (4)
H7A	0.3670	0.4272	-0.0168	0.035*
C8	0.3848 (2)	0.32888 (14)	0.09291 (10)	0.0251 (4)
C9	0.3872 (2)	0.42102 (14)	0.15843 (10)	0.0266 (4)
C10	0.3855 (2)	0.39248 (14)	0.25204 (10)	0.0270 (4)
H10A	0.3807	0.4506	0.2942	0.032*
C11	0.3750 (2)	0.31531 (14)	0.43956 (10)	0.0261 (3)
C12	0.3814 (2)	0.25771 (14)	0.52731 (10)	0.0252 (4)
C13	0.4058 (2)	0.32638 (15)	0.60106 (11)	0.0294 (4)
H13A	0.4142	0.4047	0.5938	0.035*

C14	0.4181 (3)	0.28079 (16)	0.68528 (11)	0.0324 (4)
H14A	0.4353	0.3279	0.7354	0.039*
C15	0.4050 (2)	0.16642 (16)	0.69612 (11)	0.0316 (4)
H15A	0.4137	0.1352	0.7536	0.038*
C16	0.3794 (2)	0.09777 (15)	0.62296 (11)	0.0308 (4)
H16A	0.3694	0.0195	0.6306	0.037*
C17	0.3681 (2)	0.14279 (15)	0.53839 (11)	0.0286 (4)
H17A	0.3514	0.0954	0.4884	0.034*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0618 (8)	0.0246 (7)	0.0265 (6)	0.0064 (6)	0.0101 (5)	0.0045 (5)
O2	0.0576 (7)	0.0236 (6)	0.0238 (6)	-0.0019 (5)	0.0080 (5)	0.0045 (5)
O3	0.0604 (8)	0.0235 (6)	0.0221 (6)	0.0007 (5)	0.0030 (5)	0.0025 (5)
N1	0.0412 (7)	0.0216 (7)	0.0196 (7)	0.0017 (5)	0.0047 (5)	0.0045 (5)
C1	0.0299 (7)	0.0271 (8)	0.0192 (7)	0.0005 (6)	0.0037 (5)	0.0022 (6)
C2	0.0352 (7)	0.0234 (8)	0.0226 (8)	0.0011 (6)	0.0057 (6)	0.0038 (6)
C3	0.0302 (7)	0.0258 (8)	0.0202 (7)	-0.0002 (6)	0.0059 (5)	0.0011 (6)
C4	0.0380 (8)	0.0253 (8)	0.0271 (8)	-0.0008 (6)	0.0084 (6)	-0.0007 (6)
C5	0.0392 (8)	0.0315 (9)	0.0241 (8)	0.0001 (7)	0.0060 (6)	-0.0057 (7)
C6	0.0365 (8)	0.0393 (10)	0.0208 (7)	0.0005 (7)	0.0057 (6)	0.0008 (7)
C7	0.0369 (8)	0.0294 (9)	0.0211 (8)	0.0007 (7)	0.0054 (6)	0.0035 (6)
C8	0.0288 (7)	0.0260 (9)	0.0208 (7)	0.0001 (6)	0.0042 (5)	0.0025 (6)
C9	0.0349 (7)	0.0239 (8)	0.0214 (7)	-0.0006 (6)	0.0043 (6)	0.0037 (6)
C10	0.0378 (8)	0.0250 (8)	0.0184 (7)	-0.0014 (6)	0.0041 (6)	0.0010 (6)
C11	0.0325 (7)	0.0265 (8)	0.0195 (7)	0.0004 (6)	0.0027 (6)	0.0029 (6)
C12	0.0289 (7)	0.0270 (8)	0.0201 (8)	0.0007 (6)	0.0037 (6)	0.0046 (6)
C13	0.0373 (8)	0.0286 (9)	0.0227 (8)	0.0001 (6)	0.0045 (6)	0.0017 (6)
C14	0.0403 (9)	0.0382 (10)	0.0191 (7)	-0.0006 (7)	0.0048 (6)	-0.0015 (6)
C15	0.0351 (8)	0.0412 (10)	0.0187 (7)	0.0001 (7)	0.0032 (6)	0.0077 (7)
C16	0.0390 (8)	0.0287 (9)	0.0247 (8)	-0.0003 (7)	0.0034 (6)	0.0070 (7)
C17	0.0373 (8)	0.0282 (9)	0.0202 (7)	0.0002 (6)	0.0024 (6)	0.0017 (6)

*Geometric parameters (Å, °)*

O1—C2	1.217 (2)	C7—C8	1.393 (2)
O2—C9	1.225 (2)	C7—H7A	0.9500
O3—C11	1.212 (2)	C8—C9	1.489 (2)
N1—C11	1.381 (2)	C9—C10	1.466 (2)
N1—C1	1.3869 (19)	C10—H10A	0.9500
N1—H1A	0.8800	C11—C12	1.501 (2)
C1—C10	1.345 (2)	C12—C13	1.393 (2)
C1—C2	1.500 (2)	C12—C17	1.394 (2)
C2—C3	1.484 (2)	C13—C14	1.390 (2)
C3—C4	1.392 (2)	C13—H13A	0.9500
C3—C8	1.397 (2)	C14—C15	1.387 (3)
C4—C5	1.386 (2)	C14—H14A	0.9500
C4—H4A	0.9500	C15—C16	1.386 (3)
C5—C6	1.390 (3)	C15—H15A	0.9500

C5—H5A	0.9500	C16—C17	1.393 (2)
C6—C7	1.387 (2)	C16—H16A	0.9500
C6—H6A	0.9500	C17—H17A	0.9500
C11—N1—C1	125.79 (14)	O2—C9—C10	120.48 (15)
C11—N1—H1A	117.1	O2—C9—C8	121.05 (14)
C1—N1—H1A	117.1	C10—C9—C8	118.44 (14)
C10—C1—N1	127.04 (15)	C1—C10—C9	121.76 (15)
C10—C1—C2	121.04 (14)	C1—C10—H10A	119.1
N1—C1—C2	111.90 (14)	C9—C10—H10A	119.1
O1—C2—C3	122.60 (16)	O3—C11—N1	121.69 (14)
O1—C2—C1	119.73 (14)	O3—C11—C12	121.20 (15)
C3—C2—C1	117.67 (14)	N1—C11—C12	117.10 (15)
C4—C3—C8	120.45 (15)	C13—C12—C17	119.61 (14)
C4—C3—C2	119.49 (15)	C13—C12—C11	115.94 (15)
C8—C3—C2	120.05 (15)	C17—C12—C11	124.45 (15)
C5—C4—C3	119.67 (16)	C14—C13—C12	120.32 (16)
C5—C4—H4A	120.2	C14—C13—H13A	119.8
C3—C4—H4A	120.2	C12—C13—H13A	119.8
C4—C5—C6	120.13 (16)	C15—C14—C13	119.96 (16)
C4—C5—H5A	119.9	C15—C14—H14A	120.0
C6—C5—H5A	119.9	C13—C14—H14A	120.0
C7—C6—C5	120.31 (15)	C16—C15—C14	119.95 (15)
C7—C6—H6A	119.8	C16—C15—H15A	120.0
C5—C6—H6A	119.8	C14—C15—H15A	120.0
C6—C7—C8	120.04 (16)	C15—C16—C17	120.41 (17)
C6—C7—H7A	120.0	C15—C16—H16A	119.8
C8—C7—H7A	120.0	C17—C16—H16A	119.8
C7—C8—C3	119.36 (16)	C16—C17—C12	119.75 (16)
C7—C8—C9	120.32 (15)	C16—C17—H17A	120.1
C3—C8—C9	120.32 (14)	C12—C17—H17A	120.1
C11—N1—C1—C10	-2.6 (3)	C3—C8—C9—O2	177.33 (15)
C11—N1—C1—C2	175.82 (14)	C7—C8—C9—C10	175.06 (14)
C10—C1—C2—O1	170.53 (16)	C3—C8—C9—C10	-4.8 (2)
N1—C1—C2—O1	-8.0 (2)	N1—C1—C10—C9	-177.44 (15)
C10—C1—C2—C3	-9.6 (2)	C2—C1—C10—C9	4.3 (2)
N1—C1—C2—C3	171.88 (13)	O2—C9—C10—C1	-179.15 (15)
O1—C2—C3—C4	8.5 (2)	C8—C9—C10—C1	3.0 (2)
C1—C2—C3—C4	-171.42 (14)	C1—N1—C11—O3	2.9 (3)
O1—C2—C3—C8	-172.50 (15)	C1—N1—C11—C12	-175.70 (14)
C1—C2—C3—C8	7.6 (2)	O3—C11—C12—C13	-8.8 (2)
C8—C3—C4—C5	0.1 (2)	N1—C11—C12—C13	169.78 (14)
C2—C3—C4—C5	179.07 (15)	O3—C11—C12—C17	172.36 (16)
C3—C4—C5—C6	1.4 (3)	N1—C11—C12—C17	-9.0 (2)
C4—C5—C6—C7	-1.4 (2)	C17—C12—C13—C14	0.4 (2)
C5—C6—C7—C8	-0.1 (2)	C11—C12—C13—C14	-178.52 (15)
C6—C7—C8—C3	1.6 (2)	C12—C13—C14—C15	-0.3 (2)
C6—C7—C8—C9	-178.34 (14)	C13—C14—C15—C16	-0.2 (3)

C4—C3—C8—C7	-1.5 (2)	C14—C15—C16—C17	0.5 (3)
C2—C3—C8—C7	179.46 (14)	C15—C16—C17—C12	-0.5 (2)
C4—C3—C8—C9	178.37 (14)	C13—C12—C17—C16	0.0 (2)
C2—C3—C8—C9	-0.6 (2)	C11—C12—C17—C16	178.79 (15)
C7—C8—C9—O2	-2.8 (2)		

*Hydrogen-bond geometry (Å, °)*

<i>D—H</i> ⋯ <i>A</i>	<i>D—H</i>	<i>H</i> ⋯ <i>A</i>	<i>D</i> ⋯ <i>A</i>	<i>D—H</i> ⋯ <i>A</i>
N1—H1 <i>A</i> ⋯O2 <sup>i</sup>	0.88	2.65	3.3299 (19)	135
C4—H4 <i>A</i> ⋯O3 <sup>i</sup>	0.95	2.49	3.149 (2)	127
C17—H17 <i>A</i> ⋯O2 <sup>i</sup>	0.95	2.57	3.404 (2)	146

Symmetry code: (i)  $-x+1/2, y-1/2, -z+1/2$ .