

Crystal structure of 8-ethoxy-3-(4-nitrophenyl)-2H-chromen-2-one

Shashikanth Walki,^a S. Naveen,^b S. Kenchanna,^a
K. M. Mahadevan,^a M. N. Kumara^c and N. K. Lokanath^{d*}

^aDepartment of Chemistry, Kuvempu University, P. G. Centre, Kadur 577 548, India, ^bInstitution of Excellence, University of Mysore, Manasagangotri, Mysore 570 006, India, ^cDepartment of Chemistry, Yuvaraja's College, University of Mysore, Mysore 570 005, India, and ^dDepartment of Studies in Physics, University of Mysore, Manasagangotri, Mysore 570 006, India. *Correspondence e-mail: lokanath@physics.uni-mysore.ac.in

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In the title compound, C₁₇H₁₃NO₅, the coumarin ring system is essentially planar (r.m.s. deviation = 0.008 Å). The nitrophenyl ring makes a dihedral angle of 25.27 (9)° with the coumarin ring plane. The nitro group is almost coplanar with the phenyl ring to which it is attached, making a dihedral angle of 4.3 (3)°. The ethoxy group is inclined to the coumarin ring plane by 4.1 (2)°. Electron delocalization was found at the short bridging C—C bond with a bond length of 1.354 (2) Å. In the crystal, molecules are linked *via* C—H...O hydrogen bonds, forming sheets in the *bc* plane. The sheets are linked *via* π–π stacking [centroid–centroid distances = 3.5688 (13) and 3.7514 (13) Å], forming a three-dimensional structure.

Keywords: crystal structure; coumarin; chromen; C—H...O hydrogen bonds; π–π stacking.

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1. Related literature

For coumarin derivatives as fluorescent brighteners, see: Tian *et al.* (2000). For details of natural or synthetic coumarins which inhibit lipid peroxidation and scavenge hydroxyl radicals and superoxide anions, see: Naveen *et al.* (2007). For further details of our research on coumarins, see: Naveen *et al.* (2006*a,b*).

2. Experimental

2.1. Crystal data

C ₁₇ H ₁₃ NO ₅	<i>V</i> = 1481.7 (3) Å ³
<i>M_r</i> = 311.28	<i>Z</i> = 4
Orthorhombic, <i>P</i> 2 ₁ 2 ₁ 2 ₁	Cu <i>K</i> α radiation
<i>a</i> = 6.8118 (9) Å	<i>μ</i> = 0.87 mm ⁻¹
<i>b</i> = 13.6726 (18) Å	<i>T</i> = 296 K
<i>c</i> = 15.909 (2) Å	0.29 × 0.26 × 0.21 mm

2.2. Data collection

Bruker X8 Proteum diffractometer	6729 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2013)	2371 independent reflections
<i>T</i> _{min} = 0.786, <i>T</i> _{max} = 0.838	2225 reflections with <i>I</i> > 2σ(<i>I</i>)
	<i>R</i> _{int} = 0.040

2.3. Refinement

<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.045	Δρ _{max} = 0.25 e Å ⁻³
<i>wR</i> (<i>F</i> ²) = 0.131	Δρ _{min} = -0.21 e Å ⁻³
<i>S</i> = 1.03	Absolute structure: 957 Friedel pairs; Flack (1983)
2371 reflections	Absolute structure parameter: 0.1 (2)
210 parameters	
H-atom parameters constrained	

Table 1

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>
C4—H4...O22 ⁱ	0.93	2.53	3.453 (2)	171
C8—H8...O14 ⁱⁱ	0.93	2.31	3.226 (2)	166
C20—H20...O22 ⁱ	0.93	2.52	3.275 (3)	138

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

Data collection: *APEX2* (Bruker, 2013); cell refinement: *SAINTE* (Bruker, 2013); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *SHELXL97* and *PLATON* (Spek, 2009).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: SU5224).

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supporting information

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Crystal structure of 8-ethoxy-3-(4-nitrophenyl)-2*H*-chromen-2-one

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S1. Comment

Coumarin derivatives are found to be one of the major groups of compounds used as fluorescent brighteners (Tian *et al.*, 2000) or fluorescent dyes and they have exhibited good bleed fastness and durability for coating plastics or in acrylic lacquers. Several natural or synthetic coumarins with various hydroxyl and other substituents were found to inhibit lipid peroxidation and to scavenge hydroxyl radicals and superoxide anions (Naveen *et al.*, 2007). As a part of our ongoing research on coumarins (Naveen *et al.*, 2006a,b), we report herein on the synthesis, characterization and crystal structure of the title compound. The compound is being assessed for biological activity.

The molecular structure of the title compound is shown in Fig. 1. The coumarin ring is essentially planar with the two axially fused rings forming a dihedral angle of 0.45 (10)°, while the 4-nitrophenyl ring makes a dihedral angle of 25.27 (9) Å with the coumarin mean plane. The nitro group is almost planar to the phenyl ring to which it is attached with a dihedral angle of 4.3 (3)°. The ethoxy group is inclined to the coumarin ring plane by 4.1 (2)°.

Electron delocalization was found at the short C3—C4 bond with a bond length of 1.354 (2) Å. Here as well as in other coumarin compounds reported earlier an important asymmetry in the O—C—O bond angle was detected [O1—C2—O14 = 116.10 (14)° and O14—C2—C3 = 126.14 (15)°]. The bond angles, O1—C10—C9 and C4—C5—C6, at the junction of the two rings in the coumarin moiety are 117.97 (15)° and 123.44 (16)° respectively.

In the crystal, molecules are linked via C—H...O hydrogen bonds forming sheets in the *bc* plane. The sheets are linked via π - π interactions [Cg1...Cg3ⁱ = 3.5688 (13) Å; Cg1...Cg3ⁱⁱ = 3.75114 (13) Å; Cg1 and Cg3 are the centroids of rings O1/C2—C5/C10 and C15—C20; symmetry codes: (i) $x-1/2, -y+3/2, -z+1$; (ii) $x+1/2, -y+3/2, -z+1$] forming a three-dimensional structure (Table 1 and Fig. 2)

S2. Synthesis and crystallization

A mixture of 0.512 g m (3.083 mmol) of 3-ethoxysalicylaldehyde and 0.50 g m (3.083 mmol) of 4-nitro phenylacetonitrile were dissolved in ethanol (25 ml), followed by the addition of 0.525 g m (6.16 mmol) of piperidine and then the reaction mixture was stirred at room temperature for 3 h. The completion of the reaction was monitored by thin layer chromatography [petroleum ether and ethyl acetate (8:2 v/v)]. After completion the reaction mixture was filtered and washed with diethylether giving a yellow precipitation. This product was refluxed with 10% acetic acid for 2 s and then the crude product was filtered and washed with water. It was further purified by recrystallization using acetone as solvent to give yellow crystals of the title compound in good yield (m.p.: 475–477 K; yield: 91%). ¹H NMR(400 MHz, DMSO-d₆): δ = 8.47 (s, 1H, Ar—H), 8.34 (dd, J= 2.08 Hz, 6.94 Hz, 2H, Ar—H), 8.05 (dd, J=2.0 Hz, 6.96 Hz, 2H, Ar—H), 7.34–7.36(m, 3H, Ar—H), 4.22(q, J= 6.92 Hz, 2H, CH₂), 1.43(t, J=6.96 Hz, 3H, CH₃). ¹³C NMR (400 MHz, DMSO-d₆): δ = 45.554, 142.927, 145.554, 142.927, 129.804, 124.807, 123.375, 120.205, 119.868, 115.603, 64.470, 14.63. IR (KBr) ($\nu_{\max}/\text{cm}^{-1}$): 2925 (C—H), 1700 (C=O), 1346 (N—O), 1097 (C—O—C). Mass spectra gave a molecular ion peak at m/z =

311.4[M^r].

S3. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. The hydrogen atoms were fixed geometrically (C—H = 0.93–0.96 Å) and allowed to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C-methyl})$ and = $1.2U_{\text{eq}}(\text{C})$ for other H atoms.

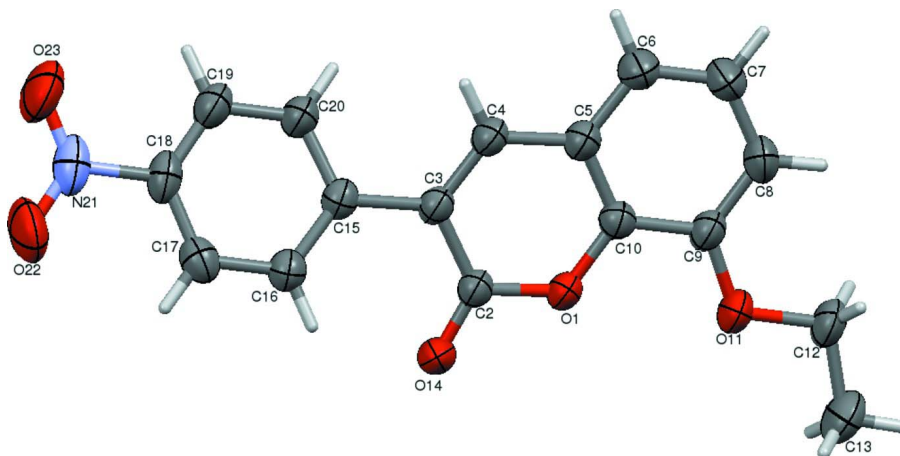


Figure 1

A view of the molecular structure of the title compound, with atom labelling. Displacement ellipsoids are drawn at the 50% probability level.

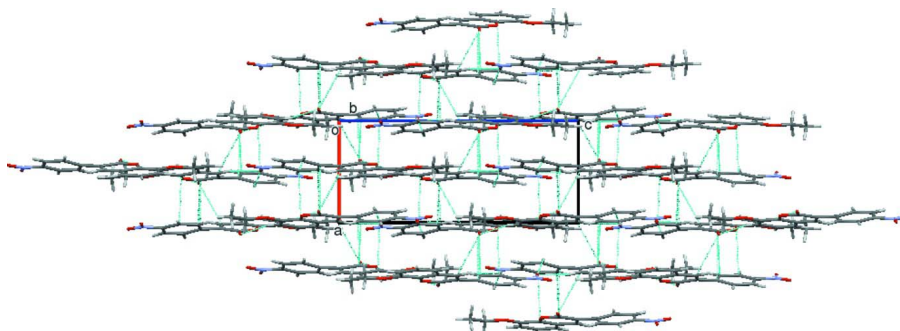


Figure 2

A view along the *b* axis of the crystal packing of the title compound.

8-Ethoxy-3-(4-nitrophenyl)-2H-chromen-2-one

Crystal data

$\text{C}_{17}\text{H}_{13}\text{NO}_5$

$M_r = 311.28$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 6.8118$ (9) Å

$b = 13.6726$ (18) Å

$c = 15.909$ (2) Å

$V = 1481.7$ (3) Å³

$Z = 4$

$F(000) = 648$

$D_x = 1.395$ Mg m⁻³

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

Cell parameters from 2371 reflections

$\theta = 5.6\text{--}64.1^\circ$

$\mu = 0.87$ mm⁻¹

$T = 296$ K

Prism, yellow

$0.29 \times 0.26 \times 0.21$ mm

Data collection

Bruker X8 Proteum
diffractometer

Radiation source: Bruker MicroStar microfocus
rotating anode

Helios multilayer optics monochromator

Detector resolution: 18.4 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2013)

$T_{\min} = 0.786$, $T_{\max} = 0.838$

6729 measured reflections

2371 independent reflections

2225 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.040$

$\theta_{\max} = 64.1^\circ$, $\theta_{\min} = 5.6^\circ$

$h = -7 \rightarrow 6$

$k = -15 \rightarrow 15$

$l = -18 \rightarrow 18$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.131$

$S = 1.03$

2371 reflections

210 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

$w = 1/[\sigma^2(F_o^2) + (0.103P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.003$

$\Delta\rho_{\max} = 0.25 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$

Extinction correction: SHELXL,

$\text{FC}^* = \text{KFC}[1 + 0.001\text{XFC}^2\text{\AA}^3/\text{SIN}(2\Theta)]^{-1/4}$

Extinction coefficient: 0.0100 (19)

Absolute structure: 957 Friedel pairs; Flack
(1983)

Absolute structure parameter: 0.1 (2)

Special details

Experimental. Commercially available chemicals were used directly as received. ¹H NMR was recorded at 400 MHz in Dimethylsulfoxide (DMSO-d₆). ¹³C NMR was recorded at 400 MHz in DMSO-d₆. Mass spectra was recorded on a Jeol SX 102=DA-6000 (10 kV) fast atom bombardment (FAB) mass spectrometer and IR spectra was recorded on a Nicolet 5700 F T—IR instrument as KBr discs.

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement on F^2 for ALL reflections except those flagged by the user for potential systematic errors. Weighted R -factors wR and all goodnesses 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 observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating $-R$ -factor-obs *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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.5699 (2)	0.71184 (8)	0.34374 (7)	0.0409 (4)
O11	0.5487 (2)	0.63066 (10)	0.19274 (7)	0.0541 (5)
O14	0.6120 (2)	0.84884 (8)	0.41311 (7)	0.0515 (5)
O22	0.5085 (3)	0.97085 (14)	0.82793 (11)	0.0869 (7)
O23	0.5615 (4)	0.83031 (16)	0.88337 (9)	0.0860 (7)
N21	0.5394 (3)	0.88318 (16)	0.82255 (11)	0.0613 (7)
C2	0.5811 (3)	0.76190 (12)	0.41836 (10)	0.0384 (5)
C3	0.5552 (3)	0.70612 (11)	0.49641 (10)	0.0371 (5)
C4	0.5162 (3)	0.60920 (12)	0.49082 (10)	0.0404 (5)

C5	0.5066 (3)	0.55829 (13)	0.41241 (11)	0.0418 (5)
C6	0.4682 (4)	0.45725 (14)	0.40574 (12)	0.0552 (7)
C7	0.4586 (4)	0.41549 (14)	0.32849 (12)	0.0600 (7)
C8	0.4848 (4)	0.47095 (14)	0.25537 (12)	0.0555 (7)
C9	0.5232 (3)	0.56989 (14)	0.25975 (11)	0.0440 (6)
C10	0.5335 (3)	0.61330 (13)	0.33952 (11)	0.0385 (5)
C12	0.5377 (4)	0.58696 (15)	0.10978 (10)	0.0560 (7)
C13	0.5582 (5)	0.66749 (19)	0.04725 (12)	0.0751 (9)
C15	0.5625 (3)	0.75681 (13)	0.57930 (11)	0.0378 (5)
C16	0.5182 (3)	0.85604 (13)	0.59012 (11)	0.0431 (5)
C17	0.5132 (3)	0.89769 (14)	0.66921 (12)	0.0489 (6)
C18	0.5499 (3)	0.83947 (15)	0.73796 (11)	0.0484 (6)
C19	0.5985 (4)	0.74317 (15)	0.73029 (11)	0.0528 (7)
C20	0.6066 (3)	0.70218 (14)	0.65155 (10)	0.0486 (6)
H4	0.49480	0.57420	0.54010	0.0480*
H6	0.44980	0.41960	0.45380	0.0660*
H7	0.43410	0.34880	0.32400	0.0720*
H8	0.47620	0.44070	0.20310	0.0670*
H12A	0.41270	0.55400	0.10250	0.0670*
H12B	0.64210	0.53950	0.10250	0.0670*
H13A	0.45240	0.71320	0.05420	0.1120*
H13B	0.55450	0.64080	-0.00850	0.1120*
H13C	0.68110	0.70040	0.05580	0.1120*
H16	0.49170	0.89450	0.54330	0.0520*
H17	0.48550	0.96380	0.67580	0.0590*
H19	0.62580	0.70580	0.77770	0.0630*
H20	0.64200	0.63680	0.64600	0.0580*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0562 (8)	0.0396 (7)	0.0269 (6)	0.0029 (6)	-0.0021 (6)	0.0005 (5)
O11	0.0836 (11)	0.0500 (8)	0.0286 (6)	0.0015 (7)	-0.0015 (6)	-0.0034 (5)
O14	0.0842 (11)	0.0360 (6)	0.0344 (6)	-0.0016 (6)	-0.0087 (7)	0.0027 (5)
O22	0.1200 (16)	0.0747 (11)	0.0659 (10)	-0.0020 (11)	0.0099 (11)	-0.0370 (9)
O23	0.1136 (16)	0.1095 (13)	0.0350 (8)	0.0045 (13)	-0.0005 (9)	-0.0103 (9)
N21	0.0599 (11)	0.0818 (14)	0.0423 (10)	-0.0075 (10)	0.0042 (9)	-0.0213 (9)
C2	0.0450 (10)	0.0401 (9)	0.0301 (8)	0.0039 (7)	-0.0032 (8)	-0.0012 (7)
C3	0.0418 (10)	0.0411 (9)	0.0285 (8)	0.0045 (8)	-0.0017 (8)	-0.0005 (7)
C4	0.0491 (10)	0.0430 (9)	0.0291 (8)	0.0027 (9)	0.0004 (7)	0.0012 (7)
C5	0.0506 (10)	0.0424 (9)	0.0325 (8)	0.0024 (8)	-0.0013 (8)	-0.0007 (7)
C6	0.0810 (15)	0.0430 (9)	0.0417 (10)	-0.0023 (10)	0.0016 (11)	0.0015 (8)
C7	0.0931 (17)	0.0382 (10)	0.0487 (10)	-0.0053 (10)	-0.0022 (12)	-0.0052 (8)
C8	0.0799 (15)	0.0470 (10)	0.0397 (9)	0.0034 (10)	-0.0059 (10)	-0.0128 (8)
C9	0.0552 (12)	0.0485 (10)	0.0282 (8)	0.0061 (9)	-0.0020 (8)	-0.0017 (7)
C10	0.0433 (9)	0.0383 (9)	0.0339 (8)	0.0044 (8)	-0.0014 (8)	-0.0014 (7)
C12	0.0765 (15)	0.0636 (12)	0.0279 (9)	0.0073 (11)	-0.0024 (9)	-0.0081 (8)
C13	0.107 (2)	0.0833 (15)	0.0351 (11)	-0.0015 (16)	0.0019 (13)	-0.0004 (10)

C15	0.0382 (9)	0.0433 (9)	0.0318 (8)	0.0011 (7)	-0.0010 (7)	-0.0026 (7)
C16	0.0500 (10)	0.0432 (9)	0.0360 (9)	-0.0011 (8)	-0.0014 (8)	-0.0033 (7)
C17	0.0528 (11)	0.0435 (10)	0.0504 (11)	-0.0023 (9)	0.0037 (10)	-0.0108 (8)
C18	0.0473 (11)	0.0645 (12)	0.0334 (9)	-0.0062 (10)	0.0010 (8)	-0.0138 (8)
C19	0.0649 (13)	0.0621 (12)	0.0314 (9)	0.0026 (10)	-0.0051 (9)	-0.0008 (8)
C20	0.0651 (13)	0.0475 (10)	0.0333 (9)	0.0055 (10)	-0.0048 (9)	-0.0014 (8)

Geometric parameters (Å, °)

O1—C2	1.372 (2)	C15—C16	1.401 (3)
O1—C10	1.372 (2)	C15—C20	1.403 (2)
O11—C9	1.363 (2)	C16—C17	1.382 (3)
O11—C12	1.451 (2)	C17—C18	1.376 (3)
O14—C2	1.210 (2)	C18—C19	1.363 (3)
O22—N21	1.220 (3)	C19—C20	1.373 (2)
O23—N21	1.217 (3)	C4—H4	0.9300
N21—C18	1.474 (3)	C6—H6	0.9300
C2—C3	1.468 (2)	C7—H7	0.9300
C3—C4	1.354 (2)	C8—H8	0.9300
C3—C15	1.491 (2)	C12—H12A	0.9700
C4—C5	1.430 (2)	C12—H12B	0.9700
C5—C6	1.410 (3)	C13—H13A	0.9600
C5—C10	1.394 (3)	C13—H13B	0.9600
C6—C7	1.357 (3)	C13—H13C	0.9600
C7—C8	1.400 (3)	C16—H16	0.9300
C8—C9	1.380 (3)	C17—H17	0.9300
C9—C10	1.403 (3)	C19—H19	0.9300
C12—C13	1.490 (3)	C20—H20	0.9300
C2—O1—C10	122.84 (13)	N21—C18—C19	119.02 (17)
C9—O11—C12	117.00 (15)	C17—C18—C19	122.13 (17)
O22—N21—O23	123.3 (2)	C18—C19—C20	119.05 (17)
O22—N21—C18	118.08 (18)	C15—C20—C19	121.42 (18)
O23—N21—C18	118.6 (2)	C3—C4—H4	119.00
O1—C2—O14	116.10 (14)	C5—C4—H4	119.00
O1—C2—C3	117.77 (14)	C5—C6—H6	120.00
O14—C2—C3	126.14 (15)	C7—C6—H6	120.00
C2—C3—C4	118.45 (14)	C6—C7—H7	119.00
C2—C3—C15	120.19 (14)	C8—C7—H7	119.00
C4—C3—C15	121.30 (15)	C7—C8—H8	120.00
C3—C4—C5	122.85 (15)	C9—C8—H8	120.00
C4—C5—C6	123.44 (16)	O11—C12—H12A	110.00
C4—C5—C10	117.19 (16)	O11—C12—H12B	110.00
C6—C5—C10	119.36 (16)	C13—C12—H12A	110.00
C5—C6—C7	119.30 (18)	C13—C12—H12B	110.00
C6—C7—C8	121.24 (18)	H12A—C12—H12B	109.00
C7—C8—C9	120.88 (17)	C12—C13—H13A	109.00
O11—C9—C8	125.63 (16)	C12—C13—H13B	109.00

O11—C9—C10	116.32 (16)	C12—C13—H13C	109.00
C8—C9—C10	118.04 (17)	H13A—C13—H13B	110.00
O1—C10—C5	120.86 (15)	H13A—C13—H13C	109.00
O1—C10—C9	117.97 (15)	H13B—C13—H13C	109.00
C5—C10—C9	121.17 (17)	C15—C16—H16	119.00
O11—C12—C13	107.35 (16)	C17—C16—H16	119.00
C3—C15—C16	123.50 (16)	C16—C17—H17	121.00
C3—C15—C20	118.97 (15)	C18—C17—H17	121.00
C16—C15—C20	117.46 (16)	C18—C19—H19	120.00
C15—C16—C17	121.10 (17)	C20—C19—H19	121.00
C16—C17—C18	118.76 (18)	C15—C20—H20	119.00
N21—C18—C17	118.84 (18)	C19—C20—H20	119.00
C10—O1—C2—O14	-179.25 (17)	C6—C5—C10—C9	-0.1 (3)
C10—O1—C2—C3	0.8 (3)	C4—C5—C10—C9	-178.89 (19)
C2—O1—C10—C5	-0.3 (3)	C4—C5—C6—C7	178.9 (2)
C2—O1—C10—C9	179.35 (18)	C10—C5—C6—C7	0.1 (4)
C12—O11—C9—C8	-1.1 (3)	C4—C5—C10—O1	0.8 (3)
C12—O11—C9—C10	-179.77 (19)	C5—C6—C7—C8	-0.4 (4)
C9—O11—C12—C13	176.8 (2)	C6—C7—C8—C9	0.6 (4)
O23—N21—C18—C17	176.1 (2)	C7—C8—C9—C10	-0.5 (4)
O22—N21—C18—C19	175.2 (2)	C7—C8—C9—O11	-179.2 (2)
O22—N21—C18—C17	-3.8 (3)	O11—C9—C10—C5	179.02 (18)
O23—N21—C18—C19	-4.9 (3)	C8—C9—C10—O1	-179.4 (2)
O1—C2—C3—C4	-1.9 (3)	C8—C9—C10—C5	0.3 (3)
O1—C2—C3—C15	-178.98 (17)	O11—C9—C10—O1	-0.7 (3)
O14—C2—C3—C15	1.1 (3)	C3—C15—C16—C17	175.17 (19)
O14—C2—C3—C4	178.2 (2)	C20—C15—C16—C17	-1.7 (3)
C2—C3—C15—C20	-157.80 (19)	C3—C15—C20—C19	-174.2 (2)
C2—C3—C15—C16	25.4 (3)	C16—C15—C20—C19	2.8 (3)
C2—C3—C4—C5	2.5 (3)	C15—C16—C17—C18	-0.9 (3)
C4—C3—C15—C16	-151.6 (2)	C16—C17—C18—N21	-178.37 (19)
C4—C3—C15—C20	25.2 (3)	C16—C17—C18—C19	2.6 (3)
C15—C3—C4—C5	179.53 (19)	N21—C18—C19—C20	179.4 (2)
C3—C4—C5—C6	179.3 (2)	C17—C18—C19—C20	-1.6 (4)
C3—C4—C5—C10	-1.9 (3)	C18—C19—C20—C15	-1.2 (4)
C6—C5—C10—O1	179.6 (2)		

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C4—H4 \cdots O22 ⁱ	0.93	2.53	3.453 (2)	171
C8—H8 \cdots O14 ⁱⁱ	0.93	2.31	3.226 (2)	166
C20—H20 \cdots O22 ⁱ	0.93	2.52	3.275 (3)	138

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